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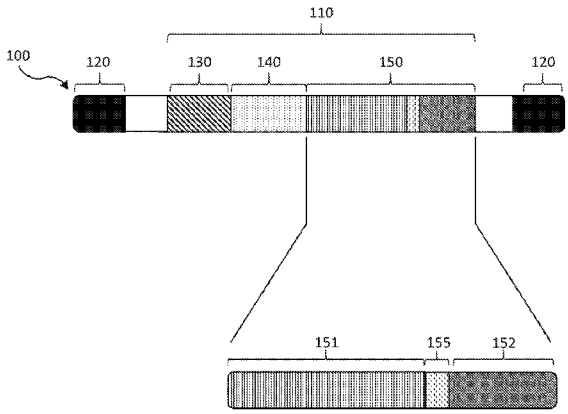
FIG. 2



(57) Abstract: The invention provides compositions and methods for the preparation, manufacture and therapeutic use of viral vectors, such as adeno-associated virus (AAV) particles having viral genomes encoding one or more antibodies or antibody fragments or antibody-like polypeptides, for the prevention and/or treatment of diseases and/or disorders.

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COMPOSITIONS FOR THE TREATMENT OF DISEASE

CROSS REFERENCE TO RELATED APPLICATIONS

[0001] This application claims priority to US Provisional Patent Application No. 62/329,468, filed on April 29, 2016, entitled Compositions for the Treatment of Disease, and US Provisional Patent Application No. 62/329,479, filed on April 29, 2016, entitled Compositions for the Treatment of Disease, the contents of each of which are herein incorporated by reference in their entireties.

REFERENCE TO THE SEQUENCE LISTING

[0002] The present application is being filed along with a Sequence Listing in electronic format. The Sequence Listing file, entitled 20571302PCTSL.txt, was created on April 25, 2017, and is 14,209,554 bytes in size. The information in electronic format of the Sequence Listing is incorporated herein by reference in its entirety.

FIELD OF THE **INVENTION**

[0003] The invention relates to compositions and methods for vectored antibody delivery (VAD).

BACKGROUND **OF** THE INVENTION

[0004] Antibody-based therapies have been developed for a wide variety of diseases, disorders and conditions, including infectious and non-infectious diseases. The U.S. Food and Drug Administration (FDA) has approved antibodies for treatment of cancers, autoimmune and immune system disorders, ocular diseases, nervous system diseases, inflammations, and infections, amongst many others. Naturally, antibodies are components of the adaptive immune response and they function by recognizing specific foreign antigens and stimulating humoral immunity responses. As a consequence, antibodies may be applied to the treatment, prevention, management, diagnosis and research of diseases, disorders and/or conditions.

[0005] Antibodies have relatively short half-lives and this presents an ongoing and long-felt challenge for antibody-based therapies. In order to achieve a sufficiently high concentration of an antibody for long lasting therapeutic effects, antibody therapies are traditionally delivered by repeated administration, e.g. by multiple injections. This dosing regimen results in an inconsistent level of antibody throughout the treatment period, limited efficiency per administration, high cost of administration and consumption of the antibody. Hence, there remains a need in the art for delivery of antibodies and antibody-based therapeutics through alternative routes or modalities of administration.

[0006] One such alternative route of administration is by expression vector (e.g. plasmid or viral vector), including but not limited to, adeno-associated viral vectors (AAVs). Adeno-associated viral vectors are widely used in gene therapy approaches due to a number of advantageous features. As dependoparvoviruses, AAV are non-replicating in infected cells and therefore not associated with any known disease. Further, AAVs may be introduced to a wide variety of host cells, do not integrate into the genome of the host cell, and are capable of infecting both quiescent and dividing cells. AAVs transduce non-replicating and long-lived cells *in vivo*, resulting in long term expression of the protein of interest. Further, AAVs can be manipulated with cellular and molecular biology techniques to produce non-toxic particles carrying a payload encoded in the AAV viral genome that can be delivered to a target tissue or set of ceils with limited or no side-effects. Given the foregoing, the use of AAVs for vectored antibody delivery (VAD) would allow for longer lasting efficacy, fewer dose treatments, and more consistent levels of the antibody throughout the treatment period.

[0007] In vectored antibody deliver}' (VAD) an AAV is used as the delivery modality for a nucleic acid sequence encoding the antibody, which results in *in vivo* expression of the encoded payload, e.g., functional antibody.

[0008] The mechanism underlying VAD is thought to proceed through the following steps. First the AAV vector enters the ceil via endocytosis, then escapes from the endosomal compartment and is transported to the nucleus wherein the viral genome is released and converted into a double-stranded episomal molecule of DNA by the host. The transcriptionally active episome results in the expression of encoded antibodies that may then be secreted from the cell into the circulation. VAD may therefore enable continuous, sustained and long-term delivery of antibodies administered by a single injection of an AAV particle.

[0009] Previous studies of an AAV-mediated antibody technique known as vectored immunoprophylaxis (VIP) have focused on neutralization of human immunodeficiency virus (HIV) (see, e.g. Johnson et al, 2009, Nature Med., 15, 901 - 906, Saunders et al., 2015, J. Virol., 89(16), 8334-8345, Balasz et al., 2012, Nature 481, 81-84, the contents of which are incorporated herein by reference in their entirety). Balasz *et al.* reported a long-term, even lifelong, expression of monoclonal antibody at high concentration from a single intramuscular administration in mice that resulted in full protection against HIV infection. AAV-mediated VIP has also been demonstrated against influenza strains (see, e.g. Balasz, et al. Nat. Biotechnol., 2013, 31(7):647-52) and *Plasmodium Falciparum*, a sporozoite causing malaria infection (see, e.g. Deal at al., 2014, PNAS, 111 (34), 12528-12532), as well as cancer, RSV and drug addiction (see, e.g. review- by Schnepp and Johnson, Microbiol. Spectrum 2(4), 2014). Though promising,

these studies emphasize efforts to merely prevent disease. There still remains a need for improved methods of prevention, and new antibody-mediated therapies for research, diagnosis, and treatment of disease.

[00010] The present invention addresses this need by providing novel AAV particles having viral genomes engineered to encode antibodies and antibody-based compositions and methods of using these constructs (e.g., VAD) for the treatment, prevention, diagnosis and research of diseases, disorders and/or conditions. The present invention further embraces optimized AAV particles for delivery' of nucleic acids (e.g., viral genomes) encoding antibodies and antibody-based compositions to a subject in need thereof.

SUMMARY OF THE INVENTION

[0010] The invention provides AAV particles comprising a capsid and a viral genome, said viral genome comprising at least one inverted terminal repeat (ITR) region and a payload region, said payload region comprising a regulatory sequence operably linked to at least a first nucleic acid segment, said first nucleic acid segment encoding one or more polypeptides given in Tables 3-42, variants and fragments thereof. The capsid of the AAV particle may be any of the serotypes described herein and/or described in Table 1.

[0011] In one aspect the first nucleic acid segment may encode one or more polypeptides such as, but not limited to, an antibody heavy chain, an antibody light chain, a linker, and combinations thereof. The first nucleic acid segment may encode one or more polypeptides which is humanized. As a non-limiting example, the first nucleic acid segment encodes from 5' to 3', an antibody heavy chain, a linker, and an antibody light chain. As another non-limiting example, the first nucleic acid segment encodes from 5' to 3', an antibody light chain, a linker, and an antibody heavy chain. As yet another non-limiting example, the first nucleic acid segment encodes one or more antibody heavy chains. As yet another non-limiting example, the first nucleic acid segment encodes one or more antibody light chains.

[0012] In one aspect, the first nucleic acid segment encodes an antibody, having at least 95% identity to any of the sequences of Tables 3-42 (SEQ ID NO: 2948-9220).

[0013] In one aspect, the regulatory sequence may comprise a promoter such as but not limited to, human elongation factor la-subunit (EF1α), cytomegalovirus (CMV) immediate-early enhancer and/or promoter, chicken β-actin (CBA) and its derivative CAG, β glucuronidase (GUSB), or ubiquitin C (UBC). Tissue-specific expression elements can be used to restrict expression to certain cell types such as, but not limited to, muscle specific promoters, B cell promoters, monocyte promoters, leukocyte promoters, macrophage promoters, pancreatic acinar cell promoters, endothelial cell promoters, lung tissue promoters, astrocyte promoters, or nervous

system promoters which can be used to restrict expression to neurons, astrocytes, or oligodendrocytes.

[0014] In one aspect, the linker in the viral genome is selected from one or more of the linkers given in Table 2.

[0015] In one aspect, the AAV particles described herein may comprise a viral genome which is single stranded.

[0016] In one aspect, the AAV particles described herein may comprise a viral genome which is self-complementary.

[0017] In one aspect, the AAV particles described herein may comprise a viral genome comprising at least one intron sequence.

[0018] In one aspect, the AAV particles described herein may comprise a viral genome comprising at least one staffer sequence to adjust the length of the viral genome to increase efficacy and/or effciency.

[0019] In one aspect, the AAV particles described herein may comprise at least one region which has been codon optimized. As a non-limiting example, the viral genome may be codon optimized. As another non-limiting example, the first nucleic acid segment is codon-optimized.

[0020] In one aspect, the AAV particles described herein may comprise a viral genome with 2 ITR regions. At least one of the ITR regions may be derived from the same or different parental serotype of the capsid. As anon-limiting example, at least one ITR region is derived from AAV2.

[0021] In one aspect, the AAV particles comprise a viral genome which comprises a second nucleic acid segment. The second nucleic acid segment may encode an aptamer, siRNA, saRNA, ribozyme, microRNA, raRNA or combination thereof.

[0022] In one aspect, the AAV particles comprise a viral genome which comprises a second nucleic acid segment encoding an siRNA designed to target the mRNA that encodes the target of the antibody encoded by the first nucleic acid segment.

[0023] In one aspect, the AAV particles comprise a viral genome which comprises a second nucleic acid segment encoding a microRNA, the microRNA is selected to target the mRNA that encodes the target of the antibody encoded by the first nucleic acid segment.

[0024] In one aspect, the AAV particles comprise a viral genome which comprises a second nucleic acid segment encoding an mRNA, the mRNA encodes one or more peptides inhibitors of the same target of the antibody encoded by the first nucleic acid segment.

[0025] In one aspect, the AAV panicles comprise a viral genome which comprises a third nucleic acid segment. The third nucleic acid segment may encode a nuclear export signal, a

polynucleotide or polypeptide which acts as a regulator of expression of the viral genome in which it is encoded, a polynucleotide or polypeptide which acts as a regulator of expression of the payload region of the viral genome in which it is encoded and/or a polynucleotide or polypeptide which acts as a regulator of expression of the first nucleic acid segment of the payload region of the viral genome in which it is encoded.

[0026] The invention provides AAV particles comprising a capsid and a viral genome, said viral genome comprising at least one inverted terminal repeat (ITR) region and a payload region comprising a regulatory sequence operabiy linked to at least a first nucleic acid segment, the first nucleic acid segment encoding a bispecific antibody derived from any of the sequences listed in Tables 3-42 or portions or fragments thereof.

[0027] The invention provides methods of producing a functional antibody in a subject in need thereof, comprising adininistering to a subject the AAV particles described herein. The level or amount of the functional antibody in the target cell or tissue after administration to the subject may be from about .001 ug/mL to 100 mg/mL. The functional antibody may be encoded by a single first nucleic acid segment of a viral genome within the AAV particle. The functional antibody may be encoded by two different viral genomes, the two different viral genomes may be packaged in separate capsids.

[0028] The invention provides a pharmaceutical composition comprising an AAV particle described herein in a pharmaceutically acceptable excipient. As a non-limiting example, the pharmaceutically acceptable excipient is saline. As a non-limiting example, the pharmaceutically acceptable excipient is 0.001% pliironic in saline.

[0029] The invention provides methods of producing a functional antibody in a subject in need thereof, comprising administering to a subject the AAV particles described herein by a delivery route such as, but not limited to, enteral (into the intestine), gastroenteral, epidural (into the dura mater), oral (by way of the mouth), transdermal, intracerebral (into the cerebrum), mtracerebroventricular (into the cerebral ventricles), epicutaneous (application onto the skin), intradermal, (into the skin itself), subcutaneous (under the skin), nasal administration (through the nose), intravenous (into a vein), intravenous bolus, intravenous drip, intra-arterial (into an artery), intramuscular (into a muscle), intracardiac (into the heart), intraosseous infusion (into the bone marrow), intrathecal (into the spinal canal), intraparenchymal (into brain tissue), intraperitoneal, (infusion or injection into the peritoneum), intravesical infusion, intravitreaL (through the eye), intracavemous injection (into a pathologic cavity) intracavitary (into the base of the penis), intravaginal administration, intrauterine, extra-amniotic administration, transdermal (diffusion through the intact skin for systemic distribution), transmucosal (diffusion

through a mucous membrane), transvaginal, insufflation (snorting), sublingual, sublabial, enema, eye drops (onto the conjunctiva), or in ear drops, auricular (in or by way of the ear), buccal (directed toward the cheek), conjunctival, cutaneous, dental (to a tooth or teeth), electro-osmosis, endoeervicai, endosinusial, endotracheal, extracorporeal, hemodialysis, infiltration, interstitial, intra-abdominal, inira-amniotic, intra-artieular, mtrabiliary, mtrabronchiai, mtrabursai, intracartiiaginous (within a cartilage), intracaudal (within the cauda equine), intracisternal (withm the cisterna magna cerebeliomedularis), mtracorneal (within the cornea), dental itttracoronal, intracoronary (within the coronaty atleries), intracorporus caveroosum (within the dilatable spaces of the corporus cavernosa of the penis), intradiscal (within a disc), intraductal (within a duct of a gland), intraduodenal (within the duodenum), intradural (within or beneath the dura), intraepidernial (to the epidermis), intraesophageal (to the esophagus), intragastric (within the stomach), intragingival (within the gingivae), intraileal (withm the distal portion of the small intestine), intralesionai (within or introduced directly to a localized lesion), intraluminal (withm a lumen of a tube), intralyniphatic (withm the lymph), intramedullary (within the marrow cavity of a bone), mtrameningeai (withm the meninges), intramyocardial (within the myocardium), intraocular (within the eye), intraovarian (within the ovary), intrapericardial (within the peri cardium), intrapleural (within the pleura), intraprostatic (within the prostate gland), intrapulmonary (within the lungs or its bronchi), mtrasmal (within the nasal or periorbital sinuses), intraspinal (within the vertebral column), intrasynoviai (withm the synovial cavity of a joint), intratendinous (within a tendon), intratesticuiar (within the testicle), intrathecal (within the cerebrospinal fluid at any level of the cerebrospinal axis), intrathoracic (within the thorax), intratubular (within the tubules of an organ), intratumor (within a tumor), intratympanic (within the aurus media), intravascular (within a vessel or vessels), intraventricular (within a ventricle), iontophoresis (by means of electric current where ions of soluble salts migrate into the tissues of the body), irrigation (to bathe or flush open wounds or body cavities), laryngeal (directly upon the larynx), nasogastric (through the nose and into the stomach), occlusive dressing technique (topical route administration which is then covered by a dressing which occludes the area), ophthalmic (to the external eye), oropharyngeal (directly to the mouth and pharynx), parenteral, percutaneous, periarticular, peridural, perineural, periodontal, rectal, respiratory (within the respiratory tract by inhaling orally or nasally for local or systemic effect), retrobulbar (behind the pons or behind the eyeball), soft tissue, subarachnoid, subconjunctival, submucosal, topical, transplacental (through or across the placenta), transtracheal (through the wall of the trachea), transtympanic (across or through the tympanic cavity), ureteral (to the

ureter), urethral (to the urethra), vaginal caudal block, diagnostic, nerve block, biliary perfusion, cardiac perfusion, photopheresis and spinal

[0030] The invention provides methods of treating and/or preventing a disease or disorder in a subject comprising administering to the subject an AAV particle described herein. The administration may be at a prophyiacticaliy effective dose such as, but not limited to, from about 1 ug/mL to about 500 ug/mL of expressed polypeptide or 1x10e4 to 1x10eI6 VG/mL from the pharmaceutical composition. The pharmaceutical composition may be administered at least once. The pharmaceutical composition may be administered daily, weekly, monthly or yearly. The pharmaceutical composition may be co-administered as part of a combination therapy.

[0031] The invention provides methods of producing an antibody in a subject by administering the AAV particles described herein, where the antibody is not a virus neutralizing antibody.

[0032] The invention provides methods of producing an antibody in a subject by administering the AAV particles described herein, where the antibody is not an HIV or HCV virus neutralizing antibody.

BRIEF DESCRIPTION OF THE DRAWINGS

[0033] The foregoing and other objects, features and advantages will be apparent from the following description of particular embodiments of the invention, as illustrated in the accompanying drawings. The drawings are not necessarily to scale, emphasis instead being placed upon illustrating the principles of various embodiments of the invention.

[0034] FIG. 1 is a schematic of vectored antibody delivery.

[0035] FIG. 2 is a schematic of a viral genome of the invention.

[0036] FIG. 3 is a schematic of payload regions. FIG. 3 discloses SEQ ID NO: 922.

DETAILED DESCRIPTION OF THE INVENTION

1. COMPOSITIONS OF THE INVENTION

[0037] According to the present invention, compositions for delivering functional antibodies and/or antibody-based compositions by adeno-associated viruses (AAVs) are provided. AAV particles of the invention may be provided via any of several routes of administration, to a cell, tissue, organ, or organism, *in vivo, ex vivo* or *in vitro.*

[0038] As used herein, an "AAV particle" is a virus which comprises a viral genome with at least one payioad region and at least one inverted terminal repeat (ITR) region.

[0039] As used herein, "viral genome" or "vector genome" refers to the nucleic acid sequence(s) encapsulated in an AAV particle. Viral genomes comprise at least one payioad

region encoding polypeptides of the invention, e.g., antibodies, antibody-based compositions or fragments thereof.

[0040] As used herein, a "payload" or "payload region" is any nucleic acid molecule which encodes one or more polypeptides of the invention. At a minimum, a payload region comprises nucleic acid sequences that encode an antibody, an antibody-based composition, or a fragment thereof, but may also optionally comprise one or more functional or regulatory elements to facilitate transcriptional expression and/or polypeptide translation.

[0041] The nucleic acid sequences and polypeptides disclosed herein may be engineered to contain modular elements and/or sequence motifs assembled to enable expression of the antibodies or antibody-based compositions of the invention. In some embodiments, the nucleic acid sequence comprising the payload region may comprise one or more of a promoter region, an intron, a Kozak sequence, an enhancer or a polyadenylation sequence. Payload regions of the invention typically encode antibodies or antibody based compositions, which may include an antibody heavy chain domain, an antibody light chain domain, both antibody heavy and light chain domains, or fragments of the foregoing in combination with each other or in combination with other polypeptide moieties. In some cases, payload regions may also encode one or more linkers or joining regions between antibody heavy and light chain domains or fragments. The order of expression, structural position, or concatemer count (heavy chain, light chain, or linker) may be different within or among different payload regions. The identity, position and number of linkers expressed by payload regions may also vary.

[0042] The payload regions of the invention may be delivered to one or more target cells, tissues, organs or organisms within the viral genome of an AAV particle.

Adeno-associated viruses (AAVs) and AAV particles

[0043] Viruses of the Parvoviridae family are small non-enveloped icosahedral capsid viruses characterized by a single stranded DNA genome. Parvoviridae family viruses consist of two subfamilies: Parvovirinae, which infect vertebrates, and Densovirinae, which infect invertebrates. Due to its relatively simple structure, easily manipulated using standard molecular biology techniques, this virus family is useful as a biological tool. The genome of the virus may be modified to contain a minimum of components for the assembly of afunctional recombinant virus, or viral particle, which is loaded with or engineered to express or deliver a desired payload, which may be delivered to a target cell, tissue, organ, or organism.

[0044] The parvoviruses aid other members of the Parvoviridae family are generally described in Kenneth I. Berns, "Parvoviridae: The Viruses and Their Replication," Chapter 69 in

FIELDS VIROLOGY (3d Ed. 1996), the contents of which are incorporated by reference in their entirety.

[0045] The Parvoviridae family comprises the Dependovirus genus which includes adeno-associated viruses (AAV) capable of replication in vertebrate hosts including, but not limited to, human, primate, bovine, canine, equine, and ovine species.

[0046] The AAV vector genome is a linear, single-stranded DNA (ssDNA) molecule approximately 5,000 nucleotides (nt) in length. The AAV viral genome can comprise a payload region and at least one inverted terminal repeat (ITR) or ITR region. ITRs traditionally flank the coding nucleotide sequences for the **non**-structural proteins (encoded by Rep genes) and the structural proteins (encoded by capsid genes or Cap genes). While not wishing to be bound by theory, an AAV viral genome typically comprises two ITR sequences. The AAV vector genome comprises a characteristic T-shaped hairpin structure defined by the self-complementary terminal 145 nt of the 5' and 3" ends of the ssDNA which form an energetically stable double stranded region. The double stranded hairpin structures comprise multiple functions including, but not limited to, acting as an origin for DNA replication by functioning as primers for the endogenous DNA polymerase complex of the host viral replication cell.

[0047] In addition to the encoded heterologous payload, AAV vectors may comprise the viral genome, in whole or in part, of any naturally occurring and/or recombinant AAV serotype nucleotide sequence or variant. AAV variants may have sequences of significant homology at the nucleic acid (genome or capsid) and amino acid levels (capsids), to produce constructs which are generally physical and functional equivalents, replicate by similar mechanisms, and assemble by similar mechanisms. Chiorini et al., J. Vir. 71: 6823-33(1997); Srivastava et al., 3. Vir. 45:555-64 (1983), Chionni et al., J. Vir. 73:1309-1319 (1999): Rutledge et al, J. Vir. 72:309-319 (1998); **and** Wu et al, J. Vir. 74: 8635-47 (2000), the contents of each of which are incorporated herein by reference in their entirety.

[0048] In one embodiment, AAV particles of the present invention are recombinant AAV viral vectors which are replication defective, lacking sequences encoding functional Rep and Cap proteins within their viral genome. These defective AAV vectors may lack most or all parental coding sequences and essentially cany only one or two AAV ITR sequences and the nucleic acid of interest for delivery to a cell, a tissue, an organ or an organism.

[0049] In one embodiment, the viral genome of the AAV particles of the present invention comprise at least one control element which provides for the replication, transcription and translation of a coding sequence encoded therein. Not all of the control elements need always be present as long as the coding sequence is capable of being replicated, transcribed and/or

translated in an appropriate host cell. Non-limiting examples of expression control elements include sequences for transcription initiation and/or termination, promoter and/or enhancer sequences, efficient RNA processing signals such as splicing and polyadenylation signals, sequences that stabilize cytoplasmic mRNA, sequences that enhance translation efficacy (e.g., Kozak consensus sequence), sequences that enhance protein stability, and/or sequences that enhance protein processing and/or secretion.

[0050] According to the present invention, AAV particles for use in therapeutics and/or diagnostics comprise a virus that has been distilled or reduced to the minimum components necessary for transduction of a nucleic acid payload or cargo of interest. In this manner, AAV particles are engineered as vehicles for specific delivery while lacking the deleterious replication and/or integration features found in wild-type viruses.

[0051] AAV vectors of the present invention may be produced recombinantly and may be based on adeno-associated virus (AAV) parent or reference sequences. As used herein, a "vector" is any molecule or moiety which transports, transduces or otherwise acts as a carrier of a heterologous molecule such as the nucleic acids described herein.

[0052] In addition to single stranded AAV viral genomes (e.g.. ssAAVs), the present invention also provides for self-complementary AAV (scAAVs) viral genomes. scAAV vector genomes contain DNA strands which anneal together to form double stranded DNA. By skipping second strand synthesis, scAAVs allow for rapid expression in the cell.

[0053] In one embodiment, the AAV particle of the present invention is an scAAV.

[0054] In one embodiment, the AAV particle of the present invention is an ssAAV.

[0055] Methods for producing and/or modifying AAV particles are disclosed in the art such as pseudotyped AAV vectors (PCX Patent Publication Nos. WO200028004; WO2001 23001; **WO20041 12727;** WO 2005005610 and WO 2005072364, the content of each of which is incorporated herein by reference in its entirety).

[0056] AAV particles may be modified to enhance the efficiency of delivery. Such modified AAV particles can be packaged efficiently and be used to successfully infect the target cells at high frequency and with minimal toxicity. In some embodiments, the capsids of the AAV particles are engineered according to the methods described in US Publication Number US 20130195801, the contents of which are incorporated herein by reference in their entirety.

[0057] In one embodiment, the AAV particles comprising a payload region encoding the polypeptides of the invention may be introduced into mammalian ceils.

AAV serotypes

[0058] AAV particles of the present invention may comprise or be derived from any natural or recombinant AAV serotype. According to the present invention, the AAV particles may utilize or be based on a serotype selected from any of the following AAV1, AAV2, AAV2G9, **AAV3,** AAV3a, AAV3b, AAV3-3, AAV4, AAV4-4, AAV5, AAV6, AAV6.1, AAV6.2, AAV6.1.2, AAV7, AAV7.2, AAV8, **AAV9, AAV9.11,** AAV9.13, AAV9.16, AAV9.24, AAV9.45, AAV9.47, AAV9.61, AAV9.68, AAV9.84, AAV9.9, AAV10, AAVll, AAV12, AAV16.3, AAV24.1, AAV2.7.3, AAV42.12, **AAV42-lb,** AAV42-2, AAV42-3a, **AAV42-3b,** AAV42-4, AAV42-5a, **AAV42-5b, AAV42-6b,** AAV42-8, AAV42-10, AAV42-11, AAV42-1~, AAV42-13, AAV42-15, AAV42-aa, **AAV43-1,** AAV43-12, AAV43-20, AAV43-21, AAV43-23, AAV43-25, AAV43-5, AAV44.1, AAV44.2, AAV44.5, AAV223.1, AAV223.2, AAV223.4, AAV223.5, AAV223.6, AAV223.7, AAVl-7/rh.48, AAY i-S/rh.49, AAV2-15/rh.62, AAV2-3/rh.61, AAV2-4/rh.50, AAV2-5/rh.51, AAV3.1/hu.6, AAV3.l/hu.9, AAV3-9/rh.52, AAV3-1l/rh.53, AAV4-8/rl l.64, AAV4-9/rh.54, **AAV4-19/A.55,** AAV5-3/rh.57, AAV5-22/srh.58, AAV7.3/11U.7, AAV16.8/hu.10, AAV16.12/hu.ll, AAV29.3/bb.L, AAV29.5/bb.2, AAV106.1/hus.37, AAVl 14.3/hu.40, AAV127.2/hu.41, AAV127.5/hu.42, AAV128.3/hu.44, AAV130.4/hu.48, AAV145.1/hu.53, AAV145.5/lm.54, AAV145.6/hu.55, AAV161.10/hu.60, AAVl6l.6/1m6L, AAV33.12/hu.l7, AAV33.4/hu.l 5, AAV33.8/hiil6, AAV52/hu.l9, AAV52.1/hu.20, AAV58.2/hu.25, AAVA3.3, AAVA3.4, AAVA3.5, AAVA3.7, AAVC1, AAVC2, AAVC5, AAV-DJ, AAV-DJ8, AAVF3, AAVF5, AAVH2, AAVrh.72, AAVhu.8, AAVrh.68, AAVrh.70, AAVpU. AAVpi.3, AAVpi.2, AAVrh.60, AAVrh.44, AAVrh.65, AAVrh.55, AAVrh.47, AAVrh.69, AAVrh.45, AAVrh.59, AAVhu.12, AAVH6, AAVLK03, AAVH-l/hu.1, AAVH-5/hu.3, AAVLG-10/rh.40, AAVLG-4/rh.38, AAVLG-9/hu.39, AAVN721-8/rh.43, AAVCh.5, AAVCh.5Rl, AAVcy.2. AAVcy.3, AAVcy.4, AAVcy.5. AAVCy.SRL AAVCy.5R2, AAVCy.5R3, AAVCy.5R4, AAVcy.6, AAVn-i, AAVhu.2, AAVhu.3, AAVhu.4, AAVhuA, AAVhuA, AAVhuA, AAVhu.9, AAVhu.lO, AAVhu.ll, AAVhu.13, **AAVhu.15,** AAVhu.16, AAVhu.17, AAVhu.18. **AAVhu.20,** AAVhu.21, AAVhu.22, AAVhu.23.2, AAVhu.24, AAVhu.25, AAVhu.27, AAVhu.28, AAVhu.29, AAVhu.29R, AAVhu.31, AAVhu.32, AAVhu.34, AAVhu.35, AAVhu.37, AAVhu.39, **AAVhu.40,** AAVhus.41, AAVhu.42, AAVhu.43, AAVhu.44, AAVhu.44Rl, AAVhu.44R2, AAVhu.44R3, AAVhu.45, AAVhu.46, AAVhu.47, AAVhu.48, AAVhu.48RI, AAVhu.48R2, AAVhu.48R3, AAVhu.49, AAVhu.51, AAVhu.52, AAVhu.54, AAVhu.55, AAVhu.56, AAVhu.57, AAVhu.58, AAVhu.60, AAVhu.61, AAVhu.63, AAVhu.64, AAVhu.66, AAVhu.67, AAVhu.14/9, AAVhu.t 19, AAVrh.2, AAVrh.2R, AAVrh.8, AAVrh.8R, AAVrh.10, AAVrh.12, AAVrh.13, AAVrh.13R, AAVrh.14, AAVrh.17, AAVrh.18, AAVrh.19, A.AVrh.20,

AAVrh.21, AAVrh.22, AAVrh.23, AAVfh.24, AAVfh.25, AAVrh.31, AAVrh.32, AAVrh.33, AAVrh.34, AAVrh.35, AAVrh.36, AAVrh.37, AAVrh.37R2, AAVrh.38, AAVrh.39, AAVrh.40, AAVrh.46, AAVrh.48, AAVrh.48.1, AAVrh.48.1.2, AAVrh.48.2, AAVrh.49, AAVrh.51, AAVrh.52, AAVrh.53, AAVrh.54, AAVrh.56, AAVrh.57, AAVrh.58, AAVrh.61, AAVrh.64, AAVrh.64R1, AAVrh.64R2, AAVrh.67, AAVrh.73, AAVrh.74, AAVrhSR, AAVrhSR A586R mutant, AAVrh8R R533A mutant, AAAV, BAAV, caprine AAV, bovine AAV, AAVhE1.1, AAVhErl.5, AAVhER1.14, AAViiErl.8, AAVhErl.16, AAVhErl.18, AAVhErl.35, AAVhErl .7, **AAVhErl.36,** AAVhEr2.29, **AAVhEr2.4,** AAVhEr2.16, AAVhEr2.30, AAVbEr2.31, AAVhEi2.36, AAVhERl.23, AAVhEr3. 1, AAV2.5T , AAV-PAEC, AAV-LKOl, AAV-LK02, AAV-LK03, AAV-LK04, AAV-LK05, AAV-LK06, AAV-LK07, AAV-LK08, AAV-LK09, AAV-LK10, AAV-LK11, AAV-LK12, AAV-LK13, AAV-LK14, AAV-LK15, AAV-LK16, AAV-LK17, AAV-LK18, AAV-LK19, AAV-PAEC2, AAV-PAEC4, AAV-PAEC6, AAV-PAEC7, AAV-PAEC8, **AAV-PAEC11,** AAV-PAEC12, AAV-2-pre-miRNA-101 , AAV-8h, AAV-8b, AAV-h, AAV-b, AAV SM 10-2 , AAV Shuffle 100-1 , AAV Shuffle 100-3, AAV Shuffle 100-7, AAV Shuffle 10-2, AAV Shuffle 10-6, AAV Shuffle 10-8, AAV Shuffle 100-2, AAV SM 10-1, AAV SM 10-8 , AAV SM 100-3, AAV SM 100-10, BNP61 AAV, BNP62 AAV, BNP63 AAV, AAVrh.50, AAVrh.43, AAVrh.62, AAVrh.48, AAVhu.!9, AAVhu.lL , AAVhu.53, AAV4-8/rh.64, AAVLG-9/liu.39, AAV54.5/hu.23, AAV54.2/hu.22, AAV54.7/hu.24, AAV54.1/hu.21, AAV54.4R/hu.27, AAV46.2/hu.28, AAV46.6/hu.29, AAV128.1/hu.43, true type AAV (ttAAV), UPENN AAV 10, Japanese AAV 10 serotypes, AAV CBr-7.1, AAV CBr-7.10, AAV CBr-7.2, AAV CBr-7.3, AAV CBr-7.4, AAV CBr-7.5, AAV **CBr-7.7,** AAV CBr-7.8, AAV CBr-B7.3, AAV CBf-B7.4, AAV CBr-El, AAV CBr-E2, AAV CBr-E3, AAV CBr-E4, AAV CBr-E5, AAV CBr-e5, AAV CBr-E6, AAV CBr-E7, AAV CBr-E8, AAV CHt-1, AAV CHt-2, AAV CHt-3, AAV CHt-6.1, AAV CHt-6.10, AAV CHt-6.5, AAV CHt-6.6, AAV CHt-6.7, AAV CHi-6.8, AAV CHt-Pl, AAV CHt-P2, AAV CHt-P5, AAV CHt-P6, AAV CHt-P8, AAV CHt-P9, AAV CKd-1, AAV CKd-10, AAV Ckc!-2, AAV CKd-3, AAV CKd-4, AAV CKd-6, AAV CKd-7, AAV CKd-8, AAV CKd-B1, AAV CKd-B2, AAV CKd-B3, AAV CKd-B4, AAV CKd-B5, AAV CKd-B6, AAV CKd-B7, AAV CKd-B8, AAV CKd-H1, AAV CKd-H2, AAV CKd-H3, AAV CKd-H4, AAV CKd-H5, AAV CKd-H6, AAV CKd-N3, AAV CKd-N4, AAV CKd-N9, AAV CLg-Fl, AAV CLg-F2, AAV CLg-F3, AAV CLg-F4, AAV CLg-F5, AAV CLg-F6, AAV **CLg-F7,** AAV CLg-F8, AAV CLv-l, AAV CLvl-1, AAV Clvl-10, AAV CLvl-2, AAV CLv-12, AAV **CLvl-3,** AAV CLv-13, AAV CLvl-4, AAV Civ 1-7, AAV Civl-8, AAV Civ 1-9, AAV CLv-2, AAV CLv-3, AAV *CLv-4*, AAV CLv-6, AAV CLv-8, AAV CLv-DL, AAV CLv-D2, AAV CLv-D3, AAV CLv-D4, AAV CLv~D5, AAV

CLv-D6, AAV CLV-D7, AAV CLv-D8. AAV CLv-El, AAV **CLv-Kl,** AAV **CLv-K3,** AAV CLv-K6, AAV CLv-E4, AAV CLv-E5, AAV CLv-L6, AAV CLv-Ml, AAV CLv-Ml 1, AAV CLv-M2, AAV CLv-MS, AAV CLv-M6, AAV CEv~M7, AAV CLv-M8, AAV CLv-M9, AAV CLv-Rl, AAV **CLv-R2,** AAV CLv-R3, AAV CLv-R4, AAV CLv-R5, AAV CLv-R6, AAV CLv-R7, AAV CLv-R8, AAV CLv-R9, AAV CSp-1, AAV **CSp-10,** AAV CSp-1 1, AAV CSp-2, AAV CSp-3, AAV CSp-4, AAV CSp-6, AAV CSp-7, AAV CSp-8, AAV CSp-8.10, AAV CSp-8.2, AAV CSp-8.4, AAV CSp-8.5, AAV CSp-8.6, AAV CSp-8.7, AAV CSp-8.8, AAV CSp-8.9, AAV CSp-9, AAV.hu.48R3, AAV.VR-355, AAV3B. AAV4, AAV5, AAVFl/HSCl. AAVF11/HSCl 1, AAVFI2/HSCI2, AAVF13/HSC13, AAVF14/FISC14, **AAVF15/HSC15,** AAVF16/HSC16, AAVF17/HSC17, AAVF2/HSC2, AAVF3/HSC3, AAVF4/HSC4, AAVF5/HSC5, AAVF6/HSC6, **AAVF7/HSC7,** AAVF8/HSC8, AAVF9/HSC9, PHP.B, PHP.A, G2B-26, G2B-13, TH 1.1-32 and/or TH 1.1-3.5 and variants thereof.

[0059] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Publication **No.** US200301 38772, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV1 (SEQ ID NO: 6 and 64 of US20030138772), AAV2 (SEQ ID **NO:** 7 and 70 of US20030138772), AAV3 (SEQ ID **NO:** 8 and 71 ofUS20030138772), AAV4 (SEQ ID **NO:** 63 of US20030138772) AAV5 (SEQ ID **NO:** 114 of US20030138772), AAV 6 (SEQ ID NO: 65 of US20030138772), AAV7 (SEQ ID NO: 1-3 of US200301 38772), AAV 8 (SEQ ID **NO:** 4 and 95 of US200301 38772), AAV9 (SEQ ID **NO:** 5 and 100 of US20030I38772), AAV 10 (SEQ ID **NO:** 117 of US200301 38772), AAVl 1 (SEQ ID **NO:** 118 of US20030138772), AAV12 (SEQ ID **NO:** 119 of US20030138772), AAVrhlO (amino acids 1 to 738 of SEQ ID **NO:** 81 of US200301 38772), AAV 16.3 (IJS20030138772 SEQ TD **NO:** 10), AAV29.3/bb. i (US20030138772 SEQ ID **NO:** 11), AAV29.4 (US20030138772 SEQ ID NO: 12), AAV29.5/bb.2 (IJS200301 38772 SEQ ID NO: 13), AAVl. 3 (US20030138772 SEQ ID NO: 14), AAV13.3 (US20030138772 SEQ ID NO: 15), AAV24.1 (US20030138772 SEQ ID NO: 16), AAV27.3 (IJS20030138772 SEQ ID NO: 17), AAV7.2 (US200301 38772 SEQ ID NO: 18), AAVC1 (US20030I38772 SEQ ID NO: 19), AAVC3 (US20030I38772 SEQ ID NO: 20), AAVC5 (US200301 38772 SEQ ID NO: 21), AAVFi (US20030138772 SEQ ID NO: 22), AAVF3 (US200301 38772 SEQ ID **NO:** 23), AAVF5 (IJS20030138772 SEQ ID **NO:** 24), AAVH6 (US20030138772 SEQ ID NO: 25), AAVH2 (US20030I38772 SEQ ID NO: 26), AAV42-8 (US200301 38772 SEQ ID NO: 27), AAV42-15 (US200301 38772 SEQ ID NO: 28), AAV42-5b (US200301 38772 SEQ ID **NO:** 29), AAV42-lb (US20030138772 SEQ ID **NO:** 30), AAV42-13 (US20030138772 SEQ ID NO: 31), AAV42-3a (US20030138772 SEQ ID NO: 32), AAV42-4 (US20030138772 SEQ ID NO: 33), AAV42-5a (US20030 138772 SEQ ID NO: 34),

AAV42-10 (US20030138772 SEQ ID NO: 35), **AAV42-3b (US20030138772** SEQ ID NO: 36), AAV42-11 (US20030138772 SEQ ID NO: 37), AAV42-6b **(US20030138772** SEQ ID NO: 38), **AAV43-1** (US20030138772 SEQ ID NO: 39), AAV43-5 (US20030138772 SEQ ID NO: 40), **AAV43-12** (US20030138772 SEQ ID NO: 41), AAV43-20 (US20030138772 SEQ ID NO: 42), AAV43-21 (US20030138772 SEQ ID NO: 43), AAV43-23 (US20030138772 SEQ ID NO: 44). AAV43-25 (US20030138772 SEQ ID NO: 45), AAV44.1 (US20030138772 SEQ ID NO: 46), AAV44.5 (US20030138772 SEQ ID NO: 47), AAV223.1 (US20030138772 SEQ ID NO: 48), AAV223.2 (US20030138772 SEQ ID NO: 49), AAY223.4 (US20030138772 SEQ ID NO: 50), AAV223.5 (IJS20030138772 SEQ ID NO: 51), AAV223.6 (US20030138772 SEQ ID NO: 52), AAV223.7 **(US20G30138772** SEQ ID NO: 53), AAV A3.4 (US2003G138772 SEQ ID NO: 54), AAV A3.5 (US20030138772 SEQ ID NO: 55), AAV A3.7 **(US20030138772** SEQ ID NO: 56), AAV A3.3 (US20030138772 SEQ ID NO: 57), AAV42.12 (US20030138772 SEQ ID NO: 58), AAV44.2 (US20030138772 SEQ ID NO: 59), AAV42-2 (US20030138772 SEQ ID NO: 9), or variants thereof.

[0060] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Publication No. US20150159173, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV2 (SEQ ID NO: 7 and 23 of US20150159173), rh20 (SEQ ID NO: 1 of US20150159173), rh32/33 (SEQ ID NO: 2 of IJS20150159173), rh39 (SEQ ID NO: 3, 20 and 36 of US20150159173), rh46 (SEQ ID NO: 4 and 22 of US20150159173), rh73 (SEQ ID NO: 5 of US20150159173), rh74 (SEQ ID NO: 6 of US20150159173), AAV6.1 (SEQ ID NO: 29 of US20150159173), rh.8 (SEQ ID NO: 41 of US20150159173), rh.48.1 (SEQ ID NO: 44 of US20150159173), hu.44 (SEQ ID NO: 45 of US20150159173). hi!.29 (SEQ ID NO: 42 of US20150159173). hi!.48 (SEQ ID NO: 38 of US20150159173). rh54 (SEQ ID NO: 49 of US20150159173), AAV2 (SEQ ID NO: 7 of US20150159173), cy.5 (SEQ ID NO: 8 and 24 of IJS20150159173), rh.lO (SEQ ID NO: 9 and 25 of US20150159173), rh.13 (SEQ ID NO: 10 and 26 of IJS20150159173), AAV1 (SEQ ID NO: 11 and 27 of US20150159173). AAV3 (SEQ ID NO: 12 and 28 of 11820150159173), AAV6 (SEQ ID NO: 13 and 29 of US20150159173), **AAV7** (SEQ ID NO: 14 and 30 of US20150159173), AAV8 (SEQ ID NO: 15 and 31 of US20150159173), hu.13 (SEQ ID NO: 16 and 32 of 11820150159173), hu.26 (SEQ ID NO: 17 and 33 of US20150159173), hu.37 (SEQ ID NO: 18 and 34 of US20150159173), hu.53 (SEQ ID NO: 19 and 35 of US20150159173), rh.43 (SEQ ID NO: 21 and 37 of US20150159173), rh2 (SEQ ID NO: 39 of US20150159173), rh.37 (SEQ ID NO: 40 of US20150159173), rh.94 (SEQ ID NO: 43 of US20150159173), rh.48 (SEQ ID NO: 44 of US20150159173), ch.5 (SEQ ID NO 46 of US20150159173), rh.67 (SEQ ID NO:

47 of US20150159173), fh.58 (SEQ ID NO: 48 of US20150159173), or variants thereof including, but not limited to Cy5Ri, Cy5R2, Cy5R3, Cy5R4, rh.13R, rh.37R2, rh.2R, rh.8R, rh.48.1, rh.48.2, rh.48.1.2, hu.44R1, hu.44R2, hu.44R3, hu.29R, ch.5Rl, rh64Ri, rb64R2, AAV6.2, AAV6.1, AAV6.12, hu.48Rl, hu.48R2, and hu.48R3.

[0061] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent No. US 7198951, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV9 (SEQ ID NO: 1-3 of US 7198951), AAV2 (SEQ ID NO: 4 of US 7198951), AAV1 (SEQ ID NO: 5 of US 7198951), AAV3 (SEQ ID NO: 6 of US 7198951), and AAV8 (SEQ ID NO: 7 of US7198951).

[0062] In some embodiments, the AAV serotype may be, or have, a mutation in the AAV9 sequence as described by N Pulicherla et al. (Molecular Therapy 19(6): 1070-1078 (2011), herein incorporated by reference in its entirety), such as but not limited to, AAV9.9, AAV9.11, AAV9.13, AAV9.16, AAV9.24, AAV9.45, AAV9.47, AAV9.61, AAV9.68, AAV9.84.

[0063] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent No. US 6156303, the contents of which are herem incorporated by reference in their entirety, such as, but not limited to. AAV3B (SEQ ID NO: I and 10 of US 6156303), AAV6 (SEQ ID NO: 2, 7 and 11 of US 6156303), AAV2 (SEQ ID NO: 3 and 8 of US 6156303), **AAV3A** (SEQ ID NO: 4 and 9, of US 6156303), or derivatives thereof.

[0064] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Publication No. US20140359799, the contents of which are herein incorporated by-reference in their entirety, such as, but not limited to, AAV8 (SEQ ID NO: 1 of US20140359799), AAVDJ (SEQ ID NO: 2 and 3 of US20140359799), or variants thereof.

[0065] In some embodiments, the serotype may be AAVDJ or a variant thereof, such as AAVDJ8 (or AAV-DJ8), as described by Grimm et al. (Journal of Virology 82(12): 5887-5911 (2008), herein incorporated by reference in its entirety). The amino acid sequence of AAVDJ8 may comprise two or more imitations in order to remove the heparin binding domain (HBD). As anon-limiting example, the AAV-DJ sequence described as SEQ ID NO: I in US Patent No. 7,588,772, the contents of which are herein incorporated by reference in their entirety, may comprise two imitations: (1) R587Q where arginine (R; Arg) at amino acid 587 is changed to glutamine (Q; Gln) and (2) R590T where arginine (R; Arg) at. ammo acid 590 is changed to threonine (T; Tbr). As another non-limiting example, may comprise three mutations: (1) K406R where lysine (K: Lys) at amino acid 406 is changed to arginine (R; Arg), (2) R587Q where arginine (R, Arg) at amino acid 587 is changed to glutamine (Q; Gin) and (3) R590T where arginine (R; Arg) at amino acid 590 is changed to threonine (T; Tbr).

[0066] In some embodiments, the AAV serotype may be, or have, a sequence of AAV4 as described in International Publication No. WO199801 l244, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to AAV4 (SEQ ID NO: 1-20 of WO19980ll244).

[0067] In some embodiments, the AAV serotype may be, or have, a mutation in the AAV2 sequence to generate AAV2G9 as described in International Publication No. WO20l4144229 and herein incorporated by reference in its entirety.

[0068] In some embodiments, the AAV serotype may be, or have, a sequence as described in international Publication No. WO2005033321, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to AAV3-3 (SEQ ID NO: 217 of WO2005033321), AAV1 (SEQ ID NO: 219 and 202 of WO2005033321), AAV106.1/hu.37 (SEQ ID No: 10 of WO2005033321), AAV114.3/hu.40 (SEQ ID No: 11 of WO2005033321), AAVl27.2/hu.41 (SEQ ID NO:6 and 8 of WO2005033321), AAVl28.3/hu.44 (SEQ ID No: 81 of WO2005033321), AAV130.4/hu.48 (SEQ ID NO: 78 of WO2005033321), AAV145.1/hu.53 (SEQ ID No: 176 and 177 of WO2005033321), AAVl45.6/hu.56 (SEQ ID NO: 168 and 192 of WO2005033321), AAVl6.12/hu.11 (SEQ ID NO: 153 and 57 of WO2005033321), AAVl6.8/hu.10(SEQ ID NO: 156 and 56 of WO2005033321), AAV16I.10/hu.60 (SEQ ID No: 170 of WO2005033321), AAVl61.6/hu.61 (SEQ ID No: 174 of WO2005033321), AAV1-7/rh.48 (SEQ ID NO: 32 of WO2005033321), AAVI-8/rh.49 (SEQ ID NOs: 103 and 25 of WO2005033321), AAV2 (SEQ ID NO: 211 and 221 of WO2005033321), AAV2-i5/rh.62 (SEQ ID No: 33 and 114 of WO2005033321), AAV2-3/rh.61 (SEQ ID NO: 21 of WO2005033321), AAV2-4/rh.50 (SEQ ID No: 23 and 108 of WO2005033321), AAV2-5/rh.51 (SEQ ID NO: 104 and 22 of WO2005033321), AAV3.1/hu.6 (SEQ ID NO: 5 and 84 of WO2005033321), AAV3.1/hu.9 (SEQ ID NO: 155 and 58 of WO2005033321), AAV3-11/rh.53 (SEQ ID NO: 186 and 176 of WO2005033321), AAV3-3 (SEQ ID NO: 200 of WO2005033321), AAV33.12/hu.17 (SEQ ID NO:4 of WO2005033321), AAV33.4/hu.15 (SEQ ID No: 50 of WO2005033321), AAV33.8/hu.16 (SEQ ID No: 51 of WO2005033321), AAV3-9/rh.52 (SEQ ID NO: 96 and 18 of WO2005033321), AAV4-19/rh.55 (SEQ ID NO: 117 of WO2005033321), AAV4-4 (SEQ ID NO: 201 and 218 of WO2005033321), AAV4-9/rh.54 (SEQ ID NO: 116 of WO2005033321), AAV5 (SEQ ID NO: 199 and 216 of WO2005033321), AAV52.1/hu.20 (SEQ ID NO: 63 of WO200503332l), AAV52/1iu.19 (SEQ ID NO: 133 of WO2005033321), AAV5-22/rh.58 (SEQ ID No: 27 of WO2005033321), AAV5-3/rh.57 (SEQ ID NO: 105 of WO2005033321), **AAV5-**3/rh.57 (SEQ ID No: 26 of WO2005033321), AAV58.2/hu.25 (SEQ ID No: 49 of WO2005033321), AAV6 (SEQ ID NO: 203 and 220 of WO2005033321), AAV7 (SEQ ID NO:

222 and 213 of WO2005033321), AAV7.3/hu.7 (SEQ ID No: 55 of WO2005033321), AAV8 (SEQ ID NO: 223 and 214 of WO2005033321), AAVH-1/hu.l (SEQ ID No: 46 of WO2005033321), AAVH-5/hu.3 (SEQ ID No: 44 of WO200503332I), AAVhu. 1 (SEQ ID NO: 144 of WO2005033321), AAVhu. 10 (SEQ ID NO: 156 of W0200503 3321), AAVhu. 11 (SEQ ID NO: 153 of WO2005033321), AAVhu. 12 (WO2005033321 SEQ ID NO: 59), AAVhu. 13 (SEQ ID NO: 129 of WO2005033321), AAVhu. 14/AAV9 (SEQ ID NO: 123 and 3 of WO2005033321), AAVhu. 15 (SEQ ID NO: 147 of WO2005033321), AAVhu.l6 (SEQ ID NO: 148 of WO2005033321), AAVhu. 17 (SEQ ID NO: 83 of WO2005033321), AAVhu. 18 (SEQ ID NO: 149 ofWO2005033321), AAVhu. 19 (SEQ ID NO: 133 of WO2005033321), AAVhu.2 (SEQ ID NO: 143 of WO2005033321), AAVhu.20 (SEQ ID NO: 134 of WO2005033321), AAVhu.21 (SEQ ID NO: 135 of WO2005033321), AAVhu.22 (SEQ ID NO: 138 of WO2005033321), AAVhu.23.2 (SEQ ID NO: 137 of WO200503332 l), AAVhu.24 (SEQ ID NO: 136 of WO2005G33321), AAVhu.25 (SEQ ID NO: 146 of WO20G5033321), AAVhu.27 (SEQ ID NO: 140 of WO2005033321), AAVhu.29 (SEQ ID NO: 132 of WO2005033321), AAVhu.3 (SEQ ID NO: 145 of WO2005033321), AAVhu.31 (SEQ ID NO: 121 of WO200503332I), AAVhu.32 (SEQ ID NO: 122 of WO2005033321), AAVhu.34 (SEQ ID NO: 125 of WO2005033321), AAVhu.35 (SEQ ID NO: 164 of WO2005033321), AAVhu.37 (SEQ ID NO: 88 of WO2005033321), AAVhu.39 (SEQ ID NO: 102 of WO2005033321), AAVhu.4 (SEQ ID NO: 141 of WO2005033321), AAVhu.40 (SEQ ID NO: 87 of WO2005033321), AAVhu.41 (SEQ ID NO: 91 of WO2005033321), AAVhu.42 (SEQ ID NO: 85 of WO2005033321), AAVhu.43 (SEQ ID NO: 160 of WO2005033321), AAVhu.44 (SEQ ID NO: 144 ofWO2005033321), AAVhu.45 (SEQ ID NO: 127 of WO2005033321), AAVhu.46 (SEQ ID NO: 159 of WO2005033321), AAVhu.47 (SEQ ID NO: 128 of WO2005033321), AAVhu.48 (SEQ ID NO: 157 of WO2005033321), AAVhu.49 (SEQ ID NO: 189 of WO2005033321), AAVhu.51 (SEQ ID NO: 190 of WO2005033321), AAVhu.52 (SEQ ID NO: 191 of WO2005033321), AAVhu.53 (SEQ ID NO: 186 of WO2005033321), AAVhu.54 (SEQ ID NO: 188 of WO2005033321), AAVhu.55 (SEQ ID NO: 187 of WO2005033321), AAVhu.56 (SEQ ID NO: 192 of WO2005033321), AAVhu.57 (SEQ ID NO: 193 of WO2005033321), AAVhu.58 (SEQ ID NO: 194 of WO2005033321), AAVhu.6 (SEQ ID NO: 84 of WO2005033321), AAVhu.60 (SEQ ID NO: 184 of WO2005033321), AAVhu.61 (SEQ ID NO: 185 of WO2005033321), AAVhu.63 (SEQ ID NO: 195 of WO2005033321), AAVhu.64 (SEQ ID NO: 196 of WO2005033321), AAVhu.66 (SEQ ID NO: 197 of WO2005033321), AAVhu.67 (SEQ ID NO: 198 of WO2005033321), AAVhu.7 (SEQ ID NO: 150 of WO2005033321), AAVhu.8 (WO2005033321 SEQ ID NO: 12), AAVhu.9 (SEQ ID NO: 155 of WO2005033321), A.AVLG-

10/fh.40 (SEQ ID No: 14 of WO2005033321), AAVLG-4/rh.38 (SEQ ID NO: 86 of WO2005033321), AAVLG-4/rh.38 (SEQ ID No: 7 of WO2005033321), AAVN721-8/rh.43 (SEQ ID NO: 163 of WO2005033321), AAVN724-8/fb.43 (SEQ ID No: 43 of WO2005033321), AAVpi.l **(WO2005033321** SEQ ID NO: 28), AAVpi.2 (WO2005033321 SEQ ID NO: 30). AAVpi.3 (WO2005033321 SEQ ID NO: 29), AAVrh.38 (SEQ ID NO: 86 of WO2005033321), AAVrh.40 (SEQ ID NO: 92 of WO2005033321), AAVrh.43 (SEQ ID NO: 163 of WO2005033321), AAVrh.44 (WO2005033321 SEQ ID NO: 34), AAVrh.45 (WO2005G3332I SEQ ID NO: 41), AAVrh.47 (WO200503332I SEQ ID NO: 38), AAVrh.48 (SEQ ID NO: 115 of WO2005033321), AAVrh.49 (SEQ ID NO: 103 of WO2005033321), AAVrb.50 (SEQ ID NO: 108 of WG2005033321), AAVrh.51 (SEQ ID NO: 104 of WO2005033321), AAVrh.52 (SEQ ID NO: 96 of WO2005033321), AAVrh.53 (SEQ ID NO: 97 ofWO2005033321), AAVrh.55 (WO2005033321 SEQ ID NO: 37), AAVrh.56 (SEQ ID NO: 152 of WO2005033324), AAVrh.57 (SEQ ID NO: 105 of WO200503332I), AAVrh.58 (SEQ ID NO: 106 of WO2005033321), AAVrh.59 (WO2005033321 SEQ ID NO: 42), AAVrh.60 (WO200503332I SEQ ID NO: 31), AAVrh.61 (SEQ ID NO: 107 of WO2005033321), AAVrb.62 (SEQ ID NO: 114 of WO2005033321), AAVrh.64 (SEQ ID NO: 99 of WO200503332I), AAVrb.65 (WO2005033321 SEQ ID NO: 35), AAVrh.68 (WO2005033321 SEQ ID NO: 16), AAVrh.69 (WO2005033321 SEQ ID NO: 39), AAVrh.70 (WO2005033321 SEQ ID NO: 20). AAVrh.72 (WO2005033321 SEQ ID NO: 9), or variants thereof including, but not limited to, AAVcy.2, AAVcy.3, AAVcy.4, AAVcy.5, AAVcy.6, AAVrhJ2, AAVrh.17, AAVrh.18, AAVrh.19, AAVrh.21, AAVrh.22, AAVrh.23, AAVrh.24, AAVih.25, AAYrh.25/42 15, AAVrh.31, AAVrh.32, AAVrh.33, AAVrh.34, AAVrh.35, AAVrh.36, AAVrh.37, AAVrh1A Non-limiting examples of variants include SEQ ID NO: 13, 15. 17, 19, 24, 36, 40, 45. 47. 48, 51-54, 60-62, 64-77, 79, 80, 82, 89, 90. 93-95, 98, 100, 101. 109-113, 118-120, 124, 126, 131, 139, 142, 151,154, 158, 161, 162, 165-183, 202, 204-212, 215, 219, 224-236, of WO2005033321, the contents of which are herein incorporated by reference in thetr entirety.

[0069] In some embodiments, the AAV serotype may be, or have, a sequence as described in International Publication No. WO2015168666, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAVrhSR (SEQ ID NO: 9 of WO2015168666), AAVrhSR A586R mutant (SEQ ID NO: 10 of WO2015168666). AAVrhSR R533A mutant (SEQ ID NO: 11 of WO2015168666), or variants thereof.

[0070] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent No. US9233131, the contents of which are herein incorporated by reference in their entirety, such as. but not limited to, AAVbEl. 1 (SEQ ID NO:44 of US9233131),

AAVhErl.5 (SEQ ID NO:45 of US9233131). AAYhER1.14 (SEQ ID NO:46 of US9233131). AAVhEr1.8 (SEQ ID NO:47 of US9233131), AAVhEr1 .I6 (SEQ ID NO:48 of US9233131), AAVhErl.18 (SEQ ID SO:49 of US9233131), AAVhErl.35 (SEQ ID NO;50 of US9233131), AAVhErl.7 (SEQ ID NO:5: of US9233131), AAVhErl.36 (SEQ ID NO:52 of TJS9233131), AAVhEr2.29 (SEQ ID NO:53 of US9233131), AAVhEr2.4 (SEQ ID NO:54 of US9233131), AAVhEr2.i6 (SEQ ID NO:55 of US9233131), AAVhEr2.30 (SEQ ID NO:56 of US9233I31), AAVhEr2.31 (SEQ ID NO:58 of US9233131), AAVhEr2.36 (SEQ ID NQ:57 of US9233131), AAVhER1.23 (SEQ ID NO: 53 of US9233131), AAVhEr3.1 (SEQ ID NO:59 of US9233131), AAV2.5T (SEQ ID NO:42 **of US9233131),** or variants thereof

[0071] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent Publication No. US20150376607, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV-PAEC (SEQ ID NO:I of US201503766G7), AAV-LK01 (SEQ ID NO:2 of US20150376607), AAV-LK02 (SEQ ID NO:3 of US20150376607), AAV-LK03 (SEQ ID NO:4 of US201 50376607), **AAV-LK04** (SEQ ID NO:5 of US201 50376607), AAV-LK.05 (SEQ ID NO:6 of US20150376607), AAV-LK06 (SEQ ID NO: 7 of US20150376607), AAV-LK07 (SEQ ID NO:8 of US201 50376607), AAV-LK08 (SEQ ID NG:9 of US20t 50376607), AAV-LK09 (SEQ ID NO. i0 of US20150376607), AAV-LK10 (SEQ ID NO: 11 of US201 50376607), AAV-LK11 (SEQ ID NO: 12 of US201 50376607). AAV-LK12 (SEQ ID NO: 13 of US201 50376607), AAV-LK13 (SEQ ID NO. 14 of US20150376607), AAV-LKI4 (SEQ ID NO:]5 of US20150376607), AAV-LK15 (SEQ ID NO:16 of US201 50376607), AAV-LK16 (SEQ ID NO: 17 of US20150376607), AAV-LK17 (SEQ ID NO: 18 of US201 50376607), AAV-LK1 8 (SEQ ID NO: 19 of US201 50376607), AAV-LK19 (SEQ ID NO: 20 of US20150376607), AAV-PAEC2 (SEQ ID NO:2i of US201503 76607), AAV-PAEC4 (SEQ ID NO:22 of US201 50376607), AAV-PAEC6 (SEQ ID NO:23 of US20150376607), AAV-PAEC 7 (SEQ ID NO:24 of US20150376607), AAV-PAEC8 (SEQ ID NO:25 of US20 150376607), AAV-PAEC 11 (SEQ ID NO:26 of US20150376607), AAV-PAEC 12 (SEQ ID NO:27, of US201 50376607), or variants thereof

|0072] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent No. US9163261, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV-2-pre-miRNA-l 01 (SEQ ID NO: 1 US9163261), or variants thereof.

[0073] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent Publication No. US201 503 76240, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV-8h (SEQ ID NO: 6

of US201 50376240), AAV-8b (SEQ ID NO: 5 of US201 50376240), AAV-h (SEQ ID NO: 2 of US201.50376240), AAV-b (SEQ ID NO: 1 of US20I50376240), or variants thereof.

[0074] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent Publication No. US2G160017295, the contents of which are herem incorporated by reference in their entirety, such as, but not limited to, AAV SM 10-2 (SEQ ID NO: 22 of US20160017295), AAV Shuffle 100-1 (SEQ ID NO: 23 of US201600I7295), AAV Shuffle 100-3 (SEQ ID NO: 24 of US20160017295), AAV Shuffle 100-7 (SEQ ID NO: 25 of US20160017295), AAV Shuffle 10-2 (SEQ ID NO: 34 of US20160017295), AAV Shuffle 10-6 (SEQ ID NO: 35 of US20160017295), AAV Shuffle 10-8 (SEQ ID NO: 36 of US20160017295), AAV Shuffle 100-2 (SEQ ID NO: 37 of US201600I7295), AAV SM 10-1 (SEQ ID NO: 38 of US20160017295), AAV SM 10-8 (SEQ ID NO: 39 of US20160017295), AAV SM 100-3 (SEQ ID NO: 40 of US20I6001729.5), AAV SM 100-10 (SEQ ID NO: 41 of US201 6001 7295), or variants thereof.

[0075] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent Publication No. US20150238550, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, BNP61 AAV (SEQ ID NO: 1 of US20I50238550), BNP62 AAV (SEQ ID NO: 3 of US20150238550), BNP63 AAV (SEQ ID NO: 4 of US20150238550), or variants thereof

[0076] In some embodiments, the AAV serotype may be or may have a sequence as described in United States Patent Publication No. US201 503156 12, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAVrh.50 (SEQ ID NO: 108 of US20150315612), AAVrh.43 (SEQ ID NO: 163 of US20150315612), AAVrh.62 (SEQ ID NO: 114 of US20150315612), AAVrh.48 (SEQ ID NO: 115 of US201503I5612), **AAVhu.19** (SEQ ID NO: 133 of US20I50315612), AAVhu.11 (SEQ ID NO: 153 of US20150315612), AAVhu.53 (SEQ ID NO: 186 of US20150315612), AAV4-8/rh.64 (SEQ ID No: 15 of US20150315612), AAVLG-9/hu.39 (SEQ ID No: 24 of US201503156I2). AAV54.5/hu.23 (SEQ ID No: 60 of US201503156I2), AAV54.2/rm.22 (SEQ ID No: 67 of US201 50315612), AAV54.7/hu.24 (SEQ ID No: 66 of US20150315612), AAV54.1/hii.2i (SEQ ID No: 65 of US201 503 15612), AAV54.4R/hu.27 (SEQ ID No: 64 of US20150315612), AAV46.21m.28 (SEQ ID No: 68 of US201503156I2), AAV46.6/hu.29 (SEQ ID No: 69 of US20150315612), AAV128.1/hu.43 (SEQ ID No: 80 of US20150315612), or variants teeof

[0077] In some embodiments, the AAV serotype may be, or have, a sequence as described in International Publication No. WO2015 121501, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, true type AAV (ttAAV) (SEQ ID NO: 2 of

WO2015121501), "UPenn AAV10" (SEQ ID NO: 8 of WO2015121501), "Japanese AAV10" (SEQ ID NO: 9 of WO2015121501), or variants thereof.

[0078] According to the present invention, AAV capsid serotype selection or use may be from a variety of species. In one embodiment, the AAV may be an avian AAV (AAAV). The AAAV serotype may be, or have, a sequence as described in United States Patent No. US 9238800, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAAV (SEQ ID NO: 1, 2, 4, 6, 8, 10, 12, and 14 of US 9,238,800), or variants thereof.

[0079] In one embodiment, the AAV may be a bovine AAV (BAAV). The BAAV serotype may be, or have, a sequence as described in United States Patent No. US 9,193,769, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, BAAV (SEQ ID NO: 1 and 6 of US 9193769), or variants thereof. The BAAV serotype may be or have a sequence as described in United States Patent No. US7427396, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, BAAV (SEQ ID NO: 5 and 6 of US7427396), or variants thereof.

[0080] In one embodiment, the AAV may be a caprine AAV. The caprine AAV serotype may be, or have, a sequence as described in United States Patent No. US7427396, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, caprine AAV (SEQ ID NO: 3 of US7427396), or variants thereof.

[0081] In other embodiments, the AAV may be engineered as a hybrid AAV from two or more parental serotypes. In one embodiment, the AAV may be AAV2G9 which comprises sequences from AAV2 and AAV9. The AAV2G9 AAV serotype may be, or have, a sequence as described in United States Patent Publication No. US20160017005, the contents of which are herein incorporated by reference in its entirety.

[0082] In one embodiment, the AAV may be a serotype generated by the AAV9 capsid library with mutations in amino acids 390-627 (VP1 numbering) as described by Pulicherla et al. (Molecular Therapy 19(6):1070-1078 (2011), the contents of which are herein incorporated by reference in their entirety. The serotype and corresponding nucleotide and amino acid substitutions may be, but is not limited to, AAV9.1 (G1594C; D532H). AAV6.2 (T1418A and T1436X; V473D and I479K), AAV9.3 (T1238A; F4I3Y), AAV9.4 (T1250C and A1617T; F417S), AAV9.5 **(A1235G,** A1314T, A1642G, C1760T; Q412R, T548A, A587V), AAV9.6 (T1231A; F411I), AAV9.9 **(G1203A,** G1785T; W595C), AAV9.10 (A1500G, T1676C; M559T), AAV9.11 (A1425T. A1702C, A1769T; T568P, Q590L), AAV9.13 (A1369C. A1720T; N457H, T574S), AAV9.14 (T1340A, T1362C, T1560C, G1713A; L447H), AAV9.16 (A1775T:

Q592L), AAV9.24 (T1507C, T1521G; W503R). AAV9.26 (A1337G, A1769C; Y446C, Q590P), AAV9.33 (A1667C; D556A), AAV9.34 (A1534G, C1794T; N512D), AAV9.35 (A1289T, T1450A, **C1494T, A1515T,** C1794A, G1816A; Q430L, **Y484N,** N98K, V606I), AAV9.40 (A1694T, E565V), AAV9.41 (A1348T, T1362C; T450S), AAY9.44 (A1684C, A1701T, A1737G; N562H, K567N), AAY9.45 (A1492T, C1804T; N498Y, L602F), AAV9.46 **(G1441C,** T1525C, T1549G; G481R, W509R, L517V), 9.47 (G1241A, G1358A, A1669G, C1745T; S414N, G453D, K557E, T582I), AAV9.48 (C1445T, A1736T; P482L, Q579L), AAV9.50 (AI638T, C1683T, T1805A; Q546H, L602H), AAV9.53 (G1301A, A1405C, C1664T, G1811T; R134Q, S469R, A555V, G604V), AAV9.54 (C1531A, T1609A; L511I, L537M), AAV9.5.5 (T1605A; F535L), AAV9.58 (C1475T, CI579A; T492I, H527N), AAV.59 (T1336C; Y446H), AAY9.61 (A1493T; N498I), AAV9.64 (C1531A, A1617T; L5111), AAV9.65 (C1335T, TI530C, C1568A; A523D), AAV9.68 (C1510A; P504TS, AAV9.80 (G1441A„G481R), AAV9.83 (C1402A, A1500T; P468T, E500D), AAV9.87 (T1464C, T1468C; S490P), AAV9.90 (A1196T; Y399F), AAV9.91 (T1316G, A1583T, C1782G, T1806C; L439R, K528I), AAV9.93 (A1273G, A1421G, A1638C, C1712T, G1732A, A1744T, A1832T; S425G, Q474R. Q546H, P571L, G578R, T582S, D611V), AAV9.94 (AI675T; M559L) and AAV9.95 (T1605A; F535L).

[0083] In some **embodiments,** the AAV serotype may be, or have, a sequence as described in International Publication No. WO20 16049230, the contents of which are herein incorporated by reference **in** their entirety, such as, but not **limited** to AAVF1/HSC1 (SEQ ID NO: 2 **and** 20 of WO20 16049230), AAVF2/HSC2 (SEQ ID NO: 3 and 21 of WO2016049230), AAVF3/HSC3 (SEQ ID NO: 5 and 22 of WO201 6049230), AAVF4/HSC4 (SEQ ID NO: 6 and 23 of WO20 16049230), AAVF5/HSC5 (SEQ ID NO: 11 and 25 **of** WO201 6049230), AAVF6/HSC6 (SEQ ID NO: 7 and 24 of WO2016049230), AAVF7/HSC7 (SEQ ID NO: 8 and 27 of WO2016049230), AAVF8/HSC8 (SEQ ID NO: 9 **and** 28 of WO201 6049230), AAVF9/HSC9 (SEQ ID NO: 10 and 29 of WO2016049230), AAVF1 1/HSCl 1 (SEQ ID NO: 4 and 26 of WO20 16049230), AAVF12/HSC12 (SEQ ID NO: 12 **and** 30 of WO20 16049230), AAVF13/HSC13 (SEQ ID NO: 14 and 31 of WO20 16049230), AAVF14/HSC14 (SEQ ID NO: 15 and 32 of WO2016049230), AAVF15/HSC15 (SEQ ID NO: 16 and 33 of WO201 6049230), AAVF16/HSC16 (SEQ ID NO: 17 and 34 of WO201 6049230). AAVF17/HSC17 (SEQ ID NO: 13 and 35 of WO2016049230), or variants or derivatives thereof

[0084] In some embodiments, the AAV serotype may be, or have, a sequence as described in United States Patent No. LIS 8734809, the contents of which are herein incorporated by reference **in** their entirety, such as, but not limited to, AAV CBr-E1 (SEQ ID NO: 13 and 87 of LTS8734809), AAV CBr-E2 (SEQ ID NO: 14 **and** 88 **of** US8734809), AAV CBr-E3 (SEQ ID

NO: 15 and 89 of US8734809), AAV CBr-E4 (SEQ ID NO: 16 and 90 of US8734809), AAV CBr-E5 (SEQ ID NO: 17 and 91 of US8734809), AAV CBr-c5 (SEQ ID NO: 18 and 92 of US8734809), AAV CBr-E6 (SEQ ID NO; 19 and 93 of US8734809), AAV CBr-E7 (SEQ ID NO: 20 and 94 of US8734809), AAV CBr-E8 (SEQ ID NO: 21 and 95 of US8734809), AAV CLv-Dl (SEQ ID NO: 22 and 96 of US8734809), AAV CLv-D2 (SEQ ID NO: 23 and 97 of LTS8734809), AAV **CLv-D3** (SEQ ID NO: 24 and 98 of US8734809), AAV **CLv-D4** (SEQ ID NO: 25 and 99 of IJS8734809), AAV CLv-D5 (SEQ ID NO: 26 and 100 of US8734809), AAV CLv-D6 (SEQ ID NO: 27 and 101 of US8734809), AAV **CLv-D7** (SEQ ID NO: 28 and 102 of US8734809), AAV CLv-D8 (SEQ ID NO: 29 and 103 of US8734809), AAV CLv-El (SEQ ID NO: 13 and 87 of US8734809), AAV **CLv-Rl** (SEQ ID NO: 30 and 104 of US8734809), AAV CLV-R2 (SEQ ID NO: 31 and 105 of US8734809), AAV CLv-R3 (SEQ ID NO: 32 and 106 of US8734809), AAV **CLv-R4** (SEQ ID NO: 33 and 107 of US8734809), AAV CLv-R5 (SEQ ID NO: 34 and 108 of US8734809), AAV CLv-R6 (SEQ ID NO: 35 and 109 of US8734809), AAV **CLv-R7** (SEQ ID NO: 36 and 110 of US8734809), AAV CLv-R8 (SEQ ID NO: 37 and lllof US8734809), AAV CLv-R9 (SEQ ID NO: 38 and 112 of US8734809), AAV CLg-Fl (SEQ ID NO: 39 and 113 of US8734809), AAV CLg-F2 (SEQ ID NO: 40 and 114 of US8734809), AAV CLg-F3 (SEQ ID NO: 41 and 115 of US8734809), AAV CLg-F4 (SEQ ID NO: 42 and 116 of **US8734809),** AAV CLg-F5 (SEQ ID NO: 43 and 117 of US8734809), AAV CLg-F6 (SEQ ID NO: 43 and 117 of US8734809). AAV CLg-F7 (SEQ ID NO: 44 and 118 of US8734809). AAV CLg-F8 (SEQ ID NO: 43 and 11.7 of US8734809), AAV CSp-1. (SEQ ID NO: 45 and 119 of LIS8734809), AAV CSpAO (SEQ ID NO: 46 and 120 of US8734809), AAV CSp-11 (SEQ ID NO: 47 and 121 of US8734809), AAV CSp-2 (SEQ ID NO: 48 and 122 of US8734809), AAV CSp-3 (SEQ ID NO: 49 and 123 of US8734809), AAV CSp-4 (SEQ ID NO: 50 and 124 of **US8734809),** AAV CSp-6 (SEQ ID NO: 51 and 125 of LTS8734809), AAV CSp-7 (SEQ ID NO: 52 and 126 **of US8734809),** AAV CSp-8 (SEQ ID NO: 53 and 127 of US8734809), AAV CSp-9 (SEQ ID NO: 54 and 128 of US8734809), AAV CHi-2 (SEQ ID NO: 55 and 129 of US8734809), AAV CHt-3 (SEQ ID NO: 56 and 130 of US8734809), AAV CKd-1 (SEQ ID NO: 57 and 131 of US8734809), AAV CKd-10 (SEQ ID NO: 58 and 132 of US8734809), AAV CKd-2 (SEQ ID NO: 59 and 133 of US8734809), AAV CKd-3 (SEQ ID NO: 60 and 134 of US8734809), AAV CKd-4 (SEQ ID NO: 61 and 135 of US8734809), AAV CKd-6 (SEQ ID NO: 62 and 136 of US8734809), AAV CKd-7 (SEQ ID NO: 63 and .137 of US8734809), AAV CKd-8 (SEQ ID NO: 64 and 138 of US8734809), AAV CLv-1 (SEQ ID NO: 35 and 139 of US8734809), AAV CLv-12 (SEQ **ID** NO: 66 and 140 of US8734809), AAV CLv-13 (SEQ ID NO: 67 and 141 of US8734809), AAV CLv-2 (SEQ ID NO: 68 and 142 of US8734809), AAV

CLv-3 (SEQ ID NO: 69 and 143 of US8734809), AAV CLv-4 (SEQ ID NO: 70 and 144 of US8734809), AAV CLv-6 (SEQ ID NO: 71 and 145 of US8734809), AAV CLv-8 (SEQ ID NO: 72 and 146 of US8734809), AAV CKd-Bl (SEQ ID NO: 73 and 147 of US8734809), AAV CKd-B2 (SEQ ID NO: 74 and 148 of US8734809), AAV CKd-B3 (SEQ ID NO: 75 and 149 of **US8734809),** AAV CKd-B4 (SEQ ID NO: 76 and 150 of US8734809), AAV CKd-B5 (SEQ **ID** NO: 77 and 151 of US8734809), AAV CKd-B6 (SEQ ID NO: 78 and 152 of US8734809), AAV CKd-B7 (SEQ **ID** NO: 79 and 153 of US8734809), AAV CKd-B8 (SEQ ID NO: 80 and 154 of US8734809), AAV CKd-H1 (SEQ ID NO: 81 and 155 of IJS8734809), AAV CKd-H2 (SEQ ID NO: 82 and 156 of US8734809), AAV CKd-H3 (SEQ ID NO: 83 and 157 of US8734809), AAV CKd-H4 (SEQ ID NO: 84 and 158 of US8734809), AAV CKd-H5 (SEQ ID NO: 85 and 159 of **US8734809),** AAV CKd-H6 (SEQ ID NO: 77 and 151 of US8734809), AAV CHt-1 (SEQ ID NO: 86 and 160 of US8734809), AAV CLvl-1 (SEQ **ID** NO: 171 of US8734809), AAV **CLvl-2** (SEQ ID NO: 172 of US8734809), AAV CLvl-3 (SEQ ID NO: 173 of US8734809), AAV **CLvl-4** (SEQ ID NO: 174 of US8734809), AAV Clvl-7 (SEQ ID NO: 175 of US8734809), AAV Civ'l-8 (SEQ ID NO: 176 of US8734809), AAV Civ1-9 (SEQ ID NO: 177 of US8734809), AAV Civl-10 (SEQ ID NO: 178 of US8734809), AAV.VR-355 (SEQ ID NO: 181 of US8734809), AAV.hu.48R3 (SEQ **ID** NO: 183 of US8734809), or variants or derivatives thereof.

[0085] In some embodiments, the AAV serotype may be, or have, a sequence as described in International Publication No. WO20 16065001, the contents of which are herein incorporated by-reference in their entirety, such as, but not limited to AAV CHt-P2 (SEQ ID NO: 1 and 51 of WO2016065001), AAV CII1-P5 (SEQ ID NO: 2 and 52 of WO2016065001), AAV CHt-P9 (SEQ ID NO: 3 and 53 of WO2016065001), AAV CBr-7.1 (SEQ ID NO: 4 and 54 of WO2016065001), AAV CBr-7.2 (SEQ ID NO: 5 and 55 of WO2016065001), AAV CBr-7.3 (SEQ ID NO: 6 and 56 of WO2016065001), AAV CBr-7.4 (SEQ ID NO: 7 and 57 of WO2016065001), AAV CBr-7.5 (SEQ ID NO: 8 and 58 of WO2016065001), AAV CBr-7.7 (SEQ ID NO: 9 and 59 of WO2016065001), AAV CBr-7.8 (SEQ ID NO: 10 and 60 of WO2016065001), AAV CBr-7.10 (SEQ ID NO: 11 and 61 of WO2016065001), AAV CKd-N3 (SEQ ID NO: 12 and 62 of WO2016065001), AAV CKd-N4 (SEQ ID NO: 13 and 63 of WO2016065001), AAV CKd-N9 (SEQ ID NO: 14 and 64 of WO2016065001), AAV CLv-L4 (SEQ ID NO: 15 and 65 of WO2016065001), AAV CLv~L5 (SEQ ID NO: 16 and 66 of WO2016065001), AAV CLv-L6 (SEQ ID NO: 17 and 67 of WO2016065001), AAV CLv-Kl (SEQ ID NO: 18 and 68 of WO2016065001), AAV CLv-K3 (SEQ ID NO: 19 and 69 of WO2016065001), AAV CLv-K6 (SEQ ID NO: 20 and 70 of WO2016065001), AAV CLv-M1

(SEQ ID NO: 21 and 71 of WO20 16065001), AAV CLv-M11 (SEQ ID NO: 22 and 72 of WO2016065001), AAV CLv-M2 (SEQ ID NO: 23 and 73 of WO2016065001), AAV CLv-M5 (SEQ ID NO: 24 and 74 of WO2016065001), AAV CLv-M6 (SEQ ID NO: 25 and 75 of WO2016065001), AAV **CLv-M7** (SEQ ID NO: 26 and 76 of WO2016065001), AAV **CLv-M8** (SEQ ID NO: 27 and 77 of WO2016065001), AAV CLv-M9 (SEQ ID NO: 28 and 78 of WO20 16065001), AAV CHt-Pl (SEQ ID NO: 29 and 79 of WO20 16065001), AAV CHt-F6 (SEQ ID NO: 30 and 80 of WO20 16065001), AAV CHt-P8 (SEQ ID NO: 31 and 81 of WO2016065001), AAV CHt-6.1 (SEQ ID NO: 32 and 82 of WO20 16065001), AAV CHt-6.10 (SEQ ID NO: 33 and 83 of WO20 16065001), AAV CHt-6.5 (SEQ ID NO: 34 and 84 of WO20 16065001), AAV CHt-6.6 (SEQ ID NO: 35 and 85 of WO20 16065001). AAV CHt-6.7 (SEQ ID NO: 36 and 86 of WO20 16065001), AAV CHt-6.8 (SEQ ID NO: 37 and 87 of WO20I6065001), AAV CSp-8.10 (SEQ ID NO: 38 and 88 of WO20I6065001), AAV CSp-8.2 (SEQ ID NO: 39 and 89 of WO2016065001), AAV CSp-8.4 (SEQ ID NO: 40 and 90 of WO2016065001), AAV CSp-8.5 (SEQ ID NO: 41 and 91 of WO2016065001), AAV CSp-8.6 (SEQ ID NO: 42 and 92 of WO2016065001), AAV CSp-8.7 (SEQ ID NO: 43 and 93 of WO20 16065001), AAV CSp-8.8 (SEQ ID NO: 44 and 94 of WO20 16065001), AAV CSp-8.9 (SEQ ID NO: 45 and 95 of WO20 16065001), AAV CBr-B7.3 (SEQ ID NO: 46 and 96 of WO2016065001), AAV CBr-B7.4 (SEQ ID NO: 47 and 97 of WO2016065001), AAV3B (SEQ ID NO: 48 and 98 of WO20 16065001), AAV4 (SEQ ID NO: 49 and 99 of WO20 16065001), AAV5 (SEQ ID NO: 50 and 100 of WO20 16065001), or variants or derivatives thereof.

[0086] In one embodiment, the AAV may be a serotype selected from any of those found in Table 1.

[0087] In one embodiment, the AAV may comprise a sequence, fragment or variant thereof, of the sequences in Table 1.

[0088] In one embodiment, the AAV may be encoded by a sequence, fragment or variant as described in Table 1.

Table 1„ AAV Serotypes

Serotype	SEQ ID NO	Reference Information
AAV1	1	US20150159173 SEQ ID NO: 11, US20150315612 SEQ ID NO: 202
AAV1	2	US20160017295 SEQ ID NO: 1US20030138772 SEQ ID NO: 64, US20150159173 SEQ ID NO: 27, US20150315612 SEQ ID NO: 219, US7198951 SEQ ID NO: 5
AAV1	3	US20030138772 SEQ ID NO: 6
AAV1.3	4	US20030138772 SEQ ID NO: 14
AAV10	5	US20030138772 SEQ ID NO: 117
AAV10	6	WO2015121501 SEQ ID NO: 9

AAV10	7	WO2015121501 SEQ ID NO: 8
AAV11	8	US20030138772 SEQ ID NO: 118
AAV12	9	US20030138772 SEQ ID NO: 119
AAV2	10	US20150159173 SEQ ID NO: 7, US201503 15612 SEQ ID NO: 211
AAV2	11	US20030138772 SEQ ID NO: 70, US20150 159173 SEQ ID NO: 23, US20 1503 156 I2 SEQ ID NO: 221. US20 160017295 SEQ ID NO: 2, b S6156303 SEQ ID NO: 4, US7198951 SEQ ID NO: 4, WO2015 121501 SEQ ID NO: 1
AAV2	12	US6156303 SEQ ID NO: 8
AAV2	13	US20030138772 SEQ ID NO: 7
AAV2	14	US6156303 SEQ ID NO: 3
AAV2.5T	15	US9233 131 SEQ ID NO: 42
AAV223. 10	16	US20030 138772 SEQ ID NO: 75
AAV223.2	17	US20030138772 SEQ ID NO: 49
AAV223.2	18	US20030138772 SEQ ID NO: 76
AAV223.4	19	US20030 138772 SEQ ID NO: 50
AAV223.4	20	US20030138772 SEQ ID NO: 73
AAV223.5	21	US20030 138772 SEQ ID NO: 51
AAV223.5	22	US20030138772 SEQ ID NO: 74
AAV223.6	23	US20030138772 SEQ ID NO: 52
AAV223.6	24	US20030 138772 SEQ ID NO: 78
AAV223.7	25	US20030138772 SEQ ID NO: 53
AAV223.7	26	US20030 138772 SEQ ID NO: 77
AAV29.3	27	US20030138772 SEQ ID NO: 82
AAV29.4	28	US20030138772 SEQ ID NO: 12
AAV29.5	29	US20030138772 SEQ ID NO: 83
AAV29.5 (AAVbb.2)	30	US20030138772 SEQ ID NO: 13
AAV3	31	US20150 159173 SEQ ID NO: 12
AAV3	32	US20030138772 SEQ ID NO: 71, US20150 159 173 SEQ ID NO: 28, US20 160017295 SEQ ID NO: 3, US719895 1 SEQ ID NO: 6
AAV3	33	US20030 138772 SEQ ID NO: 8
AAV3.3b	34	US20030138772 SEQ ID NO: 72
AAV3-3	35	US201503 15612 SEQ ID NO: 200
AAV3-3	36	US201503 15612 SEQ ID NO: 217
AAV3a	37	US6156303 SEQ ID NO: 5
AAV3a	38	IJS6 156303 SEQ ID NO: 9
AAV3b	39	US6156303 SEQ ID NO: 6
AAV3b	40	US6156303 SEQ ID NO: 10
AAV3b	41	US6156303 SEQ ID NO: 1
AAV4	42	US20 140348794 SEQ ID NO: 17
AAV4	43	US20140348794 SEQ ID NO: 5
AAV4	44	US20 140348794 SEQ ID NO: 3
AAV4	45	US20 140348794 SEQ ID NO: 14
AAV4	46	US20 140348794 SEQ ID NO: 15
AAV4	47	US20 140348794 SEQ ID NO: 19
AAV4	48	US20140348794 SEQ ID NO: 12

AAV4	49	US20140348794 SEQ ID NO: 13
AAV4	50	US20140348794 SEQ ID NO: 7
AAV4	51	US20140348794 SEQ ID NO: 8
AAV4	52	US20140348794 SEQ ID NO: 9
AAV4	53	US20140348794 SEQ ID NO: 2
AAV4	54	US20140348794 SEQ ID NO: 10
AAV4	55	US20140348794 SEQ ID NO: 11
AAV4	56	US20140348794 SEQ ID NO: 18
AAV4	57	US20030138772 SEQ ID NO: 63, US20160017295 SEQ ID NO: 4, US20140348794 SEQ ID NO: 4
AAV4	58	US20140348794 SEQ ID NO: 16
AAV4	59	US20140348794 SEQ ID NO: 20
AAV4	60	US20140348794 SEQ ID NO: 6
AAV4	61	US20140348794 SEQ ID NO: 1
AAV42.2	62	US20030138772 SEQ ID NO: 9
AAV42.2	63	US20030138772 SEQ ID NO: 102
AAV42.3b	64	US20030138772 SEQ ID NO: 36
AAV42.3B	65	US20030138772 SEQ ID NO: 107
AAV42.4	66	US20030138772 SEQ ID NO: 33
AAV42.4	67	US20030138772 SEQ ID NO: 88
AAV42.8	68	US20030138772 SEQ ID NO: 27
AAV42.8	69	US20030138772 SEQ ID NO: 85
AAV43.1	70	US20030138772 SEQ ID NO: 39
AAV43.1	71	US20030138772 SEQ ID NO: 92
AAV43.12	72	US20030138772 SEQ ID NO: 41
AAV43.12	73	US20030138772 SEQ ID NO: 93
AAV43.20	74	US20030138772 SEQ ID NO: 42
AAV43.20	75	US20030138772 SEQ ID NO: 99
AAV43.21	76	US20030138772 SEQ ID NO: 43
AAV43.21	77	US20030138772 SEQ ID NO: 96
AAV43.23	78	US20030138772 SEQ ID NO: 44
AAV43.23	79	US20030138772 SEQ ID NO: 98
AAV43.25	80	US20030138772 SEQ ID NO: 45
AAV43.25	81	US20030138772 SEQ ID NO: 97
AAV43.5	82	US20030138772 SEQ ID NO: 40
AAV43.5	83	US20030138772 SEQ ID NO: 94
AAV4-4	84	US20150315612 SEQ ID NO: 201
AAV4-4	85	US20150315612 SEQ ID NO: 218
AAV44.1	86	US20030138772 SEQ ID NO: 46
AAV44.1	87	US20030138772 SEQ ID NO: 79
AAV44.5	88	US20030138772 SEQ ID NO: 47
AAV44.5	89	US20030138772 SEQ ID NO: 80
AAV4407	90	US20150315612 SEQ ID NO: 90
AAV5	91	US7427396 SEQ ID NO: 1
AAV5	92	US20030138772 SEQ ID NO: 114

AAV5	93	US20160017295 SEQ ID NO: 5, US7427396 SEQ ID NO: 2, US20150315612 SEQ ID NO: 216
AAV5	94	US20150315612 SEQ ID NO: 199
AAV6	95	US20150159173 SEQ ID NO: 13
AAV6	96	US20030138772 SEQ ID NO: 65, US20150159173 SEQ ID NO: 29, US20160017295 SEQ ID NO: 6, US6156303 SEQ ID NO: 7
AAV6	97	US6156303 SEQ ID NO: 11
AAV6	98	US6156303 SEQ ID NO: 2
AAV6	99	US20150315612 SEQ ID NO: 203
AAV6	100	US20150315612 SEQ ID NO: 220
AAV6.1	101	US20150159173
AAV6.12	102	US20150159173
AAV6.2	103	US20150159173
AAV7	104	US20150159173 SEQ ID NO: 14
AAV7	105	US20150315612 SEQ ID NO: 183
AAV7	106	US20030138772 SEQ ID NO: 2, US20150159173 SEQ ID NO: 30, US20150315612 SEQ ID NO: 181, US20160017295 SEQ ID NO: 7
AAV7	107	US20030138772 SEQ ID NO: 3
AAV7	108	US20030138772 SEQ ID NO: 1, US20150315612 SEQ ID NO: 180
AAV7	109	US20150315612 SEQ ID NO: 213
AAV7	110	US20150315612 SEQ ID NO: 222
AAV8	111	US20150159173 SEQ ID NO: 15
AAV8	112	US20150376240 SEQ ID NO: 7
AAV8	113	US20030138772 SEQ ID NO: 4, US20150315612 SEQ ID NO: 182
AAV8	114	US20030138772 SEQ ID NO: 95, US20140359799 SEQ ID NO: 1, US20150159173 SEQ ID NO: 31, US20160017295 SEQ ID NO: 8, US7198951 SEQ ID NO: 7, US20150315612 SEQ ID NO: 223
AAV8	115	US20150376240 SEQ ID NO: 8
AAV8	116	US20150315612 SEQ ID NO: 214
AAV-8b	117	US20150376240 SEQ ID NO: 5
AAV-8b	118	US20150376240 SEQ ID NO: 3
AAV-8h	119	US20150376240 SEQ ID NO: 6
AAV-8h	120	US20150376240 SEQ ID NO: 4
AAV9	121	US20030138772 SEQ ID NO: 5
AAV9	122	US7198951 SEQ ID NO: 1
AAV9	123	US20160017295 SEQ ID NO: 9
AAV9	124	US20030138772 SEQ ID NO: 100, US7198951 SEQ ID NO: 2
AAV9	125	US7198951 SEQ ID NO: 3
AAV9 (AAVhu.14)	126	US7906111 SEQ ID NO: 3; WO2015038958 SEQ ID NO: 11
AAV9 (AAVhu.14)	127	US7906111 SEQ ID NO: 123; WO2015038958 SEQ ID NO: 2
AAVA3.1	128	US20030138772 SEQ ID NO: 120
AAVA3.3	129	US20030138772 SEQ ID NO: 57
AAVA3.3	130	US20030138772 SEQ ID NO: 66
AAVA3.4	131	US20030138772 SEQ ID NO: 54
AAVA3.4	132	US20030138772 SEQ ID NO: 68
AAVA3.5	133	US20030138772 SEQ ID NO: 55

AAVA3.5	134	US20030138772 SEQ ID NO: 69
AAVA3.7	135	US20030138772 SEQ ID NO: 56
AAVA3.7	136	US20030138772 SEQ ID NO: 67
AAV29.3 (AAVbb.1)	137	US20030138772 SEQ ID NO: 11
AAVC2	138	US20030138772 SEQ ID NO: 61
AAVCh.5	139	US20150159173 SEQ ID NO: 46, US20150315612 SEQ ID NO: 234
AAVcy.2 (AAV13.3)	140	US20030138772 SEQ ID NO: 15
AAV24.1	141	US20030138772 SEQ ID NO: 101
AAVcy.3 (AAV24.1)	142	US20030138772 SEQ ID NO: 16
AAV27.3	143	US20030138772 SEQ ID NO: 104
AAVcy.4 (AAV27.3)	144	US20030138772 SEQ ID NO: 17
AAVcy.5	145	US20150315612 SEQ ID NO: 227
AAV7.2	146	US20030138772 SEQ ID NO: 103
AAVcy.5 (AAV7.2)	147	US20030138772 SEQ ID NO: 18
AAV16.3	148	US20030138772 SEQ ID NO: 105
AAVcy.6 (AAV16.3)	149	US20030138772 SEQ ID NO: 10
AAVcy.5	150	US20150159173 SEQ ID NO: 8
AAVcy.5	151	US20150159173 SEQ ID NO: 24
AAVCy.5R1	152	US20150159173
AAVCy.5R2	153	US20150159173
AAVCy.5R3	154	US20150159173
AAVCy.5R4	155	US20150159173
AAVDJ	156	US20140359799 SEQ ID NO: 3, US7588772 SEQ ID NO: 2
AAVDJ	157	US20140359799 SEQ ID NO: 2, US7588772 SEQ ID NO: 1
AAVDJ-8	158	US7588772; Grimm et al 2008
AAVDJ-8	159	US7588772; Grimm et al 2008
AAVF5	160	US20030138772 SEQ ID NO: 110
AAVH2	161	US20030138772 SEQ ID NO: 26
AAVH6	162	US20030138772 SEQ ID NO: 25
AAVhE1.1	163	US9233131 SEQ ID NO: 44
AAVhEr1.14	164	US9233131 SEQ ID NO: 46
AAVhEr1.16	165	US9233131 SEQ ID NO: 48
AAVhEr1.18	166	US9233131 SEQ ID NO: 49
AAVhEr1.23 (AAVhEr2.29)	167	US9233131 SEQ ID NO: 53
AAVhEr1.35	168	US9233131 SEQ ID NO: 50
AAVhEr1.36	169	US9233131 SEQ ID NO: 52
AAVhEr1.5	170	US9233131 SEQ ID NO: 45
AAVhEr1.7	171	US9233131 SEQ ID NO: 51
AAVhEr1.8	172	US9233131 SEQ ID NO: 47
AAVhEr2.16	173	US9233131 SEQ ID NO: 55

AAVhEr2.30	174	US9233131 SEQ ID NO: 56
AAVhEr2.31	175	US9233131 SEQ ID NO: 58
AAVhEr2.36	176	US9233131 SEQ ID NO: 57
AAVhEr2.4	177	US9233131 SEQ ID NO: 54
AAVhEr3.1	178	US9233131 SEQ ID NO: 59
AAVhu.1	179	US20150315612 SEQ ID NO: 46
AAVhu.1	180	US20150315612 SEQ ID NO: 144
AAVhu.10 (AAV16.8)	181	US20150315612 SEQ ID NO: 56
AAVhu.10 (AAV16.8)	182	US20150315612 SEQ ID NO: 156
AAVhu.11 (AAV16.12)	183	US20150315612 SEQ ID NO: 57
AAVhu.11 (AAV16.12)	184	US20150315612 SEQ ID NO: 153
AAVhu.12	185	US20150315612 SEQ ID NO: 59
AAVhu.12	186	US20150315612 SEQ ID NO: 154
AAVhu.13	187	US20150159173 SEQ ID NO: 16, US20150315612 SEQ ID NO: 71
AAVhu.13	188	US20150159173 SEQ ID NO: 32, US20150315612 SEQ ID NO: 129
AAVhu.136.1	189	US20150315612 SEQ ID NO: 165
AAVhu.140.1	190	US20150315612 SEQ ID NO: 166
AAVhu.140.2	191	US20150315612 SEQ ID NO: 167
AAVhu.145.6	192	US20150315612 SEQ ID No: 178
AAVhu.15	193	US20150315612 SEQ ID NO: 147
AAVhu.15 (AAV33.4)	194	US20150315612 SEQ ID NO: 50
AAVhu.156.1	195	US20150315612 SEQ ID No: 179
AAVhu.16	196	US20150315612 SEQ ID NO: 148
AAVhu.16 (AAV33.8)	197	US20150315612 SEQ ID NO: 51
AAVhu.17	198	US20150315612 SEQ ID NO: 83
AAVhu.17 (AAV33.12)	199	US20150315612 SEQ ID NO: 4
AAVhu.172.1	200	US20150315612 SEQ ID NO: 171
AAVhu.172.2	201	US20150315612 SEQ ID NO: 172
AAVhu.173.4	202	US20150315612 SEQ ID NO: 173
AAVhu.173.8	203	US20150315612 SEQ ID NO: 175
AAVhu.18	204	US20150315612 SEQ ID NO: 52
AAVhu.18	205	US20150315612 SEQ ID NO: 149
AAVhu.19	206	US20150315612 SEQ ID NO: 62
AAVhu.19	207	US20150315612 SEQ ID NO: 133
AAVhu.2	208	US20150315612 SEQ ID NO: 48
AAVhu.2	209	US20150315612 SEQ ID NO: 143

AAVhu.20	210	US20150315612 SEQ ID NO: 63
AAVhu.20	211	US20150315612 SEQ ID NO: 134
AAVhu.21	212	US20150315612 SEQ ID NO: 65
AAVhu.21	213	US20150315612 SEQ ID NO: 135
AAVhu.22	214	US20150315612 SEQ ID NO: 67
AAVhu.22	215	US20150315612 SEQ ID NO: 138
AAVhu.23	216	US20150315612 SEQ ID NO: 60
AAVhu.23.2	217	US20150315612 SEQ ID NO: 137
AAVhu.24	218	US20150315612 SEQ ID NO: 66
AAVhu.24	219	US20150315612 SEQ ID NO: 136
AAVhu.25	220	US20150315612 SEQ ID NO: 49
AAVhu.25	221	US20150315612 SEQ ID NO: 146
AAVhu.26	222	US20150159173 SEQ ID NO: 17, US20150315612 SEQ ID NO: 61
AAVhu.26	223	US20150159173 SEQ ID NO: 33, US20150315612 SEQ ID NO: 139
AAVhu.27	224	US20150315612 SEQ ID NO: 64
AAVhu.27	225	US20150315612 SEQ ID NO: 140
AAVhu.28	226	US20150315612 SEQ ID NO: 68
AAVhu.28	227	US20150315612 SEQ ID NO: 130
AAVhu.29	228	US20150315612 SEQ ID NO: 69
AAVhu.29	229	US20150159173 SEQ ID NO: 42, US20150315612 SEQ ID NO: 132
AAVhu.29	230	US20150315612 SEQ ID NO: 225
AAVhu.29R	231	US20150159173
AAVhu.3	232	US20150315612 SEQ ID NO: 44
AAVhu.3	233	US20150315612 SEQ ID NO: 145
AAVhu.30	234	US20150315612 SEQ ID NO: 70
AAVhu.30	235	US20150315612 SEQ ID NO: 131
AAVhu.31	236	US20150315612 SEQ ID NO: 1
AAVhu.31	237	US20150315612 SEQ ID NO: 121
AAVhu.32	238	US20150315612 SEQ ID NO: 2
AAVhu.32	239	US20150315612 SEQ ID NO: 122
AAVhu.33	240	US20150315612 SEQ ID NO: 75
AAVhu.33	241	US20150315612 SEQ ID NO: 124
AAVhu.34	242	US20150315612 SEQ ID NO: 72
AAVhu.34	243	US20150315612 SEQ ID NO: 125
AAVhu.35	244	US20150315612 SEQ ID NO: 73
AAVhu.35	245	US20150315612 SEQ ID NO: 164
AAVhu.36	246	US20150315612 SEQ ID NO: 74
AAVhu.36	247	US20150315612 SEQ ID NO: 126
AAVhu.37	248	US20150159173 SEQ ID NO: 34, US20150315612 SEQ ID NO: 88
AAVhu.37 (AAV106.1)	249	US20150315612 SEQ ID NO: 10, US20150159173 SEQ ID NO: 18
AAVhu.38	250	US20150315612 SEQ ID NO: 161
AAVhu.39	251	US20150315612 SEQ ID NO: 102
AAVhu.39 (AAVLG-9)	252	US20150315612 SEQ ID NO: 24
AAVhu.4	253	US20150315612 SEQ ID NO: 47

AAVhu.4	254	US20150315612 SEQ ID NO: 141
AAVhu.40	255	US20150315612 SEQ ID NO: 87
AAVhu.40 (AAV114.3)	256	US20150315612 SEQ ID No: 11
AAVhu.41	257	US20150315612 SEQ ID NO: 91
AAVhu.41 (AAV127.2)	258	US20150315612 SEQ ID NO: 6
AAVhu.42	259	US20150315612 SEQ ID NO: 85
AAVhu.42 (AAV127.5)	260	US20150315612 SEQ ID NO: 8
AAVhu.43	261	US20150315612 SEQ ID NO: 160
AAVhu.43	262	US20150315612 SEQ ID NO: 236
AAVhu.43 (AAV128.1)	263	US20150315612 SEQ ID NO: 80
AAVhu.44	264	US20150159173 SEQ ID NO: 45, US20150315612 SEQ ID NO: 158
AAVhu.44 (AAV128.3)	265	US20150315612 SEQ ID NO: 81
AAVhu.44R1	266	US20150159173
AAVhu.44R2	267	US20150159173
AAVhu.44R3	268	US20150159173
AAVhu.45	269	US20150315612 SEQ ID NO: 76
AAVhu.45	270	US20150315612 SEQ ID NO: 127
AAVhu.46	271	US20150315612 SEQ ID NO: 82
AAVhu.46	272	US20150315612 SEQ ID NO: 159
AAVhu.46	273	US20150315612 SEQ ID NO: 224
AAVhu.47	274	US20150315612 SEQ ID NO: 77
AAVhu.47	275	US20150315612 SEQ ID NO: 128
AAVhu.48	276	US20150159173 SEQ ID NO: 38
AAVhu.48	277	US20150315612 SEQ ID NO: 157
AAVhu.48 (AAV130.4)	278	US20150315612 SEQ ID NO: 78
AAVhu.48R1	279	US20150159173
AAVhu.48R2	280	US20150159173
AAVhu.48R3	281	US20150159173
AAVhu.49	282	US20150315612 SEQ ID NO: 209
AAVhu.49	283	US20150315612 SEQ ID NO: 189
AAVhu.5	284	US20150315612 SEQ ID NO: 45
AAVhu.5	285	US20150315612 SEQ ID NO: 142
AAVhu.51	286	US20150315612 SEQ ID NO: 208
AAVhu.51	287	US20150315612 SEQ ID NO: 190
AAVhu.52	288	US20150315612 SEQ ID NO: 210
AAVhu.52	289	US20150315612 SEQ ID NO: 191
AAVhu.53	290	US20150159173 SEQ ID NO: 19
AAVhu.53	291	US20150159173 SEQ ID NO: 35
AAVhu.53 (AAV145.1)	292	US20150315612 SEQ ID NO: 176
AAVhu.54	293	US20150315612 SEQ ID NO: 188
AAVhu.54 (AAV145.5)	294	US20150315612 SEQ ID No: 177

AAVhu.55	295	US20150315612 SEQ ID NO: 187
AAVhu.56	296	US20150315612 SEQ ID NO: 205
AAVhu.56 (AAV145.6)	297	US20150315612 SEQ ID NO: 168
AAVhu.56 (AAV145.6)	298	US20150315612 SEQ ID NO: 192
AAVhu.57	299	US20150315612 SEQ ID NO: 206
AAVhu.57	300	US20150315612 SEQ ID NO: 169
AAVhu.57	301	US20150315612 SEQ ID NO: 193
AAVhu.58	302	US20150315612 SEQ ID NO: 207
AAVhu.58	303	US20150315612 SEQ ID NO: 194
AAVhu.6 (AAV3.1)	304	US20150315612 SEQ ID NO: 5
AAVhu.6 (AAV3.1)	305	US20150315612 SEQ ID NO: 84
AAVhu.60	306	US20150315612 SEQ ID NO: 184
AAVhu.60 (AAV161.10)	307	US20150315612 SEQ ID NO: 170
AAVhu.61	308	US20150315612 SEQ ID NO: 185
AAVhu.61 (AAV161.6)	309	US20150315612 SEQ ID NO: 174
AAVhu.63	310	US20150315612 SEQ ID NO: 204
AAVhu.63	311	US20150315612 SEQ ID NO: 195
AAVhu.64	312	US20150315612 SEQ ID NO: 212
AAVhu.64	313	US20150315612 SEQ ID NO: 196
AAVhu.66	314	US20150315612 SEQ ID NO: 197
AAVhu.67	315	US20150315612 SEQ ID NO: 215
AAVhu.67	316	US20150315612 SEQ ID NO: 198
AAVhu.7	317	US20150315612 SEQ ID NO: 226
AAVhu.7	318	US20150315612 SEQ ID NO: 150
AAVhu.7 (AAV7.3)	319	US20150315612 SEQ ID NO: 55
AAVhu.71	320	US20150315612 SEQ ID NO: 79
AAVhu.8	321	US20150315612 SEQ ID NO: 53
AAVhu.8	322	US20150315612 SEQ ID NO: 12
AAVhu.8	323	US20150315612 SEQ ID NO: 151
AAVhu.9 (AAV3.1)	324	US20150315612 SEQ ID NO: 58
AAVhu.9 (AAV3.1)	325	US20150315612 SEQ ID NO: 155
AAV-LK01	326	US20150376607 SEQ ID NO: 2
AAV-LK01	327	US20150376607 SEQ ID NO: 29
AAV-LK02	328	US20150376607 SEQ ID NO: 3
AAV-LK02	329	US20150376607 SEQ ID NO: 30
AAV-LK03	330	US20150376607 SEQ ID NO: 4
AAV-LK03	331	WO2015121501 SEQ ID NO: 12, US20150376607 SEQ ID NO: 31
AAV-LK04	332	US20150376607 SEQ ID NO: 5
AAV-LK04	333	US20150376607 SEQ ID NO: 32
AAV-LK05	334	US20150376607 SEQ ID NO: 6

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AAV-LK06	336	US20150376607 SEQ ID NO: 7
AAV-LK06	337	US20150376607 SEQ ID NO: 34
AAV-LK07	338	US20150376607 SEQ ID NO: 8
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AAV-LK08	340	1JS20150376607 SEQ ID NO: 9
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AAV-LK11	346	US20150376607 SEQ ID NO: 12
AAV-LK11	347	US20150376607 SEQ ID NO: 39
AAV-LK12	348	US20150376607 SEQ ID NO: 13
AAV-LK12	349	US20150376607 SEQ ID NO: 40
AAV-LK13	350	1JS20150376607 SEQ ID NO: 14
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AAV-LK15	354	US20150376607 SEQ ID NO: 16
AAV-LK15	355	1JS20150376607 SEQ ID NO: 43
AAV-LK16	356	US20150376607 SEQ ID NO: 17
AAV-LK16	357	US20150376607 SEQ ID NO: 44
AAV-LK17	358	US20150376607 SEQ ID NO: 18
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AAV-LK19	362	US20150376607 SEQ ID NO: 20
AAV-LK19	363	US20150376607 SEQ ID NO: 47
AAV-PAEC	364	US20150376607 SEQ ID NO: 1
AAV-PAEC	365	US20150376607 SEQ ID NO: 48
AAV-PAEC11	366	US20150376607 SEQ ID NO: 26
AAV-PAEC11	367	US20150376607 SEQ ID NO: 54
AAV-PAEC12	368	US20150376607 SEQ ID NO: 27
AAV-PAEC12	369	US20150376607 SEQ ID NO: 51
AAV-PAEC13	370	US20150376607 SEQ ID NO: 28
AAV-PAEC13	371	US20150376607 SEQ ID NO: 49
AAV-PAEC2	372	US20150376607 SEQ ID NO: 21
AAV-PAEC2	373	US20150376607 SEQ ID NO: 56
AAV-PAEC4	374	US20150376607 SEQ ID NO: 22
AAV-PAEC4	375	US20150376607 SEQ ID NO: 55
AAV-PAEC6	376	1JS20150376607 SEQ ID NO: 23

AAV-PAEC6	377	US20150376607 SEQ ID NO: 52
AAV-PAEC7	378	US20150376607 SEQ ID NO: 24
AAV-PAEC7	379	US20150376607 SEQ ID NO: 53
AAV-PAEC8	380	US20150376607 SEQ ID NO: 25
AAV-PAEC8	381	US20150376607 SEQ ID NO: 50
AAVpi.1	382	US20150315612 SEQ ID NO: 28
AAVpi.1	383	US20150315612 SEQ ID NO: 93
AAVpi.2	384	US20150315612 SEQ ID NO: 30
AAVpi.2	385	US20150315612 SEQ ID NO: 95
AAVpi.3	386	US20150315612 SEQ ID NO: 29
AAVpi.3	387	US20150315612 SEQ ID NO: 94
AAVrh.10	388	US20150159173 SEQ ID NO: 9
AAVrh.10	389	US20150159173 SEQ ID NO: 25
AAV44.2	390	US20030138772 SEQ ID NO: 59
AAVrh.10 (AAV44.2)	391	US20030138772 SEQ ID NO: 81
AAV42.1B	392	US20030138772 SEQ ID NO: 90
AAVrh.12 (AAV42.1b)	393	US20030138772 SEQ ID NO: 30
AAVrh.13	394	US20150159173 SEQ ID NO: 10
AAVrh.13	395	US20150159173 SEQ ID NO: 26
AAVrh.13	396	US20150315612 SEQ ID NO: 228
AAVrh.13R	397	US20150159173
AAV42.3A	398	US20030138772 SEQ ID NO: 87
AAVrh.14 (AAV42.3a)	399	US20030138772 SEQ ID NO: 32
AAV42.5A	400	US20030138772 SEQ ID NO: 89
AAVrh.17 (AAV42.5a)	401	US20030138772 SEQ ID NO: 34
AAV42.5B	402	US20030138772 SEQ ID NO: 91
AAVrh.18 (AAV42.5b)	403	US20030138772 SEQ ID NO: 29
AAV42.6B	404	US20030138772 SEQ ID NO: 112
AAVrh.19 (AAV42.6b)	405	US20030138772 SEQ ID NO: 38
AAVrh.2	406	US20150159173 SEQ ID NO: 39
AAVrh.2	407	US20150315612 SEQ ID NO: 231
AAVrh.20	408	US20150159173 SEQ ID NO: 1
AAV42.10	409	US20030138772 SEQ ID NO: 106
AAVrh.21 (AAV42.10)	410	US20030138772 SEQ ID NO: 35
AAV42.11	411	US20030138772 SEQ ID NO: 108
AAVrh.22 (AAV42.11)	412	US20030138772 SEQ ID NO: 37
AAV42.12	413	US20030138772 SEQ ID NO: 113
AAVrh.23 (AAV42.12)	414	US20030138772 SEQ ID NO: 58
AAV42.13	415	US20030138772 SEQ ID NO: 86

AAVrh.24 (AAV42.13)	416	US20030138772 SEQ ID NO: 31
AAV42.15	417	US20030138772 SEQ ID NO: 84
AAVrh.25 (AAV42.15)	418	US20030138772 SEQ ID NO: 28
AAVrh.2R	419	US20150159173
AAVrh.31 (AAV223.1)	420	US20030138772 SEQ ID NO: 48
AAVC1	421	US20030138772 SEQ ID NO: 60
AAVrh.32 (AAVC1)	422	US20030138772 SEQ ID NO: 19
AAVrh.32/33	423	US20150159173 SEQ ID NO: 2
AAVrh.33 (AAVC3)	424	US20030138772 SEQ ID NO: 20
AAVC5	425	US20030138772 SEQ ID NO: 62
AAVrh.34 (AAVC5)	426	US20030138772 SEQ ID NO: 21
AAVF1	427	US20030138772 SEQ ID NO: 109
AAVrh.35 (AAVF1)	428	US20030138772 SEQ ID NO: 22
AAVF3	429	US20030138772 SEQ ID NO: 111
AAVrh.36 (AAVF3)	430	US20030138772 SEQ ID NO: 23
AAVrh.37	431	US20030138772 SEQ ID NO: 24
AAVrh.37	432	US20150159173 SEQ ID NO: 40
AAVrh.37	433	US20150315612 SEQ ID NO: 229
AAVrh.37R2	434	US20150159173
AAVrh.38 (AAVLG-4)	435	US20150315612 SEQ ID NO: 7
AAVrh.38 (AAVLG-4)	436	US20150315612 SEQ ID NO: 86
AAVrh.39	437	US20150159173 SEQ ID NO: 20, US20150315612 SEQ ID NO: 13
AAVrh.39	438	US20150159173 SEQ ID NO: 3, US20150159173 SEQ ID NO: 36, US20150315612 SEQ ID NO: 89
AAVrh.40	439	US20150315612 SEQ ID NO: 92
AAVrh.40 (AAVLG-10)	440	US20150315612 SEQ ID No: 14
AAVrh.43 (AAVN721-8)	441	US20150315612 SEQ ID NO: 43, US20150159173 SEQ ID NO: 21
AAVrh.43 (AAVN721-8)	442	US20150315612 SEQ ID NO: 163, US20150159173 SEQ ID NO: 37
AAVrh.44	443	US20150315612 SEQ ID NO: 34
AAVrh.44	444	US20150315612 SEQ ID NO: 111
AAVrh.45	445	US20150315612 SEQ ID NO: 41
AAVrh.45	446	US20150315612 SEQ ID NO: 109
AAVrh.46	447	US20150159173 SEQ ID NO: 22, US20150315612 SEQ ID NO: 19
AAVrh.46	448	US20150159173 SEQ ID NO: 4, US20150315612 SEQ ID NO: 101
AAVrh.47	449	US20150315612 SEQ ID NO: 38
AAVrh.47	450	US20150315612 SEQ ID NO: 118
AAVrh.48	451	US20150159173 SEQ ID NO: 44, US20150315612 SEQ ID NO: 115

AAVrh.48.1	452	US20150159173
AAVrh.48.1.2	453	US20150159173
AAVrh.48.2	454	US20150159173
AAVrh.48 (AAV1-7)	455	US20150315612 SEQ ID NO: 32
AAVrh.49 (AAV1-8)	456	US20150315612 SEQ ID NO: 25
AAVrh.49 (AAV1-8)	457	US20150315612 SEQ ID NO: 103
AAVrh.50 (AAV2-4)	458	US20150315612 SEQ ID NO: 23
AAVrh.50 (AAV2-4)	459	US20150315612 SEQ ID NO: 108
AAVrh.51 (AAV2-5)	460	US20150315612 SEQ ID No: 22
AAVrh.51 (AAV2-5)	461	US20150315612 SEQ ID NO: 104
AAVrh.52 (AAV3-9)	462	US20150315612 SEQ ID NO: 18
AAVrh.52 (AAV3-9)	463	US20150315612 SEQ ID NO: 96
AAVrh.53	464	US20150315612 SEQ ID NO: 97
AAVrh.53 (AAV3-11)	465	US20150315612 SEQ ID NO: 17
AAVrh.53 (AAV3-11)	466	US20150315612 SEQ ID NO: 186
AAVrh.54	467	US20150315612 SEQ ID NO: 40
AAVrh.54	468	US20150159173 SEQ ID NO: 49, US20150315612 SEQ ID NO: 116
AAVrh.55	469	US20150315612 SEQ ID NO: 37
AAVrh.55 (AAV4-19)	470	US20150315612 SEQ ID NO: 117
AAVrh.56	471	US20150315612 SEQ ID NO: 54
AAVrh.56	472	US20150315612 SEQ ID NO: 152
AAVrh.57	473	US20150315612 SEQ ID NO: 26
AAVrh.57	474	US20150315612 SEQ ID NO: 105
AAVrh.58	475	US20150315612 SEQ ID NO: 27
AAVrh.58	476	US20150159173 SEQ ID NO: 48, US20150315612 SEQ ID NO: 106
AAVrh.58	477	US20150315612 SEQ ID NO: 232
AAVrh.59	478	US20150315612 SEQ ID NO: 42
AAVrh.59	479	US20150315612 SEQ ID NO: 110
AAVrh.60	480	US20150315612 SEQ ID NO: 31
AAVrh.60	481	US20150315612 SEQ ID NO: 120
AAVrh.61	482	US20150315612 SEQ ID NO: 107
AAVrh.61 (AAV2-3)	483	US20150315612 SEQ ID NO: 21
AAVrh.62 (AAV2-15)	484	US20150315612 SEQ ID No: 33
AAVrh.62 (AAV2-15)	485	US20150315612 SEQ ID NO: 114
AAVrh.64	486	US20150315612 SEQ ID No: 15
AAVrh.64	487	US20150159173 SEQ ID NO: 43, US20150315612 SEQ ID NO: 99

AAVrh.64	488	US20150315612 SEQ ID NO: 233
AAVRh.64R 1	489	US20150159173
AAVRh.64R 2	490	US20150159173
AAVrh.65	491	US20150315612 SEQ ID NO: 35
AAVrh.65	492	US20150315612 SEQ ID NO: 112
AAVrh.67	493	US20150315612 SEQ ID NO: 36
AAVrh.67	494	US20150315612 SEQ ID NO: 230
AAVrh.67	495	US20150159173 SEQ ID NO: 47, US20150315612 SEQ ID NO: 113
AAVrh.68	496	US20150315612 SEQ ID NO: 16
AAVrh.68	497	US20150315612 SEQ ID NO: 100
AAVrh.69	498	US20150315612 SEQ ID NO: 39
AAVrh.69	499	US20150315612 SEQ ID NO: 119
AAVrh.70	500	US20150315612 SEQ ID NO: 20
AAVrh.70	501	US20150315612 SEQ ID NO: 98
AAVrh.71	502	US20150315612 SEQ ID NO: 162
AAVrh.72	503	US20150315612 SEQ ID NO: 9
AAVrh.73	504	US20150159173 SEQ ID NO: 5
AAVrh.74	505	US20150159173 SEQ ID NO: 6
AAVrh.8	506	US20150159173 SEQ ID NO: 41
AAVrh.8	507	US20150315612 SEQ ID NO: 235
AAVrh.8R	508	US20150159173, WO2015168666 SEQ ID NO: 9
AAVrh.8R A586R mutant	509	WO2015168666 SEQ ID NO: 10
AAVrh.8R R533A mutant	510	WO2015168666 SEQ ID NO: 11
BAAV (bovine AAV)	511	US9193769 SEQ ID NO: 8
BAAV (bovine AAV)	512	US9193769 SEQ ID NO: 10
BAAV (bovine AAV)	513	US9193769 SEQ ID NO: 4
BAAV (bovine AAV)	514	US9193769 SEQ ID NO: 2
BAAV (bovine AAV)	515	US9193769 SEQ ID NO: 6
BAAV (bovine AAV)	516	US9193769 SEQ ID NO: 1
BAAV (bovine AAV)	517	US9193769 SEQ ID NO: 5
BAAV (bovine AAV)	518	US9193769 SEQ ID NO: 3

BAAV (bovine AAV)	519	US9193769 SEQ ID NO: 11
BAAV (bovine AAV)	520	US7427396 SEQ ID NO: 5
BAAV (bovine AAV)	521	US7427396 SEQ ID NO: 6
BAAV (bovine AAV)	522	US9193769 SEQ ID NO: 7
BAAV (bovine AAV)	523	US9193769 SEQ ID NO: 9
BNP61 AAV	524	US20150238550 SEQ ID NO: 1
BNP61 AAV	525	US20150238550 SEQ ID NO: 2
BNP62 AAV	526	US20150238550 SEQ ID NO: 3
BNP63 AAV	527	US20150238550 SEQ ID NO: 4
caprine AAV	528	US7427396 SEQ ID NO: 3
caprine AAV	529	US7427396 SEQ ID NO: 4
true type AAV (ttAAV)	530	WO2015121501 SEQ ID NO: 2
AAAV (Avian AAV)	531	US9238800 SEQ ID NO: 12
AAAV (Avian AAV)	532	US9238800 SEQ ID NO: 2
AAAV (Avian AAV)	533	US9238800 SEQ ID NO: 6
AAAV (Avian AAV)	534	US9238800 SEQ ID NO: 4
AAAV (Avian AAV)	535	US9238800 SEQ ID NO: 8
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AAAV (Avian AAV)	541	US9238800 SEQ ID NO: 3
AAAV (Avian AAV)	542	US9238800 SEQ ID NO: 7
AAAV (Avian AAV)	543	US9238800 SEQ ID NO: 11
AAAV (Avian AAV)	544	US9238800 SEQ ID NO: 13
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AAV Shuffle 100-1	546	US20160017295 SEQ ID NO: 23

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AAV Shuffle 10-6	557	US20160017295 SEQ ID NO: 27
AAV Shuffle 10-8	558	US20160017295 SEQ ID NO: 36
AAV Shuffle 10-8	559	US201600 17295 SEQ ID NO: 28
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AAV SM 100-3	563	US201600 17295 SEQ ID NO: 32
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AAV SM 10-8	568	US20160017295 SEQ ID NO: 39
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AAVF4/HSC 4	573	WO20 16049230 SEQ ID NO: 23
AAVF5/HSC 5	574	WO20 16049230 SEQ ID NO: 25
AAVF6/HSC 6	575	WO20 16049230 SEQ ID NO: 24

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AAV CLv-R7	626	US8734809 SEQ ID NO: 36
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AAV CLv-R9	628	US8734809 SEQ ID NO: 38
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AAV CSp-11	639	US8734809 SEQ ID NO: 47
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AAV CSp-4	642	US8734809 SEQ ID NO: 50

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AAV CLv-8	664	US8734809 SEQ ID NO: 72
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AAV CKd-B3	667	US8734809 SEQ ID NO: 75
AAV CKd-B4	668	US8734809 SEQ ID NO: 76
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AAV CKd-H5	677	US8734809 SEQ ID NO: 85
AAV CKd-H6	678	US8734809 SEQ ID NO: 77
AAV CHt-1	679	US8734809 SEQ ID NO: 86

AAV CLvl-1	680	US8734809 SEQ ID NO: 171
AAV CLvl-2	681	US8734809 SEQ ID NO: 172
AAV CLvl-3	682	US8734809 SEQ ID NO: 173
AAV CLvl-4	683	US8734809 SEQ ID NO: 174
AAV Civ 1-7	684	US8734809 SEQ ID NO: 175
AAV Civ 1-8	685	US8734809 SEQ ID NO: 176
AAV Clvl-9	686	US8734809 SEQ ID NO: 177
AAV Clv1-10	687	US8734809 SEQ ID NO: 178
AAV.VR-355	688	US8734809 SEQ ID NO: 181
AAV .3ЭН.48R 3	689	US8734809 SEQ ID NO: 183
AAV CBs-E1	690	US8734809 SEQ ID NO: 87
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AAV CBr-E3	692	US8734809 SEQ ID NO: 89
AAV CBr-E4	693	US8734809 SEQ ID NO: 90
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AAV CLv-D4	702	US8734809 SEQ ID NO: 99
AAV CLv-D5	703	US8734809 SEQ ID NO: 100
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AAV CLv-D8	706	US8734809 SEQ ID NO: 103
AAV CLv-El	*ioi*	US8734809 SEQ ID NO: 87
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AAV CLv-R5	712	US8734809 SEQ ID NO: 108
AAV CLv-R6	713	US8734809 SEQ ID NO: 109
AAV CLv-R7	714	US8734809 SEQ ID NO: 110

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AAV CKd-8	744	US8734809 SEQ ID NO: 138
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AAV CKcl-B6	758	US8734809 SEQ ID NO: 152
AAV CKd-B7	759	US8734809 SEQ ID NO: 153
AAV CKd-B8	760	US8734809 SEQ ID NO: 154
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AAV CKd-N4	780	WO2016065001 SEQ ID NO: 13
AAV CKd-N9	781	WO2016065001 SEQ ID NO: 14
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AAV CLv-LS	783	WO2016065001 SEQ ID NO: 16
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AAV CLv-K3	786	WO2016065001 SEQ ID NO: 19
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AAV CSp-8.9	812	WO2016065001 SEQ ID NO: 45
AAV CBr-B7.3	813	WO2016065001 SEQ ID NO: 46
AAV CBr-B7.4	814	WO2016065001 SEQ ID NO: 47
AAV3B	815	WO2016065001 SEQ ID NO: 48
AAV4	816	WO2016065001 SEQ ID NO: 49
AAV5	817	WO2016065001 SEQ ID NO: 50
AAV CHt-P2	818	WO2016065001 SEQ ID NO: 51
AAV CHt-P5	819	WO2016065001 SEQ ID NO: 52
AAV CHt-P9	820	WO2016065001 SEQ ID NO: 53

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AAV **CBr-**7.5	825	**WO20 16065001** SEQ ID **NO:** 58
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AAV **CBr-**7.8	827	**WO20 16065001** SEQ ID **NO:** 60
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AAV **CKd-N3**	829	**WO20 16065001** SEQ ID **NO:** 62
AAV CKd-**N4**	830	**WO20 16065001** SEQ ID NO: 63
AAV **CKd-N9**	831	**WO20 16065001** SEQ ID **NO:** 64
AAV CLv-L4	832	WO2016065001 SEQ ID NO: 65
AAV **CLv-L5**	833	**WO2016065001** SEQ ID NO: 66
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AAV **CLv-**K6	837	**WO20 16065001** SEQ ID NO: 70
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AAV **CLv-**M7	843	**WO20 16065001** SEQ ID NO: 76
AAV **CLv-M8**	844	**WO20 16065001** SEQ ID **NO:** 77
AAV **CLv-**M9	845	**WO20 16065001** SEQ ID NO: 78
AAV **CHt-Pl**	846	**WO20 16065001** SEQ ID **NO**: 79
AAV CH1-P6	847	WO20 16065001 SEQ ID NO: 80
AAV CHt-P8	848	**WO20 1606500** 1 SEQ ID NO: **81**
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AAV **CHt-6.7**	853	**WO20 16065001** SEQ ID **NO:** 86
AAV CHt-6.8	854	**WO20 16065001** SEQ ID NO: 87
AAV CSp-8.10	855	**WO20 16065001** SEQ ID NO: 88
AAV **CSp-**8.2	856	**WO20 16065001** SEQ ID **NO:** 89
AAV CSp-8.4	**857**	**WO20 16065001** SEQ ID NO: 90
AAV **CSp-**8.5	858	**WO20 16065001** SEQ ID **NO:** 91
AAV CSp-8.6	859	**WO20 16065001** SEQ ID NO: 92
AAV **CSp-8.7**	860	**WO20 16065001** SEQ ID **NO:** 93
AAV CSp-8.8	861	**WO20 16065001** SEQ ID NO: 94
AAV **CSp-**8.9	862	**WO20 16065001** SEQ ID **NO:** 95
AAV CBr-B7.3	863	WO20 16065001 SEQ ID NO: 96
AAV **CBr-B7.4**	864	**WO20 16065001** SEQ ID NO: 97
AAV3B	865	**WO20 16065001** SEQ ID NO: 98
AAV4	866	**WO20 1606500** 1 SEQ ID NO: 99
AAV5	867	**WO20 16065001 SEQ** ID **NO**: **100**
AAVPHP.B or G2B-26	868	**WO20** 15038958 SEQ ID NO: 8 and **13; GeiiBaiikALU85 156. 1**
AAVPHP.B	869	**WO2015038958** SEQ ID NO: 9
AAVG2B-13	870	**WO2015038958** SEQ ID **NO**: 12
AAVTHl. l-32	871	**WO20 15038958** SEQ ID NO: 14
AAVTH1. 1-35	872	**WO2015038958** SEQ ID NO: 15

[0089] Each of the patents, applications and/or publications listed in Table 1 are herein incorporated by reference in their entirely.

[0090] In one embodiment, the AAV serotype may be, or may have a sequence as described in International Patent Publication WO2015038958, the contents of which are herein incorporated by reference in their entirety, such as, but not limited to, AAV9 (SEQ ID NO: 2 and 11 of WO2015038958 or SEQ ID NO: 127 and 126 respectively herein), PHP.B (SEQ ID NO: 8 and 9 of WO2015038958, herein SEQ ID NO; 868 and 869), G2B-13 (SEQ ID NO: 12 of WO2015038958, herein SEQ ID NO: 870), G2B-26 (SEQ ID NO: 13 of WO2015038958, herem SEQ ID NO; 868 and 869), TH1.1-32 (SEQ ID NO: 14 of WO2015038958, herein SEQ ID NO: 871), TH1.1-35 (SEQ ID NO: 15 of WO2015038958, herem SEQ ID NO: 872) or variants thereof. Further, any of the targeting peptides or amino acid inserts described in WO2015038958, may be inserted into any parent AAV serotype, such as, but not limited to.

AAV9 (SEQ ID NO: 126 for the DNA sequence and SEQ ID NO: 127 for the amino acid sequence). In one embodiment, the amino acid insert is inserted between amino acids 586-592 of the parent AAV (e.g., A.AV9). In another embodiment, the amino acid insert is inserted between amino acids 588-589 of the parent AAV sequence. The amino acid insert may be, but is not limited to, any of the following ammo acid sequences, TLAVPFK (SEQ ID NO: 1 of WO2015038958; herein SEQ ID NO; 873), KFPVALT (SEQ ID NO: 3 of WO2015038958; herein SEQ ID NO: 874), LAVPFK (SEQ ID NO: 31 of WG2015038958; herein SEQ ID NO: 875), AVPFK (SEQ ID NO: 32 of WO201 5038958; herein SEQ ID NO: 876). VPFK (SEQ ID NO: 33 of WO2015038958; herein SEQ ID NO: 877), TLAVPF (SEQ ID NO: 34 of WO2015038958; herein SEQ TO NO: 878), TLAVP (SEQ TO NO: 35 of WO2015038958; herein SEQ ID NO: 879), TLAV (SEQ ID NO: 36 of WO2015038958; herein SEQ ID NO: 880), SVSKPFL (SEQ ID NO: 28 of WO201 5038958; herein SEQ ID NO: 881), FTLTTPK (SEQ ID NO: 29 of WO2015038958; herein SEQ ID NO: 8821 MNATKNV (SEQ ID NO: 30 of WO2015038958; herein SEQ ID NO: 883), QSSQTPR (SEQ ID NO: 54 of WO201 5038958; herein SEQ ID NO: 884), ILGTGTS (SEQ ID NO: 55 of WO2015038958; herein SEQ ID NO: 885), TRTNPEA (SEQ ID NO: 56 of WO20I5038958; herein SEQ ID NO: 886), NGGTSSS (SEQ ID NO: 58 of WO2015038958; herein SEQ ID NO; 887), or YTLSQGW (SEQ ID NO: 60 of WO2015038958; herein SEQ ID NO: 888). Non-limiting examples of nucleotide sequences that may encode the amino acid inserts include the following, AAGTTTCX'TGTGGCGTTGACT (for SEQ ID NO: 3 of WO2015038958; herein SEQ ID NO: 889), ACTTTGGCGGTGCCTTTTAAG (SEQ ID NO: 24 and 49 of WO2015038958; herein SEQ ID NO: 890), AGTGTGAGTAAGCC TTTTTTG (SEQ ID NO: 25 of WO2015038958, herein SEQ ID NO: 891), TTTACGTTGACGACGCCTAAG (SEQ ID NO: 26 of WO2015038958; herein SEQ ID NO: 892), ATGA ATGCTACGAAGA ATGTG (SEQ ID NO: 27 of WO2015038958; herein SEQ ID NO: 893), CAGTC GTC GCAGAC GCCTAGG (SEQ ID NO: 48 of WO2015038958; herein SEQ ID NO: 894), ATTCTGGGGACTGGTACTTCG (SEQ ID NO: 50 and 52 of WO20I5038958; herein SEQ ID NO: 895), ACGCGGACTAATCCTGAGGCT (SEQ ID NO: 51 of WO2015038958; herein SEQ ID NO: 896), AATGGGGGGACTAGTAGTTCT (SEQ ID NO: 53 of WO201 5038958; herein SEQ ID NO: 897), or TATACTTTGTCGCAGGGTTGG (SEQ ID NO: 59 of WO2015038958; herein SEQ ID NO: 898).

Viral Genome Component: Inverted Terminal Repeats (ITRs)

[0091] The AAV particles of the present invention comprise a viral genome with at least one ITR region and a payload region. In one embodiment, the viral genome has two ITRs. 'These

two ITRs flank the payload region at the 5' and 3' ends. The ITRs function as origins of replication comprising recognition sites for replication. ITRs comprise sequence regions which can be complementary and symmetrically arranged. ITRs incorporated into viral genomes of the invention may be comprised of naturally occurring polynucleotide sequences or recombinant!}, derived polynucleotide sequences.

[0092] The ITRs may be derived from the same serotype as the capsid, selected from any of the serotypes listed in Table 1, or a derivative thereof. The ITR may be of a different serotype than the capsid. In one embodiment, the AAV particle has more than one ITR. In a non-limiting example, the AAV particle bas a viral genome comprising two ITRs. In one embodiment, the TTRs are of the same serotype as one another. In another embodiment, the ITRs are of different serotypes. Non-limiting examples inciude zero, one or both of the ITRs having the same serotype as the capsid. In one embodiment both ITRs of the viral genome of the AAV particle are AAV2 ITRs.

[0093] Independently, each ITR may be about 100 to about 150 nucleotides in length. An ITR may be about 100-105 nucleotides in length, 106-110 nucleotides in length, 111-115 nucleotides in length, 116-120 nucleotides in length, 121-125 nucleotides in length, 126-130 nucleotides in length, 131-135 nucleotides in length, 136-140 nucleotides in length, 141-145 nucleotides in length or 146-150 nucleotides in length. In one embodiment, the ITRs are 140-142 nucleotides in length. Non-limiting examples of ITR length are 102, 140, 141, 142, 145 nucleotides in length, and those having at least 95% identity thereto.

Viral Genome Component: Promoters

[0094] In one embodiment, the payload region of the viral genome comprises at least one element to enhance the transgene target specificity and expression (See e.g., Powell et al. Viral Expression Cassette Elements to Enhance Transgene Target Specificity and Expression in Gene Therapy, 2015; the contents of which are herein incorporated by reference in its entirety). Non-limiting examples of elements to enhance the transgene target specificity and expression include promoters, endogenous miRNAs, post-transcriptional regulator) elements (PREs), polyadenylation (PolyA) signal sequences and upstream enhancers (USEs), CMV enhancers and inirons.

[0095] A person skilled in the art may recognize that expression of the polypeptides of the invention in a target cell may require a specific promoter, including but not limited to, a promoter that is species specific, inducible, tissue-specific, or cell cycle-specific (Parr et al, *Nat. Med.* 3:1145-9 (1997); the contents of which are herein incorporated by reference in their entirety).

[0096] In one embodiment, the promoter is deemed to be efficient when it drives expression of the polypeptide(s) encoded in the payload region of the viral genome of the AAV particle.

[0097] In one embodiment, the promoter is a promoter deemed to be efficient when it drives expression in the ceil being targeted.

[0098] In one embodiment, the promoter drives expression of the polypeptides of the invention (e.g., a functional antibody) for a period of time in targeted tissues. Expression driven by a promoter may be for a period of I hour, 2, hours, 3 hours, 4 hours, 5 hours, 6 hours, 7 hours, 8 hours, 9 hours, 10 hours, 11 hours, 12 hours, 13 hours, 14 hours, 15 hours, 16 hours, 17 hours, 18 hours, 19 hours, 20 hours, 21 hours, 22 hours, 23 hours, 1 day, 2 days, 3 days, 4 days, 5 days, 6 days, 1 week. 8 days, 9 days, 10 days, 11 days, 12 days, 13 days, 2 weeks, 15 days, 16 days, 17 days, 18 days, 19 days, 20 days, 3 weeks, 22 days, 23 days, 24 days, 25 days, 26 days, 27 days, 28 days, 29 days, 30 days, 31 days, 1 month, 2 months, 3 months, 4 months, 5 months, 6 months, 7 months, 8 months, 9 months, 10 months, 11 months, 1 year, 13 months, 14 months, 15 months, 16 months, 17 months, 18 months, 19 months, 20 months, 21 months, 22. months, 23 months, 2 years, 3 years, 4 years, 5 years, 6 years, 7 years, 8 years, 9 years, 10 years or more than 10 years. Expression may be for 1-5 hours, 1-12 hours, 1-2 days, 1-5 days, 1-2 weeks, 1-3 weeks, 1-4 weeks, 1-2 months, 1-4 months, 1-6 months, 2-6 months. 3-6 months, 3-9 months, 4-8 months, 6-12 months, 1-2 years, 1-5 years, 2-5 years, 3-6 years, 3-8 years, 4-8 years or 5-10 years.

[0099] In one embodiment, the promoter drives expression of the polypeptides of the invention (e.g., a functional antibody) for at least 1 month, 2 months, 3 months, 4 months, 5 months, 6 months, 7 months, 8 months, 9 months, 10 months, 11 months, 1 year, 2 years, 3 years 4 years, 5 years, 6 years, 7 years, 8 years, 9 years, 10 years, 11 years, 12 years, 13 years, 14 years, 15 years, 16 years, 17 years, 18 years, 19 years, 20 years, 21 years, 22 years, 23 years, 24 years, 25 years, 26 years, 27 years, 28 years, 29 years. 30 years, 31 years, 32 years, 33 years, 34 years, 35 years, 36 years, 37 years, 38 years, 39 years, 40 years, 41 years, 42 years, 43 years, 44 years, 45 years, 46 years, 47 years, 48 years, 49 years, 50 years, 55 years, 60 years, 65 years, or more than 65 years.

[00100] Promoters may be naturally occurring or non-naturally occurring. Non-limiting examples of promoters include viral promoters, plant promoters and mammalian promoters. In some embodiments, the promoters may be human promoters. In some embodiments, the promoter may be truncated.

[00101] Promoters which drive or promote expression in most tissues include, but are not limited to, human elongation factor la-subunit (EFla), cytomegalovirus (CMV) immediate-early enhancer and/or promoter, chicken β-actin (CBA) and its derivative CAG, β glucuronidase

(GUSB), or uhiquitm C (UBC). Tissue-specific expression elements can be used to restrict expression to certain cell types such as, but not limited to, muscle specific promoters, B cell promoters, monocyte promoters, leukocyte promoters, macrophage promoters, pancreatic acinar cell promoters, endothelial cell promoters, lung tissue promoters, astrocyte promoters, or nervous system promoters which can be used to restrict expression to neurons, astrocytes, or oligodendrocytes.

[00102] Non-limiting examples of muscle-specific promoters include mammalian muscle creatine kinase (MCK) promoter, mammalian desmin (DES) promoter, mammalian troponin I (TNNI2) promoter, and mammalian skeletal alpha-actin (ASKA) promoter (see. e.g. U.S. Patent Publication US 2011/0212529, the contents of which are herein incorporated by reference in their entirety)

[00103] Non-limiting examples of tissue-specific expression elements for neurons include neuron-specific enolase (NSE), platelet-derived growth factor (PDGF), platelet-derived growth factor B-chain (PDGF-β), synapsin (Syn), methyl-CpG binding protein 2. (MeCP2), €a2Vcalmodulin-dependent protein kinase 11 (CaMKII), metabotropic giutamate receptor 2 (mGluR2), neurofilament light (NFL) or heavy (NFH). β-globin minigene ηβ2, preproenkephalin (PPE), enkephalin (Enk) and excitatory amino acid transporter 2 (EAAT2) promoters. Non-limiting examples of tissue-specific expression elements for astrocytes include glial fibrillary acidic protein (GFAP) and EAAT2 promoters. A non-iimiting example of a tissue-specific expression element for oligodendrocytes includes the myelin basic protein (MBP) promoter.

|00104] In one embodiment, the promoter may be less than 1 kb. The promoter may have a length of 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, 370. 380, 390, 400, 410. 420, 430, 440, 450. 460, 470, 480, 490. 500, 510, 520, 530, 540. 550, 560. 570, 580, 590, 600. 610, 620, 630, 640, 650, 660, 670, 680, 690, 700, 710, 720, 730. 740, 750, 760, 770, 780, 790, 800 or more than 800 nucleotides. The promoter may have a length between 200-300, 200-400, 200-500, 200-600, 200-700, 200-800, 300-400, 300-500, 300-600, 300-700, 300-800, 400-500, 400-600, 400-700, 400-800. 500-600, .500-700. 500-800, 600-700. 600-800 or 700-800.

[00105] In one embodiment, the promoter may be a combination of two or more components of the same or different starting or parental promoters such as. but not limited to, CMV and CBA. Each component may have a length of 200, 210, 220, 230, 240, 250. 260, 270, 280, 290. 300, 310, 320, 330, 340, 350, 360, 370, 380, 381, 382, 383, 384, 385, 386, 387, 388, 389, 390, 400, 410, 420, 430, 440, 450. 460, 470, 480, 490, 500, 510, 520, 530, 540, 550, 560, 570, 580, 590, 600, 610. 620, 630, 640, 650. 660, 670, 680, 690, 700, 710, 720, 730, 740, 750, 760, 770,

780, 790, 800 or more than 800. Each component may have a length between 200-300, 200-400, 200-500, 200-600, 200-700, 200-800, 300-400, 300-500, 300-600, 300-700, 300-800, 400-500, 400-600, 400-700, 400-800, 500-600, 500-700, 500-800, 600-700, 600-800 or 700-800. In one embodiment, the promoter is a combination of a 382 nucleotide CMV-enhancer sequence and a 260 nucleotide CBA-promoter sequence.

[00106] In one embodiment, the viral genome comprises a ubiquitous promoter. Non-limiting examples of ubiquitous promoters include CMV, CBA (including derivatives CAG, CBh, etc.), EF-1α, PGK, UBC, GUSB ihGBp), and UCOE (promoter of HNRPA2B1-CBX3).

Yu et al. (Molecular Pain 2011, 7:63, the contents of which are herein incorporated by reference in their entirely) evaluated the expression of eGFP under the CAG, EFIα, PGK and UBC promoters in rat DRG cells and primary DRG cells using ientiviral vectors and found that TJBC showed weaker expression than the other 3 promoters and only 10-12% glial expression was seen for all promoters. Soderblom et al. (E. Neuro 2015; the contents of which are herein incorporated by reference in its entirety) evaluated the expression of eGFP in AAV8 with CMV and UBC promoters and AAV2 with the CMV promoter after injection in the motor cortex. Intranasal administration of a plasmid containing a UBC or EFIa promoter showed a sustained airway expression greater than the expression with the CMV promoter (See e.g., Gill et al, Gene Therapy 2.001, Vol. 8, 1539-1546; the contents of which are herein incorporated by reference in their entirety). Husain et al. (Gene Therapy 2009; the contents of which are herein incorporated by reference in its entirety) evaluated an HβH construct with abGlJSB promoter, aHSV-lLAT promoter and an NSE promoter and found that the HβH construct showed weaker expression than NSE in mouse brain. Passini and Wolfe (J. Virol. 2001, 12382-12392, the contents of which are herein incorporated by reference in its entirety) evaluated the long term effects of the HβH vector following an intraventricular injection in neonatal mice and found that there was sustained expression for at least 1 year. Low expression in all brain regions was found by Xu et al. (Gene Therapy 2001, 8, 1323-1332; the contents of which are herein incorporated by reference in their entirety) when NFL and NFH promoters were used as compared to the CMV-iacZ, CMV-luc, EF, GFAP, hENK, iiAChR, PPE, PPE ÷ wpre, NSE (0.3 kb), NSE (1.8 kb) and NSE (1.8 kb + wpre). Xu et al. found that the promoter activity in descending order was NSE (1.8 kb), EF, NSE (0.3 kb), GFAP, CMV, hENK, PPE, NFL and NFH. NFL is a 650 nucleotide promoter and NFH is a 920 nucleotide promoter which are both absent in the liver but NFH is abundant in the sensory- proprioceptive neurons, brain and spinal cord and NFH is present in the heart. Scn8a is a 470 nucleotide promoter which expresses throughout the DRG, spinal cord and brain with particularly high expression seen in the hippocampal neurons and cerebellar Purkinje

cells, cortex, thalamus and hypothalamus (See e.g., Drews et al. *Identification of evolutionary conserved, functional noncoding elements in (he promoter region of the sodium channel gene SCNSA.* Mamm Genome (2007) 18:723-731 ; and Raymond et al. *Expression of Alternatively Spliced Sodium Channel a-subunit genes,* Journal of Biological Chemistry (2004) 279(44} 46234-46241; the contents of each of which are herein incorporated by reference in their entireties).

[00107] Any of promoters taught by the aforementioned Yu, Soderblom, Gill, Husam, Passmi, Xu, Drews or Raymond may be used in the present inventions.

[00108] In one embodiment, the promoter is not cell specific.

[00109] In one embodiment, the promoter is a ubiquitin c (UBC) promoter. The UBC promoter may have a size of 300-350 nucleotides. As a non-limiting example, the UBC promoter is 332 nucleotides.

[00110] In one embodiment, the promoter is a β-glucuromdase (GUSB) promoter. The GUSB promoter may have a size of 350-400 nucleotides. As a non-limiting example, the GUSB promoter is 378 nucleotides.

[00111] In one embodiment, the promoter is a neurofilament light (NFL) promoter. The NFL promoter may have a size of 600-700 nucleotides. As a non-limiting example, the NFL promoter is 650 nucleotides.

[00112] In one embodiment, the promoter is a neurofilament heavy (NFF1) promoter. The NFH promoter may have a size of 900-950 nucleotides. As a non-limiting example, the NFH promoter is 920 nucleotides.

[00113] In one embodiment, the promoter is a scn8a promoter. The scnSa promoter may have a size of 450-500 nucleotides. As anon-limiting example, the scnSa promoter is 470 nucleotides.

[00114] In one embodiment, the promoter is a phosphoglycerate kinase 1 (PGK) promoter.

[00115] In one embodiment, the promoter is a chicken β-actin (CBA) promoter.

[00116] In one embodiment, the promoter is a cytomegalovirus (CMV) promoter.

[00117] In one embodiment, the promoter is a liver or a skeletal muscle promoier. Non-hnntiiig examples of liver promoters include human a-1-antitrypsin (hAAT) and thyroxine binding globulin (TBG). Non-limiting examples of skeletal muscle promoters include Desmin, MCK or synthetic C5-12.

[00118] In one embodiment, the promoter is a RNA pol III promoter. As a non-limiting example, the RNA pol iII promoter is U6. As a non-limiting example, the RNA pol III promoter is H1.

[00119] In one embodiment, the viral genome composes two promoters. As a non-limiting example, the promoters are an EF1α promoter and a CMV promoter.

[00120] In one embodiment, the viral genome comprises an enhancer element, a promoter and/or a 5'UTR intron. The enhancer element, also referred to herein as an "enhancer," may be, but is not limited to, a CMV enhancer, the promoter may be, but is not limited to, a CMV, CBA, UBC, GUSB, NSE, Synapsin, MeCP2, and GFAP promoter and the 5'UTR/intron may be, but is not limited to, SV40, and CBA-MVM. As a non-limiting example, the enhancer, promoter and/or intron used in combination may be: (1) CMV enhancer, CMV promoter, SV40 5'UTR intron, (2) CMV enhancer, CBA promoter, SV 40 5'UTR intron, (3) CMV enhancer, CBA promoter, CBA-MVM 5'UTR intron; (4) UBC promoter; (5) GUSB promoter; (6) NSE promoter; (7) Synapsin promoter; (8) MeCP2 promoter and (9) GFAP promoter.

[00121] In one embodiment, the viral genome comprises an engineered promoter.

[00122] In another embodiment, the viral genome comprises a promoter from a naturally expressed protein.

Viral Genome Component: Untranslated Regions (IJTRs)

[00011] By definition, wild type untranslated regions (UTRs) of a gene are transcribed but not translated. Generally, the 5' UTR starts at the transcription start site and ends at the start codon and the 3' UTR starts immediately following the stop codon and continues until the termination signal for transcription.

[00012] Features typically found in abundantly expressed genes of specific target organs may be engineered into UTRs to enhance the stability and protein production. As a non-limiting example, a 5' UTR from mRNA normally expressed in the liver (e.g., albumin, serum amyloid A, Apoiipoprotein A-'B/E, transferrin, alpha fetoprotein, erythropoietin, or Factor VIII) may be used in the viral genomes of the AAV particles of the invention to enhance expression in hepatic cell lines or liver.

[00013] While not wishing to be bound by theory, wild-type 5' untranslated regions (UTRs) include features which play roles in translation initiation. Kozak sequences, which are commonly known to be involved in the process by which the ribosome initiates translation of many genes, are usually included in 5' UTRs. Kozak sequences have the consensus CCR(A"G)CCAUGG, where R is a purine (adenine or guanine) three bases upstream of the start codon (ATG), which is followed by another 'G.

[00014] In one embodiment, the 5'UTR in the viral genome includes a Kozak sequence.

[00015] In one embodiment, the 5'UTR in the viral genome does not include a Kozak sequence.

[00016] **While** not wishing to be bound **by theory,** wild-type 3' UTRs **are** known **to** have stretches of Adenosines and Uridines embedded therein. These AU rich signatures are particularly prevalent **in** genes with high rates of turnover. Based on their sequence features and functional properties, the AU rich elements (AREs) can be separated into three classes (Chen et **al,** 1995, the contents **of which** are herein incorporated by reference **in** its entirety): Class I AREs, such as, but not limited **to, c-Myc** and **MyoD,** contain several dispersed copies **of** an AUUUA motif within U-nch regions. Class 11 AREs, such as, but not limited to, **GM-CSF** and TNF-a, possess two or more overlapping UUAUUUA(U/A)(U/A) **nonamers.** Class III **ARES,** such as, but not limited **to, c-Jun** and **Myogenin,** are less well defined. These **U** rich regions **do** not contain an AUUUA motif. Most proteins binding **to** the AREs are *know*n to destabilize the messenger, whereas members of the ELAV family, most notably **HuR,** have been documented to increase the stability of **mRNA.** HuR binds to AREs **of all** the three classes. Engineering **the** HuR specific binding sites **into** the 3' **UTR of** nucleic acid molecules **will** lead **to** HuR binding and thus, stabilization of the message *in vivo.*

[00017] Introduction, removal or modification of 3' UTR **AU** rich elements (AREs) can be used **to** modulate the stability of polynucleotides. When engineering specific polynucleotides, e.g., payload regions of viral genomes, one or more copies of an ARE can be introduced to make polynucleotides less stable and thereby curtail translation and decrease production of the resultant protein. Likewise, AREs can **be** identified and removed or mutated to increase the intracellular stability and thus increase translation and production of the resultant protein.

[00018] In one embodiment, the 3' UTR of the viral genome may include an **oligo(dT)** sequence **for templated** addition of a poiy-A **tail.**

[00019] **In** one embodiment, the viral genome may include at least one **miRNA** seed, binding site or **full** sequence. **microRNAs** (or miRNA **or miR)** are 19-25 nucleotide **noncoding RNAs** that bind to the sites of nucleic acid targets and down-regulate gene expression either **by** reducing nucleic acid molecule stability or by inhibiting translation. **A microRNA** sequence comprises a "seed" region, i.e., a sequence in the region of positions 2-8 **of the** mature **microRNA,** which sequence has perfect Watson-Crick complementarity to the miRNA target sequence of the nucleic acid.

[00020] **In** one embodiment, the viral genome may be engineered **to** include, alter or remove **at** least one miRNA binding site, sequence or seed region.

[00021] Any UTR from any gene known **in** the art may be incorporated into the viral genome of the AAV particle. These UTRs, or portions thereof, may be placed **in** the same orientation as in the gene from which they were selected **or** they may be altered in orientation **or** location. **In**

one embodiment, the UTR used in the viral genome of the AAV particle may be inverted, shortened, lengthened, made with one or more other 5' UTRs or 3' UTRs known in the art. As used herein, the term "'altered" as it relates to a UTR, means that the UTR has been changed in some way in relation to a reference sequence. For example, a 3' or 5' UTR may be altered relative to a wild type or native UTR by the change in orientation or location as taught above or may be altered by the inclusion of additional nucleotides, deletion of nucleotides, swapping or transposition of nucleotides.

[00022] In one embodiment, the viral genome of the AAV particle comprises at least one artificial UTRs which is not a variant of a wild type UTR.

[ΘΘ023] In one embodiment, the viral genome of the AAV particle comprises UTRs which have been selected from a family of transcripts whose proteins share a common function, structure, feature or property.

Viral Genome Component: Polyadenyiation Sequence

[00123] In one embodiment, the viral genome of the AAV particles of the present invention comprise at least one polyadenylation sequence. Hie viral genome of the AAV particle may comprise a polyadenyiation sequence between the 3` end of the payload coding sequence and the 5' end of the 3`ITR.

[00124] In one embodiment, the polyadenyiation sequence or "polyA sequence" may range from absent to about 500 nucleotides in length. The polyadenyiation sequence may be, but is not limited to, I, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74, 75, 76, 77, 78, 79. 80, 81, 82, 83, 84. 85. 86, 87, 88, 89, 90. 91, 92, 93, 94, 95. 96. 97, 98, 99, 100, 101. 102, 103. 104, 105, 106, 107. 108, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118, 119, 120. 121, 122, 123, 124, 125, 126, 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141, 142. 143, 144, 145, 146. 147, 148, 149, 150, 151, 152, 153, 154, 155, 156, 157, 158, 159. 160, 161, 162. 163, 164, 165, 166. 167, 168, 169, 170. 171, 172, 173, 174. 175, 176, 177, 178, 179, 180, 181, 182, 183, 184, 185, 186, 187, 188, 189, 190, 191, 192, 193, 194, 195, 196, 197, 198, 199, 200, 201. 202, 203, 204, 205, 206, 207, 208, 209, 210, 211, 212, 213, 214, 215, 216, 217, 218, 219, 220, 221, 222, 223, 224, 225. 226, 227, 228, 229, 230, 231, 232, 233, 234. 235, 236. 237, 238, 239, 240. 241, 242, 243, 244, 245, 246, 247, 248, 249, 250, 251, 252, 253. 254, 255, 256, 257, 258, 259, 260, 261, 262, 263, 264, 265, 266, 267, 268, 269, 270, 271, 272, 273, 274, 275. 276, 277, 278, 279. 280, 281, 282, 283, 284, 285, 286, 287, 288, 289, 290, 291, 292. 293, 294, 295. 296, 297, 298, 299. 300, 301, 302, 303, 304, 305, 306, 307, 308, 309, 310, 311,

312, 313, 314, 315. 316, 317, 318, 319, 320. 321, 322, 323, 324, 325, 326, 327, 328. 329, 330, 331, 332, 333, 334, 335, 336, 337, 338, 339, 340, 341, 342, 343, 344, 345, 346, 347, 348, 349, 350. 351, 352, 353, 354. 355, 356, 357, 358, 359, 360, 361, 362, 363, 364, 365, 366, 367. 368, 369, 370, 371, 372, 373. 374, 375, 376, 377, 378. 379, 380, 381, 382, 383, 384, 385, 386, 387, 388, 389. 390. 391, 392. 393. 394, 395, 396, 397. 398, 399, 400, 401, 402. 403, 404, 405, 406. 407, 408, 409. 410. 411, 412, 413. 414. 415. 416, 417, 418, 419, 420. 421. 422. 423, 424, 425. 426, 427, 428, 429. 430, 431, 432, 433, 434. 435, 436, 437, 438, 439, 440, 441, 442, 443, 444, 445, 446, 447, 448. 449, 450, 451, 452, 453, 454, 455, 456, 457, 458, 459, 460, 461. 462, 463, 464. 465, 466, 467, 468. 469, 470, 471, 472. 473, 474, 475, 476. 477, 478, 479, 480, 481. 482, 483. 484, 485, 486, 487. 488, 489, 490, 491. 492. 493, 494, 495, 496, 497, 498, 499, and 500 nucleotides in length.

[00125] In one embodiment the poiyadenylation sequence is 50-100 nucleotides in length.

[00126] In one embodiment, the poiyadenylation sequence is 50-150 nucleotides in length.

[00127] In one embodimeni, the poiyadenylation sequence is 50-160 nucleotides in length.

[00128] In one embodiment, the poiyadenylation sequence is 50-200 nucleotides in length.

[00129] In one embodiment, the poiyadenylation sequence is 60-100 nucleotides in length.

[00130] In one embodiment, the poiyadenylation sequence is 60-150 nucleotides in length.

[00131] In one embodiment, the poiyadenylation sequence is 60-160 nucleotides in length.

[00132] In one embodiment, the poiyadenylation sequence is 60-200 nucleotides in length.

[00133] In one embodiment the poiyadenylation sequence is 70-100 nucleotides in length.

[00134] hi one embodiment, the poiyadenylation sequence is 70-150 nucleotides in length.

[00135] In one embodiment, the poiyadenylation sequence is 70-160 nucleoiides in length.

[00136] In one embodiment, the poiyadenylation sequence is 70-200 nucleoiides in length.

[00137] in one embodiment, the poiyadenylation sequence is 80-100 nucleotides in length.

[00138] In one embodiment, the poiyadenylation sequence is 80-150 nucleotides *m* length.

[00139] In one embodiment, the poiyadenylation sequence is 80-160 nucleotides in length.

[00140] In one embodiment the poiyadenylation sequence is 80-200 nucleotides in length.

[00141] hi one embodiment, the poiyadenylation sequence is 90-100 nucleotides in length.

[00142] In one embodiment, the poiyadenylation sequence is 90-150 nucleoiides in length.

[00143] In one embodiment, the poiyadenylation sequence is 90-160 nucleoiides in length.

[00144] In one embodiment, the poiyadenylation sequence is 90-200 nucleoiides in length.

Viral Genome Component: Linkers

[00145] Viral genomes of the invention may be engineered with one or more spacer or linker regions to separate coding or non-coding regions.

[00146] In one embodiment, the payload region of the AAV particle may optionally encode one or more linker sequences. In some cases, the linker may be a peptide linker that may be used to connect the polypeptides encoded by the payload region (i.e.. light and heavy antibody chains during expression). Some peptide linkers may be cleaved after expression to separate heavy and light chain domains, allowing assembly of mature antibodies or antibody fragments. Linker cleavage may be enzymatic. in some cases, linkers comprise an enzymatic cleavage site to facilitate intracellular or extracellular cleavage. Some payload regions encode linkers that interrupt polypeptide synthesis during translation of the linker sequence from an mRNA transcript. Such linkers may facilitate the translation of separate protein domains (e.g., heavy and light chain antibody domains) from a single transcript. in some cases, two or more linkers are encoded by a payload region of the viral genome. Non-limiting examples of linkers that may be encoded by the payload region of an AAV particle viral genome are given in Table *2*.

Table **2. Linkers**

Linker No.	Description	**SEQ ID NO**
L I	Internal **ribosome entry** site (IRES)	899
L2	Foot **and** month disease **virus** 2A (F2A)	**900**
L3	Porcine **teschovinis-1** virus 2A (P2A)	901
L4	**Furin** cleavage site (F)	902
L5	**5xG4S** (SEQ ID **NO:** 9221)	903
L6	**1,4-alpha-glucan-branching** enzyme	CHP
L7	**1,4-alpha-glucan-branching enzyme**	904
L8	1,4-beta-**N**-acetylmuramidase	FKK
L9	1,4-beta-**N**-acetylmuramidase	905
L10	**1,4-beta-N-acetylmuramidase**	906
L l t	**1,4-beta-N-acetylmurainidase**	907
L12	**1,4-beta-N-acetylmuramidase**	908
L13	**1,4-beta-N-acetylmuramidase**	909
L i4	**1,4-beta-N-acetylmuramidase**	9 10
L15	**1,4-beta-N-acetylmuramidase**	9 1 l
L16	**1,4-beta-N-acetylmuramidase**	**9 l 2**
L17	**1,4-beta-N-acetylmuramidase**	913
L18	**1,4-beta-N-acetylmuramidase**	914
L19	150aa long hypothetical transcriptional regulator	9 15
L20	150aa long hypothetical **transcriptional** regulator	916
L2 1	**1-deoxy-D-xylulose 5-phosphate reductoisomerase**	**9 17**
L22	**1-deoxy-D-xylulose 5-phosphate** reductoisomerase	918
L23	**1-deoxy-D-xylulose 5-phosphate** reductoisomerase	919
L24	**1-deoxy-D-xylulose** 5-phosphate reductoisomerase	920
L25	235aa long hypothetical **biotin-[acetyl-CoA-carboxylase]** ligase	92 1
L26	**235aa** long hypothetical **biotin-[acetyl-CoA-carboxylase]** ligase	922

L27	235aa long hypothetical biotin- [acety1-CoA-carboxylasej ligase	923
L28	2-dehydropantoate 2-reductase	j 924
L29	2-dehydropaiitoate 2-reductase	j 925
L30	2-dehydropantoate 2-reductase	926
L3 !	2-dehydropantoate 2-reductase	! 927
L32	2-dehydropantoate 2-reductase	928
L33	2-dehydropantoate 2-reductase	j 929
L34	2-dehydropantoate 2-reductase, putative	j 930
L35	2-dehydropantoate 2-reductase, putative	93 1
L36	4-alpha-glucanotrans ferase	j 932
L37	4-alpha-glucano transferase	j 933
L38	4-alpha-glucanotransferase	j 934
L39	4-diphosphocytidyl-2C-methyl-D-erythritol kinase	HAA
L40	4-diphosphocytidyl-2C-inethyl-D-er3'thr tol kinase	935
L4 !	4-diphosphocytidyl-2C-metlryl-D-erythritol kinase	936
L42	4-diplrosphocytidyl-2C-me11ryl-D-erythritol kinase	937
L43	4-diphosphocytidyl-2C-methyl-D-erythritol kinase	j 938
L44	4-hydroxyphenylpyruvate dioxygenase	j 939
L45	5-13 amino acids from the N termini of human Ck and CHI domains linker	940
L46	5-13 amino acids from the N termini of human Ck and CHi domains linker	j ERK
L47	5-13 amino acids from the N termini of human Ck and CHI domains linker	941
L48	5-13 amino acids from the N termini of human Ck and CHI domains linker	j 942
L49	5-13 amino acids from the N termini of human Ck and CHI domains linker	j 943
L50	5-13 amino acids from the N termini of human Ck and CHI domains linker	944
L5 !	5'-exonuclease	j 945
L52	5-meihyltetrahydropteroyltriglutasnate--honiocysteinemethyltransferase	ARL
L53	5-inetliyltetraliydropieroyltriglutartiate--homocysteitiemethy1transferase	j 946
L54	5-methyltetrahydropteroyltriglutamaie--homocysteinemeihyltraiisferase	j 947
L55	5-methyltetrahydropteroyltriglutaniate--homocysteinemethyltransferase	948
L56	5-methyltetrahydropteroyltriglutamate--homocysteinemethyltransferase	! 949
L57	5'-nucleoridase	950
L58	5'-nucleotidase	951
L59	5'-nucleotidase	j 952
L60	5'-nucleotidase	j 953
L6 1	704aa long hypothetical glycosyltransferase	! 954
L62	704aa long hypothetical glycosyltransferase	955
L63	80 kDa nuclear cap binding protein	j 956
L64	80 kDa nuclear cap binding protein	957
L65	80 kDa nuclear cap binding protein	958
L66	80 kDa nuclear cap binding protein	! 959
L67	Acetaldehyde dehydrogenase (acylating)	960
L68	Acetaldehyde dehydrogenase (acylating)	j 961
L69	Aceto lactate synthase isozyme II small subunit	962
L70	Acetylcholine receptor protein, alpha chain	j 963
L7 1	Acetylcholine receptor protein, beta chain	j 964

L72	**Aconitate** hydratase 2	965
L73	**Aconitate hydratase** 2	966
L74	Aconitate hydratase 2	967
L75	Aconitate hydratase 2	968
L76	**Aconitate** hydratase 2	969
L77	Acriflavine resistance protein **B**	DWY
L78	**Acriflavine** resistance protein B	GGS
L79	**Acriflavine** resistance protein B	IDQ
L80	Acriflavine resistance protein B	NKV
L81	Acriflavine **resistance** protein **B**	SEA
L82	Acriilavuie resistance protein **B**	970
L83	Acriflavine resistance protein B	971
L84	Acriflavine resistance protein B	972
L85	Acriflavine resistance protein B	973
L86	**Acriflavine resistance protein B**	974
L87	Acriflavine resistance protein **B**	975
L88	Acriflavine resistance protein **B**	976
L89	Acriflavine resistance protein B	977
L90	Acriflavine resistance protein B	978
L91	**Acriflavine resistance protein B**	979
L92	Acriflavine resistance protein **B**	980
L93	Acriflavine resistance protein **B**	981
L94	Acriflavine resistance protein B	982
L95	Acriflavine resistance protein B	983
L96	**Acriflavine resistance protein B**	984
L97	Acriflavine resistance protein B	985
L98	Acriflavine resistance protein **B**	986
L99	Acriflavine resistance protein B	987
L100	Acriflavine resistance protein B	988
L101	Acriflavine resistance protein B	989
L102	Acriflavine resistance protein B	990
L103	Acriflavine resistance protein B	991
L104	Acriflavine resistance protein B	992
L105	Acriflavine resistance protein B	993
L106	**Acyl-CoA** ihioesterase **II**	994
L107	Acyl-CoA tlnoesterase IT	995
L108	**Acyl-CoA thioesterase II**	996
L109	Acyl-CoA thioesterase IE	997
L110	Acyl-CoA tlnoesterase **II**	998
L111	**Acyl-coenzyme** A tlnoesterase 4	999
L112	**Acyl-coenzyme A** ihioesterase 4	1000
L113	Acyi-coenzyme A tlnoesterase 4	1001
L114	Acyl-coenzyme A thioesterase 4	1002
L115	**Acyl-coenzyme** A thioesterase 4	1003
L116	Adenine giycosylase	1004

L I 17	Adenylate cyclase	1005
L 118	**Aerolysin**	1006
L I 19	Aeroiysm	1007
L I20	**Agglutinin**	DWK
L12 1	**Agglutinin isolectin** 1	1008
L 122	**Agglutinin** isolecitti **I**	1009
L123	Aldehyde **ferredoxin oxidoreductase**	1010
L124	Aldehyde oxidoreductase	1011
L125	Aldehyde oxidoreductase	1012
L126	Aldehyde oxidoreductase	1013
L i27	Aldehyde oxidoreductase	1014
L128	Aldehyde oxidoreductase	1015
L129	Aikyl hydroperoxide reductase subunit F	1016
L130	**Aikyl** hydroperoxide reductase **subunit** F	1017
L13 1	Aikyl hydroperoxide reductase subunit F	1018
L i32	Aikyl hydroperoxide reductase **subunit** F	1019
L133	Aikyl hydroperoxide reductase subunit F	1020
L134	Aikyl hydroperoxide reductase subunit F	1021
L135	**Aikyl** hydroperoxide reductase **subunit** F	1022
L I36	Aikyl hydroperoxide reductase subunit F	1023
L137	Aikyl hydroperoxide reductase **subunit** F	1024
L138	Aikyl hydroperoxide reductase subunit F	1025
L139	Allantoicase	1026
L i40	**Allantoicase**	1027
L 14 1	**Alliin** lyase 1	SAV
L i42	**Alliin lyase** 1	1028
L143	Alliin lyase 1	1029
L144	Alliin lyase 1	1030
L145	Alliin lyase 1	1031
L I46	Alpha amylase	1032
L i47	Alpha **amylase**	1033
L148	Alpha-actinin 1	1034
L149	**Alpha-actinin** 1	1035
L150	**Alpha-adaptin** C	1036
L15 1	**Alpha-amylase**	1037
L i52	Alpha-glueuro nidase	LSD
L153	**Alpha-glucuronidase**	1038
L i54	**Alpha-glucuronidase**	1039
L155	Alpha-glucuronidase	1040
L I56	**Alplia-glucuro**nidase	1041
L i57	**Alpha-glueuro**nidase	1042
L158	Alpha-glucuronidase	1043
L i59	Alpha-glucuronidase	1044
L I60	Alpha-glucuronidase	1045
L161	**Alplia-glucuro**nidase	1046

L162	Alpha-glucuronidase	1047
L163	Alpha-glucuronidase	1048
L164	Alplia-glueuronidase	1049
L165	Alpha-glucuio tisdase	1050
L166	Alpha -glucuronid ase	1051
L167	Alpha-glucuroiiidase	1052
L168	Alpha-glucuronidase	1053
L169	Alplia-glueuronidase	1054
L170	Alpha-glucuronidase	1055
L171	Alpha-glucuronidase	1056
L172	Alpha-glucuroiiidase	1057
L173	Alpha-glucuronidase	1058
L174	Alpha-L-arabinoftiranDsidase B	1059
L175	Alpha-niannosidase	1060
L176	Alr2269 protein	1061
L177	AMP nucleosidase	1062
L178	AMP nucleosidase	1063
L179	AMP nucleosidase	1064
L180	Angiopoietin-1 receptor	DAG
L181	Angiopoietin-1 receptor	NSG
L182	Angjopoietin- 1 receptor	TSA
L183	Angio poietin- 1 receptor	VPR
L184	Angiopoietin-1 receptor	1065
L185	Angiopoietin-1 receptor	1066
L186	Angiopoietin-1 receptor	1067
L187	Angiopoietin-1 receptor	1068
L188	Angio poietin- 1 receptor	1069
L189	Angiopoietin-1 receptor	1070
L190	Angiopoietin-1 receptor	1071
L191	Angiopoietin-1 receptor	1072
L192	Angiopoietin-1 receptor	1073
L193	Angio poietin- 1 receptor	1074
L194	Angiopoietin-1 receptor	1075
L195	Angiopoietin-1 receptor	1076
L196	Angiopoietin-1 receptor	1077
L197	Angiopoietin-1 receptor	1078
L198	Angio poietin- 1 receptor	1079
L199	Augjopoietin- 1 receptor	1080
L200	Angiopoietin-1 receptor	1081
L201	Angiopoietin-1 receptor	1082
L202	Angiopoietin-1 receptor	1083
L203	Angio poietin- 1 receptor	1084
L204	Angjopoietin- 1 receptor	1085
L205	Annexin A2	QNK
L206	Annexin A2	1086

L207	**Annexin** A2	1087
L208	Anthranilate **phosphoribosyltransferase**	1088
L209	AP-2 complex **subunit** beta-2	1089
L210	Archaeosine **tRNA-guanine transglycosylase**	LGI
L211	Archaeosine **tRNA-guanine** transglycosylase	1090
L212	Archaeosine **tRNA-guanine** transglycosylase	1091
L213	**Archaeosine** tRNA-guanine transglycosylase	1092
L214	Archaeosine **tRNA-guanine** transglycosylase	1093
L215	Archaeosine tRNA-guanine transglycosylase	1094
L216	Archaeosine **tRNA-guanine** transglycosylase	1095
L217	Archaeosine **tRNA-guanine** transglycosylase	1096
L218	**Archeal exosome RNA** binding protein np4	1097
L219	Archeal exosome RNA binding protein rrp4	1098
L220	Archeal exosome RNA binding protein rrp4	1099
L221	Arginyl-tRNA synthetase	**IDY**
L222	**Arginyl-tRNA** synthetase	**1100**
L223	Arginyl-tRNA synthetase	**1101**
L224	Arginyl-tRNA synthetase	1102
L225	Arrestin	1103
L226	**Arrestin**	**1104**
L227	Arsenite oxidase	**1105**
L228	Artificial linker	PCS
L229	Artificial linker	**ATK**
L230	Artificial linker	ASK
L231	Artificial Sinker	1106
L232	**Artificial** linker	**1107**
L233	Artificial linker	1108
L234	Artificial linker	1109
L235	Artificial linker	1110
L236	Artificial linker	**1111**
L237	ATP phosphoribosyltransferase	ANR
L238	**ATP-dependent DNA helicase**	YDP
L239	ATP-dependent DNA helicase	**1112**
L240	ATP-dependent DNA helicase	1113
L241	ATP-dependent DNA helicase	**1114**
L242	ATP-dependent DNA helicase	1115
L243	ATP-dependent DNA helicase	1116
L244	ATP-dependent DNA helicase	1117
L245	ATP-dependent DNA helicase	1118
L246	ATP-dependent DNA helicase	1119
L247	**AT-rich DNA-binding** protein	1120
L248	AT-rich **DNA-binding** protein	**1121**
L249	**Axonin-** 1	DEG
L250	**Axonin-** 1	ECF
L251	Axonin- 1	1122

L252	Axoniri- 1	1123
L253	Axonin- 1	1124
L254	Axoiiin-1	1125
L255	Axonin- !	1126
L256	Axonin- 1	1127
L257	Axonin- 1	1128
L258	Bacilysin biosynthesis protein BacB	1129
L259	Bacilysin biosynthesis protein BacB	1130
L260	Bacilysin biosynthesis protein BacB	1131
L261	Bacilysin biosynthesis protein BacB	1132
L262	Bacilysin biosynthesis protein BacB	1133
L263	Bacteriophage **Mu** transposase	1134
L264	Bacteriophage Mu transposase	1135
L265	Benzoyl-CoA-dihy drodiol lyase	1136
L266	Benzoyl-CoA-dihydtrodiol lyase	1137
L267	Benzoyl-CoA-dihy drodiol lyase	1138
L268	Benzoyl- CoA-dihydrodiol lyase	1139
L269	Benzoyl-CoA-dihy drodiol lyase	1140
L270	Benzoylformate decarboxylase	1141
L271	BenzoyIformate decarboxylase	1142
L272	Benzoylformate decarboxylase	1143
L213	Beta-amylase	1144
L274	Beta-galactosidase	AiS
L275	Beta-galactosidase	1145
L276	Beta-galactosidase	1146
L277	Beta-galactosidase	1147
L278	Beta-galactosidase	1148
L279	Beta-galactosidase	1149
L280	Beta-galactosidase	1150
L281	Beta-galactosidase	1151
L282	Beta-galactosidase	1152
L283	Beta-galactosidase	1153
L284	Beta-galactosidase	1154
L285	Beta-galactosidase	1155
L286	Beta-galactosidase	1156
L287	Beta-galactosidase	1157
L288	Beta-galactosidase	1158
L289	Beta-galactosidase	1159
L290	Beta-galactosidase	1160
L291	Beta-galactosidase	1161
L292	Beta-galactosidase	1162
L293	Beta-galactosidase	1163
L294	Beta-galactosidase	1164
L295	Beta-galactosidase	1165
L296	Beta-galactosidase	1166

L297	**Beta-N-acety Hexosaminidase**	**QRE**
L298	**Beta-N-acetylhexosaminidase**	1167
L299	**Beta-N-acetylhexosaniinidase**	1168
L300	**Beta-N-acetylhexosaminidase**	1169
L301	**Bifiinctional** NMN **adenylyltransferase/Niidix** hydrolase	**1170**
L302	Bifunetional purine biosynthesis protein PURll	1171
L303	Bi!iverdin reductase A	EHV
L304	Biliverdin reductase A	**LME**
L305	Biliverdin reductase **A**	**1172**
L306	**Biliverdin** reductase A	**1173**
L307	**Biodegradative arginine** decarboxylase	**TVQ**
L308	Biodegradative arginine decarboxylase	1174
L309	Biodegradative arginine decarboxylase	1175
L310	Biodegradative arginine decarboxylase	**1176**
L311	Biodegradative arginine decarboxylase	**1177**
L312	Biodegradative **arginine** decarboxylase	1178
L313	Biodegradative arginine decarboxylase	1179
L314	Biodegradative arginine decarboxylase	1180
L315	Biodegradative arginine decarboxylase	1181
L316	Biodegradative arginine decarboxylase	1182
L317	Biodegradative **arginine** decarboxylase	1183
L318	Biodegradative arginine decarboxylase	1184
L319	Biodegradative arginine decarboxylase	1185
L320	Biotin carboxylase	1186
L321	**Bowinan-Birk** trypsin **inhibitor**	**1187**
L322	**Bpt4** gene 59 helicase **assembly** protein	KQI
L323	**B**RCA 1-associated RING domain protein 1	1188
L324	BRCAl-associated RING domain protein 1	1189
L325	BRCA **1-associated** RING domain protein 1	1190
L326	Breast cancer 2	1191
L327	Breast cancer 2	1192
L328	Breast cancer 2	1193
L329	Breast cancer 2	1194
L330	Breast cancer 2	1195
L331	Breast cancer 2	1196
L332	**Butyrate** response factor 2	1197
L333	**C4b-binding** protein	**YKR**
L334	**C4b-binding** protein	1198
L335	C5a peptidase	1199
L336	**C5a** peptidase	1200
L337	**C5a** peptidase	1201
L338	**C5a peptidase**	1202
L339	**C5a** peptidase	1203
L340	C5a peptidase	1204
L341	**C5a** peptidase	1205

L342	C5a peptidase	1206
L343	C5a peptidase	1207
L344	C5a peptidase	1208
L345	C5a peptidase	1209
L346	C5a peptidase	1210
L347	C5a peptidase	1211
L348	Calcium-binding protein	1212
L349	CarA	1213
L350	CarA	1214
L351	Carbamoyl phosphate synthetase (small chain)	1215
L352	Carbamoyl phosphate synthetase (small chain)	1216
L353	Carbamoyl phosphate synthetase (small chain)	1217
L354	Carbamoyl phosphate synthetase (small chain)	1218
L355	Carbamoyl phosphate synthetase (small chain)	1219
L356	Carbon monoxide dehydrogenase/acetyl-CoA synthase subunitalpha	1220
L357	Carboxypeptidase Gp180 residues 503-882	HRG
L358	Catabolite activation-like protein	1221
L359	Catabolite activation-like protein	1222
L360	Catechol 2,3-dioxygenase	1223
L361	Cation-independent mannose 6-phosphate receptor	1224
L362	CD3 epsilon and gamma ectodomain fragment complex	1225
L363	CD3 epsilon and gamma ectodomain fragment complex	1226
L364	Cell filamentation protein	SNP
L365	Cell filamentation protein	1227
L366	Cell filamentation protein	1228
L367	Cellular coagulation factor XIII zymogen	DIT
L368	Cellular coagulation factor XIII zymogen	NSD
L369	Cellular coagulation factor XIII zymogen	TDT
L370	Cellular coagulation factor XIII zymogen	1229
L371	Cellular coagulation factor XIII zymogen	1230
L372	Cellular coagulation factor XIII zymogen	1231
L373	Cellular coagulation factor XIII zymogen	1232
L374	Cellular coagulation factor XIII zymogen	1233
L375	Cellular coagulation factor XIII zymogen	1234
L376	Cellular coagulation factor XIII zymogen	1235
L377	Cellular coagulation factor XIII zymogen	1236
L378	Cellular coagulation factor XIII zymogen	1237
L379	Cellular coagulation factor XIII zymogen	1238
L380	Cellular coagulation factor XIII zymogen	1239
L381	Cellular coagulation factor XIII zymogen	1240
L382	Cellular coagulation factor XIII zymogen	1241
L383	Cellular coagulation factor XIII zymogen	1242
L384	Cellular coagulation factor XIII zymogen	1243
L385	Cellular coagulation factor XIII zymogen	1244
L386	Cellular coagulation factor XIII zymogen	1245

L387	Cellular coagulation factor XIII **zymogen**	1246
L388	Cellular coagulation factor XIII **zymogen**	1247
L389	Celiulase	1248
L390	**Cellulase**	1249
L391	Celiulase	1250
L392	Cellulase	1251
L393	Celiulase	1252
L394	Celiulase	1253
L395	Celiulase	1254
L396	**Cellulase**	1255
L397	Cellulase	1256
L398	Celiulase **linker**	1257
L399	Cellulase linker	1258
L400	Celiulase linker	1259
L401	Celiulase linker	1260
L402	**Chaperone** protein FimC	**KLR**
L403	**Chaperone** protein FimC	QAA
L404	**Chaperone** protein FimC	1261
L405	Chaperone protein FimC	1262
L406	Chaperone protein HscB	RHP
L407	Chaperone protein HscB	1263
L408	CheB methylesterase	1264
L409	CheB methylesterase	1265
L410	CheB methylesterase	1266
L411	**Chelatase, putative**	1267
L412	**Chemotaxis** receptor **methyltransferase** cheR	1268
L413	**Chemotaxis** receptor methyltransferase cheR	1269
L414	Chemotaxis receptor methyltransferase cheR	1270
L415	Cholesteroi oxidase	1271
L416	**Cholesterol** oxidase	1272
L417	Cholesteroi oxidase	1273
L418	Cholesterol oxidase	1274
L419	Cholesteroi oxidase	1275
L420	Cholesterol oxidase	1276
L421	Cholesteroi oxidase	1277
L422	Cholesteroi oxidase	1278
L423	Cholesterol oxidase	1279
L424	Cholesterol oxidase	1280
L425	Cholesteroi oxidase	1281
L426	Cholesteroi oxidase	1282
L427	Chromatin **structure-remodeling** complex protein RSC4	**KNL**
L428	**Chromatin** structure-remodeling **complex** protein RSC4	1283
L429	Chromatin **structure-remodeling** complex protein RSC4	1284
L430	**Chromatin** structure-remodeling complex protein RSC4	1285
L431	**Chromodomain-helicase-DNA-binding** protein 1	1286

L432	Chromodomain-helicase-DNA-binding protein 1	1287
L433	Cleavable disulfide	1288
L434	Cleavable disulfide	1289
L435	Cleavable disulfide	1290
L436	Cleavable disulfide	1291
L437	Cleavable disulfide	1292
L438	Cleavable disulfide	1293
L439	Cleavable disulfide	1294
L440	Cleavable disulfide	1295
L441	Cleavable disulfide	1296
L442	Cleavable disulfide	1297
L443	Cleavable disulfide	1298
L444	Colicin Ia	1299
L445	Collagen adhesin	1300
L446	Complement C3 beta chain	1301
L447	Complement C3 beta chain	1302
L448	Complement C3 beta chain	1303
L449	Complement C3 beta chain	1304
L450	Complement decay-accelerating factor	EIY
L451	Complement factor H	KRP
L452	Complement receptor type 2	1305
L453	Conserved hypothetical protein	1306
L454	Conserved hypothetical protein MTH1747	DIR
L455	Conserved hypothetical protein MTH1747	1307
L456	Conserved hypothetical protein MTH1747	1308
L457	Conserved hypothetical protein MTH1747	1309
L458	Conserved hypothetical protein MTH1747	1310
L459	Conserved hypothetical protein MTH1747	1311
L460	Conserved hypothetical protein MTH1747	1312
L461	Conserved hypothetical protein MTH1747	1313
L462	Conserved protein (MTH177)	1314
L463	Creatine amidinohydrolase	1315
L464	Cruciferin	1316
L465	Cruciferin	1317
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L55 1	DNA damage-binding protein 1	1396
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L553	DNA damage-binding protein 1	1398
L554	DNA damage-binding protein 1	1399
L555	DNA damage-binding protein 1	1400
L556	DNA damage-binding protein 1	140 1
L557	DNA damage-binding protein 1	1402
L558	DNA damage-binding protein 1	1403
L559	DNA damage-binding protein 1	1404
L560	DNA damage-binding protein 1	1405
L561	DNA damage-binding protein 1	1406
L562	DNA damage-binding protein 1	1407
L563	DNA damage-binding protein 1	1408
L564	DNA damage-binding protein 1	1409
L565	DNA damage-binding protein 1	1410
L566	DNA damage-binding protein 1	141 1

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L620	DNA polymerase I	ELE
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L628	DNA **primase**	1464
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L634	DNA primase/helicase	1469
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L642	DNA **topoisomerase** 2	**IPI**
L643	DNA topoisomerase 2	**KEL**
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L648	DNA topoisomerase 2	1480
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L656	DNA topoisomerase **II,** alpha isozyme	PDL

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L765	**Fibronectin**	1591
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L767	Fibronectin	1593
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L770	Flagellar hook protein FlgE	1596
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L773	Flagellar hook protein FlgE	1599
L774	Flagellar hook protein FlgE	**1600**
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L777	Flexible G/S rich linker	S
L778	Flexible **G/S rich** linker	**GG**
L779	Flexible G/S rich **linker**	GS
L780	Flexible G/S rich linker	GGS
L781	Flexible G/S rich linker	GGG
L782	Flexible G/S rich linker	1602
L783	Flexible G/S **rich** linker	1603
L784	Flexible G/S rich linker	1604
L785	Flexible G/S rich linker	1605
L786	Flexible G/S rich linker	1606
L787	Flexible G/S rich linker	1607
L788	Flexible G/S **rich** linker	1608
L789	Flexible G/S rich linker	1609
L790	Flexible G/S rich linker	1610
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L792	Flexible G/S rich **linker**	1612
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L798	FolC bifunctional protein	1618
L799	FolC **bifunctional** protein	1619
L800	FolC bifunctional protein	1620
L801	FolC bifunctional protein	1621
L802	FolC **bifunctional** protein	1622
L803	FolC bifunctional protein	1623
L804	FolC **bifunctional** protein	1624
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L806	Formate dehydrogenase (large **subunit)**	**YDK**
L807	Formate dehydrogenase (large **subunit)**	1626
L808	Formate dehydrogenase (large subunit)	1627
L809	**Formate** dehydrogenase (large subunit)	1628
L810	Formate dehydrogenase (large subunit)	1629
L811	Formate dehydrogenase (large subunit)	1630
L812	Formate dehydrogenase (large **subunit)**	1631
L813	Formate dehydrogenase (large subunit)	1632
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L817	**Formate** dehydrogenase (large subunit)	1636
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L820	Formate dehydrogenase, nitrate-inducible, major subunit	1639
L821	Formate dehydrogenase, nitrate-inducible, major subunit	1640
L822	**Formate** dehydrogenase, nitrate-inducible, major subunit	1641
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L830	Formate dehydrogenase, nitrate-inducible, major subunit	1649
L831	Formate dehydrogenase, nitrate-inducible, major subunit	1650
L832	**Formate** dehydrogenase, nitrate-inducible, major subunit	1651
L833	**Fumarylacetoacelate** hydrolase	1652
L834	Galactose oxidase	GSV
L835	Galactose oxidase	**GWK**
L836	Galactose oxidase	IAE

L837	**Galactose** oxidase	KR**Q**
L838	Galactose oxidase	QDT
L839	Galactose oxidase	TPN
L840	**Galactose** oxidase	1653
L841	Galactose oxidase	1654
L842	Galactose oxidase	1655
L843	Galactose oxidase	1656
L844	Galactose oxidase	1657
L845	Galactose oxidase	1658
L846	Galactose oxidase	1659
L847	Galactose oxidase	1660
L848	Galactose oxidase	1661
L849	Galactose oxidase	1662
L850	Galactose oxidase	1663
L85 1	Galactose oxidase	1664
L852	Galactose oxidase	1665
L853	Galactose oxidase	1666
L854	Galactose oxidase	1667
L855	Galactose oxidase	1668
L856	Galactose oxidase	1669
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L861	Galactose oxidase	1674
L862	Galactose oxidase	1675
L863	Galactose oxidase	1676
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L866	Gelation factor	DSS
L867	Gelation factor	1679
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L873	**Glucodextraiiase**	1685
L874	Glucodextranase	1686
L875	**Gluco**sainine-**frue**tose-6-phosphate aminotransferase	YEQ
L876	**Glucosamine-fructose-6-phosphate aminotransferase**	1687
L877	**Gliicosamine-fnictose-6-phosphate** aminotransferase	1688
L878	Glucosamine-fractose-6-phosphate **aminotransferase**	1689
L879	**Glucosamine-friictose-6-phospnate aminotransferase**	1690
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L897	Glutamate synthase [NADPH] large chain	1706
L898	Glutamate synthase [NADPH] large chain	1707
L899	Glutamate synthase [NADPH] large chain	1708
L900	Glutamate synthase [NADPH] large chain	1709
L901	Glutamate synthase [NADPH] large chain	1710
L902	Glutamate synthase [NADPH] large chain	1711
L903	Glutamine synthetase	1712
L904	Glutamine synthetase	1713
L905	Glutamyl-tRNA synthetase	1714
L906	Glutamy S-tRNA synthetase	1715
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L908	Glutamy 1-tRNA synthetase	1717
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L970	Hybrid cluster protein	1777
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L972	Hybrid **cluster protein**	1779
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L983	Hypothetical protein	AEP
L984	Hypothetical protein	**1790**
L985	Hypothetical protein APE0525	PTL
L986	Hypothetical protein APE0525	**1791**
L987	Hypothetical protein **LOC449832**	1792
L988	Hypothetical protein **LOC449832**	1793
L989	Hypothetical protein PA4388	1794
L990	Hypothetical protein PA520 1	ASE
L991	Hypothetical protein PA5201	**QDP**
L992	Hypothetical protein PA520 1	**VKL**
L993	Hypothetical protein **PA5201**	1795
L994	Hypothetical protein PA520 1	1796
L995	Hypothetical protein PA520 1	1797
L996	Hypothetical protein PA5201	1798
L997	Hypothetical **protein PA520** 1	1799
L998	Hypothetical protein **PA5201**	1800
L999	Hypothetical protein PA520 1	1801
L1000	Hypothetical protein PA520 1	1802
L100 1	Hypothetical protein PA5201	1803
L I**002**	Hypothetical **protein PA520** 1	1804
L1003	Hypothetical protein **PA5201**	1805
L1004	Hypothetical protein PA520 1	1806
L1005	Hypothetical protein PA5201	1807
L I006	Hypothetical protein PA5201	1808
L I**007**	Hypothetical **protein PA520** 1	1809
L1008	Hypothetical protein **PA5201**	1810
L1009	Hypothetical protein PA520 1	**181 1**
L1010	Hypothetical protein PA5201	1812
L101 1	Hypothetical protein PA5201	1813
L 10 12	Hypothetical **protein PA520** 1	1814
L10L3	Hypothetical protein **PH0495**	**ASN**
L10 14	Hypothetical protein **PH0495**	**1815**
L1015	Hypothetical protein PH0495	1816
L1016	Hypothetical protein PH0495	1817

L1017	Hypothetical protein PH0495	1818
L1018	Hypothetical protein PH0510	1819
L1019	Hypothetical protein PH0510	1820
L1020	Hypothetical protein PH1313	1821
L1021	Hypothetical protein PH1313	1822
L1022	Hypothetical protein SLR0953	1823
L1023	Hypothetical protein SLR0953	1824
L1024	Hypothetical protein SLR0953	1825
L1025	Hypothetical protein SLR0953	1826
L1026	Hypothetical protein SLR0953	1827
L1027	Hypothetical protein YIGZ	1828
L1028	Hypothetical protein YIGZ	1829
L1029	Hypothetical protein YJIA	1830
L1030	Hypothetical protein YJIA	1831
L1031	Hypothetical protein YJIA	1832
L1032	Hypothetical protein YJIA	1833
L1033	Hypothetical protein YJIA	1834
L1034	Hypothetical tRNA/rRNA methyltransferase YJFH	1835
L1035	Hypothetical tRNA/rRNA methyltransferase YJFH	1836
L1036	IclR transcriptional regulator	1837
L1037	MR transcriptional regulator	1838
L1038	IclR transcriptional regulator	1839
L1039	MR transcriptional regulator	1840
L1040	Integrase	1841
L1041	Interferon, alpha-inducible protein (clone IFI-15k)	1842
L1042	Interleukin-1 receptor, type I	AIF
L1043	Interteukin-1 receptor, type I	1843
L1044	Interleukin-1 receptor, type I	1844
L1045	Interleukin-1 receptor, type I	1845
L1046	Interleukin-12 subunit p40	FFI
L1047	Interleukin-12 subunit p40	1846
L1048	Interleukin-12 subunit p40	1847
L1049	Interleukin-12 subunit p40	1848
L1050	Interleukin-12 subunit p40	1849
L1051	Interleukin-12 subunit p40	1850
L1052	Interleukin-12 subunit p40	1851
L1053	Interleukin-12 subunit p40	1852
L1054	Interleukin-2 receptor alpha chain	1853
L1055	Interleukin-2 receptor alpha chain	1854
L1056	Internalin B	VTQ
L1057	Internalin B	1855
L1058	Intenialin B	1856
L1059	Internalin B	1857
L1060	Internalin B	1858
L1061	Internalin B	1859

L1062	Internal!!! B	1860
L1063	**Internalin** B	1861
L I064	Internal in B	¦862
L I065	**Internalin** B	1863
L¦066	Internalin B	¦864
L1067	Internalin B	1865
L1068	**Internalin** B	1866
L I069	Intimin	SLV
L¦070	**Intimin**	1867
L¦07 i	Intimin	¦868
L1072	Intimin	1869
L1073	Intron-encoded DNA endoniiclease **I-anil**	1870
L¦074	**Intron-encoded** DNA **endonuclease** I-anil	1871
L¦075	**Invasin**	KST
L¦076	**Invasin**	¦872
L1077	Invasin	1873
L1078	Invasin	¦874
L¦079	Invasin	1875
L¦080	Invasin	1876
L108 I	Invasin	¦877
L1082	Invasin	1878
L1083	Invasin	¦879
L1084	Invasin	1880
L i085	Invasin	1881
L1086	Invasin	¦882
L I087	Invasin	1883
L1088	Iron **hydrogenase** 1	GAE
L1089	**Iron hydrogenase** 1	1884
L i090	**Iron hydrogenase** 1	1885
L109 1	Iron hydrogenase 1	¦886
L I092	Iron hydrogenase 1	1887
L1093	Iron hydrogenase 1	¦888
L1094	**Iron hydrogenase** 1	1889
L1095	Iron hydrogenase 1	1890
L1096	Iron hydrogenase 1	¦891
L I097	Iron hydrogenase 1	1892
L1098	Iron hydrogenase 1	¦893
L1099	Iron hydrogenase 1	1894
L¦ 100	Iron hydrogenase 1	1895
L I 10 1	Iron hydrogenase 1	¦896
L I 102	Iron **transport** protein	1897
L I 103	Isoilavanone **4'-0-methyltransferase**	¦898
L¦ 104	**Isoilavanone 4'-0-methyltransferase**	1899
L 1105	Junctional adhesion **molecule** 1	1900
L I 106	Junctional adhesion molecule 1	¦901

L1107	Junctional adhesion molecule 1	1902
L1108	Kanamycin nucleotidyltransferase	1903
L1109	Kanamycin nucleotidyltransferase	1904
L1110	Kanamycin nucleotidyltransferase	1905
L1111	Kanamycin nucleotidyltransferase	1906
L1112	Kelch-like protein 11	1907
L1113	Kexin	ISE
L1114	Kexin	1908
L1115	Kexin	1909
L1116	Kexin	1910
L1117	Kexin	1911
L1118	Kexin	1912
L1119	Kexin	1913
L1120	Kexin	1914
L1121	Ku70	1915
L1122	Ku70	1916
L1123	Ku70	1917
L1124	Ku70	1918
L1125	Ku80	1919
L1126	Laccase-1	1920
L1127	Laccase-1	1921
L1128	Laccase-1	1922
L1129	Laccase-1	1923
L1130	Laminin	DKC
L1131	L-aspartate dehydrogenase	SAS
L1132	L-aspartate dehydrogenase	1924
L1133	L-aspartate dehydrogenase	1925
L1134	Leucine dehydrogenase	1926
L1135	Leucine dehydrogenase	1927
L1136	Light chain of HyHel10 antibody fragment (fab)	1928
L1137	Lin2111 protein	1929
L1138	Lin2111 protein	1930
L1139	Lipopolysaccharide-responsive and beige-like anchor protein	1931
L1140	Lipopolysaccharide-responsive and beige-like anchor protein	1932
L1141	Lipovitellin (LV-1N, LV-1C)	1933
L1142	Lipovitellin (LV-1N, LV-1C)	1934
L1143	Lipovitellin (LV-1N, LV-1C)	1935
L1144	Lipovitellin (LV-1N, LV-1C)	1936
L1145	Lipovitellin (LV-1N, LV-1C)	1937
L1146	Lipoxygenase-1	1938
L1147	Lipoxygenase-1	1939
L1148	Low affinity immunoglobulin gamma Fc region receptor II-A	1940
L1149	Luciferase	1941
L1150	LysR-type regulatory protein	1942
L1151	Macrolide-specific efflux protein MacA	ATE

L I 152	Macrolide-specific efflux protein Mac A	1943
L I 153	Macrolide-specific efflux protein MacA	1944
L I 154	Magnesium transporter, putative	1945
L I 155	Main hemagglutinin component	1946
L I 156	Major centromere autoanrigen B	1947
L I 157	Major surface antigen p30	1948
L 1158	Major surface antigen p30	1949
L I 159	Major vault protein	1950
L I 160	Major vault protein	195 1
L I 16 1	Maltose phosphorylase	1952
L I 162	Maltose phosphorylase	1953
L 1163	Maltose phosphorylase	1954
L I 164	Maltose phosphorylase	1955
L I 165	Maltose phosphorylase	1956
L I 166	Manganese-dependent inorganic pyrophosphatase	1957
L I 167	Manganese-dependent inorganic pyrophosphatase	1958
L I 168	Mamian-binding lectin	1959
L I 169	Mannan-binding lectin	I960
L I 170	Mannan-binding lectin	1961
L I 17 1	Mannitol dehydrogenase	HNA
L I 172	Mannitol dehydrogenase	1962
L I 173	Membrane cofactor protein	RET
L I 174	Membrane cofactor protein	1963
L I 175	Membrane-associated prostaglandin E synthase-2	1964
L I 176	Membrane-associated prostaglandin E synthase-2	1965
L I 177	Membrane-associated prostaglandin E synthase-2	1966
L I 178	Membrane-associated prostaglandin E synthase-2	1967
L I 179	Membrane-associated prostaglandin E synthase-2	1968
L I 180	Membrane-bound lytic murein transglycosylase A	1969
L I 18 1	Methionyl-tRNA synthetase	1970
L I 182	Methyl-accepting chemotaxis protein	VRP
L I 183	Methyl-accepting chemotaxis protein	1971
L 1184	Methyl-accepting chemotaxis protein	1972
L 1185	Methyl-accepting chemotaxis protein	1973
L I 186	Methyl-coenzyme M reductase	1974
L I 187	Methyl-coenzy me M reductase	1975
L I 188	Methyl-coenzyme M reductase	1976
L I 189	Methyl-coenzyme M reductase	1977
L I 190	Methylene tetrahydromethanopterin dehydrogenase	1978
L I 19 1	Methylene tetrahydromethanopterin dehydrogenase	1979
L I 192	Mg2+- transporter MgtE	1980
L I 193	Mg2+ transporter M gtE	1981
L I 194	Mg2+ transporter MgtE	1982
L I 195	Mitochondrial aconitase	1983
L I 196	Mitochondrial aconitase	1984

L1197	Modification methylase TaqI	EGK
L1198	Modification methylase TaqI	PAT
L1199	Modification methylase TaqI	1985
L1200	Modification methylase TaqI	1986
L1201	Modification methylase TaqI	1987
L1202	Modification methylase TaqI	1988
L1203	Modification methylase TaqI	1989
L1204	Modification methylase TaqI	1990
L1205	Modification methylase TaqI	1991
L1206	Modification methylase TaqI	1992
L1207	Multidrug-efflux transporter 1 regulator	1993
L1208	Muramoyl-pentapeptide carboxypeptidase	1994
L1209	MutL	1995
L1210	MutL	1996
L1211	MutL	1997
L1212	MutL	1998
L1213	MutL	1999
L1214	MutL	2000
L1215	MutL	2001
L1216	MutL	2002
L1217	MutL	2003
L1218	MutM (Fpg) protein	2004
L1219	MutM (Fpg) protein	2005
L1220	MutM (Fpg) protein	2006
L1221	MutM (Fpg) protein	2007
L1222	Myotubularin-related protein 2	THW
L1223	Myotubularin-related protein 2	2008
L1224	Myotubularin-related protein 2	2009
L1225	Myotubularin-related protein 2	2010
L1226	Myotubularin-related protein 2	2011
L1227	Myotubularin-related protein 2	2012
L1228	N utilization substance protein A	EIP
L1229	N utilization substance protein A	2013
L1230	N utilization substance protein A	2014
L1231	N utilization substance protein A	2015
L1232	N-acetylglucosamine kinase	CAY
L1233	N-acetylglucosamine kinase	ISP
L1234	N-acetylglucosamine kinase	2016
L1235	N-acyl-D-glutamate deacylase	2017
L1236	N-acyl-D-glutamate deacylase	2018
L1237	N-acyl-D-glutamate deacylase	2019
L1238	N-acyl-D-glutamate deacylase	2020
L1239	N-acyl-D-glutamate deacylase	2021
L1240	N-acyl-D-glutamate deacylase	2022
L1241	N-acyl-D-glutamate deacylase	2023

L1242	**NAD -dependent malic enzyme**	2024
L1243	**NAD-dependent** malic enzyme	2025
L1244	**NADH** peroxidase	ADT
L1245	NADH peroxidase	**AVG**
L1246	NADH peroxidase	TLI
L1247	NADH peroxidase	2026
L1248	NADH peroxidase	2027
L1249	NADH peroxidase	2028
L1250	NADH peroxidase	2029
L1251	NADH peroxidase	2030
L1252	NADH peroxidase	2031
L1253	NADH **pyrophosphatase**	2032
L1254	Naphthalene 1,2-dioxygenase **alpha subunit**	2033
L1255	**Naphthalene** 1,2-dioxygenase alpha subunit	2034
L1256	**NEDD8-activaiing enzyme E1 catalytic** subunit	2035
L1257	**NEDD8-activating** enzyme **E1 regulatory subunit**	2036
L1258	NEDD8-activating **enzyme E1 regulatory** subunit	2037
L1259	NEDDS-activating enzyme E1 regulatory subunit	2038
L1260	**Nei** endonuclease **VIII-Like** 1	2039
L1261	**Nei** endonuclease VIII-Like 1	2040
L1262	**Nei** endonuclease VIII-Like 1	2041
L1263	**Nei endonuclease VIII-Like** 1	2042
L1264	Neural cell adhesion molecule 2	2043
L1265	Neural cell adhesion molecule 2	2044
L1266	Neural cell adhesion molecule 2	2045
L1267	Neural cell adhesion molecule 2	2046
L1268	Neural cell adhesion molecule 2	2047
L1269	Neuroplastin	2048
L1270	Neuroplastin	2049
L1271	Neuroplastin	2050
L1272	**Neutrophil cytosol factor** 1	2051
L1273	Nickel responsive regulator	2052
L1274	**NifU-like** protein 2, **chloroplast**	2053
L1275	Nitric oxide reductase	ILM
L1276	Nitric oxide reductase	2054
L1277	Nitric oxide reductase	2055
L1278	**Nitric** oxide reductase	2056
L1279	Nitric oxide reductase	2057
L1280	Nitric oxide reductase	2058
L1281	**NK** receptor	2059
L1282	Nuclear factor **of** activated **t-cells, cytoplasmic2**	2060
L1283	**Nucleolin RBD 12**	2061
L1284	**O-GlcNAcase** Nag.1	2062
L1285	Orange carotenoid protein	EGV
L1286	Orange carotenoid protein	2063

L1287	Orange **carotenoid protein**	2064
L1288	**Orn/Lys/Arg** decarboxylase family protein	j LEL
L1289	**Orn/Lys/Arg decarboxylase family** protein	j 2065
L1290	**Orn/Lys/Arg** decarboxylase **family** protein	2066
L1291	**Orn/Lys/Arg** decarboxylase family protein	j 2067
L1292	**Orn/Lys/Arg** decarboxylase family protein	2068
L1293	**Orn/Lys/Arg** decarboxylase family protein	j 2069
L1294	**Orn/Lys/Arg** decarboxylase **family** protein	j 2070
L1295	**Orn/Lys/Arg** decarboxylase family protein	2071
L1296	**Osteoclast-stimularing** factor **1**	j 2072
L1297	Oxygen-independent **coproporphyrinogen** III oxidase	j 2073
L1298	Oxygen-independent coproporphyrinogen III oxidase	j 2074
L1299	**Oxygen-independent** coproporphyrinogen III oxidase	j 2075
L1300	Oxygen-independent coproporphyrinogen III oxidase	2076
L1301	Oxygen-independent **coproporphyrinogen** III oxidase	j 2077
L1302	Oxygen-independent coproporphyrinogen III oxidase	2078
L1303	Oxygen-independent **coproporphyrinogen** III oxidase	j 2079
L1304	Oxygen-independent coproporphyrinogen III oxidase	j 2080
L1305	Oxygen-independent coproporphyrinogen III oxidase	2081
L1306	Oxygen-independent **coproporphyrinogen** III oxidase	j 2082
L1307	**Paraneoplastic encephalomyelitis** antigen **HuD**	2083
L1308	Paraneoplastic **encephalomyelitis** antigen **HuD**	j 2084
L1309	Penicillin binding protein **4**	j 2085
L13 10	Penicillin binding protein **4**	2086
L13 11	Penicillin binding protein **4**	j 2087
L13 12	Penicillin binding protein **4**	2088
L13 13	Penicillin binding protein **4**	j 2089
L13 14	Penicillin binding protein **4**	j 2090
L13 15	Penicillin binding protein **4**	2091
L13 16	**Peptide-N(4)-(N-ace1yl-beta-D-glucosaminyl)asparagine amidase** F	j DGV
L13 17	Peptide-N**(4)**-(N-acetyl-be!a-**D**-g1ucosaminyi)asparagine **amidase F**	2092
LL3 18	Peptide-N**(4)**-(N-acetyl-beta-D-ghicosafninyl)asparagine amidase F	j 2093
L13 19	**Peptide-N(4)-(N-ace1yl-beta-D-glucosaminyl)asparagine** amidase F	j 2094
L1320	Peroxisomal primary amine oxidase	j 2095
L132 1	Peroxisomal **primary** amine oxidase	j 2096
L1322	Peroxisome biogenesis factor **1**	j 2097
L1323	Pesticidial crystal protein **Cry2Aa**	j 2098
L1324	**Pesticidial** crystal protein **Cry2**Aa	j 2099
L1325	Pesticidial crystal protein **Cry2Aa**	j 2100
L1326	Phase **1** flagellin	j DLT
L1327	Phase **1** flagellin	2101
L1328	Phase **1 flagellin**	j 2102
L1329	Phase **1** flagellin	2103
L1330	Phase **1** flagellin	j 2104
L133 1	**Phase 1** flagellin	j 2105

L1332	Phase **1** flageliin	2106
L1333	Phase 1 flageliin	2107
L1334	**Phase 1** flageliin	2108
LI335	Phase 1 flageliin	2109
L1336	Phase i flageliin	2110
L1337	Phase **1** flageliin	**2111**
L1338	Phase 1 flageliin	2112
L1339	**Phenj**rialanj-1-tRNA synthetase beta **chain**	LGL
L1340	Phenyiaianyi-fRNA synthetase beta chain	**2113**
L134 I	**Phenylalanyl-tRNA** synthetase beta chain	2114
Li342	Phenylalaiiyl-tRNA **synthetase** beta chain	**2115**
L1343	**Phenylalanyl-tRNA** synthetase beta chain	2116
L1344	**Phenylalanyl-tRNA** synthetase beta chain	2117
Li345	Phenyiaianyi-fRNA synthetase beta chain	2118
L1346	Phenylalanyl-tRNA synthetase beta chain	2119
Li347	Pheriylaiaiiyl-tRNA **synthetase** beta chain	2120
LI348	**Phenylalanyl-tRNA** synthetase beta **chain**	**2121**
L1349	Phenyiaianyi-fRNA synthetase beta chain	2122
Li350	Phenyiaianyi-fRNA synthetase beta chain	2123
L135 I	Phenylalanyl-tRNA synthetase beta chain	2124
L1352	Pheriylaiaiiyl-tRNA **synthetase** beta chain	2125
L1353	**Phosphatase**	2126
L1354	Phosphatase	2127
Li355	**Phosphatase**	2128
L1356	**Phosphatidyiinositol** transfer proteirs Sec14p	YGT
LI357	**Phosphatidyl inositol** transfer protein **Secl4p**	2129
L1358	**Phosphatidyiinositol** transfer protein **Sec l4p**	2130
L1359	**Phosphatidylserine synthase**	2131
Li360	Phosphatidylserine synthase	2132
L136 1	Phosphatidylserine **synthase**	2133
LI362	Phosphogiycoiate phosphatase	2134
LI363	**Phosphogly colate** phosphatase	2135
L1364	Phosphogiycoiate **phosphatase**	2136
L1365	Phosphogiycoiate phosphatase	2137
Li366	**Phospholipase** D	2138
LI367	**Pho**spholipase D	2139
LI368	Phospholipase **D**	2140
Li369	Phosphoribosylamine—glycine ligase	**2141**
L1370	**Phosphoribosylamine—glycine** ligase	2142
L137 1	Phosphotransferase system, **enzyme** I	2143
LI372	Photo¾'stem II d1protease	2144
L1373	Photosystem II **dl** protease	2145
L1374	**Photosystem** 11 **dl** protease	2146
L1375	Photosystem II dl protease	2147
L1376	Photosystem II dl protease	2148

L1377	Phthalaie dioxygenase reductase	2149
L1378	P-hydroxybenzoate hydroxylase	DGL
L1379	P-hydroxybenzoate hydroxylase	IDL
L I380	P-hydroxybenzoate hydroxylase	RLK
L138 I	P-hydroxybenzoaie hydroxylase	2150
L1382	P-hydroxybenzoate hydroxylase	2151
L1383	P-hydroxybenzoate hydroxylase	2152
L1384	P-hydroxybenzoate hydroxylase	2153
L1385	P-hydroxybenzoate hydroxylase	2154
L1386	P-hydroxybenzoate hydroxylase	2155
L1387	P-hydroxybenzoate hydroxylase	2156
L1388	P-hydroxybenzoate hydroxylase	2157
L1389	P-hydroxybenzoate hydroxylase	2158
L1390	P-hydroxybenzoate hydroxylase	2159
L139 I	P-hydroxybenzoate hydroxylase	2160
L1392	P-hydroxybenzoate hydroxylase	2161
L1393	P-hydroxybenzoate hydroxylase	2162
L1394	P-hydroxybenzoate hydroxylase	2163
L1395	P-hydroxybenzoate hydroxylase	2164
L1396	P-hydroxybenzoate hydroxylase	2165
L1397	P-hydroxybenzoate hydroxylase	2166
L1398	Phytase	LNF
L1399	Phytase	QSN
L1400	Phytase	2167
L140 1	Phytase	2168
L I402	Phytase	2169
L1403	Phytase	2170
L1404	Phytase	2171
L1405	Phytase	2172
L I406	Phytase	2173
L I407	Phytase	2174
L1408	Pirin	LKS
L1409	Pirin	SGE
L1410	Pirin	2175
L141 1	Pirin	2176
L 14 12	Pirin	; 2177
L14L3	Pirin	2178
L14 14	Pirin	2179
L1415	Pirin	2180
L1416	Poly(A) polymerase	2181
L 14 17	Poly(A) polymerase	2182
L1418	Poly (A) polymerase	2183
L14 19	Poly (A) polymerase	2184
L1420	Poly (A) polymerase	2185
L142 1	Poly(A) polymerase	2186

L1422	**Poly(A)** polymerase	2187
L1423	Poly (A) polymerase	2188
L1424	Poly(A) polymerase	2189
LI425	Poly(A) polymerase	2190
L1426	Poly (A) polymerase	2191
L1427	**Poly(A)** polymerase	2192
L1428	Poly (rC)-binding protein 2	2193
L1429	Polymerase x	2194
L1430	Polymerase x	2195
L143 I	Polypeptide **N-acetylgalactosaminyltransferase** 2	2196
L1432	Polypeptide N-acetylgalactosaminyltraiisferase 2	2197
L1433	Polyphosphate kinase	2198
L1434	**Polyphosphate** kinase	2199
L1435	Polyphosphate kinase	2200
L1436	**Polypyrimidine tract-binding** protein	2201
L1437	Porcine **pancreatic** spasmolytic **polypeptide**	2202
L1438	Possible **3-mercaptopyruvate** sulfurtransferase	LFR
L1439	Possible **3-mercaptopyruvate sulfurtransferase**	**YGM**
L1440	Possible 3-mercaptopyruvate sulfurtransferase	2203
L144 I	Possible 3-mercaptopyruvate **sulfurtransferase**	2204
L1442	Possible **3-mercaptopyruvate** sulfurtransferase	2205
L1443	Postsynaptic density protein 95	2206
L1444	Postsynaptic **density** protein 95	2207
L1445	Predicted sugar **phosphatases** of the HAD **superfamily**	IAI
L1446	Predicted sugar phosphatases of the HAD superfamily	2208
LI447	Predicted **sugar phosphatases** of the HAD **superfamily**	2209
L1448	Predicted sugar phosphatases of the HAD superfamily	2210
L1449	Predicted sugar phosphatases **of** the HAD **superfamily**	2211
L1450	Predicted sugar **phosphatases** of the HAD **superfamily**	2212
L145 1	Predicted sugar phosphatases of the HAD **superfamily**	2213
LI452	Predicted **sugar phosphatases** of the HAD **superfamily**	2214
L1453	Predicted sugar phosphatases of the HAD **superfamily**	2215
L1454	Preprotein **translocase** SecA	**ITF**
L1455	Preprotein translocase SecA	LID
L1456	Preprotein translocase SecA	2216
LI457	Preprotein **translocase** SecA	2217
L1458	Preprotein translocase SecA	2218
L1459	Preprotein **translocase** SecA	2219
L1460	Preprotein translocase SecA	2220
LI**461**	Preprotein translocase SecA	2221
L1462	Preprotein translocase SecA	2222
L1463	Preprotein translocase SecA	2223
L1464	Preprotein **translocase** SecA	2224
L1465	Preprotein translocase SecA	2225
LI**466**	Preprotein translocase SecA	2226

L1467	Preprotein translocase SecA	2227
L1468	Preprotein translocase SecA	2228
L1469	Preprotein translocase SecA	2229
L1470	Preprotein translocase SecA	2230
L1471	Preprotein translocase SecA	2231
L1472	Preprotein translocase SecA	2232
L1473	PrfA	ING
L1474	Probable 16s rRNA-processing protein RimM	2233
L1475	Probable biphenyl-2,3-diol 1,2-dioxygenase BphC	2234
L1476	Probable chorismate mutase	LLA
L1477	Probable chorismate mutase	2235
L1478	Probable chorismate mutase	2236
L1479	Probable ferredoxin-dependent nitrite reductase NirA	VPL
L1480	Probable ferredoxin-dependent nitrite reductase NirA	WGI
L1481	Probable ferredoxin-dependent nitrite reductase NirA	2237
L1482	Probable ferredoxin-dependent nitrite reductase NirA	2238
L1483	Probable ferredoxin-dependent nitrite reductase NirA	2239
L1484	Probable ferredoxin-dependent nitrite reductase NirA	2240
L1485	Probable ferredoxin-dependent nitrite reductase NirA	2241
L1486	Probable ferredoxin-dependent nitrite reductase NirA	2242
L1487	Probable ferredoxin-dependent nitrite reductase NirA	2243
L1488	Probable ferredoxin-dependent nitrite reductase NirA	2244
L1489	Probable ferredoxin-dependent nitrite reductase NirA	2245
L1490	Probable ferredoxin-dependent nitrite reductase NirA	2246
L1491	Probable ferredoxin-dependent nitrite reductase NirA	2247
L1492	Probable ferredoxin-dependent nitrite reductase NirA	2248
L1493	Probable galactokinase	2249
L1494	Probable galactokinase	2250
L1495	Probable galactokinase	2251
L1496	Probable galactokinase	2252
L1497	Probable galactokinase	2253
L1498	Probable galactokinase	2254
L1499	Probable galactokinase	2255
L1500	Probable galactokinase	2256
L1501	Probable galactokinase	2257
L1502	Probable galactokinase	2258
L1503	Probable galactokinase	2259
L1504	Probable galactokinase	2260
L1505	Probable glutathione S-transferase	2261
L1506	Probable GST-related protein	2262
L1507	Probable HPr(Ser) kinase/phosphatase	2263
L1508	Probable thiosulfate sulfur transferase	2264
L1509	Probable thiosulfate sulfur transferase	2265
L1510	Probable thiosulfate sulfur transferase	2266
L1511	Probable thiosulfate sulfur transferase	2267

L15 12	Probable thiosulfate **sulfur** transferase	2268
L15 13	Probable **thiosulfate** sulfur transferase	j 2269
L15 14	Probable thiosulfate sulfur transferase	j 2270
L15 15	Probable **thiosulfate** sulfur transferase	227 **i**
L15 16	Probable **tRNA pseudouridine** synthase D	j 2272
L15 17	Probable tRNA pseudouridine synthase **D**	2273
L15 18	Probable tRNA pseudouridine synthase D	j 2274
L15 19	Probable tRNA pseudouridine synthase D	j 2275
L1520	Probable tRNA pseudouridine synthase D	2276
L152 I	Probable **tRNA** pseudouridine synthase D	j 2277
L1522	Programed cell death protein 8	SKE
L1523	Programed cell death protein 8	j TLQ
L1524	Programed cell death protein 8	j 2278
L1525	Programed cell death protein 8	2279
L1526	**Programed** cell death protein 8	j 2280
L1527	Programed cell death protein 8	2281
L1528	**Programed** cell death protein 8	2282
L1529	Programed cell death protein 8	j 2283
Li530	Programed cell death protein 8	2284
L153 I	**Programed** cell death protein 8	j 2285
L1532	Programed cell death protein 8	2286
L1533	Programed cell death protein 8	2287
L1534	Programed cell death protein 8	j 2288
L1535	Programed cell death protein 8	2289
L1536	**Programed** cell death protein 8	j 2290
L1**537**	Programed cell death protein 8	**229 1**
L1538	Programed cell death protein 8	! 2292
L1539	Programed cell death protein 8	j 2293
L1540	Programed cell death protein 8	2294
L154 1	Programed cell death protein 8	j 2295
L1**542**	Proline oxidase	2296
L1543	**Prolyl-tRNA synthetase**	**j** 2297
L1544	Prostaglandin G/H synthase 1	j PEI
L1545	Prostaglandin G/H **synthase** 1	j 2298
L1546	Protease	j 2299
L1**547**	Protease	2300
L1548	Protease	j 2301
L1549	Protease DegS	2302
L1550	Protease DegS	j 2303
L155 1	Protease DegS	j 2304
L1**552**	Protease DegS	2305
L1553	Protease III	j NAR
L1554	Protease III	RNP
L1555	Protease III	j 2306
L1556	Protease III	j 2307

L1557	Protease III	2308
L1558	Protease III	2309
L1559	Protease III	2310
L1560	Protease III	2311
L1561	Protease III	2312
L1562	Protease III	2313
L1563	Protease III	2314
L1564	Protease III	2315
L1565	Protease III	2316
L1566	Protease HI	2317
L1567	Protease III	2318
L1568	Protease III	2319
L1569	Protease III	2320
L1570	Protease III	2321
L1571	Protease III	2322
L1572	Protease III	2323
L1573	Protease III	2324
L1574	Protease III	2325
L1575	Protection of telomeres 1	2326
L1576	Protection of **telomeres** 1	2327
L1577	**Protein** (CD58)	2328
L1578	Protein (CRP1)	2329
L1579	Protein (DNA polymerase)	2330
L1580	Protein (DNA polymerase)	2331
L1581	Protein (DNA polymerase)	2332
L1582	Protein (electron transfer **flavoprotein)**	2333
L1583	Protein (electron transfer **flavoprotein)**	2334
L1584	Protein (Ffh)	2335
L1585	Protein (Ffh)	2336
L1586	Protein (Fill)	2337
L1587	Protein (Ffh)	2338
L1588	Protein (Ffh)	2339
L1589	Protein (Fold restriction endonuclease)	2340
L1590	Protein (Fokl restriction endonuclease)	2341
L1591	Protein (Fokl restriction endonuclease)	2342
L1592	Protein (Fokl restriction endonuclease)	2343
L1593	Protein (Fokl restriction endonuclease)	2344
L1594	Protem (Fokl restriction endonuclease)	2345
L1595	Protein (Fokl restriction endonuclease)	2346
L1596	Protein (Fokl restriction endonuclease)	2347
L1597	Protein (Fokl **restriction endonuclease)**	2348
L1598	Protein (neural cell adliesion molecule)	2349
L1599	Protem (neural cell adhesion molecule)	2350
L1600	Protein (neural cell adhesion molecule)	2351
L1601	Protein (nine-haem **cytochrome** c)	FTH

L1602	Protein **(nine-haem** cytochrome c}	2352
L1603	Protein **(nine-haem** cytochrome c)	2353
L1**604**	Protein (nine-haem **cytochrome c)**	2354
L1605	Protein **(nine-haem** cytochrome c)	2355
L1606	Protein (nine-haem cytochrome c)	2356
L1607	Protein **(nine-haem** cytochrome c}	2357
L1608	Protein **(nine-haem** cytochrome c)	2358
L1**609**	Protein (nine-haem **cytochrome** c)	2359
L1610	Protein **(protease/helicase** NS3)	2360
L1611	Protein **(protease/helicase NS3)**	2361
L1612	Protein (protease/helicase **NS3)**	2362
L1613	Protein **(protease/helicase NS3)**	2363
L1614	Protein disulfide oxidoreductase	2364
L1615	Protein disulfide **oxidoreductase**	2365
L1616	Protein **disulfide-isomerase** A4	2366
L1617	**Protein** kinase PKR	2367
L1618	Protein kinase PKR	2368
L1619	Protein TolB	**VNK**
L1620	Protein TolB	2369
L1621	Protein TolB	2370
L1622	Protein TolB	2371
L1623	Protein TolB	2372
L1624	Protein TolB	2373
L1625	Protein TolB	2374
L1626	Protein translation elongation factor 1A	2375
L1627	Protein transport protein **Sec24**	**DRN**
L1628	Protein transport protein **Sec24**	2376
L1629	Protein transport protein Sec24	2377
L1630	Protein transport protein **Sec24**	2378
L1631	Protein transport protein **Sec24**	2379
L1632	Protein transport protein **Sec24**	2380
L1633	Protein transport protein **Sec24**	2381
L1634	Protein transport protein Sec24	2382
L1635	Protein transport protein Sec24	2383
L1636	**Pseudouridine** synthase CBF5	AIQ
L1637	**Pseudouridine** synthase CBF5	2384
L1638	Pseudouridine synthase **CBF5**	2385
L1639	**Putative acetylglutamate** synthase	2386
L1640	Putative **acetylglutamate** synthase	2387
L1641	Putative acetylglutamate synthase	2388
L1642	Putative family 31 glucosidase **Yicl**	2389
L1643	Putative **family** 31 **glucosidase Yicl**	2390
L1644	**Putative** family 31 glucosidase Yicl	2391
L1645	Putative glutathione transferase	2392
L1**646**	Putative **glutathione** transferase	2393

L1647	**Putative glutathione** transferase	2394
L1648	Putative **GNTR-family** transcriptional regulator	2395
L1649	Putative **GNTR-family** transcriptional regulator	2396
L I**650**	Putative GNTR-family **transcriptional regulator**	; 2397
L165 I	**Putative HTH-type transcriptional** regulator **PH006** l	2398
L1652	Putative HTH-type transcriptional regulator PH151 9	2399
L1653	Putative HTH-type **transcriptional** regulator PH15 19	2400
L1654	Putative nietallopeptidase	2401
L1655	Putative N-acetyimaimosamine kinase	2402
L1656	Putative **N-acetyltnannosamine** kinase	2403
L1657	Putative **N-acetylmannosaniine** kinase	2404
L1658	Putative **NADP** oxidoreductase BF3 122	2405
L1659	Putative NADP oxidoreductase BF3 122	2406
L1660	Putative NADP oxidoreductase BF3122	2407
L1661	Putative NADP oxidoreductase BF3 122	2408
L1662	Putative oxidoreductase	2409
L1663	Putative secreted **alpha-galactosiclase**	**PLP**
L1664	Putative secreted **alpha-galactosidase**	TNG
L1665	Putative secreted alpha-galactosidase	2410
L1666	Putative secreted **alpha-galactosidase**	**24**1 **1**
L1667	Putative secreted alpha-galactosidase	2412
L1668	Putative **tagalose-6-phosphate** ketose/aldose isomerase	DKA
L1669	Putative **tagatose-6-phosphate** ketose/aldose isoinerase	2413
L1670	Putative tagatose-6-phosphate ketose/aldose isomerase	2414
L167 i	Putative **tagatose-6-phosphate** ketose/aldose isomerase	24 5
L I**672**	Putative transcriptional regulator **GntR**	2416
L1673	Putative transcriptional repressor **(TetR/AcrR** family)	**KFR**
L1674	Putative transcriptional repressor **(TetR/AcrR** family)	2417
L1675	Putative **uncharacterized** protein	2418
L I**676**	Putative uncharacterized protein	2419
L I**677**	Putative uncharacterized protein	2420
L1678	Putative uncharacterized protein	2421
L1679	Putative uncharacterized protein	2422
L1680	Putative uncharacterized protein	2423
L168 1	Putative uncliaracterized protein	2424
L I**682**	Putative uncharacterized protein	2425
L1683	Putative uncharacterized protein	2426
L1684	Pyruvate decarboxylase	CAA
L1685	Pyruvate decarboxylase	2427
L1686	**Pyruvate** decarboxylase	2428
L I**687**	Pyruvate decarboxylase	2429
L1688	Pyravate decarboxylase	2430
L1689	Pyruvate decarboxylase	243 1
L1690	Pyruvate dehydrogenase [lipoamide] kinase isozyme 2, mitochondrial	YVP
L169 1	**Pyruvate** dehydrogenase [lipoamide] kinase isozyme 2, mitochondrial	2432

L1692	Pyruvate dehydrogenase **[lipoamide]** kinase isozyme 2, **mitochondrial**	2433
L1693	**Pyruvate** dehydrogenase E1 component **subunit** beta, mitochondrial	2434
L1694	**Pyruvate** dehydiOgenase E1 component subunit beta, mitochondrial	2435
L I695	Pyruvate dehydrogenase **E l component** subunit beta, mitochondrial	2436
L1696	**Pyruvate** phosphate **dikinase**	FNP
L1697	Pyruvate phosphate dikinase	SAL
L1698	**Pyruvate** phosphate dikinase	2437
L1699	Pyruvate **phosphate** dikinase	2438
L1700	**Pyruvate** phosphate dikinase	2439
L170 1	**Pyruvate** phosphate dikinase	2440
L1702	Pyruvate phosphate dikinase	2441
L1703	Pyruvate phosphate dikinase	2442
L1704	Pyruvate phosphate dikinase	2443
L1705	**Pyruvate** phosphate dikinase	2444
L1706	**Pyruvate** phosphate dikinase	2445
L1707	Pyruvate phosphate dikinase	2446
L1708	**Py ruvate-ferredoxin oxidoreductase**	VRL
L1709	**Pyruvate-ferredoxin** oxidoreductase	2447
L1710	Pyravate-ferredoxin oxidoreductase	2448
L171 I	**Py ruvate-ferredoxin oxidoreductase**	2449
L 17 12	**Pyruvate-ferredoxin oxidoreductase**	2450
L1713	**Py ruvate-ferredoxin** oxidoreductase	2451
L1714	Pyruvate-ferredoxin oxidoreductase	2452
L17 15	Pyruvate-ferredoxin oxidoreductase	2453
L1716	**Py** ruvate-ferredoxin oxidoreductase	2454
L 17 17	**Pyruvate-ferredoxin** oxidoreductase	2455
L1718	**Py** ruvate-ferredoxin oxidoreductase	2456
L1719	Pyruvate-ferredoxin oxidoreductase	2457
L1720	**Pyruvate-ferredoxin** oxidoreductase	2458
L172 1	**Pyruvate-ferredoxin** oxidoreductase	2459
L I722	**Pyruvate-ferredoxin** oxidoreductase	2460
L1723	Py ruvate-ferredoxin oxidoreductase	2461
L1724	Pyruvate-ferredoxin oxidoreductase	2462
L1725	Pyruvate-ferredoxin oxidoreductase	2463
L I726	**Pyruvate-ferredoxin** oxidoreductase	2464
L I727	**Pyruvate-ferredoxin** oxidoreductase	2465
L1728	**Quinohemoprotein amine** dehydrogenase 60 **kDa subunit**	2466
L1729	**Quinohemoprotein** amine dehydrogenase 60 kDa subunit	2467
L1730	Quinohemoprotein amine dehydrogenase 60 kDa subunit	2468
L173 1	Quinohemoprotein amine dehydrogenase 60 kDa subunit	2469
L I732	Quinohemoprotein amine **dehydrogenase** 60 kDa subunit	2470
L1733	Quinohemoprotein **amine** dehydrogenase 60 kDa **subunit**	2471
L1734	**Quinohemoprotein** amine dehydrogenase 60 kDa subunit	2472
L1735	Quinohemoprotein amine dehydrogenase 60 kDa subunit	2473
L1736	Quinohemoprotein amine dehydrogenase 60 kDa subunit	2474

L1737	Quinohemoprotein amine dehydrogenase 60 kDa subunit	2475
L1738	Rag1	2476
L1739	Rag1	2477
L1740	Receptor-type tyrosine-protein phosphatase Mu	2478
L1741	Receptor-type tyrosine-protein phosphatase Mu	2479
L1742	RecG	2480
L1743	RecG	2481
L1744	RecG	2482
L1745	RecG	2483
L1746	RecG	2484
L1747	RecG	2485
L1748	RecG	2486
L1749	RecG	2487
L1750	RecG	2488
L1751	RecG	2489
L1752	RecG	2490
L1753	RecG	2491
L1754	Recombination endonuclease VII	2492
L1755	Recombining binding protein suppressor of hairless	2493
L1756	Restriction endonuclease	ERV
L1757	Restriction endonuclease	2494
L1758	Restriction endonuclease	2495
L1759	Restriction endonuclease	2496
L1760	Retinaldehyde-binding protein 1	QYP
L1761	Retinaldehyde-binding protein 1	2497
L1762	Retinaldehyde-binding protein 1	2498
L1763	Retinoblastoma pocket	2499
L1764	RfcS	ITD
L1765	RfcS	LTE
L1766	RfcS	2500
L1767	RfcS	2501
L1768	RfcS	2502
L1769	RfcS	2503
L1770	RfcS	2504
L1771	Rhamnogalacturonase B	2505
L1772	Rhamnogalacturonase B	2506
L1773	Rhamnogalacturonase B	2507
L1774	Rhamnogalacturonase B	2508
L1775	Rhamnogalacturonase B	2509
L1776	Rhodniin	2510
L1777	Rhodniin	2511
L1778	Riboflavin synthase	2512
L1779	Ribonuclease D	2513
L1780	Ribonuclease D	2514
L1781	Ribonuclease D	2515

L1782	Ribonuciease TTHA0252	2516
L1783	Ribonuclease TTHA0252	2517
L1784	Ribonuciease TTHA0252	2518
L1785	Ribonuciease TTHA0252	2519
L1786	**Ribonuciease** TTHA0252	2520
L1787	**Ribonuciease** TTHA0252	2521
L1788	Ribonucleotide reductase rl protein	2522
L1789	Ribonucleotide reductase rl protein	2523
L1790	Ribonucleotide reductase rl protein	2524
L1791	Ribonucleotide reductase **rl** protein	2525
L1792	Ribonucleotide reductase **rl** protein	2526
L1793	Ribonucleotide reductase rl protein	2527
L1794	Ribosome maturation factor **RimM**	2528
L1795	**Ribulose-1,5** bisphosphate carboxylase/oxygenase large **subunit N-methyltransferase**	RHA
L1796	**Ribulose-1,5** bisphosphate carboxylase/oxygenase large **subunit** N-methyltransferase	2529
L1797	Rigid extended **P-rich**	2530
L1798	**Rigid extended P-rich**	2531
L1799	Rigid extended P-rich	2532
L1800	Rigid extended P-rich	2533
L1801	Rigid extended **P-rich**	2534
L1802	Rigid extended P-rich	2535
L1803	**Rigid** extended P-rich	2536
L1804	Rigid extended P-rich	2537
L1805	Rigid extended P-rich	2538
L1806	Rigid extended P-rich	2539
L1807	Rigid extended P-rich	2540
L1808	**Rigid** extended P-rich	2541
L1809	Rigid extended P-rich	2542
L1810	Rigid extended P-rich	2543
L1811	Rigid extended P-rich	2544
L1812	Rigid helical	2545
L1813	Rigid helical	2546
L1814	Rigid helical	2547
L1815	Rigid helical	2548
L1816	Rigid helical	2549
L1817	Rigid helical	2550
L1818	Rigid helical	2551
L1819	Rigid helical	2552
L1820	RNA binding domain of rho transcription termination factor	2553
L1821	RNA binding protein ZFa	2554
L1822	Rob transcription factor	2555
L1823	Rob transcription lactor	2556
L1824	RP2 lipase	2557
L1825	Rubrerythrin	2558
L1826	**S-adenosylmethionine** synthetase	2559

L1827	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	QFD
L1828	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2560
L1829	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2561
L1830	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2562
L1831	Sarcoplasmic/endoplasmic reiiculum calcium ATPase 1	2563
L1832	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2564
L1833	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2565
L1834	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2566
L1835	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2567
L1836	Sarcoplasmic/endoplasmic reiiculum calcium ATPase 1	2568
L1837	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2569
L1838	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2570
L1839	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2571
L1840	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2572
L1841	Sarcoplasmic/endoplasmic reiiculum calcium ATPase 1	2573
L1842	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2574
L1843	Sarcoplasmic/endoplasmic reiiculum calcium ATPase 1	2575
L1844	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2576
L1845	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2577
L1846	Sarcoplasmic/endoplasmic reiiculum calcium ATPase 1	2578
L1847	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2579
L1848	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2580
L1849	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2581
L1850	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2582
L1851	Sarcoplasmic/endoplasmic reiiculum calcium ATPase 1	2583
L1852	Sarcoplasmic/endoplasmic reticulum calcium ATPase 1	2584
L1853	Scavenger mRNA-decapping enzyme DcpS	ETC
L1854	Scavenger mRNA-decapping enzyme DcpS	NIT
L1855	Scavenger mRNA-decapping enzyme DcpS	2585
L1856	Scavenger mRNA-decapping enzyme DcpS	2586
L1857	Sec18p (residues 22 - 210)	2587
L1858	SeclSp (residues 22 - 210)	2588
L1859	Sensor protein	2589
L1860	Sensor protein	2590
L1861	Septum site-determining protein MinC	2591
L1862	Serine acetyltransferase	2592
L1863	Serine protease/NTPase/helicase NS3	2593
L1864	Serine protease/NTPase/helicase NS3	2594
L1865	Serine protease/NTPase/helicase NS3	2595
L1866	Serine rich linker	2596
L1867	Serine rich linker	2597
L1868	Serine rich linker	2598
L1869	Serine rich linker	2599
L1870	Serine rich linker	2600
L1871	Serine rich linker	2601

L1872	Serine rich linker	2602
L1873	Seryi-tRNA synthetase	2603
LI874	Siaiidase	2604
LI875	Sialidase B	SLT
L1876	Siaiidase B	VRE
L1877	Sialidase B	2605
L1878	Siaiidase B	2606
LI879	Siaiidase B	2607
L1880	Sialidase B	2608
L1881	Siaiidase B	2609
Li882	Sialidase B	2610
L1883	Signal peptidase I	SRR
L1884	Signal peptidase I	2611
L1885	Signal peptidase I	2612
L1886	Signal peptidase I	2613
Li887	Signal peptidase I	2614
L1888	Signal peptidase I	2615
L1889	Signal peptidase I	2616
Li890	Signal peptidase I	2617
L1891	Signal peptidase I	2618
Li892	Signal peptidase I	2619
L1893	Signal peptidase I	2620
Li894	Signal recognition particle protein	2621
Li895	Signal transducer and activator of transcription 1-alpha/beta	NDE
L1896	Signal transducer and activator of transcriptionl-alpha/beta	SSF
LI897	Signal transducer and activator of transcription 1-alpha/beta	2622
L1898	Signal transducer and activator of transcriptionl-alpha/beta	2623
Li899	Signal transducer and activator of transcriptionl-alpha/beta	2624
Li900	Signal transducer and activator of transcriptionl-alpha/beta	2625
Li901	Signal transduction protein CBL	2626
LI902	Signal transduction protein CBL	2627
L1903	Similar to RAD54-hke	AKP
Li904	Similar to RAD54-like	EYF
Li905	Similar to RAD54-like	RFE
Li906	Similar to RAD54~like	2628
LI907	Similar to RAD54-like	2629
L1908	Similar to RAD54-like	2630
Li909	Similar to RAD54-like	2631
L1910	Similar to RAD54-like	2632
L1911	Similar to RAD54-like	2633
L1912	Similar to RAD54-like	2634
L1913	Similar to RAD54-like	2635
L1914	Similar to RAD54-like	2636
L1915	Similar to RAD54-like	2637
L1916	SKD1 protein	LMQ

L1917	SKD1 protein	2638
L1918	SKD1 protein	2639
L1919	SKD1 protein	2640
LI920	SKD1 protein	2641
L1921	SKD1 protein	2642
L1922	Sll1358 protein	2643
L1923	Sll1358 protein	2644
LI924	Sll1358 protein	2645
L1925	Sll1358 protein	2646
L1926	Soluble IFN alpha/beta receptor	2647
Li927	Soluble IFN alpha/beta receptor	2648
L1928	Sporozoite-specific SAG protein	2649
L1929	Staphylococcal accessory regulator a homologue	2650
Li930	Staphylococcal nuclease domain-containing protein 1	2651
L1931	Staphylococcal nuclease domain-containing protein 1	2652
L1932	Staphylococcal nuclease domain-containing protein 1	2653
L1933	Staphylococcal nuclease domain-containing protein 1	2654
L1934	Staphylococcal nuclease domain-containing protein 1	2655
Li935	Staphylococcal nuclease domain-containing protein 1	2656
L1936	Stat protein	2657
Li937	Stat protein	2658
L1938	Stat protein	2659
L1939	Stat protein	2660
Li940	Stat protein	2661
L1941	Stat protein	2662
LI942	Stat protein	2663
L1943	Stat protein	2664
Li944	Stat protein	2665
Li945	Stat protein	2666
LI946	Stat protein	2667
LI947	Stat protein	2668
L1948	Stat protein	2669
L1949	Stat protein	2670
L1950	Stat protein	2671
LI951	Subtiiisin-like protease	2672
LI952	Succinyl-CoA ligase [GDP-forming] alpha-chain, mitochondrial	2673
L1953	Succinyl-CoA ligase [GDP-forming] alpha-chain, mitochondrial	2674
Li954	Succinyl-CoA ligase jGDP-forming] alplia-chain, mitochondrial	2675
L1955	Succinyl-CoA ligase [GDP-forming] alpha-chain, mitochondrial	2676
Li956	Succinyl-CoA ligase [GDP-forming] alplia-chain, mitochondrial	2677
LI957	Succinyl-CoA ligase [GDP-forming] alpha-chain, mitochondrial	2678
L1958	Succinyl-CoA synthetase beta chain	ADG
Li959	Sisccinyi-CoA synthetase beta chain	RQP
Li960	Succinyl-CoA synthetase beta chain	2679
Li961	Succinyl-CoA synthetase beta chain	2680

L1962	Succinyl-CoA **synthetase** beta chain	2681
L1963	**Succinyl-CoA** synthetase beta chain	2682
L1964	Succinyl-CoA synthetase beta **chain**	2683
L1965	Succinyl-CoA synthetase beta chain	2684
L1966	**Succinyl-CoA :3-ketoacid-coenzyme** A transferase	2685
L1967	**Sulfurtransferase**	2686
L1968	Superantigen 5MEZ-2	2687
L1969	Superoxide disniutase 1 copper chaperone	2688
L1970	Surface layer protein	2689
L1971	Surface **layer protein**	2690
L1972	**Surface** layer protein	2691
L1973	Surface layer protein	2692
L1974	Surface layer protein	2693
L1975	Surface layer protein	2694
L1976	Surface layer protein	2695
L1977	**Surface** layer protein	2696
L1978	T lymphocyte activation antigen	2697
L1979	T **lymphocyte** activation antigen	2698
L1980	T-cell receptor **alpha** chain C region	2699
L1981	Terminal oxygenase component of carbazoSe	2700
L1982	Tetanus neurotoxin	2701
L1983	Tetracycline repressor protein class D	2702
L1984	The **GTP-binding** protein Obg	2703
L1985	The GTP-binding protein Obg	2704
L1986	The GTP-binding protein Obg	2705
L1987	The GTP-binding protein Obg	2706
L1988	Thioredoxin **domain-containing** protein 4	2707
L1989	Thioredoxin domain-containing protein 4	2708
L1990	**Thiosulfate** sulfurtransferase	IDP
L1991	**Thiosulfate** sulfurtransferase	2709
L1992	**Thiosulfate sulfurtransferase**	2710
L1993	Thiosulfate **sulfurtransferase**	2711
L1994	Thiosulfate sulfurtransferase	2712
L1995	**Threonyl-tRNA** synthetase	2713
L1996	Threonyl-tRNA synthetase	2714
L1997	Threonyl-tRNA **synthetase**	2715
L1998	Threonyl-tRNA synthetase	2716
L1999	**Threonyl-tRNA** synthetase	2717
L2000	**Threonyl-tRNA** synthetase	2718
L2001	Threonyl-tRNA synthetase	2719
L2002	Threonyl-tRNA synthetase	2720
L2003	Threonyl-tRNA synthetase	2721
L2004	**Threonyl-tRNA** synthetase 1	2722
L2005	Threonyl-tRNA synthetase 1	2723
L2006	Threonyl-tRNA synthetase 1	2724

L2007	Threonyl-tRNA synthetase i	; 2725
L2008	**Threonyl-tRNA** synthetase 1	j 2726
L2009	Threonyl-tRNA synthetase 1	j 2727
L20 10	**Tlireonyl-tRNA** synthetase i	2728
L201 I	**Threonyl-tRNA** synthetase 1	2729
L20 12	Thxombospondin 1	2730
L2013	Tick-bome encephalitis virus glycoprotein	j 273 I
L2014	Titin	j 2732
L20 15	Titin	2733
L2016	**TLR1789** protein	j 2734
L20 17	**TLR 1789 protein**	j 2735
L2018	**Topoisomerase** I	j 2736
L2019	Topoisomerase I	! 2737
L2020	Toxic shock syndrome toxin-1	2738
L202 I	Toxic shock syndrome toxin-!	j 2739
L2022	Toxic shock syndrome toxin-1	2740
j.2023	Toxic shock syndrome toxin-1	j 2741
L2024	**T-plasminogen** activator Fl-G	j **VPV**
L2025	**T-plasminogen** activator Fl-G	j 2742
L2026	TpsB transporter FhaC	j 2743
L2027	**TpsB** transporter FhaC	2744
L2028	TpsB transporter FhaC	j 2745
L2029	Transcarbamylase	j 2746
L2030	**Transcarb**amylase	2747
L203 I	Transcription **antiterminator LicT**	j 2748
L2032	Transcription elongation factor **GreB**	2749
L2033	Transcription initiation factor IIa gamma chain	j 2750
L2034	Transcription initiation factor lib	j 2751
L2035	Transcription initiation factor lib	2752
L2036	Transcriptional **regulator** (NtrC family)	j 2753
L2037	**Transcriptional** regulator **AefR**	2754
L2038	**Transcriptional regulator AefR**	2755
L2039	Transcriptional regulator AefR	j 2756
L2040	Transcriptional regulator AefR	j 2757
L204 1	Transcriptional regulator AefR	j 2758
L2042	**Transcriptional** regulator, **AsnC family**	2759
L2043	Transcriptional regulator, AsnC family	j 2760
L2044	**Transcriptional** regulator, AsnC family	276 I
L2045	Transcriptional regulator, biotin repressor **family**	j 2762
L2046	Transcriptional regulator, **Crp/Fnr family**	j 2763
L2047	**Transcriptional** regulator, **GntR family**	2764
L2048	Transcriptional regulator, HTH_**3 family**	j 2765
L2049	**Transcriptional** regulator, HTH_3 family	2766
L2050	Transcriptional regulator, **HTH_3** family	j 2767
L205 1	Transcriptional regulator, HTH_3 family	j 2768

L2052	**Transcriptional** regulator, **HTH_3** family	**2769**
L2053	**Transcriptional** regulator, laci family	**2770**
L2054	Transcriptional **regulatory** protein ZraR	2771
L2055	**Transcriptional** regulatory protein ZraR	**2772**
L2056	Transcriptional **regulatory** protein ZraR	**2773**
L2057	**Transcriptional** regulatory protein ZraR	**2774**
L2058	Transcriptional regulatory protein ZraR	**2775**
L2059	Transcriptional regulatory protein ZraR	**2776**
L2060	Transcriptional regulatory protein ZraR	**2777**
L206 I	Transferrin receptor protein	**VSN**
L2062	Transferrin receptor protein	**2778**
L2063	Transferrin receptor protein	**2779**
L2064	Transferrin receptor protein	**2780**
L2065	**Transferrin** receptor protein	**2781**
L2066	Translation initiation factor 5A	**2782**
L2067	Translation **initiation** factor 5A	**2783**
L2068	Translation initiation factor **5**A	**2784**
L2069	Translation initiation factor IF2/eIF5b	**2785**
L2070	Translation initiation factor IF2/eIF5b	**2786**
L207 i	**Transposable** element mariner, cosnpiete CDS	**2787**
L2072	Tricorn protease	**2788**
L2073	Tricorn protease	**2789**
L2074	Tricom protease	**2790**
L2075	Trigger factor	**279** I
L2076	Trigger factor	**2792**
L2077	Trigger factor	**2793**
L2078	**TRNA CCA-adding** enzyme	RR!
L2079	TRNA CCA-adding enzyme	**2794**
L2080	TRNA CCA-adding enzyme	**2795**
L208 1	TRNA CCA-adding enzyme	**2796**
L2082	TRNA CCA-adding **enzyme**	**; 2797**
L2083	TRNA nucleotidyltransferase	**2798**
L2084	TRNA-splicing endonuciease	**2799**
L2085	**Ttl467** protein	LEA
L2086	**TU467** protein	**2800**
L2087	Tumor **suppressor p53-binding** protein 1	**280 1**
L2088	Tumor suppressor **p53**-binding protein 1	**2802**
L2089	Tumor **suppressor p53-binding** protein 1	**2803**
L2090	Tumor suppressor p5**3**-binding protein 1	**2804**
L209 1	Type A **flavoprotein** FprA	**2805**
L2092	Type A **flavoprotein** FprA	**2806**
L2093	Type A flavoprotein FprA	**2807**
L2094	Type A flavoprotein FprA	**2808**
L2095	Type **A** flavoprotein FprA	**2809**
L2096	Type I restriction enzyme specificity protein **MG438**	**QMH**

L2097	Type i **restriction** enzyme specificity **protein MG438**	2810
L2098	Type I restriction enzyme specificity **protein MG43 8**	281 1
L2099	Type I restriction-modification enzyme, S **subunit**	2812
L2 100	Type i **restriction-modification enzyme, S** subunit	2813
L2 10 I	Type I site-specific restriction-modification system, **R** (restriction) subunit	28 14
L2 102	Type i site-specific restriction-modification system, **R** (restriction) **subunit**	2815
L2103	Type I site-specific restriction-modification system, R (restriction) subunit	2816
12 I 04	Type II DNA topoisomerase VI subunit B	2817
L2 105	Type II DNA topoisomerase VI subunit B	2818
L2 106	Type II DNA **topoisomerase** VI subunit **B**	28 19
L2 107	Type 11 DNA topoisomerase VI **subunit B**	2820
L2108	Type II DNA topoisomerase VI subunit B	282 1
L2109	Type II DNA topoisomerase VI subunit B	2822
L2 1 10	Type II DNA topoisomerase VI subunit B	2823
L2 U I	Type II DNA **topoisomerase** VI subunit **B**	2824
L2 1 12	Type 11 DNA topoisomerase VI **subunit B**	2825
L2 1 13	Type II DNA topoisomerase VI **subunit B**	2826
L21 14	Type II DNA topoisomerase VI subunit B	2827
L2 1 15	Type VI secretion system component	2828
L2 I 16	Type VI secretion system **component**	2829
L2 1 17	Type VI secretion system **component**	2830
L2 118	**Tyrosine-protein** kinase receptor UFO	283 I
L21 19	**Tyrosine-protein** kinase receptor UFO	2832
L2 120	Tyrosine-protein kinase **ZAP-70**	2833
L2 12 I	Tyrosine-protein kinase **ZAP-70**	2834
L2 122	**Tyrosyl-DNA** phosphodiesterase	2835
L2 123	Tyrosyl-DNA **phosphodiesterase**	2836
L2124	Ubiquitin **carboxyl-terrrrinal** hydrolase **7**	2837
L2 125	**UDP-galactopyranose mutase**	2838
L2 126	**UDP-galactopyranose** mutase	2839
L2 127	**UDP-gal actopyranose mutase**	2840
L2 128	**UDP-galactopyranose** mtstase	2841
L2129	UDP-galactopyranose mutase	2842
L2130	**UDP-glucose** dehydrogenase	2843
L2 13 I	**UDP-N-acetylmuramate-L-alanine** ligase	2844
L2 132	**UDP-N-acelylmutamate-L** -alanine ligase	2845
L2 133	**UDP-N-acetylmuramoylalaiune-D-glutamate** ligase	2846
L2 134	**UDP-N-acety]muramoylalaimte--D-glutamate** ligase	2847
L2135	UDP-N-acetyimuramoylalainne-D-glutamyl-lysine-D-alanyl-D-alanine ligase, **MurF** protein	2848
L2136	**UDP-N-acetv'lmuramoylalanyl-D-glutanmte--2,6-diaminopimelate** ligase	2849
L2 137	UDP-N-acetylmuramoylalanyl-D-giutamate-2,6-diaminopimelate ligase	2850
12 138	**UDP-N-acetylmuramoy** lalany-**l-D-glutamate—2,6-diaminopimelate** ligase	285 I
L2 139	**UDP-N-acety]muranioylalanyl-D-glutamate--2,6-diaiiiinopimelate** ligase	2852
L2 140	**UDP-N-acetylmuramoy** lalany**l-D-glutamate-2,6-diaminopime]ate** ligase	2853

L2141	UDP-N-acely lniuranioylaiaityl-D-glutaimte--2,6-cliairanopimelate ligase	2854
L2142	UDP-N-acetjiniuranioylalanyl-D-gIuiainate--2,6-diamiiiopimelate ligase	2855
L2143	Uncharacterized conserved protein	2856
L2144	Uncharacterized consen ⁄ed protein	2857
L2145	Uncharacterized GST-like protein yfcF	2858
L2146	Uncharacterized GST-like proteinprotein	2859
L2147	Uncharacterized GST-like proteinprotein	2860
L2148	Uncharacterized GST-like proteinprotein	2861
L2149	Uncharacterized protein	2862
L2150	Uncharacterized protein	2863
L2151	Uncharacterized protein BT_1490	2864
L2152	Uncharacterized protein ypfl	TLR
L2153	Uncharacterized protein ypfl	VHP
L2154	Uncharacterized protein ypf	2865
L2155	Uncharacte sized protein ypfl	2866
L2156	Uncharacterized protein ypfl	2867
L2157	Uncharacterized protein ypfl	2868
L2158	Uncharacterized protein ypfl	2869
L2159	Uncharacterized protein ypf	2870
L2160	Uncharacte sized protein ypfl	2871
L2161	Uncharacterized protein ypfl	2872
L2162	Uncharacterized protein ypfl	2873
L2163	Uncharacterized protein ypfl	2874
L2164	Uncharacterized protein ypf	2875
L2165	Uncharacte sized protein ypfl	2876
L2166	Uncharacterized protein ypfl	2877
L2167	Uncharacterized protein ypfl	2878
L2168	Uncharacterized protein ypfl	2879
L2169	Unknown protein	2880
L2170	Unknown protein	2881
L2171	UPFO !31 protein ykq A	2882
L2172	UPF013 1 protein ykqA	2883
L2173	UPFO 13 1 protein ykqA	2884
L2174	UPF0348 protein MJ095 1	2885
L2175	UPF0348 protein MJ095 l	2886
L2176	UPF0348 protein MJ095 1	2887
L2177	UPF0348 protein MJ095 l	2888
L2178	UPF0348 protein MJ095 1	2889
L2179	UPF0348 protein MJ095 1	2890
L2180	UPF0348 protein MJ095 l	2891
L2181	UPF0348 protein MJ095 1	2892
L2182	URE2 protein	2893
L2183	Uridine diphosplio-N-acetylenolpyruvylglucosaininerediictase	TAK
L2184	Uridine diphospho-N-acetylenolpyrawlglucosarninereductase	2894
L2185	Uridine dipliospho-N-acetylenolpyruvylglucosaminereductase	2895

L2 186	Uridine **diphospho-N-acetylenolpymvylglucosainiiiereductase**	2896
L2187	Uridine **diphospho-N-acetylenolpyruNylglucosaminereductase**	2897
L2 188	**Urokiiiase plasminogen** activator **surface** receptor	2898
L2 189	Urokinase plasminogen activator surface receptor	2899
L2 190	Vascular cell adhesion **molecule-**1	2900
L2 191	**VCP-like ATPase**	2901
L2192	**VCP-like** ATPase	2902
L2 193	Viral **CASP8** and FADD-like apoptosis regulator	2903
L2 194	Vitamin **K-dependent** protein Z	2904
L2 195	VP1 protein	2905
L2 196	**V-type ATP** synthase alpha chain	2906
L2197	Xaa-Pro aminopeptidase	2907
L2198	Xaa-Pro aminopeptidase	2908
L2199	Xaa-Pro **aminopeptidase**	2909
L2200	Xaa-Pro aminopeptidase	2910
L2201	**Xanthine** dehydrogenase	**2911**
L2202	Xanthine dehydrogenase	2912
L2203	**Xanthine** dehydrogenase	2913
L2204	Xanthine **dehydrogenase**	2914
L2205	X-prolyl **dipeptidyl** aminopeptidase	**KSY**
L2206	X-prolyl dipeptidyl aminopeptidase	LDG
L2207	X-prolyl dipeptidyl aminopeptidase	LLE
L2208	X-prolyl dipeptidyl **aminopeptidase**	TYS
L2209	X-prolyl dipeptidyl aminopeptidase	2915
L22 10	**X-prolyl** dipeptidyl aminopeptidase	**2916**
L2211	**X-prolyl** dipeptidyl aminopeptidase	2917
L22 12	X-prolyl dipeptidyl aminopeptidase	2918
L2213	X-prolyl dipeptidyl **aminopeptidase**	2919
L22 14	X-prolyl dipeptidyl aminopeptidase	2920
L2215	X-prolyl dipeptidyl aminopeptidase	2921
L22 16	**X-prolyl** dipeptidyl aminopeptidase	2922
L22 17	X-prolyl dipeptidyl aminopeptidase	2923
L2218	X-prolyl dipeptidyl **aminopeptidase**	2924
L2219	X-prolyl dipeptidyl aminopeptidase	2925
L2220	X-prolyl dipeptidyl aminopeptidase	2926
L2221	**X-prolyl** dipeptidyl aminopeptidase	2927
L2222	X-prolyl dipeptidyl aminopeptidase	2928
L2223	**X-prolyl** dipeptidyl aminopeptidase	2929
L2224	X-prolyl dipeptidyl aminopeptidase	2930
L2225	X-prolyl dipeptidyl aminopeptidase	2931
L2226	**X-prolyl** dipeptidyl aminopeptidase	2932
L2227	X-prolyl dipeptidyl aminopeptidase	2933
L2228	**X-prolyl** dipeptidyl aminopeptidase	2934
L2229	X-prolyl dipeptidyl aminopeptidase	2935
L2230	X-prolyl dipeptidyl aminopeptidase	2936

L2231	X-prolyl dipeptidyl aminopeptidase	2937
L2232	X-prolyl dipeptidyl aminopeptidase	2938
L2233	Xylosidase/arabinosidase	2939
L2234	Xylosidase/arabinosidase	2940
L2235	Xylosidase/arabinosidase	2941
L2236	Xylosidase/arabinosidase	2942
L2237	Xylosidase/arabinosidase	2943
L2238	Xylosidase/arabinosidase	2944
L2239	Xylosidase/arabinosidase	2945
L2240	YkoF	2946
L2241	YkuI protein	2947

[00147] Internal nbosomal entry site (IRES) is a nucleotide sequence (>500 nucleotides) that allows for initiation of translation in the middle of an mRNA sequence (Kirn, J.H. et al., 2011. PLoS One 6(4): e18556; the contents of which are herein incorporated by reference in its entirety). Use of an IRES sequence ensures co-expression of genes before and after the IRES, though the sequence following the IRES may be transcribed and translated at lower levels than the sequence preceding the IRES sequence.

[ΘΘ148] 2A peptides are small "self-cleaving " peptides (18-2.2 amino acids) derived from viruses such as foot-and-mouth disease virus (F2A), porcine teschovirus-1 (P2A), *Thoseaasigna* virus (T2A), or equine rhinitis A virus (E2A). The 2A designation refers specifically to a region of picomavirus poiyproteins that lead to a nbosomal skip at the glycyl-prolyl bond in the C-terminus of the 2A peptide (Kim, J.IT. et al.. 2011. PLoS One 6(4): e18556; the contents of which are herein incorporated by reference in its entirety)- This skip results in a cleavage between the 2A peptide and its immediate downstream peptide. As opposed to IRES linkers, 2A peptides generate stoichiometric expression of proteins flanking the 2A peptide and their shorter length can be advantageous in generating viral expression vectors.

[00149] Some payload regions encode linkers comprising furin cleavage sites. Furin is a calcium dependent serine endoproiease that cleaves proteins just downstream of a basic ammo acid target sequence (Arg-X-(Arg/Lys)-Arg) (Thomas, G., 2002. Nature Reviews Molecular Cell Biology 3(10): 753-66; the contents of which are herein incorporated by reference in its entirety). Furin is enriched in the trans-goJgi network where it is involved in processing cellular precursor proteins. Furin also plays a role in activating a number of pathogens. This activity can be taken advantage of for expression of polypeptides of the invention.

[00150] In some embodiments, the payload region may encode one or more linkers comprising cathepsin, matrix metalloproteinases or legumain cleavage sites. Such linkers are described e.g. by Cizeau and Macdonald in International Publication No. WO2008052322, the contents of

which are herein incorporated in their entirety. Caihepsins are a family of proteases with unique mechanisms to cleave specific proteins. Cathepsm B is a cysteine protease and cathepsin D is an aspartyl protease. Matrix metalloproteinases are a family of calcium-dependent and zinc-containing endopeptidases. Legiimain is an enzyme catalyzing the hydrolysis of (-Asn-Xaa-) bonds of proteins and small molecule substrates.

[00151] In some embodiments, payioad regions may encode linkers that are not cleaved. Such imkers may include a simple amino acid sequence, such as a glycine rich sequence. In some cases, linkers may comprise flexible peptide linkers comprising glycine and senne residues. The linker may comprise flexible peptide linkers of different lengths, e.g. nxG4S, where n=1-10 (SEQ ID NO: 9222) and the length of the encoded linker varies between 5 and 50 amino acids. In a non-limiting example, the linker may be 5xG4S (SEQ ID NO: 9221) encoded by SEQ ID NO: 903. These flexible linkers are small and without side chains so they tend not to influence secondary protein structure while providing a flexible linker between antibody segments (George, R.A., et al, 2002. Protein Engineering 15(11): 871-9; Huston, j.S. et al., 1988. PNAS 85:5879-83, and Shan, D. et al., 1999. Journal of Immunology. 162(1 1):6589-95; the contents of each of which are herein incorporated by reference in their entirety). Furthermore, the polarity of the serine residues improves solubility and prevents aggregation problems.

[00152] In some embodiments, payioad regions of the invention may encode small and unbranehed serine-rich peptide linkers, such as those described by Huston et al. in US Patent No. US5525491, the contents of which are herein incorporated in their entirety. Polypeptides encoded by the payioad region of the invention, linked by serine-rich linkers, have increased solubility.

[00153] In some embodiments, payioad regions of the invention may encode artificial linkers, such as those described by Whitlow and Fiipula in US Patent No. US5856456 and Ladner et al. in US Patent No. US 4946778, the contents of each of which are herein incorporated by their entirety.

Viral Genome Component: Introns

[00154] In one embodiment, the payioad region comprises at least one element to enliance the expression such as one or more introns or portions thereof. Non-limiting examples of introns include, MVM (67-97 bps), F.IX truncated intron I (300 bps), β-giohin SD/immunogiobulin heavy chain splice acceptor (250 bps), adenovirus splice donor/irnmunoglohra splice acceptor (500 bps), SV40 late splice donor/splice acceptor (19S/16S) (180 bps) and hybrid adenovirus splice donor/igG splice acceptor (230 bps).

[00155] In one embodiment, the intron or intron portion may be 100-500 nucleotides in length. The intron may have a length of 80, 90, 100, 110, 120, 130, 140, 150, 160, 170, 171, 172, 173, 174, 175, 176, 177, 178, 179, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, 370, 380, 390, 400, 410, 420, 430, 440, 450, 460, 470, 480, 490 or 500. The intron may have a length between 80-100, 80-120, 80-140, 80-160, 80-180, 80-200, 80-250, 80-300, 80-350, 80-400, 80-450, 80-500, 200-300, 200-400, 200-500, 300-400, 300-500, or 400-500.

Payloads of the Invention

[00156] The AAV particles of the present disclosure comprise at least one payload region. As used herein, "payload" or "payload region" refers to one or more polynucleotides or polynucleotide regions encoded by or within a viral genome or an expression product of such polynucleotide or polynucleotide region, e.g., a transgene, a polynucleotide encoding a polypeptide or multi-polypeptide or a modulatory nucleic acid or regulatory nucleic acid. Payloads of the present invention typically encode polypeptides (e.g., antibodies or antibody-based compositions) or fragments or variants thereof.

[00157] The payload region may be constructed in such a way as to reflect a region similar to or mirroring the natural organization of an mRNA.

[00158] The payload region may comprise a combination of coding and non-coding nucleic acid sequences.

[00159] In some embodiments, the AAV payload region may encode a coding or non-coding RNA.

[00160] In one embodiment, the AAV particle comprises a viral genome with a payload region comprising nucleic acid sequences encoding more than one polypeptide of interest (e.g., an antibody). In such an embodiment, a viral genome encoding more than one polypeptide may be replicated and packaged into a viral particle. A target cell transduced with a viral particle comprising more than one polypeptide may express each of the polypeptides in a single cell.

[00161] In one embodiment, as shown in FIG. 1, an AAV particle comprises a viral genome with a payload region comprising a nucleic acid sequence encoding a heavy chain and a light chain of an antibody. The heavy chain and light chain are expressed and assembled to form the antibody which is secreted.

[00162] In one embodiment, the payload region may comprise the components as shown in FIG. 2. The payload region 110 is located within the viral genome 100. At the 5' and/or the 3' end of the payload region 110 there may be at least one inverted terminal repeat (ITR) 120. Within the payload region, there is a promoter region 130, an intron region 140 and a coding

region 150. When the coding region 150 comprises a heavy chain region 151 and light chain region 152 of an antibody, the two chains may be separated by a linker region 155.

[00163] In one embodiment, the coding region may comprise a heavy and light chain sequence and a linker. As shown in FIG. 3, the payload region may comprise a heavy chain and light chain sequence separated by a linker and/or a cleavage site. In one embodiment, the heavy and light chain sequence is sequence separated by an IRES sequence (1 and 2). In one embodiment, the heavy and light chain sequence is separated by a foot and mouth virus sequence (3 and 4). In one embodiment, the heavy and light chain sequence is separated by a foot and mouth virus sequence and a furin cleavage site (5 and 6). In one embodiment, the heavy and light chain sequence is separated by a porcine teschovirus-1 virus sequence (7 and 8). In one embodiment, the heavy and light chain sequence is separated by a porcine teschovirus-1 virus and a furin cleavage site (9 and 10). In one embodiment, the heavy and light charn sequence is separated by a 5xG4S (SEQ ID NO: 9221) sequence (11).

[00164] Where the AAV particle payload region encodes a polypeptide, the polypeptide may be a peptide or protein. A protein encoded by the AAV particle payload region may comprise an antibody, an antibody related composition, a secreted protein, an intracellular protein, an extracellular protein, and/or a membrane protein. The encoded proteins may be structural or functional. In addition to the antibodies or antibody-based composition, proteins encoded by the payload region may include, in combination, certain mammalian proteins involved in immune system regulation. The AAV viral genomes encoding polypeptides described herein may be useful in the fields of human disease, viruses, infections veterinary applications and a variety of *in vivo* and *in vitro* settings.

[00165] In some embodiments, the AAV particles are useful in the field of medicine for the treatment, prophylaxis, palliation or amelioration of neurological diseases and/or disorders.

Antibodies and Antibody-based compositions

[00166] Payload regions of the AAV particles of the invention may encode polypeptides that form one or more functional antibodies or antibody-based compositions. As used herein, the term "antibody" is referred to in the broadest sense and specifically covers various embodiments including, but not limited to monoclonal antibodies, polyclonal antibodies, multispecific antibodies (e.g. bispeeifie antibodies formed from at least two intact antibodies), and antibody fragments (e.g., diabodies) so long as they exhibit a desired biological activity (e.g., "functional")- Antibodies are primarily ammo-acid based molecules but may also comprise one or more modifications (including, but not limited to the addition of sugar moieties, fluorescent moieties, chemical tags, etc.).

[00167] As used herein, "anti body-based" or "antibody-derived" compositions are monomeric or multi-meric polypeptides which comprise at least one amino-acid region derived from a known or parental antibody sequence and at least one amino acid region derived from a non-antibody sequence, e.g., mammalian protein.

[00168] Payioad regions may encode poiypeptides that form or function as any antibody, including antibodies that are known in the art and/or antibodies that are commercially available. The encoded antibodies may be therapeutic, diagnostic, or for research purposes. Further, polypeptides of the invention may include fragments of such antibodies or antibodies that have been developed to comprise one or more of such fragments (e.g., variable domains or complementarity determining regions (CDRs)).

[00169] In one embodiment, the viral genome of the AAV particles may comprise nucleic acids which have been engineered to enable expression of antibodies, antibody fragments, or components of any of those described in US7041807 relaied to YYX epitope; US2009G175884, US2011O305630, US20130330275 related to misfoided proteins in cancer; US20040175775 related to PrP in eye fluid; US20030114360 related to copoly mers and methods of treating prion-related diseases; WO2009121176 insulin-induced gene peptide compositions, US20030022243, WO2003000853 related to protein aggregation assays; WO200078344 related to prion protein peptides and uses thereof. Each of these publications are incorporated by reference in their entireties.

Antibody generation

[00170] In some embodiments, viral genomes of the AAV panicles of the invention may encode antibodies or antibody-based compositions produced using methods known in the art. Such methods may include, but are not limited to immunization and display technologies (e.g., phage display, yeast display, and ribosomal display). Antibodies may be developed, for example, using any naturally occurring or synthetic antigen. As used herein, an "antigen" is an entity which induces or evokes an immune response in an organism. An immune response is characterized by the reaction of the cells, tissues and/or organs of an organism to the presence of a foreign entity. Such an immune response typically leads to the production by the organism of one or more antibodies against the foreign entity, e.g., antigen or a portion of the antigen. As used herein, "antigens" also refer to binding partners for specific antibodies or binding agents in a display library.

[00171] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be derived from antibodies produced using hybridoma technology. Host animals (e.g. mice, rabbits, goats, and llamas) may be immunized by an

injection with an antigenic protein to elicit lymphocytes that specifically bind to the antigen. Lymphocytes may be collected and fused with immortalized cell lines to generate hybridornas which can be cultured in a suitable culture medium to promote growth. The antibodies produced by the cultured hybridornas may be subjected to analysis to determine binding specificity of the antibodies for the target antigen. Once antibodies with desirable characteristics are identified, corresponding hybridornas may be subcloned through limiting dilution procedures and grown by-standard methods. The antibodies produced by these ceils may be isolated and purified using standard immunoglobulin purification procedures.

[001 72] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be produced using heavy and light chain variable region cDNA sequences selected from hybridornas or from other sources. Sequences encoding antibody variable domains expressed by hybridornas may be determined by extracting RNA molecules from antibody-producing hybridoma cells and producing cDN A by reverse transcriptase polymerase chain reaction (PCR). PCR may be used to amplify cDNA using primers specific for heavy and light chain sequences. PCR products may then be subcloned into piasmids for sequence analysis. Antibodies may he produced by insertion of resulting variable domain sequences into expression vectors.

[00173] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be generated using display technologies. Display technologies used to generate polypeptides of the invention may include any of the display techniques (e.g. display library- screening techniques) disclosed in International Patent Application No. WO2014074532, the contents of which are herein incorporated by reference in their entirety. In some embodiments, synthetic antibodies may be designed, selected or optimized by screening target antigens using display technologies (e.g. phage display technologies). Phage display libraries may comprise millions to billions of phage particles, each expressing unique antibody fragments on their viral coats. Such libraries may provide richly diverse resources that may be used to select potentially hundreds of antibody fragments with diverse levels of affinity for one or more antigens of interest (McCafferty, et al, 1990. Nature. 348:552-4; Edwards, B.M. et al., 2003. JMB. 334: 103-18; Schofield, D. et al., 2007. Genome Biol. 8, R254 and Pershad, K. et al., 2010 Protein Engineering Design and Selection. 23:279-88; the contents of each of which are herein incorporated by reference in their entirety). Often, the antibody fragments present in such libraries comprise scFv antibody fragments, comprising a fusion protein of V_H and V_L antibody domains joined by a flexible linker. In some cases. scFvs may contain the same sequence with the exception of unique sequences encoding variable loops of the CDRs. In some cases, scFvs

are expressed as fusion proteins, linked to viral coat proteins (e.g. the K-ierminus of the viral pIII coat protein). VL chains may be expressed separately for assembly with VH chains in the periplasm prior to complex incorporation into viral coats. Precipitated library members may be sequenced from the bound phage to obtain cDNA encoding desired scFvs. Antibody variable domains or CDRs from such sequences may be directly incorporated into antibody sequences for recombinant antibody production, or mutated and utilized for further optimization through in vitro affinity maturation.

[00174] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be produced using yeast surface display technology, wherein antibody variable domain sequences may be expressed on the cell surface of *Saccharomyces cerevisiae*. Recombinant antibodies may be developed by displaying the antibody fragment of interest as a fusion to e.g. Aga2p protein on the surface of the yeast, where the protein interacts with proteins and small molecules in a solution. scFvs with affinity towards desired receptors may be isolated from the yeast surface using magnetic separation and flow cytometry. Several cycles of yeast surface display and isolation may be done to attain scFvs with desired properties through directed evolution.

[ΘΘ175] In one embodiment, the sequence of the polypeptides to be encoded in the viral genomes of the invention (e.g., antibodies) may be designed by VERSITOPE™ Antibody Generation and other methods used by BIOATLA ® and described in United States Patent Publication No. US20 130281303, the contents of which are herein incorporated by reference in their entirety. In brief, recombinant monoclonal antibodies are derived from B-celis of a host immuno-challenged with one or more target antigens. These methods of antibody generation do not rely on immortalized cell lines, such as hybridoma, thereby avoiding some of the associated challenges i.e., genetic instability and low production capacity, producing high affinity and high diversity recombinant monoclonal antibodies. In one embodiment, the method is a natural diversity approach. In another embodiment, the method is a high diversity approach.

[ΘΘ176] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be generated using BIOATLA® natural diversity approach. In the natural diversity approach of generating recombinant monocional antibodies described in United States Patent Publication No. US2013028 1303, the original pairings of variable heavy (Vii) and variable light (**VL**) domains are retained from the host, yielding recombinant monoclonal antibodies that are naturally paired. These may be advantageous due to a higher likelihood of functionality as compared to non-natural pairings of **VH** and **VL**. To produce the recombinant monoclonal antibodies, first a non-human host (i.e., rabbit, mouse, hamster, guinea

pig. camel or goat) is immuno-chaiienged with an antigen of interest. In some embodiments, the host may he a previously challenged human patient. In other embodiments, the host may not have been imrauno-challenged. B-cells are harvested from the host and screened by fluorescence activated cell sorting (FACS), or other method, to create a library of B-cells enriched in B-celis capable of binding the target antigen. The cDNA obtained from the mRNA of a single B-ceii is then amplified to generate an immunoglobulin library of VH and VL domains. This library of immunoglobulins is then cloned into expression vectors capable of expressing the VH and VL domains, wherein the VH and VL domains remain naturally paired. The library of expression vectors is then used in an expression system to express the VH and VL domains in order to create an antibody library. Screening of the antibody library yields antibodies able to bind the target antigen, and these antibodies can be further characterized. Characterization may-include one or more of the following: isoelectric point, thermal stability, sedimentation rate, folding rate, neutralization or antigen activity, antagonist or agonistic activity, expression level, specific and non-specific binding, inhibition of enzymatic activity, rigidiry/flexibiliry, shape, charge, stability across pH, in solvents, under UV radiation, in mechanical stress conditions, or in sonic conditions, half-life and glycosylation.

[00177] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be generated using BIO ATLAS) high diversity approach. In the high diversity approach of generating recombinant monoclonal antibodies described in United States Patent Publication No. US20130281303, additional pairings of variable heavy (VH) and variable light (VL) domains are attained. To produce the recombinant monoclonal antibodies, B-cells harvested from the host are screened by fluorescence activated cell sorting (FACS), panning, or other method, to create a library of B-cells enriched in B-cells capable of binding the target antigen. The **cDNA** obtained from the mRNA of the pooled B-cells is then amplified to generate an immunoglobulin library of VH and VL domains. This library of immunoglobulins is then used in a biological display system (mammalian, yeast or bacterial cell surface display systems) to generate a population of cells displaying antibodies, fragments or derivatives comprising the VH and VL domains wherein, the antibodies, fragments or derivatives comprise VH and VL domain combinations that were not present in the B-cells in vivo. Screening of the cell population by FACS, with the target antigen, yields a subset of cells capable of binding the target antigen and the antibodies displayed on these cells can be further characterized. In an alternate embodiment of the high diversity approach, the immunoglobulin library comprises only VH domains obtained from the B-cells of the immuno-chailenged host, while the VL domain(s) are obtained from another source.

[00178] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be evolved using BTOATLA® comprehensive approaches. The methods of generating recombinant monoclonal antibodies as described in United States Patent Publication No. US20130281303, further comprises evolving the recombinant antibody by comprehensive positional evolution (CPE™), CPE™ followed by comprehensive protein synthesis (CPS™), PGR shuffling, or other method.

[00179] In one embodiment, the sequence of the polypeptides to be encoded in the viral genomes of the invention (e.g., antibodies) may be derived from any of the BIOATLA® protein evolution methods described in International Publication WO2012009026, the contents of which are herein incorporated by reference in their entirety. In this method, mutations are systematically performed throughout the polypeptide or molecule of interest, a map is created providing useful informatics to guide the subsequent evolutionary steps. Not wishing to be bound by theory, these evolutionary methods typically start with a template polypeptide and a mutant is derived therefrom, which has desirable properties or characteristics. Non-limiting examples of evolutionary techniques include polymerase chain reaction (PCR), error prone PCR, oligormdeoti de-directed mutagenesis, cassette mutagenesis, shuffling, assembly PCR, sexual PCR mutagenesis, *in vivo* mutagenesis, site-specific mutagenesis, gene reassembly, gene site saturated mutagenesis, *in vitro* mutagenesis, ligase chain reaction, oligonucleotide synthesis or any combination thereof.

[00180] In one embodiment, the BIOATLA© evolution method is Comprehensive Positional Evolution (CPE™). In CPE, naturally occurring amino acid variants are generated for each of the codons of the template polypeptide, wherein 63 different codon options exist for each amino acid variant. A set of polypeptides with single amino acid mutations are generated and the mutations are then confirmed by sequencing or other method known in the art and each amino acid change screened for improved function, neutral mutations, inhibitory mutations, expression and compatibility with the host system. An EvoMap™ is created that describes in detail the effects of each amino acid mutation on the properties and characteristics of that polypeptide. The data from the EvoMap™ may be utilized to produce polypeptides with more than one amino acid mutation, wherein the resultant multi-site mutant polypeptides can be screened for desirable characteristics.

[00181] In one embodiment, the BIOATLA® evolution method is Synergy Evolution, wherein an EvoMap™ is used to identify amino acid positions to introduce 2-20 mutations simultaneously to produce a combinatorial effect. The resulting multi-site mutant polypeptides may be screened on one or more pre-determined characteristics to identify "uprautants" wherein

the function of the mutant is improved as compared to the parent polypeptide. In one embodiment, Synergy Evolution is used to enhance binding affinity of an antibody.

[00182] In one embodiment, the BTOATLA® evolution method is Flex Evolution, wherein an EvoMap™ is used to identify fully mutable sites within a polypeptide that may then be targeted for alteration, such as introduction of glycosylation sites or chemical conjugation.

[00183] In one embodiment, the BIOATLA® evolution method is Comprehensive Positional Insertion Evolution (CPI™), wherein an ammo acid is inserted after each amino acid of a template polypeptide to generate a set of lengthened polypeptides. CP! may be used to insert 1, 2, 3, 4, or 5 amino acids at each new position. The resultant lengthened polypeptides are sequenced and assayed for one or more pre-deterroined properties and evaluated in comparison to its template or parent molecule. In one embodiment, the binding affinity and inimunogenicity of the resultant polypeptides are assayed. In one embodiment, the lengthened polypeptides are further mutated and mapped to identify polypeptides with desirable characteristics.

[00184] In one embodiment, the BIOATLA®) evolution approach is Comprehensive Positional Deletion Evolution (CPD™), wherein each amino acid of the template polypeptide is individually and systematically deleted one at a time. The resultant shortened polypeptides are then sequenced and evaluated by assay for at least one pre-determined feature. In one embodiment, the shortened polypeptides are further mutated and mapped to identify polypeptides with desirable characteristics.

[00185] In one embodiment, the BIOATLA© evolution approach is Combinatorial Protein Synthesis (CPS™), wherein mutants identified in CPE, CPI, CPD or other evolutionary technique are combined for polypeptide synthesis. These combined mutant polypeptides are then screened for enhanced properties and characteristics. In one embodiment CPS is combined with any of the aforementioned evolutionary or polypeptide synthesis methods.

[00186] In one embodiment, the sequence of the polypeptides to be encoded in the viral genomes of the invention (e.g., antibodies) may be derived from the BIOATLA® Comprehensive Integrated Antibody Optimization (CIAO!™) described in United States Patent US8859467, the contents of which are herein incorporated by reference in their entirety. The CIAO!™ method allows for simultaneous evolution of polypeptide performance and expression optimization, within a eukaryotic cell host (i.e., mammalian or yeast cell host). First, an antibody library is generated in a mammalian cell production host by antibody cell surface display, wherein the generated antibody library targets a particular antigen of interest. The antibody library is then screened by any method known in the art, for one or more properties or characteristics. One or snore antibodies of the library, with desirable properties or characteristics

are chosen for further polypeptide evolution by any of the methods known in the art, to produce a library of mutant antibodies by antibody cell surface display in a mammalian cell production host. The generated mutant antibodies are screened for one or more predetermined properties or characteristics, whereby an upmutant is selected, wherein the upmutant has enhanced or improved characteristics as compared to the parent template polypeptide.

[00187] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be humanized by the methods of BIOATLAS) as described in United States Patent Publication US201 30303399, the contents of which are herein incorporated by reference in their entirety. In this method, for generating enhanced full length humanized antibodies in mammalian cells, no back-mutations are required to retain affinity to the antigen and no CDR grafting or phage-dispiay is necessary. The generated humanized antibody has reduced immunogenic!ty and equal or greater affinity for the target antigen as compared to the parent antibody. The variable regions or CDRs of the generated humanized antibody are derived from the parent or template, whereas the framework and constant regions are derived from one or more human antibodies. To stat , the parent, or template antibody is selected, cloned and each CDR sequence identified and synthesized into a CDR fragment library. Double stranded DNA fragment libraries for V_H and V_L are synthesized from the CDR fragment encoding libraries, wherein at least one CDR fragment library is derived from the template antibody and framework (FW) fragment encoding libraries, wherein the FW fragment library is derived from a pool of human frameworks obtained from natively expressed and functional human antibodies. Stepwise liquid phase ligation of FW and CDR encoding fragments is then used to generate both Vn and VL fragment libraries. The V_H and VL fragment libraries are then cloned into expression vectors to create a humanization library, which is further transfected into cells for expression of full length humanized antibodies, and used to create a humanized antibody library. The humanized antibody library is then screened to determine expression level of the humanized antibodies, affinity or binding ability for the antigen, and additional improved or enhanced characteristics, as compared to the template or parent antibody. Non-limiting examples of characteristics that may be screened include equilibrium dissociation constant (K_D), stability, melting temperature (T_m), pI, solubility, expression level, reduced immunogenicity and improved effector function.

[00188] In one embodiment, the sequences of the polypeptides to be encoded in the viral genomes of the invention may be generated by the BIOATLA® method for preparing conditionally active antibodies as described in International Publications WO201 6033331 and WO2016036916, the contents of which are herein incorporated by reference in their entirely. As

used herein, the term "conditionally active" refers to a molecule that is active at an aberrant condition. Further, the conditionally active molecule may be virtually inactive at normal physiological conditions. Aberrant conditions may result from changes in pH, temperature, osmotic pressure, osmolality, oxidative stress, electrolyte concentration, and/or chemical or proteolytic resistance, as non-limiting examples.

[00189] The method of preparing a conditionally active antibody is described in international Publications WO2016033331 and WO2.0160369 16 and summarized herewithin. Briefly, a wild-type polypeptide is selected and the DNA is evolved to create mutant DNAs. Non-limiting examples of evolutionary techniques that may be used to evolve the DNA include polymerase chain reaction (PGR), error prone PGR, shuffling, oligonucieotide-directed mutagenesis, assembly PCR, sexual PGR mutagenesis, *in vivo* mutagenesis, site-specific mutagenesis, gene reassembly, gene site saturated mutagenesis, *in vitro* mutagenesis, ligase chain reaction, oligonucleotide synthesis or any combination thereof. Once mutant DNAs are created, they are expressed in a eukaryotic cell production host (i.e., fungal, insect, mammalian, adenoviral, plant), wherein a mutant polypeptide is produced. The miitant polypeptide and the corresponding wild-type polypeptide are then subjected to assays under both normal physiological conditions and aberrant conditions in order to identify mutants that exhibit a decrease in activity in the assay at normal physiological conditions as compared to the wild-type polypeptide and/or an increase in activity in the assay under aberrant conditions, as compared to the corresponding wild-type polypeptide. The desired conditionally active mutant may then be produced in the aforementioned eukaryotic ceil production host.

[00190] In one embodiment, the conditionally active antibody is a "mirac protein" as described by BIOATLA® in United States Patent No US8709755, the contents of which are herein incorporated by reference in their entirety. As used herein "mirac protein" refers to a conditionally active antibody that is virtually inactive at body temperature but active at lower temperatures.

[00191] In one embodiment, the sequence of the polypeptides to be encoded in the viral genomes of the invention (e.g., antibodies) may be derived based on any of the BIOATLA™ methods including, but not limited to, VERSITOPE ™ Antibody Generation, natural diversity approaches and high diversity approaches for generating monoclonal antibodies, methods for generation of conditionally active polypeptides, humanized antibodies, mirac proteins, multi-specific antibodies or cross-species active mutant polypeptides, Comprehensive integrated Antibody Optimization (CIAO!™), Comprehensive Positional Evolution (CPE™), Synergy Evolution, Flex Evolution, Comprehensive Positional insertion Evolution (CPI™),

Comprehensive Positional Deletion Evolution (CPD™), Combinatorial Protein Synthesis (CPS™), or any combination thereof. These methods are described in United States Patent Nos. IJS8859467 and US8709755 and United States Publication Nos. US20130281303, US20130303399, US20150065690, US201502521 19, US20150086562 and US20100138945, and international Publication Nos. WO201 5105888, WO201 2009026, WO2011109726, WO20 160369 16, and WO20 1603333 1, the contents of each of which are herein incorporated by reference in their entirety.

Antibody fragments and variants

[00192] In some embodiments, antibody fragments encoded by payloads of the invention comprise antigen binding regions from intact antibodies. Examples of antibody fagments may include, but are not limited to Fab, Fab', F(ab')2, and Fv fragments; diabodies; linear antibodies; single-chain antibody molecules; and multispecific antibodies formed from antibody fragments. Papain digestion of antibodies produces two identical antigen-binding fragments, called "Fab" fragments, each with a single antigen-binding site. Also produced is a residual "Fc" fragment, whose name reflects its ability to crystailize readily. Pepsin treatment yields an F(ab')?. fragment that has two antigen-binding sites and is still capable of cross-linking antigen. Compounds and/or compositions of the present invention may comprise one or more of these fragments. For the purposes herein, an "antibody" may comprise a heavy and light variable domain as well as an Fc region.

[00193] In one embodiment, the Fc region may be a modified Fc region, as described in US Patent Publication US20150065690, wherein the Fc region may have a single amino acid substitution as compared to the corresponding sequence for the wild-type Fc region, wherein the single amino acid substitution yields an Fc region with preferred properties to those of the wild-type Fc region. Non-limiting examples of Fc properties that may be altered by the single amino acid substitution include bind properties or response to pH conditions

[00194] As used herein, the term "native antibody" refers to a usually heterotetramenc glycoprotein of about 150,000 Daltons, composed of two identical light (L) chains and two identical heavy (H) chains. Genes encoding antibody heavy and light chains are known and segments making up each have been well characterized and described (Matsuda, F. et a!., 1998. The Journal of Experimental Medicine. 188(11); 2151-62 and Li, A. et a!., 2004. Blood. 103(12: 4602-9, the content of each of which are herein incorporated by reference in their entirety). Each light chain is linked to a heavy chain by one covalent disulfide bond, while the number of disulfide linkages varies among the heavy chains of different immunoglobulin isotypes. Each heavy and light chain also has regularly spaced intrachain disulfide bridges. Each heavy chain

has at one end a vanabie domain (VK) followed by a number of constant domains. Each light chain has a variable domain at one end (VL) and a constant domain at its other end; the constant domain of the light chain is aligned with the first constant domain of the heavy chain, and the light chain variable domain is aligned with the variable domain of the heavy chain.

[00195] As used herein, the term "variable domain" refers to specific antibody domains found on both the antibody heavy and light chains that differ extensively in sequence among antibodies and are used in the binding and specificity of each particular antibody for its particular antigen. Variable domains comprise hypervariable regions. As used herein, the term "hypervariable region" refers to a region within a variable domain comprising amino acid residues responsible for antigen binding. The amino acids present within the hypervariable regions determine the structure of the complementarity determining regions (CDRs) that become part of the antigen-binding site of the antibody. As used herein, the term "CDR" refers to a region of an antibody comprising a structure that is complimentary to its target antigen or epitope. Other portions of the variable domain, not interacting with the antigen, are referred to as framework (FW) regions. The antigen-binding site (also known as the antigen combining site or paratope) comprises the amino acid residues necessary to interact with a particular antigen. The exact residues making up the antigen-binding site are typically elucidated by co-crystallography with bound antigen, however computational assessments can also be used based on comparisons with other antibodies (Strohl, W.R. Therapeutic Antibody Engineering. Woodhead Publishing, Philadelphia PA. 2012. Ch. 3, p47-54, the contents of which are herein incorporated by reference in their entirety). Determining residues making up CDRs may include the use of numbering schemes including, but not limited to, those taught by Kabat [Wu, T.T. et al., 1970, JEM, 132(2):211-50 and Johnson, G. et al., 2000, Nucleic Acids Res. 28(1): 214-8, the contents of each of which are herein incorporated by reference in their entirety], Chothia [Chothia and Lesk, J. Mol. Biol. 196, 901 (1987), Chothia et al, Nature 342, 877 (1989) and Al-Lazikani, B. et al., 1997, J. Mol. Biol. 273(4):927-48, the contents of each of which are herein incorporated by reference in their entirety], Lefranc (Lefranc, M.P. et al., 2005, Immunome Res. 1:3) and Honegger (Honegger, A. and Piuckthun, A. 2001. j. Viol. Biol. 309(3):657-70, the contents of which are herein incorporated by reference in their entirety).

[00196] **VH** and **VL** domains have three CDRs each. VL CDRS are referred to herein as CDR-L1, CDR-L2 and CDR-L3, in order of occurrence when moving from N- to C- terminus along the variable domain polypeptide. VH CDRS are referred to herein as CDR-H1, CDR-H2 and CDR-FI3, in order of occurrence when moving from N- to C- terminus along the variable domain polypeptide. Each of CDRs have favored canonical structures with the exception of the CDR-H3,

which comprises amino acid sequences that may be highly variable in sequence and length between antibodies resulting in a variety of three-dimensional structures in antigen-binding domains (Nikoioudis, D. et al., 2014. Peer J. 2:e456; the contents of which are herein incorporated by reference in their entirety)- In some cases, CDR-H3s may be analyzed among a panel of related antibodies to assess antibody diversity. Various methods of determining CDR sequences are known in the art and may be applied to known antibody sequences (StrohL W.R. Therapeutic Antibody Engineering. Woodhead Publishing, Philadelphia PA. 2012. Ch. 3, p47-54, the contents of which are herein incorporated by reference in their entirety).

[00197] As used herein, the term "Fv" refers to an antibody fragment comprising the minimum fragment on an antibody needed to form a complete antigen-binding site. These regions consist of a dimer of one heavy chain and one light chain variable domain in tight, non-covalent association. Fv fragments can be generated by proteolytic cleavage, but are largely unstable. Recombinant methods are known in the art for generating stable Fv fragments, typically through insertion of a flexible linker between the light chain variable domain and the heavy chain variable domain [to form a single chain Fv (scFv)] or through the introduction of a disulfide bridge between heavy and light chain variable domains (Strohl, W.R. Therapeutic Antibody Engineering. Woodhead Publishing, Philadelphia PA. 2012. Ch. 3, p46-47, the contents of which are herein incorporated by reference in their entirety).

[00198] As used herein, the term "light chain" refers to a component of an antibody from any vertebrate species assigned to one of two clearly distinct types, called kappa and lambda based on amino acid sequences of constant domains. Depending on the amino acid sequence of the constant domain of their heavy chains, antibodies can be assigned to different classes. There are five major classes of intact antibodies: IgA, IgD, IgE, IgG, and IgM, and several of these may be further divided into subclasses (isotypes), e.g., IgG1, IgG2, IgG3, igG4, IgA, and IgA2.

[00199] As used herein, the term "single chain Fv" or "scFv" refers to a fusion protein of V_H and V_L antibody domains, wherein these domains are linked together into a smgle polypeptide chain by a flexible peptide linker. In some embodiments, the Fv polypeptide linker enables the scFv to form the desired structure for antigen binding. In some embodiments, scFvs are utilized in conjunction with phage display, yeast display or other display methods where they may be expressed in association with a surface member (e.g. phage coat protein) and used in the identification of high affinity peptides for a given antigen.

[ΘΘ200] As used herein, the term "bispecific antibody" refers to an antibody capable of binding two different antigens. Such antibodies typically comprise regions from at least two different antibodies. Bispecific antibodies may include any of those described in Riethmuller, G. 2012.

Cancer immunity. 12:12-18, Marvin, J.S. et a!., 2005. Acta Pharmacologiea Simca. 26(6): 649-58 and Schaefer, W. et al., 2011. PNAS. 108(27): 11187-92, the contents of each of which are herein incorporated by reference in their entirety.

[00201] As used herein, the term "diabody" refers to a small antibody fragment with two antigen-binding sites. Diabodies comprise a heavy chain variable domain V_H connected to a light chain variable domain V_L in the same polypeptide chain. By using a linker that is too short to allow pairing between the two domains on the same chain, the domains are forced to pair with the complementary domains of another chain and create two antigen-binding sites. Diabodies are described more fully in, for example. EP 404,097; WO 93/11161: and Hollmger et aJ. (Holiinger, P. et al, "Diabodies" Small bivalent and bispecific antibody fragments. PNAS. 1993. 90:6444-8) the contents of each of which are incorporated herein by reference in their entirety.

[ΘΘ202] The term 'intrabody" refers to a form of antibody that is not secreted from a cell in which it is produced, but instead targets one or more intracellular proteins. Intrabodies may be used to affect a multitude of cellular processes including, but not limited to intracellular trafficking, transcription, translation, metabolic processes, proliferative signaling and cell division. In some embodiments, methods of the present invention may include intrabody -based therapies. In some such embodiments, variable domain sequences and/or CDR sequences disclosed herein may be incorporated into one or more constructs for intrabody-based therapy.

[00203] As used herein, the term "monoclonal antibody" refers to an antibody obtained from a population of substantially homogeneous cells (or clones), i.e., the individual antibodies comprising the population are identical and/or bind the same epitope, except for possible variants that may arise during production of the monoclonal antibodies, such variants generally being present in minor amounts. In contrast to polyclonal antibody preparations that typically include different antibodies directed against different determinants (epitopes), each monoclonal antibody is directed against a single determinant on the antigen

[00204] The modifier "monoclonal" indicates the character of the antibody as being obtained from a substantially homogeneous population of antibodies, and is not to be construed as requiring production of the antibody by any pamcular method. The monoclonal antibodies herein include "chimeric" antibodies (immunoglobulins) in which a portion of the heavy and/or light chain is identical with or homologous to corresponding sequences in antibodies derived from a particular species or belonging to a particular antibody class or subclass, vvhiie the remainder of the chain(s) is identical with or homologous to corresponding sequences in antibodies derived from another species or belonging to another antibody class or subclass, as well as fragments of such antibodies.

[00205] As used herein, the term "humanized antibody" refers to a chimeric antibody comprising a minimal portion from one or more non-human (e.g., murine) antibody source(s) with the remainder derived from one or more human immunoglobulin sources. For the most part, humanized antibodies are human immunoglobulins {recipient antibody) in which residues from the hypervariable region from an antibody oftiie recipient are replaced by residues from the hypervariable region from an antibody of anon-human species (donor antibody) such as mouse, rat, rabbit or nonhuman primate having the desired specificity, affinity, and/or capacity.

[00206] In some embodiments, viral genomes of the present invention may encode antibody mimetics. As used herein, the terra "antibody mimetic" refers to any molecule which mimics the function or effect of an antibody and which binds specifically and with high affinity to their molecular targets. In some embodiments, antibody mimetics may be monobodies, designed to incorporate the fibronectm type III domain (Fn3) as a protein scaffold (US 6,673,901; US 6,348,584). In some embodiments, antibody mimetics may be those known in the art including, but are not limited to affibody molecules, affilms, affirms, anticalins, avimers, Centyrins, DARPINS ™, Fynomers and Kunitz and domain peptides. In other embodiments, antibody mimetics may include one or more non-pep tide regions.

[ΘΘ207] As used herein, the term "antibody variant" refers to a modified antibody (in relation to a native or starting antibody) or a biomolecule resembling a native or starting antibody in structure and/or function (e.g., an antibody mimetic). Antibody variants may be altered in their amino acid sequence, composition or structure as compared to a native antibody. Antibody-variants may include, but are not limited to, antibodies with altered isotypes (e.g., IgA, IgD, IgE, IgGi, igG?., IgG?„ IgG*, or IgM), humanized variants, optimized variants, multispecific antibody variants (e.g., bispecific variants), and antibody fragments.

[ΘΘ208] The preparation of antibodies, whether monoclonal or polyclonal, is known in the art. Techniques for the production of antibodies are well known in the art and described, e.g. in Harlow and Lane "Antibodies, A Laboratory Manual", Cold Spring Harbor Laboratory Press, 1988, Harlow and Lane "Using Antibodies: A Laboratory Manual" Cold Spring Harbor Laboratory Press, 1999 and "Therapeutic Antibody Engineering: Current and Future Advances Driving the Strongest Growth Area in the Pharmaceutical Industry" Woodhead Publishing, 2012,

Multispecific antibodies

[ΘΘ209] In some embodiments, payloads of the invention may encode antibodies that bind more than one epitope. As used herein, the terms "multibody" or "multispecific antibody" refer to an antibody wherein two or more variable regions bind to different epitopes. The epitopes may

be on the same or different targets. In certain embodiments, a multi-specific antibody is a "bispecifie antibody," which recognizes two different epitopes on the same or different antigens.

[00210] in one embodiment, multi-specific antibodies may be prepared by the methods used by BIOATLA® and described in International Patent publication VVO201 10972.6, the contents of which are herein incorporated by reference in their entirety. First a library of homologous, naturally occurring antibodies is generated by any method known in the art (i.e., mammalian cell surface display), then screened by FACSAria or other screening method, for multi-specific antibodies that specifically bind to two or more target antigens. In one embodiment, the identified multi-specific antibodies are further evolved by any method known in the art, to produce a set of modified multi-specific antibodies. These modif ed multi-specific antibodies are screened for binding to the target antigens. in one embodiment, the multi-specific antibody may be further optimized by screening the evolved modified multi-specific antibodies for optimized or desired characteristics.

|00211] In one embodiment, multi-specific antibodies may be prepared by the methods used by BIOATLA® and described in Unites States Publication No. US20I50252119, the contents of which are herein incorporated by reference in their entirety. In one approach, the variable domains of two parent antibodies, wherein the parent antibodies are monoclonal antibodies are evolved using any method known in the art in a manner that allows a single light chain to functionally complement heavy chains of two different parent antibodies. Another approach requires evolving the heavy chain of a single parent antibody to recognize a second target antigen. A third approach involves evolving the light chain of a parent antibody so as to recognize a second target antigen. Methods for polypeptide evolution are described in International Publication WO2012009026, the contents of which are herein incorporated by reference in their entirety, and include as non-limiting examples. Comprehensive Positional Evolution (CPE), Combinatorial Protein Synthesis (CPS), Comprehensive Positional insertion (CPI), Comprehensive Positional Deletion (CPD), or any combination thereof. The Fc region of the multi-specific antibodies described in United States Publication No. US20150252119 may be created using a knob-in-hole approach, or any other method that allows the Fc domain to form heterodimers. The resultant multi-specific antibodies may he further evolved for improved characteristics or properties such as binding affinity for the target antigen.

Bispecifie antibodies

[00212] In some embodiments, payloads of the invention may encode bispecifie antibodies. Bispecifie antibodies are capable of binding two different antigens. Such antibodies typically comprise antigen-binding regions from at least two different antibodies. For example, a

bispecific monoclonal antibody (BsMAb, BsAb) is an artificial protein composed of fragments of two different monoclonal antibodies, thus allowing the BsAb to bind to two different types of antigen.

[00213] In some cases, payloads encode bispecific antibodies comprising antigen-binding regions from two different anti-tan antibodies. For example, such bispecific antibodies may comprise binding regions from two different antibodies selected from Tables 3-42.

[00214] Bispecific antibody frameworks may include any of those described in Riethniuller, G., 2012, *Cancer Immunity.* 12:12-18; Marvin, J.S. *el ai,* 2005. *Acta Pharmacoiogica Sinica.* 26(6):649-58; and Schaefer, W. *et a!.,* 2011. *PNAS.* 108(27): 11187-92, the contents of each of which are herein incorporated by reference in their entirety.

[00215] New generations of BsMAb, called "trifunctionai bispecific" antibodies, have been developed. These consist of two heavy and two light chains, one each from two different antibodies, where the two Fab regions (the arms) are directed against two antigens, and the Fc region (the foot) comprises the two heavy chains and forms the third binding site.

[00216] Of the two paratopes that form the tops of the variable domains of a bispecific antibody, one can be directed against a target antigen and the other against a T-lymphocyte antigen like CDS. In the case of trifunctionai antibodies, the Fc region may additionally bind to a cell that expresses Fc receptors, like a macrophage, a natural killer (NK) cell or a dendritic cell. In sum, the targeted cell is connected to one or two cells of the immune system, which subsequently destroy it.

[00217] Other types of bispecific antibodies have been designed to overcome certain problems, such as short half-life, immunogemcity and side-effects caused by cytokine liberation. They include chemically linked Fabs, consisting only of the Fab regions, and various types of bivalent and irivaleni single-chain variable fragments (scFvs). fusion proteins mimicking the variable domains of two antibodies. The furthest developed of these newer formats are the bi-specific T-cell engagers (BiTEs) and mAb2's, antibodies engineered to contain an Fcab antigen-binding fragment instead of the Fc constant region.

[00218] Using molecular genetics, two scFvs can be engineered in tandem into a single polypeptide, separated by a linker domain, called a 'tandem scFv" (tascFv). TascFvs have been found to be poorly soluble and require refolding when produced in bacteria, or they may be manufactured in mammalian cell culture systems, which avoids refolding requirements but may result in poor yields. Construction of a tascFv with genes for two different scFvs yields a "bispecific single-chain variable fragments" (bis-scFvs). Only two tascFvs have been developed clinically by commercial firms; both are bispecific agents in active early phase development by

Micfomet for oncologic indications, and are described as "Bispecific T-ceil Engagers (BiTE)." Blinaturaomab is an anti-CD19/anti-CD3 bispecific tascFv that potentiates T-cell responses to B-cell non-Hodgkin lymphoma in Phase 2. MTl10 is an anu-EP~CAM/anti~€D3 bispecific tascFv that potentiates T-celi responses to solid tumors in Phase 1. Bispecific, tetravalent "TandAbs" are also being researched by Affimed (Nelson, A. L., *MAbs.2010.* Jan-Feb; 2(l):77-83).

[00219] In some embodiments, pay loads may encode antibodies comprising a single antigen-binding domam. These molecules are extremely small, with moleculai weights approximately one-tenth of those observed for full-sized mAbs. Further antibodies may include "nanobodies" derived from the antigen-binding variable heavy chain regions (Vims) of heavy chain antibodies found in camels and llamas, which lack light chains (Nelson, A. L., *AfAbs. 20iQ.* Jan-Feb; 2(1):77-83).

[00220] Disclosed and claimed in PCT Publication WO2014144573 to Memorial Sloan-Kettering Cancer Center are muitimerization technologies for making diraeric raultispecific binding agents *{e.g.,* fusion proteins comprising antibody components) with improved properties over multispecific binding agents without the capability of dimerkation.

[00221] In some cases, payloads of the invention may encode tetravalent bispecific antibodies (TetBiAbs as disclosed and claimed in PCT Publication WO2014144357). TetBiAbs feature a second pair of Fab fragments with a second antigen specificity attached to the C-terminus of an antibody, thus providing a molecule that is bivalent for each of the two antigen specificities. The tetravalent antibody is produced by genetic engineering methods, by linking an antibody heavy chain covalently to a Fab light chain, which associates with its cognate, co-expressed Fab heavy chain.

[00222] In some aspects, payloads of the invention may encode biosynthetic antibodies as described in U.S. Patent No. 5,091,513, the contents of which are herein incorporated by reference in their entirety. Such antibody may include one or more sequences of ammo acids constituting a region which behaves as a biosynthetic antibody binding site (BABS). The sites comprise 1) non-covalentiy associated or disulfide bonded synthetic $\mathbf{V}_H$ and $\mathbf{V}_L$ dimers, 2) V_H-$\mathbf{V}_L$ or $\mathbf{Y}_L$-$\mathbf{V}_H$ single chains wherein the $\mathbf{V}_H$ and $\mathbf{V}_L$ are attached by a polypeptide linker, or 3) individuals $\mathbf{V}_H$ or V_L domains. The binding domains comprise linked CDR and FR regions, which may be derived from separate immunoglobulins. The biosynthetic antibodies may also include other polypeptide sequences which function, *e.g.,* as an enzyme, toxin, binding site, or site of attachment to an immobilization media or radioactive atom. Methods are disclosed for producing the biosynthetic antibodies, for designing BABS having any specificity that can be elicited by in vivo generation of antibody, and for producing analogs thereof.

[00223] In some embodiments, payloads may encode antibodies with antibody acceptor frameworks taught in U.S. Patent No. 8,399,625. Such antibody acceptor frameworks may be particularly well suited accepting CDRs from an antibody of interest. In some cases. CDRs from anti-tau antibodies known in the art or developed according to the methods presented herein may be used.

Miniaturized Antibody

100224] In one embodiment, the antibody encoded by the payloads of the invention may be a "mimatunzed" antibody. Among the best examples of mAb miniaturization are the small modular imraunopharinaceuticals (SMIPs) from Trubion Pharmaceuticals. These molecules, which can be monovalent or bivalent, are recombinant single-chain molecules containing one Vi., one VH antigen-binding domain, and one or two constant "effector" domains, ail connected by linker domains. Presumably, such a molecule might offer the advantages of increased tissue or tumor penetration claimed by fragments while retaining the immune effector functions conferred by constant domains. At least three "miniaturized" SMIPs have entered clinical development. TRU-015, an anti-CD20 SMIP developed in collaboration with Wyeth, is the most advanced project, having progressed to Phase 2 for rheumatoid arthritis (RA). Earlier attempts in systemic lupus erythrematosus (SLE) and B cell lymphomas were ultimately discontinued. Trubion and Facet Biotechnology are collaborating in the development of TRU-016, an anti-CD37 SMIP, for the treatment of CLL and other lymphoid neoplasias, a project that has reached Phase 2. Wyeth has licensed the ami-CD20 SMIP SBI-087 for the treatment of autoimmune diseases, including RA, SLE and possibly multiple sclerosis, although these projects remain in the earliest stages of clinical testing. (Nelson, A. L., *MAbs. 2010.* Jan-Feb; 2(l}:77- 83).

Diaboaies

[00225] In some embodiments, payloads of the invention may encode diabodies. Diabodies are functional bispeeific single-chain antibodies (bscAb). These bivalent antigen- binding molecules are composed of non-covalent dimers of scFvs, and can be produced in mammalian ceils using recombinant methods. *(See, e.g.*. Mack *ei al Proc. Natl. Acad. Sci.,* 92: 7021-7025, 1995). Few diabodies have entered clinical development. An iodme-123-labeled diabody version of the anti-€EA chimeric antibody cT84.66 has been evaluated for pre-surgical immunoscmtigraphic detection of colorectal cancer in a study sponsored by the Beckman Research Institute of the City of Hope (CJinicaltrials.gov NCT00647I53) (Nelson, A. *h.,MAbs.2 0 \0.* Jan-Feb; 2(l):77-83).

Unibody

[00226] In some embodiments, payloads may encode a "unibody," in which the hinge region has been removed from lgG4 molecules. While lgG4 molecules are unstable and can exchange

light-heavy chain heterodimers with one another, deletion of the hinge region prevents heavy chain-heavy chain pairing entirely, leaving highly specific monovalent light/heavy heterodimers, while retaining the Fc region to ensure stability and half-life in vivo. This configuration may minimize the risk of immune activation or oncogenic growth, as IgG4 interacts poorly with FcRs and monovalent umbodies fail to promote intracellular signaling complex formation. These contentions are, however, largely supported by laboratory, rather than clinical, evidence. Other antibodies may be "miniaturized" antibodies, which are compacted 100 kDa antibodies (see, e.g., Nelson, A. L., M4¾.2010. Jan-Feb; 2(l):77-83).

Intrabodies

[00227] In some embodiments, payloads of the invention may encode intrabodies. Intrabodies are a form of antibody that is not secreted from a cell in which it is produced, but instead targets one or more intracellular proteins. Intrabodies are expressed and function intracellularly, and may be used to affect a multitude of cellular processes including, but not limited to intracellular trafficking, transcription, translation, metabolic processes, proliferative signaling and ceil division. In some embodiments, methods described herein include intrahody-based therapies. In some such embodiments, variable domain sequences and/or CDR sequences disclosed herein are incorporated into one or more constructs for intrabody-based therapy. For example, intrabodies may target one or more glycated intracellular proteins or may modulate the interaction between one or more glycated intracellular proteins and an alternative protein.

[ΘΘ228] More than two decades ago, intracellular antibodies against intracellular targets were first described (Biocca, Neuberger and Cattaneo *EMBO J.* 9: 101-108, 1990). The intracellular expression of intrabodies in different compartments of mammalian cells allows blocking or modulation of the function of endogenous molecules (Biocca, *et al., EMBO J.* 9: 101-108, 1990; Colby et al., Proc. Natl. Acad. Sci. U.S.A. 101 : 17616-21, 2004). intrabodies can alter protein folding, protein-protein, protein-DNA, protein-RNA interactions and protein modification. They can induce a phenoiypic knockout and work as neutralizing agents by direct binding to the target antigen, by diverting its intracellular trafficking or by inhibiting its association with binding partners. They have been largely employed as research tools and are emerging as therapeutic molecules for the treatment of human diseases such as viral pathologies, cancer and misfoiding diseases. The fast growing bio-market of recombinant antibodies provides intrabodies with enhanced binding specificity, stability and solubility, together with lower immunogenic! ty, for their use in therapy (Biocca, abstract in Antibody Expression and Production Cell Engineering Volume 7, 2011, pp. 179-195).

[00229] In some embodiments, intrabodies have advantages over interfering RNA (iRNA); for example, iRNA has been shown to exert multiple non-specific effects, whereas intrabodies have been shown to have high specificity and affinity to target antigens. Furthermore, as proteins, intrabodies possess a much longer active half-life than iRNA. Thus, when the active half-life of the intracellular target molecule is long, gene silencing through iRNA may be slow to yield an effect, whereas the effects of intrabody expression can be almost instantaneous. Lastly, it is possible to design intrabodies to block certain binding interactions of a particular target molecule, while sparing others.

[00230] Intrabodies are often single chain variable fragments (scFvs) expressed from a recombinant nucleic acid molecule and engineered to be retained intracellular!); *(e.g.,* retained in the cytoplasm, endoplasmic reticulum, or periplasm). intrabodies may be used, for example, to ablate the function of a protein to which the intrabody binds. The expression of intrabodies may also be regulated through the use of inducible promoters in the nucleic acid expression vector comprising the intrabody. intrabodies may be produced for use in the viral genomes of the invention using methods known in the art, such as those disclosed and reviewed in: (Marasco *ei al.,* 1993 *Proc. Nail Acad. Sci. USA,* 90: 7889-7893; Chen *et al ,* 1994, *Hum. Gene Ther.* 5:595-601; Chen *el al,* 1994, *Proc. Nail. Acad. Sci. USA,* 91: 5932-5936; Maciejewski *ei al,* 1995, *Nature Med.,* 1: 667-673; Marasco, 1995, *Immunotech,* 1: 1-19; Mhashilkar, *ei al,* 1995, *EMBO J.* 14: 1542-51; Chen *ei al.,* 1996, *Hum. Gene Therap.,* 7: 1515-1525; Marasco, *Gene Ther.* 4:11-15, 1997, Rondon and Marasco, 1997, *Annu. Rev. Microbiol.* 51:257-283; Cohen, *et a!.,* 1998, *Oncogene* 17:2445-56; Proba *et al.,* 1998, *J. Mol. Biol.* 275:245-253; Cohen *eta!.,* 1998, *Oncogene* 17:2445-2456; Bassanzadeh. *eta!.,* 1998, *FEBS Lett.* 437:81-6; Richardson *et al.,* 1998, *Gene Ther.* 5:635-44; Ohage and Steipe, 1999, *J. Mol. Biol* 291:1119-1128; Ohage *ei al,* 1999, *J. Mol. Biol.* 291:1129-1134; Wirtz and Steipe, 1999, *Protein Sci.* 8:2245-2250; Zhu *ei al,* 1999, *J. Immunol. Methods* 231:207-222; Arafat *et al.,* 2000, *Cancer Gene Ther.* 7:1250-6; der **Maur** *el al..,* 2002, *J Biol Chem.* 277:45075-85; Mhashilkar *ei al.* 2002, *Gene Ther.* 9:307-19; and Wheeler *el al,* 2003, *FASEBJ.* 17: 1733-5; and references cited therein). In particular, a CCR5 intrabody has been produced by Steinberger *etal,* 2000, *Proc. Natl. Acad. Sci. USA* 97:805-810). *See* generally Marasco, WA, 1998, "Intrabodies: Basic Research and Clinical Gene Therapy Applications" Springer: New York; and for a review of scFvs, *see* P!uckthun in "The Pharmacology of Monoclonal Antibodies, 5" 1994. vol. 113, Rosenburg and Moore *eds.* Springer-Verlag, New York, pp. 269-315.

[00231] Sequences from donor antibodies may be used to develop intrabodies. intrabodies are often recombinant!); expressed as single domain fragments such as isolated VH and VL, domains

or as a single chain variable fragment (scFv) antibody within the ceil. For example, intrabodies are often expressed as a single polypeptide to form a single chain antibody comprising the variable domains of the heavy and light chains joined by a flexible linker polypeptide. intrabodies typically lack disulfide bonds and are capable of modulating the expression or activity of target genes through their specific binding activity. Single chain antibodies can also be expressed as a single chain variable region fragment joined to the light chain constant region.

[00232] As is known in the art, an mtrabody can be engineered into recombinant polynucleotide vectors to encode sub-cellular trafficking signals at its N or C terminus to allow expression at high concentrations in the sub-cellular compartments where a target protein is located. For example, intrabodies targeted to the endoplasmic reticulum (ER) are engineered to incorporate a leader peptide and, optionally, a C-terminal ER retention signal, such as the KDEL amino acid motif (SEQ ID NO: 9223). Intrabodies intended to exert activity in the nucleus are engineered to include a nuclear localization signal. Lipid moieties are joined to intrabodies in order to tether the mtrabody to the cytosoiic side of the plasma membrane. Intrabodies can also be targeted to exert function in the cytosol. For example, cytosoiic intrabodies are used to sequester factors within the cytosol. thereby preventing them from being transported to their natural cellular destination.

[00233] There are certain technical challenges with intrabody expression. In particular, protein conformational folding and structural stability of the new!y-synthesized intrabody within the ceil is affected by reducing conditions of the intracellular environment.

[00234] Intrabodies of the invention may be promising therapeutic agents for the treatment of misfolding diseases, including Alzheimer's, Parkinson's, Huntington's and prion diseases, because of their virtually infinite ability to specifically recognize the different conformations of a protein, including pathological isoforms, and because they can be targeted to the potential sites of aggregation (both intra- and extracellular sites). These molecules can work as neutralizing agents against amy!oidogenic proteins by preventing their aggregation, and/or as molecular shunters of intracellular traffic by rerouting the protein from its potential aggregation site (Cardinale, and Biocca, *Curr. Mol. Med.* 2008, 8:2-11).

Maxibodies

[00235] In one embodiment, the payloads of the invention encode a maxibody (bivalent scFV fused to the amino terminus of the Fc (CH2-CH3 domains) of IgG.

Chimeric antigen receptors

[00236] In some embodiments, the polypeptides encoded by the viral genomes of the invention (e.g., antibodies) may be used to generate chimeric antigen receptors (CARs) as described by

BiOATLA ® in international Publications WO2016033331 and WO2016036916, the contents of which are herein incorporated by reference in their entirety. As used herein, a "chimeric antigen receptor (CAR)" refers to an artificial chimeric protein comprising at least one antigen specific targeting region (ASTR), wherein the antigen specific targeting region comprises a full-length antibody or a fragment thereof that specifically binds to a target antigen. The ASTR may comprise any of the following; a full length heavy or light chain, an Fab fragment, a single chain Fv fragment, a divalent single chain antibody, or a diabody. As a nondimiting example the ASTR of a CAR may be any of the antibodies listed in Tables 3-42, antibody-based compositions or fragments thereof. Any molecule that is capable of binding a target antigen with high affinity can be used in the ASTR of a CAR. In one embodiment, the CAR may have more than one ASTR. These ASTRs may target two or more antigens or two or more epitopes of the same antigen. In one embodiment, the CAR is conditionally active. In one embodiment, the CAR is used to produce a genetically engineered cytotoxic cell carrying the CAR and capable of targeting the antigen bound by the ASTR.

[00237] Chimeric antigen receptors (CARs) are particularly useful in the treatment of cancers, though also therapeutically effective in treatment of a wide variety of other diseases and disorders. Non-limiting examples of disease categories that may be treated with CARs or CAR-based therapeutics include autoimmune disorders, B-celi mediated diseases, inflammatory diseases, neuronal disorders, cardiovascular disease and circulatory disorders, or infectious diseases. Not wishing to be bound by theory, CARs traditionally work by targeting antigens presented on the surface of or on the inside of cells to be destroyed e.g., cancer tumor cells, by the cytotoxic cell of the CAR.

Senescent Cell Surface Protein Antibodies

[00238] In some embodiments, the AAV particles may comprise nucleic acids which have been engineered to express of antibodies that selectively bind to surface marker proteins of senescent cells. For example, the antibodies may selectively bind to proteins that are in misfolded conformation. The binding antibodies may reduce the number of senescent cells and be used to treat age-related conditions, such as, but not limited to, Alzheimer's disease, cardiovascular disease, emphysema, sarcopenia, and tumorigenesis as well as conditions more cosmetic in nature such as signs of skin aging including wrinkling, sagging, discoloration, age-related tissue dysfunction, tumor formation, and other age-related conditions.

[00239] In one embodiment, the expressed antibodies binding to epitopes of senescent ceil surface proteins may be, but are not limited to, such as prion epitopes presented by SEQ ID NOs: 1-14 of international Publication No. WO2014186878; CD44 epitopes presented by SEQ ID

NOs: 47-51 of International Publication No. WO20141 86878; TNFR epitopes presented by SEQ ID NOs: 52-56 of International Publication No. WO2014186878; NOTCH! epitope presented by SEQ ID NOs: 57-61 of international Publication No. WO2014186878; FasR epitopes presented by SEQ ID NOs: 62-66 of International Publication No. WO2014186878; epidermal growth factor epitopes presented by SEQ ID NOs: 67-81 of International Publication No. **WO2014186878;** CD38 epitopes presented by SEQ ID NOs: 82-86 of International Publication No. WO2014186878, the contents of each of which are herein incorporated by reference in their entirety.

[00240] In one embodiment, the expressed antibodies may comprise peptides binding to senescent cell surface prion proteins, such as, but not limited to. those presented by SEQ ID NOs: 15-36 of International Publication No. WO2014186878, the contents of which are herein incorporated by reference in their entirety.

[00241] In one embodiment the expressed antibody may be AMF~3a-i 18 or AMF 3d-19 (SEQ ID NO: 89-92 and 103-106 of International publication WO2014186878, respectively, the contents of which are herein incorporated by reference in their entirety) targeting senescent cell surface protein FasR. in one embodiment, the expressed antibody may be Ab c-120 (SEQ ID NO: 37-40 of international publication WO2014186878, the contents of which are herein incorporated by reference in their entirety) targeting senescent cell surface protein PrP.

Payload antibodies of the invention

[00242] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Tables 3-42.

[00243] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences listed in Tables 3-42.

[00244] In some embodiments, the payload region of the AAV particle comprises a nucleic acid sequence encoding a payload antibody with at least 50% identity to one or more payload antibody polypeptides listed in Tables 3-42. The encoded antibody polypeptide may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%, 67%, 68%, 69%, 70%, 71%, 72%, 73%. 74%, 75%, 76%, 77%, 78%, 79%, 80%. 81%, 82%, 83%, 84%, 85%, 86%. 87%, 88%, 89%, 90%, 91%, 92%, 93%. 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to one or more of the payload antibody polypeptides listed in Tables 3-42,

[00245] In one embodiment, the full sequence of the encoded antibody polypeptide may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%. 67%. 68%, 69%. 70%. 71%. 72%. 73%. 74%. 75%. 76%. 77%. 78%. 79%. 80%. 81%.

82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to one or more of the payload antibody polypeptides listed in Tables 3-42.

[00246] In one embodiment, the variable region sequence(s) of the encoded antibody polypeptide may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%, 67%, 68%, 69%, 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to one or more of the payload antibody polypeptides listed in Tables 3-42.

[00247] In one embodiment, the heavy chain of the encoded antibody polypeptide may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%, 67%, 68%, 69%, 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to one or more of the payload heavy chain antibody polypeptides listed in Tables 3-42.

[00248] In one embodiment, the light chain of the encoded antibody polypeptide may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%, 67%, 68%, 69%, 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to one or more of the payload light chain antibody polypeptides listed in Tables 3-42.

[00249] In one embodiment, the CDR region of the encoded antibody polypeptide may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%, 67%, 68%, 69%, 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to the CDRs of one or more of the payload antibody polypeptides listed in Tables 3-42.

[00250] In one embodiment, the payload antibody has 90% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00251] In one embodiment, the payload antibody has 91% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00252] In one embodiment, the payload antibody has 92% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00253] In one embodiment, the payload antibody has 93% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00254] In one embodiment, the pavioad antibody has 94% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00255] In one embodiment, the pavioad antibody has 95% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00256] In one embodiment, the payioad antibody has 96% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00257] In one embodiment, the payioad antibody has 97% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00258] In one embodiment, the payioad antibody has 98% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00259] In one embodiment, the payioad antibody has 99% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00260] In one embodiment, the payioad antibody has 100% identity to one or more of the antibody polypeptides listed in Tables 3-42.

[00261] In some embodiments, the payioad region of the AAV particle comprises a nucleic acid sequence with at least 50% identity to one or more nucleic acid sequences listed in Tables 3-42, The payioad nucleic acid sequence may have 50%, 51%, 52%, 53%, 54%, 55%, 56%, 57%, 58%, 59%, 60%, 61%, 62%, 63%, 64%, 65%, 66%, 67%, 68%, 69%, 70%, 71%, 72%, 73%, 74%, 75%, 76%, 77%, 78%, 79%, 80%, 81%, 82%, 83%, 84%, 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99%, or 100% identity to one or more nucleic acid sequences listed in Tables 3-42.

[00262] In one embodiment, the payioad nucleic acid sequence has 90% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00263] In one embodiment, the payioad nucleic acid sequence has 91% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00264] In one embodiment, the payioad nucleic acid sequence has 92% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00265] In one embodiment, the payioad nucleic acid sequence has 93% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00266] In one embodiment, the payioad nucleic acid sequence has 94% identity to one or more of the nucleic acid sequences listed in Tables 3-42,

[00267] In one embodiment, the payload nucleic acid sequence has 95% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00268] In one embodiment, the payload nucleic acid sequence has 96% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00269] In one embodiment, the payload nucleic acid sequence has 97% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00270] In one embodiment, the payload nucleic acid sequence has 98% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00271] In one embodiment, the payload nucleic acid sequence has 99% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00272] In one embodiment, the payload nucleic acid sequence has 100% identity to one or more of the nucleic acid sequences listed in Tables 3-42.

[00273] In one embodiment, the payload region of the AAV particle comprises a nucleic acid sequence encoding a polypeptide which is an antibody, an antibody-based composition, or a fragment thereof. As a non-limiting example, the antibody may he one or more of the polypeptides listed in Tables 3-42. As another non-limiting example, the antibody may be one or more of the heavy chain sequences listed in Tables 3-42. As anon-limiting example, the antibody may be one or more of the light chain sequences listed in Tables 3-42.

[00274] In one embodiment, the payload region of the AAV particle comprises a nucleic acid sequence encoding a polypeptide comprising a heavy chain and a light chain sequence listed in Tables 3-42. The payload region may also comprise a linker between the heavy and light chain sequences. The linker may be a sequence known in the art or described in Table 2.

[00275] In one embodiment, the payload region of the AAV particle comprises a nucleic acid sequence encoding a polypeptide comprising a heavy chain and a light chain sequence listed in Tables 3-42, where the heavy chain sequence is from a different antibody than the light chain sequence. The payload region may also comprise a linker between the heavy and light chain sequences. The linker may be a sequence known in the art or described in Table 2.

[00276] In one embodiment, the payload region comprises, in the 5' to 3' direction, an antibody light chain sequence, a linker and a heavy chain sequence.

[00277] In one embodiment, the payload region comprises a nucleic acid sequence encoding, in the 5' to 3' direction, an antibody light chain sequence from Tables 3-42, a linker from Table 2 and a heavy chain sequence from Tables 3-42.

[00278] In one embodiment, the payload region comprises, in the 5' to 3' direction, an antibody heavy chain sequence, a linker and a light chain sequence.

[00279] In one embodiment, the payload region comprises a nucleic acid sequence encoding, in the 5' to 3' direction, an antibody heavy chain sequence from Tables 3-42, a linker from Table 2 and a light chain sequence from Tables 3-42.

[00280] In one embodiment, the payload region comprises a nucleic acid sequence encoding a single heavy chain. As a **non-limiting** example, the heavy chain is an amino acid sequence or fragment thereof described in Tables 3-42.

[00281] Shown in Tables 3-42 are a listing of antibodies and their polynucleotides and/or polypeptides sequences. These sequences may be encoded by or included in the AAV particles of the present invention. Variants or fragments of the antibody sequences described in Tables 3-42 may be utilized in the AAV particles of the present invention.

[00282] In some embodiments, the AAV particles may comprise codon-optimized versions of the nucleic acids encoding the polypeptides listed in Tables 3-42. In some cases, the payload region of the AAV particles of the invention may encode one or more isoforms or variants of these heavy7 and light chain antibody domains. Such variants may be humanized or optimized antibody domains comprising one or more complementarity determining regions (CDRs) from the heavy and light chains listed in Tables 3-42. Methods of determining CDRs are well known in the art and are described herein. Payload regions may encode antibody variants with one or more heavy chain variable domain (VH) or light chain variable domain (VL) derived from the antibody sequences in Tables 3-42. In some cases, such variants may include bispecific antibodies. Bispecific antibodies encoded by payload regions of the invention may comprise variable domain pairs from two different antibodies.

[00283] In one embodiment, the AAV particles may comprise a heavy and a light chain of an antibody described herein and two promoters. As a non-limiting example, the AAV particles may comprise a nucleic acid sequence of a genome as described in Figure 1 or Figure 2 of US Patent Publication No. US2.0030219733, the contents of which are herein incorporated by reference in its entirety. As another non-limiting example, the AAV particles may be a dual-promoter AAV for antibody expression as described by Lewis et al. (J. of. Virology, Sept 2002, Vol. 76(17), p 8769-8775; the contents of which are herein incorporated by reference in its entirety).

Foodborne Illness and Gastroenteritis Related Antibodies

[00284] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the gastrointestinal and food illness related payload antibody polypeptides listed in Tables 3-9.

[00285] In one embodiment, the **payload** region of the AAV particle comprises one or more **nucleic** acid sequences encoding one or **more** of the payload antibody polypeptides listed **in** Table 3 against Clostridium Difficile toxins.

Table 3. Antibodies against Clostridium Difficile toxins

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
CD1	Camelid heavy chain only, Toxin A and B,		WO2015100409 SEQ ID NO: 164	2948
CD2	Camelid heavy chain only, Toxin A and B,		WO2015100409 SEQ ID NO: 165	2949
CD3	Camelid heavy chain only, Toxin A and B,		WO2015100409 SEQ ID NO: 166	2950
CD4	Camelid heavy chain only, Toxin A and B,		WO2015100409 SEQ ID NO: 167	2951
CD5	Heavy chain variable region, toxin A	PA-39	US8986697 SEQ ID NO: 1	2952
CD6	Heavy chain variable region, toxin A	PA-39	US8986697 SEQ ID NO: 2	2953
CD7	Heavy chain variable region, toxin A	PA-50	US8986697 SEQ ID NO: 5	2954
CD8	Heavy chain variable region, toxin A	PA-50	US8986697 SEQ ID NO: 6	2955
CD9	Heavy chain variable region, toxin A		US20130202618 SEQ ID NO: 1	2956
CD10	Heavy chain variable region, toxin A		US20130202618 SEQ ID NO: 2	2957
CD11	Heavy chain variable region, toxin A		US20130202618 SEQ ID NO: 5	2958
CD12	Heavy chain variable region, toxin A		US20130202618 SEQ ID NO: 6	2959
CD13	Heavy chain variable region, toxin A and/or toxin B	H1H3067N	US20130230531 SEQ ID NO: 34	2960
CD14	Heavy chain variable region, toxin A and/or toxin B	H1H3134N	US20130230531 SEQ ID NO: 18	2961
CD15	Heavy chain variable region, toxin A and/or toxin B	H1H3117N	US20130230531 SEQ ID NO: 2	2962
CD16	Heavy chain variable region, toxin A and/or toxin B	H1H3123N	US20130230531 SEQ ID NO: 66	2963
CD17	Heavy chain variable region, toxin A and/or toxin B	H1H3121N	US20130230531 SEQ ID NO: 50	2964
CD18	Heavy chain variable region, toxin A and/or toxin B	H1H3124N	US20130230531 SEQ ID NO: 82	2965
CD19	Heavy chain variable region, toxin A and/or toxin B	H1H3328P	US20130230531 SEQ ID NO: 130	2966
CD20	Heavy chain variable region, toxin A and/or toxin B	H1H3324P	US20130230531 SEQ ID NO: 98	2967
CD21	Heavy chain variable region, toxin A and/or toxin B	H1H3325P	US20130230531 SEQ ID NO: 114	2968
CD22	Heavy chain variable region, toxin A and/or toxin B	H1H3330P	US20130230531 SEQ ID NO: 146	2969
CD23	Heavy chain variable region, toxin A and/or toxin B	H1H3350P	US20130230531 SEQ ID NO: 162	2970
CD24	Heavy chain variable region, toxin A and/or toxin B	H1H3347P	US20130230531 SEQ ID NO: 274	2971
CD25	Heavy chain variable region, toxin A and/or toxin B	H1H3335P	US20130230531 SEQ ID NO: 194	2972

CD26	**Heavy chain variable** region, toxin A **and/or** toxin B	H1H3344P	**US20 13023053 1** SEQ ID NO: 258	2973
CD27	Heavy chain variable region, **toxin** A and/or toxin **B**	H1H3339P	**US20 13023053 1** SEQ ID NO: 226	2974
CD28	Heavy **chain variable** region, toxin **A and/or** toxin **B**	**H1H3337P**	**US2013023053 1** SEQ **ID** NO: **210**	2975
CD29	Heavy chain variable region, toxin A and/or toxin **B**	**H1H3343P**	**US20 13023053** SEQ ID NO: 242	2976
CD30	Heavy **chain variable** region, toxin **A** and/or toxin **B**	**H1H341 1P**	**US2013023053 1** SEQ **ID** NO: 354	2977
CD31	Heavy chain variable region, toxin A and/or toxin **B**	**H1H3354P**	**US20 13023053** SEQ ID NO: 290	2978
CD32	Heavy **chain variable** region, toxin **A** and/or toxin **B**	**H1H33 17P**	**US2013023053 1** SEQ **ID** NO: 178	2979
CD33	Heavy chain variable region, toxin A and/or toxin **B**	**H1H3355P**	**US20 13023053** SEQ ID NO: 306	2980
CD34	Heavy **chain variable** region, toxin **A** and/or toxin **B**	**H1H3394P**	**US2013023053 1** SEQ **ID** NO: 322	2981
CD35	Heavy chain variable region, toxin A and/or toxin **B**	**H1H3401P**	**US20 13023053** SEQ ID NO: 338	2982
CD36	Heavy **chain variable** region, toxin B	**PA-41**	**US8986697** SEQ ID NO: 8	2983
CD37	Heavy chain vanable region, toxin B	PA-4 1	**US8986697** SEQ ID NO: 9	2984
CD38	Heavy **chain variable** region, toxin B		US20 1302026 18 SEQ ID NO: 8	2985
CD39	Heavy chain vanable region, toxin B		**US20130202618** SEQ ID NO: 9	2986
CD40	Heavy chain, toxin A	3D8	**US86091** 11 SEQ ID NO: 1	2987
CD41	Heavy chain, toxin **A**	**IB 11**	**US8609 111** SEQ ID NO: 2	2988
CD42	Heavy chain, toxin A	33.3H2	**US86091** 11 SEQ ID NO: 3	2989
CD43	Heavy chain, toxin **A**		**US20 1400041 18 SEQ ID** NO: 89	2990
CD44	Heavy chain, toxin A		US20 140004 18 SEQ ID NO: 93	2991
CD45	Heavy chain, toxin **B**		**US20 130058962** SEQ **ID** NO: 65	2992
CD46	Heavy chain, toxin **B**	**Bezlotoxuinab**		2993
CD47	Heavy -chain-only, toxin A		**US20 130058962** SEQ ID NO: 59	2994
CD48	Heavy -chain-only, toxin **A**		US20 130058962 SEQ **ID** NO: 60	2995
CD49	Heavy -chain-only, toxin A		**US20 130058962** SEQ ID NO: 6 1	2996
CD50	Heavy -chain-only, toxin **A**		US20 130058962 SEQ ID NO: 62	2997
CDS I	Heavy -chain-only, toxin A		**US20 130058962** SEQ **ID** NO: 63	2998
CD52	Heavy -chain-only, toxin A		**US20130058962** SEQ ID NO: 64	2999
CD53	Heavy -chain-only, toxin A		US20 130058962 SEQ ID NO: 87	3000
CD54	Heavy -chain-only, toxin A		**US20130058962** SEQ ID NO: 95	3001
CD55	Heavy -chain-only, toxin B	124-152	**US86091 11** SEQ ID NO: 54	3002
CD56	Heavy -chain-only, toxin B		**US20130058962** SEQ ID NO: 66	3003

CD57	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 67	3004
CD58	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 68	3005
CD59	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 69	3006
CD60	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 70	3007
CD61	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 71	3008
CD62	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 72	3009
CD63	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 73	3010
CD64	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 74	3011
CD65	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 75	3012
CD66	Heavy-chain-only, toxin B		US20130058962 SEQ ID NO: 76; SEQ ID NO: 87; SEQ ID NO: 95	3013
CD67	Light chain variable region, toxin A	PA-39	US8986697 SEQ ID NO: 3	3014
CD68	Light chain variable region, toxin A	PA-39	US8986697 SEQ ID NO: 4	3015
CD69	Light chain variable region, toxin A	PA-50	US8986697 SEQ ID NO: 7	3016
CD70	Light chain variable region, toxin A		US20130202618 SEQ ID NO: 3	3017
CD71	Light chain variable region, toxin A		US20130202618 SEQ ID NO: 4	3018
CD72	Light chain variable region, toxin A		US20130202618 SEQ ID NO: 7	3019
CD73	Light chain variable region, toxin A and/or toxin B	H1H3067N	US20130230531 SEQ ID NO: 42	3020
CD74	Light chain variable region, toxin A and/or toxin B	H1H3134N	US20130230531 SEQ ID NO: 26	3021
CD75	Light chain variable region, toxin A and/or toxin B	H1H3117N	US20130230531 SEQ ID NO: 10	3022
CD76	Light chain variable region, toxin A and/or toxin B	H1H3123N	US20130230531 SEQ ID NO: 74	3023
CD77	Light chain variable region, toxin A and/or toxin B	H1H3121N	US20130230531 SEQ ID NO: 58	3024
CD78	Light chain variable region, toxin A and/or toxin B	H1H3124N	US20130230531 SEQ ID NO: 90	3025
CD79	Light chain variable region, toxin A and/or toxin B	H1H3328P	US20130230531 SEQ ID NO: 138	3026
CD80	Light chain variable region, toxin A and/or toxin B	H1H3324P	US20130230531 SEQ ID NO: 106	3027
CD81	Light chain variable region, toxin A and/or toxin B	H1H3325P	US20130230531 SEQ ID NO: 122	3028
CD82	Light chain variable region, toxin A and/or toxin B	H1H3330P	US20130230531 SEQ ID NO: 154	3029
CD83	Light chain variable region, toxin A and/or toxin B	H1H3350P	US20130230531 SEQ ID NO: 170	3030
CD84	Light chain variable region, toxin A and/or toxin B	H1H3347P	US20130230531 SEQ ID NO: 282	3031
CD85	Light chain variable region, toxin **A** and/or toxin B	H1H3335P	US20130230531 SEQ ID NO: 202	3032

CD86	Light chain variable region, toxin A and/or toxin B	H1H3344P	US20 13023053 1 SEQ ID NO: 266	3033
CD87	Light chain variable region, toxin A and/or toxin B	H1H3339P	US20 13023053 1 SEQ ID NO: 234	3034
CD88	Light chain variable region, toxin A and/or toxin B	H1H3337P	US2013023053 1 SEQ ID NO: 218	3035
CD89	Light chain variable region, toxin A and/or toxin B	H1H3343P	US20 13023053 1 SEQ ID NO: 250	3036
CD90	Light chain variable region, toxin A and/or toxin B	H1H341 1P	US2013023053 1 SEQ ID NO: 362	3037
CD91	Light chain variable region, toxin A and/or toxin B	H1H3354P	US20 13023053 1 SEQ ID NO: 298	3038
CD92	Light chain variable region, toxin A and/or toxin B	H1H33 17P	US2013023053 1 SEQ ID NO: 186	3039
CD93	Light chain variable region, toxin A and/or toxin B	H1H3355P	US20 13023053 1 SEQ ID NO: 314	3040
CD94	Light chain variable region, toxin A and/or toxin B	H1H3394P	US2013023053 1 SEQ ID NO: 330	3041
CD95	Light chain variable region, toxin A and/or toxin B	H1H3401P	US20 13023053 1 SEQ ID NO: 346	3042
CD96	Light chain variable region, toxin B	PA-41	US8986697 SEQ ID NO: 10	3043
CD97	Light chain variable region, toxin B		US20130202618 SEQ ID NO: 10	3044
CD98	Light chain, toxin A	3D8	1JS86091 11 SEQ ID NO: 4	3045
CD99	Light chain, toxin A	IB 11	US86091 11 SEQ ID NO: 5	3046
CD 100	Light chain, toxin A	33.3H2	US86091 11 SEQ ID NO: 6	3047
CD 101	Light chain, toxin A		US20 1400041 18 SEQ ID NO: 91	3048
CD 102	Light chain, toxin A		1JS20 1400041 18 SEQ ID NO: 95	3049
CD 103	Light chain, toxin B	124-152	US8609 111 SEQ ID NO: 58	3050
CD104	Light chain, toxin B	Bezlotoxuinab		3051
CD 105	Recombinant camelid heavy chain only, Toxin A and B		WO20 15 100409 SEQ ID NO: 87	3052
CD 106	Recombinant camelid heavy chain only, Toxin A and B		WO2015100409 SEQ ID NO: 95	3053
CD 107	Recombinant camelid heavy-chain-only, toxin A		WO20 15 100409 SEQ ID NO: 59	3054
CD 108	Recombinant camelid heavy-chain-only, toxin A		WO2015100409 SEQ ID NO: 60	3055
CD 109	Recombinant camelid heavy-chain-only, toxin A		WO20 15 100409 SEQ ID NO: 61	3056
CD 110	Recombinant camelid heavy-chain-only, toxin A		WO2015100409 SEQ ID NO: 62	3057
com	Recombinant camelid heavy-chain-only, toxin A		WO20 15 100409 SEQ ID NO: 63	3058
CD 112	Recombinant camelid heavy-chain-only, toxin A		WO2015100409 SEQ ID NO: 64	3059
CD1 13	Recombinant camelid heavy-chain-only, toxin B		WO2015100409 SEQ ID NO: 65	3060
CD 114	Recombinant camelid heavy-chain-only, toxin B		WO2015100409 SEQ ID NO: 66	3061
CD1 15	Recombinant camelid heavy-chain-only, toxin B		WO2015100409 SEQ ID NO: 67	3062
CD 116	Recombinant camelid heavy-chain-only, toxin B		WO2015100409 SEQ ID NO: 68	3063

CD 117	Recombinant **camelid heavy-chain-only,** toxin B		**WO2015100409** SEQ ID NO: 69	3064
CD1 18	**Recombinant camelid heavy-chain-only**, toxin **B**		**WO20 15 100409** SEQ ID NO: 70	3065
CD 119	**Recombinant camelid** heavy-chain-only, toxin **B**		**WO2015100409 SEQ ID NO: 7 1**	3066
CD 120	**Recombinant** camelid heavy-chain-only, toxin **B**		**WO20 15 100409** SEQ ID NO: 72	3067
CD 12 l	**Recombinant camelid** heavy-chain-only, toxin **B**		**WO2015100409 SEQ ID** NO: 73	3068
CD 122	Recombinant camelid heavy-chain-only, toxin **B**		**WO20 15 100409** SEQ ID NO: 74	3069
CD 123	**Recombinant camelid** heavy-chain-only, toxin **B**		**WO2015100409 SEQ ID** NO: 75	3070
CD 124	Recombinant camelid heavy-chain-only, toxin **B**		**WO20 15 100409** SEQ **ID** NO: 76	3071
CD 125	Toxin A	**Actoxumab**		3072
CD 126	**Toxin** A	Actoxumab		3073
CD 127	Toxin A	MK3415A **(Actoxiimab+bezlotoxumab)**	**US7625559** SEQ **ID** NO: 1	3074
CD 128	Toxin A	**MK3415A (Actoxumab+bezlotoxumab)**	**US7625559** SEQ ID NO: 4	3075
CD 129	Toxin **A**	**MK34 15A (Actoxumab+b** ezlotoxumab)	**US7625559** SEQ ID NO: 54	3076
CD 130	Toxin A	MK3415A **(Actoxumab+b** ezlotoxumab)	**US7625559** SEQ **ID** NO: 58	3077
CD13 1	Toxin A	A4.2	**US20 130230537** SEQ ID NO: 34	3078
CD 132	Toxin A	A5. 1	**US20130230537** SEQ ID NO: 35	3079
CD133	Toxin A	A19.2	**US20 130230537** SEQ ID NO: 36	3080
CD 134	Toxin A	A20. 1	**US20130230537** SEQ ID NO: 37	3081
CD135	Toxin A	A24. 1	**US20 130230537** SEQ ID NO: 38	3082
CD 136	Toxin A	A26.8	**US20130230537** SEQ ID NO: 39	3083
CD 137	Toxin B	B5.2	**US20 130230537** SEQ ID NO: 40	3084
CD138	Toxin B	B7.3	**US20130230537** SEQ ID NO: 4 1	3085
CD139	Toxin B	B13.6	**US20 130230537** SEQ ID NO: 42	3086
CD 140	Toxin B	B 15.3	**US20 130230537** SEQ ID **NO: 43**	3087
CD 141	Toxin B	B15.5	**US20 130230537** SEQ ID **NO:** 44	3088

[00286] In one embodmieni, the pavload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the pavload antibody polypeptides listed in Table 4 against Campylobacter jejuni.

Table 4. Antibodies against **Campylobacter jejuni**

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
CAMP1	Consensus	FlagV1	WO2014063253 SEQ ID NO: 7	3089
CAMP2	-	FlagV1M	WO2014063253 SEQ ID NO: 8	3090
CAMP3	-	FlagV1F23M	WO2014063253 SEQ ID NO: 9	3091
CAMP4	-	FlagV1MDSB	WO2014063253 SEQ ID NO: 10	3092
CAMP5	-	FlagV1MDSB	WO2014063253 SEQ ID NO: 11	3093
CAMP6	Consensus	FlagV6	WO2014063253 SEQ ID NO: 12	3094
CAMP7	-	FlagV6M	WO2014063253 SEQ ID NO: 13	3095
CAMP8	-	FlagV6F23M	WO2014063253 SEQ ID NO: 14	3096
CAMP9	-	FlagV6MDSB	WO2014063253 SEQ ID NO: 15	3097
CAMP10	-	FlagV6F23MDSB	WO2014063253 SEQ ID NO: 16	3098

[00287] In one embodiment, the pavioad region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 5 against bacterial infections of the intestine.

Table 5. Antibodies against bacterial infections of the intestine

Antibody No.	Description	Antibody Name	Reference **Information**	SEQ ID NO
BACG1	Antibody against Listeria monocytogenes	Antibody from CN 10349 7252	CN103497252 SEQ ID NO: 1	3099
BACG2	Bivalent monovalent antibody against Pseiidomonas, Clostridium, Staphylococcus, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Shigella, and Listeria, Clostridium, Staphylococcus, Pseiidomonas, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli. Salmonella, Shigella, and Listeria bacteria	anti-LYS3-long hinge/Cys -Tag.	US7655759 SEQ ID NO: 22	3100
BACG3	Heavy chain only, Antibody against Pseiidomonas, Clostridium, Staphylococcus, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Shigella, and Listeria, Clostridium, Staphylococcus, Pseiidomonas, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Shigella, and Listeria bacteria	LYS2 VHH	US7655759 SEQ ID NO: 18	3101
BACG4	Heavy chain only, Antibody against Pseudomonas, Clostridium. Staphylococcus, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Shigella, and Listeria, Clostridium, Staphylococcus, Pseudomonas, Pasteurella, Yersinia, Bacillus anthracis. Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Shigella, and Listeria bacteria	LYS3 VHH	US7655759 SEQ ID NO: 24	3102
BACG5	Heavy chain segment including variable region, Starhylococcus enterotoxin B	F10	US8895704 SEQ ID NO: 30	3103
BACG6	Heavy chain variable region. Antibody against, P. aeruginosa, Proteus Vulgaris, non-pathogenic E. Coli, Citrobacter freundii, Serratia marcenscens, Enterobacter cloacae, Campylobacter jejuni, Helicobacter pylori, Salmonella typliimurium, Salmonella muenchen, Proteus mirabilis and Enteropathogenic E. Coli.,	mAb 741	US8263078 SEQ ID NO: 1	3104

BACG7	Heavy chain variable region. Antibody against, P. aeruginosa, Proteus Vulgaris, non-pathogenic E. Cols, Cilrobacter freimdii, Serralia marcenscens, Enlerobacter cloacae, Campylobacter jejuni, Helicobacter pylori, Salmonella ryphimiiriurn, Salmonella muenchen, Proteus niirabilis and Enleropathogenic E. Coli.,	mAb 763	US8263078 SEQ ID NO: 2	3105
BACG8	Heavy chain variable region, antibody against flagellin from Salmonella or Pseudomonas		US8173 I30 SEQ ID NO: i	3106
BACG9	Heavy chain variable region, Antibody against Gram negative (E. coli, Salmonella, Serralia, Proteus, Enterobacter, Cilrobacter, Campylobacter and Pseudomonas)	INO 743	US201002395 83 SEQ ID NO: 1	3107
BACG10	Heavy chain variable region. Antibody against Helicobacter pyroli	Abba3	US8025880 SEQ ID NO: 18	3108
BACG1 1	Heavy chain variable region, Antibody against Helicobacter pyroli	IgHV3-48*3	US8025880 SEQ ID NO: 20	3109
BACG1 2	Heavy chain variable region, Antibody against Helicobacter pyroli	clone 5	US8025880 SEQ ID NO: 2 1	3110
BACG13	Heavy chain variable region, Antibody against Helicobacter pyroli	C4	US8025880 SEQ ID NO: 22	3111
BACG14	Heavy chain variable region, Antibody against Helicobacter pyroli	IgHVl-18*01	US8025880 SEQ ID NO: 23	3112
BACG15	Heavy chain variable region, Antibody against Helicobacter pyroli	C5	US8025880 SEQ ID NO: 24	3113
BACG16	Heavy chain variable region, antibody against many pathogens.	SWLA3	WO20030079 89 SEQ ID NO: 4	3114
BACG17	Heavy chain variable region, antibody against Streptococcus mutans, Escherichia coli, Shigella dysenteriae, Salmonella lyphimurium, Streptococcus pneumoniae, Staphylococcus aureus, and Pseudomonas aeruginosa	SWLA3	US200400528 14 SEQ ID NO: 4	3115
BACG18	Heavy chain variable region, antibody against Streptococcus mutans, Escherichia coli, Shigella dysenteriae, Salmonella typhinmrium, Streptococcus pneumoniae, Staphylococcus aureus, and Pseudomonas aeruginosa	SWLA3	US200400528 14 SEQ ID NO: 8	3116
BACG19	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica. Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Abl	WO20121622 53 SEQ ID NO: 4	3117
BACG20	Heavy cliain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea,	Ab2	WO20121 622 53 SEQ ID NO: 14	3118
BACG21	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Ab3	WO20 121622 53 SEQ ID NO: 24	3119

BACG22	Heavy chain, antibody against E cols, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Ab4	WO20121622 53 SEQ ID NO: 34	3120
BACG23	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea,	Ab5	WO20121622 53 SEQ ID NO: 44	3121
BACG24	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvoviras, adenovirus, and astrovirus, other diseases causing diarrhea,	Ab6	WO20121622 53 SEQ ID NO: 54	3122
BACG25	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica. Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Ab7	WO20121622 53 SEQ ID NO: 64	3123
BACG26	Heavy chain, antibody against E coli. Shigella, Entaamoeba histolytica, Salmonella, Campylobacier, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea,	Ab8	WO20121622 53 SEQ ID NO: 74	3124
BACG27	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Al)9	WO20121622 53 SEQ ID NO: 84	3125
BACG28	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea,	AblO	WO20121622 53 SEQ ID NO: 94	3126
BACG29	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Abl 1	WO20121622 53 SEQ ID NO: 104	3127
BACG30	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea,	Abl2	WO20121622 53 SEQ ID NO: 114	3128
BACG3 I	Heavy chain, antibody against E coli. Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvoviras, adenovirus, and astrovirus, other diseases causing diarrhea.	AM 3	WO20121622 53 SEQ ID NO: 124	3129
BACG32	Heavy chain, antibody against E coli, Shigella, Entaamoeba histolytica. Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea.	Abl4	WO20121622 53 SEQ ID NO: 134	3130
BACG33	Heavy chain, Antibody against Escherichia coli infection. Staphylococcus infection		WO20140701 17 SEQ ID NO: 3	3131
BACG34	Heavy chain, Antibody against Listeria monocytogenes or WR-tubercle bacillus	6H8	US8445643 SEQ ID NO: 5	3132

BACG35	**Heavy** chain, Antibody against Pseudomonas, Clostridium, Staphylococcus, Pasteurella, Yersinia, Bacillus **anthracis,** Neisseria, Vibno, enteroioxic E. coli, Salmonella, Shigella, and Listeria, Clostridium, Staphylococcus, Pseudomonas, Pasteurella, Yersinia, Bacillus anthracis, Neissena, Vibno, enteroioxic E. coli, Salmonella, Shigella, and Listeria bacteria		**US7655759** SEQ ID **NO:** 25	3133
BACG36	Heavy chain, **Antibody** against Pseudomonas, Clostridium, Staphylococcus, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Sliigella, and Listeria, Clostridium, Staphylococcus, Pseudoinonas, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, Shigella, and Listeria bacteria		**US7655759** SEQ ID **NO:** 26	**3**134
BACG37	Heavy **chain, Starhylococcus** enterotoxin B	**100C9**	**US8895704** SEQ ID NO: 34	3135
BACG38	Heavy **chain,** Starhylococcus enterotoxin **B**	**79G9+**	**US8895704** SEQ ID NO: 38	3136
BACG39	Heavy **chain,** Starhylococcus enterotoxin B	79G9	US8895704 SEQ ID **NO:** 126	**3137**
BACG40	Heavy chain, Starhylococcus enterotoxin B	154G12	US8895704 SEQ ID NO: 142	3138
BACG4 1	Light chain variable region, Antibody against, P. aeruginosa, Proteus Vulgaris, **non-pathogenic E. Coli, Citrobacter freundii, Serralia marcenscens, Enterobacter** cloacae, Campylobacter jejuni, Helicobacter pylori. Salmonella typhimurium, Salmonella muenchen, Proteus **niirabilis** and **Enteropalhogenic** E. Coli.	mAb 741	**US8263078** SEQ ID **NO:** 3	3139
BACG42	Light chain variable region, Antibody against, **P.** aeruginosa, Proteus Vulgaris, **non-pathogenic E. Coli,** Citrobacter freundii, Serratia marcenscens, Enterobacter cloacae, Campylobacter jejuni, Helicobacter pylori, Salmonella typhimurium, **Salmonella** muenchen, Proteus **niirabilis** and **Enteropalhogenic** E. Coli	**tnAb** 763	US8263078 SEQ ID **NO:** 4	3140
BACG43	Light chain variable region, Antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus, other diseases causing diarrhea	**Ab**1	**WO20 121622** 53 SEQ ID NO: 1	**3141**
BACG44	Light **chain variable** region, antibody against **flagellin** from **Salmonella or Pseudonionas**		**US8 173 130** SEQ ID NO: **3**	3**142**
BACG45	Light chain variable region, Antibody against Gram negative **(E.** coli, Salmonella, Serratia, Proteus, Enterobacter, Citrobacter, Campylobacter and Pseudonionas)	INO 743	**US20 1002395** 83 SEQ ID NO: 2	3143
BACG46	Light chain variable region, Antibody against Helicobacter **pyroli**	Abba3	**US8025880** SEQ ID NO: 19	3144
BACG47	Light chain variable region, Antibody against many pathogens	SWLA3	**WO20030079** 89 SEQ ID NO: 7	3145
BACG48	Light chain, Antibody against **E.** coli, **Shigaella,** Entaamoeba histolytica, Salmonella, Campylobacter, or **Clostridium** difficile or a virus selected from rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus	**Ab** 1	**US20 1200294** 822 SEQ ID NO: 2	3146

BACG49	Light chain. **Antibody** against E. **coli,** Shigaelia, **Entaamoeba histoivtica, Salmonella,** Campylobacter, or Clostridium difficile or a viras selected from **rotavirus,** RSV, HIV, norvovirus, **adenovirus,** and **astroviras**	Ab 1	**US20 1200294** 822 SEQ **ID** NO: 4	3147
BACG50	Light chain, Antibody against E. coli, **Shigaelia, Entaamoeba** histoivtica, Salmonella, Campylobacter, or **Clostridium** difficile or a virus selected **from** rotavirus, RSV, **HIV, norvovirus,** adenovirus, and **astrovirus**	Ab 2	**US20 1200294** 822 SEQ ID **NO:** 12	3148
BACG5 1	Light chain, Antibody **against E. coli,** Shigaelia, **Entaamoeba** histoivtica, Salmonella, Campylobacter, or **Clostridium** difficile or **a** virus selected from rotavirus, RSV, HIV, **norvovirus, adenovirus,** and astroviras	**Ab 2**	**US20 1200294** 822 SEQ **ID** NO: 14	3149
BACG52	Light **chain.** Antibody against E. coli, Shigaelia, Entaamoeba histoivtica, Salmonella, Campylobacter, or Clostridium difficile **or** a virus selected from rotavirus, RSV, **HIV, norvovirus. adenovirus,** and astroviras	Ab 3	**US201200294** 822 SEQ ID **NO:** 22	3150
BACG53	Light chain, Antibody against E. coli, **Shigaelia,** Entaamoeba histoivtica, Salmonella, Campylobacter, or Clostridium difficile or a viras selected from rotavirus, RSV, **HIV, norvovirus, adenovirus,** and astroviras	Ab 3	**US20 1200294** 822 SEQ **ID** **NO:** 24	3151
BACG54	Light chain. Antibody against E. coli, **Shigaelia,** Entaamoeba histoivtica, Salmonella, Campylobacter, or Clostridium difficile or a viras selected from **rotavirus,** RSV, HIV, norvovirus, adenovirus, and astroviras	**Ab** 4	US201200294 822 SEQ ID NO: 32	3152
BACG55	Light chain, Antibody against **E.** coli, Shigaelia, Entaamoeba **histoivtica,** Salmonella, Campylobacter, or Clostridium difficile or a virus selected from **rotavirus,** RSV, **HIV,** norvovirus, adenovirus, and astroviras	Ab 4	**US20 1200294** 822 SEQ ID NO: 34	3153
BACG56	**Light chain,** Antibody against E. coli, Shigaelia, **Entaamoeba** histoivtica, **Salmonella, Campylobacter,** or Clostridium difficile or **a** viras selected from rotavirus, RSV, HIV, **norvovirus,** adenovirus, and astroviras	Ab 5	**US201200294** 822 SEQ ID **NO:** 42	3154
BACG57	Light chain, Antibody against **E. coli,** Shigaelia, Entaamoeba histoivtica, Salmonella, Campylobacter, or Clostridium difficile **or** a **viras** selected from rotavirus, RSV, **HIV, norvovirus. adenovirus,** and astroviras	Ab 5	**US20 1200294** 822 SEQ ID **NO:** 44	3155
BACG58	Light chain, Antibody against E. coli, **Shigaelia,** Entaamoeba histoivtica, Salmonella, Campylobacter, or **Clostridium** difficile or a viras selected from rotavirus, RSV, HIV, norvovirus, adenovirus, and astroviras	**Ab 6**	**US20 1200294** 822 SEQ **ID** NO: 52	3156
BACG59	Light chain, Antibody against E. coli, **Shigaelia,** Entaamoeba histoivtica, Salmonella, Campylobacter, or Clostridium difficile or **a virus** selected **from** rotavirus, RSV, **HIV, norvovirus,** adenovirus, and astroviras	**Ab 6**	**US20 1200294** 822 SEQ ID **NO:** 54	3157
BACG60	Light chain, Antibody against E. **coli,** Shigaelia, Entaamoeba histoivtica, Salmonella, Campylobacter, or **Clostridium** difficile or **a** virus selected from rotavirus, RSV, **HIV,** norvovirus, adenovirus, and astroviras	Ab 7	**US20 1200294** 822 SEQ **ID** NO: 62	3158
BACG61	**Light chain,** Antibody against E. coli, Shigaelia, **Entaamoeba** histoivtica, **Salmonella, Campylobacter,** or Clostridium difficile or **a** viras selected from rotavirus, RSV, HIV, **norvovirus,** adenovirus, and astroviras	Ab 7	**US201200294** 822 SEQ ID **NO:** 64	3159
BACG62	Light chain, Antibody against **E. coli,** Shigaelia, Entaamoeba histoivtica. Salmonella, Campylobacter, or Clostridium difficile or a viras selected from rotavirus, RSV, **HIV, norvovirus, adenovirus,** and astroviras	Ab **8**	**US20 1200294** 822 SEQ ID NO: 72	3160
BACG63	Light chain. Antibody against E. coli, **Shigaelia,** Entaamoeba histoivtica, **Salmonella,** Campylobacter, or Clostridiuni difficile or a viras selected from **rotavirus,** RSV, HIV, norvovirus, adenovirus, and astroviras	Ab 8	US201200294 822 SEQ ID NO: 74	3161

BACG64	Light chain. Antibody against **E. coli,** Shigaella, **Entaamoeba** histolytica. **Salmonella,** Campylobacter, or Clostridium difficile or a virus selected from **rotavirus,** RSV, HIV, norvovirus. adenovirus, and astrovirus	Ab 9	**US20 1200294** 822 SEQ **ID** NO: 82	3162
BACG65	Light chain, Antibody against **E.** coli, **Shigaella, Entaamoeba** histolytica. Salmonella, Campylobacter, or **Clostridium** difficile or **a** virus selected **from** rotavirus, RSV, **HIV, norvovirus,** adenovirus, and astrovirus	Ab 9	**US20 1200294** 822 SEQ ID **NO:** 84	3163
BACG66	Light chain, Antibody against **E. coli,** Shigaella, **Entaamoeba histolytica,** Salmonella, Campylobacter, or Clostridium difficile or a virus selected from rotavirus, RSV, HIV, norvovirus, **adenovirus,** and astrovirus	Ab 10	**US20 1200294** 822 SEQ **ID** NO: 92	3164
BACG67	Light **chain.** Antibody against E. coli, Shigaella, Entaamoeba histolytica. Salmonella, Campylobacter, or **Clostridium** difficile or a **vims** selected from rotavirus, RSV, **HIV, norvovirus. adenovirus,** and **astrovirus**	Ab 10	**US20 1200294** 822 SEQ ID **NO:** 94	3165
BACG68	Light chain, Antibody against **E.** coli, **Shigaella,** Entaamoeba **histolytica,** Salmonella, Campylobacter, or Clostridium difficile or a **virus** selected from rotavirus, RSV, **HIV, norvovirus, adenovirus,** and astrovirus	Ab 11	**US20 1200294** 822 SEQ **ID** NO: 102	3166
BACG69	Light chain. Antibody against E. coli, **Shigaella,** Entaamoeba histolytica. Salmonella, Campylobacter, or Clostridium difficile or a virus selected from rotavirus, RSV, HIV, norvovirus, adenovirus, and astrovirus	Ab 1**i**	**US20 1200294** 822 SEQ **ID** NO: 104	3167
BACG70	Light **chain.** Antibody against **E.** coli, Shigaella. **Entaamoeba histolytica,** Salmonella, Campylobacter, or **Clostridium** difficile or a virus selected from rotavirus, RSV, **HIV,** norvovirus, adenovirus, and astrovirus	Ab 12	US201200294 822 SEQ ID NO: 112	3168
BACG71	**Light chain,** Antibody against E. coli, Shigaella, **Entaamoeba histolytica, Salmonella, Campylobacter,** or **Clostridium** difficile or **a** virus selected from rotavirus, RSV, HIV, **norvovirus,** adenovirus, and **astrovirus**	Ab 12	**US201200294** 822 SEQ ID **NO:** 114	3**169**
BACG72	Light chain, Antibody against **E.** coli, Shigaella, Entaamoeba histolytica. Salmonella, Campylobacter, or Clostridium difficile or **a virus** selected from rotavirus, RSV, **HIV, norvovirus, adenovirus,** and **astrovirus**	Ab 13	**US20 1200294** 822 SEQ ID **NO:** 122	3170
BACG73	Light chain. Antibody against **E.** coli, **Shigaella,** Entaamoeba histolytica. Salmonella, Campylobacter, or **Clostridium** difficile or a virus selected from rotavirus, RSV, HIV, norvovirus. adenovirus, and astrovirus	Ab 13	**US20 1200294** 822 SEQ **ID** NO: 124	3171
BACG74	Light chain, Antibody against **E.** coli, **Shigaella, Entaamoeba** histolytica. Salmonella, Campylobacter, or Clostridium difficile or a virus selected from rotavirus, RSV, **HIV, norvovirus,** adenovirus, and astrovirus	Ab 14	**US20 1200294** 822 SEQ ID **NO:** 132	3172
BACG75	Light chain, Antibody against **E. coli,** Shigaella, Entaamoeba **histolytica.** Salmonella, Campylobacter, or **Clostridium** difficile or **a** virus selected from rotavirus, RSV, **HIV,** norvovirus, adenovirus, and astrovirus	Ab 14	**US20 1200294** 822 SEQ **ID** NO: 134	3173
BACG76	Light **chain,** Antibody against E coli, Shigella, Entaamoeba histolytica, **Salmonella, Campylobacter, or Clostridium** difficile, rotavirus. RSV. **HIV,** norvovirus, adenovirus, **and** astrovirus, other diseases causing **diarrhea**	Ab2	**WO20121622** 53 SEQ **ID** NO: 11	3174
BACG77	Light chain, Antibody against E coli, Shigella, **Entaamoeba** histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, **HIV, norvovirus,** adenovirus, and astrovirus, other diseases causing diarrhea	Ab3	WO2012 1622 53 SEQ ID **NO:** 22	3175

BACG78	Light chain. **Antibody** against **E** co3i, Shigella, **Entaamoeba** histolytica, **Salmonella,** Campylobacter, or **Clostridium** difficile, rotavirus, RSV, HIV, **norvovirus,** adenovirus, and **astroviras,** other diseases causing diarrhea	**Ab4**	**WO20121622** 53 SEQ ID NO: 31	3176
BACG79	Light **chain,** Antibody against E coli, Shigella, **Entaamoeba** histolytica, **Salmonella,** Campylobacter, or **Clostridium** difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astroviras, other diseases causing **diarrhea**	**Ab5**	**WO20121622** 53 SEQ ID NO: 42	3177
BACG80	Light chain, Antibody against E coli, Shigella, **Entaamoeba** histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, **norvovirus,** adenovirus, and astrovims, other diseases **causing** diarrhea	Ab6	**WO20121622** 53 SEQ ID **NO:** 52	3178
BACG81	Light chain, Antibody against **E coli,** Shigella, Entaamoeba histolytica. Salmonella, Campylobacter, or **Clostridium** difficile, **rotavirus,** RSV, HIV, norvovirus, **adenovirus,** and astroviras, other diseases causing diarrhea	**Ab7**	**WO20121622** 53 SEQ ID **NO:** 61	3179
BACG82	Light chain, Aniibody against E coli, Shigella, Entaamoeba **histolytica,** Salmonella, Campylobacier, or Clostridium difficile, **rotavirus,** RSV, HIV, norvovirus, adenovirus, and **astrovims,** other diseases causing diarrhea	Ab8	**WO20121622** 53 SEQ ID NO: **71**	3**180**
BACG83	Light chain, Antibody against **E** coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, adenovirus, and astroviras, other diseases causing diarrhea	**Ab9**	**WO20 121622** 53 SEQ ID NO: 82	3181
BACG84	Light **chain,** Antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, or **Clostridium** difficile, **rotavirus,** RSV, HIV, norvovirus, **adenovirus,** and astrovims, other diseases causing diarrhea	Ab!O	**WO20121622** 53 SEQ ID **NO:** 91	3182
BACG85	Light chain. Antibody against E coli, Shigella, **Entaamoeba histolytica, Salmonella,** Campylobacter, or **Clostridium** difficile, rotavirus, RSV, HIV, **norvovirus,** adenovirus, and astroviras, other diseases causing diarrhea	**Abl l**	**WO20121622** 53 SEQ ID NO: **102**	3183
BACG86	Light chain. Antibody against E coli, Shigella, Entaamoeba histolytica, Salmonella, Campylobacter, **or Clostridium** difficile, **rotavirus,** RSV, HIV, norvovirus, adenovirus, and astroviras, other diseases causing diarrhea	Ab12	**WO20121622** 53 SEQ ID **NO: 112**	3184
BACG87	Light chain, Aniibody against E coli, Shigella, **Entaamoeba** histolytica, Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, **norvovirus,** adenovirus, and astroviras, other diseases causing diarrhea	AM 3	**WO20121622** 53 SEQ ID **NO:** 122	3**185**
BACG88	Light chain, Antibody against **E coli,** Shigella, Entaamoeba histolytica. Salmonella, Campylobacter, or Clostridium difficile, rotavirus, RSV, HIV, norvovirus, **adenovirus,** and astroviras, other diseases causing **diarrhea**	**Abl4**	**WO20121622** 53 SEQ ID **NO:** 132	3186
BACG89	Light chain, Antibody against Escherichia coli infection, Staphylococcus infection		**WO20140701 17** SEQ ID NO: 4	3187
BACG90	Light chain, Aniibody against Listeria **monocytogenes** or **WR-tubercle** bacillus	6H8	**US8445643** SEQ ID **NO:** 6	3188

BACG91	Light chain. Staphylococcus **enterotoxin** B	**F10**	**US8895704** SEQ ID **NO:** 28	3189
BACG92	Light **chain,** Staphylococcus **enterotoxin B**	**100C9**	**US8895704** SEQ ID NO: 32	3190
BACG93	Light **chain.** Staphylococcus enterotoxin B	79G9	**US8895704** SEQ ID NO: 36	3191
BACG94	Light **chain,** Staphylococcus enterotoxin B	**154G12**	**US8895704** SEQ ID NO: 134	3192
BACG95	**ScFv,** Antibody against **Clostridium perfringens,** anti-alpha toxin **1A8**	**ScFv-lA8**	Zhao, B. and **Xu, C.** "Cloning and sequencing of the **ScFv-2E3** gene **anti-**aipha toxin of Clostridium perfringens type A", Chin. **J.** Vet. Sci. 20, 246-248 (2000), **CNBI** Accession # **AAU 11282**	3193
BACG96	ScFv, Antibody against Clostridium perfringens, anti-alpha toxin 2E3	**ScFv-2E3**	Zhao, B. and Xu, **C.** "Cloning and sequencing of **the ScFv-2E3** gene **anti-**alpha toxin of Clostridium perfringens type **A", Chin.** J. Vet. Sci. **20,** 246-248 (2000), NCBI Accession # **AAU 11283**	3194
BACG97	Variable fragment, Antibody against **Pseudonionas, Clostridium,** Staphylococcus, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. **coli.** Salmonella, Shigella, and Listeria, Clostridium, Staphylococcus, **Pseudomonas, Pasteurella,** Yersinia, Bacillus **anthracis,** Neisseria, Vibrio, enterotoxic E. coli. Salmonella, Shigella, and Listeria bacteria	**aTT2**	US7655759 SEQ ID **NO:** 8	3195
BACG98	Variable fragment, antibody against Pseudomonas, Clostridium, Staphylococcus, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic **E.** coli, Salmonella, Shigella, and Listeria, **Clostridium,** Staphylococcus, Pseudomonas, Pasteurella, Yersinia, Bacillus anthracis, Neisseria, Vibrio, enterotoxic E. coli, Salmonella, **Shigella,** and Listeria bacteria,	αTTI	**US7655759** SEQ ID NO: 7	3196

[00288] In one embodiment, the payload region of the AAV particle comprises one or more **nucleic** acid sequences encoding one or **more** of the payload antibody polypeptides listed **in** Table 6 against Hepatitis A and/or Hepatitis E.

Table 6. Antibodies against Hepatitis A and Hepatitis E

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
HEPAE1	Heavy chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-216	CN1486990A; CN100497391C	3197
HEPAE2	Heavy chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-315	CN1486990A; CN100497391C	3198
HEPAE3	Heavy chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-319	CN1486990A; CN100497391C	3199
HEPAE4	Heavy chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-328	CN1486990A; CN100497391C	3200
HEPAE5	Heavy chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-404	CN1486990A; CN100497391C	3201
HEPAE6	Heavy chain variable region, HEV monoclonal antibody	13D8	US7786264 SEQ ID NO.8; US20060233822; US20100003281; EP1452541; EP2322625	3202
HEPAE7	Heavy chain variable region, HEV monoclonal antibody	16D7	US7786264 SEQ ID NO.20; US20060233822; US20100003281; EP1452541; EP2322625	3203
HEPAE8	Heavy chain variable region, HEV monoclonal antibody	8C11	US7786264 SEQ ID NO.12; US20060233822; US20100003281; EP1452541; EP2322625	3204
HEPAE9	Heavy chain variable region, HEV monoclonal antibody	8H3	US7786264 SEQ ID NO.16; US20060233822; US20100003281; EP1452541; EP2322625	3205
HEPAE10	Heavy chain variable region, HEV neutralizing antibody	HEV#31	US7148323 SEQ ID NO: 3; US20050233316; US6930176; WO2001040270	3206
HEPAE11	Heavy chain variable region, HEV neutralizing antibody	HEV#4	US7148323 SEQ ID NO: 1; US20050233316; US6930176; WO2001040270	3207
HEPAE12	Heavy chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86899.1(124aa)	3208
HEPAE13	Heavy chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by	3209

			phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86898.1(129aa)	
HEPAE14	Heavy chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86897.1(123aa)	3210
HEPAE15	Heavy chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86896.1(129aa)	3211
HEPAE16	Heavy chain, HEV antibody (mouse monoclonal antibody), E2 glycoprotein	8g12	Gu Y., et al., Structural basis for the neutralization of hepatitis E virus by a cross-genotype antibody; Cell Res. 25 (5), 604-620 (2015); NCBI Accession # 4PLJ_H (229aa)	3212
HEPAE17	Heavy chain, HEV antibody (mouse monoclonal antibody), E2 glycoprotein		Tang X., et al., Proc. Natl. Acad. Sci. U.S.A. 108 (25), 10266-10271 (2011); NCBI Accession # 3RKD_H(230aa)	3213
HEPAE18	Light chain variable region, gamma1, HAV,	HAV#14	US7635476 SEQ ID NO: 4; US7282205; US20040260067; US20070287667; WO2003040341	3214
HEPAE19	Light chain variable region, gamma1, HAV,	HAV#4	US7635476 SEQ ID NO: 1; US7282205; US20040260067; US20070287667; WO2003040341	3215
HEPAE20	Light chain variable region, gamma1, HAV,	HAV#5	US7635476 SEQ ID NO: 2; US7282205; US20040260067; US20070287667; WO2003040341	3216
HEPAE21	Light chain variable region, gamma1, HAV,	HAV#6	US7635476 SEQ ID NO: 3; US7282205; US20040260067; US20070287667; WO2003040341	3217
HEPAE22	Light chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-216	CN1486990A; CN100497391C	3218
HEPAE23	Light chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-315	CN1486990A; CN100497391C	3219
HEPAE24	Light chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-319	CN1486990A; CN100497391C	3220
HEPAE25	Light chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-328	CN1486990A; CN100497391C	3221
HEPAE26	Light chain variable region, HEV Ab, a humanized neutralizing genetically engineered antibody	HEV-Fab-404	CN1486990A; CN100497391C	3222
HEPAE27	Light chain variable region, monovalent, HAV		WO2011114353 SEQ ID NO: 25	3223
HEPAE28	Light chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to	3224

			hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86903.1(107aa)	
HEPAE29	Light chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86902.1(107aa)	3225
HEPAE30	Light chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86901.1(107aa)	3226
HEPAE31	Light chain variable region, partial, HAV	anti-HAV capsid	Kim S.J., et al., Neutralizing human monoclonal antibodies to hepatitis A virus recovered by phage display; Virology 318 (2), 598-607 (2004), NCBI Accession # AAO86900.1(107aa)	3227
HEPAE32	Light chain variable, HEV monoclonal antibody	13D8	US7786264 SEQ ID NO.6; US20060233822; US20100003281; EP1452541; EP2322625	3228
HEPAE33	Light chain variable, HEV monoclonal antibody	16D7	US7786264 SEQ ID NO.18; US20060233822; US20100003281; EP1452541; EP2322625	3229
HEPAE34	Light chain variable, HEV monoclonal antibody	8C11	US7786264 SEQ ID NO: 10; US20060233822; US20100003281; EP1452541; EP2322625	3230
HEPAE35	Light chain variable, HEV monoclonal antibody	8H3	US7786264 SEQ ID NO.14; US20060233822; US20100003281; EP1452541; EP2322625	3231
HEPAE36	Light chain variable, HEV monoclonal antibody	HEV#31	US7148323 SEQ ID NO: 4; US20050233316; US6930176; WO2001040270	3232
HEPAE37	Light chain variable, HEV monoclonal antibody	HEV#4	US7148323 SEQ ID NO: 2; US20050233316; US6930176; WO2001040270	3233
HEPAE38	Light chain, E2 glycoprotein, HEV antibody (mouse monoclonal antibody)	8g12	Gu Y., et al., Structural basis for the neutralization of hepatitis E virus by a cross-genotype antibody; Cell Res. 25 (5), 604-620 (2015); NCBI Accession # 4PLJ_L (212aa)	3234
HEPAE39	Light chain, E2 glycoprotein, HEV antibody (mouse monoclonal antibody)		Tang X., et al., Proc. Natl. Acad. Sci. U.S.A. 108 (25), 10266-10271 (2011), NCBI Accession # 3RKD_C (214aa)	3235
HEPAE40	Monovalent Heavy chain variable region, HAV		WO2011114353 SEQ ID NO: 24	3236
HEPAE41	ScFv, HAV, Monovalent human antibody		WO2011114353 SEQ ID NO: 27	3237

[00289] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in Chinese Pub. No. CN103923881, CN103923882, CN1605628, CN1318565, CN1163512, the contents of each of which are herein incorporated by reference in their entirety, against HAY.

[00290] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 7 against Norwalk virus.

Table 7. Antibodies against Norwalk virus

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
NORV1	Heavy chain variable region, Norwalk virus	B7	WO2014126921 SEQ ID NO: 8	3238
NORV2	Light chain variable region, Norwalk virus	B7	WO2014126921 SEQ ID NO: 16	3239
NORV3	Heavy chain variable region, Norwalk virus	B72	WO2014126921 SEQ ID NO: 120	3240
NORV4	Light chain variable region, Norwalk virus	B72	WO2014126921 SEQ ID NO: 128	3241
NORV5	Heavy chain variable region, Norwalk virus	C9	WO2014126921 SEQ ID NO: 88	3242
NORV6	Light chain variable region, Norwalk virus	C9	WO2014126921 SEQ ID NO: 96	3243
NORV7	Heavy chain variable region, Norwalk virus	D4	WO2014126921 SEQ ID NO: 136	3244
NORV8	Light chain variable region, Norwalk virus	D4	WO2014126921 SEQ ID NO: 144	3245
NORV9	Heavy chain variable region, Norwalk virus	D8	WO2014126921 SEQ ID NO: 24	3246
NORV10	Light chain variable region, Norwalk virus	D8	WO2014126921 SEQ ID NO: 32	3247
NORV11	Heavy chain variable region, Norwalk virus	E5	WO2014126921 SEQ ID NO: 40	3248
NORV12	Light chain variable region, Norwalk virus	E5	WO2014126921 SEQ ID NO: 48	3249
NORV13	Heavy chain variable region, Norwalk virus	F11	WO2014126921 SEQ ID NO: 72	3250
NORV14	Light chain variable region, Norwalk virus	F11	WO2014126921 SEQ ID NO: 80	3251
NORV15	Heavy chain variable region, Norwalk virus	G3	WO2014126921 SEQ ID NO: 104	3252
NORV16	Light chain variable region, Norwalk virus	G3	WO2014126921 SEQ ID NO: 112	3253
NORV17	Heavy chain variable region, Norwalk virus	G4	WO2014126921 SEQ ID NO: 56	3254
NORV18	Light chain variable region, Norwalk virus	G4	WO2014126921 SEQ ID NO: 64	3255
NORV19	Heavy chain variable region, Norwalk or MD2004 virus		WO2014183052 SEQ ID NO: 1	3256
NORV20	Heavy chain variable region, Norwalk or MD2004 virus		WO2014183052 SEQ ID NO: 2	3257
NORV21	Heavy chain variable region, Norwalk or MD2004 virus		WO2014183052 SEQ ID NO: 3	3258

NORV22	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 4	3259
NORV23	Heavy chain variable region. Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 5	3260
NORV24	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 6	3261
NORV25	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 7	3262
NORV26	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 8	3263
NORV27	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 9	3264
NORV28	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 10	3265
NORV29	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 11	3266
NORV30	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 12	3267
NORV31	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 13	3268
NORV32	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 14	3269
NORV33	Heavy chain variable region, Norwaik or MD2004 viras		WO2014183052 SEQ ID NO: 15	3270
NORV34	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 16	3271
NORV35	Heavy chain variable region, Norwaik or MD2004 viras		WO2014183052 SEQ ID NO: 17	3272
NORV36	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 18	3273
NORV37	Heavy chain variable region, Norwaik or MD2004 viras		WO2014183052 SEQ ID NO: 19	3274
NORV38	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 20	3275
NORV39	Heavy chain variable region, Norwaik or MD2004 viras		WO2014183052 SEQ ID NO: 21	3276
NORV40	Heavy chain variable region, Norwaik **or** MD2004 virus		WO2014183052 SEQ ID NO: 22	3277
NORV41	Heavy chain variable region, Norwaik or MD2004 **viras**		WO2014183052 SEQ ID NO: 23	3278
NORV42	Heavy chain variable region, Norwaik or MD2004 virus		WO2014183052 SEQ ID NO: 24	3279
NORV43	Heavy cliain variable region, Nonvalk or MD2004 **viras**		WO2014183052 SEQ ID NO: 25	3280
NORV44	Heavy chain variable region, Nonvalk or MD2004 virus		WO2014183052 SEQ ID NO: 26	3281
NORV45	Heavy cliain variable region, Nonvalk or MD2004 **viras**		WO2014183052 SEQ ID NO: 27	3282
NORV46	Heavy chain variable region, Nonvalk or MD2004 virus		WO2014183052 SEQ ID NO: 28	3283
NORV47	Heavy cliain variable region, Nonvalk or MD2004 **viras**		WO2014183052 SEQ ID NO: 29	3284
NORV48	Heavy chain variable region, Nonvalk or MD2004 virus		WO2014183052 SEQ ID NO: 30	3285

[00291] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 8 against Rotavirus.

Table 8. Antibodies **against rotavirus**

Antibody No.	Description	Reference Information	SEQ ID NO
ROTV1	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 1	3286
ROTV2	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 2	3287
ROTV3	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 3	3288
ROTV4	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 4	3289
ROTV5	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 5	3290
ROTV6	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 6	3291
ROTV7	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 7	3292
ROTV8	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 8	3293
ROTV9	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 9	3294
ROTV10	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 10	3295
ROTV11	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 11	3296
ROTV12	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 12	3297
ROTV13	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 13	3298
ROTV14	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 14	3299
ROTV15	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 15	3300
ROTV16	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 16	3301
ROTV17	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 17	3302
ROTV18	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 18	3303
ROTV19	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 19	3304
ROTV20	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 20	3305
ROTV21	Heavy chain single domain	US8105592; US20090226418 SEQ ID NO: 21	3306
ROTV22	Human VP6 polypeptide	US20030166139 SEQ ID NO: 2	3307
ROTV23	Human VP6 polypeptide	US20030166139 SEQ ID NO: 4	3308
ROTV24		Aiyegbo, M.S., et al " Human RotavirUSVp6-Specific Antibodies Mediate Intracellular Neutralization By Binding To A Quater Structure In The Transcriptional Pore", Plos One 8, 61101 (2013), NCBI Accession # 4HFW_B	3309
ROTV25		Aiyegbo, M.S., et al " Human RotavirUSVp6-Specific Antibodies Mediate Intracellular Neutralization By Binding To A Quater Structure In The Transcriptional Pore", Plos One 8, 61101 (2013), NCBI Accession # 4HFW_A	3310

[00292] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 9 against Entamoeba Histolytica.

Table 9. Antibodies against Entamoeba Histolytica

Antibody No./ Antibody Name	Description	Reference Information	SEQ ID NO
ENTH1	Heavy chain (partial sequence) gamma, Entamoeba histolytica antibody	Cheng, X.J. et al., Exp. Parasitol. 96 (1), 52-56 (2000), NCBI Accession # BAA97670.1 (220aa)	3311
ENTH2	Heavy chain (partial sequence) gamma, Entamoeba histolytica Antibody	Tachibana, H. et al., Clin. Diagn. Lab. Immunol. 6 (3), 383-387 (1999), NCBI Accession # BAA82104.1 (222aa)	3312

ENTH3	Heavy chain (partial sequence) gamma, Entamoeba histolytica Antibody	Tachibana, H. et al., Clin. Diagn. Lab. Immunol. 6 (3), 383-387 (1999), NCBI Accession # BAA82101.1 (226aa)	3313
ENTH4	Heavy chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03695.1 (220aa)	3314
ENTH5	Heavy chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03694.1 (226aa)	3315
ENTH6	Heavy chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03693.1 (221aa)	3316
ENTH7	Heavy chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03692.1 (223aa)	3317
ENTH8	Light chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03699.1 (219aa)	3318
ENTH9	Light chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03698.1 (220aa)	3319
ENTH10	Light chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03697.1 (214aa)	3320
ENTH11	Light chain (partial sequence) IgG, Entamoeba histolytica Intermediate Subunit Lectin-Specific Human Monoclonal Antibodies	Tachibana, H., et al., Infect. Immun. 77 (1), 549-556 (2009), NCBI Accession # BAH03696.1 (214aa)	3321
ENTH12	Light chain (partial sequence) kappa, Entamoeba histolytica antibody	Cheng, X.J. et al., Exp. Parasitol. 96 (1), 52-56 (2000), NCBI Accession # BAA97671.1 (214aa)	3322
ENTH13	Light chain (partial sequence) kappa, Entamoeba histolytica antibody	Tachibana, H. et al., Clin. Diagn. Lab. Immunol. 6 (3), 383-387 (1999), NCBI Accession # BAA82105.1 (215aa)	3323
ENTH14	Light chain (partial sequence) kappa, Entamoeba histolytica antibody	Tachibana, H. et al., Clin. Diagn. Lab. Immunol. 6 (3), 383-387 (1999), NCBI Accession # BAA82100.1 (214aa)	3324
ENTH15 / 350-E2	Single chain Fv Antibody 350-E2 against Entamoeba histolytica	NCBI Accession # AEY80059.1 (274aa)	3325
ENTH16 / JR4A11	Single chain Fv Antibody JR4A11 Entamoeba histolytica	NCBI Accession # AEY80058.1 (287aa)	3326

[00293] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides, fragments or variants thereof described in international Pub. No. WO200I012646, the contents of which are herein incorporated by reference in their entirety, against listeria monocytogenes, salmonella and/or leishmania.

Neglected Tropical Diseases

[00294] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the neglected tropical disease related payload antibody polypeptides listed in Tables 10-13.

[00295] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 10 against Dengue Fever Virus.

Table **10. Antibodies** against **Dengue** Fever Vims

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
DENG1	Bispecific, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m3666	US20150218255 SEQ ID NO: 96	3327
DENG2	Fab Fragment	Fab 14c10	Teoh, E.P., et al., Sci Transl Med 4 (139), 139RA83 (2012), NCBI Accession # 4CAU_E(230 aa)	3328
DENG3	Heavy chain	5j7 Fab	Fibriansah, G., et al., A highly potent human antibody neutralizes dengue virus serotype 3 by binding across three surface proteins; Nat Commun 6, 6341 (2015), NCBI Accession # 3J6U_H (135aa)	3329
DENG4	Heavy Chain	Ede1 C8	Dejnirattisai, W., et al., A new class of highly potent, broadly neutralizing antibodies isolated from viremic patients infected with dengue virus; Nat. Immunol. 16 (2), 170-177 (2015), NCBI Accession # 4UTA_H (272 aa)	3330
DENG5	Heavy Chain	Fab 2h12	Midgley, C.M., et al., J, Immunol. 188 (10), 4971-4979 (2012), NCBI Accession # 4AL8_H (217 aa)	3331
DENG6	Heavy Chain Fab Fragment Of Antibody 1f4	1f4 Fab	Fibriansah, G., et al., A potent anti-dengue human antibody preferentially recognizes the conformation of E protein monomers assembled on the virus surface; EMBO Mol Med 6 (3), 358-371 (2014), NCBI Accession # 4C2I_H (232 aa)	3332
DENG7	Heavy chain variable region	9Fl 2	WO2010093335 SEQ ID NO: 4	3333
DENG8	Heavy chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	9F12	US20150218255 SEQ ID NO: 83	3334
DENG9	Heavy chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m366	US20150218255 SEQ ID NO: 4	3335
DENG10	Heavy chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m366.6	US20150218255 SEQ ID NO: 24	3336

DENG11	Heavy chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m360.6	US20150218255 SEQ ID NO: 44	3337
DENG12	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-1	US9073981 SEQ ID NO: 13	3338
DENG13	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-2	US9073981 SEQ ID NO: 29	3339
DENG14	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-3	US9073981 SEQ ID NO: 45	3340
DENG15	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-4	US9073981 SEQ ID NO: 61	3341
DENG16	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-4	US9073981 SEQ ID NO: 65	3342
DENG17	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-5	US9073981 SEQ ID NO: 79	3343
DENG18	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-6, HMB-DV-7	US9073981 SEQ ID NO: 95	3344
DENG19	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-8	US9073981 SEQ ID NO: 117	3345
DENG20	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-9	US9073981 SEQ ID NO: 131	3346
DENG21	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-10	US9073981 SEQ ID NO: 145	3347
DENG22	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-11	US9073981 SEQ ID NO: 151	3348
DENG23	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-12	US9073981 SEQ ID NO: 165	3349
DENG24	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-13	US9073981 SEQ ID NO: 181	3350
DENG25	Heavy chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-14	US9073981 SEQ ID NO: 195	3351
DENG26	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-10	A68	US20150225474 SEQ ID NO: 19	3352
DENG27	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-11	A100	US20150225474 SEQ ID NO: 20	3353
DENG28	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-12	C58	US20150225474 SEQ ID NO: 21	3354
DENG29	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-13	C98	US20150225474 SEQ ID NO: 32	3355
DENG30	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-14	A11	US20150225474 SEQ ID NO: 33	3356
DENG31	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-15	B11	US20150225474 SEQ ID NO: 36	3357

DENG32	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-4	D88	US20150225474 SEQ ID NO: 1	3358
DENG33	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-4	mAb11	WO2014144061 SEQ ID NO: 1	3359
DENG34	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-5	F38	US20150225474 SEQ ID NO: 80	3360
DENG35	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-6	A48	US20150225474 SEQ ID NO: 16	3361
DENG36	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-7	C88	US20150225474 SEQ ID NO: 17	3362
DENG37	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-8	F108	US20150225474 SEQ ID NO: 81	3363
DENG38	Heavy chain variable region, DV-1, DV-2, DV-3, and DV-9	B48	US20150225474 SEQ ID NO: 18	3364
DENG39	Heavy chain, Antigen-binding Fragment Of Human Antibody 2d22	2d22	Fibriansah, G., et al., DENGUE VIRUS. Cryo-EM structure of an antibody that neutralizes dengue virus type 2 by locking E protein dimers; Science 349 (6243), 88-91 (2015), NCBI Accession # 5A1Z_K (128 aa)	3365
DENG40	Heavy chain, Dengue virus NS-1 protein		US7473424; US20040209244; WO2004067567; EP1592712 SEQ ID NO: 3	3366
DENG41	Heavy chain, Dengue virus serotype 2	DB32-6	US8637035 SEQ ID NO: 1	3367
DENG42	Heavy chain, Dengue virus serotype 2	DB2-3	US8637035 SEQ ID NO: 13	3368
DENG43	Heavy chain, Dengue virus serotype 2	DB13-19	US8637035 SEQ ID NO: 14	3369
DENG44	Heavy chain, Dengue virus serotype 2	DB23-3	US8637035 SEQ ID NO: 15	3370
DENG45	Heavy chain, Dengue virus serotype 2	DB25-2	US8637035 SEQ ID NO: 16	3371
DENG46	Heavy chain, Dengue virus serotype 2	DB42-3	US8637035 SEQ ID NO: 17	3372
DENG47	Heavy chain, Dengue virus type 10	1A5	US8337853 SEQ ID NO: 97	3373
DENG48	Heavy chain, Dengue virus type 11	2H7	US8337853 SEQ ID NO: 113	3374
DENG49	Heavy chain, Dengue virus type 12	2H5	US8337853 SEQ ID NO: 129	3375
DENG50	Heavy chain, Dengue virus type 13	3A2	US20130089543 SEQ ID NO: 145	3376
DENG51	Heavy chain, Dengue virus type 14	1B2	US20130089543 SEQ ID NO: 161	3377
DENG52	Heavy chain, Dengue virus type 15	1A10	US20130089543 SEQ ID NO: 177	3378
DENG53	Heavy chain, Dengue virus type 4	5H2	US7622113 SEQ ID NO: 1	3379
DENG54	Heavy chain, Dengue virus type 5	5A7	US7622113 SEQ ID NO: 17	3380
DENG55	Heavy chain, Dengue virus type 6	3C1	US7622113 SEQ ID NO: 33	3381
DENG56	Heavy chain, Dengue virus type 7	3E4	US7622113 SEQ ID NO: 49	3382
DENG57	Heavy chain, Dengue virus type 8	7G4	US7622113 SEQ ID NO: 65	3383
DENG58	Heavy chain, Dengue virus type 9	5D9	US7622113 SEQ ID NO: 81	3384
DENG59	Heavy chain, DV 1	14c10 clone 8	US20130259871 Fig 4b	3385

DENG60	Heavy chain, DV-1, DV-2, DV-3, and DV-4	Antibody 4e11	US20140056913 SEQ ID NO: 1	3386
DENG61	Heavy chain, DV-1, DV-2, DV-3, and DV-4	Variant of 4E11	US20140056913 SEQ ID NO: 21	3387
DENG62	Heavy chain, DV-1, DV-2, DV-3, and DV-4	4E5A	WO20155123362 SEQ ID NO: 29	3388
DENG63	Light Chain	5j7 Fab	Fibriansah, G., et al., A highly potent human antibody neutralizes dengue virus serotype 3 by binding across three surface proteins; Nat Commun 6, 6341 (2015), NCBI Accession # 3J6U_L (118aa)	3389
DENG64	Light Chain	Ede1 C8	Dejnirattisai, W., et al., A new class of highly potent, broadly neutralizing antibodies isolated from viremic patients infected with dengue virus; Nat. Immunol. 16 (2), 170-177 (2015), NCBI Accession # 4UTA_L (217 aa)	3390
DENG65	Light Chain	Fab 2h12	Midgley, C.M., et al., J, Immunol. 188 (10), 4971-4979 (2012), NCBI Accession # 4AL8_L (213 aa)	3391
DENG66	Light Chain Fab Fragment Of Antibody 1f4	1f4 Fab	Fibriansah, G., et al., A potent anti-dengue human antibody preferentially recognizes the conformation of E protein monomers assembled on the virus surface; EMBO Mol Med 6 (3), 358-371 (2014), NCBI Accession # 4C2I_N (239 aa)	3392
DENG67	Light chain variable region	9F12	WO2010093335 SEQ ID NO: 6	3393
DENG68	Light chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	9F12	US20150218255 SEQ ID NO: 84	3394
DENG69	Light chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m366	US20150218255 SEQ ID NO: 6	3395
DENG70	Light chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m366.6	US20150218255 SEQ ID NO: 26	3396
DENG71	Light chain variable region, DENV serotype 1, DENV serotype 2, DENV serotype 3, and DENV serotype 4	m360.6	US20150218255 SEQ ID NO: 46	3397
DENG72	Light chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-1	US9073981 SEQ ID NO: 14	3398
DENG73	Light chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-2	US9073981 SEQ ID NO: 30	3399
DENG74	Light chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-3	US9073981 SEQ ID NO: 46	3400
DENG75	Light chain variable region, DENV-1, DENV-2, DENV-3, DENV-4	HMB-DV-4	US9073981 SEQ ID NO: 62	3401

DENG76	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-5	US9073981 SEQ ID NO: 80	3402
DENG77	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-6	US9073981 SEQ ID NO: 96	3403
DENG78	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-7	US9073981 SEQ ID NO: 103	3404
DENG79	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-8	US9073981 SEQ ID NO: 118	3405
DENG80	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-9	US9073981 SEQ ID NO: 132	3406
DENG81	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-10, HMB-DV-11	US9073981 SEQ ID NO: 146	3407
DENG82	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-12	US9073981 SEQ ID NO: 166	3408
DENG83	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-13	US9073981 SEQ ID NO: 182	3409
DENG84	Light chain variable region, DENV-I, DENV-2, DENV-3, DENV-4	HMB-DV-14	US9073981 SEQ ID NO: 196	3410
DENG85	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	D88, F38, A48, C88, F108, B48, A68, A100, C58, C78, C68, D98, D188, C128, C98	US20150225474 SEQ ID NO: 2	3411
DENG86	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	C78	US20150225474 SEQ ID NO: 23	3412
DENG87	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	C68	US20150225474 SEQ ID NO: 25	3413
DENG88	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	D98	US20150225474 SEQ ID NO: 27	3414
DENG89	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	D188	US20150225474 SEQ ID NO: 29	3415
DENG90	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	C128	US20150225474 SEQ ID NO: 31	3416
DENG91	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	A11, B11	US20150225474 SEQ ID NO: 34	3417
DENG92	Light chain variable region, DV-1, DV-2, DV-3, and DV-4	mAb11	WO2014144061 SEQ ID NO: 3	3418
DENG93	Light chain, Antigen-binding Fragment Of Human Antibody 2d22	2d22	Fibriansah, G., et al., DENGUE VIRUS. Cryo-EM structure of an antibody that neutralizes dengue virus type 2 by locking E protein dimers; Science 349 (6243), 88-91 (2015), NCBI Accession # 5A1Z_L (115 aa)	3419
DENG94	Light chain, Dengue virus NS-1 protein		US7473424; US20040209244; WO2004067567; EP1592712 SEQ ID NO: 4	3420

DENG95	Light chain, Dengue vims sero type 2	DB32-6	US8637035 SEQ ID NO: 5	3421
DENG96	Light chain, Dengue virus serotype 2	DB2~3, DB-19	US8637035 SEQ ID NO: 19	3422
DENG97	Light chain, Dengue virus sero type 2.	DB23-3	US8637035 SEQ ID NO: 20	3423
DENG98	Light chairs, Dengue vims serotype 2	DB25-2	US8637035 SEQ ID NO: 21	3424
DENG99	Light chain, Dengue virus serotype 2.	DB42-3	US8637035 SEQ ID NO: 22	3425
DENG 100	Light chairs, Dengue vims serotype 4	5H2	US7622113 SEQ ID NO: 9	3426
DENG101	Light chain, Dengue virus serotype 4	5A7	US76221 13 SEQ ID NO: 25	3427
DENG 102	Light chairs, Dengue vims serotype 4	3C1	US7622113 SEQ ID NO: 41	3428
DENG103	Light chain, Dengue virus serotype 4	3E4	US76221 13 SEQ ID NO: 57	3429
DENG 104	Light chairs, Dengue vims serotype 4	7G4	US7622113 SEQ ID NO: 73	3430
DENG105	Light chain, Dengue virus serotype 4	5D9	US76221 13 SEQ ID NO: 89	3431
DENG 106	Light chain, Dengue virus serotype 4	1A5	US8337853 SEQ ID NO: 105	3432
DENG 107	Light chain. Dengue virus serotype 4	2H7	US8337853 SEQ ID NO: 121	3433
DENG108	Light chain, Dengue virus serotype 4	2H5	US8337853 SEQ ID NO: 137	3434
DENG109	Light chain. Dengue virus serotype 4	3A2	US20130089543 SEQ ID NO: 153	3435
DENG 110	Light chain, Dengue virus serotype 4	1B2	US20130089543 SEQ ID NO: 169	3436
DENG! 11	Light chain. Dengue virus serotype 4	1A10	US20130089543 SEQ ID NO: 185	3437
DENG 112	Light chain, DV 1	14c10 clone 8	US20130259871 Fig 4b	3438
DENG! 13	Light chain, DV-1, DV-2. DV-3, and DV-4	Antibody 4el 1	US20140056913 SEQ ID NO: 2	3439
DENG1 14	Light chain, DV-1, DV-2, DV-3, and DV-4	Variant of 4E1 1	US20140056913 SEQ ID NO: 22	3440
DENG1 15	Light chain, DV-1, DV-2, DV-3, and DV-4	4E5A	WO20155 123362 SEQ ID NO: 30	3441
DENG 116	scFv	9F1 2	WO2010093335 SEQ ID NO: 8	3442
DENG 117	Scfv Fragment	Ede2 A11	Dejnirattisai, W., et al., A new class of highly potent, broadly neutralizing antibodies isolated from viremic patients infected with dengue vims; Nat. Immunol. 16 (2), I70- I77 (20 15), NCBI Accession # 4UT7_L(153 aa)	3443
DENG1 18	Scfv Fragment	Ede2 A11	Dejnirattisai, W., et al., A new class of highly potent, broadly neutralizing antibodies isolated from viremic patients infected with dengue vims; Nat. Immunol. 16 (2), 170-177 (2015), NCB1 Accession # 4UT7_H (150 aa)	3444
DENG 119		Ede2 A11	Dejnirattisai, W., et al., A new class of highly potent, broadly	3445

			neutralizing antibodies isolated **from viremic patients** infected with dengue **vims;** Nat. **Immunol.** 16 (2), 170-177 (2015), NCBI Accession # **4UTB_L** (218 aa)	
DENG 120		Edel CIO	**Dejnirattisai, W.,** et al., A new class **of highly** potent, broadly neutralizing antibodies isolated from **viremic** patients infected with dengue virus; Nat. Immunol. 16 (2), 170-177 (2015), NCBI **Accession # 4UT9_L (154aa)**	3446
DENG! 21		Ede l CIO	Dejnirattisai, **W.,** et al, **A** new class of highly potent, broadly neutralizing antibodies isolated from viremic patients infected with **dengue virus;** Nat. Immunol. 16 (2), 170-177 (2015), NCBI Accession # 4UT9_H **(144** aa)	3447
DENG 122		Ede2 B7	Dejnirattisai, W., et **al.,** A **new** class **of highly** potent, broadly neutralizing antibodies isolated from viremic patients infected with dengue vims; Nat. Immunol. 16 (2), 170-177 (2015), NCBI Accession # **4UT6_L** (218 aa)	3448
DENG 123		Ede2 B7	Dejnirattisai, W., et al., **A** new class **of highly** potent, broadly neutralizing antibodies isolated from viremic patients infected with dengue virus; Nat. Immunol. 16 (2), 170-177 (2015), NCBI Accession # 4UT6_H (283 aa)	3449

[00296] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides, fragments or variants thereof described in International Pub. No. WO2013089647 and WO2013035345, US Patent No. US8637035 and US887187, US Publication No. US20050123900, and Chinese Patent Publication No. CN102757480, the contents of which are herein incorporated by reference in their entirety, against listeria monocytogenes, salmonella and/or ieishniania.

[00297] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 11 against Rabies Virus.

Table 11. Antibodies agaisist Rabies Vims

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
RABV1	Fab Heavy Chain Fd region		CN101696242 SEQ ID NO: 9	3450
RABV2	Fab Light chain		CN101696242 SEQ ID NO: 10	3451
RABV3	Heavy chain		US6890532 SEQ ID NO: 3	3452
RABV4	Heavy chain	Mab JB.1	US7071319 SEQ ID NO: 10	3453
RABV5	Heavy chain	Mab 57	US7071319 SEQ ID NO: 14	3454

RABV6	Heavy chain	CR04-098	US9005624 SEQ ID NO: 335	3455
RABV7	Heavy chain	CR57, Rafivirumab	US9005624 SEQ ID NO: 123	3456
RABV8	Heavy chairs	CR57, Rafivirumab		3457
RABV9	Heavy chain	CRJB	US9005624 SEQ ID NO: 127	3458
RABViO	Heavy chain	Foraviriimab		3459
RABV1 l	Heavy chain, Anti-rabies SOJB immunoglobulin		Prosniak, M. et al. "Development of a cocktail of recombinant-expressed human rabies virus-neutralizing monoclonal antibodies for postexposure prophylaxis of rabies", J. Infect. Dis. 188 (1). 53-56 (2003), NCBI Accession # AAO 17822. l	3460
RABV12	Heavy chain variable region		CNIO 1696242 SEQ ID NO: 4	3461
RABV13	Heavy chain variable region	SC04-00 i	US9005624 SEQ ID NO: 26	3462
RABV14	Heavy chairs variable region	SC04-004	US9005624 SEQ ID NO: 27	3463
RABV15	Heavy chain variable region	SC04-008	1JS9005624 SEQ ID NO: 28	3464
RABV16	Heavy chain variable region	SC04-010	US9005624 SEQ ID NO: 29	3465
RABV17	Heavy chain variable region	SC04-018	US9005624 SEQ ID NO: 30	3466
RABV18	Heavy chain variable region	SC04-021	US9005624 SEQ ID NO: 31	3467
RABV19	Heavy chairs variable region	SC04-026	US9005624 SEQ ID NO: 32	3468
RABV20	Heavy chain variable region	5C04-03 1	US9005624 SEQ ID NO: 33	3469
RABV21	Heavy chain variable region	SC04-038	US9005624 SEQ ID NO: 34	3470
RABV22	Heavy chain variable region	SC04-040	US9005624 SEQ ID NO: 35	3471
RABV23	Heavy chain variable region	SC04-060	US9005624 SEQ ID NO: 36	3472
RABV24	Heavy chairs variable region	SC04-073	US9005624 SEQ ID NO: 37	3473
RABV25	Heavy chain variable region	SC04-097	US9005624 SEQ ID NO: 38	3474
RABV26	Heavy chain variable region	SC04-098	US9005624 SEQ ID NO: 39	3475
RABV27	Heavy chain variable region	SC04-103	US9005624 SEQ ID NO: 40	3476
RABV28	Heavy chain variable region	SC04-104	US9005624 SEQ ID NO: 41	3477
RABV29	Heavy chain variable region	SC04-108	US9005624 SEQ ID NO: 42	3478
RABV30	Heavy chain variable region	SC04-120	US9005624 SEQ ID NO: 43	3479
RABV3 1	Heavy chain variable region	SC04-125	US9005624 SEQ ID NO: 44	3480
RABV32	Heavy chain variable region	SC04-126	1JS9005624 SEQ ID NO: 45	3481
RABV33	Heavy chain variable region	SC04-140	US9005624 SEQ ID NO: 46	3482
RABV34	Heavy chain variable region	SC04-144	US9005624 SEQ ID NO: 47	3483
RABV35	Heavy chain variable region	SC04-S.46	US9005624 SEQ ID NO: 48	3484
RABV36	Heavy chain variable region	SC04-164	US9005624 SEQ ID NO: 49	3485
RABV37	Heavy chain variable region	RVFab5	WO201 113757 SEQ ID NO: 2	3486
RABV38	Heavy chain variable region	RVFab8	WO20 11137570 SEQ ID NO: 2	3487
RABV39	Heavy chain variable region		CN101337990 SEQ ID NO: 2	3488
RABV40	Heavy chain variable region		CN101337990 SEQ ID NO: 8	3489
RABV4 l	Heavy chairs variable region	R8 VH	CN 104 l 93823 SEQ ID NO: 1	3490
RABV42	Heavy chain variable region	R5 VH	CN104193823 SEQ ID NO: 2	3491

RABV43	***Heavy*** chain variable region	R7 VH, **R9 VH**	**CN104 193823** SEQ ID NO: 3	3492
RABV44	Heavy **chain variable** region		**CN10 1235086** SEQ **ID NO:** 38	3493
RABV45	Heavy chairs, Anti-rabies SOJA **immunoglobulin**		Prosniak, **M. et al.** "**Development** of a cocktail of recombinant-expressed human rabies virus-neutralizing monoclonal antibodies for postexposure prophylaxis of rabies", J. Infect. Dis. 188 (1), 53-56 (2003), NCBI Accession # **AAO 17823. 1**	3494
RABV46	Light chain		**US6890532** SEQ **ID NO**: 4	3495
RABV47	Light chain	**Mab JB. l**	**US70713 19 SEQ** ID NO: 12	3496
RABV48	Light chain	**Mab** 57	**US70713 19** SEQ **ID** NO: 16	3497
RABV49	Light chain	CR04-001	US9005624 SEQ ID NO: 337	3498
RABV50	Light **chain**	CR57, **Rafivirumab**	**US9005624** SEQ ID NO: 125	3499
RABV5 1	Light chain	CR57, Rafivirumab		3500
RABV52	Light **chain**	CR.IB	**US9005624** SEQ ID NO: 129	3501
RABV53	Light chain	**Foravirumab**		3502
RABV54	Light chain Kappa, Anti-rabies SOJA **immunoglobulin**		Prosniak, **M.** et **al.** "Development **of a** cocktail of recombinant-expressed human rabies virus-neutralizing monoclonal antibodies for postexposure prophylaxis of rabies", J. Infect. Dis. 188 (1), 53-56 (2003), NCBI Accession # AAO 17825. l	3503
RABV55	Light chain kappa, Anti-rabies SOJA immunoglobulin [Homo sapiens]		Prosniak, M. et **al.** "Development **of** a cocktail of recombinant-expressed human rabies virus-neutralizing monoclonal antibodies for postexposure prophylaxis of rabies", J. Infect. Dis. 188 (1), 53-56 (2003), NCBI Accession # **AAO 1782 1.1**	3504
RABV56	Light chain Lambda, Anti-rabies S057 **immunoglobulin**		Prosniak, M. et **al.** "Development **of** a cocktail of recombinant-expressed human rabies virus-neutralizing monoclonal antibodies for postexposure prophylaxis **of** rabies", J. Infect. Dis. 188 (1). 53-56 (2003), NCBI Accession # AAO 17824. l	3505
RABV57	Light chain lambda, Anti-rabies SOJB **immunoglobulin**		Prosniak, M. et **al.** "Development **of** a cocktail of recombinant-expressed human rabies virus-neutralizing monoclonal antibodies for postexposure prophylaxis **of** rabies", J. Infect. Dis. 188 (1). 53-56 (2003), NCBI Accession # AAO 17826. l	3506

RABV58	Light chain variable region	SC04-001	US9005624 SEQ ID NO: 50	3507
RABV59	Light chain variable regiors	SC04-004	US9005624 SEQ ID NO: 51	3508
RABV60	Light chain variable region	SC04-008	US9005624 SEQ ID NO: 52	3509
RABV61	Light chain variable region	SC04-010	US9005624 SEQ ID NO: 53	3510
RABV62	Light chain variable region	SC04-018	US9005624 SEQ ID NO: 54	3511
RABV63	Light chain variable region	SC04-021	US9005624 SEQ ID NO: 55	3512
RABV64	Light chain variable regiors	SC04-026	US9005624 SEQ ID NO: 56	3513
RABV65	Light chain variable region	5C04-031	US9005624 SEQ ID NO: 57	3514
RABV66	Light chain variable region	SC04-038	US9005624 SEQ ID NO: 58	3515
RABV67	Light chain variable region	SC04-040	US9005624 SEQ ID NO: 59	3516
RABV68	Light chain variable region	SC04-060	US9005624 SEQ ID NO: 60	3517
RABV69	Light chain variable regiors	SC04-073	US9005624 SEQ ID NO: 61	3518
RABV70	Light chain variable region	SC04-097	US9005624 SEQ ID NO: 62	3519
RABV71	Light chain variable region	SC04-098	US9005624 SEQ ID NO: 63	3520
RABV72	Light chain variable region	SC04-103	US9005624 SEQ ID NO: 64	3521
RABV73	Light chain variable region	SC04-104	US9005624 SEQ ID NO: 65	3522
RABV74	Light chain variable region	SC04-108	US9005624 SEQ ID NO: 66	3523
RABV75	Light chain variable region	SC04-120	US9005624 SEQ ID NO: 67	3524
RABV76	Light chain variable region	SC04-125	US9005624 SEQ ID NO: 68	3525
RABV77	Light chain variable region	SC04-126	US9005624 SEQ ID NO: 69	3526
RABV78	Light chain variable region	SC04-140	US9005624 SEQ ID NO: 70	3527
RABV79	Light chain variable region	SC04-144	US9005624 SEQ ID NO: 71	3528
RABV80	Light chain variable region	SC04-146	US9005624 SEQ ID NO: 72	3529
RABV81	Light chain variable region	SC04-164	US9005624 SEQ ID NO: 73	3530
RABV82	Light chain variable region	RVFabS	WO20 1113757 SEQ ID NO: 1	3531
RABV83	Light chain variable region	RVFahS	WO20 11137570 SEQ ID NO: 1	3532
RABV84	Light chain variable region		CN101337990 SEQ ID NO: 4	3533
RABV85	Light chain variable region		CN101337990 SEQ I DNO: 10	3534
RABV86	Light chain variable region	R8VL	CN104193823 SEQ ID NO: 4	3535
RABV87	Light chain variable region	R5 VL	CN 104193823 SEQ ID NO: 5	3536
RABV88	Light chain variable region	R7 VL	CN104193823 SEQ ID NO: 6	3537
RABV89	Light chain variable region	R9 VL	CN104193823 SEQ ID NO: 7	3538
RABV90	Light chain variable region		CN10 1696242 SEQ ID NO: 8	3539
RABV91	Light chain variable region		CN101235086 SEQ ID NO: 39	3540

[00298] In one embodiment, the pay load region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 12 against Chagas Virus.

Table 12. Antibodies against Chagas Virus

Antibody No.	Description	Reference Information	SEQ ID NO
CHAG1	Heavy Chain Of The Fab Fragment, Trypanosoma cruzi trans-sialidase	Buschiazzo et al., PLoS Pathol. 8 (1), E1002474 (2012), NCBI Accession # 3OPZ_J (222aa)	3541
CHAG2	Light chain of Fab fragment, Trypanosoma cruzi trans-sialidase	Buschiazzo et al., PLoS Pathog. 8 (1), E1002474 (2012), NCBI Accession # 3OPZ_N (213aa)	3542

[00299] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 13 against Chikungunya Vims.

Table 13. Antibodies against Chikungunya Virus

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
CHIK1	Heavy chain Fab fragment	9.8b	Sun, S. et al., Structural analyses at pseudo atomic resolution of Chikungunya virus and antibodies show mechanisms of neutralization, Elife 2, E00435 (2013), NCBI Accession # 4GQ9_H (218 aa)	3543
CHIK2	Heavy chain variable	5F10F175E2	US20130189279 SEQ ID NO: 6	3544
CHIK3	Heavy chain variable	8B10F8	US20130189279 SEQ ID NO: 26	3545
CHIK4	Light Chain Fab fragment	9.8b	Sun, S. et al., Structural analyses at pseudo atomic resolution of Chikungunya virus and antibodies show mechanisms of neutralization, Elife 2, E00435 (2013), NCBI Accession # 4GQ9_L (212 aa)	3546
CHIK5	Light chain variable	5F10F175E2	US20130189279 SEQ ID NO: 8	3547
CHIK6	Light chain variable	8B10F8	US20130189279 SEQ ID NO: 28	3548

[00300] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof encoding antibodies described International Pub No. WO 1983001785 and US Patent No. US5827671, the contents of each of which are herein incorporated by reference in their entirety, against the protozoan parasite Leishniania.

[00301] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof encoding antibodies against the Buruli ulcer (Mycobacterium uicerans), Leprosy/Hansen's disease (Mycobacterium leprae). Leishmaniasis, Cysticercosis, Dracunculiasis (GtuneaWorra Disease), Echinococcosis, Fascioliasis, Human African Trypanosomiasis (African Sleeping Sickness), Lymphatic filariasis. Onchocerciasis, Schistosomiasis, Soil-transmitted Helminths (STH).

Toxins

[00302] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the toxin related payload antibody polypeptides listed in Tables 14-17.

[00303] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 14 against Ricin Toxin.

Table 14. Antibodies against Ricin Toxin

Antibody No.	Description	Atsiibody Name	Reference **Information**	SEQ **ID** NO
RICN1	Camelid heavy-chairs only	RTA ; JIV-F5	WO20 15 100409 SEQ ID NO: 324	3549
RICN2	Camelid heavy-chain only	JIV-F6	WO20 15 100409 SEQ ID NO: 126	3550
RICN3	Camelid heavy-chain only	JIV-G 12	WO20 15 100409 SEQ ID NO: 128	3551
RICN4	Camelid heavy-chain only	JIY-A7	WO20 15 100409 SEQ ID NO: 130	3552
RICN5	Camelid heavy-chain only	JIY-D9	WO20 15 100409 SEQ ID NO: 132	3553
RICN6	Camelid heavy-chain only	JIY-D10	WO20 15 100409 SEQ ID NO: 134	3554
RICN7	Camelid heavy-chain only	JIY-E1	WO2015 100409 SEQ ID NO: 136	3555
RICN8	Camelid heavy-chain only	JIY-E3	WO20 15 100409 SEQ ID NO: 138	3556
RICN9	Camelid heavy-chain only	JIY-E5	WO20 15 100409 SEQ ID NO: 140	3557
RICN10	Camelid heavy-chain only	JIY-F10	WO20 15 100409 SEQ ID NO: 142	3558
RICN11	Camelid heavy-chain only	JIY-G11	WO20 15 100409 SEQ ID NO: 144	3559
RICN12	Camelid heavy-chain only	RTB: JIW-B 1	WO2015 100409 SEQ ID NO: 146	3560
RICN13	Camelid heavy-chain only	JIW-C12	WO2015 100409 SEQ ID NO: 148	3561
RICN1 4	Camelid heavy-chain only	JIW-D 12	WO20 15 100409 SEQ ID NO: 150	3562
RICN15	Camelid heavy-chain only	JIW-G5	WO2015 100409 SEQ ID NO: 152	3563
RICN 16	Camelid heavy-chain only	JIW-G 10	WO20 15 100409 SEQ ID NO: 154	3564
RICN17	Camelid heavy-chain only	JIZ-B7	WO2015 100409 SEQ ID NO: 156	3565
RICN18	Camelid heavy-chain only	JIZ- B9	WO2015100409 SEQ ID NO: 158	3566
RICN19	Camelid heavy-chain only	JIZ-D8	WO2015 100409 SEQ ID NO: 160	3567
RICN20	Camelid heavy-chain only	JIZ-G4	WO2015 100409 SEQ ID NO: 162	3568

[00304] In one embodiment, the pay load region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 15 against Anthrax.

Table 15. Antibodies against Anthrax

Antibody No.	**Description**	**Antibody Name**	**Reference Information**	**SEQ ID NO**
ANTH1	Camelid heavy-chain only	JHD-B6	WO2015100409 SEQ ID NO: 100	3569
ANTH2	Camelid heavy-chain only	JHE-D9	WO2015100409 SEQ ID NO: 102	3570
ANTH3	Camelid heavy-chain only	JIJ-A12	WO2015100409 SEQ ID NO: 104	3571
ANTH4	Camelid heavy-chain only	JIJ-B8	WO2015100409 SEQ ID NO: 106	3572
ANTH5	Camelid heavy-chain only	JIJ-C11	WO2015100409 SEQ ID NO: 108	3573
ANTH6	Camelid heavy-chain only	JIJ-D3	WO2015100409 SEQ ID NO: 110	3574
ANTH7	Camelid heavy-chain only	JIJ-E9	WO2015100409 SEQ ID NO: 112	3575
ANTH8	Camelid heavy-chain only	JIJ-F11	WO2015100409 SEQ ID NO: 114	3576
ANTH9	Camelid heavy-chain only	JIK-B8	WO2015100409 SEQ ID NO: 116	3577
ANTH10	Camelid heavy-chain only	JI -B 10	WO2015100409 SEQ ID NO: 118	3578
ANTH11	Camelid heavy-chain only	JIK-B 12	WO2015100409 SEQ ID NO: 120	3579
ANTH12	Camelid heavy-chain only	JIK-F4	WO2015100409 SEQ ID NO: 122	3580
ANTH13	CDR		WO2003063768 SEQ ID NO: 1	3581
ANTH14	CDR		WO2003063768 SEQ ID NO: 2	3582

ANTH15	CDR		WO2003063768 SEQ ID NO: 3	3583
ANTH16	Heavy chain		US8617548 SEQ ID NO: 2	3584
ANTH17	Heavy chain	IQNPA Lambda	US7658925 SEQ ID NO: 2	3585
ANTH18	Heavy chain	IQNLF Lambda	US7658925 SEQ ID NO: 6	3586
ANTH19	Heavy chain	1A5	US20090022736 SEQ ID NO: 1	3587
ANTH20	Heavy chain	4A12	US20090022736 SEQ ID NO: 3	3588
ANTH21	Heavy chain	24B1	US20090022736 SEQ ID NO: 5	3589
ANTH22	Heavy chain	24G4	US20090022736 SEQ ID NO: 7	3590
ANTH23	Heavy chain	32E12	US20090022736 SEQ ID NO: 9	3591
ANTH24	Heavy chain	33F4	US20090022736 SEQ ID NO: 11	3592
ANTH25	Heavy chain	scFv 2LF	EP2778173 SEQ ID NO: 9	3593
ANTH26	Heavy chain		US20040258699 SEQ ID NO: 78	3594
ANTH27	Heavy chain		US20040258699 SEQ ID NO: 79	3595
ANTH28	Heavy chain		US20040258699 SEQ ID NO: 80	3596
ANTH29	Heavy chain		US20040258699 SEQ ID NO: 81	3597
ANTH30	Heavy chain		US20040258699 SEQ ID NO: 82	3598
ANTH31	Heavy chain		US20040258699 SEQ ID NO: 83	3599
ANTH32	Heavy chain		US20040258699 SEQ ID NO: 84	3600
ANTH33	Heavy chain		US20040258699 SEQ ID NO: 85	3601
ANTH34	Heavy chain		US20040258699 SEQ ID NO: 86	3602
ANTH35	Heavy chain		US20040258699 SEQ ID NO: 87	3603
ANTH36	Heavy chain		US20040258699 SEQ ID NO: 88	3604
ANTH37	Heavy chain		US20040258699 SEQ ID NO: 89	3605
ANTH38	Heavy chain		US20040258699 SEQ ID NO: 90	3606
ANTH39	Heavy chain		US20040258699 SEQ ID NO: 91	3607
ANTH40	Heavy chain		US20040258699 SEQ ID NO: 92	3608
ANTH41	Heavy chain		US20040258699 SEQ ID NO: 93	3609
ANTH42	Heavy chain		US20040258699 SEQ ID NO: 94	3610
ANTH43	Heavy chain		US20040258699 SEQ ID NO: 95	3611
ANTH44	Heavy chain		US20040258699 SEQ ID NO: 96	3612
ANTH45	Heavy chain		US20040258699 SEQ ID NO: 97	3613
ANTH46	Heavy chain		US20040258699 SEQ ID NO: 98	3614
ANTH47	Heavy chain		US20040258699 SEQ ID NO: 99	3615
ANTH48	Heavy chain		US20040258699 SEQ ID NO: 100	3616
ANTH49	Heavy chain		US20040258699 SEQ ID NO: 101	3617
ANTH50	Heavy chain		US20040258699 SEQ ID NO: 102	3618
ANTH51	Heavy chain		US20040258699 SEQ ID NO: 103	3619
ANTH52	Heavy chain		US20040258699 SEQ ID NO: 104	3620
ANTH53	Heavy chain		US20040258699 SEQ ID NO: 105	3621
ANTH54	Heavy chain		US20040258699 SEQ ID NO: 106	3622
ANTH55	Heavy chain		US20040258699 SEQ ID NO: 107	3623
ANTH56	Heavy chain		US20040258699 SEQ ID NO: 108	3624
ANTH57	Heavy chain		US20040258699 SEQ ID NO: 109	3625
ANTH58	Heavy chain		US20040258699 SEQ ID NO: 110	3626
ANTH59	Heavy chain		US20040258699 SEQ ID NO: in	3627

ANTH60	Heavy chain		US20040258699 SEQ ID NO: 112	3628
ANTH61	Heavy chain		US20040258699 SEQ ID NO: 113	3629
ANTH62	Heavy chain		US20040258699 SEQ ID NO: 114	3630
ANTH63	Heavy chain		US20040258699 SEQ ID NO: 115	3631
ANTH64	Heavy chain		US20040258699 SEQ ID NO: 116	3632
ANTH65	Heavy chain		US20040258699 SEQ ID NO: 117	3633
ANTH66	Heavy chain		US20040258699 SEQ ID NO: 118	3634
ANTH67	Heavy chain and light chain variable region	14B7 scFv	US7902344; US6916474 SEQ ID NO: 21	3635
ANTH68	Heavy chain fd region	W1	US8685396 SEQ ID NO: 1	3636
ANTH69	Heavy chain fd region	W2	US8685396 SEQ ID NO: 17	3637
ANTH70	Heavy chain region	W5	US8685396 SEQ ID NO: 33	3638
ANTH71	Heavy chain region	A63-6	US8685396 SEQ ID NO: 34	3639
ANTH72	Heavy chain region	F3-6	US8685396 SEQ ID NO: 35	3640
ANTH73	Heavy chain region	F5-1	US8685396 SEQ ID NO: 36	3641
ANTH74	Heavy chain variable regiors	ETI-204	US20120156196 SEQ ID NO: 1	3642
ANTH75	Heavy chain variable region	6.20	WO2015107307 SEQ ID NO: 1	3643
ANTH76	Heavy chain variable region	35PA83	WO2009071860 SEQ ID NO: 1	3644
ANTH77	Heavy chain variable region	anti-yDPGA antibody	US8501182 SEQ ID NO: 1	3645
ANTH78	Heavy chain variable region	4C	US8501182 SEQ ID NO: 3	3646
ANTH79	Heavy chain variable region	11D	US8501182 SEQ ID NO: 5	3647
ANTH80	Heavy chain variable region	F20G75	WO200713 1363 SEQ ID NO: 16	3648
ANTH81	Heavy chain variable region	F20G76	WO20071 31363 SEQ ID NO: 18	3649
ANTH82	Heavy chain variable region	F20G77	WO200713 1363 SEQ ID NO: 20	3650
ANTH83	Heavy chain variable region	V2 variant	US8507655 SEQ ID NO: 7	3651
ANTH84	Heavy chain variable region	6.20 variant	US8507655 SEQ ID NO: 9	3652
ANTH85	Heavy chain variable region	J24.15 variant	US8507655 SEQ ID NO: 11	3653
ANTH86	Heavy chain variable region	J24.7 variant	US8507655 SEQ ID NO: 13	3654
ANTH87	Heavy chain variable region	V2 variant human	US8507655 SEQ ID NO: 15	3655
ANTH88	Heavy chain variable region	6.20 variant human	US8507655 SEQ ID NO: 17	3656
ANTH89	Heavy chain variable region	J24. 15 variant human	US8507655 SEQ ID NO: 19	3657
ANTH90	Heavy chain variable region	J24.7 variant human	US8507655 SEQ ID NO: 21	3658
ANTH9 I	Heavy chain variable region	HisMab 5E8	US8404820 SEQ ID NO: 2	3659
ANTH92	Heavy chain variable region	HuMab 2D5	US8404820 SEQ ID NO: 8	3660
ANTH93	Heavy chain variable regiors	HuMab 2H4	US8404820 SEQ ID NO: 12	3661
ANTH94	Heavy chain variable region	HuMab 5D5-2E10	US8404820 SEQ ID NO: 16	3662
ANTH95	Heavy chain variable region	13E3	US8309090 SEQ ID NO: 2	3663
ANTH96	Heavy chain variable region	3E1	US8309090 SEQ ID NO: 6	3664
ANTH97	Heavy chain variable region	KCTC 10756BP	US82683 16 SEQ ID NO: 2	3665
ANTH98	Heavy chain variable region	M18 scFv	US7902344; US6916474 SEQ ID NO: 23	3666
ANTH99	Heavy chain variable region	21D9 MAb	US7442373 SEQ ID NO: 2	3667
ANTH100	Heavy chain variable regiors	1C6 Mab	US7442373 SEQ ID NO: 6	3668

ANTH101	Heavy chain variable region	4H7 Mab	**US7442373** SEQ ID NO: 10	3669
ANTH102	**Heavy** chain variable region	22G12 Mab	**US7442373** SEQ ID NO: 14	3670
ANTH103	Heavy chain variable region	monoclonal antibody 9-1	**WO1999055842** SEQ ID **NO:** 20	3671
ANTH104	Heavy chain variable region	**monoclonal antibody 7-1**	WO1999055842 SEQ ID NO: 21	3672
ANTH105	Heavy chain variable region	monoclonal antibody 24-2	WO1999055842 SEQ ID **NO:** 22	3673
ANTH106	Heavy chain variable region	**monoclonal** antibody 21-4	WO1999055842 SEQ ID NO: 23	3674
ANTH107	Heavy **chain variable** regiors	monoclonal antibody **10-2**	WO1999055842 SEQ ID **NO:** 24	3675
ANTH108	Heavy chain variable region	**monoclonal** antibody **22-1**	WO1999055842 SEQ ID NO: 25	3676
ANTH109	Heavy **chain variable** regiors	monoclonal antibody **13-3**	**WO1999055842** SEQ ID NO: 26	3677
ANTH1 10	Heavy chain variable region	monoclonal antibody 8-3	WO1999055842 SEQ ID NO: 27	3678
ANTH1 11	Heavy **chain variable** regiors	monoclonal antibody 6-1	**WO1999055842** SEQ ID NO: 29	3679
ANTH1 12	Heavy chain variable region	monoclonal antibody 3-1	WO1999055842 SEQ ID NO: 30	3680
ANTH113	Heavy chain variable region, Edema factor binding	EF12A	**US8961975** SEQ ID NO: 51	3681
ANTH1 14	Heavy chain variable region, **Edema** factor binding	EF13D	US8961975 SEQ ID **NO:** 33	3682
ANTH115	Heavy chain variable region, Edema factor binding	EF14H	**US8961975** SEQ ID NO: 52	3683
ANTH1 16	Heavy chain variable region, **Edema** factor binding	EF15A	US8961975 SEQ ID **NO:** 53	3684
ANTH117	Heavy chain variable region, Lethal factor	LF9D	**US8961975** SEQ ID NO: 49	3685
ANTH118	Heavy chain variable region, Lethal **factor**	**LF10E**	**US8961975** SEQ **ID NO:** 1	3686
ANTH119	Heavy chain, Antibody **against inhalational** anthrax	**Obiltoxaximab**		3687
ANTH120	Kappa light **chain**		**US20040258699** SEQ ID NO: 19	3688
ANTH121	Kappa light chain		**US20040258699** SEQ ID NO: 20	3689
ANTH122	Kappa light **chain**		**US20040258699** SEQ ID NO: 21	3690
ANTH123	Kappa light chairs		US20040258699 SEQ ID NO: 22	3691
ANTH124	Kappa **light** chain		**US20040258699** SEQ ID **NO:** 23	3692
ANTH **125**	Kappa **light** chain		**US20040258699** SEQ ID NO: 24	3693
ANTH126	Kappa light chain		**US20040258699** SEQ ID NO: 25	3694
ANTH127	Kappa light **chain**		**US20040258699** SEQ ID NO: 26	3695
ANTH128	Kappa light chairs		US20040258699 SEQ ID NO: 39	3696
ANTH129	Kappa light chain		**US20040258699** SEQ ID NO: 40	3697
ANTH130	Kappa **light** chain		**US20040258699** SEQ ID NO: 41	3698
ANTH131	Kappa light chain		**US20040258699** SEQ ID NO: 42	3699
ANTH132	Kappa light **chain**		**US20040258699** SEQ ID NO: 43	3700
ANTH133	Kappa light chairs		US20040258699 SEQ ID **NO:** 44	3701
ANTH134	Kappa light chain		**US20040258699** SEQ ID NO: 45	3702
ANTH135	Kappa **light** chain		**US20040258699** SEQ ID NO: 46	3703
ANTH136	Kappa light chain		**US20040258699** SEQ ID NO: 47	3704

ANTH137	Kappa light chain		US20040258699 SEQ ID NO: 48	3705
ANTH138	Kappa light chain		US20040258699 SEQ ID NO: 49	3706
ANTH139	Kappa light chain		US20040258699 SEQ ID NO: 50	3707
ANTH140	Kappa light chairs		US20040258699 SEQ ID NO: 51	3708
ANTH141	Kappa light chain		US20040258699 SEQ ID NO: 52	3709
ANTH142	Kappa light chain		US20040258699 SEQ ID NO: 53	3710
ANTH143	Kappa light chain		US20040258699 SEQ ID NO: 54	3711
ANTH144	Kappa light chain		US20040258699 SEQ ID NO: 55	3712
ANTH145	Kappa light chairs		US20040258699 SEQ ID NO: 56	3713
ANTH146	Kappa light chain		US20040258699 SEQ ID NO: 57	3714
ANTH147	Kappa light chain		US20040258699 SEQ ID NO: 58	3715
ANTH148	Kappa light chain		US20040258699 SEQ ID NO: 59	3716
ANTH149	Kappa light chain		US20040258699 SEQ ID NO: 60	3717
ANTH150	Kappa light chairs		US20040258699 SEQ ID NO: 61	3718
ANTH151	Lambda light chain		US20040258699 SEQ ID NO: 27	3719
ANTH152	Lambda light chain		US20040258699 SEQ ID NO: 28	3720
ANTH153	Lambda light chain		US20040258699 SEQ ID NO: 29	3721
ANTH154	Lambda light chain		US20040258699 SEQ ID NO: 30	3722
ANTH155	Lambda light chain		US20040258699 SEQ ID NO: 31	3723
ANTH156	Lambda light chain		US20040258699 SEQ ID NO: 32	3724
ANTH157	Lambda light chain		US20040258699 SEQ ID NO: 33	3725
ANTH158	Lambda light chain		US20040258699 SEQ ID NO: 34	3726
ANTH159	Lambda light chain		US20040258699 SEQ ID NO: 35	3727
ANTH160	Lambda light chain		US20040258699 SEQ ID NO: 36	3728
ANTH161	Lambda light chain		US20040258699 SEQ ID NO: 37	3729
ANTH162	Lambda light chain		US20040258699 SEQ ID NO: 38	3730
ANTH163	Lambda light chain		US20040258699 SEQ ID NO: 62	3731
ANTH164	Lambda light chain		US20040258699 SEQ ID NO: 63	3732
ANTH165	Lambda light chain		US20040258699 SEQ ID NO: 64	3733
ANTH166	Lambda light chain		US20040258699 SEQ ID NO: 65	3734
ANTH167	Lambda light chain		US20040258699 SEQ ID NO: 66	3735
ANTH168	Lambda light chain		US20040258699 SEQ ID NO: 67	3736
ANTH169	Lambda light chain		US20040258699 SEQ ID NO: 68	3737
AIMTH 170	Lambda light chain		US20040258699 SEQ ID NO: 69	3738
ANTH171	Lambda light chain		US20040258699 SEQ ID NO: 70	3739
ANTH172	Lambda light chain		US20040258699 SEQ ID NO: 71	3740
ANTH173	Lambda light chain		US20040258699 SEQ ID NO: 72	3741
ANTH174	Lambda light chain		US20040258699 SEQ ID NO: 73	3742
AIMTH 175	Lambda light chain		US20040258699 SEQ ID NO: 74	3743
ANTH176	Lambda light chain		US20040258699 SEQ ID NO: 75	3744
ANTH177	Lambda light chain		US20040258699 SEQ ID NO: 76	3745
ANTH178	Lambda light chain		US20040258699 SEQ ID NO: 77	3746
ANTH179	Light chain		US8617548 SEQ ID NO: 1	3747
AIMTH 180	Light chain	IQNPA Lkappa	US7658925 SEQ ID NO: 4	3748
ANTH181	Light chairs	IQNLF Lkappa	US7658925 SEQ ID NO: 8	3749

ANTH182	Light chain	1A5	US20090022736 SEQ ID NO: 2	3750
ANTH183	Light chain	4A12	US20090022736 SEQ ID NO: 4	3751
ANTH184	Light chain	24B1	US20090022736 SEQ ID NO: 6	3752
ANTH185	Light chain	24G4	US20090022736 SEQ ID NO: 8	3753
ANTH186	Light chain	32E12	US20090022736 SEQ ID NO: 10	3754
ANTH187	Light chain	33F4	US20090022736 SEQ ID NO: 12	3755
ANTH188	Light chain	scFv 2LF	EP2778173 SEQ ID NO: 6	3756
ANTH189	Light chain	Obiltoxaximab		3757
ANTH190	Light chain region	W1	US8685396 SEQ ID NO: 9	3758
ANTH191	Light chain region	W2	US8685396 SEQ ID NO: 25	3759
ANTH192	Light chain region	W5	US8685396 SEQ ID NO: 37	3760
ANTH193	Light chain region	A63-6	US8685396 SEQ ID NO: 38	3761
ANTH194	Light chain region	F3-6	US8685396 SEQ ID NO: 39	3762
ANTH195	Light chain region	F5-1	US8685396 SEQ ID NO: 40	3763
ANTH196	Light chain variable region	LF!1H	US8961975 SEQ ID NO: 25	3764
ANTH197	Light chain variable region	LF9D	US8961975 SEQ ID NO: 17	3765
ANTH198	Light chain variable region	LF10E	US8961975 SEQ ID NO: 9	3766
ANTH199	Light chain variable region	6.20	WO2015107307 SEQ ID NO: 2	3767
ANTH200	Light chain variable region	35PA83	WO2009071860 SEQ ID NO: 2	3768
ANTH201	Light chain variable region	anti-yDPGA antibody	US8501182 SEQ ID NO: 2	3769
ANTH202	Light chain variable region	4C	US8501182 SEQ ID NO: 4	3770
ANTH203	Light chain variable region	11D	US8501182 SEQ ID NO: 6	3771
ANTH204	Light chain variable region	F20G75	WO2007131363 SEQ ID NO: 10	3772
ANTH205	Light chain variable region	F20G76	WO2007131363 SEQ ID NO: 12	3773
ANTH206	Light chain variable region	F20G77	WO2007131363 SEQ ID NO: 14	3774
ANTH207	Light chain variable region	V2 variant	US8507655 SEQ ID NO: 8	3775
ANTH208	Light chain variable region	6.20 variant	US8507655 SEQ ID NO: 10	3776
ANTH209	Light chain variable region	J24.15 variant	US8507655 SEQ ID NO: 12	3777
ANTH210	Light chain variable region	J24.7 variant	US8507655 SEQ ID NO: 14	3778
ANTH211	Light chain variable region	V2 variant human	US8507655 SEQ ID NO: 16	3779
ANTH212	Light chain variable region	6.20 variant human	US8507655 SEQ ID NO: 18	3780
ANTH213	Light chain variable region	J24.15 variant human	US8507655 SEQ ID NO: 20	3781
ANTH214	Light chain variable region	J24.7 variant human	US8507655 SEQ ID NO: 22	3782
ANTH215	Light chain variable region	HuMab 5E8 (Major)	US8404820 SEQ ID NO: 4	3783
ANTH216	Light chain variable region	HuMab 5E8 (Minor)	US8404820 SEQ ID NO: 6	3784
ANTH217	Light chain variable region	HuMab 2D5	US8404820 SEQ ID NO: 10	3785
ANTH218	Light chain variable region	HisMab 2H4	US8404820 SEQ ID NO: 14	3786
ANTH219	Light chain variable region	HuMab 5D5-2E10	US8404820 SEQ ID NO: 18	3787
ANTH220	Light chain variable region	13E3	US8309090 SEQ ID NO: 4	3788
ANTH221	Light chain variable region	3E1	US8309090 SEQ ID NO: 8	3789
ANTH222	Light chain variable region	KCTC 10756BP	US8268316 SEQ ID NO: 7	3790

ANTH223	Light chain variable region	modified M18 sequence	US7902344; US6916474 SEQ ID NO: 25	3791
ANTH224	Light chain variable region	21D9 MAb	US7442373 SEQ ID NO: 4	3792
ANTH225	Light chain variable region	1C6 Mab	US7442373 SEQ ID NO: 8	3793
ANTH226	Light chain variable region	4H7 Mab	US7442373 SEQ ID NO: 12	3794
ANTH227	Light chain variable region	22G12 Mab	US7442373 SEQ ID NO: 16	3795
ANTH228	Light chain variable region, antibody against anthrax toxin.	ETI-204	US20120156196 SEQ ID NO: 2	3796
ANTH229	Light chain variable region. Edema factor	EF12A	US8961975 SEQ ID NO: 54	3797
ANTH230	Light chain variable region, Edema factor	EF13D	US8961975 SEQ ID NO: 41	3798
ANTH231	Light chain variable region, Edema factor	EF14H	US8961975 SEQ ID NO: 55	3799
ANTH232	Light chain variable region, Edema factor	EF15A	US8961975 SEQ ID NO: 56	3800
ANTH233	Scfv	PWB2447 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 48	3801
ANTH234	Scfv	PWC2004 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 49	3802
ANTH235	Scfv	PWD0283 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 50	3803
ANTH236	Scfv	PWD0323 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 51	3804
ANTH237	Scfv	PWD0422 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 52	3805
ANTH238	Scfv	PWD0587 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 53	3806
ANTH239	Scfv	PWD0791 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 54	3807
ANTH240	Scfv	PHD2222 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 55	3808
ANTH241	Scfv	PHD2581 scFv	US7601351; US7906119; US20110189197 SEQ ID NO: 56	3809
ANTH242		Abthrax	US20120156196 SEQ ID NO: 48	3810
ANTH243		Abthrax	US20120156196 SEQ ID NO: 49	3811
ANTH244			WO2003063768 SEQ ID NO: 4	3812
ANTH245			WO2003063768 SEQ ID NO: 5	3813

[00305] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 16 against Botulinum Toxin.

Table **16. Antibodies** against **Botulinum** Toxin

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
BOTT1	Heavy-chain-only		US20130058962 SEQ ID NO: 56	3814
BOTT2	Heavy-chain-only		US20130058962 SEQ ID NO: 57	3815
BOTT3	Heavy-chain-only		US20130058962 SEQ ID NO: 58	3816
BOTT4	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDA-D12	WO2015100409 SEQ ID NO: 20	3817

BOTT5	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDQ-A5	WO2015100409 SEQ ID NO: 22	3818
BOTT6	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDQ-B5	WO2015100409 SEQ ID NO: 24	3819
BOTT7	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDQ-C2	WO2015100409 SEQ ID NO: 26	3820
BOTT8	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDQ-F 12	WO2015100409 SEQ ID NO: 28	3821
BOTT9	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDQ-G5	WO2015100409 SEQ ID NO: 30	3822
BOTT10	Heavy-chain only binding agents specific to BoNT/A holotoxin	JDQ-H7	WO2015100409 SEQ ID NO: 32	3823
BOTT11	Heavy-chain only binding agents specific to BoNT/A holotoxin	JEQ-A5	WO2015100409 SEQ ID NO: 34	3824
BOTT12	Heavy-chain only binding agents specific to BoNT/A holotoxin	JEQ-H11	WO2015100409 SEQ ID NO: 36	3825
BOTT13	Heavy-chain only binding agent	E-9	WO2015100409 SEQ ID NO: 38	3826
BOTT14	Heavy-chain only binding agent	B2	WO2015100409 SEQ ID NO: 40	3827
BOTT15	Heavy-chain only binding agent	C5	WO2015100409 SEQ ID NO: 42	3828
BOTT16	Heavy-chain only binding agent	F9	WO2015100409 SEQ ID NO: 44	3829
BOTT17	Heavy-chain only binding agent	heavy-chain only binding agent	WO2015100409 SEQ ID NO: 46	3830
BOTT18	Heavy-chain only binding agent with tag	heavy-chain only binding agent with tag	WO2015100409 SEQ ID NO: 48	3831
BOTT19	Heavy-chain only binding agent with tag	heavy-chain only binding agent with tag	WO2015100409 SEQ ID NO: 50	3832
BOTT20	Heavy-chain only dimer binding agent with two tags	heavy-chain only dimer binding agent with two tags	WO2015100409 SEQ ID NO: 52	3833
BOTT21	Recombinant camelid heavy-chain-only antibody	H7	WO2015100409 SEQ ID NO: 56	3834
BOTT22	Recombinant camelid heavy-chain-only antibody	B5	WO2015100409 SEQ ID NO: 57	3835
BOTT23	Recombinant camelid heavy-chain-only antibody		WO2015100409 SEQ ID NO: 58	3836
BOTT24	Scfv	scFv#2	WO2015100409 SEQ ID NO: 2	3837
BOTT25	Scfv	scFv#3	WO2015100409 SEQ ID NO: 4	3838
BOTT26	Scfv	scFv#7	WO2015100409 SEQ ID NO: 6	3839
BOTT27	Scfv	scFv#8	WO2015100409 SEQ ID NO: 8	3840
BOTT28	Scfv	scFv#21	WO2015100409 SEQ ID NO: 10	3841
BOTT29	Scfv	scFv#E	WO2015100409 SEQ ID NO: 12	3842
BOTT30	Scfv	scFv#7-2E	WO2015100409 SEQ ID NO: 14	3843

[00306] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 17 against Shiga Toxin.

Table 17. **Antibodies** against Shiga **Toxin**

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
SHIG1	Camelid heavy-chain only	JET-H12	WO2015100409 SEQ ID NO: 96	3844
SHIG2	Camelid heavy-chain only	JFG-H6	WO2015100409 SEQ ID NO: 98	3845
SHIG3	Heavy chain		US2014013548 SEQ ID NO: 44	3846
SHIG4	Heavy chain		US2014013548 SEQ ID NO: 21	3847
SHIG5	Heavy chain of cαstx1	Shigamab	US20120195891 SEQ ID NO: 1	3848
SHIG6	Heavy chain of cαstx1	Shigamab	US20120195891 SEQ ID NO: 2	3849
SHIG7	Heavy chain of cαstx2	Shigamab	US20120195891 SEQ ID NO: 3	3850
SHIG8	Heavy chain of cαstx2	Shigamab	US20120195891 SEQ ID NO: 4	3851
SHIG9	Heavy chain single domain		WO2014191904 SEQ ID NO: 7	3852
SHIG10	Heavy chain single domain		WO2014191904 SEQ ID NO: 8	3853
SHIG11	Heavy chain single domain		WO2014191904 SEQ ID NO: 9	3854
SHIG12	Heavy chain single domain		WO2014191904 SEQ ID NO: 10	3855
SHIG13	Heavy chain single domain		WO2014191904 SEQ ID NO: 11	3856
SHIG14	Heavy chain single domain		WO2014191904 SEQ ID NO: 12	3857
SHIG15	Heavy chain single domain		WO2014191904 SEQ ID NO: 13	3858
SHIG16	Heavy chain single domain		WO2014191904 SEQ ID NO: 14	3859
SHIG17	Heavy chain single domain		WO2014191904 SEQ ID NO: 15	3860
SHIG18	Heavy chain single domain		WO2014191904 SEQ ID NO: 16	3861
SHIG19	Heavy chain single domain		WO2014191904 SEQ ID NO: 17	3862
SHIG20	Heavy chain single domain		WO2014191904 SEQ ID NO: 18	3863
SHIG21	Heavy chain single domain		WO2014191904 SEQ ID NO: 19	3864
SHIG22	Heavy chain single domain		WO2014191904 SEQ ID NO: 20	3865
SHIG23	Heavy chain single domain		WO2014191904 SEQ ID NO: 21	3866
SHIG24	Heavy chain single domain		WO2014191904 SEQ ID NO: 22	3867
SHIG25	Heavy chain single domain		WO2014191904 SEQ ID NO: 23	3868
SHIG26	Heavy chain single domain		WO2014191904 SEQ ID NO: 24	3869
SHIG27	Heavy chain single domain		WO2014191904 SEQ ID NO: 25	3870
SHIG28	Heavy chain single domain		WO2014191904 SEQ ID NO: 26	3871
SHIG29	Heavy chain single domain		WO2014191904 SEQ ID NO: 27	3872
SHIG30	Heavy chain single domain		WO2014191904 SEQ ID NO: 28	3873
SHIG31	Heavy chain single domain		WO2014191904 SEQ ID NO: 29	3874
SHIG32	Heavy chain single domain		WO2014191904 SEQ ID NO: 30	3875
SHIG33	Heavy chain single domain		WO2014191904 SEQ ID NO: 31	3876
SHIG34	Heavy chain single domain		WO2014191904 SEQ ID NO: 32	3877
SHIG35	Heavy chain single domain		WO2014191904 SEQ ID NO: 33	3878
SHIG36	Heavy chain single domain		WO2014191904 SEQ ID NO: 34	3879
SHIG37	Heavy chain single domain		WO2014191904 SEQ ID NO: 35	3880

SHIG38	Heavy **chain** single domain		**WO20 14** 191904 SEQ ID NO: 36	3881
SHIG39	Heavy chain single domain		WO20 !4!9i904 SEQ ID NO: 37	3882
SHIG40	Heavy chain single domain		WO20 14 19 1904 SEQ ID NO: 38	3883
SHIG41	Heavy **chain** single domain		**WO20** 14 19 **1904** SEQ ID **NO**: 39	3884
SHIG42	Heavy chain single domain		**WO20 14 191904** SEQ ID NO: 40	3885
SHIG43	Heavy **chain** single domain		**WO20 14** 191904 SEQ ID NO: 41	3886
SHIG44	Heavy chain single domain		WO20 !4!9!**904** SEQ ID NO: 42	3887
SHEG45	Heavy chain single domain		**WO20 14 191904** SEQ **ID NO:** 43	3888
SHIG46	**Heavy** chain single domain		**WO20 14 191904** SEQ ID NO: 44	3889
SHIG47	Heavy chain single domain		**WO20 14 191904** SEQ ID NO: 45	3890
SHIG48	Heavy **chain** single domain		**WO20 14** 191904 SEQ ID NO: 46	3891
SHIG49	Heavy chain single domain		**WO20 14 I9!904** SEQ ID NO: 47	3892
SHIG50	Heavy-chain-only		US20130058962 SEQ ID NO: 77	3893
SHIG51	**Heavy-chain-only**		**US20 130058962** SEQ **ID** NO: 78	3894
SHIG52	**Heavy-chain-only**		**US20 130058962** SEQ ID NO: 79	3895
SHIG53	Heavy-chain-only		**US20 130058962** SEQ ID NO: 80	3896
SHIG54	Heavy-**chain-only**		US20130058962 SEQ ID **NO:** 81	3897
SHIG55	**Heavy-chain-only**		**US20130058962** SEQ ID NO: 82	3898
SHIG56	**Heavy-chain-only**		**US20 130058962** SEQ **ID** NO: 83	3899
SHIG57	**Heavy-chain-only**		**US20 130058962** SEQ ID NO: 84	3900
SHIG58	Heavy-chain-only		US20 130058962 SEQ ID NO: 85	3901
SHIG59	**Heavy-chain-only**		**US20 130058962** SEQ ID **NO:** 86	3902
SHTG60	Light chain		**US20140 13548** SEQ **ID** NO: 42	3903
SHIG61	Light chain		**US2014013548** SEQ **ID NO:** 19	3904
SHIG62	Recombinant **camelid heavy-chain-only** antibody, STX1	JET-A9	**WO2015 100409** SEQ ID NO: 77	3905
SHIG63	Recombinant camelid heavy-chain-only antibody, STX1	JGG-D4	**WO2015 100409** SEQ ID **NO:** 78	3906
SHIG64	Recombinant camelid heavy-**chain-only** antibody, STXI, STX2	JFD-A4	**WO2015 100409** SEQ ID NO: 84	3907
SHIG65	Recombinant camelid **heavy-cliain-only** antibody, STXI, STX2	JFD-A5	**WO20 15 100409** SEQ ID NO: 85	3908
SHIG66	Recombinant camelid heavy-chain-only antibody, STXI, STX2	JGG-G6	**WO20 !5!**00409 SEQ ID NO: 86	3909
SHIG67	Recombinant camelid heavy-chain-only antibody, STX2	JEN-D !O	**WO20 15 100409** SEQ ID **NO:** 79	3910
SHIG68	Recombinant camelid **heavy-chain-only** antibody, STX2	**JGH-G1**	WO20L5 100409 SEQ ID **NO:** 80	**391 1**
SHIG69	Recombinant camelid heavy-chain-only antibody, STX2	**JEU-A6**	**WO2015 100409** SEQ ID **NO:** 81	3912
SHIG70	Recombinant camelid heavy-**chain-only** antibody, STX2	**JEU-D2**	**WO2015 100409** SEQ ID NO: 82	3913
SHIG71	Recombinant camelid heavy-chain-only antibody, STX2	JGH-G9	**WO2015 100409** SEQ ID **NO:** 83	3914

[00307] In one embodmieni, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or valiants thereof or encodes one or more polypeptides.

fragments or variants thereof described in US Pub. No. US20090280104, the contents of each of which are herein incorporated by reference in their entirety, against Shiga toxin.

Tropical Diseases

[00308] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the tropical disease related payload antibody polypeptides listed in Tables 18-20.

[00309] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 18 against Plasmodium Falciparum causing Malaria.

Table 18. Antibodies against Plasmodium Falciparum causing Malaria

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
MALA1	Heavy chain	immunoglobulin heavy chain variable region, partial	Wajanarogana, S. et al., Construction of a human functional single-chain variable fragment (scFv) antibody recognizing the malaria parasite Plasmodium falciparum, Biotechnol. Appl. Biochem. 44 (PT 1), 55-61 (2006), NCBI Accession # AAX76832.1 (129aa)	3915
MALA2	Heavy chain	anti-MSP1 MAD20 block2 ScFv Ig heavy chain variable region, partial	Sowa, K.M. et al., Isolation of a monoclonal antibody from a malaria patient-derived phage display library recognizing the Block 2 region of Plasmodium falciparum merozoite surface protein-1, Mol. Biochem. Parasitol. 112 (1), 143-147 (2001), NCBI Accession #AAK08696.1 (119aa)	3916
MALA3	Heavy chain	immunoglobulin heavy chain variable region, partial	Lundquist, R. et al., Human recombinant antibodies against Plasmodium falciparum merozoite surface protein 3 cloned from peripheral blood leukocytes of individuals with immunity to malaria demonstrate antiparasitic properties, Infect. Immun. 74 (6), 3222-3231, (2006), NCBI Accession # AAT09786.1 (113aa)	3917
MALA4	Heavy chain variable region	2A10 anti-malaria antibody	NCBI Accession # BAK41504.1 (118aa)	3918
MALA5	Heavy chain		US7811569 to Dziegiel; SEQ ID NO: 1	3919
MALA6	Heavy chain, Anti-ang-2 antibody		US7811569 to Dziegiel; SEQ ID NO: 3	3920
MALA7	Heavy chain		US7811569 to Dziegiel; SEQ ID NO: 5	3921
MALA8	Heavy chain variable region		US20150197562 SEQ ID NO: 14	3922
MALA9	Heavy chain variable region	mAb 5D5	US20150158941 SEQ ID NO: 16	3923
MALA10	Heavy chain variable region		US20140112930 SEQ ID NO: 18	3924
MALA11	Heavy chain variable region	M071Xi0199	WO2014087007; SEQ ID NO: 182	3925
MALA12	Heavy chain variable region	M071Xi2204	WO2014087007; SEQ ID NO: 186	3926

MALA13	Heavy chain variable region	M071Xi0237	WO2014087007; SEQ ID NO: 190	3927
MALA14	Heavy chain variable region	M071Xi2127	WO2014087007; SEQ ID NO: 194	3928
MALA15	Heavy chain variable region	M071Xi0092	WO2014087007; SEQ ID NO: 198	3929
MALA16	Heavy chain variable region	M071Xi2057	WO2014087007; SEQ ID NO: 202	3930
MALA17	Heavy chain variable region	M070Xi3010	WO2014087007; SEQ ID NO: 206	3931
MALA18	Heavy chain variable region	M071Xi0227	WO2014087007; SEQ ID NO: 210	3932
MALA19	Heavy chain variable region	M071Xi0081	WO2014087007; SEQ ID NO: 214	3933
MALA20	Heavy chain variable region	M071Xi0124	WO2014087007; SEQ ID NO: 218	3934
MALA21	Heavy chain variable region	M036Xi0326	WO2014087007; SEQ ID NO: 222	3935
MALA22	Heavy chain variable region	M070Xi3195	WO2014087007; SEQ ID NO: 226	3936
MALA23	Heavy chain variable region	M070Xi3062	WO2014087007; SEQ ID NO: 230	3937
MALA24	Heavy chain variable region	M071Xi2217	WO2014087007; SEQ ID NO: 234	3938
MALA25	Heavy chain variable region	M036Xi0003	WO2014087007; SEQ ID NO: 238	3939
MALA26	Heavy chain, Eba-175	R217	Chen et al., PLoS Pathol. 9 (5), E1003390 (2013), NCBI Accession # 4QEX_I (215aa)	3940
MALA27	Heavy chain, Eba-175	R218	Chen et al., PLoS Pathol. 9 (5), E1003390 (2013), NCBI Accession # 4K2U_I (233aa)	3941
MALA28	Light chain	anti-MSP1 MAD20 block2 ScFv Ig heavy chain variable region, partial	Sowa, K.M. et al., Isolation of a monoclonal antibody from a malaria patient-derived phage display library recognizing the Block 2 region of Plasmodium falciparum merozoite surface protein-1, Mol. Biochem. Parasitol. 112 (1), 143-147 (2001), NCBI Accession #AAK08697.1 (119aa)	3942
MALA29	Light chain	anti-MSP1 MAD20 block2 ScFv Ig light chain variable region, partial	Sowa, K.M. et al., Isolation of a monoclonal antibody from a malaria patient-derived phage display library recognizing the Block 2 region of Plasmodium falciparum merozoite surface protein-1, Mol. Biochem. Parasitol. 112 (1), 143-147 (2001), NCBI Accession #AAK08698.1 (110aa)	3943
MALA30	Light chain	immunoglobulin light chain variable region, partial	Wajanarogana, S. et al., Construction of a human functional single-chain variable fragment (scFv) antibody recognizing the malaria parasite Plasmodium falciparum, Biotechnol. Appl. Biochem. 44 (PT 1), 55-61 (2006) AAX76833.1 (107aa)	3944
MALA31	Kappa light chain	immunoglobulin kappa light chain variable region, partial	Lundquist, R. et al., Human recombinant antibodies against Plasmodium falciparum merozoite surface protein 3 cloned from peripheral blood leukocytes of individuals with immunity to malaria demonstrate antiparasitic properties, Infect. Immun. 74 (6), 3222-3231, (2006), NCBI Accession # AAT09787.1 (113aa)	3945

MALA32	Light chain variable region	2A10 anti-malaria antibody	NCBI Accession # BAK4 1503. 1 (108aa)	3946
MALA33	Light chain		US781 1569 to Dziegiel; SEQ ID NO: 2	3947
MALA34	Light chain, Anti-ang-2 antibody		US781 1569 to Dziegiel; SEQ ID NO: 4	3948
MALA35	Light chain		US781 1569 to Dziegiel; SEQ ID NO: 6	3949
MALA36	Light cliain variable regiots		US20150 197562 SEQ ID NO: 15	3950
MALA37	Light chain variable region		US20 150 197562 SEQ ID NO: 19	395 I
MALA38	Light chain variable region	mAb 5D5	US20150 158941 SEQ ID NO: 14	3952
MALA39	Light chain variable region		US20 1401 12930 SEQ ID NO: 20	3953
MALA40	Light chain variable region	M071Xi0199	WO2014087007; SEQ ID NO: 184	3954
MALA41	Light chain variable region	M071Xi2204	WO20 14087007; SEQ ID NO: 188	.3955
MALA42	Light chain variable region	M071Xi0237	WO2014087007; SEQ ID NO: 192	3956
MALA43	Light chain variable region	M071Xi2127	WO20 14087007; SEQ ID NO: 196	3957
MALA44	Light chain variable region	M071Xi0092	WO2014087007; SEQ ID NO: 200	3958
MALA45	Light chain variable region	M071Xi2057	WO20 14087007; SEQ ID NO: 204	.3959
MALA46	Light chain variable region	M070Xi30 10	WO2014087007; SEQ ID NO: 208	3960
MALA47	Light chain variable region	M071Xi0227	WO20 14087007; SEQ ID NO: 212	.3961
MALA48	Light chain variable region	M071Xi0081	WO2014087007; SEQ ID NO: 216	3962
MALA49	Light chain variable region	M071Xi0124	WO2014087007; SEQ ID NO: 220	3963
MALA50	Light chain variable region	M036Xi0326	WO2014087007; SEQ ID NO: 224	3964
MALA51	Light chain variable region	M070Xi3195	WO2014087007; SEQ ID NO: 228	3965
MALA52	Light chain variable region	M070Xi3062	WO2014087007; SEQ ID NO: 232	3966
MALA53	Light chain variable region	M071Xi2217	WO2014087007; SEQ ID NO: 236	3967
MALA54	Light chain variable region	M036Xi0003	WO2014087007; SEQ ID NO: 240	3968
MALA55	Light chain, Eba-175	R217	Chen et al., PLoS Pathol. 9 (5), E1003390 (2013), NCBI Accession # 4QEX_M (214aa)	3969
MALA56	Light chain, Eba-175	R2 18	Chen et al., PLoS Pathol. 9 (5), E1003390 (20 13), NCBI Accession # 4K2U_M (234aa)	3970
MALA57	Vivax apical membrane antigen 1 monoclonal antibody, seqres	F8. 12. 19	NCBI Accession # 2.I4W_L (2 13aa)	3971

[00310] In one embodiment the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 19 against Ebola and/or Marghurg Viruses.

Table 19. Antibodies against Ebola and Marburg viruses

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
EBOL1	Chain A, Sudan Ebolavirus Glycoprotein (Strain Boniface)	16f6	Bale et al., Structural basis for differential neutralization of ebolaviruses; Viruses 4 (4), 447-470 (2012), NCBI Accession # 3VE0_B (212aa)	3972
EBOL2	Chain B, Sudan Ebolavirus Glycoprotein (Strain Boniface)	16f6	Bale et al., Structural basis for differential neutralization of ebolaviruses; Viruses 4 (4), 447-470 (2012), NCBI Accession # 3VE0_A (220aa)	3973
EBOL3	Ebola Virus Glycoprotein	13f6-1-2 Fab	Lee J. E. et al., Complex of a protective antibody with its Ebola virus GP peptide epitope: unusual features of a V lambda x light chain; J. Mol. Biol. 375 (1), 202-216 (2008), NCBI Accession # 2QHR_L (218aa)	3974
EBOL4	Ebola Virus Glycoprotein	13f6-1-2 Fab	Lee J. E. et al., Complex of a protective antibody with its Ebola virus GP peptide epitope: unusual features of a V lambda x light chain; J. Mol. Biol. 375 (1), 202-216 (2008), NCBI Accession # 2QHR_H (222aa)	3975
EBOL5	Fab heavy chain Envelope Glycoprotein Gp1	Mr78	Hashiguchi, T., et al., Cell 160 (5), 904-912 (2015), NCBI Accession # 3X2D_P (226aa)	3976
EBOL6	Fab light chain, Envelope Glycoprotein Gp1	Mr78	Hashiguchi, T., et al., Cell 160 (5), 904-912 (2015), NCBI Accession # 3X2D_O (213aa)	3977
EBOL7	Fusion protein, Zaire Ebola virus, Mayinga strain glycoprotein		US20140356354 SEQ ID NO: 2	3978
EBOL8	Heavy chain Ebolavirus-Protective Antibody		Olal, D., et al., Structure of an Antibody in Complex with Its Mucin Domain Linear Epitope That Is Protective against Ebola Virus; J. Virol. 86 (5), 2809-2816 (2012), NCBI Accession # 2Y6S_H (213aa)	3979
EBOL9	Heavy chain Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 6	3980
EBOL10	Heavy chain Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 7	3981
EBOL11	Heavy chain Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 8	3982
EBOL12	Heavy chain Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 9	3983
EBOL13	Heavy chain Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 10	3984
EBOL14	Heavy chain Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 11	3985
EBOL15	Heavy chain variable region, Zaire ebolavirus (ZEBOV) glycoprotein		WO2015127136 SEQ ID NO: 71	3986
EBOL16	Heavy chain variable region, Zaire ebolavirus (ZEBOV) glycoprotein		WO2015127136 SEQ ID NO: 47	3987
EBOL17	Heavy chain variable region, Zaire ebolavirus (ZEBOV) glycoprotein		WO2015127136 SEQ ID NO: 23	3988

EBOL18	**Heavy chain variable** region, Ebola Sudan Boniface virus **(ESB) glycoprotein (CP)**	**16H1 1**	**US9097713** SEQ ID NO: 2	3989
EBOL19	Heavy chain variable region, Ebola Sudan Boniface virus (ESB) glycoprotein **(GP)**	**19B3**	US9097713 SEQ ID NO: 4	3990
EBOL20	Heavy **chain variable** region, Ebola Sudan Boniface virus (ESB) glycoprotein **(GP)**	17F6	US9097713 SEQ ID NO: 6	3991
EBOL2 1	Heavy chain variable region, Ebola Sudan Boniface **vims** (ESB) glycoprotein **(GP)**	16F6	**US9097713** SEQ ID NO: 8	3992
EBOL22	Heavy **chain variable** region, Ebola virus **GP**	**EGP 6D8 1-2**	US7335356 SEQ ID NO: 22	3993
EBOL23	Heavy chain variable region, Ebola vims **GP**	**EGP13F6-1-2**	US7335356 SEQ ID **NO:** 32	3994
EBOL24	Heavy **chain variable** region, Ebola virus **GP**	**EGP I3C6-1-1**	US7335356 SEQ ID NO: 12	3995
EBOL25	Heavy chain variable region, **Marburg** virus, Ebola virus, Sudan virus, **Bundibugyo** virus, Tai Forest viras or **Reston** virus glycoprotein		**WO20** I5127i40 SEQ ID **NO:** I4	3996
EBOL26	Heavy **chain variable** region, Marburg virus, Ebola vims, Sudan vims, Bundibugyo vims, Tai Forest viras or Reston virus glycoprotein		**WO2015 127140** SEQ ID NO: 38	3997
EBOL27	Heavy **chain variable** region, Marburg virus, Ebola virus, Sudan virus, Bundibugyo virus, Tai Forest virus or Reston vims glycoprotein		**WO2015 127 140** SEQ ID NO: 62	3998
EBOL28	Heavy **chain variable** region, Marburg viras, Ebola viras, Sudan vims, Bundibugyo virus, Tai Forest vims or Reston vims glycoprotein		**WO2015 127140** SEQ ID NO: 86	3999
EBOL29	Heaw **chain variable** region, Marburg vims, Ebola vims, Sudan virus, **Bundibugyo** virus, Tai Forest virus or Reston vims glycoprotein		**WO2015 127140** SEQ ID NO: 110	4000
EBOL30	Heavy chain variable region, Marburg virus, Ebola virus, Sudan viras, Bundibugyo viras, Tai Forest viras or Reston **virus** glycoprotein		WO2015I27i40 SEQ ID NO: **134**	4001
EBOL3 i	Heavy **chain variable** region, Marburg vims, Ebola vims, Sudan viras, Bundibugyo viras, Tai Forest vims or Reston viras glycoprotein		**WO2015 127140** SEQ ID NO: 158	4002
EBOL32	Heavy chain, Ebola virus glycoprotein.	Fab Kz52	Lee J. **E.** et **al.,** Structure of the Ebola virus glycoprotein bound to an antibody from a **human survivor;** Nature 454 (7201), 177-182 (2008), NCBI Accession *ft* 3CSY_**G (226aa)**	4003

EBOL33	Light chain variable region, Ebola Sudan Boniface virus **(ESB)** glycoprotein (CP)	16F6	**US9097713** SEQ ID NO: **10**	4004
EBOL34	Light chain variable **region,** Ebola **virus** GP	EGP **6D8 1-2**	**US733 5356** SEQ **ID NO:** 27	4005
EBOL35	Light chain variable region, Ebola virus GP	**EGP1 3F6-** 1-2	US7335356 SEQ ID NO: 37	4006
EBOL36	Light chain variable region, Ebola **virus** GP	**EGP13C6** -1-1	**US733 5356** SEQ ID **NO:** 16	4007
EBOL37	Light chain variable region, Marburg **virus,** Ebola **virus,** Sudan **virus, Bundibugyo** virus, Tai Forest virus **or Reston virus** glycoprotein		**WO20 15 127140** SEQ **ID** NO: **2**	4008
EBOL38	Light chairs variable region, Marburg virus, Ebola virus, Sudan virus, **Bundibugyo** virus, Tai Forest virus or Reston **virus** glycoprotein		**WO20 15 127 140** SEQ **ID NO:** 26	4009
EBOL39	Light chain variable region, Marburg **virus,** Ebola **virus,** Sudan virus, Bundibugyo virus, Tai Forest **virus** or Reston virus glycoprotein		**WO2015 127140** SEQ ID NO: 50	4010
EBOL40	Light chain variable region, Marburg virus, Ebola virus, Sudan virus, Bundibugyo virus, Tai Forest virus or Reston virus glycoprotein		**WO2015 127 140** SEQ ID NO: 74	4011
EBOL41	Light **chain variable** region, Marburg virus, Ebola virus, Sudan **virus, Bundibugyo** virus, Tai Forest **vims** or Reston vims glycoprotein		**WO2015 127140** SEQ ID NO: 98	4012
EBOL42	Light chain variable region, Marburg vims, Ebola vims, Sudan vims, Bundibugyo virus, Tai Forest virus or Reston vims **gly coprotein**		**WO2015 127140** SEQ ID NO: 122	4013
EBOL43	Light chain variable region, Marburg virus, Ebola virus, Sudan vims, Bundibugyo vims, Tai Forest virus or Reston virus glycoprotein		WO2015 127140 SEQ ID NO: 146	4014
EBOL44	Light chain variable region, Zaire **ebolavirus (ZEBOV)** glycoprotein		**WO2015 127136** SEQ ID NO: 59	4015
EBOL45	Light chairs variable region, Zaire ebolavirus (ZEBOV) glycoprotein		**WO20 15 127 I36** SEQ **ID NO:** 35	4016
EBOL46	Light chain variable region, Zaire ebolavirus (ZEBOV) **glycoprotein**		**WO2015 127136** SEQ ID NO: 11	4017
EBOL47	light chain, Ebola virus glycoprotein	Fab **Kz52**	Lee J. **E.** et **al.,** Structure of the Ebola virus glycoprotein bound to an antibody from a human **survivor;** Nature 454 (7201), 177-182 (2008), NCBE Accession *it* 3CSY **H (217aa)**	4018
EBOL48	Light chain, **Ebolavirus-Protective** Antibody		**Olal,** D., et **al., Structure of** an Antibody in Complex with **Its** Mucin **Domain** Linear Epitope That Is Protective against Ebola Vims; J. Virol.	4019

			86 (5), 2809-2816 (2012), NCBI Accession # 2Y6S_L (217aa)	
EBOL49	Light chain, Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 12	4020
EBOL50	Light chains, Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 13	4021
EBOL51	Light chain, Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 14	4022
EBOL52	Light chains, Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 15	4023
EBOL53	Light chain, Filovirus (Ebola or Marburg)		US20140356354 SEQ ID NO: 16	4024

[00311] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Patent No. US7335356 and EP Pub. No. EP1539238, the contents of each of which are herein incorporated by reference in their entirety, against Ebola.

[00312] In one embodiment the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 20 against Mosquito-borne disease.

Table 20, Antibodies against **Mosquito-borne** diseases

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
MOSQ1	Gamma heavy chain, partial, anti-Saint Louis encephalitis virus envelope glycoprotein immunoglobulin		Thibodeaux, B.A. "Development of a human-murine chimeric immunoglobulin M antibody for use in the serological detection of human flavivirus antibodies", Clin. Vaccine Immunol. 16 (5), 679-685,2009), NCBI Accession # ACI62179	4025
MOSQ2	Gamma heavy chain, partial, anti-Saint Louis encephalitis virus envelope glycoprotein immunoglobulin		Thibodeaux, B.A. "Development of a human-murine chimeric immunoglobulin M antibody for use in the serological detection of human flavivirus antibodies", Clin. Vaccine Immunol. 16 (5), 679-685,2009), NCBI Accession # ACI62180	4026
MOSQ3	Heavy chain variable region, Japanese encephalitis virus	anti-DLVR1/CLEC5A	US20080292644 SEQ ID NO: 69	4027
MOSQ4	Heavy chain variable region, Japanese encephalitis virus	anti-DLVR1/CLEC5A	US20080292644 SEQ ID NO: 70	4028
MOSQ5	Heavy chain variable region, Japanese encephalitis virus	anti-DLVR1/CLEC5A	US20080292644 SEQ ID NO: 71	4029
MOSQ6	Heavy chain variable region, Japanese encephalitis virus		CN103864925 SEQ ID NO: 2	4030
MOSQ7	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20480.1	4031

MOSQ8	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west rale virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20479. 1	4032
MOSQ9	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20478. 1	4033
MOSQ10	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20477. 1	4034
MOSQ1 1	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20476. 1	4035
MOSQ12	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20475. 1	4036
MOSQ 13	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20474. 1	4037
MOSQ14	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20473. 1	4038
MOSQ15	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20472. 1	4039
MOSQ16	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006). NCBI Accession # ABF2047 1.1	4040
MOSQ17	Heavy chain variable region, WNV, Dengue, St. Louis encephalitis, yellow fever virus, Japanese encephalitis virus, Murray Valley encephalitis virus	mAb 1	WO20 14144061 SEQ ID NO: 1	3359
MOSQ18	Heavy chain, WNV	CR4348	US89 1 1738 SEQ ID NO: 30	404 1
MOSQ 19	Heavy chain, WNV	CR4354	US891 1738 SEQ ID NO: 32	4042
MOSQ20	Heavy chain, WNV	CR426 1	US89 11738 SEQ ID NO: 60	4043
MOSQ2 1	Heavy chain, WNV	CR4267	US891 1738 SEQ ID NO: 62	4044
MOSQ22	Heavy chain, WNV	CR4328	US89H738 SEQ ID NO: 64	4045
MOSQ23	Heavy chain, WNV	CR4335	US89 11738 SEQ ID NO: 66	4046

MOSQ24	Heavy chain, WNV	CR4383	US891 1738 SEQ ID NO: 68	4047
MOSQ25	Heavy chain, WNV	CRM4354	US89 11738 SEQ ID NO: 148	4048
MOSQ26	Heavy chain variable region, WNV	Antibody frons US89 11738	US891 1738 SEQ iD NO: 20	4049
MOSQ27	Heavy chain variable region, WNV	E16 heavy chain version 1	US7572456 SEQ ID NO: 21	4050
MOSQ28	Heavy chain variable region, WNV	E16 heavy chain version 2	US7572456 SEQ iD NO: 22	4051
MOSQ29	Heavy chain variable region, WNV	E16 heavy chain version 3	US7572456 SEQ ID NO: 23	4052
MOSQ30	Heavy chain variable region, WNV	Antibody frons US89 11738	US891 1738 SEQ iD NO: 18	4053
MOSQ3 1	Heavy chain variable region, WNV	Sti-E16/E16p	US8663950 SEQ ID NO: 2	4054
MOSQ32	Heavy chain variable region, WNV	in -E16/Ei6p	US8663950 SEQ iD NO: 3	4055
MOSQ33	Heavy chain variable region, WNV	E16	US7527973 SEQ ID NO: 4	4056
MOSQ34	Heavy chain variable region, WNV	E24	US7527973 SEQ ID NO: 8	4057
MOSQ35	Heavy chain variable region, WNV	E34	US7527973 SEQ ID NO: 12	4058
MOSQ36	Heavy chain variable region, WNV	11	US20090130123 SEQ ID NO: 23	4059
MOSQ37	Heavy chain variable region, WNV	71	US20090130123 SEQ ID NO: 24	4060
MOSQ38	Heavy chain variable region, WNV	73	US20090130123 SEQ ID NO: 25	4061
MOSQ39	Heavy chain variable region, WNV	85	US20090130123 SEQ ID NO: 26	4062
MOSQ40	Heavy chain variable region, WNV	15	US20090130123 SEQ ID NO: 27	4063
MOSQ41	Heavy chain variable region, WNV	95	US20090 130123 SEQ ID NO: 28	4064
MOSQ42	Heavy chain variable region, WNV	84	US20090130123 SEQ ID NO: 29	4065
MOSQ43	Heavy chain variable region, WNV	10	US20090 130123 SEQ ID NO: 30	4066
MOSQ44	Heavy chain variable region, WNV	69	US20090130123 SEQ ID NO: 31	4067
MOSQ45	Heavy chain variable region, WNV	79	US20090 130123 SEQ ID NO: 32	4068
MOSQ46	Heavy chain variable region, WNV	94	US20090130123 SEQ ID NO: 33	4069
MOSQ47	Heavy chain variable region, WNV	9FI2	WO2010093335 SEQ ID NO: 4	3333
MOSQ48	Heavy chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20481.1	4070
MOSQ49	Heavy chain translation, WNV	hu-E16/E16p	US8663950 SEQ ID NO: 5	4071
MOSQ50	Heavy chain variable region, Yellow fever virus	anti-yellow fever virus vaccine strain 17D E glycoprotein	Thibodeaux, B.A. "A humanized IgG but not IgM antibody is effective in prophylaxis and therapy of yellow fever infection in an AG129/170-204 penpherai challenge mouse model" Antiviral Res. 94 (1), 1-8 (2012), NCBI Accession # AD017683	4072

MOSQ5 1	Light chain variable region, Japanese encephalitis virus	anti-DLVR1/CLECSA	US20080292644 SEQ ID NO: 66	4073
MOSQ52	Light chain variable region, Japanese encephalitis virus	***m*{*l*·** DLVR 1/CLEC5A	US20080292644 SEQ ID NO: 67	4074
MOSQ53	Light chain variable region, Japanese encephalitis virus	anii-DLVR1/CLECS A	US20080292644 SEQ ID NO: 68	4075
MOSQ54	Light chain variable region, Japanese encephalitis virus		CN103864925 SEQ ID NO: 1	4076
MOSQ55	Light chain variable region, WNV, Dengue, St. Louis encephalitis, yellow fever virus, Japanese encephalitis virus, Murray Valley encephalitis virus	mAb} 1	WO20{4{4406} SEQ ID NO: 3	3418
MOSQ56	Light chain, WNV	CR4348	US891 1738 SEQ ID NO: 34	4077
MOSQ57	Light chain, WNV	CR4354	US89H738 SEQ ID NO: 36	4078
MOSQ58	Light chain, WNV	CR4261	US89 11738 SEQ ID NO: 70	4079
MOSQ59	Light chain, WNV	CR4267	US891 1738 SEQ ID NO: 72	4080
MOSQ60	Light chain, WNV	CR4328	US89 11738 SEQ ID NO: 74	4081
MOSQ61	Light chain, WNV	CR4335	US891 1738 SEQ ID NO: 76	4082
MOSQ62	Light chain, WNV	CR4383	US89H738 SEQ ID NO: 78	4083
MOSQ63	Light chain variable region, WNV	Antibody from US89 11738	US89 11738 SEQ ID NO: 22	4084
MOSQ64	Light chain variable region, WNV	Antibody from US89 11738	US891{738 SEQ ID NO: 24	4085
MOSQ65	Light chain variable region, WNV	EI6	US7527973 SEQ ID NO: 2	4086
MOSQ66	Light chain variable region, WNV	E24	US7527973 SEQ ID NO: 6	4087
MOSQ67	Light chain variable region, WNV	E34	US7527973 SEQ ID NO: 10	4088
MOSQ68	Light chain variable region, WNV	E16 light chain version 1	US7572456 SEQ ID NO: 25	4089
MOSQ69	Light chain variable region, WNV	E16 light chain version 2	US7572456 SEQ ID NO: 26	4090
MOSQ70	Light chain variable region, WNV	1{	US20090 130123 SEQ ID NO: 34	4091
MOSQ7!	Light chain variable region, WNV	71	US20090130 {23 SEQ ID NO: 35	4092
MOSQ72	Light chain variable region, WNV	73	US20090 {30123 SEQ ID NO: 36	4093
MOSQ73	Light chain variable region, WNV	85	US20090130 {23 SEQ ID NO: 37	4094
MOSQ74	Light chain variable region, WNV	15	US20090 {30123 SEQ ID NO: 38	4095
MOSQ75	Light chain variable region, WNV	95	US20090130 {23 SEQ ID NO: 39	4096
MOSQ76	Light chain variable region, WNV	84	US20090 {30123 SEQ ID NO: 40	4097
MOSQ77	Light chain variable region, WNV	{0	US20090130123 SEQ ID NO: 41	4098

MOSQ78	Light chain variable region, WNV	69	US20090130123 SEQ ID NO: 42	4099
MOSQ79	Light cliain variable region, WNV	79	US20090130123 SEQ ID NO: 43	4100
MOSQ80	Light chain variable region, WNV	94	US20090130123 SEQ ID NO: 44	4101
MOSQ81	Light chain variable region, WNV	9FI2	WO2010093335 SEQ ID NO: 6	3393
MOSQ82	Light chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20470.1	4102
MOSQ83	Light cliain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20469.1	4103
MOSQ84	Light chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20468.1	4104
MOSQ85	Light chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006). NCBI Accession # ABF20467.1	4105
MOSQ86	Light chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20466.1	4106
MOSQ87	Light chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20465.1	4107
MOSQ88	Light chain variable region, partial sequence, WNV		Throsby, M." Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20464.1	4108
MOSQ89	Light chain variable region, partial sequence, WNV		Throsby, M. "Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20463.1	4109
MOSQ90	Light cliain variable region, partial sequence, WNV		Throsby, M. "Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20462.1	4110
MOSQ91	Light chain variable region, partial sequence, WNV		Throsby, M. "Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20461.1	4111

MOSQ92	Light chain variable region, partial sequence, WNV		Throsby, M. "Isolation and characterization of human monoclonal antibodies from individuals infected with west mie virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20460. 1	4112
MOSQ93	Light cliain variable region, partial sequence, WNV		Throsby, M. "Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20459. 1	4113
MOSQ94	Light chain variable region, partial sequence, WNV		Throsby, M. "Isolation and characterization of human monoclonal antibodies from individuals infected with west nile virus" J. Virol. 80 (14), 6982-6992 (2006), NCBI Accession # ABF20458. 1	4114
MOSQ95	Light chain translation, WNV	in -E16/Ei6p	US8663950 SEQ ID NO: 7	4115
MOSQ96	Light chain variable region, Yellow fever vims	anti-yellow fever virus vaccine strain 17D E glycoprotein	Thibodeaux, B.A. "A humanized IgG but not IgM antibody is effective in prophylaxis and therapy of yellow fever infection in an AG129/17D-204 peripheral challenge mouse model" Antiviral Res. 94 (1), 1-8 (2012), NCBI Accession # ADO17684	4116
MOSQ97	ScFv, WNV	9FI2	WO2010093335 SEQ ID NO: 8	3442
MOSQ98	Fc region, WNV, Dengue, St. Louis encephalitis, yellow fever virus, Japanese encephalitis virus, Murray Valley encephalitis virus	mAb-11	WO20 14144061 SEQ ID NO: 5	4117
MOSQ99	Fc region, WNV, Dengue, St. Louis encephalitis, yellow fever virus, Japanese encephalitis virus, Murray Valley-encephalitis virus	snAb-11-LALA	WO20 14 14406! SEQ ID NO: 6	4118
MOSQ100	ScFv, WNV	11	US20090130123 SEQ ID NO: 12	4119
MOSQ101	ScFv, WNV	71	US20090130123 SEQ ID NO: 13	4120
MOSQ102	ScFv, WNV	19	US20090 130123 SEQ ID NO: 14	4121
MOSQ103	ScFv, WNV	85	US20090130123 SEQ ID NO: 15	4122
MOSQ104	ScFv, WNV	15	US20090 130123 SEQ ID NO: 16	4123
MOSQ105	ScFv, WNV	95	US20090130123 SEQ ID NO: 17	4124
MOSQ106	ScFv, WNV	84	US20090130123 SEQ ID NO: 18	4125
MOSQ107	ScFv, WNV	10	US20090 130123 SEQ ID NO: 19	4126
MOSQ108	ScFv, WNV	69	US20090130123 SEQ ID NO: 20	4127
MOSQ 109	ScFv, WNV	79	US20090 130123 SEQ ID NO: 21	4128
MOSQ1 10	ScFv, WNV	94	US20090130123 SEQ ID NO: 22	4129
MOSQ111	ScFvs, WNV	SC04-348	US89 11738 SEQ ID NO: 26	4130
MOSQ1 12	ScFvs, WNV	SC04-354	US891 1738 SEQ ID NO: 28	4131
MOSQ1 13	ScFv, Yellow fever virus	anti-yellow fever virus E proiem scFv 7A	Daffis, S. et al. "Antibody responses against wild-type yellow fever virus and She 17D vaccine strain: charac terization with human monoclonal antibody fragments and	4132

			neutralization escape variants" Virology 337 (2), 262-272 (2005), NCBI Accession # AAT76799	
MOSQ1 14	ScFv, Yellow fever vims	anti-yellow fever virus E protein scFv R3(27)	Daffis, S. et al. "Antibody responses against wild-type yellow fever virus and the 17D vaccine strain: characterization with human monoclonal antibody fragments and neutralization escape variants" Virology 337 (2), 262-272 (2005), NCBI Accession # AAT76800	4133
MOSQ1 15	ScFv, Yellow fever virus	aim-yellow fever virus E protein scFv 5A	Daffis, S. et al. "Antibody responses against wild-type yellow fever virus and the 17D vaccine strain: characterization with human monoclonal antibody fragments and neutralization escape variants" Virology 337 (2), 262-272 (2005), NCBI Accession # AAT76801	4134
MOSQ1 16	ScFv, Yellow fever vims	anti-yellow fever virus E proiem scFv 1A	Daffis, S. et al. "Antibody responses against wild-type yellow fever virus and the 17D vaccine strain: characterization with human monoclonal antibody fragments and neutralization escape variants" Virology 337 (2), 262-272 (2005), NCBI Accession it AAT76802	4135
MOSQ1 17	ScFv, Yellow fever virus	anti-yellow fever virus E protein scFv 2A	Daffis, S. et al. "Antibody responses against wild-type yellow fever virus and the 17D vaccine strain: characterization with human monoclonal antibody fragments and neutralization escape variants" Virology 337 (2), 262-272 (2005), NCBI Accession # AAT76803	4136
MOSQ1 18	ScFv, Yellow fever virus	aim-yellow fever virus E protein scFv R3(9)	Daffis, S. et al. "Antibody responses against wild-type yellow fever virus and the 17D vaccine strain: characterization with human monoclonal antibody fragments and neutralization escape variants" Virology 337 (2), 262-272 (2005), NCBI Accession # AAT76804	4137

[00313] In one embodiment the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Patent No. US6399062 and US Pub. No. US20110171225, the contents of each of which are herein incorporated by reference in their entirety, against Malaria.

Infectious Diseases

[00314] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the infectious disease related payload antibody polypeptides listed in Tables 21-42.

[00315] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload **antibody** polypeptides listed **in** Table 21 against Influenza virus.

Table 21. **Antibodies** against **Influenza** virus

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
INFL1	Fab Fragment Heavy chain	ch65	Whittle, J.R. et al., Broadly neutralizing human antibody that recognizes the receptor-binding pocket of influenza virus hemagglutinin; Proc. Natl. Acad. Sci. U.S.A. 108 (34), 14216-14221 (2011), NCBI Accession #3SM5_H	4138
INFL2	Fab Heavy Chain	Fab Cr6261 (Somatic Heavy Chain With Germline-Reverted Light Chain)	Lingwood, D., et al., Structural and genetic basis for development of broadly neutralizing influenza antibodies; Nature 489 (7417), 566-570 (2012), NCBI Accession #4EVN_M (242aa)	4139
INFL3	Fab heavy chain	Del2d1	Krause, J.C. et al., M Bio 2 (1), E00345-E00310 (2011), NCBI Accession #3QHF_H	4140
INFL4	Fab heavy chain	Fld194 Fab	Xiong, X. et al., Structures of complexes formed by H5 influenza hemagglutinin with a potent broadly neutralizing human monoclonal antibody; Proc. Natl. Acad. Sci. U.S.A. 112 (30), 9430-9435 (2015), NCBI Accession #5A3I_C (230aa)	4141
INFL5	Fab heavy chain	H5.3	Winarski, K.L., Thornburg, N.J. et al., Vaccine-elicited antibody that neutralizes H5N1 influenza and variants binds the receptor site and polymorphic sites" PNAS 2015 112 (30) 9346-9351", NCBI Accession #4XNM_H	4142
INFL6	Fab Heavy chain	5j8	Hong, M. et al., Antibody Recognition of the Pandemic H1N1 Influenza Virus Hemagglutinin Receptor Binding Site; J. Virol. 87 (22), 12471-12480 (2013), NCBI Accession #4M5Z_H	4143
INFL7	Fab lambda heavy chain	CR6261	Ekiert, D.C. et al., Antibody recognition of a highly conserved influenza virus epitope; Science 324 (5924), 246-251 (2009), NCBI Accession #3GBN_H	4144
INFL8	Fab lambda light chain	CR6261	Ekiert, D.C. et al., Antibody recognition of a highly conserved influenza virus epitope; Science 324 (5924), 246-251 (2009), NCBI Accession #3GBN_L	4145
INFL9	Fab lambda light chain	Fab Cr6261 (Somatic Heavy Chain With Germline-Reverted Light Chain)	Lingwood, D., et al., Structural and genetic basis for development of broadly neutralizing influenza antibodies; Nature 489 (7417), 566-570 (2012), NCBI Accession #4EVN_N (217aa)	4146
INFL10	Fab light chain	Del2d1	Krause, J.C. et al., M Bio 2 (1), E00345-E00310 (2011), NCBI Accession #3QHF_L	4147
INFL11	Fab Light Chain	Fld194 Fab	Xiong, X. et al., Structures of complexes formed by H5 influenza hemagglutinin with a potent broadly neutralizing human monoclonal antibody; Proc. Natl. Acad. Sci. U.S.A. 112 (30), 9430-9435 (2015), NCBI Accession #5A3I_D (219aa)	4148

INFL 12	Fab, heavy chain	F045-092	Lee, P.S. et al., Receptor **mimicry by** antibody **F045-092** facilitates universal binding to **the** H.3 **subtype of influenza** virus; Nat **Commun** 5, 3614 (2014), NCBI Accession #40 5] W	4149
INFL 13	Fab, Light **Chain**	F045-092	Lee, P.S. et al., Receptor **mimicry** by antibody **F045-092** facilitates **universal** binding to the **H3** subtype of influenza virus; Nat Commun 5, 3614 (2014), NCBI Accession #4051 **V**	4150
INFL14	Fab, light chain	H5.3	**Winarski,** K.L., **Thornburg,** N.J. et **al.,** "Vaccine -elicited antibody **that** neutralizes **H5N** 1 influenza and variants binds the receptor site and polymorphic sites" PNAS 2015 112 (30) 9346-935 1", **NCBI Accession #4XNM** L	4151
INFL15	Gamma heavy chain variable	**8il0**	**US8858948** SEQ ID NO: 69	4152
INFL 16	Gamma heavy chain variable	23K12	**US8858948** SEQ ID NO: 100	4153
INFL 17	Heaw chain	CR6261, Diridavumab, **CR-**6261	WO 2008028946	4154
INFL 18	Heavy chairs	**Firivumab, CT-P22**	**US20 130004505**	4155
INFL19	Heavy chain	**CT-P22**	US201.30004505 SEQ ID **NO:** 4 1; **WO** 201 1/1 11966	4156
INFL20	Heavy chain	Navivumab, CT149	**WO20 13048153,** US201402.34.336 SEQ ID NO: 40	4157
1NFL2 1	Heavy chain	**ATI 0—004**	US20150010566, **WO201308 1463** SEQ ID NO: 31	4158
INFL22	Heavy chain	**AT10—003**	**US201500 10566**, **WO2013081463** SEQ ID NO: 32	4159
1NFL2.3	Heavy chain	**ATI 0—002**	US20150010566, **WO201308 1463** SEQ ID NO: 33	4160
INFL24	Heavy chain	**AT]O—00 1**	**US201500 10566, WO2013081463** SEQ ID NO: 34	4161
1NFL25	Heavy chain	**ATI 0—005**	US20150010566, **WO201308 1463** SEQ ID NO: 35	4162
INFL26	Heavy chain	**CT104**	WO20 **11]11**966, US20 **1**30004505 SEQ ID NO: **37**	4163
INFL27	Heavy chain	CT120	WO201 1111966, **US20130004505** SEQ ID NO: 41	4164
INFL28	Heavy chain	CT123	**WO201 11] 1966,** US20 130004505 SEQ ID **NO:** 45	4165
INFL29	Heavy chain	2A	**US2** 0 1400 1]982 SEQ ID **NO:** 2	4166
INFL30	Heavy **chain**	**F005-126**	WO2014049520, **US20140086927** SEQ **ID** NO: 2	4167
INFL3 1	Heaw chain	BFl- i	WO2008 156763 SEQ ID NO: 7	4168
INFL32	Heavy **chain**	**BF1-19**	WO2008 156763 SEQ ID NO: 11	4169
INFL33	Heavy chairs	**BF**1-IO	WO2008 156763 SEQ ID NO: 9	4170
1NFL34	Heavy chain		WO20 10 127252, 1JS8894997 SEQ ID NO: 3	**4171**
INFL35	Heavy chain	A18	**W013 170139** SEQ ID **NO:** 94	4172
INFL36	Heavy chain	Ab A18	**US7788200** SEQ ID NO: 15	4173
INFL37	Heavy **chain**	Ab 0 14, **Ab 028**	**US7788200** SEQ ID NO: 16	4174
INFL38	Heaw chain	Ab 071	**US7788200** SEQ ID NO: **162**	4175

INFL39	Heavy chain	Ab 072	US7788200 SEQ ID NO: 163	4176
INFL40	Heavy chain	Ab 078, Ab 079, Ab 080, Ab 081	US7788200 SEQ ID NO: 164	4177
INFL41	Heavy chain	Ab 001, Ab 009, Ab 017, Ab 160, Ab 186, Ab 187, Ab 188, Ab 189, Ab 190, Ab 191, Ab 192, Ab 193, Ab 202, Ab 211	US7788200 SEQ ID NO: 17	4178
INFL42	Heavy chain	Ab 002, Ab 010, Ab 026, Ab 203, Ab 212	US7788200 SEQ ID NO: 18	4179
INFL43	Heavy chain	Ab 003, Ab 011, Ab 027, Ab 194, Ab 195, Ab 196, Ab 197, Ab 198, Ab 199, Ab 200, Ab 204, Ab 213	US7788200 SEQ ID NO: 19	4180
INFL44	Heavy chain	Ab 086	US7788200 SEQ ID NO: 20	4181
INFL45	Heavy chain	Ab 154, Ab 155, Ab 157	US7788200 SEQ ID NO: 21	4182
INFL46	Heavy chain	Ab 157, Ab 159	US7788200 SEQ ID NO: 22	4183
INFL47	Heavy chain	Ab 210, Ab 219	US7788200 SEQ ID NO: 23	4184
INFL48	Heavy chain	Ab A001, Ab A002, Ab A003, Ab A010, Ab A011, Ab 031, Ab 037	US7788200 SEQ ID NO: 24	4185
INFL49	Heavy chain	Ab 004, Ab 005, Ab 006, Ab 012, Ab 013, Ab 032, Ab 038, Ab 043, Ab 044, Ab 045, Ab 046, Ab 047, Ab 048, Ab 049, Ab 050, Ab 051, Ab 052, Ab 067, Ab 068, Ab 069, Ab 070, Ab 073, Ab 074, Ab 075, Ab 076, Ab 077	US7788200 SEQ ID NO: 25	4186
INFL50	Heavy chain	Ab 007, Ab 008, Ab A009, Ab A14, Ab 015, Ab 033, Ab 039	US7788200 SEQ ID NO: 26	4187
INFL51	Heavy chain	Ab 016, Ab A017, Ab C18, Ab A019, Ab 034, Ab 040	US7788200 SEQ ID NO: 27	4188
INFL52	Heavy chain	F005-126	WO2014049520 SEQ ID 2	4189
INFL53	Heavy chain	8f24	WO2012045001 SEQ ID 1	4190
INFL54	Heavy chain	3E22	WO2012045001 SEQ ID 5	4191
INFL55	Heavy chain	5117	WO2012045001 SEQ ID 9	4192
INFL56	Heavy chain		WO2012045001 SEQ ID 13	4193
INFL57	Heavy chain		WO2012045001 SEQ ID 29	4194
INFL58	Heavy chain		WO2012045001 SEQ ID 33	4195

INFL59	**Heavy** chain		WO20 12045001 SEQ ID 17	4196
INFL60	Heavy chain	10A14	WO20 12045001 SEQ ID 21	4197
INFL61	Heavy chain	8D4	WO20 12045001 SEQ ID 25	4198
INFL62	Heavy chain	2B9	US91 15201 SEQ 3D NO: 6	4199
INFL63	Heavy chain	mAB 7A7	US20150239960, US20 140170163, US86733 14, US20 110027270, WO2010138564 SEQ ID NO: 6	4200
INFL64	Heavy **chain**	mAB 12D1	US20150239960, US20 140170 163, US86733 14, US20 110027270, WO2010138564 SEQ ID NO: 12	4201
1NFL65	Heavy chain	niAB 66A6	US20150239960, US20 140170163, US86733 14, US201 10027270, WO20 10138564 SEQ ID NO: 16	4202
INFL66	Heavy chain	M1D12	US201 10033473, WO2009 125395 SEQ ID NO: 17	4203
INFL67	Heavy chain	mABl. 12	WO2013030165 SEQ ID NO: 1	4204
INFL68	Heavy chain	mAB3. 1	WO20 13030 165 SEQ ID NO: 3	4205
INFL69	Heavy chain	5A7	WO2015 120097 SEQ ID NO: 7	4206
INFL70	**Heavy** chairs	TRL053	WO20 15 120097 SEQ ID NO: 17	4207
INFL71	Heavy chain	TRL579	WO20 15 120097 SEQ ID NO: 27	4208
INFL72	**Heavy chain**	TRL784	WO20 15 120097 SEQ ID NO: 37	4209
INFL73	**Heavy** chain	TRL794	WO20 15 120097 SEQ ID NO: 47	4210
INFL74	**Heavy chain**	TRL798	WO20 15 120097 SEQ ID NO: 57	4211
INFL75	Heavy chairs	TRL799	WO20 15 120097 SEQ ID NO: 67	4212
1NFL76	Heavy chain	TRL809	WO20 15 120097 SEQ ID NO: 77	4213
INFL77	**Heavy chain**	TRL811	WO20 15 120097 SEQ ID NO: 87	4214
INFL78	Heavy chain	TRL812	WO20 15 120097 SEQ ID NO: 97	4215
INFL79	**Heavy chain**	TRL8 13	WO2015 120097 SEQ ID NO: 107	4216
INFL80	Heavy chairs	TRL823	WO20 15 120097 SEQ ID NO: 117	4217
1NFL81	Heavy chain	TRL832	WO2015 120097 SEQ ID NO: 127	4218
INFL82	Heavy chain	TRL833	WO20 15 120097 SEQ ID NO: 137	4219
INFL83	Heavy chain	TRL834	WO20 15 120097 SEQ ID NO: 147	4220
INFL84	**Heavy chain**	TRL835	WO20 15 120097 SEQ ID NO: 157	4221
INFL85	Heavy chairs	TRL835	WO20 15 120097 SEQ ID NO: 158	4222
1NFL86	Heavy chain	TRL837	WO2015 120097 SEQ ID NO: 168	4223
INFL87	Heavy chain	TRL839	WO20 15 120097 SEQ ID NO: 178	4224
INFL88	Heavy chain	TRL841	WO2015 120097 SEQ ID NO: 188	4225
INFL89	Heavy **chain**	TRL842	WO20 15 120097 SEQ ID NO: 198	4226
INFL90	Heavy chairs	TRL845	WO20 15 120097 SEQ ID NO: 208	4227
INTL.91	Heavy chain	TRL846	WO2015 120097 SEQ ID NO: 217	4228
INFL92	Heavy chain	TRL847	WO20 15 120097 SEQ ID NO: 227	4229
INFL93	Heavy chain	TRL848	WO20 15 120097 SEQ ID NO: 237	4230
INFL94	Heavy **chain**	TRL849	WO20 15 120097 SEQ ID NO: 247	4231
INFL95	Heavy chain	TRL851	WO20 15 120097 SEQ ID NO: 257	4232
INTL.96	Heavy chain	TRL854	WO2015 120097 SEQ ID NO: 267	4233
INFL97	Heavy chain	TRL856	WO20 15 120097 SEQ ID NO: 277	4234
INFL98	Heavy chain	TRL858	WO20 15 120097 SEQ ID NO: 287	4235

INFL99	Heavy chain	humM2e-hBiTE-1	WO2014140368 SEQ ID NO: 8	4236
INFL100	Heavy chain	humM2e-hBiTE-2	WO2014140368 SEQ ID NO: 16	4237
INFL101	Heavy chain	humM2e-hBiTE-3	WO2014140368 SEQ ID NO: 24	4238
INFL102	Heavy chain	humM2e-hBiTE-4	WO2014140368 SEQ ID NO: 32	4239
INFL103	Heavy chain	VH of humM2e-hBiTE-5	WO2014140368 SEQ ID NO: 40	4240
INFL104	Heavy chain	humM2e-hBiTE-6	WO2014140368 SEQ ID NO: 48	4241
INFL105	Heavy chain	humM2e-hBiTE-7	WO2014140368 SEQ ID NO: 56	4242
INFL106	Heavy chain	humM2e-hBiTE-8	WO2014140368 SEQ ID NO: 64	4243
INFL107	Heavy chain	humM2e-hBiTE-9	WO2014140368 SEQ ID NO: 72	4244
INFL108	Heavy chain	murM2e-hBiTE	WO2014140368 SEQ ID NO: 80	4245
INFL109	Heavy chain	FLA5.10	US8124092 SEQ ID NO: 1	4246
INFL110	Heavy chain	FLD21.140	US8124092 SEQ ID NO: 5	4247
INFL111	Heavy chain	FLA3.14	US8124092 SEQ ID NO: 9	4248
INFL112	Heavy chain	FLD20.19	US8124092 SEQ ID NO: 13	4249
INFL113	Heavy chain	FLD84	US8124092 SEQ ID NO: 42	4250
INFL114	Heavy chain	FLD93	US8124092 SEQ ID NO: 52	4251
INFL115	Heavy chain	FLD122	US8124092 SEQ ID NO: 62	4252
INFL116	Heavy chain	FLD127	US8124092 SEQ ID NO: 72	4253
INFL117	Heavy chain	FLD129	US8124092 SEQ ID NO: 82	4254
INFL118	Heavy chain	FLD132	US8124092 SEQ ID NO: 92	4255
INFL119	Heavy chain	FLD194	US8124092 SEQ ID NO: 102	4256
INFL120	Heavy chain	mAb2	WO2015112994 SEQ ID NO: 80	4257
INFL121	Heavy chain	mAb3	WO2015112994 SEQ ID NO: 84	4258
INFL122	Heavy chain		Tsibane, T. et al., Influenza Human Monoclonal Antibody 1F1 Interacts with Three Major Antigenic Sites and Residues Mediating Human Receptor Specificity in H1N1 Viruses; PLoS Pathol. 8 (12), E1003067 (2012), NCBI Accession #4GXU_S	4259
INFL123	Heavy chain	C05	Ekiert, D.C., et al., Cross-neutralization of influenza A viruses mediated by a single antibody loop; Nature 489 (7417), 526-532 (2012), NCBI Accession #4FNL_H (247aa)	4260
INFL124	Heavy chain	CR8020	Ekiert, D.C., et al., A highly conserved neutralizing epitope on group 2 influenza A viruses; Science 333 (6044), 843-850 (2011); WO2010130636, NCBI Accession #3SDY_H	4261
INFL125	Heavy chain	CR8043	Friesen, R.H. et al., A common solution to group 2 influenza virus neutralization; Proc. Natl. Acad. Sci. U.S.A. 111 (1), 445-450 (2014), NCBI Accession #4NM8_H	4262
INFL126	Heavy chain	CR8059	Dreyfus, C. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession #4FQK_H	4263
INFL127	Heavy chain	CR8071	Dreyfus, C. et al., Highly conserved protective epitopes on influenza B	4264

			viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession #4FQJ_H (234aa)	
INFL128	Heavy chain	CR9114	WO2013079473; WO2014191435; Dreyfus, C., Laursen, N.S. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession #4FQY_H (230aa)	4265
INFL129	Heavy chain	Ch67	Schmidt, A.G., et al., Preconfiguration of the antigen-binding site during affinity maturation of a broadly neutralizing influenza virus antibody; Proc. Natl. Acad. Sci. U.S.A. 110 (1), 264-269 (2013), NCBI Accession #4HKX_A (231aa)	4266
INFL130	Heavy chain	Fab 26/9	Schulze-Gahmen, U. et al., J. Biol. Chem. 263 (32), 17100-17105 (1988); Churchill, M.E., et al., J. Mol. Biol. 241 (4), 534-556 (1994), NCBI Accession #1FRG_H	4267
INFL131	Heavy chain	Fab 3.1	Wyrzucki, A. et al., Alternative Recognition of the Conserved Stem Epitope in Influenza A Virus Hemagglutinin by a VH3-30-Encoded Heterosubtypic Antibody; J. Virol. 88 (12), 7083-7092 (2014), NCBI Accession #4PY8_I	4268
INFL132	Heavy chain	Fab 2g1	Xu, R. et al., A recurring motif for antibody recognition of the receptor-binding site of influenza hemagglutinin; Nat. Struct. Mol. Biol. 20 (3), 363-370 (2013), NCBI Accession #4HG4_N (223aa)	4269
INFL133	Heavy chain	Fab 8m2	Xu, R. et al., A recurring motif for antibody recognition of the receptor-binding site of influenza hemagglutinin; Nat. Struct. Mol. Biol. 20 (3), 363-370 (2013), NCBI Accession #4HFU_H (226aa)	4270
INFL134	Heavy chain	Fab 8f8	Xu, R. et al., A recurring motif for antibody recognition of the receptor-binding site of influenza hemagglutinin; Nat. Struct. Mol. Biol. 20 (3), 363-370 (2013), NCBI Accession #4HF5_H (233aa)	4271
INFL135	Heavy chain	Fab 2d1	Xu, R., et al., Structural basis of preexisting immunity to the 2009 H1N1 pandemic influenza virus; Science 328 (5976), 357-360 (2010), NCBI Accession #3LZF_H (230aa)	4272
INFL136	Heavy chain	Fi6v3	Corti, D. et al., A neutralizing antibody selected from plasma cells that binds to group 1 and group 2 influenza A hemagglutinins; Science 333 (6044), 850-856 (2011), NCBI Accession #3ZTJ_G	4273
INFL137	Heavy Chain	Heavy chain 3WHE_N	Iba, Y., et al., Conserved Neutralizing Epitope at Globular Head of Hemagglutinin in H3N2 Influenza	4274

			Viruses; J. Virol. (2014), NCBI Accession #3WHE_M (226aa)	
INFL138	Heavy chain	7A13	Krause et al. "Human Monoclonal Antibodies to Pandemic 1957 H2N2 and Pandemic 1968 H3N2 Influenza Viruses" J. Virol. 86 (11), 6334-6340 (2012), NCBI Accession #AFH78447	4275
INFL139	Heavy chain	2D1	WO2010127252, US8894997 SEQ ID NO: 7	4276
INFL140	Heavy chain	1F1	WO2010127252, US8894997 SEQ ID NO: 1	4277
INFL141	Heavy chain		WO2010127252, US8894997 SEQ ID NO: 4	4278
INFL142	Heavy chain	1120	WO2010127252, US8894997 SEQ ID NO: 5	4279
INFL143	Heavy chain	4D20	WO2010127252, US8894997 SEQ ID NO: 9	4280
INFL144	Heavy chain		WO2010127252, US8894997 SEQ ID NO: 11	4281
INFL145	Heavy chairs		US20140205614, US20S003 16654 SEQ ID NO: 21	4282
INFL146	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 22	4283
INFL147	Heavy chairs		US20140205614, US20S003 16654 SEQ ID NO: 23	4284
INFL148	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 24	4285
INFL149	Heavy chairs		US2014020565 4, US20 1003 16654 SEQ ID NO: 25	4286
INFL150	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 26	4287
INFL151	Heavy chairs		US2014020565 4, US20 1003 16654 SEQ ID NO: 27	4288
INFL152	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 28	4289
INFL153	Heavy chairs		US2014020565 4, US20 1003 16654 SEQ ID NO: 29	4290
INFL154	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 30	4291
INFL155	Heavy chain		US20140205614, 1JS20 1003 16654 SEQ ID NO: 31	4292
INFL156	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 32	4293
INFL157	Heavy chain		US20140205614, 1JS20 1003 16654 SEQ ID NO: 33	4294
INFL158	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 34	4295
INFL159	Heavy chain		US20140205614, 1JS20 1003 16654 SEQ ID NO: 35	4296
INFL160	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 36	4297
INFL161	Heavy chain		US20140205614, 1JS20 1003 16654 SEQ ID NO: 37	4298
INFL162	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 38	4299
INFL163	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 39	4300
INFL164	Heavy chain		US20140205614, US201003 16654 SEQ ID NO: 40	4301

INFL165	Heavy chain		US20140205614, US20100316654 SEQ ID NO: 41	4302
INFL166	Heavy chain		US20140205614, US20100316654 SEQ ID NO: 42	4303
INFL167	Heavy chain		US20140205614, US20100316654 SEQ ID NO: 43	4304
INFL168	Heavy chain		US20140205614, US20100316654 SEQ ID NO: 44	4305
INFL169	Heavy chain		US20140205614, US20100316654 SEQ ID NO: 45	4306
INFL170	Heavy chain	mAb1	WO2015112994 SEQ ID NO: 76	4307
INFL171	Heavy chain	CR8033	Dreyfus, C., Laursen, N.S. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession # 4FQL_H	4308
INFL172	Heavy chain (Partial)	monoclonal antibody PN-SIA28	Burioni, R. et al., Monoclonal antibodies isolated from human B cells neutralize a broad range of H1 subtype influenza A viruses including swine-origin Influenza virus(S-OIV); Virology (2010), NCBI Accession #ACX30936.1 (122aa)	4309
INFL173	Heavy chain (Partial)	monoclonal antibody PN-SIA49	Burioni, R. et al., Monoclonal antibodies isolated from human B cells neutralize a broad range of H1 subtype influenza A viruses including swine-origin Influenza virus(S-OIV); Virology (2010), NCBI Accession #ACX30937.1 (127aa)	4310
INFL174	Heavy chain cdr1	Ab1A2	WO2015028478 SEQ ID 6	4311
INFL175	Heavy chain cdr2	Ab1A2	WO2015028478 SEQ ID 7	4312
INFL176	Heavy chain cdr3	Ab1A2	WO2015028478 SEQ ID 8	4313
INFL177	Heavy chain constant region, Human igg1		US8992929 SEQ ID NO. 22	4314
INFL178	Heavy chain Fab	CT147	WO2013048153, US20140234336 SEQ ID NO: 38	4315
INFL179	Heavy chain Fab	CT164	WO2013048153, US20140234336 SEQ ID NO: 42	4316
INFL180	Heavy chain Fab	CT166	WO2013048153, US20140234336 SEQ ID NO: 44	4317
INFL181	Heavy chain G2	h2B9	US9115201 SEQ ID NO: 7	4318
INFL182	Heavy chain G5	h2B10	US9115201 SEQ ID NO: 8	4319
INFL183	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 1	4320
INFL184	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 2	4321
INFL185	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 3	4322
INFL186	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 4	4323
INFL187	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 5	4324
INFL188	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 6	4325
INFL189	Heavy chain variable (exemplary)	HC-VD from US2013030234	US2013030234 SEQ ID NO: 7	4326
INFL190	Heavy chain variable **(exemplary)**	HC-VD from **US20 13030234**	US2013030234 SEQ ID NO: 8	4327

INFL 191	Heavy chain variable (exemplary)	HC-VD from US20 13030234	US2013030234 SEQ ID NO: 9	4328
INFL192	Heavy chain variable (exemplary)	HC-VD from US20 13030234	US2013030234 SEQ ID NO : 10	4329
INFL 193	Heavy chairs variable (exemplary)	HC-VD from US20 13030234	US20 13030234 SEQ ID NO: 11	4330
INTL. 194	Heavy chain variable (exemplary)	HC-VD from US20 13030234	US2013030234 SEQ ID NO: 12	4331
INFL 195	Heavy chairs variable (exemplary)	HC-VD from US20 13030234	US20 13030234 SEQ ID NO: 13	4332
INTL. 196	Heavy chain variable (exemplary)	HC-VD from US20 13030234	US2013030234 SEQ ID NO: 14	4333
INFL 197	Heavy chairs variable (exemplary)	HC-VD from US20 13030234	US20 13030234 SEQ ID NO: 15	4334
INTL. 198	Heavy chain variable (exemplar)')	HC-VD from US20 13030234	US2013030234 SEQ ID NO: 16	4335
INFL 199	Heavy chairs variable region	CR6141	US20 150104459 SEQ ID NO: 199	4336
INTL. 200	Heavy chain variable region	39.18 B11	US20140 161822 SEQ ID NO: 154	4337
INFL20 1	Heavy chairs variable region	39.18 E12	US20140161822 SEQ ID NO: 158	4338
INFL202	Heavy chain variable region	GG3	WO20 14 159960 SEQ ID NO: 17	4339
1NFL203	Heavy chain variable region	N547	US8003 106 SEQ ID NO: 28	4340
INFL204	Heavy chain variable region	L66	US8003 106 SEQ ID NO: 30	4341
INFL205	Heavy chain variable region	C40	US8003 106 SEQ ID NO: 26	4342
INFL206	Heavy chain variable region	14C2	US8080244 SEQ ID NO: 6	4343
INFL207	Heavy chain variable region	hl4C2	US8080244 SEQ ID NO: 2	4344
INFL208	Heavy chain variable region	8G9	US8603467 SEQ ID NO: 2	4345
INFL209	Heavy chain variable region	13D4	US8603467 SEQ ID NO: 6	4346
INFL210	Heavy chain variable region	20A11	US8603467 SEQ ID NO: 10	4347
INFL21 1	Heavy chain variable region	VN04-2-HuGl	US20100150941 SEQ ID NO: 5	4348
INFL212	Heavy chain variable region	VN04-3-HuGl	US20100150941 SEQ ID NO: 7	4349
INFL213	Heavy chain variable region	FI6 variant 1	US8871207 SEQ ID NO: 13	4350
INFL214	Heavy chain variable region	FI6 variant 2	US8871207 SEQ ID NO: 33	4351
INFL215	Heavy chain variable region	FI6 variant 3	US8871207 SEQ ID NO: 55	4352
INFL216	Heavy chain variable region	FI6 variant 4, FI6 variant 5	US8871207 SEQ ID NO: 59	4353
INFL217	Heavy chain variable region	FI28 variant 1	US8871207 SEQ ID NO: 29	4354
INFL218	Heavy chain variable region	FI28 variant 2	US8871207 SEQ ID NO: 35	4355
INFL219	Heavy chain variable region	21B15	US8858948 SEQ ID NO: 44	4356

INFL220	Heavy chain variable region	3241_G23	US8858948 SEQ ID NO: 116	4357
INFL221	Heavy chain variable region	3244_I10	US8858948 SEQ ID NO: 120	4358
INFL222	Heavy chairs variable region	3243_J07	US8858948 SEQ ID NO: 124	4359
INTL.223	Heavy chain variable region	3259_I21	US8858948 SEQ ID NO: 128	4360
INFL224	Heavy chairs variable region	3245_O19	US8858948 SEQ ID NO: 132	4361
INFL225	Heavy chain variable region	3244_H04	US8858948 SEQ ID NO: 136	4362
INFL226	Heavy chairs variable region	3136_G05	US8858948 SEQ ID NO: 140	4363
INFL227	Heavy chain variable region	3252_C13	US8858948 SEQ ID NO: 144	4364
INFL228	Heavy chairs variable region	3255_J06	US8858948 SEQ ID NO: 148	4365
INTL.229	Heavy chain variable region	3420_I23	US8858948 SEQ ID NO: 152	4366
INFL230	Heavy chairs variable region	3139_P23	US8858948 SEQ ID NO: 156	4367
INFL231	Heavy chain variable region	3139_P23	US8858948 SEQ ID NO: 158	4368
INFL232	Heavy chain variable region	3248_P18	US8858948 SEQ ID NO: 162	4369
INFL233	Heavy chain variable region	3253_P10	US8858948 SEQ ID NO: 166	4370
INFL234	Heavy chain variable region	3260_D19	US8858948 SEQ ID NO: 170	4371
INFL235	Heavy chain variable region	3362_B11	US8858948 SEQ ID NO: 172	4372
INFL236	Heavy chain variable region	3242_P05	US8858948 SEQ ID NO: 176	4373
INFL237	Heavy chain variable region	2K11	Krause, J.C. el al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence arsd interclonal convergence" j. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AE016793	4374
INFL238	Heavy chain variable region	2010	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" j. Immunol 187 (7), 3704-3711 (2011), NCBI Accession #AE016795	4375
INFL239	Heavy chain variable region	4K8	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AE016799	4376
INFL240	Heavy chain variable region	6D9	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AE01 6801	4377

INFL241	**Heavy** chain variable region	4D20	Yu, X. et **al** "Neutralizing antibodies derived **from** the **B** cells of 1918 **influenza pandemic survivors",** Nature 455 (7212), 532-536, **NCBI** Accession **#ACI04579**	4378
INFL242	**Heavy chain** variable region	**2B12**	**Yu,** X. et al "Neutralizing antibodies **derived from** *the* B cells **of 1918** influenza pandemic **survivors",** Nature 455 (7212), 532-536, **NCBI** Accession **#ABY48866**	4379
INFL243	Heavy chain variable region	8D4	NCBI Accession **#AFI57036**	4380
INFL244	**Heavy** chain variable region	5B6	NCBI Accession **#AFI57040**	4381
INFL245	Heavy chain variable region	A66	**WO2009079259, US201 10038935, US2014001 1982** SEQ ID NO: 32	4382
INFL246	**Heavy** chain variable region	D7	WO2009079259, US201 10038935, US201400 1982 SEQ ID NO: 6	4383
INFL247	Heavy chain variable region	D8, D80	**WO2009079259, US201 10038935, US2014001 982** SEQ **ID** NO: 12	4384
INFL248	Heavy **chain** variable region	E88	WO2009079259, **US201 10038935, US20** 1400 1982 SEQ ID **NO:** 36	4385
INFL249	Heavy chain variable region	E90, **F!O**	WO2009079259, **US201 10038935, US2014001 982** SEQ **ID** NO: 18	4386
INFL250	Heavy **chain variable** region	**F10**	WO2009079259, **US201 10038935, US20** 1400 1982 SEQ ID **NO:** 12	4387
INFL251	Heavy chain variable region	G17	WO2009079259, **US201 10038935, US2014001 982** SEQ **ID** NO: 24	4388
INFL252	**Heavy chain variable** region	H40	WO2009079259, **US201 10038935, US20** 1400 1982 SEQ ID **NO:** 28	4389
INFL253	Heavy chain variable region	**CH65**	**WO20 13020074, US20 140302043** SEQ ID NO: 14	4390
INFL254	**Heavy chain variable** region	**CH66**	**WO2013020074, US20140302043** SEQ **ID NO:** 15	4391
INFL255	**Heavy** chain variable region	CH67	**WO20 13020074, US20 140302043** SEQ **ID** NO: 16	4392
INFL256	**Heavy chain variable** region	**CL860UCA**	**WO2013020074,** US20140302043 SEQ **ID NO:** 13	4393
INFL257	Heavy chairs variable region	**Antibody** 1	**WO20** 15051010 SEQ ID NO: 2	4394
INTL.258	Heavy chain variable region	Antibody 2	WO2015051010 SEQ ID **NO:** 12	4395
INFL259	Heavy chairs variable region	**Antibody** 3	**WO20** 15051010 SEQ ID NO: 22	4396
INTL.260	Heavy chain variable region	Antibody 4	WO2015051010 SEQ ID **NO:** 32	4397
INFL261	Heavy chairs variable region	**Antibody** 5	**WO20** 15051010 SEQ ID NO: 42	4398
INFL262	Heavy chain variable region	Antibody 6	WO2015051010 SEQ ID **NO:** 52	4399
INFL263	Heavy chairs variable region	Antibody 7	**WO20** 15051010 SEQ ID NO: 62	4400
INTL.264	Heavy chain variable region	Antibody 8	WO2015051010 SEQ ID **NO:** 72	4401
INFL265	**Heavy** chain variable region	Antibody 9	**WO20 15051010** SEQ ID **NO:** 82	4402
INFL266	Heavy chain variable region	Antibody 10	**WO2015051010** SEQ **ID NO:** 92	4403

INFL267	**Heavy** chain variable region	Antibody 11	**WO20 15051010** SEQ ID NO: 102	4404
INFL268	**Heavy chain** variable **region**	Antibody 12	**WO2015051010** SEQ ID NO: 112	4405
INFL269	Heavy chairs variable region	**Antibody** 13	**WO20 1505 1010** SEQ ID NO: 122	4406
INTL.270	Heavy chain variable **region**	Antibody 14	**WO2015051010** SEQ **ID NO:** 132	4407
INFL27 1	Heavy chairs variable region	**Antibody** 15	**WO20 1505 1010** SEQ ID NO: 142	4408
INFL272	Heavy chain variable region	Antibody 3-GL	**WO2015051010** SEQ **ID NO:** 152	4409
INFL273	Heavy chairs variable region	**EM4C04**	**US20 120282273** SEQ **ID** NO: 2	4410
INTL.274	Heavy chain variable region	005-2G02	**WO2013059524, US20140348851** SEQ **ID NO:** 1	**441 1**
INFL275	Heavy chairs variable region	005-2G02	**WO20** 13059524, **US20 14034885** 1 SEQ **ID** NO: **9**	4412
INTL.276	Heavy chain variable region	09-2A06	**WO2013059524, US20140348851** SEQ **ID NO: 21**	4413
INFL277	Heavy chairs variable region	09-2A06	**WO20** 13059524, **US20 14034885** 1 SEQ ID NO: 29	4414
INFL278	Heavy chain variable region	**09-3A01**	WO20L3059524, 1JS20140348851 SEQ ID NO: 41	4415
INFL279	**Heavy** chain variable region	**09-3**A O**1**	**WO20 13059524, US20140348851** SEQ ID NO: 49	4416
INFL280	Heavy chain variable region	**70-IF02**	WO20 12096994, US20 140046039 SEQ ID NO: 18	4417
INFL28 1	**Heavy** chain variable region		**US20120058124** SEQ **ID** NO: **10**	4418
INFL282	Heavy chain variable region		**US20120058124** SEQ ID **NO:** 11	4419
INFL28.3	**Heavy** chain variable region		**US20120058124** SEQ **ID** NO: **12**	4420
INFL284	Heavy chain variable region		**US20120058124** SEQ ID **NO: 13**	4421
INFL285	**Heavy** chain variable region		**US20120058124** SEQ **ID** NO: 14	4422
INFL286	Heavy chain variable region	81.39	**US20140161822,** US20 140248286, **WO20 14078268** SEQ ID **NO:** 111	4423
INFL287	**Heavy chain variable** region	81.39	**US20140161822, US20 140248286, WO20 14078268** SEQ ID NO: 115	4424
INFL288	Heavy chain variable region	39.29	**US20140161822,** US20 140248286, **WO20 14078268** SEQ ID NO: 134	4425
INFL289	**Heavy chain variable** region	39.29	**US20140161822, US20 140248286, WO20 14078268** SEQ ID NO: 138	4426
INFL290	Heavy chain variable region	39.29	**US20140161822,** US20 140248286, **WO20 14078268** SEQ ID NO: 142	4427
INFL291	**Heavy chain variable** region	39.29	**US20140161822, US20 140248286, WO20 14078268** SEQ ID NO: 148	4428
INFL292	Heavy chain variable region	36.89	**US20140161822,** US20 140248286, **WO20 14078268** SEQ ID NO: 160	4429
INFL293	**Heavy chain variable** region	9.0 1F3	**US20140161822, US20 140248286, WO20 14078268** SEQ ID NO: 164	4430
INFL294	Heavy chain variable region	23.06C2	**US20140161822, US20** 140248286, **WO20 14078268** SEQ ID NO: 168	443 1
INFL295	**Heavy chain variable** region	39.29	**US20140161822, US20 140248286, WO20 14078268** SEQ ID NO: 234	4432

INFL296	Heavy chain variable region	F16 Variant 5	WO20 1301 1347, US20140271655, US887 1207 SEQ ID NO: 59	4433
INFL297	Heavy chain variable reedon	F16 Variant 3	WO20130 U347, US2014027 1655, US8871207 SEQ ID NO: 55	4434
INFL298	Heavy chairs variable region	F16 Variant 2	WO20 100 10466 SEQ ID NO: 33	4435
INFL299	Heavy chain variable reedon	FC4 1	WO20 100 10467 SEQ ID NO 60	4436
INFL300	Heavy chairs variable region	FE43	WO20 10010467 SEQ ID NO 74	4437
INFL301	Heavy chain variable reedon	FB75, FB1 10, FBI 77	WO20 100 10467 SEQ ID NO 12 1	4438
INFL302	Heavy chairs variable region	FE17	WO20 10010467 SEQ ID NO 105	4439
INFL303	Heavy chain variable reedon	FB79	WO20 100 10467 SEQ ID NO 13 1	4440
INFL304	Heavy chairs variable region	FC1C	WO20 10010467 SEQ ID NO 139	4441
INFL305	Heavy chain variable reedon	FC6	WO20 100 10467 SEQ ID NO 45	4442
INFL306	Heavy chairs variable region	FES3	WO20 10010467 SEQ ID NO 89	4443
INFL307	Heavy chain variable region	7A7	WO2010138564 SEQ ID NO: 6	4444
INFL308	Heavy chain variable region	12DI	WO20 10 138564 SEQ ID NO: 12	4445
INFL309	Heavy chain variable region	66A6	WO2010138564 SEQ ID NO: 16	4446
INFL310	Heavy chain variable region	B-l	US8975378, US20 1103 19600, WO20 10073647 SEQ ID NO: 27	4447
INFL311	Heavy chain variable region	D1	US8975378, US201 103 19600. WO20 10073647 SEQ ID NO: 29	4448
INFL312	Heavy chain variable region	E-2	US8975378, US20 1103 19600, WO20 10073647 SEQ ID NO: 31	4449
INFL313	Heavy chain variable region	B-3	US8975378, US201 103 19600. WO20 10073647 SEQ ID NO: 33	4450
INFL314	Heavy chain variable region	5A7	WO20 13 1 14885, US20140377262 SEQ ID NO: 33	4451
INFL315	Heavy chain variable region	3A2	WO2013 114885, US20140377262 SEQ ID NO: 37	4452
INFL316	Heavy chain variable region	10C4	WO20 13 114885, US20 140377262 SEQ ID NO: 41	4453
INFL317	Heavy chain variable region	Fab49	WO2009144667, US201 10076265 SEQ ID NO: 1	4454
INFL318	Heavy chain variable region	Fab28 IgG PN-SIA28	WO20091 15972, WO201 1117848, US201 100 14187 SEQ ID NO: I	4455
INFL319	Heavy chain variable region	TCN-522	US20120207760, US8916160 SEQ ID NO: 77 1; US8900590 SEQ ID NO: 32	4456
INFL320	Heavy chain variable region	CR8019	WO20 10 130636 SEQ ID NO: 26	4457
INFL321	Heavy chain variable region	CR8020	WO20 10 130636 SEQ ID NO: 30	4458
INFL322	Heavy chain variable region	CR8021	WO20 10 130636 SEQ ID NO: 34	4459
INFL323	Heavy chain variable region	CR8038	WO20 10130636 SEQ ID NO: 38	4460
INFL324	Heavy chain variable region	CR8039	WO20 10 130636 SEQ ID NO: 42	4461

INFL325	**Heavy** chain variable **segion**	**CR8040**	**WO20 10 130636** SEQ ID NO: 46	4462
INFL326	**Heavy chain** variable **region**	**CR804 1**	**WO20 10 130636** SEQ ID NO: 50	4463
INFL327	Heavy chairs variable region	CR8043	**WO20 10** 130636 SEQ ID NO: 54	4464
INFL328	Heavy chain variable **region**	**CR8049**	**WO2010130636** SEQ **ID NO:** 58	4465
INFL329	Heavy chairs variable region	**CR8050**	WO20 10 130636 SEQ ID NO: 61	4466
INFL330	Heavy chain variable region	**CR8052**	**WO2010130636** SEQ **ID NO:** 65	4467
INFL33 1	Heavy chairs variable region	**CR8055**	WO20 10 130636 SEQ ID NO: 69	4468
INFL332	Heavy chain variable region	**CR8057**	**WO2010130636** SEQ **ID NO: 73**	4469
INFL333	Heavy chairs variable region	**CR8069**	**WO20 10** 130636 SEQ ID NO: 77	4470
INFL334	Heavy chain variable region	**CR6255**	**US200903** 11265, **US8691223, US9109017, WO2008028946A** SEQ ID NO: 59	447 1
INFL335	Heavy chain variable region	CR6257	**US200903 11265, US8691223,** US91090 17, **WO2008028946** SEQ ID NO: **6 1**	4472
INFL336	**Heavy** chain variable **region**	**CR6260**	**US200903** 11265, US8691223, **US9 1090 17, WO2008028946** SEQ ID NO: 63	4473
INFL337	Heavy chairs variable region	**CR6261**	**US200903** 11265, **US8691** 223, US91090 17, **WO2008028946** SEQ ID NO: 65	4474
INFL338	**Heavy** chain variable region	CR6262	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: **67**	4475
INFL339	**Heavy chain variable** region	**CR6268**	US200903 11265, US8691223, **US9109017, WO2008028946** SEQ ID **NO:** 69	4476
INFL340	**Heavy chain variable** region	**CR6307**	US200903 11265, US8691223, **US9109017, WO2008028946** SEQ ID NO: 71	4477
INFL341	Heavy chain variable region	**CR63 10**	US200903 11265, US8691223, **US9 109017, WO2008028946** SEQ ID NO: 73	4478
INFL342	Heavy chain variable region	CR63 14	**US200903 11265, US8691223,** US91090 17, **WO2008028946** SEQ ID NO: 75	4479
INFL343	**Heavy** chain variable **region**	CR6323	**US200903** 11265, **US8691223, US9 1090 17, WO2008028946** SEQ ID NO: **77**	4480
INFL344	Heavy chairs variable segion	CR6325	**US200903** 11265, US8691 223, **US91090 17, WO2008028946** SEQ ID NO: 79	4481
INFL345	**Heavy** chain **variable** region	CR633 1	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID **NO:** 81	4482
INFL346	**Heavy chain variable** region	**CR6344**	US200903 11265, **US8691223, US9109017, WO2008028946** SEQ **ID** NO: 83	4483

INFL347	**Heavy** chain variable region	**CR6141**	**US200903** 11265, **US8691223, US9** 1090 I7, **WO2008028946** SEQ ID **NO: 3** 17	4484
INFL348	**Heavy** chain variable region	CR6272	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: 321	4485
INFL349	**Heavy** chain variable region	**CR6296**	US200903 11265, US8691223, US9109017, **WO2008028946** SEQ ID **NO:** 325	4486
INFL350	Heavy **chain** variable region	**CR6301**	**US200903** 11265, US8691223, **US9109017, WO2008028946** SEQ ID NO: 329	4487
INFL35 1	Heavy chain variable region	**CR6327**	**US200903** 11265, US8691223, **US9109017, WO2008028946** SEQ ID NO: 333	4488
1NFL352	Heavy chain variable region	CR6328	US200903 11265, US8691223, **US9 109017, WO2008028946** SEQ ID NO: 337	4489
INFL353	Heavy chain variable region	CR6329	US200903 11265, US8691223, **US9 1090 I7, WO2008028946** SEQ ID NO: 341	4490
INFL354	Heavy chairs variable region	CR6332	**US200903** 11265, US8691 223, US9109017, **WO2008028946** SEQ ID NO: 345	4491
INFL355	**Heavy** chain variable region	CR6334	**US200903** 11265, **US8691223,** US9109017, **WO2008028946** SEQ ID NO: 349	4492
INFL356	**Heavy chain variable** region	**CR6336**	**US200903 11265**, US8691223, US9109017, **WO2008028946** SEQ ID **NO:** 353	4493
INFL357	Heavy **chain variable** region	**CR6339**	**US200903** 11265, US8691223, **US9109017, WO2008028946** SEQ ID NO: 357	4494
INFL358	Heavy chain variable region	CR6342	**US200903** 11265, US8691223, **US9109017, WO2008028946** SEQ ID NO: 361	4495
INFL359	Heavy chain variable region	CR6343	US200903 11265, US8691223, US91090 17, **WO2008028946** SEQ ID NO: 365	4496
INFL360	Heavy chain variable region	**CR9003**	**US20 1401201 13** SEQ ID NO: 2	4497
INFL36 1	**Heavy chain variable** region	**CR9004**	**US20140 1201 13** SEQ ID NO: **6**	4498
INFL362	Heavy chain variable region	CR9005	**US201401201** 13 SEQ ID NO: 10	4499
INFL363	**Heavy chain variable** region	**CR9006**	**US20140 1201 13** SEQ ID NO: **14**	4500
INFL364	Heavy chain variable region	**CR9007**	**US201401201** 13 SEQ ID NO: 18	4501
INFL365	**Heavy chain variable** region	**CR9008**	**US20140 1201 13** SEQ ID NO: 22	4502
INFL366	**Heavy** chain variable region	**CR9009**	**US201401201** 13 SEQ ID NO: 26	4503
INFL367	Heavy chain variable region	**CR90** 10	**US20140 I201** s3 SEQ ID **NO:** 30	4504
INFL368	Heavy chairs variable region	**CR901 1**	**US20** 140120 113 SEQ ID **NO:** 34	4505
INFL369	Heavy chain variable region	**CR90** 12	**US20140 I201** s3 SEQ ID **NO:** 38	4506

INFL370	Heavy chain variable region	CR9029	US201401201 13 SEQ ID NO: 42	4507
INFL371	Heavy chain variable region	CR9030	US20140 1201 13 SEQ ID NO: 46	4508
INFL372	Heavy chairs variable region	CR903 1	US20140120113 SEQ ID NO: 50	4509
INFL373	Heavy chain variable region	CR91 12	US20140 I201 S3 SEQ ID NO: 54	4510
INFL374	Heavy chairs variable region	CR9113	US20140120113 SEQ ID NO: 58	4511
INFL375	Heavy chain variable region	CR9114	US20140 I201 S3 SEQ ID NO: 62	4512
INFL376	Heavy chairs variable region	CR8033	US8852595 SEQ ID NO: 71	4513
INTL.377	Heavy chain variable region	CR8059	US8852595 SEQ ID NO: 75	4514
INFL378	Heavy chairs variable region	CR8071	US8852595 SEQ ID NO: 78	4515
INFL379	Heavy chain variable region	CR 1005 1	US8852595 SEQ ID NO: 81	4516
INFL380	Heavy chairs variable region	CR10049	US8852595 SEQ ID NO: 85	4517
1NFL381	Heavy chain variable region	CR 10023	US8852595 SEQ ID NO: 89	4518
INFL382	Heavy chain variable region	CR10032	US8852595 SEQ ID NO: 93	4519
INFL383	Heavy chain variable region	CR 11035	US8852595 SEQ ID NO: 101	4520
INFL384	Heavy chain variable region	CR 11036	US8852595 SEQ ID NO: 105	4521
INFL385	Heavy chain variable region	CR 11038	US8852595 SEQ ID NO: 109	4522
INFL386	Heavy chain variable region	CR1 1039	US8852595 SEQ ID NO: 113	4523
INFL387	Heavy chain variable region	CR803 1	US8852595 SEQ ID NO: 119	4524
INFL388	Heavy chain variable region	CR8032	US8852595 SEQ ID NO: 123	4525
INFL389	Heavy chain variable region	CR8034	US8852595 SEQ ID NO: 127	4526
INFL390	Heavy chain variable region	CR8035	US8852595 SEQ ID NO: 131	4527
INFL391	Heavy chain variable region		US8992929 SEQ ID NO: 4	4528
INFL392	Heavy chain variable region	M2e	US8420794 SEQ ID NO: 2	4529
INFL393	Heavy chain variable region		US8715743, US20140275492 SEQ ID NO: 22	4530
INFL394	Heavy chain variable region		US8715743, US20140275492 SEQ ID NO: 25	4531
INFL395	Heavy chain variable region		US8715743, US20140275492 SEQ ID NO: 36	4532
INFL396	Heavy chain variable region	4A10	Kraiise, J.C. et al. "Epiiope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-37 11 (201 1), NCBI Accession #AEO 16797	4533

INFL397	Heavy chain variable region	anti-1918 influenza HA Ig	Yu, X., et al., Neutralizing antibodies derived from the B cells of 1918 influenza pandemic survivors; Nature 455 (7212), 532-536 (2008), NCBI Accession #ACI04579.1 (129aa)	4534
INFL398	Heavy chain variable region	TCN-522 (3212_I12)	US20150086555 SEQ ID NO: 33	4535
INFL399	Heavy chain variable region	TCN-521 (3280_D18)	US20150086555 SEQ ID NO: 21	4536
INFL400	Heavy chain variable region	TCN-523 (5248_A17)	US20150086555 SEQ ID NO: 45	4537
INFL401	Heavy chain variable region	TCN-563 (5237_B21)	US20150086555 SEQ ID NO: 57	4538
INFL402	Heavy chain variable region	TCN-526 (5084_C17)	US20150086555 SEQ ID NO: 69	4539
INFL403	Heavy chain variable region	TCN-527 (5086_C06)	US20150086555 SEQ ID NO: 81	4540
INFL404	Heavy chain variable region	TCN-528 (5087_P17)	US20150086555 SEQ ID NO: 93	4541
INFL405	Heavy chain variable region	TCN-529 (5297_H01)	US20150086555 SEQ ID NO: 105	4542
INFL406	Heavy chain variable region	TCN-530 (5248_H10)	US20150086555 SEQ ID NO: 117	4543
INFL407	Heavy chain variable region	TCN-531 (5091_H13)	US20150086555 SEQ ID NO: 129	4544
INFL408	Heavy chain variable region	TCN-532 (5262_H18)	US20150086555 SEQ ID NO: 141	4545
INFL409	Heavy chain variable region	TCN-533 (5256_A17a), TCN-564 (5256_A17b)	US20150086555 SEQ ID NO: 153	4546
INFL410	Heavy chain variable region	TCN-534 (5249_B02)	US20150086555 SEQ ID NO: 161	4547
INFL411	Heavy chain variable region	TCN-535 (5246_P19), TCN-558 (5248_H10b)	US20150086555 SEQ ID NO: 173	4548
INFL412	Heavy chain variable region	TCN-536 (5095_N01)	US20150086555 SEQ ID NO: 184	4549
INFL413	Heavy chain variable region	TCN-537 (3194_D21)	US20150086555 SEQ ID NO: 195	4550
INFL414	Heavy chain variable region	TCN-538 (3206_O17)	US20150086555 SEQ ID NO: 207	4551
INFL415	Heavy chain variable region	TCN-539 (5056_A08)	US20150086555 SEQ ID NO: 219	4552
INFL416	Heavy chain variable region	TCN-540 (5060_F05)	US20150086555 SEQ ID NO: 231	4553
INFL417	Heavy chain variable region	TCN-541 (5062_M11)	US20150086555 SEQ ID NO: 243	4554
INFL418	Heavy chain variable region	TCN-542 (5079_A16)	US20150086555 SEQ ID NO: 255	4555
INFL419	Heavy chain variable region	TCN-543 (5081_G23)	US20150086555 SEQ ID NO: 267	4556
INFL420	Heavy chain variable region	TCN-544 (5082_A19)	US20150086555 SEQ ID NO: 279	4557
INFL421	Heavy chain variable region	TCN-545 (5082_I15)	US20150086555 SEQ ID NO: 291	4558
INFL422	Heavy chain variable region	TCN-546 (5089_LOS)	US20150086555 SEQ ID NO: 302	4559

INFL423	Heavy chain variable region	TCN-547 (5092_F11)	US20150086555 SEQ ID NO: 313	4560
INFL424	Heavy chain variable reedon	TCN-548 (5092_P01)	US20150086555 SEQ ID NO: 325	4561
INFL425	Heavy chairs variable region	TCN-549 (5092_P04)	US20150086555 SEQ ID NO: 335	4562
INFL426	Heavy chain variable reedon	TCN-550 (5096_F06)	US20150086555 SEQ ID NO: 346	4563
INFL427	Heavy chairs variable region	TCN-551 (5243_D01)	US20150086555 SEQ ID NO: 358	4564
INTL.428	Heavy chain variable reedon	TCN-552 (5249_123)	US20150086555 SEQ ID NO: 370	4565
INFL429	Heavy chairs variable region	TCN-553 (5261_C18)	US20150086555 SEQ ID NO: 382	4566
INFL430	Heavy chain variable reedon	TCN-554 (5277_M05)	US20150086555 SEQ ID NO: 392	4567
INFL431	Heavy chairs variable region	TCN-555 (5246_L16)	US20150086555 SEQ ID NO: 403	4568
INFL432	Heavy chain variable reedon	TCN-556 (5089_K12)	US20150086555 SEQ ID NO: 408	4569
INFL433	Heavy chairs variable region	TCN-557 (5081_A04)	US20150086555 SEQ ID NO: 420	4570
INFL434	Heavy chain variable region	TCN-559 (5097_G08)	US20150086555 SEQ ID NO: 434	4571
INFL435	Heavy chain variable region	TCN-560 (5084_P10)	US20150086555 SEQ ID NO: 446	4572
INFL436	Heavy chain variable region	TCN-504 (3251_K17)	US20150086555 SEQ ID NO: 510	4573
INFL437	Heavy chain variable region	AB1	US20120093834, WO2009121004 SEQ ID NO: 4	4574
INFL438	Heavy chain variable region	AB2	US20120093834, WO2009121004 SEQ ID NO: 45	4575
INFL439	Heavy chain variable region	AB3	US20120093834, WO2009121004 SEQ ID NO: 9	4576
INFL440	Heavy chain variable region	AB4, AB5, AB6	US20120093834, WO2009121004 SEQ ID NO: 61	4577
INFL441	Heavy chain variable region	VN04-2	WO2008033105 SEQ ID NO: 5	4578
INFL442	Heavy chain variable region	VN04-3	WO2008033105 SEQ ID NO: 7	4579
INFL443	Heavy chain variable region	1286-C05	WO2010132604, US20120128671 SEQ ID NO: 1	4580
INFL444	Heavy chain variable region	1286-A11	WO2010132604, US20120128671 SEQ ID NO: 2	4581
INFL445	Heavy chain variable region	CR8001	WO2010130636 SEQ ID NO: 2	4582
INFL446	Heavy chain variable region	CR8003	WO2010130636 SEQ ID NO: 6	4583
INFL447	Heavy chain variable region	CR8015	WO2010130636 SEQ ID NO: 10	4584
INFL448	Heavy chain variable region	CR8016	WO2010130636 SEQ ID NO: 14	4585
INFL449	Heavy chain variable region	CR8017	WO2010130636 SEQ ID NO: 18	4586
INFL450	Heavy chain variable region	CR8018	WO2010130636 SEQ ID NO: 22	4587
INFL451	Heavy chain variable region (Partial)	anti-1918 influenza HA Ig	Yu, X., et al.. Neutralizing antibodies derived from the B cells of 1918 influenza pandemic survivors; Nature	4588

			455 (7212), 532-536 (2008), NCBI Accession #ACI04581.1 (145aa)	
INFL452	Heavy chain variable region mouse IgG	1A2	WO2015028478 SEQ ID NO: 2	4589
INFL453	Heavy chain variable region mouse IgG	7B8	WO2015028478 SEQ ID NO: 4	4590
INFL454	Heavy chain variable region, partial	monoclonal antibody TCN-031	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23854.1 (120aa)	4591
INFL455	Heavy chain variable region, partial	monoclonal antibody TCN-032	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23853.1 (120aa)	4592
INFL456	Heavy chain variable region, partial	monoclonal antibody 3362_B11	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23869.1 (123aa)	4593
INFL457	Heavy chain variable region, partial	monoclonal antibody 3260_D19	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23868.1 (118aa)	4594
INFL458	Heavy chain variable region, partial	monoclonal antibody 3253_P10	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23867.1 (121aa)	4595
INFL459	Heavy chain variable region, partial	monoclonal antibody 3248_P18	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23866.1 (120aa)	4596
INFL460	Heavy chain variable region, partial	monoclonal antibody 3139_P23	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23865.1 (119aa)	4597
INFL461	Heavy chain variable region, partial	monoclonal antibody 3420_I23	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663	4598

			(2010), NCBI Accession #ADK23864.1 (121aa)	
INFL462	Heavy chain variable region, partial	monoclonal antibody 3255_J06	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23863.1 (119aa)	4599
INFL463	Heavy chain variable region, partial	monoclonal antibody 3252_C13	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession # ADK23862.1 (119aa)	4600
INFL464	Heavy chain variable region, partial	monoclonal antibody 3136_G05	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23861.1 (120aa)	4601
INFL465	Heavy chain variable region, partial	monoclonal antibody 3244_H04	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23860.1 (118aa)	4602
INFL466	Heavy chain variable region, partial	monoclonal antibody 3245_O19	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23859.1 (118aa)	4603
INFL467	Heavy chain variable region, partial	monoclonal antibody 3259_J21	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23858.1 (120aa)	4604
INFL468	Heavy chain variable region, partial	monoclonal antibody 3243_J07	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23857.1 (121aa)	4605
INFL469	Heavy chain variable region, partial	monoclonal antibody 3244_I10	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession # ADK23856.1 (121aa)	4606

INFL470	Heavy chain variable region, partial	monoclonal antibody 3241_G23	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23855.1 (122aa)	4607
INFL471	Heavy chain variable region, partial	Monoclonal antibody clone 5E4	Yasugi, M. et al., Emerging Antigenic Variants at the Antigenic Site Sb in Pandemic A(H1N1)2009 Influenza Virus in Japan Detected by a Human Monoclonal Antibody; PLoS ONE 8 (10), E77892 (2013), NCBI Accession #BAM76754.1 (141aa)	4608
INFL472	Heavy chain variable region, partial	100F4-HV	Hu, H., et al., A Human Antibody Recognizing a Conserved Epitope of H5 Hemagglutinin Broadly Neutralizes Highly Pathogenic Avian Influenza H5N1 Viruses; J. Virol. 86 (6), 2978-2989 (2012), NCBI Accession #AEL30603.1 (121aa)	4609
INFL473	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40460.1 (120aa)	4610
INFL474	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40459.1 (127aa)	4611
INFL475	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40458.1 (129aa)	4612
INFL476	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40457.1 (132aa)	4613
INFL477	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40456.1 (127aa)	4614
INFL478	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40455.1 (121aa)	4615
INFL479	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40454.1 (126aa)	4616
INFL480	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature	4617

			(2014), NCBI Accession #AIN40453.1 (120aa)	
INFL481	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40452.1 (122aa)	4618
INFL482	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40451.1 (125aa)	4619
INFL483	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40450.1 (126aa)	4620
INFL484	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40449.1 (129aa)	4621
INFL485	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40448.1 (119aa)	4622
INFL486	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40447.1 (120aa)	4623
INFL487	Heavy chain variable region, partial, Anti-stem HA immunoglobulin	anti-stem HA immunoglobulin	Pappas, L. et al., Rapid development of broadly influenza neutralizing antibodies through redundant mutations; Nature (2014), NCBI Accession #AIN40446.1 (120aa)	4624
INFL488	Heavy chain, Human igg,	Fab 39.29	Nakamura, G. et al., An in vivo human-plasmablast enrichment technique allows rapid identification of therapeutic influenza a antibodies; Cell Host Microbe 14 (1), 93-103 (2013), NCBI Accession #4KVN_H (227aa)	4625
INFL489	Heavy chain, Igg1	Fab H5m9	Zhu, X., et al., A Unique and Conserved Neutralization Epitope in H5N1 Influenza Viruses Identified by an Antibody against the A/Goose/Guangdong/1/96 Hemagglutinin; J. Virol. 87 (23), 12619-12635 (2013), NCBI Accession #4MHH_H (222aa)	4626
INFL490	Immunoglobulin heavy chain variable region, partial	T2-6A	Huang, K.-Y.A., et al., Focused antibody response to influenza linked to antigenic drift; J. Clin. Invest. (2015), NCBI Accession #AKF02484.1 (124aa)	4627
INFL491	Kappa light chain constant region, human		US8992929 SEQ ID NO. 24	4628
INFL492	Kappa light chain variable	8D4	NCBI Accession #AFI57037	4629

INFL493	Kappa light chain variable	5B6	NCBI Accession #AFI57041	4630
INFL494	Kappa light chain variable region	8i10	US8858948 SEQ ID NO: 56	4631
INFL495	Kappa light chain variable region	23K12	US8858948 SEQ ID NO: 91	4632
INFL496	Kappa light chain variable region	4K8	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AEO16800	4633
INFL497	Kappa light chain variable region	6D9	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AEO16802	4634
INFL498	Kappa light chain variable region	8G9	US8603467 SEQ ID NO: 4	4635
INFL499	Kappa light chain variable region	13D4	US8603467 SEQ ID NO: 8	4636
INFL500	Kappa light chain variable region	20A11	US8603467 SEQ ID NO: 12	4637
INFL501	Kappa light chain variable region	EM4C04	US20120282273 SEQ ID NO: 1	4638
INFL502	Kappa, light chain	Fab H5m9	Zhu, X., et al., A Unique and Conserved Neutralization Epitope in H5N1 Influenza Viruses Identified by an Antibody against the A/Goose/Guangdong/1/96 Hemagglutinin; J. Virol. 87 (23), 12619-12635 (2013), NCBI Accession # 4MHH_L (218aa)	4639
INFL503	Lambda light chain	7A13	Krause et al. "Human Monoclonal Antibodies to Pandemic 1957 H2N2 and Pandemic 1968 H3N2 Influenza Viruses" J. Virol. 86 (11), 6334-6340 (2012), NCBI Accession #AFH78448	4640
INFL504	Lambda light chain variable	2K11	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AEO16794	4641
INFL505	Lambda light chain variable	2O10	Krause, J.C. et al. "Epitope-specific human influenza antibody repertoires diversify by B cell intraclonal sequence divergence and interclonal convergence" J. Immunol. 187 (7), 3704-3711 (2011), NCBI Accession #AEO16796	4642
INFL506	Lambda light chain variable region, partial	Monoclonal antibody clone 5E4	Yasugi, M. et al., Emerging Antigenic Variants at the Antigenic Site Sb in Pandemic A(H1N1)2009 Influenza Virus in Japan Detected by a Human Monoclonal Antibody; PLoS ONE 8 (10), E77892 (2013), NCBI Accession #BAM76755.1 (126aa)	4643
INFL507	Lambda light chain variable region,	T2-6A	Huang, K.-Y.A., et al., Focused antibody response to influenza linked to antigenic	4644

	partial, Immunoglobulin		drift; J. Clin. Invest. (2015), NCBI Accession #AKF02488. 1 (113aa)	
INFL508	Light chain	CR6261, Diridavumab, CR-6261	WO2008028946	4645
INFL509	Light chain	Firivumab, CT-P22	US20130004505	4646
INFL510	Light chain	Navivumab, CT149	WO20 13048153, US20140234336 SEQ ID NO: 39	4647
INFL51 1	Light chain	AT10—004	US20150010566, WO2013081463 SEQ ID NO: 36	4648
INFL512	Light chain	AT10—003	US20150010566, WO2013081463 SEQ ID NO: 37	4649
INFL513	Light chain	AT10—002	US20150010566, WO2013081463 SEQ ID NO: 38	4650
INFL514	Light chain	AT10—001	US20150010566, WO2013081463 SEQ ID NO: 39	4651
INFL515	Light chain	AT10—005	US20150010566, WO2013081463 SEQ ID NO: 40	4652
INFL516	Light chain	CT104	WO201 1111966, US20130004505 SEQ ID NO: 36	4653
INFL517	Light chain	CT120	WO201 1111966, US20130004505 SEQ ID NO: 40	4654
INFL518	Light chain	CT123	WO201 1111966, US20130004505 SEQ ID NO: 44	4655
INFL519	Light chain	2A	US2014001 1982 SEQ ID NO: 4	4656
INFL520	Light chain	F005-S26	WO2014049520, US20140086927 SEQ ID NO: 13	4657
INFL521	Light chain	BFl-1	WO2008 156763 SEQ ID NO: 8	4658
INFL522	Light chain	BF1-19	WO2008 156763 SEQ ID NO: 12	4659
INFL523	Light chain	BFl-10	WO2008 156763 SEQ ID NO: 10	4660
INFL524	Light chain	2D1	WO2010127252, US8894997 SEQ ID NO: 8	4661
INFL525	Light chain	1F1	WO20 10 127252, US8894997 SEQ ID NO: 2	4662
INFL526	Light chain	4D20	WO2010127252, US8894997 SEQ ID NO: 10	4663
INFL527	Light chain	A18	W013 170139 SEQ ID NO: 95	4664
INFL528	Light chain	Ab A18	US7788200 SEQ ID NO: 28	4665
INFL529	Light chain	Ab 067	US7788200 SEQ ID NO: 153	4666
INFL530	Light chain	Ab 068	US7788200 SEQ ID NO: 154	4667
INFL53 1	Light chain	Ab 069, Ab 079	US7788200 SEQ ID NO: 155	4668
INFL532	Light chain	Ab 070	US7788200 SEQ ID NO: 156	4669
INFL533	Light chain	Ab 073	US7788200 SEQ ID NO: 165	4670
INFL534	Light chain	Ab 074, Ab 080	US7788200 SEQ ID NO: 166	4671
INFL535	Light chain	Ab 075	US7788200 SEQ ID NO: 167	4672
INFL536	Light chain	Ab 076	US7788200 SEQ ID NO: 168	4673
INFL537	Light chain	Ab 077, Ab 081	US7788200 SEQ ID NO: 169	4674
INFL538	Light chain	Ab 014, Ab 154, Ab 157	US7788200 SEQ ID NO: 29	4675
INFL539	Light chain	Ab 028, Ab 155	US7788200 SEQ ID NO: 30	4676
INFL540	Light chain	Ab 001, Ab 002, Ab 003	US7788200 SEQ ID NO: 31	4677
INFL541	Light chain	Ab 009, Ab 010, Ab 011	US7788200 SEQ ID NO: 32	4678

INFL542	Light chain	Ab 017, Ab B18, Ab B18, Ab 019, Ab 019	US7788200 SEQ ID NO: 33	4679
INFL543	Light chain	Ab 025, Ab 026, Ab 027, Ab 028	US7788200 SEQ ID NO: 34	4680
INFL544	Light chain	Ab 159	US7788200 SEQ ID NO: 35	4681
INFL545	Light chain	Ab 160	US7788200 SEQ ID NO: 36	4682
INFL546	Light chain	Ab 186, Ab 194	US7788200 SEQ ID NO: 37	4683
INFL547	Light chain	Ab 187, Ab 195	US7788200 SEQ ID NO: 38	4684
INFL548	Light chain	Ab 188, Ab 196	US7788200 SEQ ID NO: 39	4685
INFL549	Light chain	Ab 189, Ab 197	US7788200 SEQ ID NO: 40	4686
INFL550	Light chain	Ab 190, Ab 198	US7788200 SEQ ID NO: 41	4687
INFL551	Light chain	Ab 191, Ab 199	US7788200 SEQ ID NO: 42	4688
INFL552	Light chain	Ab 192, Ab 200	US7788200 SEQ ID NO: 43	4689
INFL553	Light chain	Ab 193	US7788200 SEQ ID NO: 44	4690
INFL554	Light chain	Ab 202, Ab 203, Ab 204, Ab 210, Ab 031, Ab 032, Ab 033, Ab 034	US7788200 SEQ ID NO: 45	4691
INFL555	Light chain	Ab 211, Ab 212, Ab 213, Ab 219, Ab 037, Ab 038, Ab 039, Ab 040	US7788200 SEQ ID NO: 46	4692
INFL556	Light chain	Ab A001, Ab 004, Ab 007, Ab 016	US7788200 SEQ ID NO: 47	4693
INFL557	Light chain	Ab A002, Ab 005, Ab 008, Ab A017	US7788200 SEQ ID NO: 48	4694
INFL558	Light chain	Ab A003, Ab 006, Ab A009, Ab C18	US7788200 SEQ ID NO: 49	4695
INFL559	Light chain	Ab A010, Ab 012, Ab A14, Ab A019	US7788200 SEQ ID NO: 50	4696
INFL560	Light chain	Ab A011, Ab 013m Ab 0135	US7788200 SEQ ID NO: 51	4697
INFL561	Light chain	Ab 044, Ab 071, Ab 072, Ab 078	US7788200 SEQ ID NO: 52	4698
INFL562	Light chain	Ab 051	US7788200 SEQ ID NO: 53	4699
INFL563	Light chain	Ab 049	US7788200 SEQ ID NO: 54	4700
INFL564	Light chain	Ab 047	US7788200 SEQ ID NO: 55	4701
INFL565	Light chain	Ab 050	US7788200 SEQ ID NO: 56	4702
INFL566	Light chain	Ab 045	US7788200 SEQ ID NO: 57	4703
INFL567	Light chain	Ab 048	US7788200 SEQ ID NO: 58	4704
INFL568	Light chain	Ab 046	US7788200 SEQ ID NO: 59	4705
INFL569	Light chain	Ab 043	US7788200 SEQ ID NO: 60	4706
INFL570	Light chain	Ab 052	US7788200 SEQ ID NO: 61	4707
INFL571	Light chain	F005-126	WO2014049520 SEQ ID 13	4708
INFL572	Light chain	8f24	WO2012045001 SEQ ID 3	4709
INFL573	Light chain	3E22	WO2012045001 SEQ ID 7	4710
INFL574	Light chain	5117	WO2012045001 SEQ ID 11	4711
INFL575	Light chain		WO2012045001 SEQ ID 15	4712
INFL576	Light chain		WO2012045001 SEQ ID 31	4713
INFL577	Light chain		WO2012045001 SEQ ID 35	4714

INFL578	Light chain		WO20 12045001 SEQ ID 19	4715
INFL579	Light chain	10A14	WO2012045001 SEQ ID 23	4716
INFL580	Light chain	8D4	WO20 12045001 SEQ ID 27	4717
INFL581	Light chain	2B9	US91 15201 SEQ ID NO: 5	4718
INFL582	Light chain	mAB 7A7	US20150239960, US20 140170163, US86733 14, US20 110027270, WO2010138564 SEQ ID NO: 7	4719
INFL583	Light chain	mAB 12D1	US20150239960, US20 140170163, US86733 14, US20 110027270, WO2010138564 SEQ ID NO: 13	4720
INFL584	Light chain	mAB 66A6	US20150239960, US20 140170163, US86733 14, US201 10027270, WO20 10138564 SEQ ID NO: 17	4721
INFL585	Light chain	M1 D12	US201 10033473, WO2009125395 SEQ ID NO: 15	4722
INFL586	Light chain	mAB 1.12	WO2013030165 SEQ ID NO: 2	4723
INFL587	Light chain	mAB3. 1	WO20 13030165 SEQ ID NO: 4	4724
INFL588	Light chain	5A7	WO20 15120097 SEQ ID NO: 8	4725
INFL589	Light chain	TRL053	WO20 15120097 SEQ ID NO: 18	4726
1NFL590	Light chain	TRL579	WO2015120097 SEQ ID NO: 28	4727
INFL591	Light chain	TRL784	WO2015120097 SEQ ID NO: 38	4728
INFL592	Light chain	TRL794	WO20 15120097 SEQ ID NO: 48	4729
INFL593	Light chain	TRL798	WO2015120097 SEQ ID NO: 58	4730
INFL594	Light chain	TRL799	WO20 15120097 SEQ ID NO: 68	4731
1NFL595	Light chain	TRL809	WO2015120097 SEQ ID NO: 78	4732
INFL596	Light chain	TRL81 1	WO2015120097 SEQ ID NO: 88	4733
INFL597	Light chain	TRL812	WO2015120097 SEQ ID NO: 98	4734
INFL598	Light chain	TRL8 13	WO2015120097 SEQ ID NO: 108	4735
INFL599	Light chain	TRL823	WO20 15120097 SEQ ID NO: 118	4736
1NFL600	Light chain	TRL832	WO2015120097 SEQ ID NO: 128	4737
INFL601	Light chain	TRL833	WO20 15120097 SEQ ID NO: 138	4738
INFL602	Light chain	TRL834	WO2015120097 SEQ ID NO: 148	4739
INFL603	Light chain	TRL837	WO2015120097 SEQ ID NO: 167	4740
INFL604	Light chain	TRL839	WO20 15120097 SEQ ID NO: 177	4741
INFL605	Light chain	TRL841	WO2015120097 SEQ ID NO: 187	4742
INFL606	Light chain	TRL842	WO20 15120097 SEQ ID NO: 197	4743
INFL607	Light chain	TRL845	WO2015120097 SEQ ID NO: 207	4744
INFL608	Light chain	TRL846	WO2015120097 SEQ ID NO: 218	4745
INFL609	Light chain	TRL847	WO20 15120097 SEQ ID NO: 228	4746
INFL610	Light chain	TRL848	WO2015120097 SEQ ID NO: 238	4747
INFL611	Light chain	TRL849	WO20 15120097 SEQ ID NO: 248	4748
INFL612	Light chain	TRL851	WO2015120097 SEQ ID NO: 258	4749
INFL613	Light chain	TRL854	WO2015120097 SEQ ID NO: 268	4750
INFL614	Light chain	TRL856	WO20 15120097 SEQ ID NO: 278	4751
INFL615	Light chain	TRL858	WO2015120097 SEQ ID NO: 288	4752
INFL616	Light chain	humM2e-hBiTE-l	WO20 14140368 SEQ ID NO: 10	4753
INFL617	Light chain	humM2e-hBiTE-2	WO2014140368 SEQ ID NO: 18	4754

INFL618	Light chain	humM2e-hBiTE-3	WO2014140368 SEQ ID NO: 26	4755
INFL619	Light chain	humM2e-hBiTE-4	WO2014140368 SEQ ID NO: 34	4756
INFL620	Light chain	VH of humM2e-hBiTE-5	WO2014140368 SEQ ID NO: 42	4757
INFL621	Light chain	humM2e-hBiTE-6	WO2014140368 SEQ ID NO: 50	4758
INFL622	Light chain	humM2e-hBiTE-7	WO2014140368 SEQ ID NO: 58	4759
INFL623	Light chain	humM2e-hBiTE-8	WO2014140368 SEQ ID NO: 66	4760
INFL624	Light chain	humM2e-hBiTE-9	WO2014140368 SEQ ID NO: 74	4761
INFL625	Light chain	murM2e-hBiTE	WO2014140368 SEQ ID NO: 82	4762
INFL626	Light chain	FLA5.10	US8124092 SEQ ID NO: 3	4763
INFL627	Light chain	FLD21.140	US8124092 SEQ ID NO: 7	4764
INFL628	Light chain	FLA3.14	US8124092 SEQ ID NO: 11	4765
INFL629	Light chain	FLD20.19	US8124092 SEQ ID NO: 15	4766
INFL630	Light chain	FLD84	US8124092 SEQ ID NO: 44	4767
INFL631	Light chain	FLD93	US8124092 SEQ ID NO: 54	4768
INFL632	Light chain	FLD122	US8124092 SEQ ID NO: 64	4769
INFL633	Light chain	FLD127	US8124092 SEQ ID NO: 74	4770
INFL634	Light chain	FLD129	US8124092 SEQ ID NO: 84	4771
INFL635	Light chain	FLD132	US8124092 SEQ ID NO: 94	4772
INFL636	Light chain	FLD194	US8124092 SEQ ID NO: 104	4773
INFL637	Light chain	mAb1	WO2015112994 SEQ ID NO: 77	4774
INFL638	Light chain	mAb2	WO2015112994 SEQ ID NO: 81	4775
INFL639	Light chain	mAb3	WO2015112994 SEQ ID NO: 85	4776
INFL640	Light chain	CT-P22	US20130004505 SEQ ID NO: 40; WO 2011/111966	4777
INFL641	Light chain	C05	Ekiert, D.C., et al., Cross-neutralization of influenza A viruses mediated by a single antibody loop; Nature 489 (7417), 526-532 (2012), NCBI Accession #4FNL_L (214aa)	4778
INFL642	Light chain	CR8020	Ekiert, D.C., et al., A highly conserved neutralizing epitope on group 2 influenza A viruses; Science 333 (6044), 843-850 (2011); WO2010130636, NCBI Accession #3SDY_L	4779
INFL643	Light chain	CR8033	Dreyfus, C., Laursen, N.S. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession # 4FQL_L	4780
INFL644	Light chain	CR8043	Friesen, R.H. et al., A common solution to group 2 influenza virus neutralization; Proc. Natl. Acad. Sci. U.S.A. 111 (1), 445-450 (2014), NCBI Accession #4NM8_L	4781
INFL645	Light chain	CR8059	Dreyfus, C. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession #4FQK_L	4782
INFL646	Light chain	CR8071	Dreyfus, C. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348	4783

			(2012), NCBI Accession # 4FQJ_L (216aa)	
INFL647	Light chain	CR9114	WO2013079473; WO2014191435; Dreyfus, C., Laursen, N.S. et al., Highly conserved protective epitopes on influenza B viruses; Science 337 (6100), 1343-1348 (2012), NCBI Accession #4FQY_L(216aa)	4784
INFL648	Light chain	Ch67	Schmidt, A.G., et al., Preconfiguration of the antigen-binding site during affinity maturation of a broadly neutralizing influenza virus antibody; Proc. Natl. Acad. Sci. U.S.A. 110 (1), 264-269 (2013), NCBI Accession #4HKX_B (214aa)	4785
INFL649	Light chain	Fab 26/9	Schulze-Gahmen, U. et al., J. Biol. Chem. 263 (32), 17100-17105 (1988); Churchill, M.E., et al., J. Mol. Biol. 241 (4), 534-556 (1994), NCBI Accession #1FRG_L	4786
INFL650	Light chain	Fab 3.1	Wyrzucki, A. et al., Alternative Recognition of the Conserved Stem Epitope in Influenza A Virus Hemagglutinin by a VH3-30-Encoded Heterosubtypic Antibody; J. Virol. 88 (12), 7083-7092 (2014), NCBI Accession #4PY8_J	4787
INFL651	Light chain	Fab 2g1	Xu, R. et al., A recurring motif for antibody recognition of the receptor-binding site of influenza hemagglutinin; Nat. Struct. Mol. Biol. 20 (3), 363-370 (2013), NCBI Accession #4HG4_M (214aa)	4788
INFL652	Light chain	Fab 8m2	Xu, R. et al., A recurring motif for antibody recognition of the receptor-binding site of influenza hemagglutinin; Nat. Struct. Mol. Biol. 20 (3), 363-370 (2013), NCBI Accession #4HFU_L (215aa)	4789
INFL653	Light chain	Fab 8f8	Xu, R. et al., A recurring motif for antibody recognition of the receptor-binding site of influenza hemagglutinin; Nat. Struct. Mol. Biol. 20 (3), 363-370 (2013), NCBI Accession # 4HF5_L (218aa)	4790
INFL654	Light chain	Fab 2d1	Xu, R., et al., Structural basis of preexisting immunity to the 2009 H1N1 pandemic influenza virus; Science 328 (5976), 357-360 (2010), NCBI Accession #3LZF_L (217aa)	4791
INFL655	Light chain	Fi6v3	Corti, D. et al., A neutralizing antibody selected from plasma cells that binds to group 1 and group 2 influenza A hemagglutinins; Science 333 (6044), 850-856 (2011), NCBI Accession #3ZTN_L	4792
INFL656	Light chain	Light chain from 3WHE_N	Iba, Y., et al., Conserved Neutralizing Epitope at Globular Head of Hemagglutinin in H3N2 Influenza	4793

			Viruses; J. **Virol.** (2014), **NCBI** Accession **#3WHE_N (220aa)**	
INFL657	Light chain (partial)	monoclonal **antibody** PN-SIA28	Burioni, R. et **al.,** Monoclonal antibodies isolated **from** human **B** cells **neutralize** a broad range **of H1** subtype influenza A virases including swine-origin Influenza **vinis(S-OTV);** Virology (2010), NCBI Accession **#ACX30939. 1 (105aa)**	4794
INFL658	Light chain (partial)	monoclonal antibody PN-SIA49	Burioni, R. et al, **Monoclonal** antibodies isolated **from** human **B** cells **neutralize** a broad range **of H1** subtype influenza **A** virases including swine-origin Influenza **vinis(S-OTV);** Virology (2010), NCBI Accession **#ACX30938. 1 (105aa)**	4795
INFL659	Light **chain;** Fab	5j8	Hong, **M.** et al, Antibody Recognition of the **Pandemic H1N1** influenza **Virus** Hemagglutinin Receptor Binding Site; J. Virol 87 (22), **12471- 12480 (2013),** NCBI **Accession #4M5Z_L**	4796
INFL660	Light chain **CDR 1**	**Ab1A2**	WO2015028478 SEQ ID 9	4797
INFL661	Light **chain** CDR 2	Ab1A2	**WO2015028478** SEQ ID 10	4798
INFL662	Light chain CDR **3**	Ab1A2	**WO2015028478** SEQ ID **11**	4799
INFL663	Light chain Fab	CT147	WO2013048153, **US20140234336** SEQ ID NO: 37	4800
INFL664	Light chain Fab	**CT164**	**WO2013048153,** US20140234336 SEQ ID NO: **41**	4801
INFL665	Light chain Fab	CT166	WO2013048153, **US20140234336** SEQ ID NO: 43	4802
INFL666	Light **chain** Human igg	Fab 39.29	**Nakamura,** G. et **al.,** An in vivo **human-piasmablast** enrichment technique allows rapid identification **of** therapeutic influenza **a** antibodies; Cell Host Microbe **14** (1), 93-103 (2013), NCBI **Accession #4KVN_L** (215aa)	4803
INFL667	Light chain K3	**h2B1**	**US91 15201** SEQ ID NO: 9	4804
INFL668	Light chain K3	**h2B2**	**US91 15201** SEQ ID NO: 10	4805
INFL669	Light **chain partial** region	**4A10**	**Krause,** J.C. et **al. "Epitope-specific human** influenza antibody repertoires **diversify** by B cell **intraclonal** sequence divergence and **interclonal** convergence" J. Immunol 187 (7), **3704-3711** (2011), NCBI Accession #AE016798	4806
1NFL670	Light chain variable **(exemplary)**	**LC-VD** from **US20 13030234**	US201.30.30234 SEQ **ID** NO: .33	4807
INFL671	Light chain **variable (exemplary)**	**LC-VD** from US20 13030234	**US2013030234** SEQ **ID NO:** 34	4808
1NFL672	Light chain variable (exemplary)	**LC-VD** from **US20 13030234**	US201.30.30234 SEQ **ID** NO: .35	4809
INFL673	Light chain variable **(exemplary)**	**LC-VD** from US20 13030234	**US2013030234** SEQ **ID** NO: 36	4810
1NFL674	Light chain variable **(exemplary)**	**LC-VD** from **US20 13030234**	US201.30.30234 SEQ **ID** NO: .37	4811
INFL675	Light **chain variable** (exemplary)	**LC-VD** from US20 13030234	**US2013030234** SEQ **ID** NO: 38	4812
1NFL676	Light chain variable **(exemplary)**	**LC-VD** from **US20 13030234**	US201.30.30234 SEQ **ID** NO: .39	4813
INFL677	Light **chain variable** (exemplary)	**LC-VD** from US20 13030234	**US2013030234** SEQ **ID** NO: 40	4814

INFL678	Light chain variable (exemplary)	LC-VD from US20 13030234	US2013030234 SEQ ID NO: 41	4815
INFL679	Light chain variable (exemplary)	LC-VD from US20 13030234	US2013030234 SEQ ID NO: 42	4816
INFL680	Light chain variable (exemplary)	LC-VD from US20 13030234	US20 13030234 SEQ ID NO: 43	4817
INFL681	Light chain variable reedon	39.18 B11	US20140 16 1822 SEQ ID NO: 156	4818
INFL682	Light chain variable region	GG3	WO20 141 59960 SEQ ID NO: 25	4819
INFL683	Light chain variable reedon	N547	US8003 106 SEQ ID NO: 29	4820
INFL684	Light chain variable region	L66	US8003 106 SEQ ID NO: 31	4821
INFL685	Light chain variable reedon	C40	US8003 106 SEQ ID NO: 27	4822
INFL686	Light chain variable region	14C2	US8080244 SEQ ID NO: 7	4823
INFL687	Light chain variable region	h14C2	US8080244 SEQ ID NO: 1	4824
INFL688	Light chain variable region	VN04-2-HuGl	US20 100150941 SEQ ID NO: 6	4825
1NFL689	Light chain variable region	VN04-3-HuGl	US20100 I50941 SEQ ID NO: 8	4826
INFL690	Light chain variable region	F16 variant 1, FE6 variant 2	US8871207 SEQ ID NO: 14	4827
1NFL691	Light chain variable region	FI6 variant 3, FI6 variant 4	US8871207 SEQ ID NO: 57	4828
INFL692	Light chain variable region	F16 variant 5	US8871207 SEQ ID NO: 61	4829
1NFL693	Light chain variable region	FI28 variant 1, FI28 variant 2	US8871207 SEQ ID NO: 30	4830
INFL694	Light chain variable region	21B15	US8858948 SEQ ID NO: 46	4831
1NFL695	Light chain variable region	3241_G23	US8858948 SEQ ID NO: 118	4832
INFL696	Light chain variable region	3244_110	US8858948 SEQ ID NO: 122	4833
INFL697	Light chain variable region	3243_J07	US8858948 SEQ ID NO: 126	4834
INFL698	Light chain variable region	3259_J21	US8858948 SEQ ID NO: 130	4835
INFL699	Light chain variable region	3245_019	US8858948 SEQ ID NO: 134	4836
INFL700	Light chain variable region	3244_H04	US8858948 SEQ ID NO: 138	4837
INFL701	Light chain variable region	3136_G05	US8858948 SEQ ID NO: 142	4838
INFL702	Light chain variable region	3252_C13	US8858948 SEQ ID NO: 146	4839
INFL703	Light chain variable region	3255_J06	US8858948 SEQ ID NO: 150	4840
INFL704	Light chain variable region	3420J23	US8858948 SEQ ID NO: 154	4841
INFL705	Light chain variable region	3248_P18	US8858948 SEQ ID NO: 160	4842
INFL706	Light chain variable reedon	3253_P10	US8858948 SEQ ID NO: 164	4843

INFL707	Light chain variable region	3260_D19	**US8858948** SEQ ID NO: **168**	4844
INFL708	Light chain variable reedon	**3362_B 11**	**US8858948** SEQ ID NO: 174	4845
INFL709	Light chain variable region	**3242JP05**	US8858948 SEQ ID **NO:** 178	4846
INFL7 I0	Light chain variable reedon	A66	**WO2009079259, US201** 10038935, **US201400** 1982 SEQ ID **NO:** 34	4847
INFL71	Light chain variable region	D7, **H98**	**WO2009079259, US20** 10038935, **US2014001 1982** SEQ ID **NO:** 8	4848
INFL7 I2	Light chain variable reedon	**D8**	**WO2009079259, US201** 10038935, **US201400** 1982 SEQ ID **NO:** 14	4849
INFL713	Light chain variable region	D80	**WO2009079259, US20** 10038935, **US2014001 1982** SEQ ID **NO:** 16	4850
INFL714	Light chain variable reedon	**E88**	**WO2009079259, US201** 10038935, **US201400 11982** SEQ ID **NO:** 38	4851
INFL715	Light chain variable region	E90	**WO2009079259, US20** 10038935, **US2014001 1982** SEQ ID **NO: 22**	4852
INFL7 I6	Light chain variable reedon	F!O	**WO2009079259, US201** 10038935, **US201400** 1982 SEQ ID **NO: 20**	4853
INFL717	Light chain variable region	**G17**	**WO2009079259, US20** 10038935, **US2 0 1400 11982** SEQ ID **NO:** 26	4854
INFL718	Light chain variable region	**H40**	WO2009079259, **U**S201 10038935, **US2014001 1982** SEQ ID NO: 30	4855
INFL719	Light chain variable region	**H98**	**WO2009079259,** US201 10038935, **US2014001 1982** SEQ ID **NO: 10**	4856
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INFL722	Light chain variable region	**CH67**	WO20 13020074, US20 140302043 SEQ ID NO: 12	4859
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INFL724	Light chain variable region	Antibody **1**	**WO2015051010** SEQ ID **NO:** 7	4861
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INFL727	Light **chain** variable region	Antibody 4	**WO2015051010** SEQ ID **NO:** 37	4864
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INFL734	Light chain variable region	Antibody 11	**WO2015051010** SEQ ID NO: **107**	4871
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INFL737	Light chain variable region	Antibody 14	WO2015051010 SEQ ID NO: 137	4874
INFL738	Light chain variable region	Antibody 15	WO20 1505 1010 SEQ ID NO: 147	4875
INFL739	Light chain variable region	Antibody 3-GL	WO2015051010 SEQ ID NO: 157	4876
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INFL822	Light **chain** variable region	CR6310	US200903 11265, US8691223, US9109017, WO2008028946 SEQ ID **NO:** 99	4959
INFL823	Light chain variable region	**CR6314**	US200903 11265, US8691223, **US9109017, WO2008028946** SEQ ID NO: **101**	4960
INFL824	Light **chain variable** region	CR6323	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: 103	4961
1NFL825	Light chain variable region	CR6325	US200903 11265, US8691223, **US9 109017, WO2008028946** SEQ ID NO: 105	4962
INFL826	Light chain variable region	CR6331	**US200903 11265, US8691223, US9** 09017, **WO2008028946** SEQ ID NO: 107	4963
INFL827	Light chain variable region	CR6344	**US200903** 11265, **US8691**223, US91090 17, **WO2008028946** SEQ ID NO: 109	4964
INFL828	Light chain vanable region	**CR6141**	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: 319	4965
INFL829	Light **chain** variable region	**CR6272**	**US200903 11265**, US8691223, **US9109017,** WO2008028946 SEQ ID **NO:** 323	4966
INFL830	Light **chain variable** region	**CR6296**	US200903 11265, **US8691223, US9109017,** WO2008028946 SEQ ID NO: 327	4967
INTL.831	Light drain variable region	CR6301	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: 331	4968
INFL832	Light chain variable region	CR6327	**US200903 11265, US8691223,** US91090 17, **WO2008028946** SEQ ID NO: 335	4969
INFL833	Light chain variable region	CR6328	**US200903** 11265, **US8691223, US9** 09017, **WO2008028946** SEQ ID NO: 339	4970
INFL834	Light chain variable region	CR6329	**US200903** 11265, **US8691**223, US91090 17, **WO2008028946** SEQ ID NO: 343	4971
INFL835	Light chain vanable region	CR6332	**US200903** 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: 347	4972
INFL836	Light **chain** variable region	**CR6334**	**US200903 11265**, US8691223, **US9109017,** WO2008028946 SEQ ID NO: 351	4973
INFL837	Light **chain variable** region	**CR6336**	US200903 11265, **US8691223, US9109017, WO2008028946** SEQ ID NO: 355	4974
1NFL838	Light chain variable region	CR6339	US200903 11265, US8691223, **US9 109017, WO2008028946** SEQ ID NO: 359	4975
INFL839	Light chain variable region	CR6342	**US200903 11265, US8691223,** US91090 17, **WO2008028946** SEQ ID NO: 363	4976

INFL840	Light chain variable region	CR6343	US200903 I 1265, US8691223, US9 1090 I7, WO2008028946 SEQ ID NO: 367	4977
INFL84 1	Light chain vanable region	CR9003	US20 1401201 13 SEQ ID NO: 4	4978
1NFL842	Light chain variable region	CR9004	US20140 I20 I 13 SEQ ID NO: 8	4979
INFL843	Light chain vanable region	CR9005	US20140120H3 SEQ ID NO: 12	4980
1NFL844	Light chain variable region	CR9006	US20140 I20 I 13 SEQ ID NO: 16	4981
INFL845	Light chain vanable region	CR9007	US201401201 13 SEQ ID NO: 20	4982
1NFL846	Light chain variable region	CR9008	US20140 I20 I 13 SEQ ID NO: 24	4983
INFL847	Light chain vanable region	CR9009	US201401201 13 SEQ ID NO: 28	4984
1NFL848	Light chain variable region	CR9010	US20140 I20 I 13 SEQ ID NO: 32	4985
INFL849	Light chain vanable region	CR901 1	US20140120H3 SEQ ID NO: 36	4986
INFL850	Light chain variable region	CR9012	US201401201 13 SEQ ID NO: 40	4987
INFL85 1	Light chain variable region	CR9029	US201401201 13 SEQ ID NO: 44	4988
INFL852	Light chain variable region	CR9030	US201401201 13 SEQ ID NO: 48	4989
INFL853	Light chain variable region	CR903 1	US201401201 13 SEQ ID NO: 52	4990
INFL854	Light chain variable region	CR91 12	US201401201 13 SEQ ID NO: 56	4991
INFL855	Light chain variable region	CR91 13	US201401201 13 SEQ ID NO: 60	4992
INFL856	Light chain variable region	CR91 14	US201401201 13 SEQ ID NO: 64	4993
INFL857	Light chain variable region	CR8033	US8852595 SEQ ID NO: 73	4994
INFL858	Light chain variable region	CR8059	US8852595 SEQ ID NO: 77	4995
INFL859	Light chain variable region	CR8071	US8852595 SEQ ID NO: 79	4996
INFL860	Light chain variable region	CR1005 1	US8852595 SEQ ID NO: 83	4997
INFL861	Light chain variable region	CR 10049	US8852595 SEQ ID NO: 87	4998
INFL862	Light chain variable region	CR10023	US8852595 SEQ ID NO: 9 1	4999
INFL863	Light chain variable region	CR10032	US8852595 SEQ ID NO: 95	5000
INFL864	Light chain variable region	CR1 1035	US8852595 SEQ ID NO: 103	5001
INFL865	Light chain variable region	CR 11036	US8852595 SEQ ID NO: 107	5002
INFL866	Light chain variable region	CR1 1038	US8852595 SEQ ID NO: 111	5003
INFL867	Light chain variable region	CR 11039	US8852595 SEQ ID NO: 115	5004
INFL868	Light chain variable region	CR803 1	US8852595 SEQ ID NO: 12 1	5005

INFL869	Light chain variable region	CR8032	US8852595 SEQ ID NO: 125	5006
INFL870	Light chain variable reedon	CR8034	US8852595 SEQ ID NO: 129	5007
INFL87 i	Light chain variable region		US8992929 SEQ ID NO: 2	5008
INFL872	Light chain variable reedon	M2e	US8420794 SEQ ID NO: 1	5009
INFL873	Light chain variable region		US87 !5743, US20 !40275492 SEQ ID NO: 16	5010
INTL.874	Light chain variable reedon		US8715743, US20140275492 SEQ ID NO: !9	501 !
INFL875	Light chain variable region		US87 !5743, US20 !40275492 SEQ ID NO: 32	5012
INTL.876	Light chain variable region	anti-!918 influenza HA Ig	Yu, X., et al.. Neutralizing antibodies derived from the B cells of 1918 influenza pandemic survivors; Nature 455 (72 12), 532-536 (2008), NCBI Accession #ACI04582. 1 (12 laa)	5013
INFL877	Light chain variable region	anti-!918 influenza HA Ig	Yu, X., et al, Neutralizing antibodies derived from the B cells of 1918 influenza pandemic survivors; Nature 455 (72 !2), 532-536 (2008), NCBI Accession #ACI04580. ! (1 18aa)	5014
INFL878	Light chain variable region	4D20	Yu, X. et al "Neutralizing antibodies derived from the B cells of 1918 influenza pandemic survivors", Nature 455 (7212), 532-536, NCBI Accession #ACI04580	5015
INFL879	Light chain variable region	2B12	Yu, X. et al "Neutralizing antibodies derived from the B cells of 1918 influenza pandemic survivors", Nature 455 (72 12), 532-536, NCBI Accession #ABY48869	5016
INFL880	Light chain variable region	TCN-535 (5246_P I9)	US20150086555 SEQ ID NO: 180	5017
INFL881	Light chain variable region	TCN-536 (5095_N01)	US20150086555 SEQ ID NO: 191	5018
INFL882	Light chain variable region	TCN-537 (3 194_D21)	US20150086555 SEQ ID NO: 202	5019
INFL883	Light chain variable region	TCN-538 (3206_017)	US20150086555 SEQ ID NO: 214	5020
INFL884	Light chain variable region	TCN-539 (5056_AOS)	US20150086555 SEQ ID NO: 226	5021
INFL885	Light chain variable region	TCN-540 (5060_F05)	US20150086555 SEQ ID NO: 238	5022
INFL886	Light chain variable region	TCN-541 (5062_M11)	US20150086555 SEQ ID NO: 250	5023
INFL887	Light chain variable region	TCN-542 (5079_A16)	US20150086555 SEQ ID NO: 262	5024
INFL888	Light chain variable region	TCN-543 (5081_G23)	US20150086555 SEQ ID NO: 274	5025
INFL889	Light chain variable region	TCN-544 (5082_A19)	US20150086555 SEQ ID NO: 286	5026
INFL890	Light chain variable region	TCN-545 (5082_115)	US20150086555 SEQ ID NO: 298	5027
INFL89 !	Light chain variable region	TCN-546 (5089_LOS)	US20150086555 SEQ ID NO: 309	5028

INFL892	Light chain variable region	TCN-547 (5092_F11)	US20150086555 SEQ ID NO: 320	5029
INFL893	Light chain variable region	TCN-548 (5092_P01)	US20150086555 SEQ ID NO: 331	5030
INFL894	Light chain variable region	TCN-549 (5092_P04)	US20150086555 SEQ ID NO: 342	5031
INFL895	Light chain variable region	TCN-550 (5096_F06)	US20150086555 SEQ ID NO: 353	5032
INFL896	Light chain variable region	TCN-551 (5243_D01)	US20150086555 SEQ ID NO: 365	5033
INFL897	Light chain variable region	TCN-552 (5249_I23)	US20150086555 SEQ ID NO: 377	5034
INFL898	Light chain variable region	TCN-553 (5261_C18)	US20150086555 SEQ ID NO: 389	5035
INFL899	Light chain variable region	TCN-554 (5277_M05)	US20150086555 SEQ ID NO: 399	5036
INFL900	Light chain variable region	TCN-555 (5246_L16)	US20150086555 SEQ ID NO: 405	5037
INFL901	Light chain variable region	TCN-556 (5089_K12)	US20150086555 SEQ ID NO: 415	5038
INFL902	Light chain variable region	TCN-557 (5081_A04)	US20150086555 SEQ ID NO: 427	5039
INFL903	Light chain variable region	TCN-559 (5097_G08)	US20150086555 SEQ ID NO: 441	5040
INFL904	Light chain variable region	TCN-560 (5084_P10)	US20150086555 SEQ ID NO: 453	5041
INFL905	Light chain variable region	TCN-564 (5256_A17b)	US20150086555 SEQ ID NO: 519	5042
INFL906	Light chain variable region	CR8001	WO2010130636 SEQ ID NO: 4	5043
INFL907	Light chain variable region	CR8003	WO2010130636 SEQ ID NO: 8	5044
INFL908	Light chain variable region	CR8015	WO2010130636 SEQ ID NO: 12	5045
INFL909	Light chain variable region	CR8016	WO2010130636 SEQ ID NO: 16	5046
INFL910	Light chain variable region	CR8017	WO2010130636 SEQ ID NO: 20	5047
INFL911	Light chain variable region	TCN-522 (3212_I12)	US20150086555 SEQ ID NO: 40	5048
INFL912	Light chain variable region	TCN-521 (3280_D18)	US20150086555 SEQ ID NO: 28	5049
INFL913	Light chain variable region	TCN-523 (5248_A17), TCN-533 (5256_A17a), TCN-534 (5249_B02)	US20150086555 SEQ ID NO: 52	5050
INFL914	Light chain variable region	TCN-563 (5237_B21)	US20150086555 SEQ ID NO: 64	5051
INFL915	Light chain variable region	TCN-526 (5084_C17)	US20150086555 SEQ ID NO: 76	5052
INFL916	Light chain variable region	TCN-527 (5086_C06)	US20150086555 SEQ ID NO: 88	5053
INFL917	Light chain variable region	TCN-528 (5087_P17)	US20150086555 SEQ ID NO: 100	5054
INFL918	Light chain variable region	TCN-529 (5297_H01)	US20150086555 SEQ ID NO: 112	5055

INFL919	Light chain variable region	TCN-530 (5248_H10), TCN-558 (5248_H10b)	US20150086555 SEQ ID NO: 124	5056
INFL920	Light chain variable region	TCN-531 (5091_H13)	US20150086555 SEQ ID NO: 136	5057
INFL921	Light chain variable region	TCN-532 (5262_H18)	US20150086555 SEQ ID NO: 148	5058
INFL922	Light chain variable region	TCN-534 (5249_B02)	US20150086555 SEQ ID NO: 168	5059
INFL923	Light chain variable region	TCN-504 (3251_K17)	US20150086555 SEQ ID NO: 524	5060
INFL924	Light chain variable region	AB1	US20120093834, WO2009121004 SEQ ID NO: 71	5061
INFL925	Light chain variable region	AB2	US20120093834, WO2009121004 SEQ ID NO: 140	5062
INFL926	Light chain variable region	AB3	US20120093834, WO2009121004 SEQ ID NO: 81	5063
INFL927	Light chain variable region	AB4	US20120093834, WO2009121004 SEQ ID NO: 158	5064
INFL928	Light chain variable region	AB5	US20120093834, WO2009121004 SEQ ID NO: 159	5065
INFL929	Light chain variable region	AB6	US20120093834, WO2009121004 SEQ ID NO: 160	5066
INFL930	Light chain variable region	VN04-2	WO2008033105 SEQ ID NO: 6	5067
INFL931	Light chain variable region	VN04-3	WO2008033105 SEQ ID NO: 8	5068
INFL932	Light chain variable region	1286-C05	WO2010132604, US20120128671 SEQ ID NO: 3	5069
INFL933	Light chain variable region	1286-C05	WO2010132604, US20120128671 SEQ ID NO: 4	5070
INFL934	Light chain variable region	1286-C05	WO2010132604, US20120128671 SEQ ID NO: 5	5071
INFL935	Light chain variable region	1286-A11	WO2010132604, US20120128671 SEQ ID NO: 6	5072
INFL936	Light chain variable region mouse IgG	1A2	WO2015028478 SEQ ID NO: 3	5073
INFL937	Light chain variable region mouse IgG	7B8	WO2015028478 SEQ ID NO: 5	5074
INFL938	Light chain variable region, partial	monoclonal antibody TCN-031	US8900590, US2012039899, Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23871.1 (106aa)	5075
INFL939	Light chain variable region, partial	monoclonal antibody TCN-032	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23870.1 (107aa)	5076
INFL940	Light chain variable region, partial	monoclonal antibody 3362_B11	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663	5077

			(2010), NCBI Accession # ADK23886.1 (107aa)	
INFL941	Light chain variable region, partial	monoclonal antibody 3260_D19	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23885.1 (106aa)	5078
INFL942	Light chain variable region, partial	monoclonal antibody 3253_P10	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23884.1(107aa)	5079
INFL943	Light chain variable region, partial	monoclonal antibody 3248_P18	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23883.1 (106aa)	5080
INFL944	Light chain variable region, partial	monoclonal antibody 3139_P23	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23882.1(107aa)	5081
INFL945	Light chain variable region, partial	monoclonal antibody 3420_I23	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23881.1(108aa)	5082
INFL946	Light chain variable region, partial	monoclonal antibody 3255_J06	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23880.1(108aa)	5083
INFL947	Light chain variable region, partial	monoclonal antibody 3252_C13	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23879.1 (108aa)	5084
INFL948	Light chain variable region, partial	monoclonal antibody 3136_G05	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23878.1 (108aa)	5085

INFL949	Light chain variable region, partial	monoclonal antibody 3244_H04	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23877.1 (107aa)	5086
INFL950	Light chain variable region, partial	monoclonal antibody 3245_O19	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23876.1 (107aa)	5087
INFL951	Light chain variable region, partial	monoclonal antibody 3259_J21	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23875.1 (107aa)	5088
INFL952	Light chain variable region, partial	monoclonal antibody 3243_J07	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23874.1 (108aa)	5089
INFL953	Light chain variable region, partial	monoclonal antibody 3244_I10	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23873.1 (108aa)	5090
INFL954	Light chain variable region, partial	monoclonal antibody 3241_G23	Grandea, A.G. et al., Human antibodies reveal a protective epitope that is highly conserved among human and nonhuman influenza A viruses; Proc. Natl. Acad. Sci. U.S.A. 107 (28), 12658-12663 (2010), NCBI Accession #ADK23872.1 (108aa)	5091
INFL955	Light chain variable region, partial	100F4-LV	Hu, H., et al., A Human Antibody Recognizing a Conserved Epitope of H5 Hemagglutinin Broadly Neutralizes Highly Pathogenic Avian Influenza H5N1 Viruses; J. Virol. 86 (6), 2978-2989 (2012), NCBI Accession #AEL30604.1 (112aa)	5092
INFL956	Light Chain, Fab Fragment	ch65	Whittle, J.R. et al., Broadly neutralizing human antibody that recognizes the receptor-binding pocket of influenza virus hemagglutinin; Proc. Natl. Acad. Sci. U.S.A. 108 (34), 14216-14221 (2011), NCBI Accession #3SM5_L	5093
INFL957	Light chain	1I20	WO2010127252, US8894997 SEQ ID NO: 6	5094
INFL958	Light chain		WO2010127252, US8894997 SEQ ID NO: 12	5095

INFL959	Monoclonal antibody heavy chain	Neutralizing Human Monoclonal Antibody With 1968 H3 Ha	Wu, Y. et al., A potent broad-spectrum protective human monoclonal antibody crosslinking two hemagglutinin monomers of influenza A virus; Nat Commun 6, 7708 (2015), NCBI Accession #4UBD_C	5096
INFL960	Monoclonal antibody light chain	Neutralizing Human Monoclonal Antibody With 1968 H3 Ha	Wu, Y. et al., A potent broad-spectrum protective human monoclonal antibody crosslinking two hemagglutinin monomers of influenza A virus; Nat Commun 6, 7708 (2015), NCBI Accession #4UBD_D	5097
INFL96 i	Mutated heavy chain variable	8G9 mutated	US8603467 SEQ ID NO: 42	5098
1NFL962	Mutated heavy chairs variable (VH-LV)	13D4 mutated	US8603467 SEQ ID NO: 46	5099
INFL963	Mutated heavy chain variable (VH-SV)	i3D4 mutated	US8603467 SEQ ID NO: 44	5100
INTL.964	Nanobody	202-C8	US201 !0i82897, WO2009 !47248 SEQ ID NO: 138	5101
INFL965	Nanobody	203-B12	US201 10182897, WO2009147248 SEQ ID NO: 2439	5102
INFL966	Nanobody	203-H9	US201 !0182897, WO2009 !47248 SEQ ID NO: 2445	5103
INFL967	Scfv	JM7_B-G7	WO20 12072788 SEQ ID NO: 7	5104
INFL968	Scfv	JM7_S-F8	WO20 12072788 SEQ ID NO: 15	5105
INFL969	Scfv	JM7JH-F1	WO20 12072788 SEQ ID NO: 17	5106
INFL970	Scfv	JM7_S-A9	WO20 12072788 SEQ ID NO: 19	5107
INTL.971	Scfv	JM7_S-AIO	WO20 12072788 SEQ ID NO: 21	5108
INFL972	Scfv	JM7_B-I-I	WO20 12072788 SEQ ID NO: 23	5109
1NFL973	Scfv	JM6_SC-H1	WO20 12072788 SEQ ID NO: 25	5110
INFL974	Scfv	jM6_SC_D3	WO20 12072788 SEQ ID NO: 27	5111
INFL975	Scfv	H2526	Schmidt, A.G. et ai., Viral receptor-binding site antibodies with diverse germline origins; Cell 161(5), 1026-1034 (2015),~NCBI Accession #4YJZ_L	5112
1NFL976	Scfv fragment	AVC4	WO20 10040572 A2 Fig. 6	5113
INFL977	Scfv fragment	AVD1	WO2010040572A2 Fig. 8	5114
INFL978	Scfv fragment	AVE2	WO20 10040572 A2 Fig. 10	5115
INFL979	Scfv fragment	AVA6	WO2010040572A2 Fig. 12	5116
INFL980	Scfv fragment	AVC-4	WO20 10040572 A2 Fig. 14	5117
1NFL981	Scfv heavy chain variable region	SC06-141	US20150 104459 SEQ ID NO: 309	5118
INFL982	Scfv heavy chain variable region	SC06-255	US20150104459 SEQ ID NO: 313	5119
1NFL983	Scfv heavy chain variable region	SC06-257	US20150 104459 SEQ ID NO: 317	5120
INFL984	Scfv heavy chain variable region	SC6-260	US20150104459 SEQ ID NO: 321	5121
INFL985	Scfv heavy chain variable region	SC06-261	US20150104459 SEQ ID NO: 325	5122
INFL986	Scfv heavy chain variable region	SC06-262	US20150104459 SEQ ID NO: 329	5123
INFL987	Scfv heavy chain variable region	SC06-268	US20150104459 SEQ ID NO: 333	5124

INFL988	Scfv heavy chain variable region	SC06-272	US20150104459 SEQ ID NO: 337	5125
INFL989	ScfV heavy chain variable reedon	SC06-296	US20150 104459 SEQ ID NO: 341	5126
ENFL990	Scfv heavy chain variable region	SC06-301	US20 150104459 SEQ ID NO: 345	5127
INTL.991	Scfv heavy chain variable reedon	SC06-307	US20150 104459 SEQ ID NO: 349	5128
ENFL992	Scfv heavy chain variable region	SC06-3 10	US20 150104459 SEQ ID NO: 353	5129
INFL993	Scfv heavy chain variable reedon	SC06-314	US20150 104459 SEQ ID NO: 357	5130
ENFL994	Scfv heavy chain variable region	SC06-323	US20 150104459 SEQ ID NO: 361	5131
INFL995	Scfv heavy chain variable reedon	SC06-325	US20150 104459 SEQ ID NO: 365	5132
ENFL996	Scfv heavy chain variable region	SC06-327	US20 150104459 SEQ ID NO: 369	5133
INTL.997	Scfv heavy chain variable reedon	SC06-328	US20150 104459 SEQ ID NO: 373	5134
ENFL998	Scfv heavy chain variable region	SC06-329	US20 150104459 SEQ ID NO: 377	5135
1NFL999	Scfv heavy chain variable region	SC06-33 1	US20150 104459 SEQ ID NO: 381	5136
INFL IOOO	Scfv heavy chain variable region	SC06-332	US20150104459 SEQ ID NO: 385	5137
INFL1001	Scfv heavy chain variable region	SC06-334	US20150 104459 SEQ ID NO: 389	5138
INFL 1002	Scfv heavy chain variable region	SC06-336	US20150104459 SEQ ID NO: 393	5139
INFL1003	Scfv heavy chain variable region	SC06-339	US20150 104459 SEQ ID NO: 397	5140
INFL 1004	Scfv heavy chain variable region	SC06-342	US20150104459 SEQ ID NO: 401	5141
INFL1005	Scfv heavy chain variable region	SC06-343	US20150 104459 SEQ ID NO: 405	5142
INFL 1006	Scfv heavy chain variable region	SC06-344	US20150104459 SEQ ID NO: 409	5143
INFL1007	Scfv heavy chain variable region	CR6255	US20150104459 SEQ ID NO: 417	5144
INFL1008	Scfv heavy chain variable region	CR6257	US20150104459 SEQ ID NO: 423	5145
INFL1009	Scfv heavy chain variable region	CR6260	US20150104459 SEQ ID NO: 429	5146
INFL1010	Scfv heavy chain variable region	CR6261	US20150104459 SEQ ID NO: 435	5147
INFL101 1	Scfv heavy chain variable region	CR6262	US20150104459 SEQ ID NO: 441	5148
INFL1012	Scfv heavy chain variable region	CR6268	US20150104459 SEQ ID NO: 447	5149
INFL1013	Scfv heavy chain variable region	CR6272	US20150104459 SEQ ID NO: 453	5150
INFL1014	Scfv heavy chain variable region	CR696	US20150104459 SEQ ID NO: 459	5151
INFL1015	Scfv heavy chain variable region	CR6301	US20150104459 SEQ ID NO: 465	5152
INFL1016	Scfv heavy chain variable region	CR6307	US20150104459 SEQ ID NO: 471	5153

INFL 1017	Scfv heavy chain variable region	CR63 10	US20 150104459 SEQ ID NO: 477	5154
INFL1018	Scfv heavy chain variable reedon	CR63 14	US20150 104459 SEQ ID NO: 483	5155
INFL 10 I9	Scfv heavy chain variable region	CR6323	US20 150104459 SEQ ID NO: 489	5156
INTL. 1020	Scfv heavy chain variable reedon	CR6325	US20150 104459 SEQ ID NO: 495	5157
INFL 102 1	Scfv heavy chain variable region	CR6327	US20 150104459 SEQ ID NO: 501	5158
INTL. 1022	Scfv heavy chain variable reedon	CR6328	US20150 104459 SEQ ID NO: 507	5159
INFL 1023	Scfv heavy chain variable region	CR6329	US20 150104459 SEQ ID NO: 513	5160
INTL. 1024	Scfv heavy chain variable reedon	CR6331	US20150 104459 SEQ ID NO: 519	5161
INFL 1025	Scfv heavy chain variable region	CR6332	US20 150104459 SEQ ID NO: 525	5162
INTL. 1026	Scfv heavy chain variable reedon	CR6334	US20150 104459 SEQ ID NO: 531	5163
INFL 1027	Scfv heavy chain variable region	CR6336	US20 150104459 SEQ ID NO: 537	5164
INFL1028	Scfv heavy chain variable region	CR6339	US20150 104459 SEQ ID NO: 543	5165
INFL1029	Scfv heavy chain variable region	CR6342	US20 150104459 SEQ ID NO: 550	5166
1NFL1030	Scfv heavy chain variable region	CR6343	US20150 104459 SEQ ID NO: 556	5167
INFL 103 1	Scfv heavy chain variable region	CR6344	US20 150104459 SEQ ID NO: 562	5168
1NFL1032	Scfv light chain variable region	SC06-141	US20150 104459 SEQ ID NO: 310	5169
INFL 1033	Scfv light chain variable region	SC06-255	US20150104459 SEQ ID NO: 314	5170
1NFL1034	Scfv light chain variable region	SC06-257	US20150 104459 SEQ ID NO: 318	5171
INFL 1035	Scfv light chain variable region	SC6-260	US20150104459 SEQ ID NO: 322	5172
INFL1036	Scfv light chain variable region	SC06-261	US20150104459 SEQ ID NO: 326	5173
INFL1037	Scfv light chain variable region	SC06-262	US20150104459 SEQ ID NO: 330	5174
INFL1038	Scfv light chain variable region	SC06-268	US20150104459 SEQ ID NO: 334	5175
INFL1039	Scfv light chain variable region	SC06-272	US20150104459 SEQ ID NO: 338	5176
INFL1040	Scfv light chain variable region	SC06-296	US20150104459 SEQ ID NO: 342	5177
INFL1041	Scfv light chain variable region	SC06-301	US20150104459 SEQ ID NO: 346	5178
INFL1042	Scfv light chain variable region	SC06-307	US20150104459 SEQ ID NO: 350	5179
INFL1043	Scfv light chain variable region	SC06-3 10	US20150104459 SEQ ID NO: 354	5180
INFL1044	Scfv light chain variable region	SC06-3 14	US20150104459 SEQ ID NO: 358	5181
INFL1045	Scfv light chain variable region	SC06-323	US20150104459 SEQ ID NO: 362	5182

INFL 1046	Scfv light chain variable region	SC06-325	US20150104459 SEQ ID NO: 366	5183
INFL1047	Scfv light chain variable reedon	SC06-327	US20150 104459 SEQ ID NO: 370	5184
INFL 1048	Scfv Sight clmin variable region	SC06-328	US20 150104459 SEQ ID NO: 374	5185
INTL. 1049	Scfv Sight chain variable reedon	SC06-329	US20150 104459 SEQ ID NO: 378	5186
INFL 1050	Scfv Sight clmin variable region	SC06-33 1	US20 150104459 SEQ ID NO: 382	5187
INTL. 105 1	Scfv Sight chain variable reedon	SC06-332	US20150 104459 SEQ ID NO: 386	5188
INFL 1052	Scfv Sight clmin variable region	SC06-334	US20 150104459 SEQ ID NO: 390	5189
INTL. 1053	Scfv Sight chain variable reedon	SC06-336	US20150 104459 SEQ ID NO: 394	5190
INFL 1054	Scfv Sight clmin variable region	SC06-339	US20 150104459 SEQ ID NO: 398	5191
INTL. 1055	Scfv Sight chain variable reedon	SC06-342	US20150 104459 SEQ ID NO: 402	5192
INFL 1056	Scfv Sight clmin variable region	SC06-343	US20 150104459 SEQ ID NO: 406	5193
INFL1057	Scfv light chain variable region	SC06-344	US20150 104459 SEQ ID NO: 410	5194
[NFL 1058	Scfv light chain variable region	CR6141	US20 150104459 SEQ ID NO: 414	5195
INFL1059	Scfv light chain variable region	CR6255	US20150 104459 SEQ ID NO: 420	5196
[NFL 1060	Scfv light chain variable region	CR6257	US20 150104459 SEQ ID NO: 426	5197
INFL106 1	Scfv light chain variable region	CR6260	US20150 104459 SEQ ID NO: 432	5198
[NFL 1062	Scfv light chain variable region	CR6261	US20 150104459 SEQ ID NO: 438	5199
INFL1063	Scfv light chain variable region	CR6262	US20150 104459 SEQ ID NO: 444	5200
[NFL 1064	Scfv light chain variable region	CR6268	US20 150104459 SEQ ID NO: 450	5201
INFL1065	Scfv light chain variable region	CR6272	US20150104459 SEQ ID NO: 456	5202
INFL1066	Scfv light chain variable region	CR696	US20150104459 SEQ ID NO: 462	5203
INFL1067	Scfv light chain variable region	CR6301	US20150104459 SEQ ID NO: 468	5204
INFL1068	Scfv light chain variable region	CR6307	US20150104459 SEQ ID NO: 474	5205
INFL1069	Scfv light chain variable region	CR63 10	US20150104459 SEQ ID NO: 480	5206
INFL1070	Scfv light chain variable region	CR63 14	US20150104459 SEQ ID NO: 486	5207
INFL1071	Scfv light chain variable region	CR6323	US20150104459 SEQ ID NO: 492	5208
INFL1072	Scfv light chain variable region	CR6325	US20150104459 SEQ ID NO: 498	5209
INFL1073	Scfv light chain variable region	CR6327	US20150104459 SEQ ID NO: 504	5210
INFL1074	Scfv light chain variable region	CR6328	US20150104459 SEQ ID NO: 510	5211

INFL 1075	Scfv light chain variable region	CR6329	US20 150104459 SEQ ID NO: 516	5212
INFL 1076	Scfv light chain variable region	CR633 1	US20150 104459 SEQ ID NO: 522	52 13
INFL 1077	Scfv Sight cliain variable region	CR6332	US20 150104459 SEQ ID NO: 528	52 14
INTL. 1078	Scfv light chain variable region	CR6334	US20150 104459 SEQ ID NO: 534	52 15
INFL 1079	Scfv Sight cliain variable region	CR6336	US20 150104459 SEQ ID NO: 540	52 16
INTL. 1080	Scfv light chain variable region	CR6339	US20150 104459 SEQ ID NO: 547	52 17
INFL 1081	Scfv light cliain variable region	CR6342	US20 150104459 SEQ ID NO: 553	52 18
INTL. 1082	Scfv light chain variable region	CR6343	US20150 104459 SEQ ID NO: 559	52 19
INFL 1083	Scfv light cliain variable region	CR6344	US20 150104459 SEQ ID NO: 565	5220
INTL. 1084	Vhch antibody	641 1-9	Schmidt, A.G. et al., Viral receptor-binding site antibodies with diverse germline origins; Cell 16 1 (5), 1026-1034 (2015), NCBI Accession #4YK4_Z	522 1
INFL 1085	Vlcl antibody	641 I-9	Schmidt, A.G. et al., Viral receptor-binding site antibodies with diverse germline origins; Cell 161 (5), 1026-1034 (2015), NCBI Accession #4YK4_Y	5222

[00316] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Patent No. US8003106 and US8540995, international Patent Publication No. WO20 15028478, WO201 2045001, US Publication No. LTS20150239960 and 11820130251715, the contents of each of which are herein incorporated by reference in their entirety, against influenza.

[00317] In one embodiment, the pay load region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 22 against Respiratory Syncytial Virus.

Table 22. Antibodies against Respiratory Syncytial Virus

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
RSV1	Heavy chain variable, F and G Proteins	clone 888	US20110189171; US7879329SEQ ID NO: 43	5223
RSV2	Heavy chain variable, F and G Proteins	mAb 824	US20110189171; US7879329 SEQ ID NO: 178	5224
RSV3	Heavy chain variable, F and G Proteins	clone 735	US20110189171; US7879329 SEQ ID NO: 1	5225
RSV4	Heavy chain variable, F and G Proteins	clone 736	US20110189171; US7879329 SEQ ID NO: 2	5226
RSV5	Heavy chain variable, F and G Proteins	clone 744	US20110189171; US7879329 SEQ ID NO: 3	5227
RSV6	Heavy chain variable, F and G Proteins	clone 793	US20110189171; US7879329 SEQ ID NO: 4	5228
RSV7	Heavy chain variable, F and G Proteins	clone 795	US20110189171; US7879329 SEQ ID NO: 5	5229

RSV8	**Heavy chain variable,** F and G Proteins	clone 796	**US201 10189171; US7879329** SEQ ID NO: 6	5230
RSV9	Heavy chain variable, F and G Proteins	clone 799	**US201 10189171; US7879329** SEQ ID NO: 7	5231
RSV10	Heavy chain variable, **F** and G Proteins	clone 800	**US20110189171; US78793 29** SEQ ID NO: 8	5232
RSVH	Heavy chains variable, F **and** G Proteins	clone 801	US20110189171; **US7879329** SEQ ID NO: 9	5233
RSV12	Heavy chain variable, **F** and G Proteins	clone 804	**US20110189171; US78793 29** SEQ ID NO: **10**	5234
RSV13	Heavy chains variable, F **and** G Proteins	clone 810	US20110189171; **US7879329** SEQ ID NO: **11**	5235
RSV14	Heavy chain variable, **F** and G Proteins	clone 811	**US20110189171; US78793 29** SEQ ID NO: **12**	5236
RSV15	Heavy chains variable, F **and** G Proteins	clone 812	US20110189171; **US7879329** SEQ ID NO: 13	5237
RSV16	Heavy chain variable, **F** and G Proteins	clone 814	**US20110189171; US78793 29** SEQ ID NO: **14**	5238
RSV17	Heavy chains variable, F **and** G Proteins	clone 816	US20110189171; **US7879329** SEQ ID NO: 15	5239
RSV18	Heavy chain variable, **F** and G Proteins	clone 817	**US20110189171; US78793 29** SEQ ID NO: 16	5240
RSV19	**Heavy** chain variable, F and G Proteins	clone 818	**US20110189171; US7879329** SEQ ID NO: 17	5241
RSV20	Heavy chain variable, **F** and **G** Proteins	clone **819**	US20110189171; **US7879329** SEQ ID NO: 18	5242
RSV21	**Heavy** chain variable, F and G Proteins	clone 824	**US20110189171; US7879329** SEQ ID NO: 19	5243
RSV22	Heavy chain variable, **F** and **G** Proteins	clone 825	US20110189171; **US7879329** SEQ ID NO: 20	5244
RSV23	**Heavy** chain variable, F and G Proteins	clone 827	**US20110189171; US7879329** SEQ ID NO: 21	5245
RSV24	Heavy chain variable, **F** and **G** Proteins	clone 829	US20110189171; **US7879329** SEQ ID NO: 22	5246
RSV25	**Heavy** chain variable, F and G Proteins	clone 830	**US20110189171; US7879329** SEQ ID NO: 23	5247
RSV26	Heavy chain variable, **F** and **G** Proteins	clone 831	US20110189171; **US7879329** SEQ ID NO: 24	5248
RSV27	**Heavy chain variable,** F and G Proteins	clone 835	**US201 10189171; US7879329** SEQ ID NO: 25	5249
RSV28	Heavy chain variable, F and G Proteins	clone 838	**US201 10189171; US7879329** SEQ ID NO: 26	5250
RSV29	**Heavy chain variable,** F and G Proteins	clone 841	**US201 10189171; US7879329** SEQ ID NO: 27	5251
RSV30	Heavy chain variable, F and G Proteins	clone 853	**US201 10189171; US7879329** SEQ ID NO: 28	5252
RSV31	**Heavy chain variable,** F and G Proteins	clone 855	**US201 10189171; US7879329** SEQ ID NO: 29	5253
RSV32	Heavy chain variable, F and G Proteins	clone 856	**US201 10189171; US7879329** SEQ ID NO: 30	5254
RSV33	**Heavy chain variable,** F and G Proteins	clone 857	**US201 10189171; US7879329** SEQ ID NO: 31	5255
RSV34	Heavy chain variable, F and G Proteins	clone 858	**US201 10189171; US7879329** SEQ ID NO: 32	5256
RSV35	**Heavy chain variable,** F and G Proteins	clone 859	**US201 10189171; US7879329** SEQ ID NO: 33	5257
RSV36	Heavy chain variable, F and G Proteins	clone 861	**US201 10189171; US7879329** SEQ ID NO: 34	5258

RSV37	Heavy chain variable, F and G Proteins	clone 863	US20110189171; US7879329 SEQ ID NO: 35	5259
RSV38	Heavy chain variable, F and G Proteins	clone 868	US20110189171; US7879329 SEQ ID NO: 36	5260
RSV39	Heavy chain variable, F and G Proteins	clone 870	US20110189171; US7879329 SEQ ID NO: 37	5261
RSV40	Heavy chain variable, F and G Proteins	clone 871	US20110189171; US7879329 SEQ ID NO: 38	5262
RSV41	Heavy chain variable, F and G Proteins	clone 880	US20110189171; US7879329 SEQ ID NO: 39	5263
RSV42	Heavy chain variable, F and G Proteins	clone 881	US20110189171; US7879329 SEQ ID NO: 40	5264
RSV43	Heavy chain variable, F and G Proteins	clone 884	US20110189171; US7879329 SEQ ID NO: 41	5265
RSV44	Heavy chain variable, F and G Proteins	clone 886	US20110189171; US7879329 SEQ ID NO: 42	5266
RSV45	Heavy chain variable, F and G Proteins	clone 894	US20110189171; US7879329 SEQ ID NO: 44	5267
RSV46	heavy chain variable, F protein of RSV, MPV, or PVM	3210 variant 1	WO2013140247 SEQ ID NO: 13	5268
RSV47	heavy chain variable, F protein of RSV, MPV, or PVM	3210 variant 2, 3210 variant 3, 3210 variant 6	WO2013140247 SEQ ID NO: 17	5269
RSV48	heavy chain variable, F protein of RSV, MPV, or PVM	2430 variant 1	WO2013140247 SEQ ID NO: 29	5270
RSV49	heavy chain variable, F protein of RSV, MPV, or PVM	2430 variant 2, 2430 variant 5	WO2013140247 SEQ ID NO: 33	5271
RSV50	heavy chain variable, F protein of RSV, MPV, or PVM	3210 variant 4, 3210 variant 5	WO2013140247 SEQ ID NO: 49	5272
RSV51	heavy chain variable, F protein of RSV, MPV, or PVM	2430 variant 3, 2430 variant 4	WO2013140247 SEQ ID NO: 59	5273
RSV52	Heavy chain variable, CDR Grafted, F Protein,		US20140093501 SEQ ID NO: 31	5274
RSV53	Heavy chain, F Protein	AM22	US8568726 SEQ ID NO: 16	5275
RSV54	Heavy chain, F Protein	RSVF2-5	US8221759 SEQ ID NO: 1	5276
RSV55	Heavy chain, F Protein		EP1259547; US8153133 SEQ ID NO: 4	5277
RSV56	Heavy chain, F Protein	MEDI-493/Pavilizumab-N-VL (Brand name Synagis)	EP1259547; US8153133 SEQ ID NO: 2	5278
RSV57	Heavy chain, F Protein		EP1259547; US8153133 SEQ ID NO: 36	5279
RSV58	Heavy chain, F Protein	clone 18	EP1259547; US8153133 SEQ ID NO: 37	5280
RSV59	Heavy chain, F Protein	clone 19	EP1259547; US8153133 SEQ ID NO: 39	5281
RSV60	Heavy chain, F Protein	clone 20	EP1259547; US8153133 SEQ ID NO: 41	5282
RSV61	Heavy chain, F Protein	clone 21	EP1259547; US8153133 SEQ ID NO: 43	5283
RSV62	Heavy chain, F Protein	clone 22	EP1259547; US8153133 SEQ ID NO: 45	5284
RSV63	Heavy chain, F Protein	clone 23	EP1259547; US8153133 SEQ ID NO: 47	5285
RSV64	Heavy chain, F Protein	clone 24	EP1259547; US8153133 SEQ ID NO: 49	5286
RSV65	Heavy chain, F Protein	clone 25	EP1259547; US8153133 SEQ ID NO: 51	5287

RSV66	Heavy chain, F Protein	clone 26	EP1259547; US8153 133 SEQ ID NO: 53	5288
RSV67	Heavy chain variable region, F Protein		US20140093501 SEQ ID NO: 17	5289
RSV68	Heavy chain variable region, F Protein	MAM308F	US20 14009350 I SEQ ID NO: 18	5290
RSV69	Heavy chai rs variable region, F Protein	hmCOR	US20140093501 SEQ ID NO: 30	529 !
RSV70	Heavy chain variable region, F Protein	MAS) 1129	US20 14009350 I SEQ ID NO: 32	5292
RSV71	Heavy chai rs variable region, F Protein	RSV G8	US7867497 SEQ ID NO: 2	5293
RSV72	Heavy chain variable region, F Protein	Clone I	US20 I20 135006 SEQ ID NO: 18	5294
RSV73	Heavy chai rs variable region, F Protein	Clone 2	US20120 135006 SEQ ID NO: 20	5295
RSV74	Heavy chain variable region, F Protein	Clone 3	US20 I20 135006 SEQ ID NO: 22	5296
RSV75	Heavy chai rs variable region, F Protein	Clone 22	US20120 135006 SEQ ID NO: 24	5297
RSV76	Heavy chain variable region, F Protein	Clone 23	US20120135006 SEQ ID NO: 26	5298
RSV77	Heavy chain variable region, F Protein	RSV13-9	WO20090881 59 SEQ ID NO: 4	5299
RSV78	HV3 heavy chain variable, F Protein		US20140093501 SEQ ID NO: 16	5300
RSV79	Constant heavy region, F protein	B4HuVK	EP63 6182; WO 1993 0202 10; SEQ ID NO: 6	530 1
RSV80	Constant heavy region, F protein	B 13/B14HuVK	EP63 6182; WO19930202 10; SEQ ID NO: 8	5302
RSV81	Heavy chain, F protein	58c5	1JS20 1400447 I9 SEQ ID NO: 1	5303
RSV82	Heavy chain, F protein	sc5	US20140044719 SEQ ID NO: 9	5304
RSV83	Heavy chain, F protein		US201 10027294 SEQ ID NO: 74	5305
RSV84	Heavy chain, F protein		US201 10027294 SEQ ID NO: 75	5306
RSV85	Heavy chain, F protein		US201 10027294 SEQ ID NO: 76	5307
RSV86	Heavy chain, F protein		US201 10027294 SEQ ID NO: 77	5308
RSV87	Heavy chain, F protein		US201 10027294 SEQ ID NO: 78	5309
RSV88	Heavy chain, F protein		US201 10027294 SEQ ID NO: 79	5310
RSV89	Heavy chain, F protein		US201 10027294 SEQ ID NO: 80	5311
RSV90	Heavy chain, F protein	*G1-1*	US20050 175986 SEQ ID NO: 5	5312
RSV9 1	Heavy chain, F protein	A construct	US20050175986 SEQ ID NO: 7	5313
RSV92	Heavy chain, F protein	B construct	US20050 175986 SEQ ID NO: 8	5314
RSV93	Heavy chain, F protein	hul 9A	US200500 19758; WO1998019704 SEQ ID NO: 5	5315
RSV94	Heavy chain, F protein	hul9B	US20050019758; WO19980 19704 SEQ ID NO: 6	5316
RSV95	Heavy chain, F protein	hul9C	US20050019758; WO1998019704 SEQ ID NO: 7	5317
RSV96	Heavy chain, F protein	hul9D	US20050019758; WO19980 19704 SEQ ID NO: 8	5318

RSV97	Heavy chain, F protein	B4HuVH	EP636182; WO1993020210; SEQ ID NO: 5	5319
RSV98	Heavy chain, F protein	B13/B14HuVK	EP636182; WO1993020210; SEQ ID NO: 7	5320
RSV99	Heavy chain, F protein	RSV19	EP636182; WO1993020210; SEQ ID NO: 10	5321
RSV100	Heavy chain, F protein		WO19922004381	5322
RSV101	Heavy chain, F protein		WO19922004381	5323
RSV102	Heavy chain variable region, F Protein	P1212	US20140044719 SEQ ID NO: 122	5324
RSV103	Heavy chain variable region, F Protein	P12f4	US20140044719 SEQ ID NO: 131	5325
RSV104	Heavy chain variable region, F Protein	P11d4	US20140044719 SEQ ID NO: 137	5326
RSV105	Heavy chain variable region, F Protein	A1e9	US20140044719 SEQ ID NO: 144	5327
RSV106	Heavy chain variable region, F Protein	A12a6	US20140044719 SEQ ID NO: 149	5328
RSV107	Heavy chain variable region, F Protein	A13c4	US20140044719 SEQ ID NO: 155	5329
RSV108	Heavy chain variable region, F Protein	A17d4	US20140044719 SEQ ID NO: 161	5330
RSV109	Heavy chain variable region, F Protein	A4B4	US20140044719 SEQ ID NO: 167	5331
RSV110	Heavy chain variable region, F Protein	A8c7	US20140044719 SEQ ID NO: 172	5332
RSV111	Heavy chain variable region, F Protein	IX-493L1FR	US20140044719 SEQ ID NO: 176	5333
RSV112	Heavy chain variable region, F Protein	M3H9	US20140044719 SEQ ID NO: 181	5334
RSV113	Heavy chain variable region, F Protein	B21M	US20110027294 SEQ ID NO: 49	5335
RSV114	Heavy chain variable region, F Protein	101F	US20110027294 SEQ ID NO: 4	5336
RSV115	Heavy chain variable region, F Protein	HNK20	EP1720908; WO2005079479 SEQ ID NO: 1	5337
RSV116	Heavy chain variable region, F Protein	P1212	US20140044719 SEQ ID NO: 123	5338
RSV117	Heavy chain variable region, F Protein	P12f4	US20140044719 SEQ ID NO: 132	5339
RSV118	Heavy chain variable region, F Protein	P11d4	US20140044719 SEQ ID NO: 138	5340
RSV119	Heavy chain variable region, F Protein	A1e9	US20140044719 SEQ ID NO: 145	5341
RSV120	Heavy chain variable region, F Protein	A12a6	US20140044719 SEQ ID NO: 150	5342
RSV121	Heavy chain variable region, F Protein	A13c4	US20140044719 SEQ ID NO: 156	5343
RSV122	Heavy chain variable region, F Protein	A17d4	US20140044719 SEQ ID NO: 162	5344
RSV123	Heavy chain variable region, F Protein	A4B4	US20140044719 SEQ ID NO: 168	5345
RSV124	Heavy chain variable region, F Protein	A8c7	US20140044719 SEQ ID NO: 173	5346
RSV125	Heavy chain variable region, F Protein	IX-493L1FR	US20140044719 SEQ ID NO: 177	5347
RSV126	Heavy chain variable region, F Protein	H1 H3564P	WO2014159822 SEQ ID NO: 2	5348

RSV127	Heavy chain variable region, F Protein	H1 H3565P	WO2014 159822 SEQ ID NO: 18	5349
RSV128	Heavy chain variable region, F Protein	HI H3566P	WO20 14 159822 SEQ ID NO: 34	5350
RSV129	Heavy chain variable region, F Protein	H1 H3567P	WO20 14 !59822 SEQ ID NO: 50	535 1
RSV130	Heavy chai rs variable region, F Protein	H1 H3581 P	WO2014159822 SEQ ID NO: 66	5352
RSV13 1	Heavy chain variable region, F Protein	H1 H3583P	WO20 14 !59822 SEQ ID NO: 82	5353
RSV132	Heavy chai rs variable region, F Protein	H1 H3589P	WO2014159822 SEQ ID NO: 98	5354
RSV133	Heavy chain variable region, F Protein	H1 H359 I P	WO20 14 !59822 SEQ ID NO: 114	5355
RSV134	Heavy chai rs variable region, F Protein	H1 H3592P	WO2014159822 SEQ ID NO: 130	5356
RSV135	Heavy chain variable region, F Protein	H1 H3597P	WO20 14 !59822 SEQ ID NO: 146	5357
RSV136	Heavy chai rs variable region, F Protein	H i H3598P	WO2014159822 SEQ ID NO: 162	5358
RSV137	Heavy chain variable region, F Protein	H1 H3603P	WO20 14 !59822 SEQ ID NO: 178	5359
RSV138	Heavy chain variable region, F Protein	H1 H3604P	WO20 14 159822 SEQ ID NO: 194	5360
RSV139	Heavy chain variable region, F Protein	H1 H3605P	WO20 14 159822 SEQ ID NO: 210	5361
RSV140	Heavy chain variable region, F Protein	H1 H3607P	WO20 14 159822 SEQ ID NO: 226	5362
RSV141	Heavy chain variable region, F Protein	H 1 H3608P2	WO20 14 159822 SEQ ID NO: 242	5363
RSV142	Heavy chain variable region, F Protein	H1 H3592P2	WO20 14 159822 SEQ ID NO: 258	5364
RSV143	Heavy chain variable region, F Protein	H1 H3592P3	WO20 14 159822 SEQ ID NO: 274	5365
RSV144	Heavy chain variable region, F Protein	H1 M3621 N	WO20 14 159822 SEQ ID NO: 290	5366
RSV145	Heavy chain variable region, F Protein	H1 M3622N	WO20 14 159822 SEQ ID NO: 306	5367
RSV146	Heavy chain variable region, F Protein	H1 M2634N	WO2014159822 SEQ ID NO: 322	5368
RSV147	Heavy chain variable region, F Protein	H1 M3627N	WO20 14 159822 SEQ ID NO: 338	5369
RSV148	Heavy chain variable region, F Protein	Clone No. 735	US20 120009623 SEQ ID NO: 1	5370
RSV149	Heavy chain variable region, F Protein	Clone No. 736	US20 120009623 SEQ ID NO: 2	5371
RSV150	Heavy chain variable region, F Protein	Clone No. 744	US20 120009623 SEQ ID NO: 3	5372
RSV15 1	Heavy chain variable region, F Protein	Clone No. 793	US20 120009623 SEQ ID NO: 4	5373
RSV152	Heavy chain variable region, F Protein	Clone No. 795	US20 120009623 SEQ ID NO: 5	5374
RSV153	Heavy chain variable region, F Protein	Clone No. 796	US20 120009623 SEQ ID NO: 6	5375
RSV154	Heavy chain variable region, F Protein	Clone No. 799	US20 120009623 SEQ ID NO: 7	5376
RSV155	Heavy chain variable region, F Proteirs	Clone No. 800	US20 120009623 SEQ ID NO: 8	5377

RSV156	Heaw_ chain variable region F Protein	Clone No. 801	US20 120009623 SEQ ID NO: 9	5378
RSV157	Heavy chain variable region, F Protein	Clone No. 804	US20120009623 SEQ ID NO: 10	5379
RSV158	Heavy chain variable region, F Protein	Clone No. 810	US20120009623 SEQ ID NO: 11	5380
RSV159	Heavy chain variable region, F Protein	Clone No. 811	US20120009623 SEQ ID NO: 12	5381
RSV160	Heavy chain variable region, F Protein	Clone No. 812	US20120009623 SEQ ID NO: 13	5382
RSV161	Heavy chain variable region, F Protein	Clone No. 814	US20120009623 SEQ ID NO: 14	5383
RSV162	Heavy chain variable region, F Protein	Clone No. 816	US20120009623 SEQ ID NO: 15	5384
RSV163	Heavy chain variable region, F Protein	Clone No. 817	US20120009623 SEQ ID NO: 16	5385
RSV164	Heavy chain variable region, F Protein	Clone No. 818	US20120009623 SEQ ID NO: 17	5386
RSV165	Heavy chain variable region, F Protein	Clone No. 819	US20120009623 SEQ ID NO: 18	5387
RSV166	Heavy chain variable region, F Protein	Clone No. 824	US20120009623 SEQ ID NO: 19	5388
RSV167	Heavy chain variable region, F Protein	Clone No. 825	US20120009623 SEQ ID NO: 20	5389
RSV168	Heavy chain variable region, F Protein	Clone No. 827	US20120009623 SEQ ID NO: 21	5390
RSV169	Heavy chain variable region, F Protein	Clone No. 829	US20120009623 SEQ ID NO: 22	5391
RSV170	Heavy chain variable region, F Protein	Clone No. 830	US20120009623 SEQ ID NO: 23	5392
RSV171	Heavy chain variable region, F Protein	Clone No. 831	US20120009623 SEQ ID NO: 24	5393
RSV172	Heavy chain variable region, F Protein	Clone No. 835	US20120009623 SEQ ID NO: 25	5394
RSV173	Heavy chain variable region, F Protein	Clone No. 838	US20120009623 SEQ ID NO: 26	5395
RSV174	Heavy chain variable region, F Protein	Clone No. 841	US20120009623 SEQ ID NO: 27	5396
RSV175	Heavy chain variable region, F Protein	Clone No. 853	US20120009623 SEQ ID NO: 28	5397
RSV176	Heavy chain variable region, F Protein	Clone No. 855	US20120009623 SEQ ID NO: 29	5398
RSV177	Heavy chain variable region, F Protein	Clone No. 856	US20120009623 SEQ ID NO: 30	5399
RSV178	Heavy chain variable region, F Protein	Clone No. 857	US20120009623 SEQ ID NO: 31	5400
RSV179	Heavy chain variable region, F Protein	Clone No. 858	US20120009623 SEQ ID NO: 32	5401
RSV180	Heavy chain variable region, F Protein	Clone No. 859	US20120009623 SEQ ID NO: 33	5402
RSV181	Heavy chain variable region, F Protein	Clone No. 861	US20120009623 SEQ ID NO: 34	5403
RSV182	Heavy chain variable region, F Protein	Clone No. 863	US20120009623 SEQ ID NO: 35	5404
RSV183	Heavy chain variable region, F Protein	Clone No. 868	US20120009623 SEQ ID NO: 36	5405
RSV184	Heavy chain variable region, F Proteirs	Clone No. 870	US20120009623 SEQ ID NO: 37	5406

RSV185	Heavy chain variable region, F Protein	Clone No. 871	US20120009623 SEQ ID NO: 38	5407
RSV186	Heavy chain variable region, F Protein	Clone No. 880	US20120009623 SEQ ID NO: 39	5408
RSV187	Heavy chain variable region, F Protein	Clone No. 881	US20120009623 SEQ ID NO: 40	5409
RSV188	Heavy chain variable region, F Protein	Clone No. 884	US20120009623 SEQ ID NO: 41	5410
RSV189	Heavy chain variable region, F Protein	Clone No. 886	US20120009623 SEQ ID NO: 42	5411
RSV190	Heavy chain variable region, F Protein	Clone No. 888	US20120009623 SEQ ID NO: 43	5412
RSV191	Heavy chain variable region, F Protein	Clone No. 894	US20120009623 SEQ ID NO: 44	5413
RSV192	Heavy chain variable region, F Protein	Gλ-1	US20050175986 SEQ ID NO: 4	5414
RSV193	Super humanized heavy chain based on HNK20, F protein	SHVh1	EP1720908; WO2005079479 SEQ ID NO: 3	5415
RSV194	Super humanized heavy chain based on HNK20, F protein	SHVh2	EP1720908; WO2005079479 SEQ ID NO: 4	5416
RSV195	Super humanized heavy chain based on HNK20, F protein	SHVh3	EP1720908; WO2005079479 SEQ ID NO: 5	5417
RSV196	Super humanized heavy chain based on HNK20, F protein	SHVM	EP1720908; WO2005079479 SEQ ID NO: 6	5418
RSV197	Super humanized heavy chain based on HNK20, F protein	SHVh5	EP1720908; WO2005079479 SEQ ID NO: 7	5419
RSV198	Super humanized heavy chain based on HNK20, F protein	SHVM	EP1720908; WO2005079479 SEQ ID NO: 8	5420
RSV199	Super humanized heavy chain based on HNK20, F protein	SHVh7	EP1720908; WO2005079479 SEQ ID NO: 9	5421
RSV200	Heavy chain variable region, F Protein	B4	EP636182; WO1993020210; SEQ ID NO: 3	5422
RSV201	Heavy chain variable region, F Protein	B13/14	EP636182; WO1993020210; SEQ ID NO: 4	5423
RSV202	Heavy chain variable region, F Protein	RF-1	EP854730; WO1996040252; FIG 7B	5424
RSV203	Heavy chain variable region, F Protein	RF-2	EP854730; WO1996040252; FIG 8B	5425
RSV204	Heavy chain, G Protein	1F12	US8273354 SEQ ID NO: 28	5426
RSV205	Heavy chain, G Protein	3G12	US8273354 SEQ ID NO: 29	5427
RSV206	Heavy chain, G Protein	1A5	US8273354 SEQ ID NO: 30	5428
RSV207	Heavy chain, G Protein	3D3	US8273354 SEQ ID NO; 31	5429
RSV208	Heavy chain, G Protein	1G1	US8273354 SEQ ID NO: 32	5430
RSV209	Heavy chain, G Protein	2B11	US8273354 SEQ ID NO: 33	5431
RSV210	Heavy chain, G Protein	5D8	US8273354 SEQ ID NO: 34	5432
RSV211	Heavy chain, G Protein	2D10	US8273354 SEQ ID NO: 35	5433
RSV212	Heavy chain, G Protein	3F9	US8273354 SEQ ID NO; 36	5434
RSV213	Heavy chain, G Protein	1D4	US8273354 SEQ ID NO: 37	5435
RSV214	Heavy chain, G Protein	1G8	US8273354 SEQ ID NO: 38	5436
RSV215	Heavy chain, G Protein	6A12	US8273354 SEQ ID NO: 39	5437
RSV216	Heavy chain, G Protein	10C6	US8273354 SEQ ID NO: 40	5438
RSV217	Heavy chain, G Protein	Hu 131-2G	US8273354 SEQ ID NO: 41	5439
RSV218	Heavy chain, G Protein	AT46	US20150004155 SEQ ID NO: 109	5440

RSV219	Heavy chain, G Protein	AT32	US20150004155 SEQ ID NO: no	5441
RSV220	Heavy chain, G Protein	AT33	US20150004155 SEQ ID NO: 111	5442
RSV221	Heavy chain, G Protein	AT34	US20150004155 SEQ ID NO: 112	5443
RSV222	Heavy chains, G Protein	AT35	US20150004155 SEQ ID NO: 113	5444
RSV223	Heavy chain, G Protein	AT36	US20150004155 SEQ ID NO: 114	5445
RSV224	Heavy chains, G Protein	AT37	US20150004155 SEQ ID NO: 115	5446
RSV225	Heavy chain, G Protein	AT39	US20150004155 SEQ ID NO: 116	5447
RSV226	Heavy chains, G Protein	AT40	US20150004155 SEQ ID NO: 117	5448
RSV227	Heavy chain, G Protein	AT42	US20150004155 SEQ ID NO: 118	5449
RSV228	Heavy chains, G Protein	AT43	US20150004155 SEQ ID NO: 119	5450
RSV229	Heavy chain, G Protein	AT44	US20150004155 SEQ ID NO: 120	5451
RSV230	Heavy chain, G Protein	AT45	US20150004155 SEQ ID NO: 121	5452
RSV231	Heavy chain, G Protein	AT47	US20150004155 SEQ ID NO: 122	5453
RSV232	Heavy chain, G Protein	AT49	US20150004155 SEQ ID NO: 123	5454
RSV233	Heavy chain, G Protein	AT50	US20150004155 SEQ ID NO: 124	5455
RSV234	Heavy chain, G Protein	AT51	US20150004155 SEQ ID NO: 125	5456
RSV235	Heavy chain variable region, G Protein	CB058.1	WO2014170257 SEQ ID NO: 37	5457
RSV236	Heavy chain variable region, G Protein	CB048.3	WO2014170257 SEQ ID NO: 39	5458
RSV237	Heavy chain variable region, G Protein	CB010.7	WO2014170257 SEQ ID NO: 41	5459
RSV238	Heavy chain variable region, G Protein	CB003.1	WO2014170257 SEQ ID NO: 43	5460
RSV239	Heavy chain variable region, G Protein	CB028.2	WO2014170257 SEQ ID NO: 45	5461
RSV240	Heavy chain variable region, G Protein	CB002.1	WO2014170257 SEQ ID NO: 47	5462
RSV241	Heavy chain variable region, G Protein	CB017.3L	WO2014170258 SEQ ID NO: 73	5463
RSV242	Heavy chain variable region, G Protein	CB017.5L	WO2014170258 SEQ ID NO: 75	5464
RSV243	Heavy chain variable region, G Protein	CB028.1	WO2014170258 SEQ ID NO: 77	5465
RSV244	Heavy chain variable region, G Protein	CB030.1	WO2014170258 SEQ ID NO: 79	5466
RSV245	Heavy chain variable region, G Protein	CB047.1	WO2014170258 SEQ ID NO: 81	5467
RSV246	Heavy chain variable region, G Protein	CB047.2	WO2014170258 SEQ ID NO: 83	5468
RSV247	Heavy chain variable region, G Proteirs	CB065.1	WO2014170258 SEQ ID NO: 85	5469

RSV248	Heavy chain variable region, G Protein	CB071. IL	WO20 14 170258 SEQ ID NO: 87	5470
RSV249	Heavy chain variable region, G Protein	CB072. 1L	WO20 14 170258 SEQ ID NO: 89	547 1
RSV250	Heavy chain variable region, G Protein	CB073. IL	WO20 14 170258 SEQ ID NO: 91	5472
RSV25 !	Heavy chains variable region, G Protein	CB076.2L	WO2014170258 SEQ ID NO: 93	5473
RSV252	Heavy chain variable region, G Protein	CB079. 1	WO20 14 170258 SEQ ID NO: 95	5474
RSV253	Heavy chains	AM14	US20140377279 SEQ ID NO: 78	5475
RSV254	Heavy chain	AM 16	US20140377279 SEQ ID NO: 85	5476
RSV255	Heavy chains	AM23	US20140377279 SEQ ID NO: 92	5477
RSV256	Heavy chain	D25	US20140377279 SEQ ID NO: 7	5478
RSV257	Heavy chain	AFFF	US7635568 SEQ ID NO: 210	5479
RSV258	Heavy chain	P1212	US7635568 SEQ ID NO: 212	5480
RSV259	Heavy chain	P12f4	US7635568 SEQ ID NO: 214	5481
RSV260	Heavy chain	P1ld4	US7635568 SEQ ID NO: 216	5482
RSV261	Heavy chain	Ale9	US7635568 SEQ ID NO: 218	5483
RSV262	Heavy chain	A12a6	US7635568 SEQ ID NO: 220	5484
RSV263	Heavy chain	A13c4	US7635568 SEQ ID NO: 222	5485
RSV264	Heavy chain	A17d4	US7635568 SEQ ID NO: 224	5486
RSV265	Heavy chain	A4B4	US7635568 SEQ ID NO: 226	5487
RSV266	Heavy chain	A8c7	US7635568 SEQ ID NO: 228	5488
RSV267	Heavy chain	1X-493L1FR	US7635568 SEQ ID NO: 230	5489
RSV268	Heavy chain	H3-3F4	US7635568 SEQ ID NO: 232	5490
RSV269	Heavy chain	M3H9	US7635568 SEQ ID NO: 234	5491
RSV270	Heavy chain	Y10H6	US7635568 SEQ ID NO: 236	5492
RSV271	Heavy chain	DG	US7635568 SEQ ID NO: 238	5493
RSV272	Heavy chains	AFFF(l)	US7635568 SEQ ID NO: 240	5494
RSV273	Heavy chain	6H8	US7635568 SEQ ID NO: 242	5495
RSV274	Heavy chain	L1-7E5	US7635568 SEQ ID NO: 244	5496
RSV275	Heavy chain	L2-15B10	US7635568 SEQ ID NO: 246	5497
RSV276	Heavy chain	A13al 1	US7635568 SEQ ID NO: 248	5498
RSV277	Heavy chain	A1h5	US7635568 SEQ ID NO: 250	5499
RSV278	Heavy chain	A4B4(1)	US7635568 SEQ ID NO: 252	5500
RSV279	Heavy chain	A4B4L 1FR-S28R (MEDI-524, Motavizumab, Nuniax)	US7635568 SEQ ID NO: 254	5501
RSV280	Heavy chain	A4B4-F52S	US7635568 SEQ ID NO: 256	5502
RSV281	Heavy chain		US7364737 SEQ ID NO: 1	5503
RSV282	Heavy chain		US7364737 SEQ ID NO: 2	5504
RSV283	Heavy chain variable region	J variant	WO20 15 108967 SEQ ID NO: 12	5505
RSV284	Heavy chain variable region	L variant	WO2015 108967 SEQ ID NO: 13	5506

RSV285	Heavy chain variable region	LA variant	WO2015108967 SEQ ID NO: 14	5507
RSV286	Heavy chain variable region	1G7	WO2015108967 SEQ ID NO: 15	5508
RSV287	Heavy chain variable region	1F5	WO2015108967 SEQ ID NO: 16	5509
RSV288	Heavy chain variable region	2D10	WO2015108967 SEQ ID NO: 17	5510
RSV289	Heavy chain variable region	1G7-GLM	WO2015108967 SEQ ID NO: 18	5511
RSV290	Heavy chain variable region	B12-1	WO2015108967 SEQ ID NO: 19	5512
RSV291	Heavy chain variable region	E3-5	WO2015108967 SEQ ID NO: 20	5513
RSV292	Heavy chain variable region	E9-2	WO2015108967 SEQ ID NO: 21	5514
RSV293	Heavy chain variable region	1X-493L1FR	US7635568 SEQ ID NO: 7	5515
RSV294	Heavy chain variable region	AFFF, AFFF(1)	US7635568 SEQ ID NO: 9	5516
RSV295	Heavy chain variable region	P12f2	US7635568 SEQ ID NO: 17	5517
RSV296	Heavy chain variable region	P12f4	US7635568 SEQ ID NO: 24	5518
RSV297	Heavy chain variable region	P11d4	US7635568 SEQ ID NO: 28	5519
RSV298	Heavy chain variable region	Ale9, A1h5	US7635568 SEQ ID NO: 33	5520
RSV299	Heavy chain variable region	A12a6	US7635568 SEQ ID NO: 36	5521
RSV300	Heavy chain variable region	A13c4	US7635568 SEQ ID NO: 40	5522
RSV301	Heavy chain variable region	A17d4	US7635568 SEQ ID NO: 44	5523
RSV302	Heavy chain variable region	A4B4, A4B4(1), A4B4L1FR-S28R (MEDI-524, Motavizumab, Numax), A4B4-F52S	US7635568 SEQ ID NO: 48	5524
RSV303	Heavy chain variable region	A8c7	US7635568 SEQ ID NO: 51	5525
RSV304	Heavy chain variable region	H3-3F4, M3H9, Y10H6	US7635568 SEQ ID NO: 55	5526
RSV305	Heavy chain variable region	DG, 6H8, L1-7E5, L2-15B10	US7635568 SEQ ID NO: 78	5527
RSV306	Heavy chain variable region	A13a11	US7635568 SEQ ID NO: 67	5528
RSV307	Heavy chain variable region		US7364742 SEQ ID NO: 7	5529
RSV308	Heavy chain variable region		US7364742 SEQ ID NO: 8	5530
RSV309	Heavy chain variable region	D2E7	EP1807111; WO2006041970 SEQ ID NO: 2	5531
RSV310	Heavy chain variable region	2SD4	EP1807111; WO2006041970 SEQ ID NO: 10	5532
RSV311	Heavy chain, human metapneumovirus fusion protein with neutralizing antibody identifies a pneumovirus antigenic site,		Wen, X., "Structure of the human metapneumovirus fusion protein with neutralizing antibody identifies a pneumovirus antigenic site", Nat. Struct. Mol. Biol. 19 (4), 461-463 (2012), NCBI Accession # 4DAG_H(220 aa)	5533
RSV312	Heavy chain variable, M2 1 antigen	8A4/G9 - IgG	US20140348858 SEQ ID NO: 3	5534
RSV313	Heavy chain, Pre fusion RSV F protein	HMB2435	WO2015010792 SEQ ID NO: 13	5535

RSV314	Heavy chain, Pre fusion RSV F protein	HMB2437	WO2015010792 SEQ ID NO: 29	5536
RSV315	Heavy chain, Pre fusion RSV F protein	HMB2416	WO2015010792 SEQ ID NO: 45	5537
RSV316	Heavy chain, Pre fusion RSV F protein	HMB2437	WO2015010792 SEQ ID NO: 85	5538
RSV317	Heavy chain, Pre fusion RSV F protein	CR9501	WO2014202570 SEQ ID NO: 53	5539
RSV318	Heavy chain, Pre fusion RSV F protein	CR9502	WO2014202570 SEQ ID NO: 57	5540
RSV319	Heavy chain 1, Pre fusion RSV F protein	HMB2432	WO2015010792 SEQ ID NO: 61	5541
RSV320	Heavy chain 2, Pre fusion RSV F protein	HMB2432	WO2015010792 SEQ ID NO: 65	5542
RSV321	Heavy chain FR LG, Pre fusion RSV F protein	HMB2435	WO2015010792 SEQ ID NO: 75	5543
RSV322	light chain, F and G Proteins	clone 735	US20110189171; US7879329 SEQ ID NO: 89	5544
RSV323	light chain, F and G Proteins	clone 736	US20110189171; US7879329 SEQ ID NO: 90	5545
RSV324	light chain, F and G Proteins	clone 744	US20110189171; US7879329 SEQ ID NO: 91	5546
RSV325	light chain, F and G Proteins	clone 793	US20110189171; US7879329 SEQ ID NO: 92	5547
RSV326	light chain, F and G Proteins	clone 795	US20110189171; US7879329 SEQ ID NO: 93	5548
RSV327	light chain, F and G Proteins	clone 796	US20110189171; US7879329 SEQ ID NO: 94	5549
RSV328	light chain, F and G Proteins	clone 799	US20110189171; US7879329 SEQ ID NO: 95	5550
RSV329	light chain, F and G Proteins	clone 800	US20110189171; US7879329 SEQ ID NO: 96	5551
RSV330	light chain, F and G Proteins	clone 801	US20110189171; US7879329 SEQ ID NO: 97	5552
RSV331	light chain, F and G Proteins	clone 804	US20110189171; US7879329 SEQ ID NO: 98	5553
RSV332	light chain, F and G Proteins	clone 810	US20110189171; US7879329 SEQ ID NO: 99	5554
RSV333	light chain, F and G Proteins	clone 811	US20110189171; US7879329 SEQ ID NO: 100	5555
RSV334	light chain, F and G Proteins	clone 812	US20110189171; US7879329 SEQ ID NO: 101	5556
RSV335	light chain, F and G Proteins	clone 814	US20110189171; US7879329 SEQ ID NO: 102	5557
RSV336	light chain, F and G Proteins	clone 816	US20110189171; US7879329 SEQ ID NO: 103	5558
RSV337	light chain, F and G Proteins	clone 817	US20110189171; US7879329 SEQ ID NO: 104	5559
RSV338	light chain, F and G Proteins	clone 818	US20110189171; US7879329 SEQ ID NO: 105	5560
RSV339	light chain, F and G Proteins	clone 819	US20110189171; US7879329 SEQ ID NO: 106	5561
RSV340	light chain, F and G Proteins	clone 824	US20110189171; US7879329 SEQ ID NO: 107	5562
RSV341	light chain, F and G Proteins	clone 825	US20110189171; US7879329 SEQ ID NO: 108	5563
RSV342	light chain, F and G Proteins	clone 827	US20110189171; US7879329 SEQ ID NO: 109	5564

RSV343	light chain, F and G Proteins	clone 829	US20110189171; US7879329 SEQ ID NO: 110	5565
RSV344	light chain, F and G Proteins	clone 830	US20110189171; US7879329 SEQ ID NO: 111	5566
RSV345	light chain, F and G Proteins	clone 831	US20110189171; US7879329 SEQ ID NO: 112	5567
RSV346	light chain, F and G Proteins	clone 835	US20110189171; US7879329 SEQ ID NO: 113	5568
RSV347	light chain, F and G Proteins	clone 838	US20110189171; US7879329 SEQ ID NO: 114	5569
RSV348	light chain, F and G Proteins	clone 841	US20110189171; US7879329 SEQ ID NO: 115	5570
RSV349	light chain, F and G Proteins	clone 853	US20110189171; US7879329 SEQ ID NO: 116	5571
RSV350	light chain, F and G Proteins	clone 855	US20110189171; US7879329 SEQ ID NO: 117	5572
RSV351	light chain, F and G Proteins	clone 856	US20110189171; US7879329 SEQ ID NO: 118	5573
RSV352	light chain, F and G Proteins	clone 857	US20110189171; US7879329 SEQ ID NO: 119	5574
RSV353	light chain, F and G Proteins	clone 858	US20110189171; US7879329 SEQ ID NO: 120	5575
RSV354	light chain, F and G Proteins	clone 859	US20110189171; US7879329 SEQ ID NO: 121	5576
RSV355	light chain, F and G Proteins	clone 861	US20110189171; US7879329 SEQ ID NO: 122	5577
RSV356	light chain, F and G Proteins	clone 863	US20110189171; US7879329 SEQ ID NO: 123	5578
RSV357	light chain, F and G Proteins	clone 868	US20110189171; US7879329 SEQ ID NO: 124	5579
RSV358	light chain, F and G Proteins	clone 870	US20110189171; US7879329 SEQ ID NO: 125	5580
RSV359	light chain, F and G Proteins	clone 871	US20110189171; US7879329 SEQ ID NO: 126	5581
RSV360	light chain, F and G Proteins	clone 880	US20110189171; US7879329 SEQ ID NO: 127	5582
RSV361	light chain, F and G Proteins	clone 881	US20110189171; US7879329 SEQ ID NO: 128	5583
RSV362	light chain, F and G Proteins	clone 884	US20110189171; US7879329 SEQ ID NO: 129	5584
RSV363	light chain, F and G Proteins	clone 886	US20110189171; US7879329 SEQ ID NO: 130	5585
RSV364	light chain, F and G Proteins	clone 888	US20110189171; US7879329 SEQ ID NO: 131	5586
RSV365	light chain, F and G Proteins	clone 894	US20110189171; US7879329 SEQ ID NO: 132	5587
RSV366	Light chain variable, F protein of RSV, MPV, or PVM	3210 variant 1, 3210 variant 2, 3210 variant 5	WO2013140247 SEQ ID NO: 14	5588
RSV367	Light chain variable, F protein of RSV, MPV, or PVM	2430 variant 1, 2430 variant 2, 2430 variant 4	WO2013140247 SEQ ID NO: 30	5589
RSV368	Light chain variable, F protein of RSV, MPV, or PVM	3210 variant 3	WO2013140247 SEQ ID NO: 37	5590
RSV369	Light chain variable, F protein of RSV, MPV, or PVM	3210 variant 4, 3210 variant 6	WO2013140247 SEQ ID NO: 50	5591
RSV370	Light chain variable, F protein of RSV, MPV, or PVM	2430 variant 3, 2430 variant 5	WO2013140247 SEQ ID NO: 60	5592

RSV371	Light chain, F Protein	clone 19	**EP1259547;** US8153 133SEQ ID **NO:** 40	5593
RSV372	Light chain variable region, **CDR** Grafted, F Protein		**US20140093501** SEQ ID NO: 20	5594
RSV373	Light **chain variable** region, CDR Grafted, **F Protein**		US20140093501 SEQ **ID NO:** 34	5595
RSV374	Light chain, F Protein	**AM22**	**US8568726** SEQ ID NO: 32	5596
RSV375	Light **chain,** F Protein	**RSVF2-5**	**US822 1759** SEQ ID **NO:** 9	5597
RSV376	Light chain, F Protein		**EP1259547; US8153 133** SEQ ID NO: **3**	5598
RSV377	Light chain, F Protein	**MEDI-493/Pavilizumab-N-VL** (Brand name Synagis)	**EP1259547;** 1JS8153 133 SEQ ID NO: 1	5599
RSV378	Light **chain,** F Protein		EP1259547; US8153 133 SEQ ID NO: 35	5600
RSV379	Light chain, F Protein	ciorse 18	**EP1259547; US8153 133** SEQ ID NO: 38	5601
RSV380	Light **chain,** F Protein	clone 20	EP 1259547; US8153 133 SEQ ID NO: 42	5602
RSV381	Light chain, F Protein	ciorse 21	**EP1259547; US8153 133** SEQ ID NO: 44	5603
RSV382	Light **chain,** F Protein	clone 22	EP 1259547; US8153 133 SEQ ID NO: 46	5604
RSV383	Light chain, F Protein	ciorse 23	**EP1259547; US8153 133** SEQ ID NO: 48	5605
RSV384	Light **chain,** F Protein	clone 24	EP 1259547; US8153 133 SEQ ID NO: 50	5606
RSV385	Light chain, F Protein	ciorse 25	**EP1259547; US8153 133** SEQ ID NO: 52	5607
RSV386	Light chain, F Protein	clone 26	**EP1259547;** 1JS8153 133 SEQ ID NO: 54	5608
RSV387	Light chain variable region, **F** Protein	**huK l02**	**US20140093501** SEQ **ID** NO: 19	5609
RSV388	Light chain variable region, F Protein	**huK l02**	**US2014009350** 1 SEQ **ID** NO: 33	5610
RSV389	Light chain variable region, **F** Protein	**RSV** G8	**US7867497** SEQ ID NO: 4	**5611**
RSV390	Light chain variable region, F Protein	Clone 1	**US20120135006** SEQ **ID** NO: 17	5612
RSV391	Light chain variable region, **F** Protein	Clone 2	1JS20120135006 SEQ **ID** NO: 19	5613
RSV392	Light chain variable region, F Protein	Clone **3**	**US20120135006** SEQ **ID** NO: 21	5614
RSV393	Light chain variable region, **F** Protein	Clone 22	1JS20120135006 SEQ **ID** NO: 23	5615
RSV394	Light chain variable region, F Protein	Clone 23	**US20120135006** SEQ **ID** NO: 25	5616
RSV395	Light chain variable region, F Protein	**RSV13-9**	**WO2009088159** SEQ ID NO: 2	5617
RSV396	Light chain variable region, F Protein	MAb1308F	**US20140093501** SEQ ID NO: 21	5618
RSV397	Light chain, F Protein	**58c5**	**US20 1400447 19** SEQ ID NO: 5	5619
RSV398	Light chain, F Protein	sc5	**US20140044719** SEQ ID NO: **13**	5620
RSV399	Light **chain,** F Protein	Clone No. 735	**US20 120009623** SEQ ID NO: 89	5621

RSV400	Light chain, F Protein	Clone No. 736	US20 120009623 SEQ ID NO: 90	5622
RSV40 1	Light chain, F Protein	Clone No. 744	US20 120009623 SEQ ID NO: 9 1	5623
RSV402	Light chain, F Protein	Clone No. 793	US20 120009623 SEQ ID NO: 92	5624
RSV403	Light chain, F Protein	Clone No. 795	US20 120009623 SEQ ID NO: 93	5625
RSV404	Light chain, F Protein	Clone No. 796	US20 120009623 SEQ ID NO: 94	5626
RSV405	Light chain, F Protein	Clone No. 799	US20 [20009623 SEQ ID NO: 95	5627
RSV406	Light chain, F Protein	Clone No. 800	US20 120009623 SEQ ID NO: 96	5628
RSV407	Light chain, F Protein	Clone No. 801	US20 [20009623 SEQ ID NO: 97	5629
RSV408	Light chain, F Protein	Clone No. 804	US20 120009623 SEQ ID NO: 98	5630
RSV409	Light chain, F Protein	Clone No. 8 10	US20 [20009623 SEQ ID NO: 99	563 !
RSV4 10	Light chain, F Protein	Clone No. 8 11	US20 120009623 SEQ ID NO: 100	5632
RSV41 1	Light chain, F Protein	Clone No. 812	US20 120009623 SEQ ID NO: 101	5633
RSV412	Light chain, F Protein	Clone No. 814	US20 120009623 SEQ ID NO: 102	5634
RSV413	Light chain, F Protein	Clone No. 816	US20 120009623 SEQ ID NO: 103	5635
RSV414	Light chain, F Protein	Clone No. 817	US20 120009623 SEQ ID NO: 104	5636
RSV415	Light chain, F Protein	Clone No. 818	US20 120009623 SEQ ID NO: 105	5637
RSV416	Light chain, F Protein	Clone No. 819	US20 120009623 SEQ ID NO: 106	5638
RSV417	Light chain, F Protein	Clone No. 824	US20 120009623 SEQ ID NO: 107	5639
RSV418	Light chain, F Protein	Clone No. 825	US20 120009623 SEQ ID NO: 108	5640
RSV419	Light chain, F Protein	Clone No. 827	US20 120009623 SEQ ID NO: 109	5641
RSV420	Light chain, F Protein	Clone No. 829	US20 120009623 SEQ ID NO: 110	5642
RSV421	Light chain, F Protein	Clone No. 830	US20 120009623 SEQ ID NO: 111	5643
RSV422	Light chain, F Protein	Clone No. 83 1	US20 120009623 SEQ ID NO: 112	5644
RSV423	Light chain, F Protein	Clone No. 835	US20 120009623 SEQ ID NO: 113	5645
RSV424	Light chain, F Protein	Clone No. 838	US20 120009623 SEQ ID NO: 114	5646
RSV425	Light chain, F Protein	Clone No. 841	US20 120009623 SEQ ID NO: 115	5647
RSV426	Light chain, F Protein	Clone No. 853	US20 120009623 SEQ ID NO: 116	5648
RSV427	Light chain, F Protein	Clone No. 855	US20 120009623 SEQ ID NO: 117	5649
RSV428	Light chain, F Protein	Clone No. 856	US20 120009623 SEQ ID NO: 118	5650

RSV429	Light chain, F Protein	Clone No. 857	US20120009623 SEQ ID NO: 119	5651
RSV430	Light chain, F Protein	Clone No. 858	US20120009623 SEQ ID NO: 120	5652
RSV431	Light chain, F Protein	Clone No. 859	US20120009623 SEQ ID NO: 121	5653
RSV432	Light chain, F Protein	Clone No. 861	US20120009623 SEQ ID NO: 122	5654
RSV433	Light chain, F Protein	Clone No. 863	US20120009623 SEQ ID NO: 123	5655
RSV434	Light chain, F Protein	Clone No. 868	US20120009623 SEQ ID NO: 124	5656
RSV435	Light chain, F Protein	Clone No. 870	US20120009623 SEQ ID NO: 125	5657
RSV436	Light chain, F Protein	Clone No. 871	US20120009623 SEQ ID NO: 126	5658
RSV437	Light chain, F Protein	Clone No. 880	US20120009623 SEQ ID NO: 127	5659
RSV438	Light chain, F Protein	Clone No. 881	US20120009623 SEQ ID NO: 128	5660
RSV439	Light chain, F Protein	Clone No. 884	US20120009623 SEQ ID NO: 129	5661
RSV440	Light chain, F Protein	Clone No. 886	US20120009623 SEQ ID NO: 130	5662
RSV441	Light chain, F Protein	Clone No. 888	US20120009623 SEQ ID NO: 131	5663
RSV442	Light chain, F Protein	Clone No. 894	US20120009623 SEQ ID NO: 132	5664
RSV443	Light chain, F Protein		US20110027294 SEQ ID NO: 63	5665
RSV444	Light chain, F Protein		US20110027294 SEQ ID NO: 64	5666
RSV445	Light chain, F Protein		US20110027294 SEQ ID NO: 65	5667
RSV446	Light chain, F Protein		US20110027294 SEQ ID NO: 66	5668
RSV447	Light chain, F Protein		US20110027294 SEQ ID NO: 67	5669
RSV448	Light chain, F Protein		US20110027294 SEQ ID NO: 68	5670
RSV449	Light chain, F Protein		US20110027294 SEQ ID NO: 69	5671
RSV450	Light chain, F Protein		US20110027294 SEQ ID NO: 70	5672
RSV451	Light chain, F Protein		US20110027294 SEQ ID NO: 71	5673
RSV452	Light chain, F Protein		US20110027294 SEQ ID NO: 72	5674
RSV453	Light chain, F Protein		US20110027294 SEQ ID NO: 73	5675
RSV454	Light chain, F Protein		US20110027294 SEQ ID NO: 81	5676
RSV455	Light chain, F Protein		US20110027294 SEQ ID NO: 82	5677
RSV456	Light chain, F Protein		US20110027294 SEQ ID NO: 83	5678
RSV457	Light chain, F Protein		US20110027294 SEQ ID NO: 84	5679

RSV458	Light **chain,** F Protein		**US20 110027294** SEQ ID NO: 85	5680
RSV459	Light chain, F Protein		**US20 110027294** SEQ ID NO: 86	568
RSV460	Light chain, F Protein		**US201 10027294** SEQ **ID NO:** 87	5682
RSV46 !	Light chain, F Protein		US20 i 10027294 SEQ ID NO: 88	5683
RSV462	Light chain, F Protein		**US201 10027294** SEQ **ID NO:** 89	5684
RSV463	Light chain, F Protein		US20 i 10027294 SEQ ID NO: 90	5685
RSV464	Light chain, F Protein		**US201 10027294** SEQ **ID NO:** 91	5686
RSV465	Light chain, F Protein		US20 i 10027294 SEQ ID NO: 92	5687
RSV466	Light chain, F Protein		**US201 10027294** SEQ **ID NO: 93**	5688
RSV467	Light chain, F Protein		US20 i 10027294 SEQ ID NO: 94	5689
RSV468	Light chain, F Protein		**US201 10027294** SEQ **ID NO:** 95	5690
RSV469	Light chain, F Protein		**US20 110027294** SEQ **ID** NO: 96	569
RSV470	Light chain, F Protein		**US20 10027294** SEQ **ID** NO: 97	5692
RSV47	Light chain, F Protein		**US20 110027294** SEQ **ID** NO: 98	5693
RSV472	Light chain, F Protein		**US20 10027294** SEQ **ID** NO: 99	5694
RSV473	Light chain, F Protein		**US20 110027294** SEQ **ID** NO: 100	5695
RSV474	Light chain, F Protein		**US20 10027294** SEQ **ID** NO: 101	5696
RSV475	Light chain, F Protein		**US20 110027294** SEQ **ID** NO: 102	5697
RSV476	Light chain, F Protein		**US20 10027294** SEQ **ID** NO: 103	5698
RSV477	Light chain, F Protein		**US20 10027294** SEQ ID NO: 104	5699
RSV478	Light chain, F Protein		**US20 10027294** SEQ ID NO: 105	5700
RSV479	Light chain, F Protein		**US20 10027294** SEQ ID NO: 106	5701
RSV480	Light chain, F Protein		**US20 10027294** SEQ ID NO: 107	5702
RSV481	Light chain, F Protein		**US20 10027294** SEQ ID NO: 108	5703
RSV482	Light chain, F Protein		**US20 10027294** SEQ ID NO: 109	5704
RSV483	Light chain, F Protein		**US20 10027294** SEQ ID NO: **110**	5705
RSV484	Light chain, F Protein		**US20 10027294** SEQ ID NO: **111**	5706
RSV485	Light chain, F Protein		**US20 10027294** SEQ ID NO: **112**	5707
RSV486	Light chain, F Protein	Gλ-1A	US20050175986 SEQ ID NO: 9	5708
RSV487	Light **chain,** F Protein	A **construct**	**US20050 175986** SEQ ID NO: 11	5709

RSV488	Light chain, F Protein	B construct	US20050175986 SEQ ID NO: 12	5710
RSV489	Light chain, F Protein	hu19A	US20050019758; WO1998019704 SEQ ID NO: 10	5711
RSV490	Light chain, F Protein	hu19B	US20050019758; WO1998019704 SEQ ID NO: 11	5712
RSV491	Light chain, F Protein	hu19C	US20050019758; WO1998019704 SEQ ID NO: 12	5713
RSV492	Light chain, F Protein	hu19D	US20050019758; WO1998019704 SEQ ID NO: 13	5714
RSV493	Light chain, F Protein	RSV19	EP636182; WO1993020210; SEQ ID NO: 12	5715
RSV494	Light chain, F Protein		WO19922004381	5716
RSV495	Light chain variable region, F Protein	P1212	US20140044719 SEQ ID NO: 127	5717
RSV496	Light chain variable region, F Protein	P12f4	US20140044719 SEQ ID NO: 134	5718
RSV497	Light chain variable region, F Protein	P11d4	US20140044719 SEQ ID NO: 140	5719
RSV498	Light chain variable region, F Protein	A1e9	US20140044719 SEQ ID NO: 146	5720
RSV499	Light chain variable region, F Protein	A12a6	US20140044719 SEQ ID NO: 152	5721
RSV500	Light chain variable region, F Protein	A13c4	US20140044719 SEQ ID NO: 158	5722
RSV501	Light chain variable region, F Protein	A17d4	US20140044719 SEQ ID NO: 164	5723
RSV502	Light chain variable region, F Protein	A4B4	US20140044719 SEQ ID NO: 169	5724
RSV503	Light chain variable region, F Protein	A8c7	US20140044719 SEQ ID NO: 174	5725
RSV504	Light chain variable region, F Protein	IX-493L1FR	US20140044719 SEQ ID NO: 178	5726
RSV505	Light chain variable region, F Protein	M3H9	US20140044719 SEQ ID NO: 180	5727
RSV506	Light chain variable region, F Protein	B21M	US20110027294 SEQ ID NO: 51	5728
RSV507	Light chain variable region, F Protein	101F	US20110027294 SEQ ID NO: 6	5729
RSV508	Light chain variable region, F Protein	HNK20	EP1720908; WO2005079479 SEQ ID NO: 2	5730
RSV509	Light chain variable region, F Protein	P1212	US20140044719 SEQ ID NO: 128	5731
RSV510	Light chain variable region, F Protein	P12f4	US20140044719 SEQ ID NO: 135	5732
RSV511	Light chain variable region, F Protein	P11d4	US20140044719 SEQ ID NO: 141	5733
RSV512	Light chain variable region, F Protein	A1e9	US20140044719 SEQ ID NO: 147	5734
RSV513	Light chain variable region, F Protein	A12a6	US20140044719 SEQ ID NO: 153	5735
RSV514	Light chain variable region, F Protein	A13c4	US20140044719 SEQ ID NO: 159	5736
RSV515	Light chain variable region, F Protein	A17d4	US20140044719 SEQ ID NO: 165	5737

RSV5 16	Light chain variable region, **F** Protein	A4B4	**US20140044719** SEQ ID NO: **170**	5738
RSV517	Light chain variable region, **F** Protein	A8c7	**US201400447 19** SEQ ID NO: 175	5739
RSV5 18	Light chai n variable regiors, **F** Protein	**LX-493L IFR**	**US20140044719** SEQ ID **NO:** 179	5740
RSV519	Light chain variable region, **F** Protein	H l **H3564P**	**WO2014159822** SEQ ID NO: 10	574 l
RSV520	Light chain variable regiors, **F** Protein	H l **H3565P**	**WO20 l4 l**59822 SEQ ID **NO:** 26	5742
RSV52 !	Light chain variable region, **F** Protein	H l H3566P	**WO2014159822** SEQ ID NO: **42**	5743
RSV522	Light chain variable regiors, **F** Protein	H l **H3567P**	**WO20 l4 l**59822 SEQ ID **NO:** 58	5744
RSV523	Light chain variable region, **F** Protein	**H l H3581 P**	**WO2014159822** SEQ ID NO: **74**	5745
RSV524	Light chain variable regiors, **F** Protein	H l **H3583P**	**WO20 l4 l**59822 SEQ ID **NO:** 90	5746
RSV525	Light chain variable region, **F** Protein	H l **H3589P**	**WO2014159822** SEQ ID NO: 106	5747
RSV526	Light chain variable regiors, **F** Protein	H l H359 l P	**WO20 l4 l**59822 SEQ ID **NO:** 122	5748
RSV527	Light chain variable region, **F** Protein	H i H3592P	**WO20 14 159822** SEQ ID NO: 138	5749
RSV528	Light chain variable region, **F** Protein	H l **H3597P**	**WO20 14 159822** SEQ ID NO: 154	5750
RSV529	Light chain variable region, **F** Protein	H i H3598P	**WO20 14 159822** SEQ ID NO: 170	575 **1**
RSV530	Light chain variable region, **F** Protein	H l **H3603P**	**WO20 14 159822** SEQ ID NO: 186	5752
RSV53 **1**	Light chain variable region, **F** Protein	H i H3604P	**WO20 14 159822** SEQ ID NO: 202	5753
RSV532	Light chain variable region, **F** Protein	H l **H3605P**	**WO20 14 159822** SEQ ID NO: 218	5754
RSV533	Light chain variable region, **F** Protein	H i H3607P	**WO20 14 159822** SEQ ID NO: 234	5755
RSV534	Light chain variable region, **F** Protein	**H 1 H3608P2**	**WO20 14 159822** SEQ ID NO: 250	5756
RSV535	Light chain variable region, **F** Protein	H l H3592P2	**WO2014159822** SEQ ID NO: 266	5757
RSV536	Light chain variable region, **F** Protein	H l H3592P3	**WO20 14 159822** SEQ ID NO: 282	5758
RSV537	Light chain variable region, **F** Protein	**H 1 M3621** N	**WO2014159822** SEQ ID NO: 298	5759
RSV538	Light chain variable region, **F** Protein	H l **M3622N**	**WO20 14 159822** SEQ ID NO: **3 14**	5760
RSV539	Light chain variable region, **F** Protein	H l **M2634N**	**WO2014159822** SEQ ID NO: 330	5761
RSV540	Light chain variable region, **F** Protein	H l M3627N	**WO20 14 159822** SEQ ID NO: 346	5762
RSV541	Light chain variable region, **F** Protein	Gλ-1	**US20050175986** SEQ ID NO: 2	5763
RSV542	Light chain variable region, **F** Protein	**MAbl l29**	**US20140093501** SEQ ID NO: 35	5764
RSV543	super humanized kappa light chain based on HNK20, **F** protein	**SHV11**	EP 1720908; **WO2005079479** SEQ ID NO: **10**	5765

RSV544	super humanized kappa light chairs based on HNK20, F protein	SHV12	EP1720908; WO2005079479 SEQ ID NO: 11	5766
RSV545	super humanized kappa light chain based on HNK20, F protein	SHV13	EP1720908; WO2005079479 SEQ ID NO: 12	5767
RSV546	super humanized kappa light chain based on HNK20, F protein	SHV14	EP1720908; WO2005079479 SEQ ID NO: 13	5768
RSV547	super humanized kappa light chain based on HNK20, F protein	SHV15	EP1720908; WO2005079479 SEQ ID NO: 14	5769
RSV548	super humanized kappa light chain based on HNK20. F protein	SHV16	EP1720908; WO2005079479 SEQ ID NO: 15	5770
RSV549	Light chain variable region, F Protein	B4	EP636182; WO1993020210; SEQ ID NO: 1	5771
RSV550	Light chain variable region, F Protein	B 13/14	EP636182; WO1993020210; SEQ ID NO: 2	5772
RSV551	Light chain variable region, F Protein	RF-1	EP854730; WO1996040252; FIG 7A	5773
RSV552	Light chain variable region, F Protein	RF-2	EP854730; WO1996040252; FIG 8A	5774
RSV553	Light chain variable region Kappa, G protein	CB058.1	WO2014170257 SEQ ID NO: 38	5775
RSV554	Light chain variable region Kappa, G protein	CB048.3	WO2014170257 SEQ ID NO: 40	5776
RSV555	Light chain variable region Kappa, G protein	CB010.7	WO2014170257 SEQ ID NO: 42	5777
RSV556	Light chain variable region Kappa, G protein	CB003.1	WO2014170257 SEQ ID NO: 44	5778
RSV557	Light chain variable region Kappa, G protein	CB028.2	WO2014170257 SEQ ID NO: 46	5779
RSV558	Light chain variable region Kappa, G protein	CB002.1	WO2014170257 SEQ ID NO: 48	5780
RSV559	Light chain, G Protein	1F12	US8273354 SEQ ID NO: 42	5781
RSV560	Light chain, G Protein	3G12	US8273354 SEQ ID NO: 43	5782
RSV561	Light chain, G Protein	1A5	US8273354 SEQ ID NO: 44	5783
RSV562	Light chain, G Protein	3D3	US8273354 SEQ ID NO: 45	5784
RSV563	Light chain, G Protein	1G1	US8273354 SEQ ID NO: 46	5785
RSV564	Light chain, G Protein	2B11	US8273354 SEQ ID NO: 47	5786
RSV565	Light chain, G Protein	5D8	US8273354 SEQ ID NO: 48	5787
RSV566	Light chain, G Protein	2D so	US8273354 SEQ ID NO: 49	5788
RSV567	Light chain, G Protein	3F9	US8273354 SEQ ID NO: 50	5789
RSV568	Light chain, G Protein	1D4	US8273354 SEQ ID NO: 51	5790
RSV569	Light chain, G Protein	1G8	US8273354 SEQ ID NO: 52	5791
RSV570	Light chain, G Protein	6A12	US8273354 SEQ ID NO: 53	5792
RSV571	Light chain, G Protein	10C6	US8273354 SEQ ID NO: 54	5793
RSV572	Light chain, G Protein	Hu 13 1-2G	US8273354 SEQ ID NO: 55	5794
RSV573	Light chain, G Protein	AT46	US20150004155 SEQ ID NO: 127	5795
RSV574	Light chain, G Protein	AT32	US20150004155 SEQ ID NO: 128	5796
RSV575	Light chain, G Protein	AT33	US20150004155 SEQ ID NO: 129	5797

RSV576	Light chain, G Protein	AT34	US20150004155 SEQ ID NO: 130	5798
RSV577	Light chain, G Protein	AT35	US20150004155 SEQ ID NO: 131	5799
RSV578	Light chain, G Protein	AT36	US20150004155 SEQ ID NO: 132	5800
RSV579	Light chain, G Protein	AT37	US20150004155 SEQ ID NO: 133	5801
RSV580	Light chain, G Protein	AT39	US20150004155 SEQ ID NO: 134	5802
RSV581	Light chain, G Protein	AT40	US20150004155 SEQ ID NO: 135	5803
RSV582	Light chain, G Protein	AT42	US20150004155 SEQ ID NO: 136	5804
RSV583	Light chain, G Protein	AT43	US20150004155 SEQ ID NO: 137	5805
RSV584	Light chain, G Protein	AT44	US20150004155 SEQ ID NO: 138	5806
RSV585	Light chain, G Protein	AT45	US20150004155 SEQ ID NO: 139	5807
RSV586	Light chain, G Protein	AT47	US20150004155 SEQ ID NO: 140	5808
RSV587	Light chain, G Protein	AT49	US20150004155 SEQ ID NO: 141	5809
RSV588	Light chain, G Protein	AT50	US20150004155 SEQ ID NO: 142	5810
RSV589	Light chain, G Protein	AT51	US20150004155 SEQ ID NO: 143	5811
RSV590	Light chain variable region, G Protein	CB017.3L	WO2014170258 SEQ ID NO: 74	5812
RSV591	Light chain variable region, G Protein	CB017.5L	WO2014170258 SEQ ID NO: 76	5813
RSV592	Light chain variable region, G Protein	CB028.1	WO2014170258 SEQ ID NO: 78	5814
RSV593	Light chain variable region, G Protein	CB030.1	WO2014170258 SEQ ID NO: 80	5815
RSV594	Light chain variable region, G Protein	CB047.1	WO2014170258 SEQ ID NO: 82	5816
RSV595	Light chain variable region, G Protein	CB047.2	WO2014170258 SEQ ID NO: 84	5817
RSV596	Light chain variable region, G Protein	CB065.1	WO2014170258 SEQ ID NO: 86	5818
RSV597	Light chain variable region, G Protein	CB071.1L	WO2014170258 SEQ ID NO: 88	5819
RSV598	Light chain variable region, G Protein	CB072.1L	WO2014170258 SEQ ID NO: 90	5820
RSV599	Light chain variable region, G Protein	CB073.1L	WO2014170258 SEQ ID NO: 92	5821
RSV600	Light chain variable region, G Protein	CB076.2L	WO2014170258 SEQ ID NO: 94	5822
RSV601	Light chain variable region, G Protein	CB079.1	WO2014170258 SEQ ID NO: 96	5823
RSV602	Light chain, human metapneumovirus fusion protein with neutralizing antibody identifies a pneumovirus antigenic site		WeiiX., "Structure of the human metapneumovirus fusion protein with neutralizing antibody identifies a pneumovirus antigenic site", Nat. Struct. Mol. Biol. 19 (4),	5824

			461-463 (2012), NCBI Accession # 4DAG_L(213 aa)	
RSV603	Light chain	AM14	US20140377279 SEQ ID NO: 79	5825
RSV604	Light chain	AM16	US20140377279 SEQ ID NO: 86	5826
RSV605	Light chain	AM23	US20140377279 SEQ ID NO: 93	5827
RSV606	Light chain	D25	US20140377279 SEQ ID NO: 8	5828
RSV607	Light chain	AFFF	1JS7635568 SEQ ID NO: 211	5829
RSV608	Light chain	P12f2	US7635568 SEQ ID NO: 213	5830
RSV609	Light chain	P12f4	US7635568 SEQ ID NO: 215	5831
RSV610	Light chain	P11d4	US7635568 SEQ ID NO: 217	5832
RSV611	Light chain	Ale9	US7635568 SEQ ID NO: 219	5833
RSV612	Light chain	A12a6	US7635568 SEQ ID NO: 221	5834
RSV613	Light chain	A13c4	US7635568 SEQ ID NO: 223	5835
RSV614	Light chain	A17d4	US7635568 SEQ ID NO: 225	5836
RSV615	Light chain	A4B4	US7635568 SEQ ID NO: 227	5837
RSV616	Light chain	A8c7	US7635568 SEQ ID NO: 229	5838
RSV617	Light chain	1X-493L1FR	US7635568 SEQ ID NO: 231	5839
RSV618	Light chain	H3-3F4	US7635568 SEQ ID NO: 233	5840
RSV619	Light chain	M3H9	US7635568 SEQ ID NO: 235	5841
RSV620	Light chain	Y10H6	US7635568 SEQ ID NO: 237	5842
RSV621	Light chain	DG	US7635568 SEQ ID NO: 239	5843
RSV622	Light chain	AFFF(1)	US7635568 SEQ ID NO: 241	5844
RSV623	Light chain	6H8	US7635568 SEQ ID NO: 243	5845
RSV624	Light chain	L1-7E5	1JS7635568 SEQ ID NO: 245	5846
RSV625	Light chain	L2-15B10	US7635568 SEQ ID NO: 247	5847
RSV626	Light chain	A13al 1	US7635568 SEQ ID NO: 249	5848
RSV627	Light chain	A1h5	US7635568 SEQ ID NO: 251	5849
RSV628	Light chain	A4B4(1)	US7635568 SEQ ID NO: 253	5850
RSV629	Light chain	A4B4L 1FR-S28R	1JS7635568 SEQ ID NO: 255	5851
RSV630	Light chain	A4B4-F52S	US7635568 SEQ ID NO: 257	5852
RSV631	Light chain variable region	AFFF	US7635568 SEQ ID NO: 13	5853
RSV632	Light chain variable region	P12f2	US7635568 SEQ ID NO: 21	5854
RSV633	Light chain variable region	P12f4	US7635568 SEQ ID NO: 26	5855
RSV634	Light chain variable region	P11d4	1JS7635568 SEQ ID NO: 30	5856
RSV635	Light chain variable region	Ale9	US7635568 SEQ ID NO: 34	5857
RSV636	Light chain variable region	A12a6	US7635568 SEQ ID NO: 38	5858
RSV637	Light chain variable region	A13c4	US7635568 SEQ ID NO: 42	5859
RSV638	Light chain variable region	A17d4	US7635568 SEQ ID NO: 46	5860
RSV639	Light chain variable region	A4B4	US7635568 SEQ ID NO: 49	5861
RSV640	Light chain variable region	A8c7	US7635568 SEQ ID NO: 52	5862
RSV641	Light chain variable region	1X-493L1FR	US7635568 SEQ ID NO: 54	5863
RSV642	Light chain variable region	H3-3F4, DG	US7635568 SEQ ID NO: 56	5864
RSV643	Light chain variable region	M3H9	US7635568 SEQ ID NO: 70	5865
RSV644	Light chain variable region	Y10H6	1JS7635568 SEQ ID NO: 58	5866

RSV645	Light chain variable region	AFFF(1)	US7635568 SEQ ID NO: 60	5867
RSV646	Light chain variable region	6H8	US7635568 SEQ ID NO: 62	5868
RSV647	Light chain variable region	L1-7E5	US7635568 SEQ ID NO: 64	5869
RSV648	Light chain variable region	L2-15B10	US7635568 SEQ ID NO: 65	5870
RSV649	Light chain variable region	A13a11	US7635568 SEQ ID NO: 68	5871
RSV650	Light chain variable region	A1h5	US7635568 SEQ ID NO: 71	5872
RSV651	Light chain variable region	A4B4(1)	US7635568 SEQ ID NO: 74	5873
RSV652	Light chain variable region	A4B4L1FR-S28R	US7635568 SEQ ID NO: 11	5874
RSV653	Light chain variable region	A4B4-F52S	US7635568 SEQ ID NO: 76	5875
RSV654	Light chain variable region	6H; 11H; 21H; 22H; and 23H	US7364737 SEQ ID NO: 21	5876
RSV655	Light chain variable region	13H and 19H	US7364737 SEQ ID NO: 22	5877
RSV656	Light chain variable region	6L; 11L; 21L; and 22L	US7364737 SEQ ID NO: 23	5878
RSV657	Light chain variable region	23L	US7364737 SEQ ID NO: 24	5879
RSV658	Light chain variable region	13L and 19L	US7364737 SEQ ID NO: 25	5880
RSV659	Light chain variable region		US7364742 SEQ ID NO: 9	5881
RSV660	Light chain variable region		US7364742 SEQ ID NO: 10	5882
RSV661	Light chain variable region		US7364742 SEQ ID NO: 11	5883
RSV662	Light chain variable region		US7364742 SEQ ID NO: 12	5884
RSV663	Light chain variable region	D2E7	EP1807111; WO2006041970 SEQ ID NO: 1	5885
RSV664	Light chain variable region	2SD4	EP1807111; WO2006041970 SEQ ID NO: 9	5886
RSV665	Light chain variable, M2 1 antigen	8A4/G9 - IgG	US20140348858 SEQ ID NO: 4	5887
RSV666	Light chain, Pre fusion RSV F protein	HMB2435	WO2015010792 SEQ ID NO: 14	5888
RSV667	Light chain, Pre fusion RSV F protein	HMB2437	WO2015010792 SEQ ID NO: 30	5889
RSV668	Light chain, Pre fusion RSV F protein	HMB2416	WO2015010792 SEQ ID NO: 46	5890
RSV669	Light chain, Pre fusion RSV F protein	HMB2437	WO2015010792 SEQ ID NO: 86	5891
RSV670	Light chain, Pre fusion RSV F protein	CR9501	WO2014202570 SEQ ID NO: 61	5892
RSV671	Light chain, Pre fusion RSV F protein	CR9502	WO2014202570 SEQ ID NO: 65	5893
RSV672	Light chain 1, Pre fusion RSV F protein	HMB2432	WO2015010792 SEQ ID NO: 62	5894
RSV673	Light chain FR GL, Pre fusion RSV F protein	HMB2432	WO2015010792 SEQ ID NO: 66	5895
RSV674	Light chain 2, Pre fusion RSV F protein	HMB2435	WO2015010792 SEQ ID NO: 76	5896
RSV675	derived Ig variable region amino acid sequence, F protein	RSV19VH	EP636182; WO1993020210; SEQ ID NO: 13	5897
RSV676	derived Ig variable region amino acid sequence, F protein	pHuRSV19VH	EP636182; WO1993020210; SEQ ID NO: 14	5898
RSV677	derived Ig variable region amino acid sequence, F protein	pHuRSV19VHFNS	EP636182; WO1993020210; SEQ ID NO: 15	5899
RSV678	derived Ig variable region amino acid sequence, F protein	pHuRSV19VHNIK	EP636182; WO1993020210; SEQ ID NO: 16	5900
RSV679	derived Ig variable region amino acid sequence, F protein	pHuRSV19VK	EP636182; WO1993020210; SEQ ID NO: 17	5901

RSV680	**Nanobody** binding to **RSV** F **protein**	LG202A10	**US20 110 182897** SEQ ID NO: 126	5902
RSV68 1	Nanobody binding to RSV F protein	**LG202A12**	**US20 110 182897** SEQ ID NO: **127**	5903
RSV682	Nanobody binding to RSV F protein	**LG202A5**	**US201 10182897** SEQ **ID NO:** 128	5904
RSV683	Nanobody binding to RSV **F** protein	**LG202A9**	US20 [ ] 0] 82897 SEQ ID NO: 129	5905
RSV684	Nanobody binding to RSV F protein	**LG202B 10**	**US201 10182897** SEQ **ID NO: 130**	5906
RSV685	Nanobody binding to RSV **F** protein	**LG202B7**	US20 [10 182897 SEQ ID NO: **13** [	5907
RSV686	Nanobody binding to RSV F protein	**LG202B8**	**US201 10182897** SEQ **ID NO: 132**	5908
RSV687	Nanobody binding to RSV **F** protein	**LG202B9**	US20 [10 182897 SEQ ID NO: 133	5909
RSV688	Nanobody binding to RSV F protein	**LG202C 1**	**US201 10182897** SEQ **ID NO:** 134	59 10
RSV689	Nanobody binding to RSV **F** protein	**LG202C 1I**	US20 [10 182897 SEQ ID NO: **135**	59 [[
RSV690	Nanobody binding to RSV F _proiem	**LG202C2**	**US201 10182897** SEQ **ID NO:** 136	59 12
RSV69 1	Nanobody binding to RSV **F** protein	LG202C7	ıı S20 **110 182897** SEQ **ID** NO: 137	5913
RSV692	Nanobody binding to RSV F _protein	**LG202C8**	**US20 [10 182897** SEQ **ID** NO: 138	591 4
RSV693	Nanobody binding to RSV **F** protein	**LG202C9**	ıı S20 **110 182897** SEQ **ID** NO: 139	5915
RSV694	Nanobody binding to RSV F _protein	**LG202D5**	**US20 [10 182897** SEQ **ID** NO: 140	59 16
RSV695	Nanobody binding to RSV **F** protein	**LG202D7**	ıı S20 **110 182897** SEQ **ID** NO: 141	5917
RSV696	Nanobody binding to RSV F protein	**LG202D8**	**US20 [10 182897** SEQ **ID** NO: 142	591 8
RSV697	Nanobody binding to RSV **F** protein	**LG202E1 1**	ıı S20 **110 182897** SEQ **ID** NO: 143	5919
RSV698	Nanobody binding to RSV F protein	**LG202E2**	**US20 [10 182897** SEQ **ID** NO: **144**	5920
RSV699	Nanobody binding to RSV F protein	LG202E5	**US20 [10 182897** SEQ ID NO: 145	5921
RSV700	Nanobody binding to RSV F protein	**LG202E6**	**US20 [10 182897** SEQ ID NO: [46	5922
RSV701	Nanobody binding to RSV F protein	LG202E7	**US20 [10 182897** SEQ ID NO: 147	5923
RSV702	Nanobody binding to RSV F protein	**LG202F10**	**US20 [10 182897** SEQ ID NO: 148	5924
RSV703	Nanobody binding to RSV F protein	LG202F12	**US20 [10 182897** SEQ ID NO: 149	5925
RSV704	Nanobody binding to RSV F protein	**LG202F3**	**US20 [10 182897** SEQ ID NO: [50	5926
RSV705	Nanobody binding to RSV F protein	LG202F4	**US20 [10 182897** SEQ ID NO: 151	5927
RSV706	Nanobody binding to RSV F protein	**LG202F8**	**US20 [10 182897** SEQ ID NO: 152	5928
RSV707	Nanobody binding to RSV F protein	**LG202G1 1**	**US20 [10 182897** SEQ ID NO: 153	5929
RSV708	Nanobody binding to RSV F protein	**LG202G3**	**US20 [10 182897** SEQ ID NO: [54	5930

RSV709	**Nanobody** binding to **RSV** F **protein**	LG202G8	**US20 110 182897** SEQ ID NO: 155	593 1
RSV710	Nanobody binding to RSV F protein	LG202H2	**US201 10182897** SEQ ID NO: 156	5932
RSV7 ! 1	Nanobody binding to RSV F protein	**LG202H8**	**US201 10182897** SEQ **ID NO:** 157	5933
RSV712	Nanobody binding to RSV **F** protein	**LG I9 1B9**	US20 i]0]82897 SEQ ID NO: 158	**5934**
RSV7 13	Nanobody binding to RSV F protein	**LG191D3**	**US201 10182897** SEQ **ID NO:** 159	5935
RSV7]4	Nanobody binding to RSV **F** protein	**LG I92A8**	US20 i 10 182897 SEQ ID NO: **160**	5936
RSV7 15	Nanobody binding to RSV F protein	**LG192B 1**	**US201 10182897** SEQ **ID NO:** 161	5937
RSV716	Nanobody binding to RSV **F** protein	**LG I92C 10**	US20 i 10 182897 SEQ ID NO: **162**	5938
RSV7 17	Nanobody binding to RSV F protein	**LG192C4**	**US201 10182897** SEQ **ID NO:** 163	5939
RSV718	Nanobody binding to RSV **F** protein	**LG I92C6**	US20 i 10 182897 SEQ ID NO: **164**	5940
RSV7 19	Nanobody binding to RSV F _proiem	**LG192D3**	**US201 10182897** SEQ **ID NO:** 165	5941
RSV720	Nanobody binding to RSV **F** protein	**LG 19 1E4**	ıı S20 **110 182897** SEQ ID NO: 166	5942
RSV72 1	Nanobody binding to RSV F _protein	**LG 192F2**	**US20] 10 182897** SEQ **ID** NO: 167	5943
RSV722	Nanobody binding to RSV **F** protein	**LG 192H1**	ıı S20 **110 182897** SEQ ID NO: 168	5944
RSV723	Nanobody binding to RSV F _protein	**LG 192H2**	**US20] 10 182897** SEQ **ID** NO: 169	5945
RSV724	Nanobody binding to RSV **F** protein	**LG20610B**	ıı S20 **110 182897** SEQ ID NO: 170	5946
RSV725	Nanobody binding to RSV F protein	**LG20610C**	**US20] 10 182897** SEQ **ID** NO: 171	5947
RSV726	Nanobody binding to RSV **F** protein	**LG20610D**	ıı S20 **110 182897** SEQ ID NO: 172	5948
RSV727	Nanobody binding to RSV F protein	**LG20610E**	**US20] 10 182897** SEQ **ID** NO: 173	5949
RSV728	Nanobody binding to RSV F protein	**LG20610F**	**US20] 10 182897** SEQ ID NO: 174	5950
RSV729	Nanobody binding to RSV F protein	**LG2061 1D**	**US20] 10 182897** SEQ ID NO: 175	595 1
RSV730	Nanobody binding to RSV F protein	**LG2061 1H**	**US20] 10 182897** SEQ ID NO: 176	5952
RSV73 1	Nanobody binding to RSV F protein	**LG20612F**	**US20] 10 182897** SEQ ID NO: 177	5953
RSV732	Nanobody binding to RSV F protein	LG2062A	**US20] 10 182897** SEQ ID NO: 178	5954
RSV733	Nanobody binding to RSV F protein	**LG2062C**	**US20] 10 182897** SEQ ID NO: 179	5955
RSV734	Nanobody binding to RSV F protein	LG2062E	**US20] 10 182897** SEQ ID NO: 180	5956
RSV735	Nanobody binding to RSV F protein	**LG2062F**	**US20] 10 182897** SEQ ID NO: **18 1**	5957
RSV736	Nanobody binding to RSV F protein	LG2062G	**US20] 10 182897** SEQ ID NO: 182	5958
RSV737	Nanobody binding to RSV F protein	**LG2062H**	**US20] 10 182897** SEQ ID NO:]83	5959

RSV738	Nanobody binding to RSV F protein	LG2063A	US20110182897 SEQ ID NO: 184	5960
RSV739	Nanobody binding to RSV F protein	LG2063B	US20110182897 SEQ ID NO: 185	5961
RSV740	Nanobody binding to RSV F protein	LG2063C	US20110182897 SEQ ID NO: 186	5962
RSV741	Nanobody binding to RSV F protein	LG2063D	US20110182897 SEQ ID NO: 187	5963
RSV742	Nanobody binding to RSV F protein	LG2063E	US20110182897 SEQ ID NO: 188	5964
RSV743	Nanobody binding to RSV F protein	LG2063F	US20110182897 SEQ ID NO: 189	5965
RSV744	Nanobody binding to RSV F protein	LG2064D	US20110182897 SEQ ID NO: 190	5966
RSV745	Nanobody binding to RSV F protein	LG2064G	US20110182897 SEQ ID NO: 191	5967
RSV746	Nanobody binding to RSV F protein	LG2065A	US20110182897 SEQ ID NO: 192	5968
RSV747	Nanobody binding to RSV F protein	LG2065E	US20110182897 SEQ ID NO: 193	5969
RSV748	Nanobody binding to RSV F protein	LG2066A	US20110182897 SEQ ID NO: 194	5970
RSV749	Nanobody binding to RSV F protein	LG2066D	US20110182897 SEQ ID NO: 195	5971
RSV750	Nanobody binding to RSV F protein	LG2067B	US20110182897 SEQ ID NO: 196	5972
RSV751	Nanobody binding to RSV F protein	LG2067C	US20110182897 SEQ ID NO: 197	5973
RSV752	Nanobody binding to RSV F protein	LG2067E	US20110182897 SEQ ID NO: 198	5974
RSV753	Nanobody binding to RSV F protein	LG2067G	US20110182897 SEQ ID NO: 199	5975
RSV754	Nanobody binding to RSV F protein	LG2067H	US20110182897 SEQ ID NO: 200	5976
RSV755	Nanobody binding to RSV F protein	LG2071 1A	US20110182897 SEQ ID NO: 201	5977
RSV756	Nanobody binding to RSV F protein	LG2071 1B	US20110182897 SEQ ID NO: 202	5978
RSV757	Nanobody binding to RSV F protein	LG2071 1D	US20110182897 SEQ ID NO: 203	5979
RSV758	Nanobody binding to RSV F protein	LG2071 1E	US20110182897 SEQ ID NO: 204	5980
RSV759	Nanobody binding to RSV F protein	LG2071 1F	US20110182897 SEQ ID NO: 205	5981
RSV760	Nanobody binding to RSV F protein	LG2071 1G	US20110182897 SEQ ID NO: 206	5982
RSV761	Nanobody binding to RSV F protein	LG2071 1H	US20110182897 SEQ ID NO: 207	5983
RSV762	Nanobody binding to RSV F protein	LG2071A	US20110182897 SEQ ID NO: 208	5984
RSV763	Nanobody binding to RSV F protein	LG2071B	US20110182897 SEQ ID NO: 209	5985
RSV764	Nanobody binding to RSV F protein	LG2071C	US20110182897 SEQ ID NO: 210	5986
RSV765	Nanobody binding to RSV F protein	LG207D1	US20110182897 SEQ ID NO: 211	5987
RSV766	Nanobody binding to RSV F protein	LG2071E	US20110182897 SEQ ID NO: 212	5988

RSV767	Nanobody binding to RSV F protein	LG2071F	US20110182897 SEQ ID NO: 213	5989
RSV768	Nanobody binding to RSV F protein	LG2074A	US20110182897 SEQ ID NO: 214	5990
RSV769	Nanobody binding to RSV F protein	LG2074B	US20110182897 SEQ ID NO: 215	5991
RSV770	Nanobody binding to RSV F protein	LG2074D	US20110182897 SEQ ID NO: 216	5992
RSV771	Nanobody binding to RSV F protein	LG2074H	US20110182897 SEQ ID NO: 217	5993
RSV772	Nanobody binding to RSV F protein	LG2075A	US20110182897 SEQ ID NO: 218	5994
RSV773	Nanobody binding to RSV F protein	LG2075B	US20110182897 SEQ ID NO: 219	5995
RSV774	Nanobody binding to RSV F protein	LG2075C	US20110182897 SEQ ID NO: 220	5996
RSV775	Nanobody binding to RSV F protein	LG2075D	US20110182897 SEQ ID NO: 221	5997
RSV776	Nanobody binding to RSV F protein	LG2075E	US20110182897 SEQ ID NO: 222	5998
RSV777	Nanobody binding to RSV F _proiem	LG2076A	US20110182897 SEQ ID NO: 223	5999
RSV778	Nanobody binding to RSV F protein	LG2076B	US20110182897 SEQ ID NO: 224	6000
RSV779	Nanobody binding to RSV F _protein	LG2076C	US20110182897 SEQ ID NO: 225	6001
RSV780	Nanobody binding to RSV F protein	LG2076D	US20110182897 SEQ ID NO: 226	6002
RSV781	Nanobody binding to RSV F _protein	LG2076E	US20110182897 SEQ ID NO: 227	6003
RSV782	Nanobody binding to RSV F protein	LG2076F	US20110182897 SEQ ID NO: 228	6004
RSV783	Nanobody binding to RSV F protein	LG2079A	US20110182897 SEQ ID NO: 229	6005
RSV784	Nanobody binding to RSV F protein	LG2079B	US20110182897 SEQ ID NO: 230	6006
RSV785	Nanobody binding to RSV F protein	LG2079C	US20110182897 SEQ ID NO: 231	6007
RSV786	Nanobody binding to RSV F protein	LG2079D	US20110182897 SEQ ID NO: 232	6008
RSV787	Nanobody binding to RSV F protein	LG2G79E	US20110182897 SEQ ID NO: 233	6009
RSV788	Nanobody binding to RSV F protein	LG2079F	US20110182897 SEQ ID NO: 234	6010
RSV789	Nanobody binding to RSV F protein	LG2079G	US20110182897 SEQ ID NO: 235	6011
RSV790	Nanobody binding to RSV F protein	LG2079H	US20110182897 SEQ ID NO: 236	6012
RSV791	Nanobody binding to RSV F protein	LG213B7	US20110182897 SEQ ID NO: 237	6013
RSV792	Nanobody binding to RSV F protein	LG213D6	US20110182897 SEQ ID NO: 238	6014
RSV793	Nanobody binding to RSV F protein	LG213D7	US20110182897 SEQ ID NO: 239	6015
RSV794	Nanobody binding to RSV F protein	LG213E6	US20110182897 SEQ ID NO: 240	6016
RSV795	Nanobody binding to RSV F protein	LG213H7	US20110182897 SEQ ID NO: 241	6017

RSV796	**Nanobody** binding to **RSV F protein**	**LG214A8**	**US20 110 182897** SEQ ID NO: 242	60 8
RSV797	Nanobody binding to RSV **F** protein	**LG214C10**	**US201 10182897** SEQ ID NO: 24.3	6019
RSV798	Nanobody binding to RSV **F** protein	**LG2 4D 0**	**US201 10182897** SEQ **ID NO:** 244	6020
RSV799	Nanobody binding to RSV **F** protein	**LG214E8**	US20 10182897 SEQ ID NO: 245	602
RSV800	Nanobody binding to RSV **F** protein	**LG2 14F8**	**US201 10182897** SEQ **ID NO:** 246	6022
RSV80 !	Nanobody binding to RSV **F** protein	**LG214H 0**	US20 10 182897 SEQ ID NO: 247	6023
RSV802	Nanobody binding to RSV **F** protein	**RSVPMP5C I**	**US201 10182897** SEQ **ID NO:** 248	6024
RSV803	Nanobody binding to RSV **F** protein	**RSVPMP8A1**	**US201** 10 82897 SEQ **ID** NO: 249	6025
RSV804	Nanobody binding to RSV **F** protein	**RSVPMP8G**	US20 10182897 SEQ ID **NO:** 250	6026
RSV805	Nanobody binding to RSV **F** protein	**RSVPMP25B3**	US20 10 182897 SEQ ID NO: **25**	6027
RSV806	Nanobody binding to RSV **F** _proiém	**RSVPMP8C8**	**US201 10182897** SEQ **ID NO:** 252	6028
RSV807	Nanobody binding to RSV **F** protein	**RSVPMP5A6**	**US201 10182897** SEQ **ID** NO: 253	6029
RSV808	Nanobody binding to RSV **F** _protein	**RSVPMP8E1 1**	**US20 110 182897** SEQ **ID** NO: 254	60.30
RSV809	Nanobody binding to RSV **F** protein	**RSVPMP8F1 1**	**US201 10182897** SEQ **ID** NO: 255	603 **1**
RSV8 10	Nanobody binding to RSV **F** _protein	**RSVPMP13F1**	**US201 10182897** SEQ **ID** NO: 256	60.32
RSV81 1	Nanobody binding to RSV **F** protein	**RSVPMP15B8**	S20 **110 182897** SEQ **ID** NO: 257	603.3
RSV8 12	Nanobody binding to RSV **F** protein	**RSVPMP15G 11**	**US201 10182897** SEQ **ID** NO: 258	60.34
RSV813	Nanobody binding to RSV **F** protein	**RSVPMP17C 10**	**US201 0 182897** SEQ **ID** NO: 259	6035
RSV8 14	Nanobody binding to RSV **F** protein	**RSVPMP21E7**	**US20 10 182897** SEQ **ID** NO: 260	60.36
RSV815	Nanobody binding to RSV **F** protein	**RSVPMP2 1F8**	**US201 10182897** SEQ ID NO: 261	6037
RSV816	Nanobody binding to RSV **F** protein	**RSVPMP5A2**	**US201 10182897** SEQ ID NO: 262	6038
RSV817	Nanobody binding to RSV **F** protein	**RSVPMP5B2**	**US20 10 182897** SEQ ID NO: 263	6039
RSV818	Nanobody binding to RSV **F** protein	**RSVPMP5C3**	**US20 10 182897** SEQ ID NO: 264	6040
RSV819	Nanobody binding to RSV **F** protein	**RSVPMP5D2**	**US201 10182897** SEQ ID NO: 265	6041
RSV820	Nanobody binding to RSV **F** protein	**RSVPMP5E2**	**US20 10 182897** SEQ ID NO: 266	6042
RSV821	Nanobody binding to RSV **F** protein	**RSVPMP5F3**	**US201 10182897** SEQ ID NO: 267	6043
RSV822	Nanobody binding to RSV **F** protein	**RSVPMP5G3**	**US201 10182897** SEQ ID NO: 268	6044
RSV823	Nanobody binding to RSV **F** protein	**RSVPMP5H2**	**US201 10182897** SEQ ID NO: 269	6045
RSV824	Nanobody binding to RSV **F** protein	**RSVPMP5H3**	**US201 10182897** SEQ ID NO: 270	6046

RSV825	**Nanobody** binding to **RSV F protein**	**RSVPMP8C 1**	**US20 110 182897** SEQ ID NO: 271	6047
RSV826	Nanobody binding to RSV **F** protein	**RSVPMP8F2**	**US201 10182897** SEQ ID NO: 272	6048
RSV827	Nanobody binding to RSV **F** protein	**RSVPMP8G4**	US20 I 10182897 SEQ ID **NO:** 273	6049
RSV828	Nanobody binding to RSV **F** protein	RSVPMP13A I	US201 I0 I82897 SEQ ID NO: 274	6050
RSV829	Nanobody binding to RSV **F** protein	RSVPMP13A4	US201 10182897 SEQ ID **NO:** 275	605 I
RSV830	Nanobody binding to RSV **F** protein	**RSVPMP13B I**	US20 I 10 182897 SEQ ID NO: 276	6052
RSV83 1	Nanobody binding to RSV **F** protein	RSVPMP I 3B2	**US201 10182897** SEQ **ID NO:** 277	6053
RSV832	Nanobody binding to RSV **F** protein	**RSVPMP13C I**	US20 I 10 182897 SEQ ID NO: 278	6054
RSV833	Nanobody binding to RSV **F** protein	RSVPMP13C3	**US201 10182897** SEQ **ID NO:** 279	6055
RSV834	Nanobody binding to RSV **F** protein	**RSVPMP13D6**	US201 10 I 82897 SEQ ID NO: 280	6056
RSV835	Nanobody binding to RSV **F** _proiem	**RSVPMP13E2**	**US201 10182897** SEQ **ID NO:** 281	6057
RSV836	Nanobody binding to RSV **F** protein	**RSVPMP13E3**	ıı S20 **110 182897** SEQ **ID** NO: 282	6058
RSV837	Nanobody binding to RSV **F** _protein	**RSVPMP15A5**	**US201 10182897** SEQ **ID** NO: 283	6059
RSV838	Nanobody binding to RSV **F** protein	**RSVPMP15A6**	**US201 10182897** SEQ **ID** NO: 284	6060
RSV839	Nanobody binding to RSV **F** _protein	**RSVPMP15B2**	**US20 I 10 182897** SEQ **ID** NO: 285	6061
RSV840	Nanobody binding to RSV **F** protein	**RSVPMP15B3**	ıı S20 **110 182897** SEQ **ID** NO: 286	6062
RSV841	Nanobody binding to RSV **F** protein	**RSVPMP15E5**	**US20 I 10 182897** SEQ **ID** NO: 287	6063
RSV842	Nanobody binding to RSV **F** protein	**RSVPMP17C2**	ıı S20 **110 182897** SEQ **ID** NO: 288	6064
RSV843	Nanobody binding to RSV **F** protein	**RSVPMP17D4**	**US201 10182897** SEQ **ID** NO: 289	6065
RSV844	Nanobody binding to RSV **F** protein	**RSVPMP17G4**	**US201 10182897** SEQ ID NO: 290	6066
RSV845	Nanobody binding to RSV **F** protein	**RSVPMP19B2**	**US20 I 10 182897** SEQ ID NO: 29 I	6067
RSV846	Nanobody binding to RSV **F** protein	**RSVPMP25A4**	**US201 10182897** SEQ ID NO: 292	6068
RSV847	Nanobody binding to RSV **F** protein	**RSVPMP25A9**	**US201 10182897** SEQ ID NO: 293	6069
RSV848	Nanobody binding to RSV **F** protein	**RSVPMP25B5**	**US20 I 10 182897** SEQ ID NO: 294	6070
RSV849	Nanobody binding to RSV **F** protein	**RSVPMP25G2**	**US201 10182897** SEQ ID NO: 295	6071
RSV850	Nanobody binding to RSV **F** protein	**RSVPMP25H5**	**US201 10182897** SEQ ID NO: 296	6072
RSV85 1	Nanobody binding to RSV **F** protein	**RSVPMP25E1 1**	**US20 I 10 182897** SEQ ID NO: 297	6073
RSV852	Nanobody binding to RSV **F** protein	**RSVPMP8G3**	**US201 10182897** SEQ ID NO: 298	6074
RSV853	Nanobody binding to RSV **F** protein	**RSVPMP13B5**	**US20 I 10 182897** SEQ ID NO: 299	6075

RSV854	**Nanobody** binding to **RSV** F **protein**	**RSVPMP15F2**	**US201 10182897** SEQ ID NO: 300	6076
RSV855	Nanobody binding to RSV F protein	**RSVPMP19E2**	**US201 10182897** SEQ ID NO: **301**	6077
RSV856	Nanobody binding to RSV F protein	**RSVPMP25D 1**	**US201 10182897** SEQ **ID NO:** 302	6078
RSV857	Nanobody binding to RSV **F** protein	RSVPMP5A1	**US201** 10182897 SEQ **ID** NO: 303	6079
RSV858	Nanobody binding to RSV F protein	**RSVPMP5G2**	US201 10182897 SEQ ID **NO:** 304	6080
RSV859	Nanobody binding to RSV **F** protein	**RSVPMP5H1**	US201 10182897 SEQ ID NO: 305	6081
RSV860	Nanobody binding to RSV F protein	**RSVPMP6B I**	**US201 10182897** SEQ **ID NO:** 306	6082
RSV861	Nanobody binding to RSV **F** protein	**RSVPMP8H2**	US201 10182897 SEQ ID NO: 307	6083
RSV862	Nanobody binding to RSV F protein	**RSVPMP8H3**	US201 10182897 SEQ ID **NO:** 308	6084
RSV863	Nanobody binding to RSV **F** protein	**RSVPMP13A3**	US201 10182897 SEQ ID NO: 309	6085
RSV864	Nanobody binding to RSV F protein	RSVPMP13C5	**US201 10182897** SEQ **ID NO:** 310	6086
RSV865	Nanobody binding to RSV **F** protein	**RSVPMP13H 1**	US201 **10182897** SEQ **ID** NO: 311	6087
RSV866	Nanobody binding to RSV F protein	**RSVPMP13H2**	**US201 10182897** SEQ **ID** NO: 312	6088
RSV867	Nanobody binding to RSV **F** protein	**RSVPMP15E6**	US20 **110 182897** SEQ **ID** NO: 313	6089
RSV868	Nanobody binding to RSV F protein	**RSVPMP17A3**	**US201 10182897** SEQ **ID** NO: 314	6090
RSV869	Nanobody binding to RSV **F** protein	**RSVPMP25G8**	US201 **10182897** SEQ ID NO: 315	6091
RSV870	Nanobody binding to RSV F protein	**RSVPMP6D1**	**US201 10182897** SEQ **ID** NO: 316	6092
RSV871	Nanobody binding to RSV **F** protein	**RSVPMP8D5**	US201 **10182897** SEQ ID NO: 317	6093
RSV872	Nanobody binding to RSV F protein	**RSVPMP13B4**	**US20 10 182897** SEQ **ID** NO: 318	6094
RSV873	Nanobody binding to RSV F protein	**RSVPMP13B6**	**US20 10 182897** SEQ ID NO: 319	6095
RSV874	Nanobody binding to RSV F protein	**RSVPMP13E6**	**US20 10 182897** SEQ ID NO: 320	6096
RSV875	Nanobody binding to RSV F protein	**RSVPMP13F4**	**US201 10182897** SEQ ID NO: 321	6097
RSV876	Nanobody binding to RSV F protein	**RSVPMP15H3**	**US201 10182897** SEQ ID NO: 322	6098
RSV877	Nanobody binding to RSV F protein	**RSVPMP17E5**	**US20 10 182897** SEQ ID NO: 323	6099
RSV878	Nanobody binding to RSV F protein	**RSVPMP19D3**	**US201 10182897** SEQ ID NO: 324	6100
RSV879	Nanobody binding to RSV F protein	**RSVPMP19F3**	**US201 10182897** SEQ ID NO: 325	6101
RSV880	Nanobody binding to RSV F protein	**RSVPMP25C4**	**US20 10 182897** SEQ ID NO: 326	6102
RSV881	Nanobody binding to RSV F protein	**RSVPMP25E3**	**US20 10 182897** SEQ ID NO: 327	6103
RSV882	Nanobody binding to RSV F protein	**RSVPMP5G4**	**US201 10182897** SEQ ID NO: 328	6104

RSV883	**Nanobody** binding to **RSV** F **protein**	**RSVPMP6G5**	**US201 10182897** SEQ ID NO: 329	6105
RSV884	Nanobody binding to RSV F protein	**RSVPMP8E6**	**US20 110 182897** SEQ ID NO: 330	6106
RSV885	Nanobody binding to RSV F protein	**RSVPMP13A10**	US20110182897 SEQ ID **NO: 331**	6107
RSV886	Nanobody binding to RSV **F** protein	RSVPMP2 1H10	US201 10182897 SEQ ID NO: 332	6108
RSV887	Nanobody binding to RSV F protein	**RSVPMP5A8**	US201 10182897 SEQ ID **NO:** 333	6109
RSV888	Nanobody binding to RSV **F** protein	**RSVPMP5A10**	US201 10 182897 SEQ ID NO: 334	6110
RSV889	Nanobody binding to RSV F protein	RSVPMP1 4A6	US201 10182897 SEQ ID **NO:** 335	6111
RSV890	Nanobody binding to RSV **F** protein	**RSVPMP16A6**	US201 10182897 SEQ ID NO: 336	6112
RSV891	Nanobody binding to RSV F protein	**RSVPMP22D6**	US201 10182897 SEQ ID **NO:** 337	6113
RSV892	Nanobody binding to RSV **F** protein	**RSVPMP8E2**	US20110182897 SEQ ID NO: 338	6114
RSV893	Nanobody binding to RSV F _proiem	**RSVPMP8C6**	**US201 10182897** SEQ **ID NO:** 339	6115
RSV894	Nanobody binding to RSV **F** protein	**RSVPMP5C6**	US20 **110 182897** SEQ **ID** NO: 340	6116
RSV895	Nanobody binding to RSV F _protein	**RSVPMP6D4**	**US201 10182897** SEQ **ID** NO: 341	6117
RSV896	Nanobody binding to RSV **F** protein	**RSVPMP8B10**	US20 **110 182897** SEQ **ID** NO: 342	6118
RSV897	Nanobody binding to RSV F _protein	**RSVPMP8E10**	**US20110 182897** SEQ **ID** NO: 343	**6119**
RSV898	Nanobody binding to RSV **F** protein	**RSVPMP15A7**	**US201 10182897** SEQ **ID** NO: 344	6120
RSV899	Nanobody binding to RSV F protein	**RSVPMP15E10**	**US20110 182897** SEQ **ID** NO: 345	6121
RSV900	Nanobody binding to RSV **F** protein	**RSVPMP13C7**	US20 **110 182897** SEQ **ID** NO: 346	6122
RSV901	Nanobody binding to RSV F protein	**RSVPMP15A9**	**US201 10182897** SEQ **ID** NO: 347	6123
RSV902	Nanobody binding to RSV F protein	**RSVPMP15F1 1**	**US201 10182897** SEQ ID NO: 348	6124
RSV903	Nanobody binding to RSV F protein	**RSVPMP15A1**	**US201 10182897** SEQ ID NO: 349	6125
RSV904	Nanobody binding to RSV F protein	**RSVPMP6H2**	**US201 10182897** SEQ ID NO: 350	6126
RSV905	Nanobody binding to RSV F protein	**RSVPMP17A9**	**US201 10182897** SEQ ID NO: 351	6127
RSV906	Nanobody binding to RSV F protein	**RSVPMP7G1**	**US201 10182897** SEQ ID NO: 352	6128
RSV907	Nanobody binding to RSV F protein	**RSVPMP5A9**	**US201 10182897** SEQ ID NO: 353	6129
RSV908	Nanobody binding to RSV F protein	**RSVPMP7B2**	**US20110 182897** SEQ ID NO: 354	6130
RSV909	Nanobody binding to RSV F protein	**RSVPMP22A4**	**US201 10182897** SEQ ID NO: 355	**6131**
RSV910	Nanobody binding to RSV F protein	**RSVPMP22E10**	**US20110 182897** SEQ ID NO: 356	6132
RSV91 1	Nanobody binding to RSV F protein	**RSVPMP22H4**	**US201 10182897** SEQ ID NO: 357	6133

RSV912	**Nanobody** binding to **RSV** F **protein**	**RSVPMP15C5**	**US20110182897** SEQ ID NO: 358	6134
RSV913	Nanobody binding to RSV F protein	**RSVNC39**	**US20110182897** SEQ ID NO: 359	6135
RSV914	Nanobody binding to RSV F protein	**RSVPMP7B9**	**US20110182897** SEQ **ID NO:** 360	6136
RSV915	Nanobody binding to RSV **F** protein	**RSVPMP15E11**	US20110182897 SEQ ID NO: **361**	6137
RSV916	Nanobody binding to RSV F protein	**RSVPMP7E7**	**US20110182897** SEQ **ID NO:** 362	6138
RSV917	Nanobody binding to RSV **F** protein	**RSVPMP14H3**	**US20110182897** SEQ **ID** NO: 363	6139
RSV918	Nanobody binding to RSV F protein	**RSVPMP24D6**	US20110182897 SEQ ID **NO:** 364	6140
RSV919	Nanobody binding to RSV **F** protein	RSVPMP23E5	US20110182897 SEQ ID NO: 365	6141
RSV920	Nanobody binding to RSV F protein	**RSVPMP8A6**	US20110182897 SEQ ID **NO:** 366	6142
RSV921	Nanobody binding to RSV **F** protein	**RSVPMP14E2**	US20110182897 SEQ ID NO: 367	6143
RSV922	Nanobody binding to RSV F protein	**RSVPMP25F3**	**US20110182897** SEQ **ID NO:** 368	6144
RSV923	Nanobody binding to RSV **F** protein	**RSVPMP19A6**	**US20110182897** SEQ **ID** NO: 369	6145
RSV924	Nanobody binding to RSV F protein	**RSVPMP23G1**	**US20110182897** SEQ **ID** NO: 370	6146
RSV925	Nanobody binding to RSV **F** protein	**RSVPMP15H8**	**US20110182897** SEQ **ID** NO: 371	6147
RSV926	Nanobody binding to RSV F protein	**RSVNC41**	**US20110182897** SEQ **ID** NO: 372	6148
RSV927	Nanobody binding to RSV **F** protein	**RSVPMP6A8**	**US20110182897** SEQ **ID** NO: 373	6149
RSV928	Nanobody binding to RSV F protein	**RSVPMP25H9**	**US20110182897** SEQ **ID** NO: 374	6150
RSV929	Nanobody binding to RSV **F** protein	**RSVPMP8B1**	US20110182897 SEQ **ID** NO: 375	**6151**
RSV930	Nanobody binding to RSV F protein	**RSVPMP17E1**	**US20110182897** SEQ **ID** NO: 376	6152
RSV931	Nanobody binding to RSV F protein	**RSVPMP21A4**	**US20110182897** SEQ ID NO: 377	6153
RSV932	Nanobody binding to RSV F protein	**RSVPMP25A11**	**US20110182897** SEQ ID NO: 378	6154
RSV933	Nanobody binding to RSV F protein	**RSVPMP25C8**	**US20110182897** SEQ ID NO: 379	6155
RSV934	Nanobody binding to RSV F protein	**RSVNC23**	**US20110182897** SEQ ID NO: 380	6156
RSV935	Nanobody binding to RSV F protein	**RSVPMP20A11**	**US20110182897** SEQ ID NO: 381	6157
RSV936	Nanobody binding to RSV F protein	**RSVPMP20A9**	**US20110182897** SEQ ID NO: 382	6158
RSV937	Nanobody binding to RSV F protein	**RSVPMP1F7**	**US20110182897** SEQ ID NO: 383	6159
RSV938	Nanobody binding to RSV F protein	**RSVPMP20D6**	**US20110182897** SEQ ID NO: **384**	6160
RSV939	Nanobody binding to RSV F protein	**RSVPMP1F1**	**US20110182897** SEQ ID NO: 385	6161
RSV940	Nanobody binding to RSV F protein	**RSVPMP3D3**	**US20110182897** SEQ ID NO: 386	6162

RSV941	Nanobody binding to RSV F protein	RSVPMP3E6	US20110182897 SEQ ID NO: 387	6163
RSV942	Nanobody binding to RSV F protein	RSVPMP1C8	US20110182897 SEQ ID NO: 388	6164
RSV943	Nanobody binding to RSV F protein	RSVPMP1A2	US20110182897 SEQ ID NO: 389	6165
RSV944	Nanobody binding to RSV F protein	RSVPMP1C5	US20110182897 SEQ ID NO: 390	6166
RSV945	Nanobody binding to RSV F protein	RSVPMP20G5	US20110182897 SEQ ID NO: 391	6167
RSV946	Nanobody binding to RSV F protein	RSVPMP4D8	US20110182897 SEQ ID NO: 392	6168
RSV947	Nanobody binding to RSV F protein	RSVPMP20B6	US20110182897 SEQ ID NO: 393	6169
RSV948	Nanobody binding to RSV F protein	RSVPMP1D4	US20110182897 SEQ ID NO: 394	6170
RSV949	Nanobody binding to RSV F protein	RSVPMP20A8	US20110182897 SEQ ID NO: 395	6171
RSV950	Nanobody binding to RSV F protein	RSVPMP20E7	US20110182897 SEQ ID NO: 396	6172
RSV951	Nanobody binding to RSV F protein	RSVPMP20G8	US20110182897 SEQ ID NO: 397	6173
RSV952	Nanobody binding to RSV F protein	RSVPMP2D3	US20110182897 SEQ ID NO: 398	6174
RSV953	Nanobody binding to RSV F protein	RSVPMP2G5	US20110182897 SEQ ID NO: 399	6175
RSV954	Nanobody binding to RSV F protein	RSVPMP2A6	US20110182897 SEQ ID NO: 400	6176
RSV955	Nanobody binding to RSV F protein	RSVPMP3A2	US20110182897 SEQ ID NO: 401	6177
RSV956	Nanobody binding to RSV F protein	RSVPMP4A8	US20110182897 SEQ ID NO: 402	6178
RSV957	Nanobody binding to RSV F protein	RSVPMP4F9	US20110182897 SEQ ID NO: 403	6179
RSV958	Nanobody binding to RSV F protein	RSVPMP1A6	US20110182897 SEQ ID NO: 404	6180
RSV959	Nanobody binding to RSV F protein	RSVPMP3C2	US20110182897 SEQ ID NO: 405	6181
RSV960	Nanobody binding to RSV F protein	RSVPMP4H9	US20110182897 SEQ ID NO: 406	6182
RSV961	Nanobody binding to RSV F protein	RSVPMP4B10	US20110182897 SEQ ID NO: 407	6183
RSV962	Nanobody binding to RSV F protein	203B1	US20110182897 SEQ ID NO: 2431	6184
RSV963	Nanobody binding to RSV F protein	203B2	US20110182897 SEQ ID NO: 2432	6185
RSV964	Nanobody binding to RSV F protein	2G3G1	US20110182897 SEQ ID NO: 2433	6186
RSV965	Nanobody binding to RSV F protein	203H1	US20110182897 SEQ ID NO: 2434	6187
RSV966	Nanobody binding to RSV F protein	2G2E4	US20110182897 SEQ ID NO: 2435	6188
RSV967	Nanobody binding to RSV F protein	189E2	US20110182897 SEQ ID NO: 2436	6189
RSV968	Nanobody binding to RSV F protein	203A12	US20110182897 SEQ ID NO: 2437	6190
RSV969	Nanobody binding to RSV F protein	203A9	US20110182897 SEQ ID NO: 2438	6191

RSV970	Nanobody binding to RSV F protein	203B12	US20110182897 SEQ ID NO: 2439	6192
RSV971	Nanobody binding to RSV F protein	203D2	US20110182897 SEQ ID NO: 2440	6193
RSV972	Nanobody binding to RSV F protein	203D9	US20110182897 SEQ ID NO: 2441	6194
RSV973	Nanobody binding to RSV F protein	203G3	US20110182897 SEQ ID NO: 2442	6195
RSV974	Nanobody binding to RSV F protein	203G9	US20110182897 SEQ ID NO: 2443	6196
RSV975	Nanobody binding to RSV F protein	203G10	US20110182897 SEQ ID NO: 2444	6197
RSV976	Nanobody binding to RSV F protein	203H9	US20110182897 SEQ ID NO: 2445	6198
RSV977	Nanobody binding to RSV F protein	203H10	US20110182897 SEQ ID NO: 2446	6199
RSV978	Nanobody binding to RSV F protein	202E4	US20110182897 SEQ ID NO: 2447	6200
RSV979	Nanobody binding to RSV F protein	189E2	US20110182897 SEQ ID NO: 2448	6201
RSV980	Nanobody binding to RSV F protein	PRSVPMP20C3	US20110182897 SEQ ID NO: 2574	6202
RSV981	Nanobody binding to RSV F protein	PRSVPMP20C5	US20110182897 SEQ ID NO: 2575	6203
RSV982	Nanobody binding to RSV F protein	PRSVPMP20B2	US20110182897 SEQ ID NO: 2576	6204
RSV983	Nanobody binding to RSV F protein	PRSVPMP20C1	US20110182897 SEQ ID NO: 2577	6205
RSV984	Nanobody binding to RSV F protein	PRSVPMP 1G8	US20110182897 SEQ ID NO: 2578	6206
RSV985	Nanobody binding to RSV F protein	PRSVNMP1A4	US20110182897 SEQ ID NO: 2579	6207
RSV986	Nanobody binding to RSV F protein	PRSVPMP 13E 12	US20110182897 SEQ ID NO: 2580	6208
RSV987	Nanobody binding to RSV F protein	PRSVPMP5C6	US20110182897 SEQ ID NO: 2581	6209
RSV988	Nanobody binding to RSV F protein	LG203E7	US20110182897 SEQ ID NO: 2682	6210
RSV989	Nanobody binding to RSV F protein	LG203G8	US20110182897 SEQ ID NO: 2683	6211
RSV990	Nanobody binding to RSV F protein	LG211A10	US20110182897 SEQ ID NO: 2684	6212
RSV991	Nanobody binding to RSV F protein	LG211A8	US20110182897 SEQ ID NO: 2685	6213
RSV992	Nanobody binding to RSV F protein	LG211B10	US20110182897 SEQ ID NO: 2686	6214
RSV993	Nanobody binding to RSV F protein	LG211B8	US20110182897 SEQ ID NO: 2687	6215
RSV994	Nanobody binding to RSV F protein	LG211C12	US20110182897 SEQ ID NO: 2688	6216
RSV995	Nanobody binding to RSV F protein	LG211C8	US20110182897 SEQ ID NO: 2689	6217
RSV996	Nanobody binding to RSV F protein	LG211D10	US20110182897 SEQ ID NO: 2690	6218
RSV997	Nanobody binding to RSV F protein	LG211D8	US20110182897 SEQ ID NO: 2691	6219
RSV998	Nanobody binding to RSV F protein	LG211E10	US20110182897 SEQ ID NO: 2692	6220

RSV999	Nanobody binding to RSV F protein	LG21 1E12	US20 110 182897 SEQ ID NO: 2693	622 1
RSV1000	Nanobody binding to RSV F protein	LG21 1E8	US20 110 182897 SEQ ID NO: 2694	6222
RSVIOO !	Nanobody binding to RSV F protein	LG2 i 1H8	US201 10182897 SEQ ID NO: 2695	6223
RSV1 002	Nanobody binding to RSV F protein	LG212A ! 0	US20 ! ! 0 ! 182897 SEQ ID NO: 2696	6224
RSV1003	Nanobody binding to RSV F protein	LG2 12A12	US201 10182897 SEQ ID NO: 2697	6225
RSV1 004	Nanobody binding to RSV F protein	LG212A2	US20 ! 10 182897 SEQ ID NO: 2698	6226
RSV1005	Nanobody binding to RSV F protein	LG2 12A8	US201 10182897 SEQ ID NO: 2699	6227
RSV1 006	Nanobody binding to RSV F protein	LG212B 12	US20 ! 10 182897 SEQ ID NO: 2700	6228
RSV1007	Nanobody binding to RSV F protein	LG2 12B2	US201 10182897 SEQ ID NO: 270 !	6229
RSV1 008	Nanobody binding to RSV F protein	LG212C 12	US20 ! 10 182897 SEQ ID NO: 2702	6230
RSV1009	Nanobody binding to RSV F _proiem	LG2 ! 2D ! 0	US201 10182897 SEQ ID NO: 2703	623 !
RSV10 10	Nanobody binding to RSV F protein	LG212D12	ıı S20 110 182897 SEQ ID NO: 2704	6232
RSV101 1	Nanobody binding to RSV F _protein	LG212D2	US20 ! 10 182897 SEQ ID NO: 2705	6233
RSV10 12	Nanobody binding to RSV F protein	LG212E10	ıı S20 110 182897 SEQ ID NO: 2706	6234
RSV1013	Nanobody binding to RSV F _protein	LG212E12	US20 ! 10 182897 SEQ ID NO: 2707	6235
RSV10 14	Nanobody binding to RSV F protein	LG212E6	ıı S20 110 182897 SEQ ID NO: 2708	6236
RSV1015	Nanobody binding to RSV F protein	LG212F10	US20 ! 10 182897 SEQ ID NO: 2709	6237
RSV10 16	Nanobody binding to RSV F protein	LG212F1 2	ıı S20 110 182897 SEQ ID NO: 2710	6238
RSV1017	Nanobody binding to RSV F protein	LG212F6	US20 ! 10 182897 SEQ ID NO: 27 ! 1	6239
RSV1018	Nanobody binding to RSV F protein	LG212F8	US20 ! 10 182897 SEQ ID NO: 2712	6240
RSV1019	Nanobody binding to RSV F protein	LG212G10	US20 ! 10 182897 SEQ ID NO: 27 ! 3	6241
RSV1020	Nanobody binding to RSV F protein	LG212G2	US20 ! 10 182897 SEQ ID NO: 2714	6242
RSV1021	Nanobody binding to RSV F protein	LG212H10	US20 ! 10 182897 SEQ ID NO: 27 ! 5	6243
RSV1022	Nanobody binding to RSV F protein	LG212H2	US20 ! 10 182897 SEQ ID NO: 2716	6244
RSV1023	Nanobody binding to RSV F protein	LG212H8	US20 ! 10 182897 SEQ ID NO: 27 ! 7	6245
RSV1024	Nanobody binding to RSV F protein	IV121	US201 10182897 SEQ ID NO: 3064	6246
RSV1025	Nanobody binding to RSV F protein	IV122	US20 ! 10 182897 SEQ ID NO: 3065	6247
RSV1026	Nanobody binding to RSV F protein	IV123	US20 ! 10 182897 SEQ ID NO: 3066	6248
RSV1027	Nanobody binding to RSV F protein	IV126	US20 ! 10 182897 SEQ ID NO: 3067	6249

RSV1028	**Nanobody** binding to **RSV** F **protein**	IV127	**US20110182897** SEQ ID NO: 3068	6250
RSV1029	**Nanobody** binding to RSV F protein	**IV131**	**US20110182897** SEQ ID NO: 3069	6251
RSV1030	**Nanobody** binding **to** RSV F protein	IV132	**US20110182897** SEQ **ID NO:** 3070	6252
RSV1031	Nanobody binding to RSV **F** protein	**IV133**	US20110182897 SEQ **ID** NO: 3071	6253
RSV1032	**Nanobody** binding **to** RSV F protein	IV134	**US20110182897** SEQ **ID NO:** 3072	6254
RSV1033	Nanobody binding to RSV **F** protein	**TV135**	US20110182897 SEQ **ID** NO: 3073	6255
RSV1034	**Nanobody** binding **to** RSV F protein	IV136	**US20110182897** SEQ **ID NO:** 3074	6256
RSV1035	Nanobody binding to RSV **F** protein	**TV140**	US20110182897 SEQ **ID** NO: 3075	6257
RSV1036	**Nanobody** binding **to** RSV F protein	IV **144**	**US20110182897** SEQ **ID NO:** 3076	6258
RSV1037	Nanobody binding to RSV **F** protein	**TV156**	US20110182897 SEQ **ID** NO: 3077	6259
RSV1038	**Nanobody** binding **to** RSV F _proiem	IV **157**	**US20110182897** SEQ **ID NO:** 3078	6260
RSV1039	Nanobody binding to RSV **F** protein	EV160	US20**110182897** SEQ **ID NO:** 3079	6261
RSV1040	Nanobody binding to RSV F _protein	IV **124**	**US20110182897** SEQ **ID** NO: 3080	6262
RSV1041	Nanobody binding to RSV **F** protein	EV125	US20**110182897** SEQ **ID** NO: 3081	6263
RSV1042	Nanobody binding to RSV F _protein	IV **145**	**US20110182897** SEQ **ID** NO: 3082	6264
RSV1043	Nanobody binding to RSV **F** protein	EV146	US20**110182897** SEQ **ID** NO: 3083	6265
RSV1044	Nanobody binding to RSV F protein	IV147	**US20110182897** SEQ **ID** NO: 3084	6266
RSV1045	Nanobody binding to RSV **F** protein	IV **151**	US20**110182897** SEQ **ID** NO: 3085	6267
RSV1046	Nanobody binding to RSV F protein	IV **153**	**US20110182897** SEQ **ID** NO: 3086	6268
RSV1047	Nanobody binding to RSV F protein	**IV154**	**US20110182897** SEQ ID NO: 3087	6269
RSV1048	Nanobody binding to RSV F protein	**IV155**	**US20110182897** SEQ ID NO: 3088	6270
RSV1049	Nanobody binding to RSV F protein	**IV1**	**US20110182897** SEQ ID NO: 3089	6271
RSV1050	Nanobody binding to RSV F protein	IV2	**US20110182897** SEQ ID NO: 3090	6272
RSV1051	Nanobody binding to RSV F protein	**IV3**	**US20110182897** SEQ ID NO: 3091	6273
RSV1052	Nanobody binding to RSV F protein	IV4	**US20110182897** SEQ ID NO: 3092	6274
RSV1053	Nanobody binding to RSV F protein	**IV6**	**US20110182897** SEQ ID NO: 3093	6275
RSV1054	Nanobody binding to RSV F protein	IV7	**US20110182897** SEQ ID NO: 3094	6276
RSV1055	Nanobody binding to RSV F protein	**IV9**	**US20110182897** SEQ ID NO: 3095	6277
RSV1056	Nanobody binding to RSV F protein	**IV10**	**US20110182897** SEQ ID NO: 3096	6278

RSV1057	Nanobody binding to RSV F protein	IV 11	US20 1 10 182897 SEQ ID NO: 3097	6279
RSV1058	Nanobody binding to RSV F protein	IV12	US201 10182897 SEQ ID NO: 3098	6280
RSV1059	Nanobody binding to RSV F protein	IV 16	US201 10182897 SEQ ID NO: 3099	6281
RSV1 060	Nanobody binding to RSV F protein	IV24	US20 1 10 182897 SEQ ID NO: 3 100	6282
RSV106 1	Nanobody binding to RSV F protein	IV26	US201 10182897 SEQ ID NO: 3 10 1	6283
RSV1 062	Nanobody binding to RSV F protein	IV30	US20 1 10 182897 SEQ ID NO: 3 102	6284
RSV1063	Nanobody binding to RSV F protein	IV34	US201 10182897 SEQ ID NO: 3 103	6285
RSV1 064	Nanobody binding to RSV F protein	IV14	US20 1 10 182897 SEQ ID NO: 3 104	6286
RSV1065	Nanobody binding to RSV F protein	IV 15	US201 10182897 SEQ ID NO: 3 105	6287
RSV1 066	Nanobody binding to RSV F protein	IV 17	US20 1 10 182897 SEQ ID NO: 3 106	6288
RSV1067	Nanobody binding to RSV F _proiem	IV 18	US201 10182897 SEQ ID NO: 3 107	6289
RSV1068	Nanobody binding to RSV F protein	IV 29	ıı S20 110 182897 SEQ ID NO: 3 108	6290
RSV1069	Nanobody binding to RSV F _protein	IV3 1	US20 1 10 182897 SEQ ID NO: 3 109	6291
RSV1070	Nanobody binding to RSV F protein	EV3.3	ıı S20 110 182897 SEQ ID NO: 3 110	6292
R SV 107 1	Nanobody binding to RSV F _protein	IV35	US20 1 10 182897 SEQ ID NO: 3 111	6293
RSV1072	Nanobody binding to RSV F protein	EV36	ıı S20 110 182897 SEQ ID NO: 3 112	6294
RSV1073	Nanobody binding to RSV F protein	IV40	US20 1 10 182897 SEQ ID NO: 3 113	6295
RSV1074	Nanobody binding to RSV F protein	EV42	ıı S20 110 182897 SEQ ID NO: 3 114	6296
RSV1075	Nanobody binding to RSV F protein	IV8	US20 1 10 182897 SEQ ID NO: 3 115	6297
RSV1076	Nanobody binding to RSV F protein	IV21	US20 1 10 182897 SEQ ID NO: 3 116	6298
RSV1077	Nanobody binding to R SV F protein	IV23	US20 1 10 182897 SEQ ID NO: 3 117	6299
RSV1078	Nanobody binding to RSV F protein	IV45	US20 1 10 182897 SEQ ID NO: 3 118	6300
RSV1079	Nanobody binding to R SV F protein	IV47	US20 1 10 182897 SEQ ID NO: 3 119	6301
RSV1080	Nanobody binding to RSV F protein	IV48	US20 1 10 182897 SEQ ID NO: 3 120	6302
RSV1081	Nanobody binding to R SV F protein	IV50	US20 1 10 182897 SEQ ID NO: 3 121	6303
RSV1082	Nanobody binding to RSV F protein	IV22	US20 1 10 182897 SEQ ID NO: 3 122	6304
RSV1083	Nanobody binding to R SV F protein	IV37	US20 1 10 182897 SEQ ID NO: 3 123	6305
RSV1084	Nanobody binding to RSV F protein	IV38	US20 1 10 182897 SEQ ID NO: 3 124	6306
RSV1085	Nanobody binding to R SV F protein	IV5	US20 1 10 182897 SEQ ID NO: 3 125	6307

RSV1086	Nanobody binding to RSV F protein	IV27	US20110182897 SEQ ID NO: 3126	6308
RSV1087	Nanobody binding to RSV F protein	IV25	US20110182897 SEQ ID NO: 3127	6309
RSV1088	Nanobody binding to RSV F protein	IV28	US20110182897 SEQ ID NO: 3128	6310

[00318] In one embodiment, the payload region of the AAV particle composes one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides. fragments or variants thereof described in US Publication No. US20140363427, and international Publication No. WO2004083373, the contents of each of which are herein incorporated by reference in their entirety, against RSV F or RSV G protein.

[00319] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 23 against Hepatitis B, Hepatitis C and/or Hepatitis D.

Table 23. Antibodies against Hepatitis B, C, D viruses

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
HEPBD1	Anti-preS1 immunoglobulin, HBV Ab		Park, S.G., et al., Hepatitis B virus-neutralizing anti-pre-S1 human antibody fragments from large naive antibody phage library; Antiviral Res. 68 (3), 109-115 (2005); NCBI Accession # AAW82034.1 (107aa)	6311
HEPBD2	Anti-preS1 immunoglobulin, HBV Ab		Park, S.G., et al., Hepatitis B virus-neutralizing anti-pre-S1 human antibody fragments from large naive antibody phage library; Antiviral Res. 68 (3), 109-115 (2005); NCBI Accession # AAW82035.1 (132aa)	6312
HEPBD3	Anti-preS1 immunoglobulin, HBV Ab		Park, S.G., et al., Hepatitis B virus-neutralizing anti-pre-S1 human antibody fragments from large naive antibody phage library; Antiviral Res. 68 (3), 109-115 (2005); NCBI Accession # AAW82033.1(111aa)	6313
HEPBD4	Anti-preS1 immunoglobulin, HBV Ab		Park, S.G., et al., Hepatitis B virus-neutralizing anti-pre-S1 human antibody fragments from large naive antibody phage library; Antiviral Res. 68 (3), 109-115 (2005); NCBI Accession # AAW82032.1 (142aa)	6314
HEPBD5	HCV Ab	Hu5b3.v3	Pantua, H., et al., Glycan shifting on hepatitis C virus(HCV) e2 glycoprotein is a mechanism for escape from broadly neutralizing antibodies; J. Mol. Biol. 425 (11), 1899-1914 (2013) NCBI Accession # 4HS8_H(228aa)	6315
HEPBD6	HCV Ab	Igh526	Kong L., et al., Structure of Hepatitis C Virus Envelope Glycoprotein E1 Antigenic Site 314-324 in Complex with Antibody IGH526; J. Mol. Biol. 427 (16), 2617-2628	6316

			(2015) NCBI Accession # 4N0Y_H(231aa)	
HEPBD7	Heavy chain partial, HCV Ab		Esposito, G., et al., Recombinant human antibodies specific for hepatitis C virus proteins; Arch. Virol. 142 (3), 601-610 (1997) NCBI Accession # CAA54914 (122aa)	6317
HEPBD8	Heavy chain variable gene, Chimeric HBV Ab		EP0521348	6318
HEPBD9	Heavy chain variable region partial, HCV Ab		Keck, Z.Y., et al., Human monoclonal antibody to hepatitis C virUSE1 glycoprotein that blocks virus attachment and viral infectivity; J. Virol. 78 (13), 7257-7263 (2004) NCBI Accession # AAS47839 (142aa)	6319
HEPBD10	Heavy chain variable region, HBV Ab	E183/A2	US20120308580 SEQ ID NO: 33; WO 2011062562; CN102781961, EP2501723	6320
HEPBD11	Heavy chain variable region, HBV Ab		US20100260712 SEQ ID NO: 1; WO2009069917	6321
HEPBD12	Heavy chain variable region, HBV Ab		WO2015107126 SEQ ID NO: 2	6322
HEPBD13	Heavy chain variable region, HBV Ab	HB48-33, HB48-35, HB48-59	US8840895 SEQ ID NO: 1	6323
HEPBD14	Heavy chain variable region, HBV Ab	HFW141	US7435414 SEQ ID NO: 35; US20060014937; WO2005100400; CN1980956	6324
HEPBD15	Heavy chain variable region, HBV Ab		US7112664 SEQ ID NO: 8; US6680053, US6924368, US20020061581, US20040191259, US20050249753, WO2001092529	6325
HEPBD16	Heavy chain variable region, HBV Ab	Ab17.1.4 1	USRE39586 SEQ ID NO: 4; US6146629; WO1997047653	6326
HEPBD17	Heavy chain variable region, HBV Ab	Ab 19.79.5	USRE40831 SEQ ID NO: 4	6327
HEPBD18	Heavy chain variable region, HBV Ab		US20150232537 SEQ ID NO: 7; WO2014048910; CA2884388; CN104662041A; EP2900692	6328
HEPBD19	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 45	6329
HEPBD20	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 54	6330
HEPBD21	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 63	6331
HEPBD22	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 72	6332
HEPBD23	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 81	6333
HEPBD24	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 90	6334
HEPBD25	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 99	6335
HEPBD26	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 108	6336
HEPBD27	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 117	6337
HEPBD28	Heavy chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 126	6338

HEPBD29	Heavy **chain variable** region, HBV Ab		**WO2013** 165972 SEQ ID NO: 135	6339
HEPBD30	Heavy **chain variable** region, **HBV** Ab		**WO2013 165972** SEQ ID **NO:** 144	6340
HEPBD31	Heavy chain variable region, HBV Ab		**WO2013** 165972 SEQ ID NO: 153	6341
HEPBD32	Heavy **chain variable** region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 162	6342
HEPBD33	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 171	6343
HEPBD34	Heavy **chain variable** region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 180	6344
HEPBD35	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 189	6345
HEPBD36	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO;** 198	6346
HEPBD37	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 207	6347
HEPBD38	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 405	6348
HEPBD39	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 409	6349
HEPBD40	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 412	6350
HEPBD41	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID** NO: 418	6351
HEPBD42	Heavy chain variable region, HBV Ab		**WO2013 165972** SEQ **ID NO:** 421	6352
HEPBD43	Heavy chain variable region, HBV Ab		**WO2009069916** SEQ **ID** NO: 1	6353
HEPBD44	Heavy chain variable region, HBV Ab	PE1-1	WO 19940 11495	6354
HEPBD45	Heavy chain variable region, HBV Ab	ZM1-1	WO 19940 11495	6355
HEPBD46	Heavy chain variable region, HBV Ab	**ZMl-2**	WO 19940 11495	6356
HEPBD47	Heavy chain variable region, HBV Ab	**MD3-4**	WO 19940 11495	6357
HEPBD48	Heavy chain variable region, HBV Ab	A2E2	**CN102757492** SEQ ID **NO:** 2	6358
HEPBD49	Heavy chain variable region, HBV Ab	C9G9	**CN102757492** SEQ ID NO: 6	6359
HEPBD50	Heavy chain variable region, HBV Ab		**CN104530228** SEQ ID **NO: 3**	6360
HEPBD51	Heavy chain variable region, HBV Ab		CN104530228 SEQ ID NO: 4	6361
HEPBD52	Heavy chain variable region, HBV Ab	**Ab19**	US8580256 SEQ ID NO: 2	6362
HEPBD53	Heavy chain variable region, HBV Ab	**AM7**	US8580256 SEQ ID **NO:** 4	6363
HEPBD54	Heavy chain variable region, HBV Ab	**KR127**	US8420353 SEQ ID NO: 28	6364
HEPBD55	Heavy chain variable region, HBV Ab	DP7	US8420353 SEQ ID **NO:** 32	6365
HEPBD56	Heavy chain variable region, HBV Ab	HZI	**US8420353** SEQ ID NO: 36	6366
HEPBD57	Heavy chain variable region, HCV **Ab**	**MBL-HCV1** (Antibody	US855 1484 SEQ ID NO: 1	6367

		produced by clone **83-128)**		
HEPBD58	Heavy **cliain variable** regíon, HC**V Ab**	MBL-HCVl (Antibody produced by clone **95-2)**	**US855 1484** SEQ ID NO: 3	6368
HEPBD59	Heaw chain variable region, **HCV** Ab	**MBL-HCVl** (Antibody produced by clone 95- 14)	US855 1484 SEQ ID **NO:** 5	6369
HEPBD60	Heavy chain variable region, HCV Ab	Clone 13	**US7250 166** SEQ ID **NO:** 1	6370
HEPBD6 !	Heavy **chain variable** regíon, HCV **Ab**	Clone 98	**US7250166** SEQ ID NO: **3**	6371
HEPBD62	Heavy chain variable region, HCV Ab	Clone **1:4**	**US7250 166** SEQ ID **NO:** 5	6372
HEPBD63	Heavy **chain variable** regíon, HCV **Ab**	Clone **1:8**	**US7250166** SEQ ID NO: 7	6373
HEPBD64	Heavy chain variable region, HCV Ab	Clone **1:9**	**US7250 166** SEQ ID NO: 9	6374
HEPBD65	Heavy **chain variable** region, HCV Ab	Clone 1: 10	**US7250166** SEQ ID NO: **1**1	6375
HEPBD66	Heavy chain variable region, HCV Ab	Clone 4:6	**US7250166** SEQ ID NO: 13	6.376
HEPBD67	Heavy chain variable region, HCV Ab	Clone 6a: 5	1JS7250166 SEQ ID NO: 15	6377
HEPBD68	Heavy chain variable region, HCV Ab	Clone 2a: 2	US7250166 SEQ ID NO: 17	6.378
HEPBD69	Heavy chain variable region, HCV Ab	Clone **2a:4**	1JS7250166 SEQ ID NO: 19	6379
HEPBD70	Heavy chain variable region, HCV Ab	Clone 2a: 5	US7250166 SEQ ID NO: 2 1	6.380
HEPBD7 1	Heavy chain variable region, HCV Ab	Clone 2a: 13	**US7250166** SEQ ID NO: 23	6381
HEPBD72	Heavy chain variable region, HCV Ab	Clone 2a: 14	US7250166 SEQ ID NO: 25	6.382
HEPBD73	Heavy chain variable region, HCV Ab	Clone **2a:23**	**US7250166** SEQ ID NO: 27	6383
HEPBD74	Heaw chain variable region, HCV Ab	Clone 2a:23	**US7250166** SEQ ID **NO:** 29	6384
HEPBD75	**Heavy** chain variable region, HCV Ab	Clone 2a: 25	**US7250166** SEQ ID NO: 31	6385
HEPBD76	Heaw chain variable region, HCV Ab	Clone 2a: 30	**US7250166** SEQ ID **NO:** 33	6386
HEPBD77	**Heavy** chain variable region, HCV Ab	Clone 2a: 32	**US7250166** SEQ ID NO: 35	6387
HEPBD78	Heaw chain variable region, HCV Ab	Clone 2a: 33	**US7250166** SEQ ID **NO:** 37	6388
HEPBD79	**Heavy** chain variable region, HCV Ab	Clone 2a: 37	**US7250166** SEQ ID NO: 39	6389
HEPBD80	Heaw chain variable region, HCV Ab	Clone 2a: 40	**US7250166** SEQ ID **NO:** 4 1	6390
HEPBD81	**Heavy** chain variable region, HCV Ab	**Clone 2b :1**	**US7250166** SEQ ID NO: 43	6391
HEPBD82	Heaw chain variable region, HCV **Ab**	Clone 2b: 3	US7250166 SEQ ID **NO:** 45	6392
HEPBD83	Heavy chain variable region, HCV Ab	Clone 2b: 4	**US7250166** SEQ ID NO: 47	6393

HEPBD84	Heavy **chain variable** region, HCV Ab	Clone 2b:5	US7250166 SEQ ID NO: 49	6394
HEPBD85	Heavy **cliain variable** region, HCV Ab	Clone 2b:**7**	US7250166 SEQ ID NO: 51	6395
HEPBD86	Heavy chain variable region, HCV Ab	Clone 2b:9	US7250166 SEQ ID **NO:** 53	6396
HEPBD87	Heavy **chain variable** region, HCV **Ab**	Clone 2b:10	**US7250166** SEQ ID NO: 55	6397
HEPBD88	Heavy chain variable region, HCV Ab	**anti-NS3** Fab	US7314919 SEQ ID NO: **1**	6398
HEPBD89	Heavy **chain variable** region, HCV **Ab**	Antibody produced **by** clone 95-14	**US855 1484** SEQ ID NO: 32	6399
HEPBD90	Heavy chain variable region, HCV Ab	Antibody produced by clone 95-38	1JS855 1484 SEQ ID NO: 33	6400
HEPBD91	Heavy chain variable region, HCV Ab	Antibody produced by clone 95-25	US855 1484 SEQ ID NO: 34	6401
HEPBD92	Heavy chain variable region, HCV Ab	Antibody produced by clone 95.42	**US855 1484** SEQ ID NO: 35	6402
HEPBD93	Heavy chain variable region, HCV Ab	**Antibody** produced by clone 95-43	US855 1484 SEQ ID NO: 36	6403
HEPBD94	Heavy chain variable region, HCV Ab	Antibody produced **by** clone 95-49	US855 1484 SEQ ID **NO:** 37	6404
HEPBD95	Heavy **cliain variable** region, HCV **Ab**	Antibody produced by clone 95-54	US855 1484 SEQ ID NO: 38	6405
HEPBD96	Heavy chain variable region, HCV **Ab**	Antibody produced **by** clone 95-58	**US855 1484** SEQ ID NO: 39	6406
HEPBD97	Heavy chain variable region, HCV Ab	Antibody produced by clone 95-62	1JS855 1484 SEQ ID NO: 40	6407
HEPBD98	Heavy chain variable region, HCV Ab	**HC-84. 1**	**US20130084301** SEQ ID NO: 55	6408
HEPBD99	Heavy cliain variable region, HCV **Ab**	**HC-84.20**	**US20130084301** SEQ ID NO: 56	6409
HEPBD 100	Heavy chain variable region, HCV Ab	HC-84.2 1	**US20130084301** SEQ ID NO: 57	6410
HEPBD101	Heavy cliain variable region, HCV **Ab**	HC-84.22	**US20130084301** SEQ ID NO: 58	**641 1**
HEPBD 102	Heavy chain variable region, HCV Ab	HC-23	US20 13008430**1** SEQ ID NO: 59	6412
HEPBD103	Heavy cliain variable region, HCV **Ab**	HC-84.24	**US20130084301** SEQ ID NO: 60	6413
HEPBD 104	Heavy chain variable region, HCV Ab	HC-84.25	**US20130084301** SEQ ID NO: 61	6414
HEPBD105	Heavy cliain variable region, HCV **Ab**	HC-84.26	**US20130084301** SEQ ID NO: 62	6415
HEPBD 106	Heavy chain variable region, HCV Ab	HC-84.27.	**US20130084301** SEQ ID NO: 63	6416
HEPBD107	Heavy chain variable region, HCV **Ab**	**AOT3**	**US20120009196** SEQ **ID** NO: 1	6417
HEPBD 108	Heavy chain variable region, HCV Ab	CI1-3	**US20 120009 196** SEQ **ID NO: 3**	6418

HEPBD109	Heavy chain variable region, HCV Ab	C11-7	US20120009196 SEQ ID NO: 5	6419
HEPBD110	Heavy chain variable region, HCV Ab	C11-9	US20120009196 SEQ ID NO: 7	6420
HEPBD111	Heavy chain variable region, HCV Ab	C11-14	US20120009196 SEQ ID NO: 9	6421
HEPBD112	Heavy chain variable region, HCV Ab		WO2014065822 SEQ ID NO: 3	6422
HEPBD113	Heavy chain variable region, HCV Ab		WO2014065822 SEQ ID NO: 7	6423
HEPBD114	Heavy chain variable region, HCV Ab	rmPA-29	WO2007143701 SEQ ID NO: 2	6424
HEPBD115	Heavy chain variable region, HCV Ab	Hc33.1	Li Y. et al., Structural basis for penetration of the glycan shield of hepatitis C virUSe2 glycoprotein by a broadly neutralizing human antibody; J. Biol. Chem. 290 (16), 10117-10125 (2015) NCBI Accession # 4XVJ_H (141aa)	6425
HEPBD116	Heavy chain variable region, HCV Ab		Martin, F., et al., Affinity selection of a camelized V(H) domain antibody inhibitor of hepatitis C virUSNS3 protease; Protein Eng. 10 (5), 607-614 (1997NCBI Accession # 1OL0_B (121aa)	6426
HEPBD117	Heavy chain variable region, HCV Ab		US20150118242 SEQ ID NO: 2	6427
HEPBD118	Heavy chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 1; WO2014010890; CA2878155, CN104487090, EP2858674	6428
HEPBD119	Heavy chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 2; WO2014010890; CA2878155, CN104487090, EP2858674	6429
HEPBD120	Heavy chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 3; WO2014010890; CA2878155, CN104487090, EP2858674	6430
HEPBD121	Heavy chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 4; WO2014010890; CA2878155, CN104487090, EP2858674	6431
HEPBD122	Heavy chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 5; WO2014010890; CA2878155, CN104487090, EP2858674	6432
HEPBD123	Heavy chain variable region, Monoclonal HBV antibody	c18/A2	US20120308580 SEQ ID NO: 2; WO 2011062562; CN102781961, EP2501723	6433
HEPBD124	Heavy chain variable region, Neutralizing monoclonal HBV antibody		US20110097270 SEQ ID NO: 1	6434
HEPBD125	Heavy chain, full HBV Ab		US20150232537 SEQ ID NO: 9; WO2014048910; CA2884388; CN104662041A; EP2900692	6435

HEPBD126	Heavy chain, HBV Ab	HBFab21	CN103588874 SEQ ID NO: 2	6436
HEPBD127	Heavy chain, HCV Ab	Fab clone 1:5	US6747136 SEQ ID NO: 1	6437
HEPBD128	Heavy chain, HCV Ab	Fab clone 1:7	US6747136 SEQ ID NO: 2	6438
HEPBD129	Heavy chain, HCV Ab	Fab clone 1:11	US6747136 SEQ ID NO: 3	6439
HEPBD130	Heavy chain, HCV Ab	Fab clone L3	US6747136 SEQ ID NO: 4	6440
HEPBD131	Heavy chain, HCV Ab	Fab clone L1	US6747136 SEQ ID NO: 5	6441
HEPBD132	Heavy chain, HCV Ab	Fab clone A8	US6747136 SEQ ID NO: 6	6442
HEPBD133	Heavy chain, HCV Ab	Fab clone A12	US6747136 SEQ ID NO: 7	6443
HEPBD134	Heavy chain, HCV Ab	HCV-AB 68	US7241445 SEQ ID NO: 3	6444
HEPBD135	Heavy chain, HCV Ab	e8	US7727529 SEQ ID NO: 1	6445
HEPBD136	Heavy chain, HCV Ab	e10	US7727529 SEQ ID NO: 3	6446
HEPBD137	Heavy chain, HCV Ab	e20	US7727529 SEQ ID NO: 5	6447
HEPBD138	Heavy chain, HCV Ab	e137	US7727529 SEQ ID NO: 7	6448
HEPBD139	Heavy chain, HCV Ab	e301	US7727529 SEQ ID NO: 9	6449
HEPBD140	Heavy chain, HCV Ab	e509	US7727529 SEQ ID NO: 11	6450
HEPBD141	Heavy chain, HCV Ab	5D2	US20090104207 SEQ ID NO: 7	6451
HEPBD142	Heavy chain, HCV Ab	Mab V	WO2013186752 SEQ ID NO: 3	6452
HEPBD143	Heavy chain, HCV Ab	Mab VI	WO2013186752 SEQ ID NO: 5	6453
HEPBD144	Heavy chain, HCV Ab		WO2007143701 SEQ ID NO: 12	6454
HEPBD145	Heavy chain, HCV Ab	HuPA29VH#1	WO2007143701 SEQ ID NO: 15	6455
HEPBD146	Heavy chain, HCV Ab	HuPA29VH#2	WO2007143701 SEQ ID NO: 16	6456
HEPBD147	Heavy chain, HCV Ab	HuPA29VH#3	WO2007143701 SEQ ID NO: 17	6457
HEPBD148	Heavy chain, HCV Ab	PA29	WO2007143701 SEQ ID NO: 28	6458
HEPBD149	Heavy chain, HCV Ab	Ap33	Kong, L., et al., Structure of Hepatitis C Virus Envelope Glycoprotein E2 Antigenic Site 412 to 423 in Complex with Antibody AP33; J. Virol. 86 (23), 13085-13088 (2012) NCBI Accession # 4G6A_H (224aa)	6459
HEPBD150	Heavy chain, HCV Ab	Single Chain Fv Fragment 1:7	Gilmartin, A.A., et al., Protein Eng. Des. Sel. 25 (2), 59-66 (2012) NCBI Accession # 3U6R_A(149aa)	6460
HEPBD151	Heavy Chain, HCV Fab	Ar3c	Kong, L., et al., Hepatitis C virUSE2 envelope glycoprotein core structure; Science 342 (6162), 1090-1094 (2013) NCBI Accession # 4MWF_A (233aa)	6461
HEPBD152	Heavy Chain, HCV Fab	Mrct10.v362	Pantua, H., et al., Glycan shifting on hepatitis C virus (HCV) e2 glycoprotein is a mechanism for escape from broadly neutralizing antibodies; J. Mol. Biol. 425 (11), 1899-1914 (2013) NCBI Accession # HS6_H (226aa)	6462
HEPBD153	Heavy Chain, Hcv1 HCV Ab	Hcv1, P2(1) Form	Kong, L., et al., Structural basis of hepatitis C virus neutralization by broadly neutralizing antibody HCV1; Proc. Natl. Acad. Sci. U.S.A. 109 (24), 9499-9504 (2012) NCBI Accession # 4DGV_H (226aa)	6463
HEPBD154	Heavy Chain, Hcv1 HCV Ab	Hcv1, C2 Form	Kong, L., et al., Structural basis of hepatitis C virus neutralization by broadly neutralizing antibody HCV1; Proc. Natl. Acad. Sci. U.S.A. 109 (24), 9499-9504	6464

			(2012) NCBI Accession # 4DGY_H (226aa)	
HEPBD155	Heavy gamma chain variable, HCV Ab	P18-9E	US8592559 SEQ ID NO: 13	6465
HEPBD156	Heavy-chain-only, HCV Ab	VHH D03	WO2014053634 SEQ ID NO: 4	6466
HEPBD157	Heavy-chain-only, HCV Ab	VHH C09	WO2014053634 SEQ ID NO: 5	6467
HEPBD158	Heavy-chain-only, HCV Ab	Bl 1	WO2014053634 SEQ ID NO: 6	6468
HEPBD159	Heavy-chain-only, HCV Ab	D04	WO2014053634 SEQ ID NO: 7	6469
HEPBD160	Light chain full, HBV Ab		US20150232537 SEQ ID NO: 10; WO2014048910; CA2884388; CN104662041A; EP2900692	6470
HEPBD161	Light chain kappa, partial, HCV Ab		Esposito, G., et al., Recombinant human antibodies specific for hepatitis C virus proteins; Arch. Virol. 142 (3), 601-610 (1997) NCBI Accession # CAA54913.1(110aa)	6471
HEPBD162	Light chain variable domain, monoclonal HBV antibody	c18/A2	US20120308580 SEQ ID NO: 1; WO 2011062562; CN102781961, EP2501723	6472
HEPBD163	Light chain variable domain, neutralizing monoclonal HBV antibody,		US20110097270 SEQ ID NO: 9	6473
HEPBD164	Light chain variable gene, Chimeric HBV Ab		EP0521348	6474
HEPBD165	Light chain variable region, HBV Ab	E183/A2	US20120308580 SEQ ID NO: 32; WO 2011062562; CN102781961, EP2501723	6475
HEPBD166	Light chain variable region, HBV Ab	HB48-33	US8840895 SEQ ID NO: 2	6476
HEPBD167	Light chain variable region, HBV Ab	HB48-35	US8840895 SEQ ID NO: 3	6477
HEPBD168	Light chain variable region, HBV Ab	HB48-59	US8840895 SEQ ID NO: 4	6478
HEPBD169	Light chain variable region, HBV Ab	LFW22-31	US7435414 SEQ ID NO: 36; US20060014937; WO2005100400; CN1980956	6479
HEPBD170	Light chain variable region, HBV Ab	LFW22-312	US7435414 SEQ ID NO: 37; US20060014937; WO2005100400; CN1980956	6480
HEPBD171	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 216	6481
HEPBD172	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 225	6482
HEPBD173	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 234	6483
HEPBD174	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 243	6484
HEPBD175	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 252	6485
HEPBD176	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 261	6486
HEPBD177	Light chain variable region, HBV Ab		WO2013165972 SEQ ID NO: 270	6487

HEPBD 178	Light chain variable region, HBV Ab		WO2013 165972 SEQ ID NO: 279	6488
HEPBD179	Light chain variable region, HBV Ab		WO2013 165972 SEQ ID NO: 288	6489
HEPBD 180	Light chain variable region, HBV Ab		WO20 13 165972 SEQ ID NO: 297	6490
HEPBD181	Light chain variable region, HBV Ab		WO20 13 165972 SEQ ID NO: 306	6491
HEPBD 182	Light chain variable region, HBV Ab		WO20 13165972 SEQ ID NO: 3 15	6492
HEPBD183	Light chain variable region, HBV Ab		WO20 13 165972 SEQ ID NO: 324	6493
HEPBD 184	Light chain variable region, HBV Ab		WO20 13 165972 SEQ ID NO: 333	6494
HEPBD185	Light chain variable region, HBV Ab		WO2013 165972 SEQ ID NO: 342 and 351	6495
HEPBD 186	Light chain variable region, HBV Ab		WO20 13165972 SEQ ID NO: 360	6496
HEPBD187	Light chain variable region, HBV Ab		WO2013 165972 SEQ ID NO: 369	6497
HEPBD 188	Light chain variable region, HBV Ab		WO20 13165972 SEQ ID NO: 378	6498
HEPBD189	Light chain variable region, HBV Ab		WO2013 165972 SEQ ID NO: 387	6499
HEPBD 190	Light chain variable region, HBV Ab		WO20 13 165972 SEQ ID NO: 396	6500
HEPBD191	Light chain variable region, HBV Ab		WO2009069916 SEQ ID NO: 2	6501
HEPBD 192	Light chain variable region, HBV Ab	PE1-1	WO 19940 11495	6502
HEPBD193	Light chain variable region, HBV Ab	ZM1-1	WO 19940 1 1495	6503
HEPBD 194	Light chain variable region, HBV Ab	ZM1-2	WO 19940 11495	6504
HEPBD195	Light chain variable region, HBV Ab	MD3-4	WO 19940 1 1495	6505
HEPBD 196	Light chain variable region, HBV Ab	A2E2	CN 102757492 SEQ ID NO: 4	6506
HEPBD197	Light chain variable region, HBV Ab	C9G9	CN102757492 SEQ ID NO: 8	6507
HEPBD198	Light chain variable region, HBV Ab		CN104530228 SEQ ID NO: 1	6508
HEPBD199	Light chain variable region, HBV Ab		CN104530228 SEQ ID NO: 2	6509
HEPBD200	Light chain variable region, HBV Ab	Ab19	US8580256 SEQ ID NO: 1	6510
HEPBD201	Light chain variable region, HBV Ab	Ab17	US8580256 SEQ ID NO: 3	651 1
HEPBD202	Light chain variable region, HBV Ab	KR127	US8420353 SEQ ID NO: 4	6512
HEPBD203	Light chain variable region, HBV Ab	KR127	US8420353 SEQ ID NO: 2	6513
HEPBD204	Light chain variable region, HBV Ab	KR127	US8420353 SEQ ID NO: 30	6514
HEPBD205	Light chain variable region, HBV Ab	DPK12	US8420353 SEQ ID NO: 34	6515
HEPBD206	Light chain variable region, HBV Ab	HZ1	US8420353 SEQ ID NO: 38	6516

HEPBD207	Light chain variable region, HBV Ab		US71 12664 SEQ ID NO: 7; US6680053, US6924368, US2002006 l58l, US20040 191259, US200 50249753, WO2001092529	6517
HEPBD208	Light chain variable region, HBV Ab	Ab17.1.4 1	USRE39586 SEQ ID NO: 2; US6146629; WO1997047653	6518
HEPBD209	Light chain variable region, HBV Ab	Ab 19.79.5	USRE4083 l SEQ ID NO: 2	6519
HEPBD210	Light chain variable region, HBV Ab		US20150232537 SEQ ID NO: 8; WO20 140489 10; CA2884388; CN 104662041A; EP2900692	6520
HEPBD2 11	Light chain variable region, HBV Ab		US20100260712 SEQ ID NO: 2; WO2009069917	652 1
HEPBD2 12	Light chain variable region, HBV Ab		WO2015 107 126 SEQ ID NO: 4	6522
HEPBD2 13	Light chain variable region, HCV Ab	MBL-HCV1 (Antibody produced by clone 83-128)	US855 1484 SEQ ID NO: 2	6523
HEPBD2 14	Light chain variable region, HCV Ab	MBL-HCV 1 (Antibody produced by clone 95-2)	1JS855 1484 SEQ ID NO: 4	6524
HEPBD2 !5	Light chain variable region, HCV Ab	MBL-HCV! (Antibody produced by clone 073- 1)	US855 l484 SEQ ID NO: 6	6525
HEPBD216	Light chain variable region, HCV Ab	Clone 13	US7250166 SEQ ID NO: 2	6526
HEPBD217	Light chain variable region, HCV Ab	Clone 98	US7250166 SEQ ID NO: 4	6527
HEPBD218	Light chain variable region, HCV Ab	Clone 1:4	US7250166 SEQ ID NO: 6	6528
HEPBD219	Light chain variable region, HCV Ab	Clone 1:8	US7250166 SEQ ID NO: 8	6529
HEPBD220	Light chain variable region, HCV Ab	Clone 1:9	US7250166 SEQ ID NO: 10	6530
HEPBD22 l	Light chain variable region, HCV Ab	Clone 1:10	US7250166 SEQ ID NO: 12	653 1
HEPBD222	Light chain variable region, HCV Ab	Clone 4:6	US7250166 SEQ ID NO: 14	6532
HEPBD223	Light chain variable region, HCV Ab	Clone 6a: 5	US7250166 SEQ ID NO: 16	6533
HEPBD224	Light chain variable region, HCV Ab	Clone 2a: 2	US7250166 SEQ ID NO: 18	6534
HEPBD225	Light chain variable region, HCV Ab	Clone 2a: 4	US7250166 SEQ ID NO: 20	6535
HEPBD226	Light chain variable region, HCV Ab	Clone 2a: 5	US7250166 SEQ ID NO: 22	6536
HEPBD227	Light chain variable region, HCV Ab	Clone 2a: 13	US7250166 SEQ ID NO: 24	6537
HEPBD228	Light chain variable region, HCV Ab	Clone 2a: 14	US7250166 SEQ ID NO: 26	6538
HEPBD229	Light chain variable region, HCV Ab	Clone 2a:23	US7250166 SEQ ID NO: 28	6539
HEPBD230	Light chairs variable region, HCV Ab	Clone 2a: 23	US7250166 SEQ ID NO: 30	6540
HEPBD23 l	Light chain variable region, HCV Ab	Clone 2a: 25	US7250 166 SEQ ID NO: 32	654 l

HEPBD232	Light chain variable region, HCV Ab	Clone 2a:30	**US7250166** SEQ ID NO: 34	6542
HEPBD233	Light chain **variable** region, HCV Ab	Clone 2a:32	**US7250166** SEQ ID NO: 36	6543
HEPBD234	Light chain **variable** region, HCV Ab	Clone 2a:33	**US7250 166** SEQ ID NO: 38	6544
HEPBD235	Light chain variable region, HCV **Ab**	Clone **2a:37**	US7250166 SEQ ID NO: 40	6545
HEPBD236	Light chain **variable** region, HCV Ab	Clone **2a:40**	**US7250 166** SEQ ID NO: 42	6546
HEPBD237	Light chain variable region, HCV **Ab**	Clone 2b:1	**US7250166** SEQ ID NO: 44	6547
HEPBD238	Light **chain variable** region, HCV Ab	Clone 2b:3	US7250166 SEQ ID NO: 46	6548
HEPBD239	Light chain variable region, HCV **Ab**	Clone 2b:4	**US7250166** SEQ ID NO: 48	6549
HEPBD240	Light **chain variable** region, HCV Ab	Clone 2b:5	US7250166 SEQ ID NO: 50	6550
HEPBD241	Light chain variable region, HCV **Ab**	Clone 2b:7	**US7250166** SEQ ID NO: 52	6551
HEPBD242	Light **chain variable** region, HCV Ab	Clone 2b:9	US7250166 SEQ ID NO: 54	6552
HEPBD243	Light chain variable region, HCV Ab	Clone 2b:10	1JS7250166 SEQ ID NO: 56	6553
HEPBD244	Light chain variable region, HCV Ab	**anti-NS3** Fab	US7314919 SEQ ID NO: 6	6554
HEPBD245	Light chain variable region, HCV Ab		**US7507408** SEQ ID NO: **2**	6555
HEPBD246	Light chain variable region, HCV Ab		US7507408 SEQ ID NO: 4	6556
HEPBD247	Light chain variable region, HCV Ab		1JS7507408 SEQ ID NO: 6	6557
HEPBD248	Light chain variable region, HCV Ab	Antibody produced by clone 95-14	US8551484 SEQ ID NO: 44	6558
HEPBD249	Light chain variable region, HCV Ab	Antibody produced **by** clone 95-38	US8551484 SEQ ID **NO:** 53	6559
HEPBD250	Light **chain variable** region, HCV **Ab**	**HC-84.1**	**US20130084301** SEQ ID NO: 64	6560
HEPBD251	Light **chain variable** region, HCV Ab	HC-84.20	**US20 13008430**1 SEQ **ID NO:** 65	6561
HEPBD252	Light chain variable region, HCV **Ab**	**HC-84.21**	**US20 130084301** SEQ **ID** NO: 66	6562
HEPBD253	Light chain **variable** region, HCV Ab	HC-84.22	**US20 13008430**1 SEQ **ID NO:** 67	6563
HEPBD254	Light chain variable region, HCV **Ab**	**HC-23**	**US20 130084301** SEQ **ID** NO: 68	6564
HEPBD255	Light chain **variable** region, HCV Ab	**HC-84.24**	**US20 13008430**1 SEQ **ID NO:** 69	6565
HEPBD256	Light chain variable region, HCV **Ab**	HC-84.25	**US20 130084301** SEQ **ID** NO: 70	6566
HEPBD257	Light chain **variable** region, HCV Ab	**HC-84.26**	**US2013008430**1 SEQ **ID NO: 71**	6567
HEPBD258	Light chain variable region, HCV Ab	HC-84.27.	**US20 130084301** SEQ **ID** NO: 72	6568
HEPBD259	Light chain variable region, HCV Ab	AOT3	**US20120009 196** SEQ **ID** NO: 2	6569

HEPBD260	Light chain variable region, HCV Ab	C11-3	US20120009196 SEQ ID NO: 4	6570
HEPBD261	Light chain variable region, HCV Ab	C11-7	US20120009196 SEQ ID NO: 6	6571
HEPBD262	Light chain variable region, HCV Ab	C11-9	US20120009196 SEQ ID NO: 8	6572
HEPBD263	Light chain variable region, HCV Ab	C11-14	US20120009196 SEQ ID NO: 10	6573
HEPBD264	Light chain variable region, HCV Ab		WO2014065822 SEQ ID NO: 5	6574
HEPBD265	Light chain variable region, HCV Ab		WO2014065822 SEQ ID NO: 9	6575
HEPBD266	Light chain variable region, HCV Ab	Antibody light chain from WO2007143701	WO2007143701 SEQ ID NO: 1	6576
HEPBD267	Light chain variable region, HCV Ab	Hc33.1	Li Y. et al., Structural basis for penetration of the glycan shield of hepatitis C virUSe2 glycoprotein by a broadly neutralizing human antibody; J. Biol. Chem. 290 (16), 10117-10125 (2015) NCBI Accession # 4XVJ_L (115aa)	6577
HEPBD268	Light chain variable region, HCV Ab		US20150118242 SEQ ID NO: 3	6578
HEPBD269	Light chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 6; WO2014010890; CA2878155, CN104487090, EP2858674	6579
HEPBD270	Light chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 7; WO2014010890; CA2878155, CN104487090, EP2858674	6580
HEPBD271	Light chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 8; WO2014010890; CA2878155, CN104487090, EP2858674	6581
HEPBD272	Light chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 9; WO2014010890; CA2878155, CN104487090, EP2858674	6582
HEPBD273	Light chain variable region, Human HBV antibody that binds to the surface antigen (HBsAg)		US20150166637 SEQ ID NO: 10; WO2014010890; CA2878155, CN104487090, EP2858674	6583
HEPBD274	Light chain variable region, partial, HCV Ab		Keck, Z.Y., et al., Human monoclonal antibody to hepatitis C virUSE1 glycoprotein that blocks virus attachment and viral infectivity; J. Virol. 78 (13), 7257-7263 (2004) NCBI Accession # AAS47840 (147aa)	6584
HEPBD275	Light chain, HCV Ab	Hu5b3.v3	Pantua, H., et al., Glycan shifting on hepatitis C virus (HCV) e2 glycoprotein is a mechanism for escape from broadly neutralizing antibodies; J. Mol. Biol. 425	6585

			(11), 1899-1914 (2013) NCBI Accession # 4HS8_L (218aa)	
HEPBD276	Light chain, HCV Ab	Ap33	Kong, L., et al., Structure of Hepatitis C Virus Envelope Glycoprotein E2 Antigenic Site 412 to 423 in Complex with Antibody AP33; J. Virol. 86 (23), 13085-13088 (2012) NCBI Accession # 4G6A_L (218aa)	6586
HEPBD277	Light chain, HBV Ab	HBFab21	CN103588874 SEQ ID NO: 1	6587
HEPBD278	Light chain, HCV Ab	Fab clone 1:5	US6747136 SEQ ID NO: 8	6588
HEPBD279	Light chain, HCV Ab	Fab clone 1:7	US6747136 SEQ ID NO: 9	6589
HEPBD280	Light chain, HCV Ab	Fab clone 1:11	US6747136 SEQ ID NO: 10	6590
HEPBD281	Light chain, HCV Ab	Fab clone L3	US6747136 SEQ ID NO: 11	6591
HEPBD282	Light chain, HCV Ab	Fab clone L1	US6747136 SEQ ID NO: 12	6592
HEPBD283	Light chain, HCV Ab	Fab clone A8	US6747136 SEQ ID NO: 13	6593
HEPBD284	Light chain, HCV Ab	Fab clone A12	US6747136 SEQ ID NO: 14	6594
HEPBD285	Light chain, HCV Ab	HCV#1	US6924362 SEQ ID NO: 1	6595
HEPBD286	Light chain, HCV Ab	HCV#4	US6924362 SEQ ID NO: 2	6596
HEPBD287	Light chain, HCV Ab	HCV#7	US6924362 SEQ ID NO: 3	6597
HEPBD288	Light chain, HCV Ab	HCV#12	US6924362 SEQ ID NO: 4	6598
HEPBD289	Light chain, HCV Ab	HCV#13	US6924362 SEQ ID NO: 5	6599
HEPBD290	Light chain, HCV Ab	HCV-AB 68	US7241445 SEQ ID NO: 4	6600
HEPBD291	Light chain, HCV Ab	e8	US7727529 SEQ ID NO: 2	6601
HEPBD292	Light chain, HCV Ab	e10	US7727529 SEQ ID NO: 4	6602
HEPBD293	Light chain, HCV Ab	e20	US7727529 SEQ ID NO: 6	6603
HEPBD294	Light chain, HCV Ab	e137	US7727529 SEQ ID NO: 8	6604
HEPBD295	Light chain, HCV Ab	e301	US7727529 SEQ ID NO: 10	6605
HEPBD296	Light chain, HCV Ab	e509	US7727529 SEQ ID NO: 12	6606
HEPBD297	Light chain, HCV Ab	5D2	US20090104207 SEQ ID NO: 8	6607
HEPBD298	Light chain, HCV Ab	MabV	WO2013186752 SEQ ID NO: 4	6608
HEPBD299	Light chain, HCV Ab	Mab VI	WO2013186752 SEQ ID NO: 6	6609
HEPBD300	Light chain, HCV Ab		WO2007143701 SEQ ID NO: 11	6610
HEPBD301	Light chain, HCV Ab	HuPA29VH#1	WO2007143701 SEQ ID NO: 18	6611
HEPBD302	Light chain, HCV Ab	HuPA29VH#2	WO2007143701 SEQ ID NO: 19	6612
HEPBD303	Light chain, HCV Ab	PA29	WO2007143701 SEQ ID NO: 29	6613
HEPBD304	Light chain, HCV Ab	Single Chain Fv Fragment 1:7	Gilmartin, A.A., et al., Protein Eng. Des. Sel. 25 (2), 59-66 (2012) NCBI Accession # 3U6R_B (143aa)	6614
HEPBD305	Light chain, HCV Ab	Igh526	Kong L., et al., Structure of Hepatitis C Virus Envelope Glycoprotein E1 Antigenic Site 314-324 in Complex with Antibody IGH526; J. Mol. Biol. 427 (16), 2617-2628 (2015) NCBI Accession # 4N0Y_L (218aa)	6615
HEPBD306	Light chain, HCV Fab	Ar3c	Kong, L., et al., Hepatitis C virUSE2 envelope glycoprotein core structure; Science 342 (6162), 1090-1094 (2013) NCBI Accession # 4MWF_B (214aa)	6616
HEPBD307	Light chain, HCV Fab	Mrct10.v362	Pantua, H., et al., Glycan shifting on hepatitis C virus (HCV) e2 glycoprotein is a mechanism for escape from broadly	6617

			neutralizing antibodies; J. Mol. Biol. 425 (11), 1899-1914 (2013) NCBI Accession # 4HS6_L (218aa)	
HEPBD308	Light chain, Hcv1 HCV Ab	Hcv1, C2 Form	Kong, L., et al., Structural basis of hepatitis C virus neutralization by broadly neutralizing antibody HCV1; Proc. Natl. Acad. Sci. U.S.A. 109 (24), 9499-9504 (2012) NCBI Accession # 4DGY_L (213aa)	6618
HEPBD309	Light chain, Hcv1 HCV Ab	Hcv1, P2(1) Form	Kong, L., et al., Structural basis of hepatitis C virus neutralization by broadly neutralizing antibody HCV1; Proc. Natl. Acad. Sci. U.S.A. 109 (24), 9499-9504 (2012) NCBI Accession # 4DGV_L (213aa)	6619
HEPBD310	Light kappa chain variable, HCV Ab	P18-9E	US8592559 SEQ ID NO: 14	6620
HEPBD311	PEGylated anti-E2 heavy chain, HCV Ab		WO2006028634 SEQ ID NO: 1	6621
HEPBD312	PEGylated anti-E2 heavy chain, HCV Ab		WO2006028634 SEQ ID NO: 2	6622
HEPBD313	PEGylated anti-E2 heavy chain, HCV Ab		WO2006028634 SEQ ID NO: 3	6623
HEPBD314	PEGylated anti-E2 heavy chain, HCV Ab		WO2006028634 SEQ ID NO: 4	6624
HEPBD315	PEGylated anti-E2 heavy chain, HCV Ab		WO2006028634 SEQ ID NO: 8	6625
HEPBD316	single chain, HBV Ab		US6562599 SEQ ID NO: 4	6626
HEPBD317	single chain, HBV Ab		US6562599 SEQ ID NO: 6	6627

[00320] In one embodiment, the payload region of the AAV particle composes one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Patent No. US7241445, and US8858947, the contents of each of which are herein incorporated by reference in their entirety, against HCV.

[00321] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Publication No. US20150072885 and US20110046354, US Patent No. US5204095, European Publication No. EP023292L, EP0038642, and EP0186371, and International Publication No. WO1994011495, the contents of each of which are herein incorporated by reference in their entirety, against HBV.

[00322] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Patent No. US6020195, the contents of which are herem incorporated by reference in their entirety, against HGV (hepatitis G vims).

[00323] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 24 against Herpes Virus.

Table 24. Antibodies against Herpesvirus

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
HERP1	Chain A, HSV	E317 Fab	Lee, C. et al., "Structural basis for the antibody neutralization of herpes simplex virus" Acta Crystallogr. D Biol. Crystallogr. 69 (PT 10), 1935-1945 (2013), NCBI Accession # 3W9D_A	6628
HERP2	Chain B, HSV	E317 Fab	Lee, C. et al., "Structural basis for the antibody neutralization of herpes simplex virus" Acta Crystallogr. D Biol. Crystallogr. 69 (PT 10), 1935-1945 (2013), NCBI Accession # 3W9D_B	6629
HERP3	Chain C, HSV	E317 Fab	Lee, C. et al., "Structural basis for the antibody neutralization of herpes simplex virus" Acta Crystallogr. D Biol. Crystallogr. 69 (PT 10), 1935-1945 (2013), NCBI Accession # 3W9D_C	6630
HERP4	Chain D, HSV	E317 Fab	Lee, C. et al., "Structural basis for the antibody neutralization of herpes simplex virus" Acta Crystallogr. D Biol. Crystallogr. 69 (PT 10), 1935-1945 (2013), NCBI Accession # 3W9D_D	6631
HERP5	Chimeric anti-EBVs gp350 antibody		Tanner, J.E., "Peptides Designed To Spatially Depict the Epstein-Barr Virus Major Virion Glycoprotein gp350 Neutralization Epitope Elicit Antibodies That Block Virus-Neutralizing Antibody 72A1 Interaction with the Native gp350 Molecule"", J. Virol. 89 (9), 4932-4941 (2015), NCBI Accession #AJR20276	6632
HERP6	Chimeric anti-EBVs gp350 antibody		Tanner, J.E., "Peptides Designed To Spatially Depict the Epstein-Barr Virus Major Virion Glycoprotein gp350 Neutralization Epitope Elicit Antibodies That Block Virus-Neutralizing Antibody 72A1 Interaction with the Native gp350 Molecule"", J. Virol. 89 (9), 4932-4941 (2015), NCBI Accession #AJR20275	6633
HERP7	CMV	AE11F/3-20L1	Lantto, J. et al., Binding characteristics determine the neutralizing potential of antibody fragments specific for antigenic domain 2 on glycoprotein B of human cytomegalovirus, Virology 305 (1), 201-209 (2003), NCBI Accession # AAN87569.1 (256 aa)	6634
HERP8	Fv, EBV	G5	Fang, C.Y., "Modulation of Epstein-Barr virus latent membrane protein 1 activity by intrabodies", Intervirology 50 (4), 254-263 (2007), NCBI Accession #ABA55015	6635
HERP9	Fv, EBV	A4	Fang, C.Y., "Modulation of Epstein-Barr virus latent membrane protein 1 activity by intrabodies", Intervirology 50 (4), 254-263 (2007), NCBI Accession #ABA55014	6636
HERP10	Fv, EBV	B8	Fang, C.Y., "Modulation of Epstein-Barr virus latent membrane protein 1 activity by intrabodies", Intervirology 50 (4), 254-263 (2007), NCBI **Accession #ABA55013**	6637

HERP1 1	Fv, **EBV**	F5	Fang, C.Y., "Modulation of **Epstein-Barr** virus latent membrane protein **1** activity by **intrabodies", Intervirology** 50 (4), 254-263 (2007), NCBI Accession #ABA550 12	6638
HERP 12	Fv, EBV	E2	Fang, C.Y., "Modulation of Epstein-Barr virus latent membrane protein 1 activity by intrabodies", **Tntervirology** 50 (4), 254-263 (2007), NCBI Accession #ABA5501 1	6639
HERP 13	Fv, EBV	H3	Fang, C.Y., **"Modulation of Epstein-Barr vims latent** membrane protein **1** activity by intrabodies", **Intervirology** 50 **(4),** 254-263 (2007), NCBI Accession **#ABA55010**	6640
HERP14	Heavy chain, FLAGhis tagged sequence, VZV	DDF-VZV1	US20100074906 SEQ ID NO: 20	6641
HERP 15	Heavy chain **variable domain,** clone **11, HSV**	ACHDV1	**Burioni,** R. et al. "Recombinant human Fab to glycoprotein **D** neutralizes **iirfectivity** and prevents cell-to-cell transmission **of herpes** simplex viruses 1 and 2 in vitro", Proc. **Natl.** Acad. Sci. U.S.A. 91 **(1),** 355-359 (1994), NCBI Accession # AAB29447	6642
HERP 16	Heavy chain variable **domain,** clone 13, HSV	**ACHDV1**	Burioni, **R. et al. "Recombinant** human Fab to glycoprotein D **neutralizes iirfectivity** and prevents cell-to-cell transmission **of herpes** simplex viruses 1 and 2 in vitro", Proc. **Natl.** Acad. Sci. U.S.A. 91 **(1),** 355-359 (1994), NCBI Accession # AAB29449	6643
HERP **17**	Heavy chain variable **domain,** clone 15, HSV	ACHDV2	Burioni, **R. et al.** "Recombinant human Fab to glycoprotein D **neutralizes iirfectivity** and prevents cell-to-cell transmission **of herpes** simplex viruses 1 and 2 in vitro", Proc. **Natl.** Acad. Sci. U.S.A. 91 (I), 355-359 (1994), NCBI Accession # AAB29456	6644
HERP 18	Heavy chain variable domain, clone 15, HSV	**ACHDV1**	Burioni, **R. et al.** "Recombinant human Fab to glycoprotein D neutralizes **iirfectivity** and prevents **cell-to-ceU transmission of herpes** simplex viruses 1 and 2 **in** vitro", Proc. **Natl.** Acad. Sci. U.S.A. 91 (I), 355-359 (1994), NCBI Accession # AAB29450	6645
HERP 19	**Heavy** chain variable domain, clone 18, HSV	**ACHDV1**	Burioni, **R.** et **al.** "Recombinant human Fab **to** glycoprotein D neutralizes **iirfectivity** and prevents **cell-to-ceU transmission of herpes** simplex viruses 1 and 2 **in** vitro", Proc. **Natl.** Acad. Sci. U.S.A. 91 (I), 355-359 (1994), NCBI Accession # **AAB29448**	6646
HERP20	**Heavy** chain variable domain, clone 2, HSV	**ACHDV1**	Burioni, **R.** et **al.** "Recombinant human Fab **to** glycoprotein D neutralizes **iirfectivity** and prevents **cell-to-cell** transmission **of herpes** simplex viruses 1 **and 2 in vitro",** Proc. **Natl.** Acad. Sci. U.S.A. 91 (1), 355-359 (1994), NCBI Accession # AAB29455	6647
HERP21	**Heavy chain** variable region, **CMV**	1F7	US82025 18 SEQ ID NO: 5	6648
HERP22	**Heavy chain** variable region, CMV	Humanized 57.4	**WO20 14200898** SEQ ID NO: 633	6649
HERP23	Heavy **chain** variable **region,** CMV	**Humanized** 57.4	**WO20 14200898** SEQ **ID NO:** 634	6650

HERP24	Heavy chain variable region, CMV	Humanized 58.5	WO2014200898 SEQ ID NO: 637	6651
HERP25	Heavy chain variable region, CMV	Humanized 58.5	WO2014200898 SEQ ID NO: 638	6652
HERP26	Heavy chain variable region, CMV	Humanized 272.7	WO2014200898 SEQ ID NO: 641	6653
HERP27	Heavy chain variable region, CMV	Humanized 276.10	WO2014200898 SEQ ID NO: 644	6654
HERP28	Heavy chain variable region, CMV	Humanized 276.10	WO2014200898 SEQ ID NO: 645	6655
HERP29	Heavy chain variable region, CMV	Sm5-1	Li, B., Construction and characterization of a high-affinity humanized SM5-1 monoclonal antibody, Biochem. Biophys. Res. Commun. 357 (4), 951-956 (2007), NCBI Accession # ABI22831.1	6656
HERP30	Heavy chain variable region, CMV		Schoppel, K. et al., Antibodies specific for the antigenic domain 1 of glycoprotein B (gpUL55) of human cytomegalovirus bind to different substructures, Virology 216 (1), 133-145 (1996), NCBI Accession # AAB26953.1 (163 aa)	6657
HERP31	Heavy chain variable region, CMV		Schoppel, K. et al., Antibodies specific for the antigenic domain 1 of glycoprotein B (gpUL55) of human cytomegalovirus bind to different substructures, Virology 216 (1), 133-145 (1996), NCBI Accession # AAB26952.1 (161 aa)	6658
HERP32	Heavy chain variable region, CMV		Schoppel, K. et al., Antibodies specific for the antigenic domain 1 of glycoprotein B (gpUL55) of human cytomegalovirus bind to different substructures, Virology 216 (1), 133-145 (1996), NCBI Accession # AAB26951.1 (158 aa)	6659
HERP33	Heavy chain variable region, CMV		Potzsch, S., B Cell Repertoire Analysis Identifies New Antigenic Domains on Glycoprotein B of Human Cytomegalovirus which Are Target of Neutralizing Antibodies, NCBI Accession # AEF33814.1	6660
HERP34	Heavy chain variable region, CMV, a complex of human cytomegalovirus (hCMV) proteins UL130 and UL131A	1F11	US9149524 SEQ ID NO: 7	6661
HERP35	Heavy chain variable region, CMV, a complex of human cytomegalovirus (hCMV) proteins UL130 and UL131A	2F4	US9149524 SEQ ID NO: 17	6662

HERP36	Heavy chain variable region, CMV, a complex of human cytomegalovirus (hCMV) proteins UL 130 and UL13 1A	5A2	US9149524 SEQ ID NO: 39	6663
HERP37	Heavy chain variable region CMV, AD 1 region of HCMV glycoprotein gB	EV2038	US8492529 SEQ ID NO: 10	6664
HERP38	Heavy chain variable region, EBV		1JS20 150064 174 SEQ ID 1	6665
HERP39	Heavy chain variable region, EBV		US20 150064 174 SEQ ID 2	6666
HERP40	Heavy chain variable region, gH glycoprotein of HCMV	HCMV 16	WO1994009136, Fig. !	6667
HERP41	Heavy chain variable region, HSV		Nejatollahi, F. and Bagheri, V., "Isolation of neutralizing human specific single-chain antibodies against Herpes Simplex Virus type 1 glycoprotein D", unpublished, NCBI Accession # AGO590 15	6668
HERP42	Heavy chain variable region, HSV I&2	E3 i7	US843 1118 SEQ ID NO: 1; US8252906	6669
HERP43	Heavy chain variable region, HSV I&2	E425	US843 1118 SEQ ID NO: 3; US8252906	6670
HERP44	Heavy chain variable region, HSV 1&2	Y573	US843 1118 SEQ ID NO: 4!; US8252906	6671
HERP45	Heavy chain variable region, VZV		1JS5506132 SEQ ID NO: 4	6672
HERP46	Heavy chain variable region, VZV	DDF-VZV2	US20100074906 SEQ ID NO: 26	6673
HERP47	Heavy chain without a signal sequence, CMV, AD 1 region of HCMV glycoprotein gB	EV2038	US8492529 SEQ ID NO: 6	6674
HERP48	Heavy chain, CMV	8f9	McLean, G.R. et al., Recognition of human cytomegalovirus by human primary immunoglobulins identifies an innate foundation to an adaptive immune response, J. Immunol. 174 (8), 4768-4778 (2005), NCBI Accession # CAE54374.1	6675
HERP49	Heavy chain, CMV	Mab 109	Simpson, J.A. et al., Neutralizing monoclonal antibodies that distinguish three antigenic sites on human cytomegalovirus glycoprotein H have conformationalry distinct binding sites, J.	6676

			Virol. 67 (1), 489-496 (1993), **NCBI** Accession # **AAB24505. 1** (119 **aa)**	
HERP50	**Heavy** chain, **CMV**	Mab **115**	Simpson, **J**.A. et al., Neutralizing monoclonal antibodies thai **distinguish** three antigenic sites on human cytomegalovirus glycoprotein H **have** conformational!}' distinct binding sites, **J**. Virol. 67 (1), 489-496 (1993), NCBI Accession # **AAB24504. 1** (117 **aa)**	6677
HERP5 1	Heavy chain, CMV	Mab **33**	Simpson, **J.A.** et **al.,** Neutralizing monoclonal antibodies thai **distinguish** three antigenic sites on human cytomegalovirus glycoprotein H **have** confonnationaliy distinct binding sites, **J**. Virol. 67 (1), 489-496 (1993), NCBI Accession # **AAB24503. 1 (120 aa)**	6678
HERP52	Heavy chain, CMV	Mab 5	Simpson, **J.A.** et **al.,** Neutralizing monoclonal antibodies thai **distinguish** three antigenic sites on human cytomegalovirus glycoprotein H have **conformational!}'** distinct binding sites, **J**. Virol. 67 (1), 489-496 (1993), NCBI **Accession** # **AAB24502. 1** (120 **aa)**	6679
HERP53	**Heavy** chain, **CMV,** a combination of the **hCMV** proteins UL1 **28.** UL 130 and **UL13 1A**	6G4	**WO20 10007463** SEQ ID NO: 7	6680
HERP54	Heavy chain, HHV-6		US20140093526 SEQ ID 12	6681
HERP55	**Heavy chain, HSv l&2**	FabHSV 8.	US61563 13 SEQ ID NO: 2	6682
HERP56	Heavy chain, **HSv l&2**	64-683	US5646041 SEQ ID **NO:** 2; EP876478	6683
HERP57	Heavy **chain,** HSV **I&2**	**H005157**	**US20140302062** SEQ ID **NO:** 3	6684
HERP58	**Heavy** chain, **HSV l&2**	H005158	**US20140302062** SEQ ID NO: 4	6685
HERP59	Heavy **chain,** HSV **I&2**	H005159	**US20140302062** SEQ ID **NO:** 5	6686
HERP60	**Heavy** chain, **HSV l&2**	H005160	**US20140302062** SEQ ID NO: **6**	6687
HERP6 1	Heavy **chain,** HSV **I&2**	H005188	**US20140302062** SEQ ID **NO:** 7	6688
HERP62	**Heavy** chain, **HSV l&2**	H005190	**US20140302062** SEQ ID NO: 8	6689
HERP63	Heavy **chain,** HSV **I&2**	H005192	**US20140302062** SEQ ID **NO:** 9	6690
HERP64	Light chain variable region, gH glycoprotein **of HCMV**	**HCMV16**	**WO1994009136, Fig.** 2	6691
HERP65	Light **chain** recombinant, **VZV**	DDF-VZV1	US20100074906 SEQ ID NO: 22	6692
HERP66	Light chain variable region, CMV	1F7	**US82025 18** SEQ ID NO: 10	6693
HERP67	Light chain variable region, **CMV**	**Humanized** 57.4	**WO20 14200898** SEQ ID NO: 63 **1**	6694

HERP68	Light chain variable region, CMV	Humanized 57.4	WO2014200898 SEQ ID NO: 632	6695
HERP69	Light chain variable region, CMV	Humanized 58.5	WO2014200898 SEQ ID NO: 635	6696
HERP70	Light chain variable region, CMV	Humanized 58.5	WO2014200898 SEQ ID NO: 636	6697
HERP71	Light chain variable region, CMV	Humanized 272.7	WO2014200898 SEQ ID NO: 639	6698
HERP72	Light chain variable region, CMV	Humanized 272.7	WO2014200898 SEQ ID NO: 640	6699
HERP73	Light chain variable region, CMV	Humanized 276.10	WO2014200898 SEQ ID NO: 642	6700
HERP74	Light chain variable region, CMV	Humanized 276.10	WO2014200898 SEQ ID NO: 643	6701
HERP75	Light chain variable region, CMV	Sm5-1	Li, B., Construction and characterization of a high-affinity humanized SM5-1 monoclonal antibody, Biochem. Biophys. Res. Commun. 357 (4), 951-956 (2007), NCBI Accession # ABI22832.1	6702
HERP76	Light chain variable region, CMV	8f9	Schoppel, K. et al., Antibodies specific for the antigenic domain 1 of glycoprotein B (gpUL55) of human cytomegalovirus bind to different substructures, Virology 216 (1), 133-145 (1996), NCBI Accession # AAB26956.1 (146 aa)	6703
HERP77	Light chain variable region, CMV		Schoppel, K. et al., Antibodies specific for the antigenic domain 1 of glycoprotein B (gpUL55) of human cytomegalovirus bind to different substructures, Virology 216 (1), 133-145 (1996), NCBI Accession # AAB26955.1 (141 aa)	6704
HERP78	Light chain variable region, CMV		Schoppel, K. et al., Antibodies specific for the antigenic domain 1 of glycoprotein B (gpUL55) of human cytomegalovirus bind to different substructures, Virology 216 (1), 133-145 (1996), NCBI Accession # AAB26954.1 (152 aa)	6705
HERP79	Light chain variable region, CMV		Potzsch, S., B Cell Repertoire Analysis Identifies New Antigenic Domains on Glycoprotein B of Human Cytomegalovirus which Are Target of Neutralizing Antibodies, NCBI Accession # AEF33824.1	6706
HERP80	Light chain variable region, CMV, a combination of the hCMV proteins UL128, UL130 and UL131A	1F11	US9149524 SEQ ID NO: 8	6707
HERP81	Light chain variable region, CMV, a	2F4	US9149524 SEQ ID NO: 18	6708

	combination of the hCMV proteins UL1 28. UL 130 and UL131A			
HERP82	Light chain variable region, CMV, a combination of the hCMV proteins UL1 28, UL 130 and UL13 1A	5A2	US9149524 SEQ ID NO: 40	6709
HERP83	Light chain variable region, CMV, AD 1 region of HCMV glycoprotein gB	EV2038	US8492529 SEQ ID NO: 12	6710
HERP84	Light chain variable region, EBV		US20 150064 174 SEQ ID 3	6711
HERP85	Light chain variable region, EBV		US20 150064 174 SEQ ID 4	6712
HERP86	Light chain variable region, HSV		Nejatollahi, F. and Bagheri, V., "Isolation of neutralizing human specific single-chain antibodies against Herpes Simplex Vims type 1 glycoprotein D", unpublished", NCBI Accession # AGO590 16	6713
HERP87	Light chain variable region, HSV 1&2	E317	US843 1118 SEQ ID NO: 2; US8252906	6714
HERP88	Light chain variable region, HSV 1&2	E425	US843 1118 SEQ ID NO: 4; US8252906	6715
HERP89	Light chain variable region, HSV I&2	Y571	US843 1118 SEQ ID NO: 42; US8252906	6716
HERP90	Light chain variable region, **vzv**		US5506132 SEQ ID NO: 2	6717
HERP91	Light chain variable region, VZV	DDF-VZV2	US20100074906 SEQ ID NO: 24	6718
HERP92	Light chain without a signal sequence, CMV, AD1 region of HCMV glycoprotein gB	EV2038	US8492529 SEQ ID NO: 8	6719
HERP93	Light chain. CMV	8f9	McLean, G.R. et al. Recognition of human cytomegalovirus by human primary immunoglobulins identifies an innate foundation to an adaptive immune response, J. Immunol. 174 (8), 4768-4778 (2005), NCBE Accession # CAE54366. 1	6720
HERP94	Light chain. CMV	Mab 109	Simpson, J.A. et al., Neutralizing monoclonal antibodies that distinguish three antigenic sites on human cytomegalovirus glycoprotein H have conformational!}' distinct binding sites, J.	6721

			Virol. 67 (1), 489-496 (1993), NCBI Accession # AAB24501.1 (111 aa)	
HERP95	Light chain, CMV	Mab 115	Simpson, J.A. et al., Neutralizing monoclonal antibodies that distinguish three antigenic sites on human cytomegalovirus glycoprotein H have conformationally distinct binding sites, J. Virol. 67 (1), 489-496 (1993), NCBI Accession # AAB24500.1 (107 aa)	6722
HERP96	Light chain, CMV	Mab 33	Simpson, J.A. et al., Neutralizing monoclonal antibodies that distinguish three antigenic sites on human cytomegalovirus glycoprotein H have conformationally distinct binding sites, J. Virol. 67 (1), 489-496 (1993), NCBI Accession # AAB24499.1 (107 aa)	6723
HERP97	Light chain, CMV	Mab 5	Simpson, J.A. et al., Neutralizing monoclonal antibodies that distinguish three antigenic sites on human cytomegalovirus glycoprotein H have conformationally distinct binding sites, J. Virol. 67 (1), 489-496 (1993), NCBI Accession # AAB24498.1 (107 aa)	6724
HERP98	Light chain, CMV, a combination of the hCMV proteins UL128, UL130 and UL131A	6G4	WO2010007463 SEQ ID NO: 8	6725
HERP99	Light chain, HHV-6		US20140093526 SEQ ID 10	6726
HERP100	Light chain, HSV 1&2	64-683	US5646041 SEQ ID NO: 4; EP876478	6727
HERP101	Light chain, HSV 1&2	K003927	US20140302062 SEQ ID NO: 10	6728
HERP102	Light chain, HSV 1&2	K003928	US20140302062 SEQ ID NO: 11	6729
HERP103	Light chain, HSV 1&2	K003929	US20140302062 SEQ ID NO: 12	6730
HERP104	Light chain, HSV 1&2	K003930	US20140302062 SEQ ID NO: 13	6731
HERP105	Light chain, HSV 1&2	K003946	US20140302062 SEQ ID NO: 14	6732
HERP106	Light chain, HSV 1&2	K003948	US20140302062 SEQ ID NO: 15	6733
HERP107	Light chain, HSV 1&2	K003949	US20140302062 SEQ ID NO: 16	6734
HERP108	Light chain, HSV 1&2	L001844	US20140302062 SEQ ID NO: 17	6735
HERP109	Single chain Fv antibody, glycoprotein B recombinant, CMV		Lantto, J. et al., Non-germ-line encoded residues are critical for effective antibody recognition of a poorly immunogenic neutralization epitope on glycoprotein B of human cytomegalovirus, Eur. J. Immunol. 32 (6), 1659-1669 (2002), NCBI Accession # AAM92769.1 (255 aa)	6736

[00324] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, **fragments** or variants thereof descnbed in International Publication No. WO2010109874, **and**

WO1997026329, the contents of each of which are herein incorporated by reference in their entirety, against HSV.

[00325] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in international Publication No. WO199503546, the contents of which are herein incorporated by reference in their entirety, against *WIN* .

[00326] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 25 against Coronavirus.

Table 25. Antibodies against Coronaviruses

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
CORV1	Heavy chain partial sequence, Human anti-SARS antibody, Ig		Liu, J., unpublished, NCBI Accession # BAE94186.1(228aa)	6737
CORV2	Heavy chain partial sequence Human SARS neutralization antibody, Ig	H12	AAX19356.1(127aa)	6738
CORV3	Heavy chain variable partial sequence, Human neutralizing SARS antibody		Leung et al., PLoS Med. 3 (7), E237 (2006), NCBI Accession # ABA54614.1(113aa)	6739
CORV4	Heavy chain variable region, Human anti-SARS antibody	M396	Prabakaran et al., J. Biol. Chem. 281 (23), 15829-15836 (2006), NCBI Accession # 2G75_D (213aa)	6740
CORV5	Heavy chain variable region, Human neutralizing SARS antibody		Prabakaran et al., J. Biol. Chem. 281 (23), 15829-15836 (2006), NCBI Accession # 2DD8_L (213aa)	6741
CORV6	Heavy chain variable region, Humanized neutralizing murine monoclonal MERS		CN103864924 SEQ ID NO: 1	6742
CORV7	Heavy chain variable region, MERS		CN104447986 SEQ ID NO: 1	6743
CORV8	Heavy chain variable region, Neutralizing antibody (binds to the spike protein (S) of SARS-cov)		US7750123 SEQ ID NO: 12; WO2005060520; CN1914226; US20050249739	6744
CORV9	Heavy chain variable region, SARS antibody	s110.4	US20110159001 SEQ ID NO: 62; WO2009128963; EP2242768; CN102015767	6745
CORV10	Heavy chain variable region, SARS antibody	s124.5	US20110159001 SEQ ID NO: 66; WO2009128963; EP2242768; CN102015767	6746
CORV11	Heavy chain variable region, SARS antibody	s215.17	US20110159001 SEQ ID NO: 70; WO2009128963; EP2242768; CN102015767	6747
CORV12	Heavy chain variable region, SARS antibody	s218.9	US20110159001 SEQ ID NO: 74; WO2009128963; EP2242768; CN102015767	6748

CORV13	Heavy chain variable region, SARS antibody	s223.4	US20110159001 SEQ ID NO: 78; WO2009128963; EP2242768; CN102015767	6749
CORV14	Heavy chain variable region, SARS antibody	s225.12	US20110159001 SEQ ID NO: 82; WO2009128963; EP2242768; CN102015767	6750
CORV15	Heavy chain variable region, SARS antibody	s231.19	US20110159001 SEQ ID NO: 86; WO2009128963; EP2242768; CN102015767	6751
CORV16	Heavy chain variable region, SARS antibody	s230.14+15	US20110159001 SEQ ID NO: 90; WO2009128963; EP2242768; CN102015767	6752
CORV17	Heavy chain variable region, SARS antibody	s227.14	US20110159001 SEQ ID NO: 94; WO2009128963; EP2242768; CN102015767	6753
CORV18	Heavy chain variable region, SARS antibody	s109.8	US20110159001 SEQ ID NO: 98; WO2009128963; EP2242768; CN102015767	6754
CORV19	Heavy chain variable region, SARS antibody	Fab58	CN1513874	6755
CORV20	Heavy chain variable region, SARS antibody	Fab59	CN1513874	6756
CORV21	Heavy chain variable region, SARS human monoclonal antibody	3C7	US7728110 SEQ ID NO: 60; WO2008060331; EP2035454A2, US20080248043	6757
CORV22	Heavy chain variable region, SARS human monoclonal antibody	F26G18	US7622112 SEQ ID NO: 5; WO2005054469; US20080248043; US20080081047	6758
CORV23	Heavy chain variable region A, humanized antibody binding to S2 domain of SARS		WO2006095180 SEQ ID NO: 24	6759
CORV24	Heavy chain, Humanized neutralizing murine monoclonal MERS		CN103864924 SEQ ID NO: 3	6760
CORV25	Heavy chain, MERS	m336	WO2015057942 SEQ ID NO: 1	6761
CORV26	Heavy chain, MERS	m337	WO2015057942 SEQ ID NO: 9	6762
CORV27	Heavy chain, MERS	m338	WO2015057942 SEQ ID NO: 16	6763
CORV28	Heavy chain, MERS	2e 6	CN104447986 SEQ ID NO: 3	6764
CORV29	Heavy chain, MERS	M336	Ying et al., Nat Commun 6, 8223 (2015), NCBI Accession # 4XAK_H(252aa)	6765
CORV30	Human anti-SARS antibody		Leung et al., PLoS Med. 3 (7), E237 (2006), NCBI Accession # ABA54613.1(117aa)	6766
CORV31	Human monoclonal MERS	Mers-27	CN104628848 SEQ ID NO: 1	6767
CORV32	Human monoclonal MERS	Mers-27	CN104628848 SEQ ID NO: 3	6768
CORV33	Human monoclonal MERS	Mers-4	CN104628849 SEQ ID NO: 1	6769
CORV34	Human monoclonal MERS	Mers-4	CN104628849 SEQ ID NO: 3	6770
CORV35	Kappa light chain partial sequence, human SARS neutralization antibody, Ig	H12	AAX19355.1(108aa)	6771
CORV36	Light chain partial sequence, Human anti-SARS antibody, Ig		Liu, J., unpublished, NCBI Accession # BAE94187.1(219aa)	6772
CORV37	Light chain variable domain, MERS		CN104447986 SEQ ID NO: 2	6773

CORV38	Light chain variable partial sequence, Human neutralizing SARS antibody	80R	Hwang et al., J. Biol. Chem. 281 (45), 34610-34616 (2006), NCBI Accession # 2GHW_D (247aa)	6774
CORV39	Light chain variable region, A humanized antibody binding to S2 domain of SARS		WO2006095180 SEQ ID NO: 25	6775
CORV40	Light chain variable region, human anti-SARS antibody	M396	Prabakaran et al., J. Biol. Chem. 281 (23), 15829-15836 (2006), NCBI Accession # 2G75_C (245aa)	6776
CORV41	Light chain variable region, Human neutralizing SARS antibody		Prabakaran et al., J. Biol. Chem. 281 (23), 15829-15836 (2006), NCBI Accession # 2DD8_H(245aa)	6777
CORV42	Light chain variable region, Humanized neutralizing murine monoclonal MERS		CN103864924 SEQ ID NO: 2	6778
CORV43	Light chain variable region, neutralizing antibody (binds to the spike protein (S) of SARS-cov)		US7750123 SEQ ID NO: 20; WO2005060520; CN1914226; US20050249739	6779
CORV44	Light chain variable region, SARS antibody	s110.4	US20110159001 SEQ ID NO: 64; WO2009128963; EP2242768; CN102015767	6780
CORV45	Light chain variable region, SARS antibody	s124.5	US20110159001 SEQ ID NO: 68; WO2009128963; EP2242768; CN102015767	6781
CORV46	Light chain variable region, SARS antibody	s215.17	US20110159001 SEQ ID NO: 72; WO2009128963; EP2242768; CN102015767	6782
CORV47	Light chain variable region, SARS antibody	s218.9	US20110159001 SEQ ID NO: 76; WO2009128963; EP2242768; CN102015767	6783
CORV48	Light chain variable region, SARS antibody	s223.4	US20110159001 SEQ ID NO: 80; WO2009128963; EP2242768; CN102015767	6784
CORV49	Light chain variable region, SARS antibody	s225.12	US20110159001 SEQ ID NO: 84; WO2009128963; EP2242768; CN102015767	6785
CORV50	Light chain variable region, SARS antibody	s231.19	US20110159001 SEQ ID NO: 88; WO2009128963; EP2242768; CN102015767	6786
CORV51	Light chain variable region, SARS antibody	s230.14+15	US20110159001 SEQ ID NO: 92; WO2009128963; EP2242768; CN102015767	6787
CORV52	Light chain variable region, SARS antibody	s227.14	US20110159001 SEQ ID NO: 96; WO2009128963; EP2242768; CN102015767	6788
CORV53	Light chain variable region, SARS antibody	s109.8	US20110159001 SEQ ID NO: 101; WO2009128963; EP2242768; CN102015767	6789
CORV54	Light chain variable region, SARS antibody	Fab58	CN1513874	6790
CORV55	Light chain variable region, SARS antibody	Fab59	CN1513874	6791
CORV56	Light chain variable region, SARS human monoclonal antibody	3C7	US7728110 SEQ ID NO: 58; WO2008060331; EP2035454A2, US20080248043	6792

CORV57	Light chain variable region, SARS human monoclonal antibody	F26G18	US7622112 SEQ ID NO: 14; WO2005054469; US20080248043; US20080081047	6793
CORV58	Light chain, Humanized neutralizing murine monoclonal MERS		CN103864924 SEQ ID NO: 4	6794
CORV59	Light chain, MERS	m336	WO2015057942 SEQ ID NO: 2	6795
CORV60	Light chain, MERS	m337	WO2015057942 SEQ ID NO: 10	6796
CORV61	Light chain, MERS	m338	WO2015057942 SEQ ID NO: 17	6797
CORV62	Light chain, MERS	2E 6	CN104447986 SEQ ID NO: 4	6798
CORV63	Light chain, MERS	M336	Ying et al., Nat Commun 6, 8223 (2015), NCBI Accession # 4XAK_L (214aa)	6799
CORV64	Variable heavy chain-constant heavy chain 1, Humanized neutralizing murine monoclonal MERS	4C2Fab	CN103864924 SEQ ID NO: 7	6800
CORV65	Variable light chain-constant light chain 1, Humanized neutralizing murine monoclonal MERS	4C2Fab	CN103864924 SEQ ID NO: 9	6801

[00327] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Patent No. US7629443, US Publication No. US20080254440, Chinese Publication No. CN 10361 3666, CN1570638, CN101522208, CN1 673231, CN 1590409, CN1557838. and CN 1488645, the contents of each of which are herein incorporated by reference in their entirety, against SARS.

[00328] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 26 against John Cunningham Virus.

Table 26. **Antibodies against** John **Cunningham** Virus

Antibody No.	**Description**	Antibody Name	**Reference Information**	SEQ **ID** NO
JCV1	Heavy chain	14C8	US20150056188 SEQ ID NO: 1	6802
JCV2	Heavy chain	16H5	US20150056188 SEQ ID NO: 5	6803
JCV3	Heavy chain	18C9	US20150056188 SEQ ID NO: 9	6804
JCV4	Heavy chain	34C6	US20150056188 SEQ ID NO: 13	6805
JCV5	Heavy chain	18C9 N55S	US20150056188 SEQ ID NO: 16	6806
JCV6	Heavy chain	18C9 N55Q	US20150056188 SEQ ID NO: 18	6807
JCV7	Heavy chain	18C9 N55D	US20150056188 SEQ ID NO: 20	6808
JCV8	Heavy chain	18C9 N55H	US20150056188 SEQ ID NO: 22	6809
JCV9	Heavy chain	18C9 N55T	US20150056188 SEQ ID NO: 24	6810
JCV10	Heavy chain	18C9 N55A	US20150056188 SEQ ID NO: 26	6811
JCV11	Heavy chain	18C9 N55L	US20150056188 SEQ ID NO: 28	6812
JCV12	Heavy chain	18C9 N55X	US20150056188 SEQ ID NO: 30	6813
JCV13	Heavy chain	18C9 G56A	US20150056188 SEQ ID NO: 32	6814

JCV14	Heavy chain	18C9 G56V	US20150056 188 SEQ ID NO: 34	6815
JCV15	Heavy chain	18C9 G56P	US20150056188 SEQ ID NO: 36	6816
JCV16	Heavy chain	18C9 G56X	US20150056188 SEQ ID NO: 38	6817
JCV17	Heavy chain	399-h (C35A V50A)	US20150050271 SEQ ID NO: 20	6818
JCV18	Heavy chain	Antibody from US20 13005027 i	US20150050271 SEQ ID NO: 66	6819
JCV19	Heavy chain	HO	US20150050271 SEQ ID NO: 51	6820
JCV20	Heavy chain	H1	US20150050271 SEQ ID NO: 52	6821
JCV21	Heavy chain	H3	US20150050271 SEQ ID NO: 54	6822
JCV22	Heavy chain	H4	US20150050271 SEQ ID NO: 55	6823
JCV23	Heavy chain	H5	US20150050271 SEQ ID NO: 56	6824
JCV24	Heavy chain	H6	US20150050271 SEQ ID NO: 57	6825
JCV25	Heavy chain	H7	US20150050271 SEQ ID NO: 58	6826
JCV26	Heavy cbaisi	H8	US20150050271 SEQ ID NO: 59	6827
JCV27	Heavy chain	H9	US20150050271 SEQ ID NO: 60	6828
JCV28	Heavy chain	LO	US20150050271 SEQ ID NO: 48	6829
JCV29	Heavy chain	jCv H vis	US20150050271 SEQ ID NO: 43	6830
JCV30	Heavy chain	IGH V3~30-3x01	US20150050271 SEQ ID NO: 44	6831
JCV31	Heavy chain	HO	US20150050271 SEQ ID NO: 19	6832
JCV32	Heavy chain	HO V50G	US20150050271 SEQ ID NO: 21	6833
JCV33	Heavy chain	H1	US20150050271 SEQ ID NO: 22	6834
JCV34	Heavy chain	H2	US20150050271 SEQ ID NO: 23	6835
JCV35	Heavy chain	H3	US20150050271 SEQ ID NO: 24	6836
JCV36	Heavy chain	H4	US20150050271 SEQ ID NO: 25	6837
JCV37	Heavy chain	H5	US20150050271 SEQ ID NO: 26	6838
JCV38	Heavy chain	H6	US20150050271 SEQ ID NO: 27	6839
JCV39	Heavy chain	H7	US20150050271 SEQ ID NO: 28	6840
JCV40	Heavy chain	H8	US20150050271 SEQ ID NO: 29	6841
JCV41	Heavy chain	H9	US20150050271 SEQ ID NO: 30	6842
JCV42	Heavy chaiii vanable region	GREI	US20150191530 SEQ ID NO: 1	6843
JCV43	Heavy chain **variable** region	R399	US20150050271 SEQ ID NO: 6	6844
JCV44	Light chain	14G8	US20150056188 SEQ ID NO: 3	6845
JCV45	Ligla **chain**	16H5	US20150056188 SEQ ID NO: 7	6846
JCV46	Light chain	18C9	US20150056 188 SEQ ID NO: 11	6847
JCV47	Light chain	34C6	US20150056188 SEQ ID NO: 14	6848
JCV48	Ligbi chain	18C9 C96L	US20150056 188 SEQ ID NO: 40	6849
JCV49	Light chain	18C9 G96S	US20150056188 SEQ ID NO: 42	6850
JCV50	Ligla **chain**	18C9 C96A	US20150056188 SEQ ID NO: 44	6851
JCV51	Light chain	18C9 C96X	US20150056 188 SEQ ID NO: 46	6852
JCV52	Light chain	399-i (N31G), L	US20150050271 SEQ ID NO: 15	6853
JCV53	Ligbi chain	Antibody front US2015005027 1	US20150050271 SEQ ID NO: 67	6854
JCV54	Light chain	H2	US20150050271 SEQ ID NO: 53	6855
JCV55	Ligbi chain	L i	US20150050271 SEQ ID NO: 49	6856

JCV56	Light chain	L2	US20150050271 SEQ ID NO: 50	6857
JCV57	Light chain	1GKV1D-13x01	US20150050271 SEQ ID NO: 39	6858
JCV58	LiglU chain	LO	US20150050271 SEQ ID NO: 11	6859
JCV59	Light chain	L1	US20150050271 SEQ ID NO: 12	6860
JCV60	Light chain	L2	US20150050271 SEQ ID NO: 13	6861
JCV61	Light chain	L2 N31A	US20150050271 SEQ ID NO: 14	6862
JCV62	Light chain variable region	GRE1	US20150191530 SEQ ID NO: 2	6863
JCV63	LiglU chain variable region	R399	US20150050271 SEQ ID NO: 1	6864
JCV64	Light chain variable region	R411,jcv41 1_vh	US20150050271 SEQ ID NO: 38	6865

[ΘΘ329] In one embodiment the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 27 against Poxvirus.

Table **27. Antibodies against Poxyirus**

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
POXV1	Heavy chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 2	6866
POXV2	Heavy chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 6	6867
POXV3	Heavy chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 10	6868
POXV4	Heavy chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 14	6869
POXV5	Heavy chain, H3L envelope protein	H3L binding antibody	US20140186370 SEQ ID NO: 14	6870
POXV6	Light chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 4	6871
POXV7	Light chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 8	6872
POXV8	Light chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 12	6873
POXV9	Light chain variable region, B5R envelope protein	B5R binding antibody	US8623370 SEQ ID NO: 16	6874
POXV10	Light chain, H3L envelope protein	H3L binding antibody	US20140186370 SEQ ID NO: 16	6875

[ΘΘ330] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 28 against Enterovirus 71.

Table 28. Antibodies **against Enterovirus** 71

Antibody No.	Descripiiots	Antibody **Name**	Reference Information	SE**Q ID** NO
ENTV1	Heavy chain variable region		CN102718864A SEQ ID NO: 2	6876
ENTV2	Heavy chain variable region	E18	WO2015092668 SEQ ID NO: 1	6877
ENTV3	Heavy chain variable region	E19	WO2015092668 SEQ ID NO: 3	6878
ENTV4	Heavy chain variable region	E20	WO2015092668 SEQ ID NO: 5	6879
ENTV5	Heavy chain variable region	E19 humanized VH1	WO20 15092668 SEQ ID NO: 19	6880
ENTV6	Heavy chain variable region	E19 humanized VH2	WO20 15092668 SEQ ID NO: 20	6881

ENTV7	Heavy chain variable region	E19 humanized VLB	WO20 15092668 SEQ ID NO: 21	6882
ENTV8	Heavy chain variable region	E19 humanized VH4	WO20 15092668 SEQ ID NO: 22	6883
ENTV9	Light chain variable region		CN1027 18864A SEQ ID NO: 1	6884
ENTV10	Light chain variable region	E18	WO20 15092668 SEQ ID NO: 2	6885
ENTVl i	Light chain variable region	E19	WO2015092668 SEQ ID NO: 4	6886
ENTV12	Light chain variable region	E20	WO2015092668 SEQ ID NO: 6	6887
ENTV13	Light chain variable region	E18 VL2	WO2015092668 SEQ ID NO: 15	6888
ENTV14	Light chain variable region	E19 humanized VL1	WO2015092668 SEQ ID NO: 16	6889
ENTV!5	Light chain variable region	E19 humanized VL2	WO20 15092668 SEQ ID NO: 17	6890
ENTV16	Light chain variable region	E19 humanized VL3	WO2015092668 SEQ ID NO: 18	6891

[00331] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in Chinese Publication No. CN104357400, the contents of which are herein incorporated by reference in their entirety, against EV71.

[00332] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants encoding MAB979, fragments or variants thereof for treating a disease and/or disorder or preventing a disease and/or disorder. As a non-limiting example, the disease and/or disorder is EV71.

[00333] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 29 against Rubella Virus.

Table 29. Antibodies against Rubella Virus

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
RUBV1	Heavy chain variable region	DDF-RuV1	US20100143376 SEQ ID NO: 2	6892
RUBV2	Heavy chain variable region	DDF-RuV2	US20100143376 SEQ ID NO: 9	6893
RUBV3	Light chain variable region	DDF-RuV1	US20100143376 SEQ ID NO: 7	6894
RUBV4	Light chain variable region	DDF-RuV2	US20100143376 SEQ ID NO: 14	6895

[00334] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 30 against Human Papilloma Virus.

Table 30. Antibodies against Human Papilloma Virus

Antibody No.	Description	Reference Information	SEQ ID NO
HPV1	Heavy chain variable region	WO2015096269 SEQ ID NO: 1	6896
HPV2	Light chain variable region	WO2015096269 SEQ ID NO: 2	6897

[00335] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides.

fragments or valiants thereof described in US Publication No. US201 30337438, the contents of which are herein incorporated by reference in their entirety, against HBV.

[00336] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the broadly neutralizing payload antibody polypeptides listed in Table 31 against viruses.

Table **31.** Broadly Neiitralizmg **Antibodies** for Viruses

Antibody No.	**Description**	Antibody Name	Reference **Information**	SEQ ID NO
V1R1	Heavy chain variable region, hepatitis, influenza, HIV, herpes, paramyxovirus, poxvirus, rhabdovirus or arenavirus	3G4	US76 11704 SEQ ID NO: 2	6898
VIR2	Heavy chain variable region, hepatitis, influenza, HIV, herpes, paramyxovirus, poxvirus, rhabdovirus or arenavirus	3G4	US76 11704 SEQ ID NO: 4	6899
V1R3	Heavy chain variable region, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B, Sendai, feline leukemia, Reo, polio, human serum parvo-like, simian 40, respiratory syncytial, mouse mammary rumor, Varicella-Zoster, Sight chain variable region. Dengue, rubella, measles, adenovirus, human T-cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stomatitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue	679	US7429381 SEQ ID NO: 4	6900
VIR4	Heavy chain variable region, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B, Sendai, feline leukemia, Reo, polio, human serum parvo-like, simian 40, respiratory syncytial, mouse mammary tumor, Varicella-Zoster, Dengue, rubella, measles, adenovirus, human T-cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stomatitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue, light chain variable region	Mu-9V	US742938! SEQ ID NO: 10	690 I
VTR5	Heavy chain variable region, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B, Sendai, feline leukemia, Reo, polio, human serum parvo-like, simian 40, respirator}' syncytial, mouse mammary tumor, Varicella-Zoster, Dengue, rubella, measles, adenovirus, human T-cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stomatitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue, light chain variable region	humanized Mu-9	US7429381 SEQ ID NO: 14	6902
VIR6	Heavy chain variable region, Human cytomegalovirus, HCMV, human T-cell leukemia virus type 1, HIV-I, simian immunodeficiency virus, Ebola vims, Herpesvirus saimiri virus, influenza virus, and vaccinia virus	Fab-2 Clone3	US20 120269801 SEQ ID NO: 6	6903
VIR7	Heavy chain variable region, Human cytomegalovirus, HCMV, human T-cell leukemia virus type 1, HIV-I, simian immunodeficiency virus, Ebola virus, Herpesvirus saimiri virus, influenza virus, and vaccinia virus	Fab-3 Clone 7	US20 12026980 I SEQ ID NO: 10	6904
VIR8	Light chain variable region, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B,	Mu-9V	US7429381 SEQ ID NO: 8	6905

	Sendai, feline leukemia, Reo, polio, human serum parvo-like, simian 40, respiratory syncytial. mouse mammary iumor, Varicella-Zoster, Dengue, rubella, measles, adenovirus, human T-cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stomatitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue, light chain variable region			
VIR9	Light chain variable region, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B, Sendai, feline leukemia, Reo, polio, human serum parvo-iike, simian 40, respiratory syncytial, mouse mammary tumor, Varicella-Zoster, Dengue, rubella, measles, adenovirus, human T~ cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stoniaiitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue, light chain variable region	humanized Mu-9	US7429381 SEQ ID NO: 12	6906
VIR1O	Light chain variable region, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B, Sendai, feline leukemia, Reo, polio, human serum parvo-like, simian 40, respiratory syncytial, mouse mammary tumor, Varicella-Zoster, Dengue, rubella, measles, adenovirus, human T-cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stomatitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue	679	US7429381 SEQ ID NO: 2	6907
VIR 11	Light chain variable region, Human cytomegalovirus, HCMV, human T-cell leukemia virus type 1, HIV-1, simian immunodeficiency virus, Ebola vims, Herpesvirus saimiri virus, influenza virus, and vaccinia virus	Fab-3 Clone 7	US20 12026980 1 SEQ ID NO: 8	6908
VIR12	Light chain variable, Human cytomegalovinis, HCMV, human T-cell leukemia virus type 1, HIV-1, simian immunodeficiency virus, Ebola virus, Herpesvirus saimiri virus, influenza viras. and vaccinia virus, region	Fab-2 Clone3	US20 120269801 SEQ ID NO: 4	6909
VIR13	ScFv, hepatitis, influenza, HIV, herpes, paramyxovirus, poxvirus, rhabdovirus or arenavirus	3A2	US76 11704 SEQ ID NO: 6	6910
VIR 14	ScFv, HIV, herpes, cytomegalovirus, rabies, influenza, hepatitis B, Sendai, feline leukemia, Reo, polio, human serum parvo-like, simian 40, respiratory syncytial, mouse mammary tumor, Varicella-Zoster, Dengue, rubella, measles, adenovirus, human T-cell leukemias, Epstein-Barr, murine leukemia, mumps, vesicular stomatitis, Sindbis, lymphocytic choriomeningitis, wart and blue tongue	679	US7429381 SEQ ID NO: 6	691 l

[00337] In one embodiment, the pavioad region of the AAV particle composes one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 32 against Pseudomonas Aeruginosa.

Table 32. Antibodies against **Pseudomonas Aeruginosa**

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
PSEU1	Bivalent nanobody	260 (1E11-40GS-2B10)	US20150044215 SEQ ID NO: 118	6912

PSEU2	Bivalent nanobody	272 (11B09-40GS-10C05)	US20150044215 SEQ ID NO: 119	6913
PSEU3	Bivalent nanobody	308 (6B05-40GS-1E11)	US20150044215 SEQ ID NO: 120	6914
PSEU4	Bivalent nanobody	264 (1E11-40GS-2B02)	US20150044215 SEQ ID NO: 121	6915
PSEU5	Bivalent nanobody	302 (5H01-40GS-7C10)	US20150044215 SEQ ID NO: 122	6916
PSEU6	Bivalent nanobody	234 (7C10-40GS-5H01)	US20150044215 SEQ ID NO: 123	6917
PSEU7	Bivalent nanobody	064 (13F07-40GS-7C10)	US20150044215 SEQ ID NO: 124	6918
PSEU8	Bivalent nanobody	275 (2G09-40GC-5H10)	US20150044215 SEQ ID NO: 125	6919
PSEU9	Bivalent nanobody	083 (7C10-40GS-11B09)	US20150044215 SEQ ID NO: 126	6920
PSEU10	Bivalent nanobody	087 (1E11-40GS-7C10)	US20150044215 SEQ ID NO: 127	6921
PSEU11	Bivalent nanobody	269 (6B05-40GS-13F07)	US20150044215 SEQ ID NO: 128	6922
PSEU12	Bivalent nanobody	256 (13F07-40GS-5H01)	US20150044215 SEQ ID NO: 129	6923
PSEU13	Bivalent nanobody	277 (5H01-40GS-11B09)	US20150044215 SEQ ID NO: 130	6924
PSEU14	Bivalent nanobody	257 (13F07-40GS-2B10)	US20150044215 SEQ ID NO: 131	6925
PSEU15	Bivalent nanobody	285 (13F07-40GS-2B02)	US20150044215 SEQ ID NO: 132	6926
PSEU16	Bivalent nanobody	115 (11B09-40GS-13F07)	US20150044215 SEQ ID NO: 133	6927
PSEU17	Bivalent nanobody	258 (13F07-40GS-14E10)	US20150044215 SEQ ID NO: 134	6928
PSEU18	Bivalent nanobody	283 (7E09-40GS-6B05)	US20150044215 SEQ ID NO: 135	6929
PSEU19	Bivalent nanobody	271 (7C10-40GS-14E10)	US20150044215 SEQ ID NO: 136	6930
PSEU20	Bivalent nanobody	259 (1E11-40GS-5H01)	US20150044215 SEQ ID NO: 137	6931
PSEU21	Bivalent nanobody	319 (13F07-40GS-6B05)	US20150044215 SEQ ID NO: 138	6932
PSEU22	Bivalent nanobody	335 (5H01-40GS-1E11)	US20150044215 SEQ ID NO: 139	6933
PSEU23	Bivalent nanobody	261 (5H01-40GS-2B10)	US20150044215 SEQ ID NO: 140	6934
PSEU24	Bivalent nanobody	262 (7E09-40GS-7C10)	US20150044215 SEQ ID NO: 141	6935
PSEU25	Constant heavy chain		US20150044215 SEQ ID NO: 148	6936
PSEU26	Constant light chain		US20150044215 SEQ ID NO: 149	6937
PSEU27	Heavy chain	Panobacumab	US8197816 SEQ ID NO: 8	6938
PSEU28	Heavy chain		US20130156696 SEQ ID NO: 2	6939
PSEU29	Heavy chain		US7494653 SEQ ID NO: 2	6940
PSEU30	Heavy chain variable region	KB0001	US8044181 SEQ ID NO: 3	6941
PSEU31	Heavy chain variable region	KB0001	US8044181 SEQ ID NO: 5	6942
PSEU32	Heavy chain variable region	KB0001	US8044181 SEQ ID NO: 7	6943
PSEU33	Heavy chain variable region	KB0001	US8044181 SEQ ID NO: 9	6944
PSEU34	Heavy chain variable region	KB0001	US8044181 SEQ ID NO: 11	6945
PSEU35	Heavy chain variable region	1F3	US9085611 SEQ ID NO: 11	6946
PSEU36	Heavy chain variable region	2A4	US9085611 SEQ ID NO: 13	6947
PSEU37	Heavy chain variable region		US9085611 SEQ ID NO: 27	6948
PSEU38	Heavy chain variable region	mAbs LST-001	US8653242 SEQ ID NO: 29	6949

PSEU39	**Heavy chain variable** region	mAbs LST-002	US8653242 SEQ ID NO: 49	6950
PSEU40	Heavy chain variable region	mAbs LST-005	**US8653242** SEQ ID NO: 52	695 1
PSEU41	Heavy **chain variable** reeion	niAbs LST-006	**US8653242** SEQ ID NO: 54	6952
PSEU42	Heavy chain variable region	mAbs LST-007	US8653242 SEQ ID **NO:** 13	6953
PSEU43	Heavy **chain variable** reeion	niAbs LST-008	**US8653242** SEQ ID NO: 15	6954
PSEU44	Heavy chain variable region	310B06	US7597893 SEQ ID **NO:** 8	6955
PSEU45	Heavy **chain variable** reeion	**Cam-003**	**US20140227285** SEQ **ID NO:** 1	6956
PSEU46	Heavy chain variable region	**Catn-004**	US20140227285 SEQ ID **NO: 3**	6957
PSEU47	Heavy **chain variable** region	**Cam-005**	**US20140227285** SEQ **ID NO:** 4	6958
PSEU48	Heavy chain variable region	**WapR-00**	US20 140227285 SEQ ID NO: **5**	6959
PSEU49	Heavy **chain variable** region	**WapR-002**	**US20140227285** SEQ **ID NO:** 7	6960
PSEU50	**Heavy** chain **variable** region	**WapR-003**	**US20 140227285** SEQ **ID** NO: 9	6961
PSEU5 1	Heavy chain variable region	**WapR-004**	**US20140227285** SEQ **ID** NO: 11	6962
PSEU52	**Heavy** chain **variable** region	**WapR-007**	**US20 140227285** SEQ **ID** NO: 13	6963
PSEU53	Heavy chain variable region	**WapR-016**	**US20140227285** SEQ **ID** NO: 15	6964
PSEU54	**Heavy** chain **variable** region	1584	**US20 130045207** SEQ **ID** NO: 8	6965
PSEU55	Heavy chain variable region	1573	**US20130045207** SEQ **ID NO:** 16	6966
PSEU56	**Heavy** chain **variable** region	1572	**US20 130045207** SEQ **ID** NO: 24	6967
PSE1J57	Heavy chain variable region	1587	**US20130045207** SEQ **ID NO:** 32	6968
PSEU58	**Heavy chain** variable region	3099	US20 130022604 SEQ ID NO: 8	6969
PSEU59	Heavy chain variable region	2745	**US20 130022604** SEQ ID NO: 16	6970
PSEU60	**Heavy chain** variable region	2459	US20 130022604 SEQ ID NO: 24	6971
PSEU61	Heavy chain variable region	2316	**US20 130022606** SEQ ID NO: 8	6972
PSEU62	**Heavy chain** variable region	1838	US20 130022606 SEQ ID NO: 16	6973
PSEU63	Heavy chain variable region	2314	**US20 130022606** SEQ ID NO: 24	6974
PSEU64	**Heavy chain** variable region	2326	US20 130022606 SEQ ID NO: 32	6975
PSEU65	Heavy chain variable region	2328	**US20 130022606** SEQ ID NO: 40	6976
PSEU66	**Heavy chain** variable region	2438	US20130022606 SEQ ID NO: 48	6977
PSEU67	Heavy chain variable region	1774	**US20130004500** SEQ ID NO: 8	6978

PSEU68	Heavy chain variable region	1660	US20130004500 SEQ ID NO: 16	6979
PSEU69	Heavy chain variable region	1923	US20130004500 SEQ ID NO: 24	6980
PSEU70	Heavy chain variable reeion	1656	US20130004499 SEQ ID NO: 8	6981
PSEU71	Heavy chain variable region	1640	US20130004499 SEQ ID NO: 16	6982
PSEU72	Heavy chain variable reeion	2459	US20130004499 SEQ ID NO: 24	6983
PSEU73	Heavy chain variable region		US20120114657 SEQ ID NO: 8	6984
PSEU74	Heavy chain variable reeion	Anti-It-2	US20110177087 SEQ ID NO: 13	6985
PSEU75	Heavy chain variable region	Anti-It-3	US20110177087 SEQ ID NO: 14	6986
PSEU76	Heavy chain variable region	Anti-It-4	US20110177087 SEQ ID NO: 15	6987
PSEU77	Heavy chain variable region	Anti-It-5	US20110177087 SEQ ID NO: 16	6988
PSEU78	Heavy chain variable region	Anti-It-6	US20110177087 SEQ ID NO: 17	6989
PSEU79	Heavy chain variable region	Anti-170003	US201 10177087 SEQ ID NO: 18	6990
PSE1J80	Heavy chain variable region	Anti- 170006	1JS20110177087 SEQ ID NO: 19	6991
PSEU8 1	Heavy chain variable region	Anti-PaOl	US201 10177087 SEQ ID NO: 20	6992
PSE1J82	Heavy chain variable region	Anti-IATS0 16	US2.01101177087 SEQ ID NO: 21	6993
PSEU83	Heavy chain variable region		US20090191 186 SEQ ID NO: I	6994
PSE1J84	Heavy chain variable region		US20090 191186 SEQ ID NO: 11	6995
PSEU85	Heavy chain variable region		US20090191 186 SEQ ID NO: 3	6996
PSE1J86	Heavy chain variable region		US20090 191186 SEQ ID NO: 7	6997
PSEU87	Heavy chain variable region		US20090191 186 SEQ ID NO: 9	6998
PSEU88	Heavy chain variable region		US20090191 186 SEQ ID NO: 5	6999
PSEU89	Heavy chain variable region		US20090191 186 SEQ ID NO: 13	7000
PSEU90	Heavy chain variable region		US20090191 186 SEQ ID NO: 21	7001
PSEU91	Heavy chain variable region		US20090191 186 SEQ ID NO: 17	7002
PSEU92	Heavy chain variable region		US20090191 186 SEQ ID NO: 26	7003
PSEU93	Heavy chain variable region		US20090191 186 SEQ ID NO: 25	7004
PSEU94	Heavy chain variable region		US20090191 186 SEQ ID NO: 23	7005
PSEU95	Heavy chain variable region		US20090191 186 SEQ ID NO: 29	7006
PSEU96	Heavy chain variable region		US20090191 186 SEQ ID NO: 35	7007

PSEU97	Heavy chain variable region	V2L2	WO2014074528 SEQ ID NO: 216	7008
PSEU98	Heavy chain variable region	V2L2-MD	WO2014074528 SEQ ID NO: 255	7009
PSEU99	Heavy chain variable region	V2L2-MD and V2L2-GL	WO2014074528 SEQ ID NO: 256	7010
PSEU100	Heavy chain variable region	V2L2-GL	WO2014074528 SEQ ID NO: 257	7011
PSEU101	Heavy chain variable region	2409	WO2013024905 SEQ ID NO: 16	7012
PSEU102	Heavy chain variable region	2453	WO2013024905 SEQ ID NO: 24	7013
PSEU103	Heavy chain variable region	S20	US7972845 SEQ ID NO: 2	7014
PSEU104	Heavy chain variable region	Fab 13.37	US20150044215 SEQ ID NO: 142	7015
PSEU105	Heavy chain variable region	Fab 26.24	US20150044215 SEQ ID NO: 144	7016
PSEU106	Heavy chain variable region	Fab 35.36	US20150044215 SEQ ID NO: 146	7017
PSEU107	Heavy chain variable region	KB0001	US8044181 SEQ ID NO: 1	7018
PSEU108	Heavy chain, LPS serotype IATS-O11,		Horn, M.P. et al. "Preclinical In Vitro and In Vivo characterization of the fully human monoclonal IgM antibody KBPA101 specific for Pseudomonas aeruginosa serotype IATS-O11", Antimicrob. Agents Chemother. 54 (6), 2338-2344 (2010)	7019
PSEU109	J chain	Panobacumab		7020
PSEU110	Light chain	Panobacumab	US8197816 SEQ ID NO: 7	7021
PSEU111	Light chain		US20130156696 SEQ ID NO: 4	7022
PSEU112	Light chain		US7494653 SEQ ID NO: 4	7023
PSEU113	Light chain variable region	1F3	US9085611 SEQ ID NO: 12	7024
PSEU114	Light chain variable region	2A4	US9085611 SEQ ID NO: 14	7025
PSEU115	Light chain variable region		US9085611 SEQ ID NO: 28	7026
PSEU116	Light chain variable region	mAbs LST-001	US8653242 SEQ ID NO: 18	7027
PSEU117	Light chain variable region	mAbs LST-006	US8653242 SEQ ID NO: 53	7028
PSEU118	Light chain variable region	mAbs LST-008	US8653242 SEQ ID NO: 14	7029
PSEU119	Light chain variable region	mAbs LST-008	US8653242 SEQ ID NO: 16	7030
PSEU120	Light chain variable region	310BO6	US7597893 SEQ ID NO: 7	7031
PSEU121	Light chain variable region	Cam-003, Cam-004, Cam-005	US20140227285 SEQ ID NO: 2	7032
PSEU122	Light chain variable region	WapR-001	US20140227285 SEQ ID NO: 6	7033
PSEU123	Light chain variable region	WapR-002	US20140227285 SEQ ID NO: 8	7034
PSEU124	Light chain variable **region**	WapR-003	US20140227285 SEQ ID NO: 10	7035

PSEU125	Light chain variable region	WapR-004, WapR-004RAD	US20140227285 SEQ ID NO: 12	7036
PSEU126	Light chain variable region	WapR-007	US20 140227285 SEQ ID NO: 14	7037
PSEU 127	Light chain variable region	WapR-0 16	US20140227285 SEQ ID NO: 16	7038
PSEU128	Light chain variable region	1584	US20 130045207 SEQ ID NO: 7	7039
PSEU 129	Light chain variable region	L573	US20130045207 SEQ ID NO: 15	7040
PSEU130	Light chain variable region	1572	US20 130045207 SEQ ID NO: 23	7041
PSEU 13 1	Light chain variable region	1587	US20130045207 SEQ ID NO: 3 1	7042
PSEU132	Light chain variable region	3099	US20 130022604 SEQ ID NO: 7	7043
PSEU 133	Light chain variable region	2745	US20 130022604 SEQ ID NO: 15	7044
PSEU134	Light chain variable region	2459	US20 130022604 SEQ ID NO: 23	7045
PSEU 135	Light chain variable region	2316	US20 130022606 SEQ ID NO: 7	7046
PSEU 136	Light chain variable region	1838	US20130022606 SEQ ID NO: 15	7047
PSEU 137	Light chain variable region	2314	US20130022606 SEQ ID NO: 23	7048
PSEU138	Light chain variable region	?3?6	US20130022606 SEQ ID NO: 31	7049
PSEU 139	Light chain variable region	2328	US20 130022606 SEQ ID NO: 39	7050
PSEU 140	Light chain variable region	2438	US20130022606 SEQ ID NO: 47	705 1
PSEU 14 1	Light chain variable region	1774	US20130004500 SEQ ID NO: 7	7052
PSEU142	Light chain variable region	1660	US20130004500 SEQ ID NO: 15	7053
PSEU 143	Light chain variable region	1923	US20130004500 SEQ ID NO: 23	7054
PSEU 144	Light chain variable region	1656	US20130004499 SEQ ID NO: 7	7055
PSEU 145	Light chain variable region	1640	US20130004499 SEQ ID NO: 15	7056
PSEU 146	Light chain variable region	2459	US20130004499 SEQ ID NO: 23	7057
PSEU 147	Light chain variable region		US201201 14657 SEQ ID NO: 7	7058
PSEU148	Light chain variable region	Anti-It-2	US201 10177087 SEQ ID NO: 22	7059
PSEU 149	Light chain variable region	Anti-It-3	US201 10177087 SEQ ID NO: 23	7060
PSEU 150	Light chain variable region	Anti-It-4	US201 10177087 SEQ ID NO: 24	7061
PSEU 15 1	Light chain variable region	Anti-It-5	US201 10177087 SEQ ID NO: 25	7062
PSEU 152	Light chain variable region	Anti-It-6	US201 10177087 SEQ ID NO: 26	7063
PSEU 153	Light chain variable region	Anti-170003	US201 10177087 SEQ ID NO: 27	7064

PSEU154	Light chain variable region	Anti- 170006	US201 10 177087 SEQ ID NO: 28	7065
PSEU155	Light chain variable region	Anti-PaOl	US201 10177087 SEQ ID NO: 29	7066
PSEU156	Light chain variable region	Anti-IATS0 16	US201 10 177087 SEQ ID NO: 30	7067
PSEU157	Light chain variable region		US20090191 186 SEQ ID NO: 2	7068
PSEU158	Light chain variable region		US20090 191 186 SEQ ID NO: 12	7069
PSEU 159	Light chain variable region		US20090191 186 SEQ ID NO: 8	7070
PSEU160	Light chain variable region		US20090 191 186 SEQ ID NO: 10	707 1
PSEU 161	Light chain variable region		US2009019H 86 SEQ ID NO: 6	7072
PSEU162	Light chain variable region		US20090 191 186 SEQ ID NO: 37	7073
PSEU163	Light chain variable region		US2009019H 86 SEQ ID NO: 18	7074
PSEU 164	Light chain variable region		US20090 191 186 SEQ ID NO: 24	7075
PSEU165	Light chain variable region		US20090191 186 SEQ ID NO: 20	7076
PSEU 166	Light chain variable region		US20090 191 186 SEQ ID NO: 36	7077
PSEU 167	Light chain variable region		US20090191 186 SEQ ID NO: 28	7078
PSEU 168	Light chain variable region		US20090 191 186 SEQ ID NO: 30	7079
PSEU169	Light chain variable region		US20090191 186 SEQ ID NO: 34	7080
PSEU170	Light chain variable region		US20090 191 186 SEQ ID NO: 32	7081
PSEU171	Light chain variable region	V2L2	WO20 14074528 SEQ ID NO: 217	7082
PSEU172	Light chain variable region	2409	WO2013024905 SEQ ID NO: 15	7083
PSEU173	Light chain variable region	2453	WO20 13024905 SEQ ID NO: 23	7084
PSEU 174	Light chain variable region	S20	US7972845 SEQ ID NO: 4	7085
PSEU175	Light chain variable region	Fab 13.37	US20150044215 SEQ ID NO: 143	7086
PSEU 176	Light chain variable region	Fab 26.24	US20150044215 SEQ ID NO: 145	7087
PSEU 177	Light chain variable region	Fab 35.36	US20150044215 SEQ ID NO: 147	7088
PSEU 178	Light chain variable region majority	**mAbs** LST-002	US8653242 SEQ ID NO: 32	7089
PSEU179	Light chain variable region majority	rnAhs LST-006	US8653242 SEQ ID NO: 55	7090
PSEU 180	Light chain variable region minority	**mAbs** LST-002	US8653242 SEQ ID NO: 51	7091
PSEU181	Light chain variable region minority	rnAhs LST-007	US8653242 SEQ ID NO: 56	7092
PSEU 182	Light chain, Anti-P. Aeuginosa LPS serotype IATS-01 1,		Horn, M.P. et al. "Preclinical In Vitro arsd In Vivo characterization of the fully human monoclonal	7093

			IgM antibody KBPA101 specific for Pseudomonas aeruginosa serotype IATS-O11", Antimicrob. Agents Chemother. 54 (6), 2338-2344 (2010)	
PSEU183	Light kappa chain variable region	KB0001	US8044181 SEQ ID NO: 10	7094
PSEU184	Light kappa chain variable region	KB0001	US8044181 SEQ ID NO: 2	7095
PSEU185	Light kappa chain variable region	KB0001	US8044181 SEQ ID NO: 4	7096
PSEU186	Light kappa chain variable region	KB0001	US8044181 SEQ ID NO: 6	7097
PSEU187	Light kappa chain variable region	KB0001	US8044181 SEQ ID NO: 8	7098
PSEU188	Monovalent nanobody	5H01	US20150044215 SEQ ID NO: 1	7099
PSEU189	Monovalent nanobody	7C10	US20150044215 SEQ ID NO: 2	7100
PSEU190	Monovalent nanobody	1E11	US20150044215 SEQ ID NO: 3	7101
PSEU191	Monovalent nanobody	2B02	US20150044215 SEQ ID NO: 4	7102
PSEU192	Monovalent nanobody	2B10	US20150044215 SEQ ID NO: 5	7103
PSEU193	Monovalent nanobody	2G09	US20150044215 SEQ ID NO: 6	7104
PSEU194	Monovalent nanobody	6B05	US20150044215 SEQ ID NO: 7	7105
PSEU195	Monovalent nanobody	10C05	US20150044215 SEQ ID NO: 8	7106
PSEU196	Monovalent nanobody	11B09	US20150044215 SEQ ID NO: 9	7107
PSEU197	Monovalent nanobody	14E10	US20150044215 SEQ ID NO: 10	7108
PSEU198	Monovalent nanobody	7E09	US20150044215 SEQ ID NO: 11	7109
PSEU199	Monovalent nanobody	13F07	US20150044215 SEQ ID NO: 12	7110
PSEU200	Monovalent nanobody	3B11	US20150044215 SEQ ID NO: 13	7111
PSEU201	Monovalent nanobody	4C03	US20150044215 SEQ ID NO: 14	7112
PSEU202	Monovalent nanobody	4G10	US20150044215 SEQ ID NO: 15	7113
PSEU203	Monovalent nanobody	12B02	US20150044215 SEQ ID NO: 16	7114
PSEU204	Monovalent nanobody	14B10	US20150044215 SEQ ID NO: 17	7115
PSEU205	Monovalent nanobody	3E10	US20150044215 SEQ ID NO: 18	7116
PSEU206	Monovalent nanobody	5E02	US20150044215 SEQ ID NO: 19	7117
PSEU207	Scfv-Fc	W4-M1	WO2014074528 SEQ ID NO: 78	7118
PSEU208	Scfv-Fc	W4-M5	WO2014074528 SEQ ID NO: 79	7119
PSEU209	Scfv-Fc	W4-M6	WO2014074528 SEQ ID NO: 80	7120
PSEU210	Scfv-Fc	W4-M7	WO2014074528 SEQ ID NO: 81	7121
PSEU211	Scfv-Fc	W4-M8	WO2014074528 SEQ ID NO: 82	7122
PSEU212	Scfv-Fc	W4-M9	WO2014074528 SEQ ID NO: 83	7123
PSEU213	Scfv-Fc	W4-M11	WO2014074528 SEQ ID NO: 84	7124
PSEU214	Scfv-Fc	W4-M12	WO2014074528 SEQ ID NO: 85	7125
PSEU215	Scfv-Fc	W4-M14	WO2014074528 SEQ ID NO: 86	7126
PSEU216	Scfv-Fc	W4-M15	WO2014074528 SEQ ID NO: 87	7127
PSEU217	Scfv-Fc	W4-M16	WO2014074528 SEQ ID NO: 88	7128
PSEU218	Scfv-Fc	W4-M17	WO2014074528 SEQ ID NO: 89	7129
PSEU219	Scfv-Fc	W4-M19	WO2014074528 SEQ ID NO: 90	7130
PSEU220	Scfv-Fc	W4-M20	WO2014074528 SEQ ID NO: 91	7131

PSEU22 1	Scfv-Fc	W4-M4	WO2014074528 SEQ ID NO: 92	7 132
PSEU222	Scfv-Fc	W4-M I0	WO20 14074528 SEQ ID NO: 93	7133
PSEU223	Scfv-Fc	W4-HC 1-LCP	WO20 14074528 SEQ ID NO: 94	7 134
PSEU224	Scfv-Fc	W4-HC1-LC7	WO2014074528 SEQ ID NO: 95	7135
PSEU225	Scfv-Fc	W4-HC2-LC7	WO20 14074528 SEQ ID NO: 96	7136
PSEU226	Scfv-Fc	W4-HC3-LCP	WO2014074528 SEQ ID NO: 97	7 137
PSEU227	Scfv-Fc	W4-HC4-LCP	WO20 14074528 SEQ ID NO: 98	7138
PSEU228	Scfv-Fc	W4-HC5-LCP	WO20 14074528 SEQ ID NO: 99	7 139
PSEU229	Scfv-Fc	W4-HC5-LC7	WO20 14074528 SEQ ID NO: 100	7140
PSEU230	Scfv-Fc	W4-HC7-LCP	WO20 14074528 SEQ ID NO: 101	7141
PSEU23 1	Scfv-Fc	W4-VH1-VL8	WO20 1407452 8 SEQ ID NO: 102	7 142
PSEU232	Scfv-Fc	W4-VH2-VLP	WO20 14074528 SEQ ID NO: 103	7143
PSEU233	Scfv-Fc	W4-VH2-VL8	WO20 14074528 SEQ ID NO: 104	7 144
PSEU234	Scfv-Fc	W4-VH3-VL7	WO20 14074528 SEQ ID NO: 105	7145
PSEU235	Scfv-Fc	W4-VH3-VL8	WO20 14074528 SEQ ID NO: 106	7146
PSEU236	Scfv-Fc	W4-VH5-VL8	WO20 1407452 8 SEQ ID NO: 107	7 147
PSEU237	Scfv-Fc	W4-VH6-VL7	WO20 14074528 SEQ ID NO: 108	7148
PSEU238	Scfv-Fc	W4-VH6-VL8	WO20 14074528 SEQ ID NO: 109	7 149
PSEU239	Scfv-Fc	W4-VH6-VLP	WO20 14074528 SEQ ID NO: 110	7150
PSEU240	Scfv-Fc	W4-VH7-VLP	WO20 14074528 SEQ ID NO: 111	715 1
PSEU241	Scfv-Fc	W4-VH7-VL7	WO20 1407452 8 SEQ ID NO: 112	7152
PSEU242	Scfv-Fc	W4-VH7-VL8	WO20 14074528 SEQ ID NO: 113	7153
PSEU243	Scfv-Fc	W4-VH9-VLP	WO20 14074528 SEQ ID NO: 114	7 154
PSEU244	Scfv-Fc	W4-VH10-VLP	WO20 14074528 SEQ ID NO: 115	7155
PSEU245	Scfv-Fc	W4-VH1 1-VLP	WO20 14074528 SEQ ID NO: 116	7156
PSEU246	Scfv-Fc	W4-VH12-VLP	WO2014074528 SEQ ID NO: 117	7157
PSEU247	Scfv-Fc	W4-VH15-VLP	WO20 14074528 SEQ ID NO: 118	7158
PSEU248	Scfv-Fc	W4-VH 16-VLP	WO20 14074528 SEQ ID NO: 119	7 159
PSEU249	Scfv-Fc	W4-VH20-VLP	WO20 14074528 SEQ ID NO: 120	7160
PSEU250	Scfv-Fc	W4-VH3 1-VLP	WO20 14074528 SEQ ID NO: 121	7 161
PSEU25 1	Scfv-Fc	W4-VH37-VLP	WO2014074528 SEQ ID NO: 122	7162
PSEU252	Scfv-Fc	W4-VH4 1-VLP	WO20 14074528 SEQ ID NO: 123	7163
PSEU253	Scfv-Fc	W4-VH42-VLP	WO20 14074528 SEQ ID NO: 124	7 164
PSEU254	Scfv-Fc	W4-VH35-VLP	WO20 14074528 SEQ ID NO: 125	7165
PSEU255	Scfv-Fc	W4-VH36-VLP	WO20 14074528 SEQ ID NO: 126	7 166
PSEU256	Scfv-Fc	W4-VH52-VLP	WO2014074528 SEQ ID NO: 127	7167
PSEU257	Scfv-Fc	W4-VH53-VLP	WO20 14074528 SEQ ID NO: 128	7168
PSEU258	Scfv-Fc	W4-VH54-VLP	WO2014074528 SEQ ID NO: 129	7 169
PSEU259	Scfv-Fc	W4-VH55-VLP	WO20 14074528 SEQ ID NO: 130	7170
PSEU260	Scfv-Fc	W4-VH56-VLP	WO20 14074528 SEQ ID NO: 131	7 171
PSEU261	Scfv-Fc	W4-VH57-VLP	WO2014074528 SEQ ID NO: 132	7172
PSEU262	Scfv-Fc	W4-VH58-VLP	WO20 14074528 SEQ ID NO: 133	7173
PSEU263	Scfv-Fc	W4-VH60-VLP	WO2014074528 SEQ ID NO: 134	7 174
PSEU264	Scfv-Fc	W4-VH6 1-VLP	WO20 14074528 SEQ ID NO: 135	7175
PSE1J265	Scfv-Fc	W4-VH62-VLP	WO20 14074528 SEQ ID NO: 136	7 176

PSEU266	Scfv-Fc	W4-VH63-VLP	WO2014074528 SEQ ID NO: 137	7177
PSEU267	Scfv-Fc	W4-VH64-VLP	WO20 14074528 SEQ ID NO: 138	7178
PSEU268	Scfv-Fc	W4-VH65-VLP	WO20 14074528 SEQ ID NO: 139	7179
PSEU269	Scfv-Fc	W4-VH66-VLP	WO2014074528 SEQ ID NO: 140	7180
PSEU270	Scfv-Fc	W4-VH67-VLP	WO20 14074528 SEQ ID NO: 141	7181
PSEU27 1	Scfv-Fc	W4-VH69-VLP	WO2014074528 SEQ ID NO: 142	7182
PSEU272	Scfv-Fc	W4-VH70-VLP	WO20 14074528 SEQ ID NO: 143	7183
PSEU273	Scfv-Fc	W4-VH72-VLP	WO20 14074528 SEQ ID NO: 144	7184
PSEU274	Scfv-Fc	W4-VH79-VLP	WO20 14074528 SEQ ID NO: 145	7185
PSEU275	Scfv-Fc	W4-VH80-VLP	WO20 14074528 SEQ ID NO: 146	7186
PSEU276	Scfv-Fc	W4-M9	WO2014074528 SEQ ID NO: 152	7187
PSEU277	Scfv-Fc	Psl0170	WO20 14074528 SEQ ID NO: 245	7188
PSEU278	Scfv-Fc	Psl0304	WO20 14074528 SEQ ID NO: 246	7189
PSEU279	Scfv-Fc	Psi0348	WO20 14074528 SEQ ID NO: 247	7190
PSEU280	Scfv-Fc	PsI0573	WO20 14074528 SEQ ID NO: 248	7191
PSEU28 1	Scfv-Fc	Psi0574	WO20 14074528 SEQ ID NO: 249	7192
PSEU282	Scfv-Fc	PsI0582	WO20 14074528 SEQ ID NO: 250	7193
PSEU283	Scfv-Fc	Psl0584	WO20 14074528 SEQ ID NO: 251	7194
PSEU284	Scfv-Fc	Psi0585	WO20 14074528 SEQ ID NO: 252	7195
PSEU285	Scfv-Fc	PsI0589	WO20 14074528 SEQ ID NO: 253	7196

[00338] In one embodiment, ihe payioad region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payioad antibody polypeptides listed **in** Table 33 against Streptococcus **bacteria.**

Table 33. **Antibodies against Streptococcus bacteria**

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
STRP1	Heavy chain variable region, Diabody for Streptococcus		US7625561 SEQ ID NO: 5	7197
STRP2	Heavy chain variable region, Diabody for Streptococcus		US7625561 SEQ ID NO: 3	7198
STRP3	Heavy chain variable region, Diabody for Streptococcus		US7625561 SEQ ID NO: 7	7199
STRP4	Heavy chain variable region, partial, Streptococcus pneumoniae	DP-54	Lucas, A.H. "Combinatorial library cloning of human antibodies to Slreptococcus pneumoniae capsular polysaccharides: variable region primary structures and evidence for somatic mutation of Fab fragments specific for capsular serotypes 6B, 14, and 23F ' Infect. Immun. 69 (2), 853-864 (2001), NCBI Accession # AAD48823	7200
STOPS	Heavy chain variable region, partial, Streptococcus pneumoniae	DP-35	Lucas, A.H. "Combinatorial library cloning of human antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region primary structures and evidence for somatic mutation of Fab fragments specific for capsular serotypes 6B, 14, and 23F" Infect. Immun. 69 (2),	7201

			853-864 (2001), NCBI Accession # AAD48825	
STRP6	**Heavy** chain variable region, **partial,** Streptococcus **pneumoniae**	DP-47	Lucas, A.H. "Combinatorial **library** cloning **of** human antibodies **to** Streptococcus pneumoniae capsular **polysaccharides:** variable region primary **structures** and evidence **for** somatic **mutation of Fab fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. **Immun.** 69 (2), 853-864 (2001), NCBI Accession *it* **AAD48827**	7202
STRP7	Heavy **chain variable** region, partial, Streptococcus **pneumoniae**	DP-47	Lucas, **A.H.** "Combinatorial library cloning of **human** antibodies to Streptococcus pneumoniae **capsular polysaccharides:** variable region **primary** structures **and** evidence for somatic **mutation** of Fab **fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. Immun. 69 (2), 853-864 (2001), NCBI Accession # AAD48828	7203
STRP8	**Heavy** chain **variable** region, partial, Streptococcus pneumoniae	LSG6.1	Lucas, **A.H.** "Combinatorial **library** cloning of **human** antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region **primary structures** and evidence for somatic mutation of Fab fragments specific for capsular serotypes 6B, 14. and 23F" Infect. Immun. 69 **(2),** 853-864 (2001), NCBI Accession # AAD48830	7204
STRP9	Heavy chain variable region, partial, Streptococcus **pneumoniae**	LSG6.1	Lucas, A.H. **"Combinatorial library** cloning of **human** antibodies to Streptococcus **pneumoniae** capsular polysaccharides: variable region **primary** structures and evidence for somatic mutation of Fab fragments specific for capsular serotypes 6B, 14, and 23F" Infect. **Immun.** 69 (2), 853-864 (2001), NCBI Accession # AAD48832	7205
STRP10	**Heavy chain** variable region, partial, Streptococcus pneumoniae	DP-47	Lucas, A.H. "Combinatorial **library** cloning of human antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region **primary structures** and evidence for somatic mutation of Fab **fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. Immun. 69 (2), 853-864 (2001), NCBI Accession # AAD48835	7206
STRP11	Heavy chain variable region, Streptococcus **agalacriae, Legionella pneximophilia,** Streptococcus pyogenes, Escherichia coli, Neisseria gonorrhoeae, Neisseria **meningitidis, Pneumococciis, Hemophilis** influenzae B,	**humanize** d **Mu-9**	US7429381 SEQ ID NO: 14	6902

	Treponema pallidum, Lyme disease spirochetes, Pseudomonas aeruginosa, **Mycobacterium** leprae, **Bmcella** abortus and **Mycobacterium** tube rculosis.			
STRP12	**Heavy** chain **variable** region, Streptococcus pneumoniae	Anti-PsaA 7-1G9	**US20070003561** SEQ ID NO: 16	7207
STRP13	Heavy chain variable region, Streptococcus pneumoniae	Anii-PsaA 1-15E5	**US20070003561** SEQ ID NO: 32	7208
STRP14	**Heavy** chain variable region, Streptococcus pneumoniae	Anti-PsaA 9A7	**US20070003561** SEQ ID NO: 48	7209
STRP15	Heavy chain variable region, Streptococcus pneumoniae	23f Fab 023.102, chain B	**Bryson,** S., "**Multitasking** Immunoglobulin **V-Genes** And Somatic Div Cdr3 Loops Generate Binding Sites For **Chemically** Di Antigens From Bacterial And Viral Pathogens" Unpublished", **NCBI Accession** # 4HIE_B	7210
STRP16	**Heavy chain variable** region, Streptococcus **pneumoniae,** Escherichia coli, or Pseudomonas **aeruginosa**	5.12.14	**US5686070** SEQ ID NO: 22	7211
STRP17	**Heavy** chain variable region, Streptococcus pneumoniae, Escherichia coli, or Pseudomonas **aeruginosa**	6G4.2.5	**US5686070** SEQ ID NO: 50	7212
STRP18	Heavy chain variable region, Streptococcus pneumoniae, **Escherichia** coli, **or** Pseudomonas aeruginosa	**chimeric** 6G4.2.5	**US5686070** SEQ ID NO: 58	7213
STRP19	Heavy chain, Streptococcus agalactiae, Legionella **pneumophilia,** Streptococcus pyogenes, Escherichia coli, Neisseria gonorrhoeae, Neisseria meningitidis, **Pneumococcus, Hemophilus** influenzae B, Treponema **pallidum,** Lyme disease spirochetes, Pseudomonas aeruginosa, **Mycobacterium** leprae, Brucella abortus and Mycobacterium tube rculosis.	**Mab679**	US7429381 SEQ ID NO: 4	6900
STRP20	Light chain variable **region, Diabody for** Streptococcus		**US7625561 1** SEQ ID NO: 6	7214
STRP21	Light chain variable region, Diabody for Streptococcus		**US7625561** SEQ ID NO: 8	7215
STRP22	Light chain variable **region,** Diabody for Streptococcus		**US7625561 1** SEQ ID NO: 4	7216
STRP23	Light chain variable region, partial, Streptococcus pneumoniae	A2	Lucas, A.H. "Combinatorial **library** cloning of **human** antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region **primary** stnictares and evidence for somatic mutation of Fab **fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. **Immun.** 69 (2),	7217

			853-864 (2001), NCBI Accession # AAD48824	
STRP24	Light chain variable **region,** partial, Streptococcus **pneumoniae**	B3	Lucas, **A.H.** "Combinatorial library cloning **of** human antibodies **to** Streptococcus pneumoniae capsular **polysaccharides:** variable region primary structures and evidence for somatic **mutation of Fab fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. **Immun.** 69 (2), 853-864 (2001), NCBI Accession *it* **AAD48822**	7218
STRP25	Light chain variable region, partial, Streptococcus **pneumoniae**	A23	Lucas, **A.H. "Combinatorial** library cloning of human antibodies to Streptococcus pneumoniae **capsular polysaccharides:** variable region **primary** structures and evidence for somatic **mutation** of Fab fragments specific for capsular serotypes 6B, 14, and 23F" Infect. Immun. 69 (2), 853-864 (2001), NCBI Accession # AAD48826	7219
STRP26	Light chain variable region, partial, Streptococcus pneumoniae	L2	Lucas, **A.H.** "Combinatorial **library** cloning of **human** antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region **primary structures** and evidence for somatic mutation of Fab fragments specific for capsular serotypes 6B, 14. and 23F" Infect. Immun. 69 **(2),** 853-864 (2001), NCBI Accession # AAD48829	7220
STRP27	Light **chain** variable region, partial, Streptococcus **pneumoniae**	DPL5	Lucas, A.H. **"Combinatorial library** cloning of human antibodies to Streptococcus **pneumoniae** capsular polysaccharides: variable region primary structures and evidence for somatic mutation of Fab fragments specific for capsular serotypes 6B, 14, and 23F" Infect. **Immun.** 69 (2), 853-864 (2001), NCBI Accession # AAD4883 I	722 1
STRP28	Light chain variable region, partial, Streptococcus pneumoniae	DPL5	Lucas, A.H. "Combinatorial **library** cloning of human antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region **primary** structures and evidence for somatic mutation of Fab **fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. Immun. 69 (2), 853-864 (2001), NCBI Accession # AAD48833	7222
STRP29	Light chain variable region, partial, Streptococcus pneumoniae	L2	Lucas, A.H. "Combinatorial **library** cloning of **human** antibodies to Streptococcus pneumoniae capsular polysaccharides: variable region primary structures and evidence for somatic **mutation of Fab fragments** specific for capsular serotypes 6B, 14, and 23F" Infect. Immun. 69 (2),	7223

			853-864 (2001), NCBI Accession # AAD48834	
STRP30	Light chain variable **region,** Streptococcus **pneumoniae**	Anii-PsaA 7-1G9	**US20070003561** SEQ ID **NO:** 8	7224
STRP31	Light chain variable region, Streptococcus pneumoniae	Anti-PsaA 1-15E5	**US20070003561** SEQ ID NO: 24	7225
STRP32	Light chain variable region, Streptococcus pneumoniae	Anii-PsaA 9A7	**US20070003561** SEQ **ID NO:** 40	7226
STRP33	Light chain variable region, Streptococcus pneumoniae, Escherichia coli, or **Pseudomonas aeruginosa**	5.12.14	**US5686070** SEQ ID NO: 20	**7227**
STRP34	Light chain variable region, Streptococcus pneumoniae, Escherichia coli, or Pseudomonas aeruginosa	6G4.2.5	**US5686070** SEQ ID NO: 48	7228
STRP35	Light chain variable region, Streptococcus pneumoniae, Escherichia coli, or **Pseudomonas aeruginosa**	chime sic 6G4.2.5	**US5686070** SEQ **ID NO:** 56	7229
STRP36	Light chain, Streptococcus **agalactiae,** Legionella **pneumophilia,** Streptococcus pyogenes, **Escherichia** coli, Neisseria **gonorrhoeae.** Neisseria meningitidis, **Pneumococcus,** Hemophilis **influenzae** B, Treponema pallidum, Lyme disease spirochetes, **Pseudomonas** aeruginosa, **My**cobacte **rium** leprae. Brucella abostus arsd **Mycobacterium** rube **rculosis**	**Mab679**	**US742938** 1 SEQ ID NO: 2	6907
STRP37	**scFv,** Streptococcus **agalactiae, Legionella pneumophila,** Streptococcus pyogenes, **Escherichia** coli, Neisseria gonorrhoeae, **Neisseria** meningitidis, Pneumococcus, Hemophilis influenzae B, Treponema pallidum, Lyme disease spirochetes, Pseudomonas aeruginosa, Mycobacterium leprae, **Bmcella** abostus arsd **Mycobacterium** rube rculosis	Mab679	US742938 1 SEQ **ID NO:** 6	6911
STRP38	**scFv,** Streptococcus agalactiae, Legionella **pneumophila,** Streptococcus pyogenes, **Escherichia** coli, Neisseria gonorrhoeae, Neisseria meningitidis, Pneumococcus, Hemophilis influenzae B, Treponema palirdusn, **Lyme** disease spirochetes, Pseudomonas aeruginosa, Mycobacterium leprae, **Brucella**	**Mu-9V**	**US742938 1** SEQ ID NO: 8	6905

	abortus and Mycobacterium tube rculosis			
STRP39	**scFv,** Streptococcus agalactiae, **Legionella pneumophila,** Streptococcus pyogenes, Escherichia coli, Neisseria gonorrhoeae, Neisseria meningitidis, **Piieumococcus, Hemophilis** influenzae B, Treponema pallidum, Lyme disease spirochetes, **Pseudomonas** aeruginosa, Mycobacterium leprae, **Brucella** abortus and Mycobacterium rube rculosis	**Mu-9V**	**US742938 1** SEQ ID NO: **10**	6901
STRP40	**scFv,** Streptococcus agalactiae, Legionella pneumophilia, Streptococcus pyogenes, Escherichia coli, Neisseria gonorrhoeae, **Neisseria** meningitidis, **Pneumococcus,** Hemophilis influenzae B, Treponema pallidum, Lyme disease spirochetes, Pseudomonas aeruginosa, Mycobacterium leprae, Brucella abortus and Mycobacterium tube rculosis	humanize d **Mu-9**	**US742938** 1 SEQ ID NO: 12	6906

[00339] In one embodiment the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in US Pub No. US200401 98960 and IJS20130195876, the contents of each of which are herein incorporated by reference in their entirety, against *Streptococcus Pneumoniae* infection.

[00340] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants encoding Afeliraomah, fragments or variants thereof for treating a disease and/or disorder or preventing a disease and/or disorder. As a non-limiting example, the disease and/or disorder is sepsis.

[00341] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants encoding Nebacumab, fragments or variants thereof for treating a disease and/or disorder or preventing a disease and/or disorder. As a non-limiting example, the disease and/or disorder is sepsis.

[00342] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 34 against Staphylococcal bacteria and related bacteria.

Table 34. Antibodies against Staphylococcal bacteria and related bacteria

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
STPH i	Heavy chain variable region, S. aureus		US8609102 SEQ ID NO: 2	7230
STPH2	Heavy chain vanable region, S. aureus		US8609102 SEQ ID NO: 6	7231
STPH3	Heavy chain variable region, S. aureus or S. epidemridis, E. co3i, Yersinia pestis (Y. pestis), Y. entercolitica, Xanthomnosias axonopodis (X. axonopodis), Pseudonmonas fuorescerns (P. iluorescens), Actinobacillus actinomycetemcoiriitans (A. actinomycetemcomitans), A. pleuropneumoniae, Ralstonia solanacearuni (R. solanacearum), Bordetella pertussis (B. pertussis), B. parapertussis or B. bronchiseptica	SAR279356	US7786255 SEQ ID NO: 1	7232
STPH4	Heavy chain variable region, S. aureus or S. epiderniidis, E. coli, Yersinia pestis (Y. pestis), Y. entercolitica, Xanthomnonas axonopodis (X. axonopodis), Pseudonmonas fuorescerns (P. Iluorescens), Actinobacillus actinoniycetemcomitans (A. actinomycetemcomitans), A. pleuropneumoniae, Ralstonia solanacearum (R. solanacearum), Bordetella pertussis (B. pertussis), B. parapertussis or B. bronchiseptica	SAR279356	US20110002932 SEQ ID NO: 1	7233
STPH5	Heavy chain variable region, S. epiderniidis	108-1	US8475798 SEQ ID NO: 18	7234
STPH6	Heavy cluain variable region, S. epiderniidis	108-36	US8475798 SEQ ID NO: 22	7235
STPH7	Heavy chain variable region, S. epiderniidis	110-15	US8475798 SEQ ID NO: 26	7236
STPH 8	Heavy chain variable region, S. ejjidermidis	108-1VH-Hu	US8475798 SEQ ID NO: 28	7237
STPH9	Heavy chain vanable region, S. epiderniidis	108-36VH-Hu	US8475798 SEQ ID NO: 30	7238
STPH 10	Heavy chain variable region, S. ejjidermidis	110-15VH-Hu	US8475798 SEQ ID NO: 32	7239
STPH 11	Heavy chain vanable region, Staphylococcal sepsis	Pagibaximab	US8372958 SEQ ID NO: 87	7240
STPH 12	Heavy chain variable region, Staphylococcal sepsis	Pagibaximab	US8372958 SEQ ID NO: 12	7241
STPH 13	Heavy chain vanable region, Staphylococcal sepsis	Pagibaximab	US8372958 SEQ ID NO: 17	7242
STPH 14	Heavy chain variable region, Staphylococci such as S. aureus and S. epidermidis, E, coli such as E. coli strains 0157:H7 and CFT073, Yersinia pestis, Yersinia entercolitica, Xanthomonas axonopodis, Pseudomonas fluorescens Call of which	P628	US8912314 SEQ ID NO: 3	7243

	are sequenced species with complete **pgaABCD** iocs), and **Actinobacillus** actinomycetemcomitans (AA), Actmoba **cillus** pleuropneumoniae (Ap), Ralstonia solanacearum (e.g., **niegaplasmid** form), **Bordetella** pertussis, **Bordetella** parapertussis and Bordetella bronchiseptica			
STPH15	Heavy chain variable region, Staphylococci such as S. aureus and S. epidermidis, E. **coli** such as **E.** coli strains **0157:H7** and CFT073, Yersinia pestis, Yersinia entercolitica, **Xanthomonas** axonopodis, **Pseudomonas** fluorescens **(all of** which are sequenced species with **complete** pgaABCD loci), and Actinobacillus **actinomycetemcomitans** (AA), Actinoba cillus **pleuropneumoniae** (Ap), Ralstonia **solanacearum** (e.g., **megaplasmid** form), Bordetella **pertussis,** Bordetella parapertussis and Bordetella **bronchiseptica**	F630	**US89123 14 SEQ** ID NO: 5	7244
STPH I6	Heavy chain variable region, Staphylococci **such as** S. aureus and S. epidermidis, E. coli such as E. coli strains 0157:H7 and CFT073, Yersinia pestis, Yersinia entercolitica, Xanthomonas axonopodis, Pseudomonas fluorescens (all of **which** are sequenced species with complete pgaABCD loci), and Actinobacillus **actinomycetemcomitans** (AA), Actinoba cillus **pleuropneumoniae (Ap),** Ralstonia **solanacearum** (e.g., **megaplasmid** form), Bordetella pertussis, Bordetella parapertussis and Bordetella **bronchiseptica**	FS98	**US89123** 14 SEQ **ID NO: 55**	7245
STPH17	**Heavy chain** variable region. Staphylococci such as S. aureus and S. epidermidis, **E.** coli such as E. coli strains **0157:H7** and **CFT073,** Yersinia pestis, Yersinia entercolitica, Xanthomonas axonopodis, Pseudomonas fluorescens **(all** of which are sequenced species with complete pgaABCD loci), and Actinobacillus **actinomycetemcomitans** (AA), Actinoba cillus **pleuropneumoniae (Ap),** Ralstonia solanacearum (e.g., megaplasmid **form),** Bordetella pertussis, Bordetella parapertussis and Bordetella bronchiseptica	F628	**US89123 14** SEQ ID NO: 58	7246
STPH18	Heavy chain variable region, Staphylococci such as S. aureus and S. **epidermidis,** E. coli such as E. coli strains 0157:H7 and **CFT073,** Yersinia **pestis,** Yersinia entercolitica, Xanthomonas axonopodis, Pseudomonas fluorescens (ail of which	F598	**US89 I2314 SEQ ID NO:** i	7247

	are sequenced species with complete **pgaABCD** iocs), and **Actinobacillus** actinomycetemcornitans (AA), Actinoba cilkis pleuropneumoniae (Ap), Ralstonia soianaeearum (e.g., **megaplasmid** form), **Bordetella** pertussis, **Bordetella** parapertussis and Bordetella bronchiseptica			
STPH19	Heavy chain variable region, Staphylococcus **epidermidis**	I08-3BVH-Hu	US8475798 SEQ ID NO: 34	7248
STPH20	**Heavy chain,** MRSA, MSSA	2B2	**US8735554** SEQ ID NO: 3	7249
STPH21	Heavy chain, **MRSA,** MSSA	2G7	**US8735554** SEQ ID NO: **5**	7250
STPH22	**Heavy chain,** MRSA, MSSA	3B12	**US8735554** SEQ ID NO: 7	7251
STPH23	Heavy chain, S. aureus	DF1.i	US8715673 SEQ **ID** NO: 2	7252
STPH24	Heavy chain, S. aureus	DF1	US8715673 SEQ ID NO: 4	7253
STPH25	**Heavy** chain, S. aureus	DF2	US8715673 SEQ ID NO: 35	7254
STPH26	Heavy chain, S. aureus	DF3	US8715673 SEQ ID NO: 36	7255
STPH27	**Heavy chain,** S. aureus	DF4	US8715673 SEQ ID NO: 37	7256
STPH28	Heavy chain, S. aureus	DPS	US8715673 SEQ ID NO: 38	7257
STPH29	Heavy chain, S. aureus	DF6	US8715673 SEQ ID NO: 39	7258
STPH30	**Heavy** chain, S. aureus	DF7	US8715673 SEQ ID NO: 40	7259
STPH31	Heavy chain, S. aureus	DF8	US8715673 SEQ ID NO: 41	7260
STPH32	**Heavy chain,** S. aureus	DF9	US8715673 SEQ ID NO: 42	7261
STPH33	Heavy chain, S. aureus	DF10	US8715673 SEQ ID NO: 43	7262
STPH34	Heavy chain, S. aureus	DF11	US8715673 SEQ ID NO: 44	7263
STPH35	**Heavy** chain, S. aureus	DF12	US8715673 SEQ ID NO: 45	7264
STPH36	Heavy chain, S. aureus	DF13	US8715673 SEQ ID NO: 46	7265
STPH37	**Heavy chain,** S. aureus	DF14	US8715673 SEQ ID NO: 47	7266
STPH38	Heavy chain, S. aureus	DP15	US8715673 SEQ ID NO: 48	7267
STPH39	Heavy chain, S. aureus	DF15	US8715673 SEQ ID NO: 49	7268
STPH40	**Heavy** chain, S. aureus	DP17	US8715673 SEQ ID NO: 50	7269
STPH41	Heavy chain, S. aureus	DF18	US8715673 SEQ ID NO: 51	7270
STPH42	**Heavy chain,** S. aureus	DF19	US8715673 SEQ ID NO: 52	7271
STPH43	Heavy chain, S. aureus	DF20	US8715673 SEQ ID NO: 53	7272
STPH44	Heavy chain, S. aureus and S. **epidermiclis**	CR2430	US8460666 SEQ ID NO: 26	7273
STPH45	Heavy chain, S. aureus and S. **epidermidis**	CR5132	US8460666 SEQ ID NO: **28**	7274
STPH46	Heavy chain, S. aureus and S. **ejridermidis**	CR5133	US8460666 SEQ ID NO: 30	7275
STPH47	**Heavy** chain, S. aureus and S. epidermidis	CR6166	US8460666 SEQ ID NO: 117	7276
STPH48	Heavy chain, S. aureus and S. **ejridermidis**	CR6171	US8460666 SEQ ID NO: 119	7277
STPH49	**Heavy** chain, S. aureus and S. epidermidis	CR6176	US8460666 SEQ **ID** NO: 121	7278
STPH50	Heavy chain, S. aureus and S. **ejridermidis**	CR6187	US8460666 SEQ ID NO: 123	7279
STPH51	**Heavy** chain, S. aureus and S. epidermidis	CR6193	US8460666 SEQ ID NO: 125	7280
STPH52	Heavy chain, S. aureus and S. **epidermidis**	CR6249	US8460666 SEQ ID NO: 127	7281

STPH53	Heavy chain, S. aureus and S. epidermidis	CR6273	US8460666 SEQ ID NO: 129	7282
STPH54	Heavy chain, S. aureus and S. epidermidis	CR6389	US8460666 SEQ ID NO: 131	7283
STPH55	Heavy chain, S. aureus and S. epidermidis	CR6403	US8460666 SEQ ID NO: 133	7284
STPH56	Heavy chain, S. aureus arsd S. epidermidis	CR6406	US8460666 SEQ **ID** NO: 135	7285
STPH57	Heavy chain, S. aureus and S. epidermidis	CR6410	US8460666 SEQ ID NO: 137	7286
STPH58	Heavy chain, S. aureus arsd S. epidermidis	CR6446	US8460666 SEQ **ID** NO: 139	7287
STPH59	Heavy chain, S. aureus and S. epidermidis	CR6450	US8460666 SEQ ID NO: 141	7288
STPH60	Heavy chain, S. aureus arsd S. epidermidis	CR6452	US8460666 SEQ **ID** NO: 143	7289
STPH61	Heavy chain, S. aureus and S. epidermidis	CR6453	US8460666 SEQ ID NO: 145	7290
STPH62	Heaw chain, S. aureus arsd S. epidermidis	CR6464	US8460666 SEQ **ID** NO: 147	7291
STPH63	Heavy chain, S. aureus and S. ejjidermidis	CR6471	US8460666 SEQ ID NO: 149	7292
STPH64	Heaw chain, S. aureus and S. epidermidis	CR6516	US8460666 SEQ ID NO: 151	7293
STPH65	Heavy chain, S. aureus and S. ejjidermidis	CR6517	US8460666 SEQ ID NO: 153	7294
STPH66	Heaw chain, S. aureus and S. epidermidis	CR6526	US8460666 SEQ **ID** NO: 155	7295
STPH67	Heavy chain, S. aureus and S. ejjidermidis	CR6528	US8460666 SEQ ID NO: 157	7296
STPH68	Heaw chain, S. aureus and S. epidermidis	CR6531	US8460666 SEQ **ID** NO: 159	7297
STPH69	Heavy chain, S. aureus and S. epidermidis	CR6533	US8460666 SEQ ID NO: 161	7298
STPH70	Heaw chain, S. aureus and S. epidermidis	CR6536	US8460666 SEQ **ID** NO: 163	7299
STPH71	Heavy chain, S. aureus and S. epidermidis	CR6537	US8460666 SEQ ID NO: 165	7300
STPH72	Heaw chain, S. aureus and S. epidermidis	CR6538	US8460666 SEQ ID NO: 167	7301
STPH73	Heavy chain, S. aureus and S. epidermidis	CR6540	US8460666 SEQ ID NO: 169	7302
STPH74	Heaw chain, S. aureus and S. epidermidis	CR6544	US8460666 SEQ ID NO: 171	7303
STPH75	Heavy chain, S. aureus and S. epidermidis	CR6566	US8460666 SEQ ID NO: 173	7304
STPH76	Heaw chain, S. aureus and S. epidermidis	CR6625	US8460666 SEQ **ID** NO: 175	7305
STPH77	Heavy chain, S. aureus, Enterococcus	CR5140	US862S776 SEQ ID NO: 395	7306
STPH78	Heaw chain, S. aureus, Enterococcus	CR5159	US8628776 SEQ ID NO: 82	7307
STPH79	Heavy chain, S. aureus, Enterococcus	CR5179	US8628776 SEQ **ID** NO: 399	7308
STPH80	Heavy chain, S. aureus, Enterococcus	CR6016	US8628776 SEQ ID NO: 88	7309
STPH81	Heaw chain, S. aureus, Enterococcus	CR6049	US8628776 SEQ **ID** NO: 92	7310
STPH82	Heavy chain, S. aureus, Enterococcus	CR6071	US862S776 SEQ ID NO: 94	7311
STPH83	Heaw chain, S. aureus, Enterococcus	CR6078	US8628776 SEQ ID NO: 96	7312
STPH84	Heavy chain, S. aureus, Enterococcus	CR6086	US8628776 SEQ ID NO: 407	7313

STPH85	**Heavy chain,** S. aureus, Enteroeoccus	CR6089	U58628776 SEQ ID NO: 213	7314
STPH86	Heavy chain, S. aureus, Enteroeoccus	CR6191	US8628776 SEQ ID NO: 411	7315
STPH87	Heavy chain, S. aureus, Enteroeoccus	CR6198	US8628776 SEQ ID NO: 415	7316
STPH88	**Heavy chain,** S. aureus, Enteroeoccus	CR6242	US8628776 SEQ ID NO: 417	7317
STPH89	Heavy chain, S. aureus, Enteroeoccus	CR6252	US8628776 SEQ ID NO: 100	7318
STPH90	**Heavy chain,** S. aureus, Enteroeoccus	CR6389	US8628776 SEQ ID NO: 423	7319
STPH9 i	Heavy chain, S. aureus, Enteroeoccus	CR6402	US8628776 SEQ ID NO: 427	7320
STPH92	Heavy chain, S. aureus, Enteroeoccus	CR6415	US8628776 SEQ ID NO: 431	7321
STPH93	**Heavy** chain, S. aureus, Enteroeoccus	CR6429	US8628776 SEQ ID NO: 435	7322
STPH94	Heavy chain, **S.** aureus, Enteroeoccus	**CR5140**	US8628776 SEQ ID NO: 439	7323
STPH95	**Heavy chain,** S. aureus, Enteroeoccus	CR5159	US8628776 SEQ ID NO: 102	7324
STPH96	Heavy chain, S. aureus, Enteroeoccus	CR5179	US8628776 SEQ ID NO: 443	7325
STPH97	Heavy chain, S. aureus, Enteroeoccus	CR6016	US8628776 SEQ ID NO: 108	7326
STPH98	**Heavy** chain, S. aureus, Enteroeoccus	CR6049	US8628776 SEQ ID NO: 112	7327
STPH99	Heavy chain, S. aureus, Enteroeoccus	CR6071	US8628776 SEQ ID NO: 114	7328
STPH100	**Heavy chain,** S. aureus, Enteroeoccus	CR6078	U88628776 SEQ ID NO: 116	7329
STPH101	Heavy chain, S. aureus, Enteroeoccus	CR6086	US8628776 SEQ ID NO: 451	7330
STPH I02	Heavy **chain, S**. aureus, Enteroeoccus	CR6089	US8628776 SEQ ID NO: 217	7331
STPH103	**Heavy** chain, S. aureus, Enteroeoccus	CR6191	US8628776 SEQ ID NO: 455	7332
STPH104	Heavy chain, S. aureus, Enteroeoccus	CR6I98	US8628776 SEQ ID NO: 459	7333
STPH105	**Heavy chain,** S. aureus, Enteroeoccus	CR6242	US8628776 SEQ ID NO: 461	7334
STPH106	Heavy chain, S. aureus, Enteroeoccus	CR6252	US8628776 SEQ ID NO: 120	7335
STPH I07	Heavy **chain, S**. aureus, Enteroeoccus	CR6389	US8628776 SEQ ID NO: 467	7336
STPH108	**Heavy** chain, S. aureus, Enteroeoccus	CR6402	US8628776 SEQ ID NO: 471	7337
STPH109	Heavy chain, S. aureus, Enteroeoccus	CR6415	US8628776 SEQ ID NO: 475	7338
STPH110	**Heavy chain,** S. aureus, Enteroeoccus	CR6429	US8628776 SEQ ID NO: 479	7339
STPHl 11	Heavy chain, S. **aureus,** S. **epidermidis,** S. **caprae,** S. **saprophytics,** S. capitis, or **metmcillin-resistanl** S. aureus **(MRSA)**	F1 **antibody variant**	US8617556 SEQ ID NO: 7	7340
STPH1 I2	Heavy chain, S. aureus, S. epidermidis, S. caprae, S. saprophyticus, S. capitis, or **methicillin-resistant** S. aureus (MRSA)	**F1** antibody variant	US8617556 SEQ ID NO: 9	7341
STPH113	**Heavy** chain, S. aureus, S. epidermidis, S. caprae, S. saprophyticus, S. capitis, or **methicillin-resistant** S. aureus **(MRSA)**	rFl	US8617556 SEQ ID NO: 55	7342
STPH114	**Heavy chain,** S. aureus, S. epidermidis, S. caprae, S. saprophyticus, S. capitis, or methicillin-resistant S. aureus (MRSA)	rF! **A 114C**	US8617556 SEQ ID NO: 56	7343
STPH I15	Heavy chain, S. aureus, S. **epidermidis,** S. caprae, S. saprophyticus, S. capitis, or methicillin-resistant S. aureus (MRSA)	**rF** l	OS8617556 SEQ ID NO: 63	7344
STPH116	Heavy chain, S. aureus, S. **epidermidis,** S. caprae, S. saprophyticus, S. capitis, or methicillin-resistant S. aureus (MRSA)	**rFl A 114C**	US8617556 SEQ ID NO: 62	7345
STPH1 17	Light **chain**		US8609102 **SEQ** ID NO: 4	7346
STPH118	Light chain variable region, S. aureus		US8609102 SEQ ID NO: 8	7347
STPH1 I9	Light chain variable region, S. aureus or S. **epidermidis,** E. coli, Yersinia **pestis** (Y. **pestis),** Y. entercolitica, **Xanthomnonas axonopodis** (X.	SAR279356	US7786255 SEQ ID NO: 2	7348

	axonopodis), **Pseudonmonas** fuorescerns **(P.** fluorescens), **Actinobacillus actinomycetemcomitans** (A. **actinomycetemcomitans), A. pleiiropiieumoniae, Ralstoiiia solanacearum (R. solanacearam),** Bordetella pertussis (B. pertussis), B. **parapertussis or** B. **bronchiseptica**			
STPH120	Light chain variable region, S. aureus or S. **epidermidis, E.** coli, Yersinia pestis (Y. pestis), Y. entercolitica, **Xanthomnonas** axonopodis (X. axonopodis), **Pseudonmonas fuorescerns** (P. fluorescens), Actinobacillus **actinonwcetemcomitans (A. actinomycetemcomitans), A. pleuropneumoniae,** Ralstoiiia solanacearum (R. **solanacearam),** Bordetella pertussis (B. pertussis), **B. parapertussis** or **B. bronchisejjtica**	SAR279356	**US20 110002932** SEQ ID NO: 2	7349
STPH121	Light chain variable region, S. epidermidis	108-1	US8475798 SEQ ID NO: 16	7350
STPH 122	Light chain variable region, S. **epidermidis**	108-36	US8475798 SEQ ID NO: 20	7351
STPH123	Light chain variable region, S. epidermidis	110-15	US8475798 SEQ ID NO: 24	7352
STPH 124	Light chain variable region, S. **epidermidis**	108-1VL-Hu	US8475798 SEQ ID NO: 27	7353
STPH125	Light chain variable region, S. epidermidis	108-36VL-Hu	US8475798 SEQ ID NO: 29	7354
STPH126	Light **chain variable** region, S. **epidermidis**	110-15VL-Hu	US8475798 SEQ ID NO: 31	7355
STPH127	Light chain variable region, Staphylococcal sepsis	**Pagibaxima** b	US8372958 SEQ ID NO: 89	7356
STPH128	Light **chain** variable region, Staphylococcal sepsis	Pagibaxima b	US8372958 SEQ ID NO: 10	7357
STPH129	Light chain variable region, Staphylococcal sepsis	**Pagibaxima** b	US8372958 SEQ ID NO: 16	7358
STPH130	Light **chain** variable region, Staphylococci such as S. aureus and S. **epidermidis,** E. coli such as **E.** coli strains **0157** :117 and CFT073, Yersinia pestis, Yersinia entercolitica, **Xanthomonas axonopodis, Pseudomonas** fluorescens **(all of which** are sequenced species with complete **pgaABCD** loci), and Actinobacillus **actinomycetemcomitans** (AA), **Actinoba cillus pleuropneumoniae** (Ap), Ralstoiiia solanacearum (e.g., **megaplasmid** form), Bordetella **pertussis,** Bordetella parapertussis and Bordetella **bronchiseptica**	F598	US8912314 SEQ ID NO: 2	7359
STPH 131	Light chain variable region, Staphylococci such as S. aureus and S. epidermidis, E. coli such as E.	P628	**US8912314** SEQ **ID NO:** 4	7360

	coli strains 0157:H7 and CFT073, **Yersinia** pestis, Yersinia **entercolitica, Xanthomonas** axonopodis, **Pseudomonas** fluorescens **(all** of **which** are **sequenced** species **with** complete **pgaABCD** loci), and Actinobacilhis **actinomycetemcomitans** (AA), Actinoba cillus **pleuropneumoniae (Ap), Ralstonia solanacearam** (e.g., **megaplasmid** form), **Bordetella pertussis, Bordetella** parapertussis and Bordetella broncliiseptica			
STPH132	Light chiin variable region, Staphylococci **such as** S. aureus and S. **epidermidis,** E. coli such as E. coli strains **0157:H7** and **CFT073,** Yersinia pestis, Yersinia entercolitica, **Xanthomonas** axonopodis, **Pseudomonas fluorescens (all** of which are sequenced species with complete pgaABCD loci), and **Actinobacilhis actinomycetemcomitans** (AA), Actinoba cillus **pleuropneumoniae (Ap),** Ralstonia solanacearam (e.g., megaplasmid **form),** Bordetella pertussis, Bordetella **parapertussis** and Bordetella **broncliiseptica**	F630	**US89123 14** SEQ ID NO: **6**	7361
STPH133	Light chain variable region, Staphylococci such as S. aureus and S. **epidermidis,** E. coli such as E. coli strains 0 !57:H7 and **CFT073,** Yersinia pestis, Yersinia entercolitica, Xanthomonas axonopodis, Pseudomonas fluorescens (ail of which are sequenced species with complete pgaABCD **loci),** and **Actinobacilhis** actinomycetemcomitans (AA), Actinoba cillus pleuropneumoniae **(Ap),** Ralstonia solanacearam (e.g., megaplasmid form), Bordetella pertussis, Bordetella parapertussis and Bordetella **broncliiseptica**	F598	**US89 I2314** SEQ **ID** NO: **57**	7362
STPH134	Light **chain** variable region, Staphylococci such as S. aureus and S. **epidermidis,** E. coli such as E. coli strains **0157:H7** and CFT073, Yersinia pestis, Yersinia entercolitica, Xanthomonas **axonopodis,** Pseudomonas **fluorescens (all of which** are sequenced species **with** complete pgaABCD loci), and Actinobacilius actinomycetemcomitans (AA), Actinoba cillus pleuropneumoniae (Ap), Ralstonia solanacearam (e.g., megaplasmid form), Bordetella **pertussis,** Bordetella parapertussis and Bordetella broncliiseptica	F630	**US89123** 14 SEQ ID NO: 60	7363
STPH135	Light chain, MRSA, **MSSA**	*2B2*	**US8735554** SEQ **ID** NO: **2**	7364
STPH136	Light chain. MRSA, MSSA	2G7	**US8735554** SEQ **ID** NO: 4	7365

STPH137	Light chain, MRS A, MSSA	3B12	**US8735554** SEQ ID NO: **6**	7366
STPH138	Light chain, S. aureus	DF1.1	US8715673 SEQ **ID** NO: **E**	7367
STPH139	Light chain, **S.** aureus	DF1-DF20	US8715673 SEQ ID NO: 3	7368
STPH140	Light chain, **S.** aureus and **S.** epidermidis	CR2430	USS460666 **SEQ** ID NO: 32	7369
STPH141	Light **chain, S.** aureus and **S.** epidennidis	CR5132	US8460666 SEQ ID NO: 34	7370
STPH 142	Light chain, **S.** aureus and **S.** epidennidis	CR5133	U58460666 SEQ ID NO: 36	7371
STPH143	Light chain, S. aureus and **S.** epidennidis	CR6166	US8460666 SEQ **ID** NO: 177	7372
STPH 144	Light chain, **S.** aureus and S. epidennidis	CR6171	US8460666 SEQ ID NO: 179	7373
STPH145	Light chain, **S.** aureus and **S. epidermidis**	CR6176	US8460666 SEQ ID NO: 181	7374
STPH146	Light **chain, S.** aureus and **S.** epidennidis	CR6187	US8460666 SEQ ID NO: 183	7375
STPH 147	Light chain, **S.** aureus and S. epidermidis	CR6193	US8460666 SEQ ID NO: 185	7376
STPH148	Light chain, S. aureus and **S.** epidermidis	CR6249	US8460666 SEQ **ID** NO: 187	7377
STPH149	Light chain, **S.** aureus and S. epidennidis	CR6273	US8460666 SEQ ID NO: 189	7378
STPH150	Light chain, S. aureus and **S. epidermidis**	CR6389	US8460666 SEQ **ID** NO: 191	7379
STPH151	Light **chain, S.** aureus and **S.** epidennidis	CR6403	US8460666 SEQ ID NO: 193	7380
STPH152	Light chain, **S.** aureus and S. epidermidis	CR6406	U58460666 SEQ ID NO: 195	7381
STPH153	Light chain, S. aureus and **S.** epidennidis	CR6410	US8460666 SEQ ID NO: 197	7382
STPH I54	Light chain, **S.** aureus and S. epidennidis	CR6446	US8460666 SEQ ID NO: 199	7383
STPH155	Light chain, S. aureus and **S. epidermidis**	CR6450	US8460666 SEQ ID NO: 201	7384
STPH156	Light **chain, S.** aureus and **S.** epidennidis	CR6452	US8460666 SEQ ID NO: 203	7385
STPH157	Light chain, **S.** aureus and **S.** epidermidis	CR6453	U58460666 **SEQ** ID NO: 205	7386
STPH158	Light chain, S. aureus and **S.** epidennidis	CR646-4	US8460666 SEQ ID NO: 207	7387
STPH I59	Light chain, **S.** aureus and S. epidennidis	CR647 1	US8460666 SEQ ID NO: 209	7388
STPH160	Light chain, S. aureus and **S. epidermidis**	CR6516	US8460666 SEQ ID NO: 211	7389
STPH161	Light **chain, S.** aureus and **S.** epidennidis	CR6517	US8460666 SEQ ID NO: 213	7390
STPH162	Light chain, **S.** aureus and **S.** epidermidis	CR6526	U58460666 **SEQ** ID NO: 215	7391
STPH163	Light chain, S. aureus and **S.** epidennidis	CR6528	US8460666 SEQ ID NO: 217	7392
STPH I64	Light chain, **S.** aureus and S. epidennidis	CR6531	US8460666 SEQ ID NO: 219	7393
STPH165	Light chain, S. aureus and **S. epidermidis**	CR6533	US8460666 SEQ ID NO: 221	7394
STPH166	Light chain, **S.** aureus and S. epidennidis	CR6536	US8460666 SEQ ID NO: 223	7395
STPH167	Light chain, **S.** aureus and **S.** epidermidis	CR6537	U58460666 **SEQ** ID NO: 225	7396
STPH168	Light chain, S. aureus and **S.** epidennidis	CR6538	US8460666 SEQ ID NO: 227	7397
STPH I69	Light chain, **S.** aureus and S. epidennidis	CR6540	US8460666 SEQ ID NO: 229	7398
STPH170	Light chain, S. aureus and **S. epidermidis**	CR6544	US8460666 SEQ ID NO: 231	7399
STPH 171	Light chain, **S.** aureus and S. epidennidis	CR6S66	US8460666 SEQ ID NO: 233	7400
STPH172	Light chain, **S.** aureus and **S.** epidermidis	**CR6625**	U58460666 **SEQ** ID NO: 235	7401
STPH173	Light chain, S. aureus, **Enterococcus**	CR5157	US8628776 SEQ ID NO: 397	7402
STPH174	Light chain, **S.** aureus, Enterococcus	**CR5166**	US8628776 SEQ ID NO: 84	7403
STPH175	Light chain, S. aureus, Enterococcus	CR5187	US8628776 SEQ ID NO: 86	7404
STPH 176	Light chain, **S.** aureus, Enterococcus	CR6043	US8628776 SEQ ID NO: 90	7405
STPH177	Light chain, **S.** aureus, Enterococcus	CR6050	US8628776 SEQ ID NO: 401	7406
STPH178	Light **chain, S.** aureus, Enterococcus	CR6077	US8628776 SEQ ID NO: 403	7407
STPH179	Light chain, **S.** aureus, Enterococcus	CR6079	US8628776 SEQ ID NO: 405	7408
STPH180	Light chain, S. aureus, Enterococcus	CR6087	US8628776 SEQ ID NO: 211	7409
STPH181	Light chain, **S.** aureus, Enterococcus	CR6092	US8628776 SEQ ID NO: 409	7410

STPH182	Light chain, S. aureus, Enterococcus	CR6195	US8628776 SEQ ID NO: 413	7411
STPH183	Light chain, S. aureus, Enterococcus	CR6241	US8628776 SEQ ID NO: 98	7412
STPH184	Light chain, **S.** aureus, **Enterococcus**	CR6246	US8628776 SEQ ID NO: 419	7413
STPH185	Light **chain,** S. aureus, Enterococcus	CR6388	US8628776 SEQ ID NO: 421	7414
STPH186	Light **chain,** S. aureus, Enterococcus	CR6396	US8628776 SEQ ID NO: 425	7415
STPH187	Light chain, S. aureus, Enterococcus	CR6409	US8628776 SEQ ID NO: 429	7416
STPH188	Light chain, S. aureus, Enterococcus	CR6421	US8628776 SEQ ID NO: 433	7417
STPH189	Light chain, **S.** aureus, **Enterococcus**	CR6432	US8628776 SEQ ID NO: 437	7418
STPH190	Light chain, S. **aureus, Enterococcus**	CR5157	US8628776 SEQ ID NO: 441	7419
STPH191	Light **chain,** S. aureus, Enterococcus	CR5166	US8628776 SEQ ID NO: 104	7420
STPH192	Light chain, S. aureus, Enterococcus	CRS187	U58628776 SEQ ID NO: 106	7421
STPH193	Light chain, S. aureus, Enterococcus	CR6043	US8628776 SEQ ID NO: 110	7422
STPH194	Light chain, **S.** aureus, **Enterococcus**	CR6050	US8628776 SEQ ID NO: 445	7423
STPH195	Light chain, S. aureus, Enterococcus	CR6077	US8628776 SEQ ID NO: 447	7424
STPH196	Light **chain,** S. aureus, Enterococcus	CR6079	US8628776 SEQ ID NO: 449	7425
STPH197	Light chain, S. aureus, Enterococcus	CR6087	U58628776 SEQ ID NO: 215	7426
STPH198	Light chain, S. aureus, Enterococcus	CR6092	US8628776 SEQ ID NO: 453	7427
STPH I99	Light chain, **S.** aureus, **Enterococcus**	CR6195	US8628776 SEQ ID NO: 457	7428
STPH200	Light chain, S. aureus, Enterococcus	CR6241	US8628776 SEQ ID NO: 115	7429
STPH201	Light **chain,** S. aureus, Enterococcus	CR6246	US8628776 SEQ ID NO: 463	7430
STPH202	Light chain, S. aureus, Enterococcus	CR6388	US8628776 SEQ ID NO: 465	7431
STPH203	Light chain, S. aureus, Enterococcus	CR6396	US8628776 SEQ ID NO: 469	7432
STPH204	Light chain, S. aureus, Enterococcus	CR6409	US8628776 SEQ ID NO: 473	7433
STPH205	Light chain, S. aureus, Enterococcus	CR6421	US8628776 SEQ ID NO: 477	7434
STPH206	Light **chain,** S. aureus, Enterococcus	CR6432	US8628776 SEQ ID NO: 481	7435
STPH207	Light chain, S. aureus, S. epidermidis, S. **caprae,** S. saprophyticus, S. capitis, or **methicillin-resistant** S. aureus (MRSA)	F1 antibody **variant**	US8617556 SEQ ID NO: 8	7436
STPH208	Light chain, S. aureus, S. epidermidis, S. caprae, S. saprophyticus, S. capitis, or **methicillin-resistant** S. aureus (MRSA)	F1 antibody variant	U586I7556 SEQ ID NO: **10**	7437
STPH209	Light chain, S. aureus, S. epidenmdis, S. caprae, S. saprophyticus, S. capitis, or methicillin-resistant S. aureus (MRSA)	F1 **antibody** variant	US8617556 SEQ ID NO: 11	7438
STPH210	Light **chain,** S. aureus, S. **epidermidis,** S. caprae, S. saprophyticus, S. capitis, or methicillin-resistant S. aureus (MRSA)	**rFl**	US86I7556 SEQ ID NO: 57	7439
STPH211	Light chain, S. aureus, S. epidenmdis, S. caprae, S. saprophyticus, S. capitis, or methicillin-resistant S. aureus (MRSA)	**rFl** V205C	US8617556 SEQ ID NO: 58	7440
STPH2 I2	Light chain, S. aureus, S. epidermidis, S. caprae, S. saprophyticus, S. capitis, or **methicillin-resistant** S. aureus (MRSA)	rF1	US86 I7556 SEQ ID NO: **64**	744 1
STPH213	**ScFv,** S. aureus and S. epidennidis	SC02-430	**US8460666** SEQ ID NO: 20	7442
STPH214	**ScFv,** S. aureus and S. epidermidis	SC05-132	US8460666 SEQ ID NO: 22	7443
STPH215	ScFv, S. aureus and S. epidermidis	**SC05-133**	**US8460666** SEQ ID NO: 24	7444
STPH2 I6	ScFv, S. aureus, **Enterococcus**	SC05-140	US8628776 SEQ ID NO: 351	7445
STPH217	ScFv, S. aureus, Enterococcus	SC05-157	US8628776 SEQ ID NO: 353	7446
STPH2 18	ScFv, S. aureus, **Enterococcus**	SC05-159	US8628776 SEQ ID NO: 62	7447

STPH219	ScFv, S. aureus, Enterococcus	SC05-166	US8628776 SEQ ID NO: 64	7448
STPH220	ScFv, S. aureus, Enterococcus	SC05-179	US8628776 SEQ ID NO: 355	7449
STPH221	ScFv, S. aureus, Enterococcus	SC05-187	US8628776 SEQ ID NO: 66	7450
STPH222	ScFv, S. aureus, Enterococcus	SC06-016	US8628776 SEQ ID NO: 68	7451
STPH223	ScFv, S. aureus, Enterococcus	SC.06-043	US8628776 SEQ ID NO: 70	7452
STPH224	ScFv, S. aureus, Enterococcus	SC06-049	US8628776 SEQ ID NO: 72	7453
STPH225	ScFv, S. aureus, Enterococcus	SC06-050	US8628776 SEQ ID NO: 357	7454
STPH226	ScFv, S. aureus, Enterococcus	SC06-071	US8628776 SEQ ID NO: 74	7455
STPH227	ScFv, S. aureus, Enterococcus	SC06--077	US8628776 SEQ ID NO: 359	7456
STPH228	ScFv, S. aureus, Enterococcus	SC06-078	US8628776 SEQ ID NO: 76	7457
STPH229	ScFv, S. aureus, Enterococcus	SC06-079	US8628776 SEQ ID NO: 361	7458
STPH230	ScFv, S. aureus, Enterococcus	SC06-086	US8628776 SEQ ID NO: 363	7459
STPH231	ScFv, S. aureus, Enterococcus	SC06-087	US8628776 SEQ ID NO: 207	7460
STPH232	ScFv, S. aureus, Enterococcus	SC06--089	US8628776 SEQ ID NO: 209	7461
STPH233	ScFv, S. aureus, Enterococcus	SC06-092	US862S776 SEQ ID NO: 365	7462
STPH234	ScFv, S. aureus, Enterococcus	SC06-191	US8628776 SEQ ID NO: 367	7463
STPH235	ScFv, S. aureus, Enterococcus	SC06-195	US8628776 SEQ ID NO: 369	7464
STPH236	ScFv, S. aureus, Enterococcus	SC06-198	US8628776 SEQ ID NO: 371	7465
STPH237	ScFv, S. aureus, Enterococcus	SC06-241	US8628776 SEQ ID NO: 78	7466
STPH238	ScFv, S. aureus, Enterococcus	SC.06-242	US8628776 SEQ ID NO: 373	7467
STPH239	ScFv, S. aureus, Enterococcus	SC06-246	US8628776 SEQ ID NO: 375	7468
STPH240	ScFv, S. aureus, Enterococcus	SC06-252	US8628776 SEQ ID NO: 80	7469
STPH241	ScFv, S. aureus, Enterococcus	SC06-388	US8628776 SEQ ID NO: 377	7470
STPH242	ScFv, S. aureus, Enterococcus	SC06-389	US8628776 SEQ ID NO: 379	7471
STPH243	ScFv, S. aureus, Enterococcus	SC06-396	US8628776 SEQ ID NO: 381	7472
STPH244	ScFv, S. aureus, Enterococcus	SC06-402	US8628776 SEQ ID NO: 383	7473
STPH245	ScFv, S. aureus, Enterococcus	SC06-409	US8628776 SEQ ID NO: 385	7474
STPH246	ScFv, S. aureus, Enterococcus	SC06-415	US8628776 SEQ ID NO: 387	7475
STPH247	ScFv, S. aureus, Enterococcus	SC06-421	US8628776 SEQ ID NO: 389	7476
STPH248	ScFv, S. aureus, Enterococcus	SC06-429	US8628776 SEQ ID NO: 391	7477
STPH249	ScFv, S. aureus, Enterococcus	SC06-432	US8628776 SEQ ID NO: 393	7478

[00343] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants thereof or encodes one or more polypeptides, fragments or variants thereof described in international Publication No. WO2000071585, WO201316275L, WO2015089502, WO2015088346 (e.g., SEQ ID NO: 17), US Pub No. US20030224000, US20080014202, US20140037650, US20140170134, US Patent No. US8460666, the contents of each of which are herein incorporated by reference in their entirety, against *Staphylococcus* infection.

[00344] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 35 against Clostridium Tetani.

Table 35. **Antibodies** against **Clostridium Tetani**

Antibody No.	**Description**	**Antibody** Name	Reference **Information**	SEQ **ID** NO
CTET1	Heavy chain partial		Sims, G.P. "Tetanus toxoid specific antibody heavy chain V-gene sequence", Unpublished, CNBI Accession * AAC69189.1	7479
CTET2	Heavy chain variable region	F5-20	Sims, G.P. "Tetanus toxoid specific antibody heavy chain V-gene sequence", Unpublished, CNBI Accession * AAB50736.1	7480
CTET3	Heavy chain variable region		Larrick, J.W., "Therapeutic human antibodies derived from PGR amplification of B-celi variable regions", Immunol. Rev. 130, 69-85 (1992), CNBI Accession # AAB25318.1	7481
CTET4	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36976.1	7482
CTET5	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36975.1	7483
CTET6	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36974.1	7484
CTET7	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36973.1	7485
CTET8	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36972.1	7486
CTET9	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36971.1	7487
CTET10	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL36970.1	7488
CTET11	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al, "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36969.1	7489
CTET12	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36968.1	7490
CTET13	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36967.1	7491
CTET14	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL36966.1	7492

CTET15	**Heavy cliain variable** region. **Human** immunoglobulin		de Kraif, J. et **al, "Human** immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous V(H) genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI Accession # **ACL36965. 1**	7493
CTET16	**Heavy** chain variable region, **Human** immunoglobulin		de **Kruif,** J. **et** al., "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI Accession # ACL36964. 1	7494
CTET17	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of** mgh-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI Accession # **ACL36963. 1**	7495
CTET18	Heavy chain variable region, Human **immunoglobulin**		de Kruif, J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession *ii* **ACL36962. 1**	7496
CTET19	Heavy **chain variable** region. **Human immunoglobulin**		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL3696 1.1	**7497**
CTET20	**Heavy cliain** variable region. Human immunoglobulin		de Kruif, J. et **al., "Human** immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI **Accession** # **ACL36960. 1**	7498
CTET21	**Heavy** chain variable region, Human **immunoglobulin**		de **Kruif,** J. et al, "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 **(3),** 548-558 (2009), CNBI Accession # ACL36958. !	7499
CTET22	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol. Biol.** 387 **(3),** 548-558 (2009), CNBI Accession # ACL36959. 1	7500
CTET23	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession *ii* **ACL36957. 1**	7501
CTET24	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 **(3),** 548-558 (2009), CNBI Accession # **ACL36956. 1**	7502
CTET25	**Heavy** chain variable region, Human immunoglobulin		de Kruif, J. et al, "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol. Biol.** 387 **(3),** 548-558 (2009), CNBI Accession # **ACL36955. 1**	7503
CTET26	Heavy chain variable region, Human **immunoglobulin**		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of** mgh-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI Accession # ACL36954. !	7504
CTET27	Heavy chain variable region, Human **immunoglobulin**		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid coniain **a** large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol. Biol.** 387 (3), 548-558 (2009), CNBI Accession * **ACL36953. 1**	7505
CTET28	Heavy chain variable region. **Human immunoglobulin**		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL36952. 1	7506
CTET29	**Heavy** cliain variable region. Human immunoglobulin		de Kruif, J. et **al., "Human** immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. **Mol. Biol.** 387 **(3),** 548-558 (2009), CNBI **Accession # ACL3695 1.1**	7507

CTET30	Heavy cliain variable region. Human immunoglobulin		de Kruif, J. et ai., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36950. 1	7508
CTET31	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36949. 1	7509
CTET32	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al, "Human immunoglobulin repertoires against tetanus toxoid coniain a large and diverse fraction of nigh-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36948. 1	7510
CTET33	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36947. 1	7511
CTET34	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36946. 1	7512
CTET35	Heavy cliain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36924. 1	7513
CTET36	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al, "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36925. 1	7514
CTET37	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid coniain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36926. 1	7515
CTET38	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36927. 1	7516
CTET39	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL36928. 1	7517
CTET40	Heavy chain variable region, Human inmiunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL36929. 1	7518
CTET41	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36930. 1	7519
CTET42	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL3693 1.1	7520
CTET43	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36932. 1	7521
CTET44	Heavy cliain variable region. Human immunoglobulin		de Kruif, J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession * ACL36933. 1	7522

CTET45	**Heavy cliain variable** region. **Human immunoglobulin**		de Kraif, J. et **al, "Human** immunoglobulin repertoires against tetanus toxoid contain a large **and** diverse fraction of high-affinity promiscuous V(H) genes", J. **Mol. Biol.** 387 **(3),** 548-558 (2009), CNBI **Accession** # **ACL36934. 1**	7523
CTET46	**Heavy** chain variable region, **Human immunoglobulin**		de Kraif, J. **et** al., "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. **Mol. Biol.** 387 (3), 548-558 (2009), CNBI **Accession** # **ACL36935. 1**	7524
CTET47	Heavy chain variable region, Human immunoglobulin		de Kraif, J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid **contain** a large and diverse fraction of mgh-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI Accession # **ACL36936. 1**	7525
CTET48	Heavy chain variable region, Human **immunoglobulin**		de **Kruif,** J. et **al.,** "Human immunoglobulin repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # **ACL36937.1**	7526
CTET49	Heavy **chain variable** region. **Human immunoglobulin**		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 (3), 548-558 (2009), CNBI Accession # ACL36938. 1	7527
CTET50	**Heavy cliain** variable region. Human immunoglobulin		de Kruif, J. et **al., "Human** immunoglobulin repertoires against tetanus toxoid contain a large **and** diverse fraction of high-affinity **promiscuous** V(H) genes", J. **Mol. Biol.** 387 **(3),** 548-558 (2009), CNBI **Accession** # **ACL36939. 1**	7528
CTET51	**Heavy** chain variable region, Human **immunoglobulin**		de Kraif, J. et al, "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction of **high-affinity** promiscuous V(H) genes", J. Mol. Biol. 387 **(3),** 548-558 (2009), CNBI Accession # ACL36940. 1	7529
CTET52	Heavy chain variable region, **Human** immunoglobulin		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous **V(H)** genes", J. **Mol. Biol.** 387 **(3),** 548-558 (2009), CNBI **Accession * ACL36941. 1**	7530
CTET53	Heavy chain variable region, Human immunoglobulin		de Kruif, J. et **al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. Mol. Biol. 387 (3), 548-558 (2009), CNBI Accession # **ACL36943. 1**	7531
CTET54	Heavy chain variable region. Human immunoglobulin		de Kruif, J. et al., "Human immunoglobulin repertoires against tetanus toxoid contain **a** large and diverse fraction of high-affinity promiscuous V(H) genes", J. Mol. Biol. 387 **(3),** 548-558 (2009), CNBI **Accession * ACL36942. 1**	7532
CTET55	**Heavy** chain variable region, Human **inmiunoglobulin**		de Kraif, J. **et al.,** "Human **immunoglobulin** repertoires against tetanus toxoid contain a large and diverse fraction **of** high-affinity promiscuous **V(H)** genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI **Accession * ACL36944. 1**	7533
CTET56	Heavy chain variable region, Human **immunoglobulin**		de Kraif, J. et **al.,** "Human **inmiunoglobulin** repertoires against tetanus toxoid **contain** a large and diverse fraction **of high-affinity** promiscuous **V(H)** genes", J. **Mol.** Biol. 387 **(3),** 548-558 (2009), CNBI Accession # ACL36945. 1	7534
CTET57	Light chain variable region		Larrick, J.W., "Therapeutic human antibodies derived from PGR amplification of B-cell **variable** regions", Immunol. Rev. **130,** 69-85 (1992), CNBI Accession # AAB25319.1	7535

[00345] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 36 **against Bordetelia** Pertussis **and/or Bordetelia** Parapertussis.

Table 36. **Antibodies against** Bordetella Pertisssss and Bordetella Parapertussis

Antibody No.	**Description**	Antibody Name	Reference information	SEQ ID NO
BORTl	Heavy chain	42. i 1.D4	WO20 14 160098 SEQ ID NO: 47	7536
BORT2	Heavy chain	42. 12.G2	WO20 l4 160098 SEQ ID NO: 5l	7537
BORT3	Heavy chain	42. L2.Ai2	WO20 14 160098 SEQ ID NO: 55	7538
BORT4	Heavy chain	42.12.A9	WO20 14 160098 SEQ ID NO: 59	7539
BORT5	Heavy chain	42. 18.E12	WO20 14 160098 SEQ ID NO: 63	7540
BORT6	Heavy chain	55. I2.A8	WO20 14 160098 SEQ ID NO: 67	7541
BORT7	Heavy chain	55.15.H5	WO20 l4 160098 SEQ ID NO: 7l	7542
BORT8	Heavy chain	55.17.D8	WO20 14 160098 SEQ ID NO: 75	7543
BORT9	Heavy chain	55.22.E7	WO20 14 160098 SEQ ID NO: 79	7544
BORT10	Light chain	42. 1 1.D4	WO20 14 160098 SEQ ID NO: 49	7545
BORTl 1	Light chain	42. 12.G2	WO20 14 160098 SEQ ID NO: 53	7546
BORTl 2	Light chain	42. 12.A12	WO20 l4 160098 SEQ ID NO: 57	7547
BORTl 3	Light chain	42. I2.A9	WO20 14 160098 SEQ ID NO: 6l	7548
BORTl 4	Light chain	42. 18.E12	WO20 14 160098 SEQ ID NO: 65	7549
BORTl 5	Light chain	55.12.A8	WO20 14 160098 SEQ ID NO: 69	7550
BORTl 6	Light chain	55. 15.H5	WO20 14 160098 SEQ ID NO: 73	7551
BORTl 7	Light chain	55.17.DS	WO20 l4 160098 SEQ ID NO: 77	7552
BORTl 8	Light chain	55.22.E7	WO20 14 160098 SEQ ID NO: 81	7553
BORTl 9	Single chain variable fragment antibody type 1 a1, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesins filamentous hemagglutinin and pertaciin" Infect. Immun. 75 (11), 5476-5482 (2007), NCBI Accession # ABB13478.1	7554
BORT20	Single chain variable fragment antibody type 18 a18, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesins filamentous hemagglutinin and pertactin" Infect. Immun 75 (11), 5476-5482 (2007), NCBI Accession # ABB 13483. 1	7555
BORT2 1	Single chain variable fragment antibody type 2 a2, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesins filamentous hemagglutinin and pertactin" Infect. Immun. 75 (11), 5476-5482 (2007), NCBI Accession # ABB 13479. 1	7556
BORT22	Single chain variable fragment antibody type 4 b4, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesins filamentous hemagglutinin and pertactin" Infect. Immun. 75 (11), 5476-5482 (2007), NCBI Accession # ABB13480.1	7557
BORT23	Single chain variable fragment antibody type 5 c5, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesins filamentous hemagglutinin and pertactin"	7558

			Infect. Immun. 75 (11), 5476-0482 (2007), NCBI Accession # ABB 13481. 1	
BORT24	Single chain variable fragment antibody type 6 d6, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesms filamentous hemagglutinin and pertactuT Infect. Immun. 75 (11), 5476-5482 (2007), NCBI Accession # ABB 13482. 1	7559
BORT25	Single chain variable fragment antibody type 7 e, single chain variable region		Hussein, A.H. et al. "Construction and characterization of single-chain variable fragment antibodies directed against the Bordetella pertussis surface adhesins filamentous hemagglutinin and pertactin" Infect. Immun. 75 (11), 5476-5482 (2007), NCBI Accession # ABB 13484. 1	7560

[00346] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 37 against Mycobacteria.

Table 37. Antibodies **against Mycobacteria**

Antibody No.	**Description**	**Antibody** Name	Reference Information	SEQ **ID** NO
MYCO1	Autod2 Single-chain variable fragment antibody, Tb antibody, anti-neutrophil cytoplasmic antibodies cross-react with mycobacterium avium subsp. Paratuberculosis antigens		Berger et al., Microbes Infect. 9 (8), 963-970 (2007), NCBI Accession # ABI8 1486. 1	7561
MYC02	autohl single-chain variable fragment antibody, Tb antibody, anti-neutrophil cytoplasmic antibodies cross-react with mycobacterium avium subsp. Paratuberculosis antigens,		Berger et al., Microbes Infect. 9 (8), 963-970 (2007), NCBI Accession # ABI8 1485. 1	7562
MYC03	Heavy chain constant region, Mycobacteria	moG2a/ moG2afull	US20130309237 SEQ ID NO: 10	7563
MYC04	Heavy chain constant region, Mycobacteria	hG1mG2a	US20130309237 SEQ ID NO: 11	7564
MYC05	Heavy chain constant region, Mycobacteria	hG3mG2a	US20130309237 SEQ ID NO: 12	7565
MYC06	Heavy chain constant region, Mycobacteria	hmG1&ll	US20130309237 SEQ ID NO: 13	7566
MYC07	Heavy chain constant region, Mycobacteria	huG3full	US20130309237 SEQ ID NO: 14	7567
MYC08	Heavy chain variable region, Mycobacteria	2FI2 IgGs	US20130309237 SEQ ID NO: 15	7568
MYC09	Heavy chain variable region, Mycobacteria	2F12 igGs	US20130309237 SEQ ID NO: 18	7569
MYCOIO	Heavy chain variable region, partial sequence, Tb antibody, mouse monoclonal mpt51	16al	Al-sayyed et al., Tuberculosis (Edinb) 87 (6), 489-497 (2007), NCBI Accession # ABS20005. 1	7570
MYCOl l	Light chain constant region, Mycobacteria	HsCK	US20130309237 SEQ ID NO: 16	7571
MYCO12	Light chain variable region, Mycobacteria	MoCK	US20130309237 SEQ ID NO: 17	7572
MYC013	Light chain variable region, partial sequence, Tb antibody, mouse monoclonal mpt51	16al	Al-sayyed et al., Tuberculosis (Edinb) 87 (6), 489-497 (2007), NCBI Accession # ABS20006. 1	7573

MYC014	Scfv, Tb antibody, an engineered single chain antibody		US20060229438 SEQ ID NO: 3	7574
MYC015	Scfv, Tb antibody, an engineered single chain antibody		US20060229438 SEQ ID NO: 4	7575
MYC016	Scfv, Tb antibody, an engineered single chain antibody		US20060229438 SEQ ID NO: 2	7576

[00347] In one embodiment, the pavioad region of the AAV particle composes one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 38 against Francisella Tularensis.

Table 38. Antibodies **against** Frasicisella Telaresisis

Antibody **No.**	Description	**Antibody** Name	**Reference** Information	SEQ ID NO
FRAN1	Chain H	Ab-52	Rynkiewicz, M.J. et al., "Structural Analysis of a Protective Epitope of the Francisella tularensis 0-Poiysaccliari.de", Biochemistry 51 (28), 5684-5694 (2012), NCBI Accession # 3UJT_H	7577
FRAN2	Chain H	N62	Lu, Z., et al., "The binding sites of monoclonal antibodies to the non-reducing end of Francisella tularensis O-antigen accommodate mainly the terminal saccharide", Immunology 140 (3), 374-389 (2013), NCBI Accession # 4KPH_H	7578
FRAN3	Chain H	Ab64	Lu, Z. et al., "B-cell epitopes in GroEL of Francisella tularensis", PLoS ONE 9 (6), E99847 (2014), NCBI Accession # 4PB9_H	7579
FRAN4	Chain H	Ab53	Lu, Z. et al., "B-cell epitopes in GroEL of Francisella tularensis", PLoS ONE 9 (6), E99847 (2014), NCBI Accession # 4PB0_H	7580
FRAN5	Chain H	N203	Lu, Z. et al., "Functional and Structural Characterization of Francisella tularensis O-Antigen Antibodies at the Low End of Antigen Reactivity", Monoclonal Antib Immunodiagn Immunother 33 (4), 235-245 (2014), NCBI Accession # 40TX_H	7581
FRAN6	Chain 1	Ab-52	Rynkiewicz, MJ. et al., "Structural Analysis of a Protective Epitope of the Francisella tularensis O-Poiysaccharide", Biochemistiy 51 (28), 5684-5694 (2012), NCBI Accession # 3UJT_I	7582
FRAN7	Chain I	N62	Lu, Z., et al, "The binding sites of monoclonal antibodies to the non-reducing end of Francisella tularensis O-antigen accommodate mainly the terminal saccharide", Immunologv 140 (3), 374-389 (2013), NCBI Accession # 4KPH_I	7583
FRAN8	Chain I	N203	Lu, Z. et al., "Functional and Structural Characterization of Francisella tularensis O-Antigen Antibodies at the Low End of Antigen Reactivity", Monoclonal Antib Immunodiagn Immunother 33 (4), 235-245 (2014), NCBI Accession # 40TX_I	7584
FRAN9	Chain L	Ab-52	Rynkiewicz, MJ. et al., "Structural Analysis of a Protective Epitope of the Francisella tularensis O-Poiysaccharide", Biochemistry 51 (28), 5684-5694 (2012), NCBI Accession # 3UJT_L	7585
FRAN10	Chain L	N62	Lu, Z., et al., "The binding sites of monoclonal antibodies to the non-reducing end of Francisella tularensis O-antigen accommodate mainly the terminal saccharide", Immunologv 140 (3), 374-389 (2013), NCBI Accession # 4KPH_L	7586
FRAN11	Chain L	Ab64	Lu, Z. et al., "B-cell epitopes in GroEL of Francisella tularensis", PLoS ONE 9 (6), E99847 (2014), NCBI Accession # 4PB9_L	7587

FRAN 12	Chain L	Ab53	Lu, Z. et al., "B-cell epitopes in GroEL of Francisella tularensis", PLoS ONE 9 (6), E99847 (2014), NCBI Accession # 4PB0_L	7588
FRAN 13	Chain L	N203	Lis, Z. et al., "Functional and Structural Characterization of Francisella tularensis O-Antigen Antibodies at the Low End of Antigen Reactivity", Monoclonal Antib Imrnunodiagn Immunother 33 (4), 235-245 (2014), NCBI Accession # 40TX_L	7589
FRAN 14	Chain M	Ab-52	Rynkiewicz, M.J. et al., "**Structural** Analysis of a Protective Epitope of the Francisella tularensis O-Polysaccharicle", Biochemistry 51 (28), 5684-5694 (2012), NCBI Accession # 3UJT_M	7590
FRAN 15	Chain M	N62	Lu, Z., et al, "The binding sites of monoclonal antibodies to the non-reducing end of Francisella tularensis O-antigen accommodate mainly the terminal saccharide", Immunology 140 (3), 374-389 (2013), NCBI Accession # 4KPH_M	7591
FRAN 16	Chain M	N203	Lu, Z. et al., "Functional and Structural Characterization of Francisella tularensis O-Antigen Antibodies at the Low End of Antigen Reactivity", Monoclonal Antib Imrnunodiagn Immunother 33 (4), 235-245 (2014), NCBI Accession # 40TX_M	7592

[00348] In one embodiment, the pavioad region of the AAV particle composes one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 39 aeainsi Bacteria.

Table 39. Antibodies against Bacteria

Antibody No.	**Description**	**Antibody Name**	Reference **Information**	SEQ ID NO
BACH	Heavy chain variable region, Enterococcus faecium, Enterococcus faecalis, Clostridium difficile		US20080038266 SEQ ID NO: 1	7593
BACI2	Heavy chain variable region, Neisseria meningitidis,	Naid60	US20060073 139 SEQ ID NO: 5	7594
BACI3	Heavy chain, Neisseria meningitidis,		Fernandez de Cossio, M.E., et al "Human monoclonal antibodies against an epitope on the class 5c outer membrane protein common to many pathogenic strains of Neisseria meningitidis", J. Infect. Dis. 166 (6), 1322-1328 (1992), AAB 18935	7595
BAC14	Heavy chain, Neisseria meningitidis.		Fernandez de Cossio, M.E., et al. "Human monoclonal antibodies against an epitope on the class 5c outer membrane protein common to many pathogenic strains of Neisseria meningitidis", J. Infect. Dis. 166 (6), 1322-1328 (1992), AAB 18934	7596
BACK	Heavy chain, Septic shock, meningococcal septic shock	Edobacomab, E5, XMMEN-0E5		7597
BACI6	Ig kappa chain V-I region WEA, Klebsiella bacteria		Goni, F. and Frangione, B., "Amino acid sequence of the Fv region of a human monoclonal IgM (protein WEA) with antibody activity against 3,4-pyruvyiated galactose in Klebsiella polysaccharides	7598

			K30 and **K33",** Proc. **Natl.** Acad. Sci. U.S.A. 80 **(15),** 4837-4841 (1983). P01610	
BACI7	**Ig** kappa **cliain** V-I region WEA, Klebsiella bacteria		Goni, F. and **Frangjone,** B., "Amino acid sequence of the Fv region of **a** human monoclonal **IgM** (protein WEA) with antibody activity against **3,4-pymvylated** galactose in Klebsiella polysaccharides K30 and K33", Proc. **Natl.** Acad. Sci. U.S.A. 80 (15), 4837-4841 (1983), **P01763**	7599
BACI8	Light chain variable region, Enterococcus **faecium,** Enterococcus faecalis, **Clostridium** difficile		US20080038266 SEQ ID NO: 16	7600
BACI9	Light clsain variable region, Neisseria meningitidis	Naid60	**US20060073 139** SEQ ID NO: 6	7601
BACIiO	Light chain, Septic shock, meningococcal septic shock,	**Edobacomab,** E5, XMMEN-0E5		7602
BACI11	scFv antibody, Anti-Burkholderia mallei		Zou, N., et **al.** "Human Single-Chain Fv Antibodies against Burkholderia mallei and Burkholderia **pseudomallei", unpublished,** NCBI Accession # **ABI97022.1**	7603
BACH 2	**scFv** antibody, **Anti-Burkholderia** mallei		Zou, N., et **al.** "Human Single-Chain Fv Antibodies against **Burkholderia** mallei and Burkholderia pseudomallei", unpublished, NCBI Accession # **AB197023.1**	7604
BACH 3	scFv antibody, **Anti-**Burkholderia mallei		Zou, N , et al. "Human Single-Chain Fv Antibodies against Burkholderia **mallei** and Burkholderia **pseudomallei",** unpublished, NCBI Accession # ABI97021.1	7605
BACI14	scFv antibody, Anti-**Burkholderia** mallei		Zou, N., et **al.** "Human Single-Chain Fv Antibodies against Burkholderia mallei and Burkholderia pseudomallei", unpublished, NCBI Accession # **ABI97018.1**	7606
BACH 5	scFv antibody, **Anti-**Burkholderia mallei		Zou, N., et **al.** "Human **Single-Chain** Fv Antibodies against Burkholderia mallei and Burkholderia **pseudomallei",** unpublished, NCBI Accession # AB19702**4.1**	7607
BACI16	scFv antibody, Anti-Burkholderia mallei		Zou, N., et **al.** "Human Single-Chain Fv Antibodies against Burkholderia mallei and Burkholderia pseudomallei", unpublished, NCBI Accession # ACZ65033.**1**	7608
BAC117	**scFv** antibody, Anti-Burkholderia mallei		Zou, N , et **al.** "**Human** Single-Chain Fv Antibodies against Burkholderia mallei and Burkholderia pseudomallei", unpublished, NCBI Accession # ACZ65032.**1**	7609
BACH 8	scFv antibody, **Anti-Burkholderia** mallei		Zou, N., et al. "Human Single-Chain Fv Antibodies against **Burkholderia** mallei and Burkholderia **pseudomallei",**	7610

			unpublished, NCBI Accession # ACZ65031.1	
BACI19	scFv antibody, Anti-Buikholderia mallei		Zou, N., et al. "Human Single-Chain Fv Antibodies against Burkliolderia mallei and Burkliolderia pseudomaller , unpublished, NCBI Accession # ACZ65030.1	7611
BACI20	scFv antibody, Anti-Burkholderia mallei		Zou, N., et al. "Human Single-Chain Fv Antibodies against Buikholderia mallei and Burkliolderia pseudomallei", unpublished, NCBI Accession # ACZ65029.1	7612
BACI21	scFv antibody, Anti-Burkholderia mallei		Zou, N., et al. "Human Single-Chain Fv Antibodies against Burkliolderia mallei and Burkliolderia pseudomallei", unpublished, NCBI Accession # ACZ65028.1	7613
BAC122	scFv antibody, Anti-Burkholderia mallei		Zou, N., et al. "Human Single-Chain Fv Antibodies against Buikholderia mallei and Burkhoideria pseudomallei", unpublished, NCBI Accession # AB197020.1	7614
BACI23	scFv antibody, Anti-Buikholderia mallei		Zou, N., et al. "Human Single-Chain Fv Antibodies against Burkliolderia mallei and Burkliolderia pseudomallei", unpublished, NCBI Accession # ABI97019.1	7615
BACI24	Single chain variable, Borrelia,	CB515	LaRocca, T.J., et al. "Bactericidal action of a complement-independent relapsing fever Borrelia resides in its variable region", J. Immunol. 180 (9), 6222-6228 (2008), NCBI Accession # ABV22509	7616

[00349] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants encoding Doxorubicin, fragments or variants thereof for treating a disease and/or disorder or preventing a disease and/or disorder. As a non-limiting example, the disease and/or disorder is bacterial infection.

[00350] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payload antibody polypeptides listed in Table 40 against Toxoplasma gondii.

Table 40. Asrtibodies **against** Toxoplasma gondii

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
TOXO1	4f11e12	Fab Heavy Chain Variable Region, Surface antigen 1 (SAG1)	Graille, M. et al., J. Mol. Biol. 354 (2), 447-458 (2005), NCBI Accession # 1YNT_D (218aa)	7617
TOXO2	4f11e12	Fab Light Chain Variable Region, Surface antigen 1 (SAG1)	Graille, M. et al., J. Mol. Biol. 354 (2), 447-458 (2005), NCBI Accession # 1YNT_C (213aa)	7618

[00351] In one embodiment, the payioad region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payioad antibody polypeptides listed in Table 41 against Candida Yeast.

Table **41.** Antibodies against Candida Yeast

Antibody No.	Description	Antibody Name	SEQ ID NO
CAND1	Efungumab	Candida	7619

[00352] In one embodiment, the payioad region of the AAV particle comprises one or more nucleic acid sequences encoding one or more of the payioad antibody polypeptides listed in Table 42.

Table 42. **laiV** Antibodies

Antibody No.	Description	Antibody Name	Reference Information	SEQ ID NO
HIV1	4e10 Antibody Germline Precursor 7 Heavy Chain Fv	4e10antibody	NCBI Accession # 4OB5_H (127aa)	7620
HIV2	4e10 Antibody Germline Precursor 7 Light Chain Fv	4e10antibody	NCBI Accession # 4OB5_L (114aa)	7621
HIV3	Antigen Binding Fragment Of Heavy Chain	Ch103	Liao et al., Co-evolution of a broadly neutralizing HIV-1 antibody and founder virus; Nature 496 (7446), 469-476 (2013), NCBI Accession # 4JAM_H (226aa)	7622
HIV4	Antigen Binding Fragment Of Light Chain	Ch103	Liao et al., Co-evolution of a broadly neutralizing HIV-1 antibody and founder virus; Nature 496 (7446), 469-476 (2013), NCBI Accession # 4JAM_L (209aa)	7623
HIV5	Fab Fragment, Heavy Chain	N12-i15	NCBI Accession # 3QEH_G (232aa)	7624
HIV6	Fab Fragment, Light Chain	N12-i15	NCBI Accession # 3QEH_H (218aa)	7625
HIV7	Fab Heavy Chain	35o22	Huang et al., Broad and potent HIV-1 neutralization by a human antibody that binds the gp41-gp120 interface; Nature 515 (7525), 138-142 (2014), NCBI Accession # 4TOY_H (243aa)	7626
HIV8	Fab Heavy Chain	8anc195	Scharf, L., et al., Cell Rep 7 (3), 785-795 (2014), NCBI Accession # 4P9H_H (244aa)	7627
HIV9	Fab Heavy Chain	B13	Chen L. et al., Science 326 (5956), 1123-1127 (2009), NCBI Accession # 3IDY_B (231aa)	7628
HIV10	Fab Heavy Chain	Ch58	Liao et al., vaccine Induction of Antibodies against a Structurally Heterogeneous Site of Immune Pressure within HIV-1 Envelope Protein Variable region 1 and 2; Immunity 38 (1), 176-186 (2013), NCBI Accession # 4HQQ_H (231aa)	7629
HIV11	Fab Heavy Chain	Ch59	Liao et al., vaccine Induction of Antibodies against a Structurally Heterogeneous Site of Immune Pressure within HIV-1 Envelope Protein Variable region 1 and 2; Immunity 38 (1), 176-186 (2013), NCBI Accession # 4HPY_H (225aa)	7630

HIV12	Fab Heavy Chain	E51	Huang C et al., Proc. Natl. Acad. Sci. U.S.A. 101 (9), 2706-2711 (2004), NCBI Accession # 1RZF_H (235aa)	7631
HIV13	Fab Heavy Chain	N26-i1 Fab	NCBI Accession # 4FZE_H(232aa)	7632
HIV14	Fab Heavy Chain	Pgt145	McLellan, J.S. et al., Structure of HIV-1 gp120 V1 V2 domain with broadly neutralizing antibody PG9; Nature 480 (7377), 336-343 (2011), NCBI Accession # 3U1S_H (267aa)	7633
HIV15	Fab Heavy Chain Of Human Anti-hiv-1 Env Antibody A32	A32 Fab	NCBI Accession # 3TNM_A (231aa)	7634
HIV16	Fab Heavy Chain Of Human Anti-hiv-1 Env Antibody C11	C11 Fab	NCBI Accession # 4FZ8_H (237aa)	7635
HIV17	Fab Light Chain	35o22	Huang et al., Broad and potent HIV-1 neutralization by a human antibody that binds the gp41-gp120 interface; Nature 515 (7525), 138-142 (2014), NCBI Accession # 4TOY_L (216aa)	7636
HIV18	Fab Light Chain	8anc195	Scharf, L., et al., Cell Rep 7 (3), 785-795 (2014), NCBI Accession # 4P9H_L (215aa)	7637
HIV19	Fab Light Chain	B13	Chen L. et al., Science 326 (5956), 1123-1127 (2009), NCBI Accession # 3IDY_C (215aa)	7638
HIV20	Fab Light Chain	Ch58	Liao et al., vaccine Induction of Antibodies against a Structurally Heterogeneous Site of Immune Pressure within HIV-1 Envelope Protein Variable region 1 and 2; Immunity 38 (1), 176-186 (2013), NCBI Accession # 4HQQ_L (216aa)	7639
HIV21	Fab Light Chain	Ch59	Liao et al., vaccine Induction of Antibodies against a Structurally Heterogeneous Site of Immune Pressure within HIV-1 Envelope Protein Variable region 1 and 2; Immunity 38 (1), 176-186 (2013), NCBI Accession # 4HPY_L (215aa)	7640
HIV22	Fab Light Chain	E51	Huang C et al., Proc. Natl. Acad. Sci. U.S.A. 101 (9), 2706-2711 (2004), NCBI Accession # 1RZF_L (213aa)	7641
HIV23	Fab Light Chain	Monoclonal Antibody Vrc03	Bartesaghi, A. et al., Perfusion structure of trimeric HIV-1 envelope glycoprotein determined by cryo-electron microscopy; Nat. Struct. Mol. Biol. 20 (12), 1352-1357 (2013), NCBI Accession # 4CC8_L (209aa)	7642
HIV24	Fab Light Chain	N26-i1 Fab	NCBI Accession # 4FZE_L (212aa)	7643
HIV25	Fab Light Chain	Pgt145	McLellan, J.S. et al., Structure of HIV-1 gp120 V1 V2 domain with broadly neutralizing antibody PG9; Nature 480 (7377), 336-343 (2011), NCBI Accession # 3U1S_L (239aa)	7644
HIV26	Fab Light Chain Of Human Anti-hiv-1 Env Antibody A32	A32 Fab	NCBI Accession # 3TNM_B (216aa)	7645
HIV27	Fab Light Chain Of Human Anti-hiv-1 Env Antibody C11	C11 Fab	NCBI Accession # 4FZ8_L (218aa)	7646
HIV28	Fab Region Of The Heavy Chain	Fab 2558	Gorny et al., PLoS ONE 6 (12), E27780 (2011), NCBI Accession # 3UJI_H (223aa)	7647
HIV29	Fab Region Of The Heavy Chain	Fab 4025	Gorny et al., PLoS ONE 6 (12), E27780 (2011), NCBI Accession # **3UJJ_H (230aa)**	7648

HIV30	Fab, Heavy Chain	3bnc60	Scheid, J.F., et al., Science 333 (6049), 1633-1637 (2011), NCBI Accession # 3RPI_A (229aa)	7649
HIV31	Fab, Heavy Chain	48d	Huang CC et al., Proc. Natl. Acad. Sci. U.S.A. 101 (9), 2706-2711 (2004), NCBI Accession # 1R27_H (219aa)	7650
HIV32	Fab, Heavy Chain	4e10Fab	Bird et al., Nat. Struct. Mol. Biol. (2014), NCBI Accession # 4NGH_H (228aa)	7651
HIV33	Fab, Heavy Chain	Ch58-ua	Nicely et al. Ebiomedicine 2 (2015), NCBI Accession # 4RIS_H (230aa)	7652
HIV34	Fab, Heavy chain	Mab 2909	Spurrier, B., et al., Structure 19 (5), 691-699 (2011), NCBI Accession # 3Q6F_J (233aa)	7653
HIV35	Fab, Heavy Chain	Monoclonal Antibody Vrc03	Bartesaghi, A. et al., Perfusion structure of trimeric HIV-1 envelope glycoprotein determined by cryo-electron microscopy; Nat. Struct. Mol. Biol. 20 (12), 1352-1357 (2013), NCBI Accession # 4CC8_I (233aa)	7654
HIV36	Fab, light chain	3bnc60	Scheid, J.F., et al., Science 333 (6049), 1633-1637 (2011), NCBI Accession # 3RPI_B (206aa)	7655
HIV37	Fab, Light Chain	48d	Huang CC et al., Proc. Natl. Acad. Sci. U.S.A. 101 (9), 2706-2711 (2004), NCBI Accession # 1RZ7_L (212aa)	7656
HIV38	Fab, Light Chain	4e10Fab	Bird et al., Nat. Struct. Mol. Biol. (2014), NCBI Accession # 4NGH_L (215aa)	7657
HIV39	Fab, Light Chain	Ch58-ua	Nicely et al. Ebiomedicine 2 (2015), NCBI Accession # 4RIS_L (216aa)	7658
HIV40	Fab, light Chain	Mab 2909	Spurrier, B., et al., Structure 19 (5), 691-699 (2011), NCBI Accession # 3Q6F_K (213aa)	7659
HIV41	Gamma heavy chain	1443_C16	US9051362 SEQ ID NO: 12	7660
HIV42	Gamma heavy chain	1471 M23	US9051362 SEQ ID NO: 139	7661
HIV43	Gamma heavy chain	1489_I13	US9051362 SEQ ID NO: 59	7662
HIV44	Gamma heavy chain	1503_H05	US9051362 SEQ ID NO: 53	7663
HIV45	Gamma heavy chain variable region	1456_A12	US9051362 SEQ ID NO: 48	7664
HIV46	Gamma heavy chain variable region	1456_P20	US9051362 SEQ ID NO: 33	7665
HIV47	Gamma heavy chain variable region	1460_G14	US9051362 SEQ ID NO: 35	7666
HIV48	Gamma heavy chain variable region	1470 M23	US9051362 SEQ ID NO: 140	7667
HIV49	Gamma heavy chain variable region	1480_I08	US9051362 SEQ ID NO: 31	7668
HIV50	Gamma heavy chain variable region	1480_I08	US9051362 SEQ ID NO: 65	7669
HIV51	Gamma heavy chain variable region	1489_I13	US9051362 SEQ ID NO: 60	7670
HIV52	Gamma heavy chain variable region	1495_C14	US9051362 SEQ ID NO: 37	7671
HIV53	Gamma heavy chain variable region	1503_H05	US9051362 SEQ ID NO: 54	7672
HIV54	Gamma heavy chain variable region	1496_C09	US9051362 SEQ ID NO: 39	7673
HIV55	Gamma heavy chain	1456_A12	US9051362 SEQ ID NO: 47	7674
HIV56	Gamma heavy chain	1460_G14	US9051362 SEQ ID NO: 20	7675
HIV57	Gamma heavy chain	1495_C14	US9051362 SEQ ID NO: 24	7676
HIV58	Gamma heavy chain	1496_C09	US9051362 SEQ ID NO: 28	7677

HEV59	Gamma heavy	**1456JP20**	US905 1362 SEQ **ID NO:** 16	7678
HIV60	Gp41-specific antibody, heavy chain		US20140348785 SEQ **ID** NO: 11	7679
HIV61	**Gp41-specific** antibody, heavy **chain consensus**		US20140348785 SEQ **ID** NO: 146	7680
HIV62	Gp41-specific **antibody,** heavy chain consensus variable region		**US20140348785** SEQ **ID** NO: 187	7681
HIV63	Gp41-specific antibody, **heavy chain** consensus variable **region**		US20140348785 SEQ **ID** NO: 188	7682
HTV64	**Gp41-specific** antibody, heavy chain variable region		**US20 140348785** SEQ **ID** NO: 153	7683
HEV65	**Gp41-specific** antibody, heavy **chain** variable region		**US20140348785** SEQ **ID** NO: 154	7684
HIV66	Gp41-specific antibody, heavy **chain** variable region		US20140348785 SEQ **ID** NO: 155	7685
HIV67	Gp41-specific **antibody,** heavy chain variable region		**US20140348785** SEQ **ID** NO: 156	7686
HIV68	Gp41-specific antibody, heavy chain variable region		US20140348785 SEQ **ID** NO: 157	7687
HIV69	Gp41-specific antibody, **heavy** chain variable region		US20140348785 SEQ **ID** NO: 158	7688
HIV70	**Gp41-specific** antibody, heavy chairs variable region		**US20140348785** SEQ **ID** NO: 159	7689
HTV7 i	**Gp41-specific** antibody, heavy chain variable region		**US20 140348785** SEQ **ID** NO: 160	7690
HEV72	**Gp41-specific** antibody, heavy **chain** variable region		**US20140348785** SEQ **ID** NO: 161	7691
HIV73	Gp41-specific **antibody,** heavy chain variable region		**US20140348785** SEQ **ID** NO: 162	7692
HIV74	Gp41-specific antibody, heavy chain variable region		US20140348785 SEQ ID **NO:** 163	7693
HIV75	Gp41-specific antibody, heavy chain variable region		US20140348785 SEQ **ID** NO: 189	7694
HIV76	Gp41-specific antibody, **heavy** chain variable region		US20140348785 SEQ **ID** NO: 190	7695
HIV77	**Gp41-specific** antibody, heavy chain variable region		**US20140348785** SEQ **ID** NO: **191**	7696
HEV78	**Gp41-specific** antibody, heavy **chain** variable region		**US20140348785** SEQ **ID** NO: 192	7697

HEV79	**Gp41-specific** antibody, **heavy chain** variable region		**US20140348785** SEQ ID NO: 200	7698
HIV80	Gp41-specific **antibody,** heavy chain variable region		**US20140348785** SEQ **ID** NO: 201	7699
HIV81	Gp41-specific **antibody,** heavy chain variable region		**US20140348785** SEQ **ID NO:** 202	7700
HIV82	Gp41-specific antibody, heavy chain variable region		**US20140348785** SEQ ID NO: 203	7701
HIV83	Gp41-specific antibody, **heavy** chain variable region		US20140348785 SEQ **ID** NO: 204	7702
HIV84	**Gp41-specific** antibody, heavy chain variable **region**		**US20140348785** SEQ **ID** NO: 205	7703
HTV85	**Gp41-specific** antibody, heavy chain variable region		**US20 140348785** SEQ **ID** NO: 206	7704
HIV86	Gp41-specific antibody, heavy **chain** variable region		US20140348785 SEQ **ID** NO: 207	7705
HIV87	Gp41-specific antibody, heavy chain variable region		**US20140348785** SEQ **ID** NO: 208	7706
HIV88	Gp41-specific antibody, heavy chain variable region		**US20140348785** SEQ **ID NO:** 209	7707
HIV89	Gp41-specific antibody, light chain		**US20140348785** SEQ ID NO: **12**	7708
HEV90	**Gp41-specific** antibody, light chain variable region		**US20140348785** SEQ ID NO: 164	7709
HIV91	Gp41-specific antibody, light chain variable region		**US20140348785** SEQ **ID** NO: 165	7710
HIV92	Gp41-specific antibody, light chain variable region		**US20140348785** SEQ **ID NO:** 166	7711
HIV93	Gp41-specific antibody, light chain variable region		**US20140348785** SEQ ID NO: **167**	7712
HIV94	Gp41-specific antibody, light **chain** variable region		US20140348785 SEQ **ID** NO: 168	7713
HIV95	**Gp41-specific** antibody, **light chain** variable region		**US20140348785** SEQ **ID** NO: 169	7714
HTV96	**Gp41-specific** antibody, light chain variable region		**US20 140348785** SEQ **ID** NO: **170**	7715
HIV97	Gp41-specific antibody, light chain variable region		US20140348785 SEQ **ID** NO: 171	7716
HIV98	Gp41-specific antibody, light chain variable region		**US20140348785** SEQ **ID** NO: 172	7717

HEV99	**Gp41-specific** antibody, light chain variable region		US**20140348785** SEQ ID NO: 173	7718
HIV100	Gp41-specific **antibody,** light chain variable region		US20140348785 SEQ ID NO: 174	7719
HIV101	Gp41-specific antibody, light chain variable region		US20140348785 SEQ **ID NO:** 175	7720
HIV102	**Gp41**-specific antibody, light chain variable region		US20140348785 SEQ ID NO: 176	7721
HIV103	Gp41-specific antibody, light **chain** variable region		US20140348785 SEQ ID NO: 177	7722
HIV104	**Gp41-specific** antibody, **light chain** variable **region**		**US20140348785** SEQ ID NO: 178	7723
HTV105	**Gp41-specific** antibody, **light** chain variable region		**US20 140348785** SEQ **ID** NO: 179	7724
HIV106	Gp41-specific antibody, light chain variable region		US20140348785 SEQ ID NO: 180	7725
HIV107	**Gp41-specific** antibody, light chain variable region		**US20140348785** SEQ ID NO: **18 1**	7726
HIV108	Gp41-specific antibody, light chain variable region		US20140348785 SEQ **ID NO:** 182	7727
HIV109	**Gp41**-specific antibody, light chain variable region		**US20140348785** SEQ **ID** NO: **183**	7728
HIV1 10	Gp41-specific antibody, **light chain** variable region		US20140348785 SEQ ID NO: 184	7729
HIV111	**Gp41-specific** antibody, **light** chain variable region		**US20140348785** SEQ ID NO: 185	7730
HEV112	**Gp41-specific** antibody, light chain variable region		US**20140348785** SEQ ID NO: 186	773 1
HIV1 13	Gp41-specific antibody, light chain variable region		US20140348785 SEQ ID NO: 197	7732
HIV 114	**Gp41-specific** antibody, light chain variable region		**US20140348785** SEQ ID NO: 198	7733
HIV115	Gp41-specific antibody, light chain variable region		US20140348785 SEQ **ID NO:** 199	7734
HIVI16	Heavy chain	**Vrc06b**	Wu, X., et **al.,** Maturation and Diversity of the **VRCOI**-Antibody Lineage over 15 Years of Chronic HIV-1 Infection; Cell 161 (3), 470-485 (2015), **NCBI** Accession # **4XNZ_E (234aa)**	7735
HIV117	Heavy Chain	2424	**Kumar, R.,** et al, Functional and **Structural Characterization of** Human V3-Specific **Monoclonal** Antibody 2424 with Neutralizing Activity- against HIV-1 JRFL; J. Virol. 89 (17),	7736

			9090-9102 (2015), NCBI Accession # 4XML_H (223aa)	
HIV118	Heavy chain	5827	US20140205607 Table S13	7737
HIV119	Heavy chain	7863	US20140205607 Table S13	7738
HIV120	Heavy Chain	8062	Gustchina, E., PLoS ONE 8 (11), E78187 (2013), NCBI Accession # 4KHX_H(245aa)	7739
HIV121	Heavy chain	18761	US20140205607 Table S13	7740
HIV122	Heavy chain	19891	US20140205607 Table S13	7741
HIV123	Heavy chain	22425	US20140205607 Table S13	7742
HIV124	Heavy chain	28241	US20140205607 Table S13	7743
HIV125	Heavy chain	61272	US20140205607 Table S13	7744
HIV126	Heavy chain	61822	US20140205607 Table S13	7745
HIV127	Heavy chain	65030	US20140205607 Table S13	7746
HIV128	Heavy chain	70085	US20140205607 Table S13	7747
HIV129	Heavy chain	70542	US20140205607 Table S13	7748
HIV130	Heavy chain	80585	US20140205607 Table S13	7749
HIV131	Heavy chain	87722	US20140205607 Table S13	7750
HIV132	Heavy chain	96362	US20140205607 Table S13	7751
HIV133	Heavy chain	103787	US20140205607 Table S13	7752
HIV134	Heavy chain	146940	US20140205607 Table S13	7753
HIV135	Heavy chain	153849	US20140205607 Table S13	7754
HIV136	Heavy chain	1.00E+09	US20140348785 SEQ ID NO: 1	7755
HIV137	Heavy chain	104625_2	US20140205607 Table S14	7756
HIV138	Heavy chain	105239_4	US20140205607 Table S14	7757
HIV139	Heavy chain	10731_1	US20140205607 Table S14	7758
HIV140	Heavy Chain	10e8 (monoclonal)	Huang J et al., Nature 491 (7424), 406-412 (2012), NCBI Accession # 4G6F_B (236aa)	7759
HIV141	Heavy chain	120119_4	US20140205607 Table S14	7760
HIV142	Heavy chain	121325_4	US20140205607 Table S14	7761
HIV143	Heavy chain	12467_3	US20140205607 Table S14	7762
HIV144	Heavy chain	124918_2	US20140205607 Table S14	7763
HIV145	Heavy chain	127586_4	US20140205607 Table S14	7764
HIV146	Heavy chain	12A10HC	US20140328862 SEQ ID NO: 147	7765
HIV147	Heavy chain	12A12HC	US20140328862 SEQ ID NO: 148	7766
HIV148	Heavy chain	12A13HC	US20140328862 SEQ ID NO: 149	7767
HIV149	Heavy chain	12A16HC	US20140328862 SEQ ID NO: 150	7768
HIV150	Heavy chain	12A17HC	US20140328862 SEQ ID NO: 151	7769
HIV151	Heavy chain	12A1HC	US20140328862 SEQ ID NO: 152	7770
HIV152	Heavy chain	12A20HC	US20140328862 SEQ ID NO: 153	7771
HIV153	Heavy Chain	12a21	NCBI Accession # 4JPW_H (225aa)	7772
HIV154	Heavy chain	12A21HC	US20140328862 SEQ ID NO: 154	7773
HIV155	Heavy chain	12A22HC	US20140328862 SEQ ID NO: 155	7774
HIV156	Heavy chain	12A23HC	US20140328862 SEQ ID NO: 156	7775
HIV157	Heavy chain	12A27HC	US20140328862 SEQ ID NO: 157	7776
HIV158	Heavy chain	12A2HC	US20140328862 SEQ ID NO: 158	7777
HIV159	Heavy chain	12A30HC	US20140328862 SEQ ID NO: 159	7778

HEV160	Heavy chain	12A37HC	US20140328862 SEQ ID NO: 160	7779
HIV161	Heavy chain	12A46HC	US20140328862 SEQ ID NO: 161	7780
HIV 162	Heavy chain	12A4HC	US20140328862 SEQ ID NO: 162	7781
HIV163	Heavy chains	12A55HC	US20140328862 SEQ ID NO: 163	7782
HIV164	Heavy chain	12A56HC	US20140328862 SEQ ID NO: 164	7783
HEV165	Heavy chain	12A6HC	US20140328862 SEQ ID NO: 165	7784
HIV166	Heavy chain	12A7HC	US20140328862 SEQ ID NO: 166	7785
HIV 167	Heavy chain	12A9HC	US20140328862 SEQ ID NO: 167	7786
HIV168	Heavy chain	132797_4	US20140205607 Table S14	7787
HIV169	Heavy chain	135083_3	US20140205607 Table S14	7788
HEV170	Heavy chain	13826_2	US20140205607 Table S14	7789
HIV171	Heavy chain	143251_3	US20140205607 Table S14	7790
HIV172	Heavy chain	149590_4	US20140205607 Table S14	7791
HIV173	Heavy chain	149768_4	US20140205607 Table S14	7792
HIV174	Heavy chain	151901_4	US20140205607 Table S14	7793
HEV175	Heavy chain	156858_3	US20140205607 Table S14	7794
HIV 176	Heavy chain	164202-3	US20140205607 Table S14	7795
HIV177	Heavy chain	164922_3	US20140205607 Table S14	7796
HIV178	Heavy chain	165478_2	US20140205607 Table S14	7797
HIV179	Heavy chain	166726_3	US20140205607 Table S14	7798
HEV180	Heavy chain	167612_4	US20140205607 Table S14	7799
HIV 181	Heavy chain	168509_2	US20140205607 Table S14	7800
HIV182	Heavy chain	169094_4	US20140205607 Table S14	7801
HIV183	Heavy chain	17720_4	US20140205607 Table S14	7802
HIV184	Heavy chain	178037_3	US20140205607 Table S14	7803
HIV185	Heavy chains	179400_4	US20140205607 Table S14	7804
HIV 186	Heavy chain	179500_4	US20140205607 Table S14	7805
HIV187	Heavy chain	179888_3	US20140205607 Table S14	7806
HIV188	Heavy Chain	17b	Kwong, P.D., et al, structure of an HIV gp120 envelope glycoprotein in complex with the CD4 receptor and a neutralizing human antibody; Nature 393 (6686), 648-659 (1998), NCBI Accession # 1G9M_H (229aa)	7807
HIV189	Heavy chain	18278_1	US20140205607 Table S14	7808
HIV190	Heavy chain	184939_4	US20140205607 Table S14	7809
HIV 191	Heavy chain	185961_4	US20140205607 Table S14	7810
HIV192	Heavy chains	186066_4	US20140205607 Table S14	7811
HIV 193	Heavy chain	186275_2	US20140205607 Table S14	7812
HIV194	Heavy chain	186640_2	US20140205607 Table S14	7813
HIV195	Heavy chain	190244_4	US20140205607 Table S14	7814
HIV196	Heavy chain	193526_4	US20140205607 Table S14	7815
HIV197	Heavy chains	193896_4	US20140205607 Table S14	7816
HIV 198	Heavy chain	195462_4	US20140205607 Table S14	7817
HIV199	Heavy chain	196147_4	US20140205607 Table S14	7818
HIV200	Heavy chain	196283_4	US20140205607 Table S14	7819

HIV201	Heavy Chain	1b2530	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YFL_H (227aa)	7820
HIV202	Heavy chain	1F7	US6057421A Fig 8	7821
HIV203	Heavy chain	1NC9	WO2012154312 SEQ ID NO: 2471	7822
HIV204	Heavy Chain	2.2C	Acharya, P., et al., Structural Definition of an Antibody-Dependent Cellular Cytotoxicity Response Implicated in Reduced Risk for HIV-1 Infection; J. Virol. 88 (21), 12895-12906 (2014), NCBI Accession # 4R4N_H (220aa)	7823
HIV205	Heavy chain	24972_4	US20140205607 Table S14	7824
HIV206	Heavy chain	28936_1	US20140205607 Table S14	7825
HIV207	Heavy chain	2F5	US8637036B2 SEQ ID NO: 5	7826
HIV208	Heavy chain	2F5 F100BW	US8637036B2 SEQ ID NO: 7	7827
HIV209	Heavy chain	2F5 L100AW	US8637036B2 SEQ ID NO: 6	7828
HIV210	Heavy chain	2F5 L100AW-V100DW	US8637036B2 SEQ ID NO: 9	7829
HIV211	Heavy chain	2F5 V100DW	US8637036B2 SEQ ID NO: 8	7830
HIV212	Heavy chain	30263_2	US20140205607 Table S14	7831
HIV213	Heavy chain	3040HC	WO2015117008 SEQ ID NO: 14	7832
HIV214	Heavy chain	3044HC	WO2015117008 SEQ ID NO: 17	7833
HIV215	Heavy chain	31458_3	US20140205607 Table S14	7834
HIV216	Heavy chain	3430HC	WO2015117008 SEQ ID NO: 15	7835
HIV217	Heavy chain	3484HC	WO2015117008 SEQ ID NO: 16	7836
HIV218	Heavy chain	3630HC	WO2015117008 SEQ ID NO: 18	7837
HIV219	Heavy chain	3A124HC	US20140328862 SEQ ID NO: 261	7838
HIV220	Heavy chain	3A125HC	US20140328862 SEQ ID NO: 262	7839
HIV221	Heavy chain	3A140HC	US20140328862 SEQ ID NO: 263	7840
HIV222	Heavy chain	3A144HC	US20140328862 SEQ ID NO: 264	7841
HIV223	Heavy chain	3A160HC	US20140328862 SEQ ID NO: 265	7842
HIV224	Heavy chain	3A18HC	US20140328862 SEQ ID NO: 266	7843
HIV225	Heavy chain	3A204HC	US20140328862 SEQ ID NO: 267	7844
HIV226	Heavy chain	3A228HC	US20140328862 SEQ ID NO: 268	7845
HIV227	Heavy chain	3A233HC	US20140328862 SEQ ID NO: 269	7846
HIV228	Heavy chain	3A244HC	US20140328862 SEQ ID NO: 270	7847
HIV229	Heavy chain	3A255HC	US20140328862 SEQ ID NO: 271	7848
HIV230	Heavy chain	3A296HC	US20140328862 SEQ ID NO: 272	7849
HIV231	Heavy chain	3A334HC	US20140328862 SEQ ID NO: 273	7850
HIV232	Heavy chain	3A366HC	US20140328862 SEQ ID NO: 274	7851
HIV233	Heavy chain	3A381HC	US20140328862 SEQ ID NO: 275	7852
HIV234	Heavy chain	3A384HC	US20140328862 SEQ ID NO: 276	7853
HIV235	Heavy chain	3A419HC	US20140328862 SEQ ID NO: 277	7854
HIV236	Heavy chain	3a426hc	US20140328862 SEQ ID NO: 343	7855
HIV237	Heavy chain	3A461HC	US20140328862 SEQ ID NO: 278	7856

HEV238	Hc;n′ş chain	3A474HC	US20 140328862 SEQ İD NO: 279	7857
HIV239	Heavy chain	3a5 15hc	US20 140328862 SEQ ID NO: 344	7858
HIV240	Heavy chain	3A5 18HC	US20 140328862 SEQ ID NO: 280	7859
HİV24 1	Heavy chairs	3A539HC	US20 [40328862 SEQ ID NO: 281	7860
HIV242	Heavy chain	3A576HC	US20 140328862 SEQ ID NO: 282	7861
HEV243	Heavy chain	3A6 13HC	US20 140328862 SEQ İD NO: 283	7862
HIV244	Heavy chain	3A64HC	US20 140328862 SEQ ID NO: 284	7863
HIV245	Heavy chain	3A650HC	US20 140328862 SEQ ID NO: 285	7864
HIV246	Heavy chain	3A67HC	US20 140328862 SEQ ID NO: 286	7865
HIV247	Heavy chain	3A779HC	US20 140328862 SEQ ID NO: 287	7866
HEV248	Heavy chain	3A8 16HC	US20 140328862 SEQ İD NO: 288	7867
HIV249	Heavy chain	3A869HC	US20 140328862 SEQ ID NO: 289	7868
HIV250	Heavy chain	3A93HC	US20 140328862 SEQ ID NO: 290	7869
HIV25 1	Heavy chain	3A966HC	US20 140328862 SEQ ID NO: 291	7870
HIV252	Heavy chain	3A978HC	US20 140328862 SEQ ID NO: 292	7871
HEV253	Heavy chain	3ANC32HC	US20 140328862 SEQ İD NO: 346	7872
HIV254	Heavy chain	3ANC3HC	US20 140328862 SEQ ID NO: 293	7873
HIV255	Heavy chain	3ANC3HC	US20 140328862 SEQ ID NO: 347	7874
HIV256	Heavy chain	3ANC41HC	US20 140328862 SEQ ID NO: 348	7875
HIV257	Heavy chain	3ANC42HC	US20 140328862 SEQ ID NO: 294	7876
HEV258	Heavy chain	3ANC42HC	US20 140328862 SEQ İD NO: 349	7877
HIV259	Heavy chain	3ANC66HC	US20 140328862 SEQ ID NO: 295	7878
HIV260	Heavy chain	3ANC66HC	US20 140328862 SEQ ID NO: 350	7879
HIV261	Heavy chain	3ANC70HC	US20 140328862 SEQ ID NO: 35 1	7880
HIV262	Heavy chain	3ANC75HC	US20 140328862 SEQ ID NO: 352	7881
HTV263	Heavy chairs	3ANC79HC	US20 140328862 SEQ ID NO: 296	7882
HIV264	Heavy chain	3ANC79HC	US20 140328862 SEQ ID NO: 353	7883
HIV265	Heavy chain	3ANC87HC	US20 140328862 SEQ ID NO: 354	7884
HIV266	Heavy chain	3ANC8HC	US20 140328862 SEQ ID NO: 355	7885
HIV267	Heavy chain	3ANC96HC	US20 140328862 SEQ ID NO: 356	7886
HTV268	Heavy chairs	3B106HC	US20 [40328862 SEQ ID NO: 357	7887
HIV269	Heavy chain	3B10HC	US20 140328862 SEQ ID NO: 297	7888
HIV270	Heavy chain	3B120HC	US20 140328862 SEQ ID NO: 298	7889
HIV271	Heavy chain	3B126HC	US20 140328862 SEQ ID NO: 299	7890
HIV272	Heavy chain	3B129HC	US20 140328862 SEQ ID NO: 300	7891
HTV273	Heavy chairs	3B142HC	US20 [40328862 SEQ ID NO: 301	7892
HIV274	Heavy chain	3B1 54HC	US20 140328862 SEQ ID NO: 302	7893
HEV275	Heavy chain	3B165HC	US20 140328862 SEQ İD NO: 303	7894
HIV276	Heavy chain	3B16HC	US20 140328862 SEQ ID NO: 358	7895
HIV277	Heavy chain	3B171HC	US20 140328862 SEQ ID NO: 304	7896
HTV278	Heavy chairs	3B17HC	US20 140328862 SEQ ID NO: 305	7897
HIV279	Heavy chain	3B180HC	US20 140328862 SEQ ID NO: 359	7898
HEV280	Heavy chain	3B183HC	US20 140328862 SEQ İD NO: 360	7899
HIV281	Heavy chain	3B186HC	US20 140328862 SEQ ID NO: 306	7900
HIV282	Heavy chain	3B191HC	US20 140328862 SEQ ID NO: 361	7901

HEV283	Heavy chain	3B193HC	US20140328862 SEQ ID NO: 307	7902
HIV284	Heavy chain	3B21HC	US20 140328862 SEQ ID NO: 362	7903
HIV285	Heavy chain	3B22HC	US20 140328862 SEQ ID NO: 308	7904
HTV286	Heavy chairs	3B27HC	US20 140328862 SEQ ID NO: 309	7905
HIV287	Heavy chain	3B29HC	US20 140328862 SEQ ID NO: 310	7906
HEV288	Heavy chain	3B2HC	US20140328862 SEQ ID NO: 311	7907
HIV289	Heavy chain	3B31HC	US20 140328862 SEQ ID NO: 312	7908
HIV290	Heavy chain	3B33HC	US20 140328862 SEQ ID NO: 313	7909
HIV291	Heavy chain	3B40HC	US20140328862 SEQ ID NO: 314	7910
HIV292	Heavy chain	3B41HC	US20 140328862 SEQ ID NO: 315	7911
HEV293	Heavy chain	3B44HC	US20140328862 SEQ ID NO: 316	7912
HIV294	Heavy chain	3B45HC	US20 140328862 SEQ ID NO: 317	7913
HIV295	Heavy chain	3b46HC	US20 140328862 SEQ ID NO: 345	7914
HIV296	Heavy chain	3B48HC	US20140328862 SEQ ID NO: 318	7915
HIV297	Heavy chain	3B50HC	US20 140328862 SEQ ID NO: 319	7916
HEV298	Heavy chain	3B51HC	US20140328862 SEQ ID NO: 320	7917
HIV299	Heavy chain	3B56HC	US20 140328862 SEQ ID NO: 321	7918
HIV300	Heavy chain	3B57HC	US20 140328862 SEQ ID NO: 322	7919
HIV301	Heavy chain	3B5HC	US20140328862 SEQ ID NO: 323	7920
HIV302	Heavy chain	3B61HC	US20 140328862 SEQ ID NO: 324	7921
HEV303	Heavy chain	3B6HC	US20140328862 SEQ ID NO: 325	7922
HIV304	Heavy chain	3B77HC	US20140328862 SEQ ID NO: 326	7923
HIV305	Heavy chain	3B79HC	US20 140328862 SEQ ID NO: 327	7924
HIV306	Heavy chain	3B84HC	US20140328862 SEQ ID NO: 328	7925
HIV307	Heavy chain	3B86HC	US20 140328862 SEQ ID NO: 329	7926
HTV308	Heavy chairs	3B8HC	US20 [40328862 SEQ ID NO: 330	7927
HIV309	Heavy chain	3B93HC	US20 140328862 SEQ ID NO: 331	7928
HIV310	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 363	7929
HIV311	Heavy chain	3BBM60	US20140328862 SEQ ID NO: 364	7930
HIV312	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 365	7931
HTV313	Heavy chairs	3BBM60	US20 140328862 SEQ ID NO: 366	7932
HIV314	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 367	7933
HIV315	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 368	7934
HIV316	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 369	7935
HIV317	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 370	7936
HTV318	Heavy chairs	3BBM60	US20 140328862 SEQ ID NO: 371	7937
HIV319	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 372	7938
HEV320	Heavy chain	3BBM60	US20140328862 SEQ ID NO: 373	7939
HIV321	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 374	7940
HIV322	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 375	7941
HTV323	Heavy chairs	3BBM60	US20 140328862 SEQ ID NO: 376	7942
HIV324	Heavy chain	3BBM60	US20 140328862 SEQ ID NO: 377	7943
HEV325	Heavy chain	3BNC10 1HC	US20140328862 SEQ ID NO: 332	7944
HIV326	Heavy chain	3BNC101HC	US20 140328862 SEQ ID NO: 378	7945
HIV327	Heavy chain	3BNC102HC	US20 140328862 SEQ ID NO: 379	7946

HEV328	Heavy chain	**3BNC104HC**	**US20140328862** SEQ ID NO: 380	7947
HIV329	Heavy chain	**3BNC105HC**	**US20 140328862** SEQ ID NO: 381	7948
HIV330	**Heavy chain**	**3BNC106HC**	**US20 140328862** SEQ ID NO: 382	7949
HIV33 1	Heavy chairs	**3BNC 107HC**	**US20 140328862** SEQ ID NO: 383	7950
HIV332	Heavy chain	**3BNC108HC**	US20 140328862 SEQ ID **NO:** 384	7951
HEV333	Heavy chain	**3BNC10HC**	**US20 140328862** SEQ ID NO: 385	7952
HIV334	Heavy chain	**3BNC114HC**	**US20 140328862** SEQ ID NO: 386	7953
HIV335	Heavy Chain	**3bncl l7**	Zhou T et **al., Immunity** 39 **(2),** 245-258 (2013), NCBI Accessiors # **4LSV_H** (226aa)	7954
HIV336	Heavy chain	**3BNC117HC**	**US20140328862** SEQ ID NO: 387	7955
HIV337	Heavy **chain**	**3BNC124HC**	**US20140328862** SEQ ID NO: 333	7956
HIV338	Heavy chairs	**3BNC 126HC**	**US20 140328862** SEQ ID NO: 388	7957
HIV339	Heavy chain	**3BNC127HC**	US20 140328862 SEQ ID NO: 389	7958
HIV340	Heavy chain	**3BNC130HC**	**US20 140328862** SEQ ID NO: 334	7959
HIV34 1	Heavy chain	**3BNC134HC**	**US20140328862** SEQ ID NO: 390	7960
HIV342	Heavy **chain**	**3BNC140HC**	**US20 140328862** SEQ ID NO: 391	7961
HIV343	Heavy chairs	**3BNC14 1HC**	**US20 140328862** SEQ ID NO: 392	7962
HIV344	Heavy chain	**3BNC142HC**	US20 140328862 SEQ ID NO: 393	7963
HIV345	Heavy chain	**3BNC148HC**	**US20 140328862** SEQ ID NO: 394	7964
HIV346	Heavy chain	**3BNC149HC**	**US20140328862** SEQ ID NO: 335	7965
HIV347	Heavy **chain**	**3BNC149HC**	**US20 140328862** SEQ ID NO: 395	7966
HIV348	Heavy chairs	**3BNC15 1HC**	**US20 140328862** SEQ ID NO: 396	7967
HIV349	Heavy chain	**3BNC153HC**	US20 140328862 SEQ ID NO: 397	7968
HIV350	Heavy chain	**3BNC156HC**	**US20140328862** SEQ ID NO: 398	7969
HIV35 1	Heavy chain	**3BNC158HC**	**US20 140328862** SEQ ID NO: 399	7970
HIV352	Heavy **chain**	**3BNC159HC**	**US20 140328862** SEQ ID NO: 400	7971
HIV353	Heavy chairs	3BNC15HC	US20 [40328862 SEQ ID NO: 401	7972
HIV354	Heavy chain	**3BNC 173HC**	US20 140328862 SEQ ID **NO:** 402	7973
HIV355	Heavy chain	**3BNC175HC**	**US20140328862** SEQ ID NO: 403	7974
HIV356	Heavy chain	**3BNC176HC**	**US20 140328862** SEQ ID NO: 404	7975
HIV357	Heavy **chain**	3BNC177HC	**US20 140328862** SEQ ID NO: 336	7976
HIV358	Heavy chain	3BNC17HC	**US20140328862** SEQ ID NO: 337	7977
HIV359	Heavy chain	**3BNC 181HC**	US20 140328862 SEQ ID **NO:** 405	7978
HEV360	Heavy chain	**3BNC186HC**	**US20140328862** SEQ ID NO: 406	7979
HIV361	Heavy chain	**3BNC18HC**	**US20 140328862** SEQ ID NO: 407	7980
HIV362	Heavy **chain**	**3BNC193HC**	**US20 140328862** SEQ ID NO: 408	7981
HIV363	Heavy chain	**3BNC196HC**	**US20140328862** SEQ ID NO: 409	7982
HIV364	Heavy chain	3BNC20HC	**US20140328862** SEQ **ID NO:** 410	7983
HIV365	Heavy chain	**3BNC29HC**	**US20140328862** SEQ ID NO: **4 11**	7984
HIV366	Heavy chain	**3BNC31HC**	**US20140328862** SEQ ID NO: 412	7985
HIV367	Heavy chain	3BNC33HC	**US20 140328862** SEQ ID NO: 413	7986
HIV368	Heavy chain	3BNC42HC	**US20140328862** SEQ ID NO: 414	7987
HIV369	Heavy chain	3BNC44HC	**US20140328862** SEQ **ID NO:** 415	7988
HEV370	Heavy chain	**3BNC45HC**	**US20140328862** SEQ ID NO: **4 16**	7989
HIV37 1	Heavy chain	**3BNC48HC**	**US20140328862** SEQ **ID** NO: 338	7990

HIV372	Heavy chain	3BNC53HC	US20140328862 SEQ ID NO: 417	7991
HIV373	Heavy chain	3BNC54HC	US20140328862 SEQ ID NO: 418	7992
HIV374	Heavy chain	3BNC55HC	US20140328862 SEQ ID NO: 419	7993
HIV375	Heavy chain	3BNC58HC	US20140328862 SEQ ID NO: 339	7994
HIV376	Heavy chain	3BNC59HC	US20140328862 SEQ ID NO: 420	7995
HIV377	Heavy chain	3BNC60HC	US20140328862 SEQ ID NO: 421	7996
HIV378	Heavy chain	3BNC62HC	US20140328862 SEQ ID NO: 422	7997
HIV379	Heavy chain	3BNC64HC	US20140328862 SEQ ID NO: 423	7998
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HIV381	Heavy chain	3BNC66HC	US20140328862 SEQ ID NO: 425	8000
HIV382	Heavy chain	3BNC6HC	US20140328862 SEQ ID NO: 426	8001
HIV383	Heavy chain	3BNC72HC	US20140328862 SEQ ID NO: 427	8002
HIV384	Heavy chain	3BNC75HC	US20140328862 SEQ ID NO: 428	8003
HIV385	Heavy chain	3BNC78HC	US20140328862 SEQ ID NO: 340	8004
HIV386	Heavy chain	3BNC79HC	US20140328862 SEQ ID NO: 429	8005
HIV387	Heavy chain	3BNC81HC	US20140328862 SEQ ID NO: 430	8006
HIV388	Heavy chain	3BNC82HC	US20140328862 SEQ ID NO: 341	8007
HIV389	Heavy chain	3BNC84HC	US20140328862 SEQ ID NO: 431	8008
HIV390	Heavy chain	3BNC86HC	US20140328862 SEQ ID NO: 432	8009
HIV391	Heavy chain	3BNC87HC	US20140328862 SEQ ID NO: 433	8010
HIV392	Heavy chain	3BNC89HC	US20140328862 SEQ ID NO: 434	8011
HIV393	Heavy chain	3BNC8HC	US20140328862 SEQ ID NO: 342	8012
HIV394	Heavy chain	3BNC91HC	US20140328862 SEQ ID NO: 435	8013
HIV395	Heavy chain	3BNC92HC	US20140328862 SEQ ID NO: 436	8014
HIV396	Heavy chain	3BNC94HC	US20140328862 SEQ ID NO: 437	8015
HIV397	Heavy chain	3BNC95HC	US20140328862 SEQ ID NO: 438	8016
HIV398	Heavy Chain	412d	Huang et al., Science 317 (5846), 1930-1934 (2007), NCBI Accession # 2QAD_H (231aa)	8017
HIV399	Heavy chain	43243_3	US20140205607 Table S14	8018
HIV400	Heavy chain	43359_2	US20140205607 Table S14	8019
HIV401	Heavy chain	43555_1	US20140205607 Table S14	8020
HIV402	Heavy chain	43567_2	US20140205607 Table S14	8021
HIV403	Heavy Chain	44-vrc13.01	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDJ_A(238aa)	8022
HIV404	Heavy chain	45-46m2	Diskin, R., et al., Restricting HIV-1 pathways for escape using rationally designed anti-HIV-1 antibodies; J. Exp. Med. 210 (6), 1235-1249 (2013), NCBI Accession # 4JKP_H (229aa)	8023
HIV405	Heavy chain	46260_1	US20140205607 Table S14	8024
HIV406	Heavy chain	47890_1	US20140205607 Table S14	8025
HIV407	Heavy Chain	4e10 Fv	Finton, K.A., et al., PLoS Pathol. 9 (9), E1003639 (2013), NCBI Accession # 4LLV_A (129aa)	8026
HIV408	Heavy chain	53821_1	US20140205607 Table S14	8027
HIV409	Heavy chain	57729_2	US20140205607 Table S14	8028

HIV410	Heavy chain	61048_1	US20140205607 Table S14	8029
HIV411	Heavy chain	69713_1	US20140205607 Table S14	8030
HIV412	Heavy chain	70679_1	US20140205607 Table S14	8031
HIV413	Heavy chain	71632_2	US20140205607 Table S14	8032
HIV414	Heavy chain	74400_3	US20140205607 Table S14	8033
HIV415	Heavy chain	74511_1	US20140205607 Table S14	8034
HIV416	Heavy chain	76927_2	US20140205607 Table S14	8035
HIV417	Heavy Chain	7b2	Santra, S., et al., PLoS Pathol. 11 (8), E1005042 (2015), NCBI Accession # 4YDV_H (252aa)	8036
HIV418	Heavy chain	7H6	US20140348785 SEQ ID NO: 3	8037
HIV419	Heavy chain	7N16	US20140348785 SEQ ID NO: 5	8038
HIV420	Heavy chain	8460_4	US20140205607 Table S14	8039
HIV421	Heavy chain	86277_2	US20140205607 Table S14	8040
HIV422	Heavy chain	86343_1	US20140205607 Table S14	8041
HIV423	Heavy chain	86984_2	US20140205607 Table S14	8042
HIV424	Heavy chain	89680_4	US20140205607 Table S14	8043
HIV425	Heavy chain	8A253HC	US20140328862 SEQ ID NO: 5	8044
HIV426	Heavy chain	8A275HC	US20140328862 SEQ ID NO: 6	8045
HIV427	Heavy chain	8ABM11	US20140328862 SEQ ID NO: 7	8046
HIV428	Heavy chain	8ABM12	US20140328862 SEQ ID NO: 8	8047
HIV429	Heavy chain	8ABM13	US20140328862 SEQ ID NO: 9	8048
HIV430	Heavy chain	8ABM14	US20140328862 SEQ ID NO: 10	8049
HIV431	Heavy chain	8ABM20	US20140328862 SEQ ID NO: 11	8050
HIV432	Heavy chain	8ABM24	US20140328862 SEQ ID NO: 12	8051
HIV433	Heavy chain	8ABM26	US20140328862 SEQ ID NO: 13	8052
HIV434	Heavy chain	8ABM27	US20140328862 SEQ ID NO: 14	8053
HIV435	Heavy chain	8ANC103HC	US20140328862 SEQ ID NO: 36	8054
HIV436	Heavy chain	8ANC105HC	US20140328862 SEQ ID NO: 15	8055
HIV437	Heavy chain	8ANC106HC	US20140328862 SEQ ID NO: 37	8056
HIV438	Heavy chain	8ANC107HC	US20140328862 SEQ ID NO: 38	8057
HIV439	Heavy chain	8ANC108HC	US20140328862 SEQ ID NO: 39	8058
HIV440	Heavy chain	8ANC109HC	US20140328862 SEQ ID NO: 40	8059
HIV441	Heavy chain	8ANC10HC	US20140328862 SEQ ID NO: 41	8060
HIV442	Heavy chain	8ANC111HC	US20140328862 SEQ ID NO: 42	8061
HIV443	Heavy chain	8ANC112HC	US20140328862 SEQ ID NO: 43	8062
HIV444	Heavy chain	8ANC113HC	US20140328862 SEQ ID NO: 44	8063
HIV445	Heavy chain	8ANC114HC	US20140328862 SEQ ID NO: 45	8064
HIV446	Heavy chain	8ANC115HC	US20140328862 SEQ ID NO: 46	8065
HIV447	Heavy chain	8ANC116HC	US20140328862 SEQ ID NO: 16	8066
HIV448	Heavy chain	8ANC117HC	US20140328862 SEQ ID NO: 47	8067
HIV449	Heavy chain	8ANC11HC	US20140328862 SEQ ID NO: 48	8068
HIV450	Heavy chain	8ANC121HC	US20140328862 SEQ ID NO: 49	8069
HIV451	Heavy chain	8ANC126HC	US20140328862 SEQ ID NO: 50	8070
HIV452	Heavy chain	8ANC127HC	US20140328862 SEQ ID NO: 17	8071

HIV453	Heavy chain	8ANC130HC	US20140328862 SEQ ID NO: 51	8072
HIV454	Heavy Chain	8anc131	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4RWY_H (227aa)	8073
HIV455	Heavy chain	8ANC131HC	US20140328862 SEQ ID NO: 18	8074
HIV456	Heavy chain	8ANC132HC	US20140328862 SEQ ID NO: 52	8075
HIV457	Heavy chain	8ANC133HC	US20140328862 SEQ ID NO: 53	8076
HIV458	Heavy Chain	8anc134	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4RX4_H (229aa)	8077
HIV459	Heavy chain	8ANC134HC	US20140328862 SEQ ID NO: 19	8078
HIV460	Heavy chain	8ANC136HC	US20140328862 SEQ ID NO: 54	8079
HIV461	Heavy chain	8ANC137HC	US20140328862 SEQ ID NO: 55	8080
HIV462	Heavy chain	8ANC139HC	US20140328862 SEQ ID NO: 56	8081
HIV463	Heavy chain	8ANC13HC	US20140328862 SEQ ID NO: 20	8082
HIV464	Heavy chain	8ANC140HC	US20140328862 SEQ ID NO: 57	8083
HIV465	Heavy chain	8ANC142HC	US20140328862 SEQ ID NO: 58	8084
HIV466	Heavy chain	8ANC143HC	US20140328862 SEQ ID NO: 59	8085
HIV467	Heavy chain	8ANC144HC	US20140328862 SEQ ID NO: 60	8086
HIV468	Heavy chain	8ANC145HC	US20140328862 SEQ ID NO: 61	8087
HIV469	Heavy chain	8ANC146HC	US20140328862 SEQ ID NO: 62	8088
HIV470	Heavy chain	8ANC147HC	US20140328862 SEQ ID NO: 63	8089
HIV471	Heavy chain	8ANC148HC	US20140328862 SEQ ID NO: 64	8090
HIV472	Heavy chain	8ANC149HC	US20140328862 SEQ ID NO: 65	8091
HIV473	Heavy chain	8ANC14HC	US20140328862 SEQ ID NO: 66	8092
HIV474	Heavy chain	8ANC150HC	US20140328862 SEQ ID NO: 67	8093
HIV475	Heavy chain	8ANC151HC	US20140328862 SEQ ID NO: 68	8094
HIV476	Heavy chain	8ANC153HC	US20140328862 SEQ ID NO: 69	8095
HIV477	Heavy chain	8ANC154HC	US20140328862 SEQ ID NO: 70	8096
HIV478	Heavy chain	8ANC155HC	US20140328862 SEQ ID NO: 71	8097
HIV479	Heavy chain	8ANC156HC	US20140328862 SEQ ID NO: 72	8098
HIV480	Heavy chain	8ANC157HC	US20140328862 SEQ ID NO: 73	8099
HIV481	Heavy chain	8ANC158HC	US20140328862 SEQ ID NO: 74	8100
HIV482	Heavy chain	8ANC160HC	US20140328862 SEQ ID NO: 75	8101
HIV483	Heavy chain	8ANC161HC	US20140328862 SEQ ID NO: 76	8102
HIV484	Heavy chain	8ANC162HC	US20140328862 SEQ ID NO: 77	8103
HIV485	Heavy chain	8ANC163HC	US20140328862 SEQ ID NO: 78	8104
HIV486	Heavy chain	8ANC164HC	US20140328862 SEQ ID NO: 79	8105
HIV487	Heavy chain	8ANC165HC	US20140328862 SEQ ID NO: 80	8106
HIV488	Heavy chain	8ANC166HC	US20140328862 SEQ ID NO: 81	8107
HIV489	Heavy chain	8ANC168HC	US20140328862 SEQ ID NO: 82	8108
HIV490	Heavy chain	8ANC169HC	US20140328862 SEQ ID NO: 83	8109
HIV491	Heavy chain	8ANC16HC	US20140328862 SEQ ID NO: 84	8110

HEV492	Heavy chain	8ANC171HC	US20140328862 SEQ ID NO: 21	8111
HIV493	Heavy chain	8ANC173HC	US20 140328862 SEQ ID NO: 85	8112
HIV494	Heavy chain	8ANC 174HC	US20 140328862 SEQ ID NO: 86	8113
HTV495	Heavy chairs	8ANC175HC	US20 140328862 SEQ ID NO: 87	8114
HIV496	Heavy chain	8ANC I76HC	US20 140328862 SEQ ID NO: 88	8115
HEV497	Heavy chain	8ANC177HC	US20140328862 SEQ ID NO: 89	8116
HIV498	Heavy chain	8ANC178HC	US20 140328862 SEQ ID NO: 90	8117
HIV499	Heavy chain	8ANC 179HC	US20 140328862 SEQ ID NO: 91	8118
HIV500	Heavy chain	8ANC17HC	US20140328862 SEQ ID NO: 92	8119
HIV50 i	Heavy chain	8ANC I8	US20 140328862 SEQ **ID** NO: 22	8120
HEV502	Heavy chain	8ANC180HC	US20140328862 SEQ ID NO: 93	8121
HIV503	Heavy chain	8ANC181HC	US20 140328862 SEQ ID NO: 94	8122
HIV504	Heavy chain	8ANC 182HC	US20 140328862 SEQ ID NO: 23	8123
HIV505	Heavy chain	8ANC184HC	US20140328862 SEQ ID NO: 95	8124
HIV506	Heavy chain	8ANC I85HC	US20 140328862 SEQ **ID** NO: 96	8125
HEV507	Heavy chain	8ANC186HC	US20140328862 SEQ ID NO: 97	8126
HIV508	Heavy chain	8ANC 187HC	US20 140328862 SEQ ID NO: 98	8127
HIV509	Heavy chain	8ANC188HC	US20 140328862 SEQ ID NO: 99	8128
HIV5 10	Heavy chain	8ANC191HC	US20 140328862 SEQ ID NO: 100	8129
HIV5 1I	Heavy chain	8ANC I92HC	US20 140328862 SEQ ID NO: 24	8130
HBV512	Heavy chain	8ANC193HC	US20 140328862 SEQ ID NO: 101	8131
HIV5 13	Heavy chain	8ANC 194HC	US20 140328862 SEQ ID NO: 102	8132
HIV5 14	Heavy chain	8ANC195HC	US20 140328862 SEQ ID NO: 103	8133
HIV5 15	Heavy chain	8ANC196HC	US20 140328862 SEQ ID NO: 104	8134
HIV5 16	Heavy chain	8ANC20HC	US20 140328862 SEQ **ID** NO: 105	8135
HBV517	Heavy chairs	8ANC2 IHC	US20 [40328862 SEQ ID NO: 106	8136
HIV5 18	Heavy chain	8ANC22HC	US20 140328862 SEQ ID NO: 25	8137
HIV5 19	Heavy chain	8ANC24HC	US20 140328862 SEQ ID NO: 107	8138
HIV520	Heavy chain	8ANC25HC	US20 140328862 SEQ ID NO: 108	8139
HIV52 1	Heavy chain	8ANC26HC	US20 140328862 SEQ ID NO: 26	8140
HTV522	Heavy chairs	8ANC27HC	US20 [40328862 SEQ ID NO: 109	8141
HIV523	Heavy chain	8ANC2HC	US20 140328862 SEQ ID NO: 27	8142
HIV524	Heavy chain	8ANC30HC	US20 140328862 SEQ ID NO: 28	8143
HIV525	Heavy chain	8ANC3 1HC	US20 140328862 SEQ ID NO: 110	8144
HIV526	Heavy chain	8ANC33HC	US20 140328862 SEQ ID NO: 111	8145
HTV527	Heavy chairs	8ANC34HC	US20 [40328862 SEQ ID NO: 112	8146
HIV528	Heavy chain	8ANC36HC	US20 140328862 SEQ ID NO: 113	8147
HEV529	Heavy chain	8ANC37HC	US20140328862 SEQ ID NO: 29	8148
HIV530	Heavy chain	8ANC38HC	US20 140328862 SEQ ID NO: 114	8149
HIV53 I	Heavy chain	8ANC39HC	US20 140328862 SEQ ID NO: 115	8150
HIV532	Heavy chairs	8ANC3HC	US20 [40328862 SEQ ID NO: 116	8151
HIV533	Heavy chain	8ANC40HC	US20 140328862 SEQ ID NO: 30	8152
HEV534	Heavy chain	8ANC41HC	US20140328862 SEQ ID NO: 31	8153
HIV535	Heavy chain	8ANC43HC	US20 140328862 SEQ ID NO: 117	8154
HIV536	Heavy chain	8ANC45HC	US20 140328862 SEQ ID NO: 32	8155

HEV5.37	Hie;n′s chain	8ANC46HC	US20 140328862 SEQ ID NO: 118	8 156
HIV538	Heavy chain	8ANC48HC	US20 140328862 SEQ ID NO: 119	8157
HIV539	Heavy chain	8ANC49HC	US20 140328862 SEQ ID NO: 120	8158
HTV540	Heavy chairs	8ANC50HC	US20 140328862 SEQ ID NO: 33	8159
HIV54 i	Heavy chain	8ANC51HC	US20 140328862 SEQ ID NO: 121	8160
HEV542	Heavy chain	8ANC53HC	US20140328862 SEQ ID NO: 34	8 161
HIV543	Heavy chain	8ANC57HC	US20 140328862 SEQ ID NO: 122	8162
HIV544	Heavy chain	8ANC58HC	US20 140328862 SEQ ID NO: 123	8163
HIV545	Heavy chain	8ANC5HC	US20 140328862 SEQ ID NO: 124	8 164
HIV546	Heavy chain	8ANC60HC	US20 140328862 SEQ ID NO: 125	8165
HEV547	Heavy chain	8ANC63HC	US20 140328862 SEQ ID NO: 126	8 166
HIV548	Heavy chain	8ANC65HC	US20 140328862 SEQ ID NO: 127	8167
HIV549	Heavy chain	8ANC67HC	US20 140328862 SEQ ID NO: 128	8168
HIV550	Heavy chain	8ANC69HC	US20 140328862 SEQ ID NO: 129	8 169
HIV55 i	Heavy chain	8ANC6HC	US20 140328862 SEQ ID NO: 130	8170
HEV552	Heavy chain	8ANC70HC	US20 140328862 SEQ ID NO: 131	8 171
HIV55.3	Heavy chain	8ANC71HC	US20 140328862 SEQ ID NO: 132	8172
HIV554	Heavy chain	8ANC72HC	US20 140328862 SEQ ID NO: 133	8173
HIV555	Heavy chain	8ANC74HC	US20 140328862 SEQ ID NO: 134	8 174
HIV556	Heavy chain	8ANC75HC	US20 140328862 SEQ ID NO: 135	8175
HEV557	Heavy chain	8ANC76HC	US20 140328862 SEQ ID NO: 136	8 176
HIV558	Heavy chain	8ANC78HC	US20 140328862 SEQ ID NO: 137	8177
HIV559	Heavy chain	8ANC79HC	US20 140328862 SEQ ID NO: 138	8178
HIV560	Heavy chain	8ANC7HC	US20 140328862 SEQ ID NO: 139	8 179
HIV56 i	Heavy chain	8ANC80HC	US20 140328862 SEQ ID NO: 140	8180
HTV562	Heavy chairs	8ANC82HC	US20 140328862 SEQ ID NO: 141	8 18 1
HIV56.3	Heavy chain	8ANC83HC	US20 140328862 SEQ ID NO: 142	8182
HIV564	Heavy chain	8ANC88HC	US20 140328862 SEQ ID NO: 35	8183
HIV565	Heavy chain	8ANC91HC	US20 140328862 SEQ ID NO: 143	8 184
HIV566	Heavy chain	8ANC92HC	US20 140328862 SEQ ID NO: 144	8185
HTV567	Heavy chairs	8ANC93HC	US20 140328862 SEQ ID NO: 145	8 186
HIV568	Heavy chain	8ANC9HC	US20 140328862 SEQ ID NO: 146	8187
HIV569	Heavy chain	94565_1	US20 140205607 Table S14	8188
HIV570	Heavy chain	95589_2	US20 140205607 Table S14	8189
HIV57 1	Heavy chain	96298_1	US20 140205607 Table S14	8190
HTV572	Heavy chairs	9815_2	US20 140205607 Table S14	8 19 1
HIV573	Heavy chain	99473_3	US20 140205607 Table S14	8192
HEV574	Heavy chain	99989_1	US20140205607 Table S14	8 193
HIV575	Heavy chain	Antibody	US20140328862 SEQ ID NO: 439	8194
HIV576	Heavy chain	Anti-HcG	Fotinoii C. et al " Structure of an Fab fragment against a C-terminal peptide of hCG at 2.0 A resolution" J. Biol. Chem. 273 (35), 225 15-225 18 (1998); NCBI Accession # 1SBS_H	8195
HEV577	Heavy Chain	B 12	Zhou T et al., Structural definition of a conserved neutralization epitope on HIV-1 gpl20; Nature 445 (7129), 732-737 (2007), NCBI Accession # 2NY7_H (2.30aa)	8 196

HIV578	Heavy Chain	C38-vrc16.01	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDK_H (234aa)	8197
HIV579	Heavy Chain	C38-vrc18.02	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDL_H (226aa)	8198
HIV580	Heavy chain	CAP256-VRC26.01	WO2015128846 SEQ ID NO: 13	8199
HIV581	Heavy chain	CAP256-VRC26.02	WO2015128846 SEQ ID NO: 17	8200
HIV582	Heavy chain	CAP256-VRC26.03	WO2015128846 SEQ ID NO: 21	8201
HIV583	Heavy chain	CAP256-VRC26.04	WO2015128846 SEQ ID NO: 25	8202
HIV584	Heavy chain	CAP256-VRC26.05	WO2015128846 SEQ ID NO: 29	8203
HIV585	Heavy chain	CAP256-VRC26.06	WO2015128846 SEQ ID NO: 33	8204
HIV586	Heavy chain	CAP256-VRC26.07	WO2015128846 SEQ ID NO: 37	8205
HIV587	Heavy chain	CAP256-VRC26.08	WO2015128846 SEQ ID NO: 41	8206
HIV588	Heavy chain	CAP256-VRC26.09	WO2015128846 SEQ ID NO: 45	8207
HIV589	Heavy chain	CAP256-VRC26.10	WO2015128846 SEQ ID NO: 49	8208
HIV590	Heavy chain	CAP256-VRC26.11	WO2015128846 SEQ ID NO: 53	8209
HIV591	Heavy chain	CAP256-VRC26.12	WO2015128846 SEQ ID NO: 57	8210
HIV592	Heavy chain	CAP256-VRC26.25	WO2015128846 SEQ ID NO: 170	8211
HIV593	Heavy chain	CAP256-VRC26.26	WO2015128846 SEQ ID NO: 178	8212
HIV594	Heavy chain	CAP256-VRC26.27	WO2015128846 SEQ ID NO: 186	8213
HIV595	Heavy chain	CAP256-VRC26-I1	WO2015128846 SEQ ID NO: 5	8214
HIV596	Heavy chain	CAP256-VRC26-I2.	WO2015128846 SEQ ID NO: 9	8215
HIV597	Heavy chain	CAP256-VRC26-UCA.	WO2015128846 SEQ ID NO: 1	8216
HIV598	Heavy chain	construct #2816, #2859	WO2015013390 SEQ ID NO: 3	8217
HIV599	Heavy chain	construct #2817	WO2015013390 SEQ ID NO: 4	8218
HIV600	Heavy chain	construct #2858, #2860	WO2015013390 SEQ ID NO: 8	8219
HIV601	Heavy Chain	Fab 2219	Stanfield, R.L., et al., J. Virol. 80 (12), 6093-6105 (2006), NCBI Accession # 2B0S_H (226aa)	8220
HIV602	Heavy Chain	Fab 2g12	Doores, K.J., et al., J. Virol. 84 (20), 10690-10699 (2010), NCBI Accession # 3OAU_H (225aa)	8221

HIV603	Heavy Chain	Fab 2g12	Stanfield, R.L. et al., Crystal structure of the HIV neutralizing antibody 2G12 in complex with a bacterial oligosaccharide analog of mammalian oligomannose; Glycobiology 25 (4), 412-419 (2015), NCBI Accession # 4RBP_H (224aa)	8222
HIV604	Heavy Chain	Fab F425-b4e8	Bell et al., J. Mol. Biol. 375 (4), 969-978 (2008), NCBI Accession # 2QSC_H (222aa)	8223
HIV605	Heavy chain	fusion protein of A32 and m9	US20080038280 SEQ ID NO: 5	8224
HIV606	Heavy chain	g20	WO2015117008 SEQ ID NO: 4	8225
HIV607	Heavy chain	g22	WO2015117008 SEQ ID NO: 7	8226
HIV608	Heavy chain	g23	WO2015117008 SEQ ID NO: 2	8227
HIV609	Heavy chain	g3	WO2015117008 SEQ ID NO: 13	8228
HIV610	Heavy chain	g4	WO2015117008 SEQ ID NO: 9	8229
HIV611	Heavy chain	g44	WO2015117008 SEQ ID NO: 11	8230
HIV612	Heavy chain	g46	WO2015117008 SEQ ID NO: 10	8231
HIV613	Heavy chain	G4D	US20130195881 SEQ ID NO: 9	8232
HIV614	Heavy chain	G4H	US20130195881 SEQ ID NO: 8	8233
HIV615	Heavy chain	g50	WO2015117008 SEQ ID NO: 12	8234
HIV616	Heavy chain	g52	WO2015117008 SEQ ID NO: 1	8235
HIV617	Heavy chain	g59	WO2015117008 SEQ ID NO: 5	8236
HIV618	Heavy chain	g62	WO2015117008 SEQ ID NO: 6	8237
HIV619	Heavy chain	g8	WO2015117008 SEQ ID NO: 3	8238
HIV620	Heavy chain	gl5	WO2015117008 SEQ ID NO: 8	8239
HIV621	Heavy chain	gVRC-H5(d74)/VRC-PG04LC	WO2013090644 SEQ ID NO: 45	8240
HIV622	Heavy chain	gVRCOH12(D74)/VRC-PG04LC	WO2013090644 SEQ ID NO: 46	8241
HIV623	Heavy Chain	l2 (unbound) From Ch103 Lineage	Fera, D. et al., Affinity maturation in an HIV broadly neutralizing B-cell lineage through reorientation of variable domains; Proc. Natl. Acad. Sci. U.S.A. 111 (28), 10275-10280 (2014), NCBI Accession # 4QHN_A (232aa)	8242
HIV624	Heavy chain	IGHV3-15*05	US20140348785 SEQ ID NO: 7	8243
HIV625	Heavy chain	LSSB2055HC	US20140328862 SEQ ID NO: 229	8244
HIV626	Heavy chain	LSSB2066HC	US20140328862 SEQ ID NO: 230	8245
HIV627	Heavy chain	LSSB2068HC	US20140328862 SEQ ID NO: 231	8246
HIV628	Heavy chain	LSSB2080HC	US20140328862 SEQ ID NO: 232	8247
HIV629	Heavy chain	LSSB2133HC	US20140328862 SEQ ID NO: 233	8248
HIV630	Heavy chain	LSSB2182HC	US20140328862 SEQ ID NO: 234	8249
HIV631	Heavy chain	LSSB218HC	US20140328862 SEQ ID NO: 235	8250
HIV632	Heavy chain	LSSB2277HC	US20140328862 SEQ ID NO: 236	8251

HIV633	Heavy chain	**LSSB2288H** C	**US20140328862** SEQ ID NO: 237	8252
HIV634	Heavy chain	LSSB2339H C	**US20140328862** SEQ **ID** NO: **168**	8253
HIV635	Heavy **chain**	**LSSB235 1H** C	**US20140328862** SEQ ID NO: 169	8254
HIV636	Heavy chain	LSSB2361H C	**US20140328862** SEQ ID NO: 170	8255
HIV637	Heavy chain	LSSB2364H C	US20140328862 SEQ ID NO: 171	8256
HIV638	Heavy chain	LSSB2367H C	**US20140328862** SEQ ID NO: 172	8257
HIV639	Heavy chain	LSSB2416H C	**US20140328862** SEQ ID NO: 173	8258
HIV640	Heavy chain	LSSB2434H C	**US20140328862** SEQ ID NO: 174	8259
HIV641	Heavy chain	LSSB2483H C	**US20140328862** SEQ ID NO: 175	8260
HIV642	Heavy chain	LSSB2490H C	**US20140328862** SEQ ID NO: 176	8261
HIV643	Heavy chain	LSSB2503H C	**US20140328862** SEQ ID NO: 177	8262
HIV644	**Heavy** chain	**LSSB2525H** C	**US20140328862** SEQ ID NO: 178	8263
HIV645	Heavy **chain**	LSSB2530H **C**	**US20140328862** SEQ ID NO: 179	8264
HIV646	**Heavy** chain	**LSSB2538H** C	**US20140328862** SEQ ID NO: 180	8265
HIV647	Heavy **chain**	LSSB2554H **C**	**US20140328862** SEQ ID NO: **18 1**	8266
HIV648	**Heavy** chain	**LSSB2573H** C	**US20140328862** SEQ ID NO: 182	8267
HIV649	Heavy **chain**	LSSB2578H **C**	**US20140328862** SEQ ID NO: 183	8268
HIV650	**Heavy** chain	**LSSB2586H** C	**US20140328862** SEQ ID NO: 184	8269
HIV65 I	Heavy **chain**	LSSB2609H **C**	**US20140328862** SEQ ID NO: 185	8270
HIV652	**Heavy** chain	**LSSB2612H** C	**US20140328862** SEQ ID NO: 186	8271
HIV653	Heavy chain	**LSSB2630H** C	**US20140328862** SEQ **ID NO:** 187	8272
HTV654	Heavy chairs	LSSB2640H C	**US20 140328862** SEQ ID NO: 188	8273
HIV655	Heavy chain	**LSSB2644H** C	**US20140328862** SEQ ID **NO:** 189	8274
HTV656	Heavy chairs	LSSB2665H C	**US20 140328862** SEQ ID NO: 190	8275
HIV657	Heavy chain	LSSB2666H C	**US20140328862** SEQ ID **NO:** 19 !	8276
HTV658	Heavy chairs	LSSB2669H C	**US20 140328862** SEQ ID NO: 192	8277
HIV659	Heavy chain	**LSSB2680H** C	**US20140328862** SEQ ID **NO:** 193	8278
HTV660	Heavy chairs	**LSSB2683H** C	**US20 140328862** SEQ ID NO: 194	8279
HIV66 1	Heavy chain	**LSSB33 1HC**	**US20140328862** SEQ ID NO: 238	8280
HEV662	Heavy chain	**LSSB344HC**	**US20 140328862** SEQ ID NO: 195	8281

HEV663	Heavy chain	LSSNEC 101 HC	US20140328862 SEQ ID NO: 239	8282
HIV664	Heavy chain	LSSNEC106 HC	US20140328862 SEQ ID NO: 240	8283
HEV665	Heavy chain	LSSNEC 107 HC	US20140328802 SEQ ID NO: 196	8284
HIV666	Heavy chain	LSSNEC108 HC	US20140328862 SEQ ID NO: 197	8285
HIV667	Heavy chain	LSSNEC109 HC	US20140328862 SEQ ID NO: 198	8286
HIV668	Heavy chain	LSSNEC1 10 HC	US20140328862 SEQ ID NO: 199	8287
HIV669	Heavy chain	LSSNEC112 HC	US20140328862 SEQ ID NO: 241	8288
HIV670	Heavy chain	LSSNEC1 15 HC	US20140328862 SEQ ID NO: 242	8289
HIV671	Heavy chain	LSSNEC1 16 HC	US20140328862 SEQ ID NO: 200	8290
HIV672	Heavy chain	LSSNEC1 17 HC	US20140328862 SEQ ID NO: 201	8291
HIV673	Heavy chain	LSSNEC1 18 HC	US20140328862 SEQ ID NO: 202	8292
HIV674	Heavy chain	LSSNEC 11H C	US20140328862 SEQ ID NO: 203	8293
HIV675	Heavy chain	LSSNEC122 HC	US20140328862 SEQ ID NO: 204	8294
HIV676	Heavy chain	LSSNEC 123 HC	US20140328862 SEQ ID NO: 205	8295
HIV677	Heavy chain	LSSNEC124 HC	US20140328862 SEQ ID NO: 243	8296
HIV678	Heavy chain	LSSNEC 125 HC	US20140328862 SEQ ID NO: 244	8297
HIV679	Heavy chain	LSSNEC126 HC	US20140328862 SEQ ID NO: 245	8298
HIV680	Heavy chain	LSSNEC 127 HC	US20140328862 SEQ ID NO: 206	8299
HIV68 1	Heavy chain	LSSNECUH C	US20140328862 SEQ ID NO: 246	8300
HIV682	Heavy chain	LSSNEC 16H C	US20140328862 SEQ ID NO: 247	8301
HIV683	Heavy chain	LSSNEC18H C	US20140328862 SEQ ID NO: 207	8302
HTV684	Heavy chairs	LSSNEC21H C	US20 140328862 SEQ ID NO: 248	8303
HIV685	Heavy chain	LSSNEC24H C	US20140328862 SEQ ID NO: 208	8304
HTV686	Heavy chairs	LSSNEC29H C	US20 140328862 SEQ ID NO: 209	8305
HIV687	Heavy chain	LSSNEC2HC	US20 140328862 SEQ ID NO: 210	8306
HEV688	Heavy chain	LSSNEC30H C	US20140328862 SEQ ID NO: 249	8307
HIV689	Heavy chain	LSSNEC33H C	US20140328862 SEQ ID NO: 211	8308
HEV690	Heavy chain	LSSNEC34H C	US20140328862 SEQ ID NO: 212	8309
HIV69 1	Heavy chain	LSSNEC3HC	US20140328862 SEQ ID NO: 213	8310
HIV692	Heavy chain	LSSNEC46H C	US20140328862 SEQ ID NO: 214	8311

HEV693	Heavy chain	LSSNEC48HC	US20140328862 SEQ ID NO: 215	8312
HIV694	Heavy chain	LSSNEC49HC	US20140328862 SEQ ID NO: 250	8313
HEV695	Heavy chain	LSSNEC52HC	US20140328862 SEQ ID NO: 216	8314
HIV696	Heavy chain	LSSNEC54HC	US20140328862 SEQ ID NO: 251	8315
HIV697	Heavy chain	LSSNEC55HC	US20140328862 SEQ ID NO: 252	8316
HIV698	Heavy chain	LSSNEC56HC	US20140328862 SEQ ID NO: 217	8317
HIV699	Heavy chain	LSSNEC57HC	US20140328862 SEQ ID NO: 253	8318
HIV700	Heavy chain	LSSNEC5HC	US20 140328862 SEQ ID NO: 254	8319
HIV70 1	Heavy chain	LSSNEC60HC	US20140328862 SEQ ID NO: 218	8320
HIV702	Heavy chain	LSSNEC66HC	US20140328862 SEQ ID NO: 219	8321
HIV703	Heavy chain	LSSNEC67HC	US20140328862 SEQ ID NO: 255	8322
HIV704	Heavy chain	LSSNEC70HC	US20140328862 SEQ ID NO: 220	8323
HIV705	Heavy chain	LSSNEC72HC	US20140328862 SEQ ID NO: 221	8324
HIV706	Heavy chain	LSSNEC74HC	US20140328862 SEQ ID NO: 256	8325
HIV707	Heavy chain	LSSNEC77HC	US20140328862 SEQ ID NO: 257	8326
HTV708	Heavy chairs	LSSNEC7HC	US20 140328862 SEQ ID NO: 222	8327
HIV709	Heavy chain	LSSNEC82HC	US20140328862 SEQ ID NO: 223	8328
HIV7 I0	Heavy chairs	LSSNEC85HC	US20 140328862 SEQ ID NO: 258	8329
HIV71 1	Heavy chain	LSSNEC89HC	US20140328862 SEQ ID NO: 224	8330
HEV712	Heavy chain	LSSNEC8HC	US20140328862 SEQ ID NO: 225	8.33 1
HIV713	Heavy chain	LSSNEC91HC	US20140328862 SEQ ID NO: 259	8332
HIV714	Heavy chain	LSSNEC92HC	US20140328862 SEQ ID NO: 260	8333
HIV715	Heavy chain	LSSNEC94HC	US20140328862 SEQ ID NO: 226	8334
HIV716	Heavy chain	LSSNEC95HC	US20140328862 SEQ ID NO: 227	8335
HIV717	Heavy chain	LSSNEC9HC	US20 140328862 SEQ ID NO: 228	8336
HIV718	Heavy chain	m l2_Fd-aa	US7803913B2 SEQ ID NO: 3	8337
HTV7 I9	Heavy chairs	ml4-Fd-aa	US78039 13B2 SEQ ID NO: 1	8338
HIV720	Heavy chain	m l6-Fd-aa	US7803913B2 SEQ ID NO: 4	8339
HEV721	Heavy chain	ml8_Fd-aa	US7803913B2 SEQ ID NO: 2	8340
HIV722	Heavy Chain	M66	Ofek, G., et al, Structural Basis for HIV-1 Neutralization by 2F5-Like Antibodies m66 and m66.6; J. Virol. 88 (5), 2426-2441 (2014), NCBE Accession ii 4NRY L (220aa)	8341
HIV723	Heavy Chain	M66.6	Ofek, G., et al., Stnictiiral Basis for HIV-1 Neutralization by 2F5-Like Antibodies 11166	8342

			and m66.6; J. Virol. 88 (5), 2426-2441 (2014), NCBI Accession # 4NRZ_H (234aa)	
HIV724	Heavy Chain	Mab 2158	Spurrier, B., et al., Functional Implications of the Binding Mode of a Human Conformation-Dependent V2 Monoclonal Antibody against HIV; J. Virol. 88 (8), 4100-4112 (2014), NCBI Accession # 4OAW_D (236aa)	8343
HIV725	Heavy chain	MV1	US20130195881 SEQ ID NO: 10	8344
HIV726	Heavy Chain	Pg16 Fab	Pancera, M., et al., Nat. Struct. Mol. Biol. 20 (7), 804-813 (2013), NCBI Accession # 4DQO_H (246aa)	8345
HIV727	Heavy Chain	Pg9	Willis, J.R., et al., J. Clin. Invest. 125 (6), 2523-2531 (2015), NCBI Accession # 4YAQ_H(248aa)	8346
HIV728	Heavy Chain	Pgt121-GI Fab	Mouquet H et al., Complex-type N-glycan recognition by potent broadly neutralizing HIV antibodies; Proc Natl Acad Sci U S A. 2012 Nov 20;109(47):E3268-77, NCBI Accession # 4FQQ_B (244aa)	8347
HIV729	Heavy Chain	Pgt122	Julien, J.P., et al., PLoS Pathol. 9 (5), E1003342 (2013), NCBI Accession # 4JY5_H (235aa)	8348
HIV730	Heavy Chain	Pgt123	Julien, J.P., et al., PLoS Pathol. 9 (5), E1003342 (2013), NCBI Accession # 4JY6_B (235aa)	8349
HIV731	Heavy Chain	Pgt124	Garces, F., et al., Structural Evolution of Glycan Recognition by a Family of Potent HIV Antibodies; Cell 159 (1), 69-79 (2014), NCBI Accession # 4R26_H (236aa)	8350
HIV732	Heavy Chain	Pgt130	Doores, K.J., et al., J. Virol. 89 (2), 1105-1118 (2015), NCBI Accession # 4RNR_A (233aa)	8351
HIV733	Heavy Chain	Pgt135	Grover et al., Science 343 (6171), 656-661 (2014), NCBI Accession # 4NZR_H (234aa)	8352
HIV734	Heavy chain	S19	US20110059015 SEQ ID NO: 6	8353
HIV735	Heavy chain	S20	US20110059015 SEQ ID NO: 8	8354
HIV736	Heavy chain	S8	US20110059015 SEQ ID NO: 4	8355
HIV737	Heavy Chain	Vrc- Pg04	Wu, X., et al., Focused evolution of HIV-1 neutralizing antibodies revealed by structures and deep sequencing; Science 333 (6049), 1593-1602 (2011)", NCBI Accession # 3SE9_H (228aa)	8356
HIV738	Heavy chain	VRC01	US8637036B2 SEQ ID NO: 1	8357
HIV739	Heavy chain	VRC01HC/VRC03LC	WO2013090644 SEQ ID NO: 2	8358
HIV740	Heavy chain	VRC02	US8637036B2 SEQ ID NO: 3	8359
HIV741	Heavy chain	VRC03	US8637036B2 SEQ ID NO: 27	8360
HIV742	Heavy chain	VRC03HC-VRC01LC	WO2013090644 SEQ ID NO: 32	8361
HIV743	Heavy chain	VRC07 G54H, S58N	US20140322163 SEQ ID NO: 258	8362
HIV744	Heavy chain	VRC07 I37V, G54H, S58N, T93A	US20140322163 SEQ ID NO: 260	8363
HIV745	Heavy chain	VRC07 I37V, G54H, T93A	US20140322163 SEQ ID NO: 259	8364
HIV746	Heavy Chain	Vrc08c	Wu, X., et al., Maturation and Diversity of the VRC01-Antibody Lineage over 15 Years of	8365

			Chronic HIV-1 Infection; Cell 161 (3), 470-485 (2015), NCBI Accession # 4XNY_H (235aa)	
HIV747	Heavy Chain	Vrc23	Georgiev, I.S., et al., Delineating antibody recognition in polyclonal sera from patterns of HIV-1 isolate neutralization; Science 340 (6133), 751-756 (2013), NCBI Accession # 4J6R_H (224aa)	8366
HIV748	Heavy chain	VRC-CH30	WO2013090644 SEQ ID NO: 22	8367
HIV749	Heavy Chain	Vrc-ch31	Zhou T et al., Immunity 39 (2), 245-258 (2013), NCBI Accession # 4LSP_H (236aa)	8368
HIV750	Heavy chain	VRC-CH32	Wu X. et al, " Focused evolution of HIV-1 neutralizing antibodies revealed by structures and deep sequencing" Science 333 (6049), 1593-1602 (2011), NCBI Accession # AEM62724	8369
HIV751	Heavy chain	VRC-CH33	WO2013090644 SEQ ID NO: 28	8370
HIV752	Heavy chain	VRC-CH34	WO2013090644 SEQ ID NO: 30	8371
HIV753	Heavy chain	VRCO7 G54H	US20140322163 SEQ ID NO: 33	8372
HIV754	Heavy chain	VRC-PG04	Wu X. et al, " Focused evolution of HIV-1 neutralizing antibodies revealed by structures and deep sequencing" Science 333 (6049), 1593-1602 (2011), NCBI Accession # AEM62752	8373
HIV755	Heavy chain	VRC-PG04b	WO2013090644 SEQ ID NO: 44	8374
HIV756	Heavy Chain	Vrc-pg20	Zhou T et al., Immunity 39 (2), 245-258 (2013), NCBI Accession # 4LSU_H (227aa)	8375
HIV757	Heavy chain	X5	US7378093B2 SEQ ID NO: 3	8376
HIV758	Heavy chain	X5	US8110192B2 SEQ ID NO: 5	8377
HIV759	Heavy chain	X5 variant	US7378093B2 SEQ ID NO: 11	8378
HIV760	Heavy Chain	Z13e1	Stanfield, R.L., et al, J. Mol. Biol. 414 (3), 460-476 (2011), NCBI Accession # 3Q1S_H(230aa)	8379
HIV761	Heavy Chain	Z258-vrc27.01	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDI_H(227aa)	8380
HIV762	Heavy Chain		NCBI Accession # 1N0X_K (230aa)	8381
HIV763	Heavy chain		US5804440A SEQ ID NO: 142	8382
HIV764	Heavy chain		US5804440A SEQ ID NO: 143	8383
HIV765	Heavy chain		US5804440A SEQ ID NO: 144	8384
HIV766	Heavy chain		US5804440A SEQ ID NO: 145	8385
HIV767	Heavy chain		US5804440A SEQ ID NO: 146	8386
HIV768	Heavy chain		US5804440A SEQ ID NO: 66	8387
HIV769	Heavy chain		US5804440A SEQ ID NO: 67	8388
HIV770	Heavy chain		US5804440A SEQ ID NO: 68	8389
HIV771	Heavy chain		US5804440A SEQ ID NO: 70	8390
HIV772	Heavy chain		US5804440A SEQ ID NO: 72	8391
HIV773	Heavy chain		US5804440A SEQ ID NO: 73	8392
HIV774	Heavy chain		US5804440A SEQ ID NO: 74	8393
HIV775	Heavy chain		US5804440A SEQ ID NO: 75	8394
HIV776	Heavy chain		US5804440A SEQ ID NO: 78	8395

HIV777	Heavy chain		WO2014063059 SEQ ID NO: 10	8396
HIV778	Heavy chain		WO2014063059 SEQ ID NO: 12	8397
HIV779	Heavy chain		WO2014063059 SEQ ID NO: 130	8398
HIV780	Heavy chain		WO2014063059 SEQ ID NO: 14	8399
HIV781	Heavy chain		WO2014063059 SEQ ID NO: 16	8400
HIV782	Heavy chain		WO2014063059 SEQ ID NO: 18	8401
HIV783	Heavy chain		WO2014063059 SEQ ID NO: 20	8402
HIV784	Heavy chain		WO2014063059 SEQ ID NO: 22	8403
HIV785	Heavy chain		WO2014063059 SEQ ID NO: 24	8404
HIV786	Heavy chain		WO2014063059 SEQ ID NO: 4	8405
HIV787	Heavy chain		WO2014063059 SEQ ID NO: 6	8406
HIV788	Heavy chain		WO2014063059 SEQ ID NO: 8	8407
HIV789	Heavy chain consensus		WO2014063059 SEQ ID NO: 2	8408
HIV790	Heavy chain constant region	G4D	US20130195881 SEQ ID NO: 6	8409
HIV791	Heavy chain constant region	G4H	US20130195881 SEQ ID NO: 5	8410
HIV792	Heavy chain constant region	MV1	US20130195881 SEQ ID NO: 7	8411
HIV793	Heavy chain constant region	TNX-355, Idalizumab	US20130195881 SEQ ID NO: 4	8412
HIV794	Heavy Chain Fab	Ch04	McLellan, J.S., et al., Nature 480 (7377), 336-343 (2011), NCBI Accession # 3U46_A (238aa)	8413
HIV795	Heavy Chain Of Anti-HIV Fab From Human 21c Antibody	21C	Diskin, R., et al., Nat. Struct. Mol. Biol. 17 (5), 608-613 (2010), NCBI Accession # 3LMJ_H (231aa)	8414
HIV796	Heavy Chain Of Anti-hiv-1 Gp120 V1v2 Antibody 830a	830a	Pan et al., J. Virol. 89 (15), 8003-8010 (2015), NCBI Accession # 4YWG_H (226aa)	8415
HIV797	Heavy chain partial	412D	Huang C. et al "Structural basis of tyrosine sulfation and VH-gene usage in antibodies that recognize the HIV type 1 coreceptor-binding site on gp120" Proc. Natl. Acad. Sci. U.S.A. 101 (9), 2706-2711 (2004), NCBI Accession # AAR88379	8416
HIV798	Heavy chain variable region	0.5γ(1C10)	US8722861B2 SEQ ID NO: 1	8417
HIV799	Heavy chain variable region	0.5δ (3D6)	US8722861B2 SEQ ID NO: 5	8418
HIV800	Heavy chain variable region	10J4 mAb	WO2015103549 SEQ ID NO: 3	8419
HIV801	Heavy chain variable region	10M6 mAb	WO2015103549 SEQ ID NO: 5	8420
HIV802	Heavy chain variable region	13110 mAb	WO2015103549 SEQ ID NO: 7	8421
HIV803	Heavy chain variable region	2N5mAb	WO2015103549 SEQ ID NO: 9	8422
HIV804	Heavy chain variable region	35022 mAb	WO2015103549 SEQ ID NO: 1	8423
HIV805	Heavy chain variable region	42F9	US8722861B2 SEQ ID NO: 7	8424
HIV806	Heavy chain variable region	4835_F12 (PGT-124)	US20140205612 SEQ ID NO: 404	8425

HIV807	Heavy chain variable region	4838_L06 (PGT-121)	US20140205612 SEQ ID NO: 66	8426
HIV808	Heavy chain variable region	4858_P08 (PGT-123)	US20140205612 SEQ ID NO: 167	8427
HIV809	Heavy chain variable region	4869-K15 (PGT-133)	US20140205612 SEQ ID NO: 419	8428
HIV810	Heavy chain variable region	4873_E03 (PGT-121)	US20140205612 SEQ ID NO: 62	8429
HIV811	Heavy chain variable region	4876_M06 (PGT-134)	US20140205612 SEQ ID NO: 434	8430
HIV812	Heavy chain variable region	4877_D15 (PGT-122)	US20140205612 SEQ ID NO: 155	8431
HIV813	Heavy chain variable region	4964_G22 (PGT-141), 4993_K13 (PGT-141)	US20140205612 SEQ ID NO: 275	8432
HIV814	Heavy chain variable region	4970_K22 (PGT-144)	US20140205612 SEQ ID NO: 306	8433
HIV815	Heavy chain variable region	4980_N08 (PGT-143)	US20140205612 SEQ ID NO: 297	8434
HIV816	Heavy chain variable region	4995_E20 (PGT-142)	US20140205612 SEQ ID NO: 291	8435
HIV817	Heavy chain variable region	4995_P16 (PGT-145)	US20140205612 SEQ ID NO: 400	8436
HIV818	Heavy chain variable region	49G2	US8722861B2 SEQ ID NO: 9	8437
HIV819	Heavy chain variable region	4O20mAb	WO2015103549 SEQ ID NO: 11	8438
HIV820	Heavy chain variable region	5114_A19 (PGT-128)	US20140205612 SEQ ID NO: 333	8439
HIV821	Heavy chain variable region	5120_N10 (PGT-139)	US20140205612 SEQ ID NO: 462	8440
HIV822	Heavy chain variable region	5131_A17 (PGT-132)	US20140205612 SEQ ID NO: 443	8441
HIV823	Heavy chain variable region	5136_H01 (PGT-131)	US20140205612 SEQ ID NO: 345	8442
HIV824	Heavy chain variable region	5138_G07 (PGT-138)	US20140205612 SEQ ID NO: 453	8443
HIV825	Heavy chain variable region	5141_B17 (PGT-126)	US20140205612 SEQ ID NO: 199	8444
HIV826	Heavy chain variable region	5145_B14 (PGT-127)	US20140205612 SEQ ID NO: 318	8445
HIV827	Heavy chain variable region	5147_N06 (PGT-130)	US20140205612 SEQ ID NO: 215	8446
HIV828	Heavy chain variable region	5329_C19 (PGT-136), 5366_P21 (PGT-136)	US20140205612 SEQ ID NO: 248	8447
HIV829	Heavy chain variable region	5343_B08 (PGT-135), 5344_E16 (PGT-135)	US20140205612 SEQ ID NO: 231	8448
HIV830	Heavy chain variable region	5345_I01 (PGT-137)	US20140205612 SEQ ID NO: 362	8449
HIV831	Heavy chain variable region	5G2	US8722861B2 SEQ ID NO: 3	8450
HIV832	Heavy chain variable region	6808_B09 (PGT-156)	US20140205612 SEQ ID NO: 546	8451

HIV833	Heavy chain variable region	6831_A21 (PGT-151)	US20140205612 SEQ ID NO: 473	8452
HIV834	Heavy chain variable region	6843_G20 (PGT-154)	US20140205612 SEQ ID NO: 516	8453
HIV835	Heavy chain variable region	6881_N05 (PGT-158).	US20140205612 SEQ ID NO: 572	8454
HIV836	Heavy chain variable region	6889_I17 (PGT-152)	US20140205612 SEQ ID NO: 489	8455
HIV837	Heavy chain variable region	6891_F06 (PGT-153)	US20140205612 SEQ ID NO: 501	8456
HIV838	Heavy chain variable region	6892_C23 (PGT-157)	US20140205612 SEQ ID NO: 559	8457
HIV839	Heavy chain variable region	6892_D19 (PGT-155)	US20140205612 SEQ ID NO: 531	8458
HIV840	Heavy chain variable region	7B9mAb	WO2015 103549 SEQ ID NO: 13	8459
HIV841	Heavy chain variable region	7K3mAb	WO2015 103549 SEQ ID NO: 15	8460
HIV842	Heavy chain variable region	B4	US7872110B2 SEQ ID NO: 2	8461
HIV843	Heavy chain variable region	B4DIVHv.l	US78721 10B2 SEQ ID NO: 5	8462
HIV844	Heavy chain variable region	B4DIVHv.2	US78721 10B2 SEQ ID NO: 6	8463
HIV845	Heavy chain variable region	B4DIVHv.3	US78721 10B2 SEQ ID NO: 7	8464
HIV846	Heavy chain variable region	B4DIVHV.4	US78721 10B2 SEQ ID NO: 8	8465
HIV847	Heavy chain variable region	bI2 IgA2 antibody	WO20 14040024 SEQ ID NO: 29	8466
HIV848	Heavy chain variable region	CHμ39.1	US5773247 SEQ ID NO: 10	8467
HIV849	Heavy chain variable region	CHμ5.5	US5773247 SEQ ID NO: 14	8468
HIV850	Heavy chain variable region	F425-Alg8 antibody	WO20 14040024 SEQ ID NO: 9	8469
HIV851	Heavy chain variable region	Fab 43	US20090191216 SEQ ID NO: 8	8470
HIV852	Heavy chain variable region	HGN194	US20110212106 SEQ ID NO: 45	8471
HIV853	Heavy chain variable region	HJ16	US20110212106 SEQ ID NO: 53	8472
HTV854	Heavy chairs variable region	HK20	US20110212106 SEQ ID NO: 29	8473
HIV855	Heavy chain variable region	IgA antibody	WO2014040024 SEQ ID NO: 11	8474
HTV856	Heavy chairs variable region	L1719A I1	US20150158934 SEQ ID NO: 175	8475
HIV857	Heavy chain variable region	L17 19A12	US20150158934 SEQ ID NO: 176	8476
HTV858	Heavy chairs variable region	L1719A9	US20150158934 SEQ ID NO: 174	8477
HIV859	Heavy chain variable region	L17 19B12	US20150158934 SEQ ID NO: 177	8478
HTV860	Heavy chairs variable region	L1719C1	US20 150 158934 SEQ ID NO: 178	8479
HIV86 1	Heavy chain variable region	L17191) 10	US20150158934 SEQ ID NO: 179	8480

HEV862	**Heavy** chain variable region	L17 19E1	**US201 50158934** SEQ ID NO: 180	8481
HIV863	Heavy chain variable region	**L17 19E11**	**US20150158934** SEQ **ID** NO: **18 1**	8482
HEV864	**Heavy** chain variable region	L17 19E12	**US201 50158934** SEQ ID NO: 182	8483
HIV865	Heavy chain variable region	**L1719F1 1**	**US20150158934** SEQ ID NO: 183	8484
HIV866	**Heavy chain variable** region	L1719H10	**US20150158934** SEQ ID NO: 185	8485
HIV867	Heavy chain variable region	**L1719H9**	**US20150158934** SEQ ID NO: 184	8486
HIV868	**Heavy chain variable** region	L1720C1	**US20150158934** SEQ ID NO: 186	8487
HIV869	Heavy chain variable region	L1720E4	**US20150158934** SEQ ID NO: 187	8488
HIV870	**Heavy chain variable** region	**L1721A3**	**US20150158934** SEQ ID NO: 188	8489
HIV871	Heavy chain variable region	L1721A5	**US20150158934** SEQ ID NO: 189	8490
HIV872	**Heavy chain variable** region	**L1721A8**	**US20150158934** SEQ ID NO: 190	8491
HIV873	Heavy chain variable region	**L172 1H4**	**US20150158934** SEQ ID NO: **191**	8492
HIV874	Heavy **chain variable** region	**L1723A10**	**US20150158934** SEQ ID NO: 193	8493
HIV875	Heavy chain variable region	**L1723A11**	**US20150158934** SEQ ID NO: 194	8494
HIV876	Heavy **chain variable** region	L1723A9	**US20150158934** SEQ ID NO: 192	8495
HIV877	Heavy chain variable region	**L1723E5**	**US20150158934** SEQ ID NO: 195	8496
HIV878	Heavy **chain variable** region	L23 **19G11**	**US20150158934** SEQ ID NO: 197	8497
HIV879	Heavy chain variable region	**L23 19G7**	**US20150158934** SEQ ID NO: **196**	8498
HIV880	Heavy **chain variable** region	L23 19H7	**US20150158934** SEQ ID NO: 198	8499
HIV881	Heavy chain variable region	**L2320E9**	**US20150158934** SEQ ID NO: 199	8500
HIV882	Heavy chain variable region	**L2320F9**	**US20150158934** SEQ **ID NO:** 200	8501
HIV883	Heavy chairs variable region	**L232 IB7**	**US20150 158934** SEQ ID **NO:** 201	8502
HIV884	Heavy chain variable region	**L2321H6**	**US20150158934** SEQ **ID NO:** 202	8503
HIV885	Heavy chairs variable region	**L8 1C11**	US20150158934 SEQ ID **NO:** 15	8504
HIV886	Heavy chain variable region	**L81C9**	**US20150158934** SEQ **ID NO:** 30	8505
HTV887	Heavy chairs variable region	**L8 1D9**	**US20150 158934** SEQ ID NO: 10	8506
HIV888	Heavy chain variable region	**L81E1**	**US20150158934** SEQ ID **NO:** 18	8507
HTV889	Heavy chairs variable region	**L8 1E7**	**US20150** 158934 SEQ ID **NO:** 16	8508
HIV890	Heavy chain variable region	**L81F1**	**US20150158934** SEQ **ID** NO: **19**	8509

HIV891	**Heavy** chain variable region	**L81G7**	**US201 50158934** SEQ ID NO: **13**	8510
HIV892	Heavy chain variable region	**L81H1**	**US20150158934** SEQ **ID** NO: 98	851 1
HEV893	**Heavy** chain variable region	**L81H2**	**US201 50158934** SEQ ID NO: 23	8512
HIV894	Heavy chain variable region	L81H7	**US20150158934** SEQ ID NO: 11	8513
HIV895	**Heavy chain variable** region	L81H9	**US20150158934** SEQ ID NO: 28	8514
HIV896	Heavy chain variable region	**L82B12A**	US20150158934 SEQ ID NO: 105	8515
HIV897	**Heavy chain variable** region	L82B1A	**US20150158934** SEQ ID NO: 99	8516
HIV898	Heavy chain variable region	L82B1D	**US20150158934** SEQ ID NO: 100	8517
HIV899	**Heavy chain variable** region	**L82B2A**	**US20150158934** SEQ ID NO: 101	8518
HIV900	Heavy chain variable region	L82B3F	**US20150158934** SEQ ID NO: 102	8519
HIV901	**Heavy chain variable** region	L82B4A	**US20150158934** SEQ ID NO: 103	8520
HIV902	Heavy chain variable region	**L82B4E**	**US20 150 158934** SEQ ID NO: 104	852 1
HIV903	Heavy **chain variable** region	L82B4F	**US20150158934** SEQ ID NO: 2 1	8522
HIV904	Heavy chain variable region	**L832G6**	**US20150158934** SEQ ID NO: 1 **13**	8523
HIV905	Heavy **chain variable** region	L833E1	**US20150158934** SEQ ID NO: 72	8524
HIV906	Heavy chain variable region	**L833F5**	**US20 150 158934** SEQ ID NO: 17	8525
HIV907	Heavy **chain variable** region	L833H I	**US20150158934** SEQ ID NO: **114**	8526
HIV908	Heavy chain variable region	**L833H3**	**US20150158934** SEQ ID NO: 1 **15**	8527
HIV909	Heavy **chain variable** region	L88B10B	**US20150158934** SEQ ID NO: 27	8528
HIV9 10	Heavy chain variable region	**L88B1 1B**	**US20150158934** SEQ ID NO: 12	8529
HIV9I !	Heavy chain variable region	**L88B12G**	**US20150158934** SEQ ID **NO:** 29	8530
HEV9 12	Heavy chairs variable region	**L88B I D**	**US20 I 50 I** 58934 SEQ ID **NO:** 20	853 1
HIV913	Heavy chain variable region	**L88B2A**	**US20150158934** SEQ ID **NO: 106**	8532
HTV9 I4	Heavy chairs variable region	L88FA2	**US20 I 50 158934** SEQ ID **NO: 26**	8533
HIV915	Heavy chain variable region	**L88FA3**	**US20150158934** SEQ ID **NO: 107**	8534
HEV9 16	Heavy chairs variable region	L88FA5	US20 I 50 158934 SEQ ID NO: 108	8535
HIV917	Heavy chain variable region	**L88FB !**	**US20150158934** SEQ ID **NO:** 25	8536
HTV9 I8	Heavy chairs variable region	**L88FC 1 I**	US20 150 158934 SEQ ID **NO:** 22	8537
HIV919	Heavy chain variable region	**L88FD 12**	US20150158934 SEQ **ID** NO: 24	8538

HEV920	Heavy chain variable region	L89B12D	US201 50158934 SEQ ID NO: 112	8539
HIV92 1	Heavy chain variable region	L89B1D	US20150158934 SEQ ID NO: 109	8540
HEV922	Heavy chain variable region	L89B2C	US201 50158934 SEQ ID NO: 110	8541
HIV923	Heavy chain variable region	L89B3E	US20150158934 SEQ ID NO: 14	8542
HIV924	Heavy chain variable region	L89B6B	US20150158934 SEQ ID NO: 111	8543
HIV925	Heavy chain variable region	L8Cb15	US20150158934 SEQ ID NO: 116	8544
HIV926	Heavy chain variable region	L8CJ3	US20150158934 SEQ ID NO: 73	8545
HIV927	Heavy chain variable region	L8Fe2	US20150158934 SEQ ID NO: 117	8546
HIV928	Heavy chain variable region	L8Fgl2	US20150158934 SEQ ID NO: 118	8547
HIV929	Heavy chain variable region	L8FJ19	US20150158934 SEQ ID NO: 119	8548
HIV930	Heavy chain variable region	L8Fol7	US20150158934 SEQ ID NO: 120	8549
HIV93 1	Heavy chain variable region	L8Fp6	US20150158934 SEQ ID NO: 12 1	8550
HIV932	Heavy chain variable region	L8Hi20	US20150158934 SEQ ID NO: 122	855 1
HIV933	Heavy chain variable region	L91 1B 11E	US20150158934 SEQ ID NO: 140	8552
HIV934	Heavy chain variable region	L9UB12B	US20150158934 SEQ ID NO: 7 1	8553
HIV935	Heavy chain variable region	L91 1B 1E	US20150158934 SEQ ID NO: 137	8554
HIV936	Heavy chain variable region	L9UB1G	US20150158934 SEQ ID NO: 65	8555
HIV937	Heavy chain variable region	L91 1B2E	US20150158934 SEQ ID NO: 138	8556
HIV938	Heavy chain variable region	L911B3D	US20150158934 SEQ ID NO: 75	8557
HIV939	Heavy chain variable region	L91 1B9A	US20150158934 SEQ ID NO: 139	8558
HIV940	Heavy chain variable region	L91 1F12B	US20150158934 SEQ ID NO: 142	8559
HIV941	Heavy chairs variable region	L911F 1B	US20 150 158934 SEQ ID NO: 141	8560
HIV942	Heavy chain variable region	L91IF1F	US20150158934 SEQ ID NO: 77	8561
HTV943	Heavy chairs variable region	L9UF4C	US20150 158934 SEQ ID NO: 33	8562
HIV944	Heavy chain variable region	L91A1	US20150158934 SEQ ID NO: 123	8563
HEV945	Heavy chairs variable region	L91B5	US20150158934 SEQ ID NO: 37	8564
HIV946	Heavy chain variable region	L91B5, 4A7	US20150158934 SEQ ID NO: 97	8565
HTV947	Heavy chairs variable region	L91B5, A12	US20150 158934 SEQ ID NO: 92	8566
HIV948	Heavy chain variable region	L91B5, A4	US20150158934 SEQ ID NO: 90	8567

HEV949	**Heavy** chain variable region	**L91B5,** A7	**US20 150158934** SEQ ID NO: 91	8568
HIV950	Heavy chain variable region	**L91B5, B2**	**US20150158934** SEQ ID NO: 93	8569
HEV95 1	**Heavy** chain variable region	**L91B5, D4**	**US20 150158934** SEQ ID NO: 94	8570
HIV952	Heavy chain variable region	**L91B5, F 11**	**US20150158934** SEQ ID NO: 96	8571
HIV953	**Heavy chain variable** region	**L91B5,** F4	**US20150158934** SEQ ID NO: 95	8572
HIV954	Heavy chain variable region	L91C2	**US20150158934** SEQ ID NO: **6 1**	8573
HIV955	**Heavy chain variable** region	**L91E1**	**US20150158934** SEQ ID NO: 45	8574
HIV956	Heavy chain variable region	L91E2	**US20150158934** SEQ ID NO: 124	8575
HIV957	**Heavy chain variable** region	**L91F10**	**US20150158934** SEQ ID NO: 69	8576
HIV958	Heavy chain variable region	L91G2	**US20150158934** SEQ ID NO: 64	8577
HIV959	**Heavy chain variable** region	L91H3	**US20150158934** SEQ ID NO: 128	8578
HIV960	Heavy chain variable region	**L91H9**	**US20150158934** SEQ ID NO: 41	8579
HIV96 1	Heavy **chain variable** region	L922B2	**US20150158934** SEQ ID NO: 143	8580
HIV962	Heavy chain variable region	**L922B4**	**US20150158934** SEQ ID NO: 144	8581
HIV963	Heavy **chain variable** region	L922E1	**US20150158934** SEQ ID NO: 145	8582
HIV964	Heavy chain variable region	**L922E2**	**US20 150 158934** SEQ ID NO: 53	8583
HIV965	Heavy **chain variable** region	L923A 1	**US20150158934** SEQ ID NO: 146	8584
HIV966	Heavy chain variable region	L923A4	**US20 150 158934** SEQ ID NO: 32	8585
HIV967	Heavy **chain variable** region	L92A1 **1**	**US20150158934** SEQ ID NO: 125	8586
HIV968	Heavy chain variable region	**L92C7**	**US20150158934** SEQ ID NO: 62	8587
HIV969	Heavy chain variable region	**L92D4**	**US20150158934** SEQ **ID NO:** 126	8588
HTV970	Heavy chairs variable region	**L92E6**	**US20** 150158934 SEQ ID **NO:** 63	8589
HIV97 1	Heavy chain variable region	**L92E7**	**US20150158934** SEQ ID **NO:** 74	8590
HTV972	Heavy chairs variable region	**L92E7,** A1	**US20** 150 158934 SEQ ID **NO:** 85	8591
HIV973	Heavy chain variable region	**L92E7,** A2	**US20150158934** SEQ **ID NO:** 86	8592
HTV974	Heavy chairs variable region	**L92E7,** A3	**US20** 150 158934 SEQ ID **NO:** 87	8593
HIV975	Heavy chain variable region	**L92E7,** A4	**US20150158934** SEQ **ID NO:** 80	8594
HTV976	Heavy chairs variable region	**L92E7,** A4	**US20** 150 158934 SEQ ID **NO:** 88	8595
HIV977	Heavy chain variable region	L92E7. A5	**US20150158934** SEQ ID NO: 89	8596

HEV978	**Heavy** chain variable region	L92E7, B5	**US20 150158934** SEQ ID NO: 78	8597
HIV979	Heavy chain variable region	L92E7. **C**	**US20150158934** SEQ **ID** NO: 79	8598
HEV980	**Heavy** chain variable region	L92E7, C3	**US20 150158934** SEQ ID NO: 82	8599
HIV981	Heavy chain variable region	L92E7, D3	**US20150158934** SEQ ID NO: 83	8600
HIV982	**Heavy chain variable** region	**L92E7,** E1	**US20150158934** SEQ ID NO: 84	8601
HIV983	Heavy chain variable region	L92E7, G4	**US20150158934** SEQ ID NO: 81	8602
HIV984	**Heavy chain variable** region	**L932A9**	**US20150158934** SEQ ID NO: 58	8603
HIV985	Heavy chain variable region	L932E10	**US20150158934** SEQ ID NO: 35	8604
HIV986	**Heavy chain variable** region	L932E8	**US20150158934** SEQ ID NO: 147	8605
HIV987	Heavy chain variable region	L932G9	**US20150158934** SEQ ID NO: 34	8606
HIV988	**Heavy chain variable** region	L933D10	**US20150158934** SEQ ID NO: 50	8607
HIV989	Heavy chain variable region	L93B3	**US20150158934** SEQ ID NO: 70	8608
HIV990	Heavy **chain variable** region	L93B4	**US20150158934** SEQ ID NO: 127	8609
HIV991	Heavy chain variable region	L93C3	**US20150158934** SEQ ID NO: **51**	8610
HIV992	Heavy **chain variable** region	L93C6	**US20150158934** SEQ ID NO: 67	8611
HIV993	Heavy chain variable region	**L93D3**	**US20150158934** SEQ ID NO: 129	8612
HIV994	Heavy **chain variable** region	L93D4	**US20150158934** SEQ ID NO: 43	8613
HIV995	Heavy chain variable region	**L93D9**	**US20150158934** SEQ ID NO: 130	8614
HIV996	Heavy **chain variable** region	L93E3	**US20150158934** SEQ ID NO: 55	8615
HIV997	Heavy chain variable region	**L93E6**	**US20150158934** SEQ ID NO: 131	8616
HIV998	Heavy chain variable region	**L93F12**	**US20150158934** SEQ **ID NO: 133**	**8617**
HTV999	Heavy chairs variable region	L93F2	**US20**150 158934 SEQ ID NO: 132	8618
HIV1000	Heavy chain variable region	**L93F2**	**US20150158934** SEQ ID **NO:** 59	**8619**
HIVlOO1	Heavy chairs variable region	L93H6	**US20150 158934** SEQ ID NO: 38	8620
HIV1002	Heavy chain variable region	**L93H9**	**US20150158934** SEQ ID **NO: 134**	8621
HEV1003	Heavy chairs variable region	**L94A 12**	**US20150158934** SEQ ID **NO:** 46	8622
HIV1004	Heavy chain variable region	**L94C2**	**US20150158934** SEQ **ID NO:** 31	8623
HTV1005	Heavy chairs variable region	**L94D 12**	**US20150158934** SEQ ID **NO:** 42	8624
HIV1006	Heavy chain variable region	**L94D4**	**US20150158934** SEQ **ID** NO: 47	8625

HEV1007	**Heavy** chain variable region	L94E3	**US20 150158934** SEQ ID NO: 39	8626
HIV1008	Heavy chain variable region	L94E4	US20150158934 SEQ **ID** NO: 54	8627
HEV1009	**Heavy** chain variable region	L94E5	**US20 150158934** SEQ ID NO: 57	8628
HIV1010	Heavy chain variable region	L94H1	**US20150158934** SEQ ID NO: 36	8629
HIV101 1	**Heavy chain variable** region	**L94H2**	US20150158934 SEQ ID NO: 40	8630
HIV1012	Heavy chain variable region	L94H5	**US20150158934** SEQ ID NO: 48	863 1
HIV1013	**Heavy chain variable** region	**L94H7**	**US20150158934** SEQ ID NO: 135	8632
HIV1014	Heavy chain variable region	**L95B10D**	US20150158934 SEQ ID NO: 136	8633
HIV1015	**Heavy chain variable** region	L95B12A	**US20150158934** SEQ ID NO: 68	8634
HIV1016	Heavy chain variable region	L95B12E	**US20150158934** SEQ ID NO: 66	8635
HIV1017	**Heavy chain variable** region	L95B8A	**US20150158934** SEQ ID NO: 60	8636
HIV1018	Heavy chain variable region	**L98FB 10**	**US20150158934** SEQ ID NO: 76	8637
HIV10 19	Heavy **chain variable** region	L9Ab16	**US20150158934** SEQ ID NO: 148	8638
HIV1020	Heavy chain variable region	L9AM9	**US20150158934** SEQ ID NO: 149	8639
HIV1021	Heavy **chain variable** region	**L9Adl3**	**US20150158934** SEQ ID NO: 15 1	8640
HIV1022	Heavy chain variable region	**L9Adl4**	**US20150158934** SEQ ID NO: 152	8641
HIV1023	Heavy **chain variable** region	**L9Ad3**	**US20150158934** SEQ ID NO: 150	8642
HIV1024	Heavy chain variable region	L9AJ2	**US20150158934** SEQ ID NO: 153	8643
HIV1025	Heavy **chain variable** region	**L9An7**	**US20150158934** SEQ ID NO: 154	8644
HIV1026	Heavy chain variable region	L9AoS 5	**US20150158934** SEQ ID NO: 155	8645
HIV1027	Heavy chain variable region	**L9Apl !**	**US20150158934** SEQ ID **NO:** 156	8646
HTV1028	Heavy chairs variable region	**L9Bb3**	US20150 158934 SEQ ID **NO:** 157	8647
HIV1029	Heavy chain variable region	**L9Bc6**	**US20150158934** SEQ ID **NO:** 158	8648
HTV1030	Heavy chairs variable region	**L9Bd8**	US20150 158934 SEQ **ID NO:** 159	8649
HIV1031	Heavy chain variable region	**L9Bd9**	**US20150158934** SEQ ID **NO:** 160	8650
HTV1032	Heavy chairs variable region	**L9Be l l**	**US20150 158934** SEQ **ID** NO: 161	865 1
HIV1033	Heavy chain variable region	**L9Bf 1** 1	**US20150158934** SEQ **ID NO:** 49	8652
HTV1034	Heavy chairs variable region	**L9Bfl 9**	**US20150158934** SEQ **ID NO:** 162	8653
HIV1035	Heavy chain variable region	L9BJ13	**US20150158934** SEQ **ID** NO: **163**	8654

HIV1036	Heavy chain variable region	L9Bm10	US20150158934 SEQ ID NO: 164	8655
HIV1037	Heavy chain variable region	L9Bm16	US20150158934 SEQ ID NO: 56	8656
HIV1038	Heavy chain variable region	L9Bp16	US20150158934 SEQ ID NO: 165	8657
HIV1039	Heavy chain variable region	L9Bp5	US20150158934 SEQ ID NO: 44	8658
HIV1040	Heavy chain variable region	L9Ca12	US20150158934 SEQ ID NO: 166	8659
HIV1041	Heavy chain variable region	L9Ca13	US20150158934 SEQ ID NO: 167	8660
HIV1042	Heavy chain variable region	L9Cd12	US20150158934 SEQ ID NO: 168	8661
HIV1043	Heavy chain variable region	L9Cf15	US20150158934 SEQ ID NO: 169	8662
HIV1044	Heavy chain variable region	L9Cl22	US20150158934 SEQ ID NO: 52	8663
HIV1045	Heavy chain variable region	L9Cm18	US20150158934 SEQ ID NO: 170	8664
HIV1046	Heavy chain variable region	L9Co22	US20150158934 SEQ ID NO: 171	8665
HIV1047	Heavy chain variable region	L9Cp5	US20150158934 SEQ ID NO: 172	8666
HIV1048	Heavy chain variable region	L9Cpl3	US20150158934 SEQ ID NO: 173	8667
HIV1049	Heavy chain variable region	Makandal monoclonal antibody (Mmab)	US20100111990 SEQ ID NO: 4	8668
HIV1050	Heavy chain variable region	NM-01	US5665569 SEQ ID NO: 17	8669
HIV1051	Heavy chain variable region	NM-01 HuVH	US5665569 SEQ ID NO: 27	8670
HIV1052	Heavy chain variable region	NM-01 HuVK	US5665569 SEQ ID NO: 29	8671
HIV1053	Heavy chain variable region	NM-01 HuVKF	US5665569 SEQ ID NO: 31	8672
HIV1054	Heavy chain variable region	PGT125	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14393	8673
HIV1055	Heavy chain variable region	PGT126	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14394	8674
HIV1056	Heavy chain variable region	PGT131	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14389	8675
HIV1057	Heavy chain variable region	PGT136	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14400	8676
HIV1058	Heavy chain variable region	PGT137	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14401	8677

HEV1059	**Heavy** chain variable region	PGT141	**Walker** L.M. **et** al "Broad neutralization coverage of HIV by multiple **highly** potent antibodies", Nature 477 (7365), 466-470 **(201 1),** NCBI Accession # **AEN14402**	8678
HIV1060	Heavy **chain** variable region	PGT142	Walker L.M. **et al** "Broad neutralization coverage **of HIV by multiple** highly potent antibodies", Nature 477 (7365), 466-470 **(201 1), NCBI** Accession # AEN14368	8679
HIV1061	Heavy chain variable region	PGT143	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 **(201 1),** NCBI Accession # AEN 14404	8680
HIV1062	Heavy chain variable region	**PGT144**	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies". Nature 477 (7365), 466-470 (201 **1),** NCBI Accession # AEN14405	8681
HIV1063	**Heavy chain variable** region	PGT151	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a Glycan-Dependent Epitope on the Perfusion **Conformation** of gp41 on Cleaved Envelope Trimers" **Immunity** 40 (5), 657-668 (2014), NCBI Accession # AIC3535	8682
HIV1064	Heavy chain variable region	PGT152	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a **Glycan-Dependent** Epitope on the Perfusion Conformation of **gp41** on Cleaved Envelope Trimers" **Immunity** 40 (5), 657-668 (2014), NCBI Accession # AIC32536	8683
HEV1065	**Heavy** chain variable region	**PGT153**	Falkowska, E. et al "Broadly Neutralizing **HIV** Antibodies Define **a** Glycan-Dependent Epitope on the Perfusion Conformation of **gp41** on Cleaved Envelope Trimers" **Immunity 40** (5), 657-668 (2014), NCBI Accession # AIC32537	8684
HIV1066	Heavy chain variable region	**PGT154**	Falkowska, E. et **al** "Broadly Neutralizing **HIV** Antibodies Define a Glycan-Dependent Epitope on the Perfusion Conformation of **gp41** on Cleaved Envelope **Trimers" Immunity** 40 (5), 657-668 (2014), NCBI Accession # AEC32521	8685
HIV1067	Heavy chairs variable region	**PGT I55**	Falkowska, E. et al "Broadly Neutralizing **HIV** Antibodies Define **a** Glycan-Dependent Epitope on the **Perfusion** Conformation **of** gp41 on Cleaved Envelope Trimers" Immunity 40 (5), 657-668 (2014), NCBI Accession # AIC32539	8686
HIV1068	Heavy **chain variable** region	PGT156	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define **a** Glycan-Dependent Epitope on the Perfusion Conformation **of gp41** on Cleaved Envelope Trimers" **Immunity** 40 (5), 657-668 (2014), NCBI Accession # AIC32540	8687
HIV1069	Heavy chain variable region	PGT157	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a Glycan-Dependent Epitope on the Perfusion **Conformation of gp41** on Cleaved Envelope Trimers" Immunity 40 (5), 657-668 (2014), NCBI Accession # AIC32541	8688
HIV1070	**Heavy chain variable** region	PGT158	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a Glycan-Dependent Epitope on the Perfusion Conformation of gp41 on Cleaved Envelope Trimers" **Immunity** 40 (5), 657-668 (2014), NCBI **Accession # AIC32542**	8689
HIV **1071**	Heavy chain variable region	**rF105**	**WO1993012232** SEQ ID NO: 4	8690
HEV1072	**Heavy** chain variable region	**ScFvX5-**CD4	US**7378093B2** SEQ ID NO: 14	8691

HEV1073	Heavy chain variable region	TNX-355, Idaliznniab	US20130195881 SEQ ID NO: 3	8692
HIV1074	Heavy chain variable region	VCR 14	US20150044137 SEQ ID NO: 13	8693
HEV1075	Heavy chain variable region	VCR 14b	US20150044137 SEQ ID NO: 14	8694
HIV1076	Heavy chain variable region	VCR 14c	US20150044137 SEQ ID NO: 15	8695
HIV1077	Heavy chain variable region	VCR 16	US20150044137 SEQ ID NO: 29	8696
HIV1078	Heavy chain variable region	VCR16b	US20150044137 SEQ ID NO: 30	8697
HIV1079	Heavy chain variable region	VCR 16c	US20150044137 SEQ ID NO: 31	8698
HIV1080	Heavy chain variable region	VCR16d	US20150044137 SEQ ID NO: 32	8699
HIV1081	Heavy chain variable region	VLP_A14	US20150158934 SEQ ID NO: 203	8700
HIV1082	Heavy chain variable region	VLP_B9	US20150158934 SEQ ID NO: 204	8701
HIV1083	Heavy chain variable region	VLP3_B21	US20150158934 SEQ ID NO: 205	8702
HIV1084	Heavy chain variable region	VRC13	US20150044137 SEQ ID NO: 5	8703
HIV1085	Heavy chain variable region	VRC13b	US20150044137 SEQ ID NO: 6	8704
HIV1086	Heavy chain variable region	VRC13c	US20150044137 SEQ ID NO: 7	8705
HIV1087	Heavy chain variable region	VRC13d	US20150044137 SEQ ID NO: 8	8706
HIV1088	Heavy chain variable region	VRC13e	US20150044137 SEQ ID NO: 9	8707
HIV1089	Heavy chain variable region	VRC13f	US20150044137 SEQ ID NO: 10	8708
HIV1090	Heavy chain variable region	VRC13g	US20150044137 SEQ ID NO: 11	8709
HIV1091	Heavy chain variable region	VRC13h	US20150044137 SEQ ID NO: 12	8710
HIV1092	Heavy chain variable region	VRC15	US20150044137 SEQ ID NO: 16	8711
HIV1093	Heavy chain variable region		US20150041190 SEQ ID NO: 56	8712
HIV1094	Heavy chairs variable region, partial	P7	NCBI Accession # AAB41043.1 (136aa)	8713
HIV1095	Heavy Chain, Fab	Ch04	McLelian, J.S. et al., Structure of HIV-1 gpl20 V1 V2 domain with broadly neutralizing antibody PG9; Nature 480 (7377), 336-343 (2011), NCBI Accession # 3TCL_A (237aa)	8714
HIV1096	Heavy Chain, Fab	N5-i5	Acharya, P., et al., Structural Definition of an Antibody-Dependent Cellular Cytotoxicity Response Implicated in Reduced Risk for HIV-1 Infection; J. Virol. 88 (21), 12895-12906 (2014), NCBI Accession # 4H8W_H (226aa)	8715
HIV 1097	Heavy Chain, Fab	N60-i3	Gohain, N., et al., Cocrystal Structures of Antibody N60-i3 and Antibody JR4 in Complex with gpl20 Define More Cluster A Epitopes Involved in Effective Antibody-Dependent Effector Function against HIV-1; J.	8716

			Virol. 89 (17), 8840-8854 (2015), NCBE Accession # **4RFO_H (229aa)**	
HIV1098	Heavy Chain, Ig **Gamma-1 Chain C** Region	**Nih45-46 Fab**	Diskin, **R.,** et **al.,** Science 334 (6060), **1289-**1293 (2011), **NCBI** Accession # 3U7Y_**H (229aa)**	8717
HIV1099	**Heavy** Chain, Ig **Gamma-1 Chain C** Region	Pgt127	Pejchal, R., et al., Science 334 (6059), **1097-**1103 **(2011),** NCBI Accession # **3TWC_H(239aa)**	8718
HTV1 I00	Heavy Chain, Ig **Gamma-1** Chain C Region	**Pgt128**	Pejchal, **R.,** et al, Science 334 (6059), **1097-**1103 **(2011),** NCBI Accession # 3TV3_**H(239aa)**	8719
HIV1101	**HIV,** heavy chain	**Suvizumab**		8720
HIV1 102	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 43	8721
HIV1 103	HIV1 gpl20 antibody, heavy **chain**	**HIV1 gp120** antibody	**WO2001000678** SEQ ID NO: 44	8722
HIV1 104	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 45	8723
HIV1 105	HIV1 gpl20 antibody, heavy **chain**	**HIV1 gp120** antibody	**WO2001000678** SEQ ID NO: 46	8724
HIV1 106	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 47	8725
HIV1 107	HIV1 gpl20 antibody, heavy **chain**	**HIV1 gp120** antibody	**WO2001000678** SEQ ID NO: 48	8726
HIV1 108	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 49	8727
HIV1 109	HIV1 gpl20 antibody, heavy chain	**HIV1 gp120** antibody	**WO2001000678** SEQ ID NO: 57	8728
HIV1110	**HIV1** gp120 antibody, heavy chain	**HIV1** gp120 antibody	**WO2001000678** SEQ ID NO: 58	8729
HIV111 **U**	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 59	8730
HIV1112	**HIV1** gp120 antibody, heavy chain	**HIV1** gp120 antibody	**WO2001000678** SEQ ID NO: 60	8731
HIVU 13	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: **61**	8732
HIV1114	**HIV1** gp120 antibody, heavy chain	**HIV1** gp120 antibody	**WO2001000678** SEQ ID NO: 62	8733
HIVU 15	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 63	8734
HIV1116	**HIV1** gp120 antibody, heavy chain	**HIV1** gp120 antibody	**WO2001000678** SEQ ID NO: 64	8735
HIVU 17	**HIV1** gp120 antibody, heavy chain	**HIV1** gpl20 antibody	**WO2001000678** SEQ ID NO: 73	8736
HTVl118	**HIV1 gp120** antibody, heavy chain	HTVl **gp120** antibody	**WO200 1000678** SEQ **ID NO:** 74	8737
HIV1119	HIV1 **gp120** antibody, heavy chain	**HIV1 gp120** antibody	**WO2001000678** SEQ **ID** NO: 75	8738
HTVl 120	**HIV1 gp120** antibody, heavy chain	HIV1 **gp120** antibody	**WO200 1000678** SEQ **ID NO:** 76	8739
HIV1121	HTVl **gp120** antibody, heavy chain	**HIV1 gp120** antibody	**WO2001000678** SEQ **ID** NO: 77	8740
HTVl122	**HIV1 gp120** antibody, heavy chain	HTVl **gp120** antibody	**WO200 1000678** SEQ **ID NO:** 78	8741
HIV1123	HIV1 **gp120** antibody, **light** chain	**HIV1 gp120** antibody	**WO200 1000678** SEQ **ID** NO: 50	8742
HTVl124	**HIV1 gp120** antibody, light chain	HIV1 **gp120** antibody	**WO200 1000678** SEQ **ID NO:** 51	8743

HEV1 125	**HIV1 gpl20** antibody, light chain	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 52	8744
HIV l l26	HIVl **gpl20** antibody, light chain	HlVl **gpl20** antibody	**WO2001000678** SEQ ID **NO:** 53	8745
H BV l l27	**HIV1 gpl20** antibody, light chain	**HIVl** gpl20 antibody	**WO2001000678** SEQ ID NO: 54	8746
HIV1 128	**HIV1** gpl20 antibody, light **chain**	HIVl **gpl20** antibody	WO2001000678 SEQ ID NO: 55	8747
HIV1 129	**HIV1** gpl20 antibody, light chain	HIVl **gpl20** antibody	**WO2001000678** SEQ ID NO: 56	8748
HIVl 130	**HIV1** gpl20 antibody, light **chain**	HIVl **gpl20** antibody	WO2001000678 SEQ ID NO: 65	8749
HIV1 131	**HIV1** gpl20 antibody, light chain	HIVl **gpl20** antibody	**WO2001000678** SEQ ID NO: 66	8750
HIV1 132	**HIV1** gpl20 antibody, light **chain**	HIVl **gpl20** antibody	WO2001000678 SEQ ID NO: 67	8751
HIV1 133	**HIV1** gpl20 antibody, light chain	HIVl **gpl20** antibody	**WO2001000678** SEQ ID NO: 68	8752
HIV1 134	**HIV1** gpl20 antibody, light **chain**	HIVl **gpl20** antibody	WO2001000678 SEQ ID NO: 69	8753
HIV1 135	**HIV1** gpl20 antibody, light chain	HIVl **gpl20** antibody	**WO2001000678** SEQ ID NO: 70	8754
HIV1 136	**HIV1** gpl20 antibody, **light chain**	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 71	8755
HIV l l37	**HIV1** gpl20 **antibody,** light chain	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 72	8756
HIV1 138	**HIV1** gpl20 antibody, **light chain**	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 79	8757
HIV l l39	**HIV1** gpl20 **antibody,** light chain	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 80	8758
HIV1 140	**HIV1** gpl20 antibody, **light chain**	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 81	8759
HIV 1141	**HIV1** gpl20 **antibody,** light chain	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 82	8760
HIV1 142	**HIV1** gpl20 antibody, **light chain**	HIVl gpl20 antibody	**WO2001000678** SEQ ID NO: 83	8761
HIV **1143**	Kappa light chain	1460_G14	**US905 1362** SEQ ID NO: 22	8762
HIVl 144	Kappa **light** chain variable region	1456_P20	US905 1362 SEQ ID **NO:** 34	8763
HIV l 145	Kappa light chain variable reedon	1460_G 14	**US905** 1362 SEQ ID NO: 36	8764
HTV1 I46	Kappa **light** chain	1456_P20	US905 1362 SEQ ID NO: 18	8765
HIV 1147	**Lambda light** chain	1456_A12	**US905I362** SEQ ID NO: 50	8766
HIV1 148	Lambda light chain	1469 **M23**	US905 1362 SEQ ID NO: 142	8767
HIV1 l49	Lambda light **chain**	1489_113	**US905 1362** SEQ ID NO: **14**	8768
HIV l l50	Lambda light chain	1495_C14	**US9051362** SEQ ID NO: 26	8769
HIVl 15 1	Lambda **light** chain variable region	1489_113	US905 1362 **SEQ** ID **NO**: 32	8770
HIV l 152	Lambda **light** chain variable reedon	1495_C14	**US905** 1362 SEQ ID NO: 38	8771
HIVl 153	Lambda **light** chain variable region	1496_C09	US905 1362 SEQ ID **NO: 40**	8772
HIV l 154	Lambda **light** chain variable region	1456_A12	**US905I362** SEQ ID NO: **5** l	8773
HEV1 l55	Lambda light chain variable region	1503_**H05**	US905 1362 SEQ ID **NO:** 56	8774
HIV **1156**	**Lambda light** chain	1496_C09	**US905 1362** SEQ ID **NO: 30**	8775

HIV1157	Light chain	2424	Kumar, R., et al., Functional and Structural Characterization of Human V3-Specific Monoclonal Antibody 2424 with Neutralizing Activity against HIV-1 JRFL; J. Virol. 89 (17), 9090-9102 (2015), NCBI Accession # 4XML_L (215aa)	8776
HIV1158	Light chain	8062	Gustchina, E., PLoS ONE 8 (11), E78187 (2013), NCBI Accession # 4KHX_L(213aa)	8777
HIV1159	Light chain	1.00E+09	US20140348785 SEQ ID NO: 2	8778
HIV1160	Light Chain	10e8 (monoclonal)	Huang J et al., Nature 491 (7424), 406-412 (2012), NCBI Accession # 4G6F_D (215aa)	8779
HIV1161	Light chain	12a12kc	US20140328862 SEQ ID NO: 453	8780
HIV1162	Light chain	12a13kc	US20140328862 SEQ ID NO: 454	8781
HIV1163	Light chain	12a16kc	US20140328862 SEQ ID NO: 455	8782
HIV1164	Light chain	12a1kc	US20140328862 SEQ ID NO: 456	8783
HIV1165	Light chain	12a20kc	US20140328862 SEQ ID NO: 457	8784
HIV1166	Light chain	12a21	NCBI Accession # 4JPW_L (210aa)	8785
HIV1167	Light chain	12a21kc	US20140328862 SEQ ID NO: 458	8786
HIV1168	Light chain	12a22kc	US20140328862 SEQ ID NO: 459	8787
HIV1169	Light chain	12a23kc	US20140328862 SEQ ID NO: 460	8788
HIV1170	Light chain	12a27kc	US20140328862 SEQ ID NO: 461	8789
HIV1171	Light chain	12a46kc	US20140328862 SEQ ID NO: 462	8790
HIV1172	Light chain	12a55kc	US20140328862 SEQ ID NO: 463	8791
HIV1173	Light chain	12a56kc	US20140328862 SEQ ID NO: 464	8792
HIV1174	Light chain	12a6kc	US20140328862 SEQ ID NO: 465	8793
HIV1175	Light chain	12a7kc	US20140328862 SEQ ID NO: 466	8794
HIV1176	Light chain	17b	Kwong, P.D., et al., structure of an HIV gp120 envelope glycoprotein in complex with the CD4 receptor and a neutralizing human antibody; Nature 393 (6686), 648-659 (1998), NCBI Accession # 1G9M_L (214aa)	8795
HIV1177	Light chain	1b2530	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YFL_L (215aa)	8796
HIV1178	Light chain	1F7	US6057421A Fig 8	8797
HIV1179	Light chain	1NC9	WO2012154312 SEQ ID NO: 2472	8798
HIV1180	Light chain	2.2C	Acharya, P., et al., Structural Definition of an Antibody-Dependent Cellular Cytotoxicity Response Implicated in Reduced Risk for HIV-1 Infection; J. Virol. 88 (21), 12895-12906 (2014), NCBI Accession # 4R4N_L (210aa)	8799
HIV1181	Light chain	2F5	US8637036B2 SEQ ID NO: 10	8800
HIV1182	Light chain	3040LC	WO2015117008 SEQ ID NO: 29	8801
HIV1183	Light chain	3044LC	WO2015117008 SEQ ID NO: 32	8802
HIV1184	Light chain	3430LC	WO2015117008 SEQ ID NO: 30	8803
HIV1185	Light chain	3484LC	WO2015117008 SEQ ID NO: 31	8804
HIV1186	Light chain	3630LC	WO2015117008 SEQ ID NO: 33	8805
HIV1187	Light chain	3A124KC	US20140328862 SEQ ID NO: 506	8806
HIV1188	Light chain	3A125KC	US20140328862 SEQ ID NO: 507	8807

HEV1189	Light chain	3A140LC	**US20 140328862** SEQ ID NO: 508	8808
HIV1190	Light chain	3A144KC	**US20 140328862** SEQ ID NO: 509	8809
HIVU91	Light **chain**	3A160KC	**US20 140328862** SEQ ID NO: 510	8810
HTV1 I92	Light chain	3A18KC	US20 140328862 SEQ ID NO: **511**	881 1
HIV1193	Light **chain**	3A204KC	**US20 140328862** SEQ ID NO: 512	8812
HEV1194	Light chain	**3A228KC**	US20 140328862 SEQ ID NO: 513	8813
HIVI 195	Light chain	3A233LC	**US20 140328862** SEQ ID NO: 514	8814
HIV1196	Light **chain**	3A244LC	**US20 140328862** SEQ ID NO: 515	8815
HIV1 197	Light chain	3A255LC	**US20 140328862** SEQ ID NO: 516	8816
HIV1198	Light **chain**	3A296KC	**US20 140328862** SEQ ID NO: 517	8817
HEV1199	Light chain	3A334LC	**US20 140328862** SEQ ID NO: 518	8818
HIV1200	Light chain	**3A366KC**	**US20 140328862** SEQ ID NO: 519	8819
HIV1201	Light chain	3A384KC	**US20 140328862** SEQ ID NO: 520	8820
HIV1202	Light chain	**3A4 19KC**	**US20 140328862** SEQ ID NO: 521	8821
HIV1203	Light **chain**	**3a426kc**	US20 140328862 SEQ ID NO: 535	8822
HEV1204	Light chain	3A461KC	US20 140328862 SEQ ID NO: 522	8823
HIV1205	Light chain	3A474KC	US20 140328862 SEQ ID NO: 523	8824
HIV1206	Light chain	3a5 15kc	**US20 140328862** SEQ ID NO: 536	8825
HIV1207	Light chain	**3A5 18KC**	**US20 140328862** SEQ ID NO: 524	8826
HIV1208	Light chain	3A539LC	**US20 140328862** SEQ ID NO: 525	**8827**
HEV1209	Light chain	3A576LC	**US20 140328862** SEQ ID NO: 526	8828
HIV1210	Light chain	3A613LC	US20 140328862 SEQ ID NO: 527	8829
HIV121 1	Light chain	3A64KC	**US20 140328862** SEQ ID NO: 528	8830
HIV1212	Light chain	3A650KC	**US20 140328862** SEQ ID NO: 529	8831
HIV **12 13**	Light chain	3A67KC	**US20 140328862** SEQ ID NO: 530	8832
HTV12 14	Light chain	**3A779KC**	US20 140328862 SEQ ID NO: 531	8833
HIV 1215	Light chain	**3A816KC**	US20 140328862 SEQ ID NO: 532	8834
HIV1216	Light chain	3A869KC	**US20 140328862** SEQ ID NO: 533	8835
HIV1217	Light chain	**3A93LC**	**US20140328862** SEQ ID NO: 534	8836
HIV12 18	Light **chain**	**3anc3kc**	**US20 140328862** SEQ ID NO: 547	8837
HTV12 19	Light chain	**3bl06kc**	**US20 140328862** SEQ ID NO: 548	8838
HIV 1220	Light chain	**3b l29kc**	US20 140328862 SEQ ID NO: 537	8839
HIV1221	Light chain	**3bl6kc**	**US20 140328862** SEQ ID NO: 549	8840
HIV1222	Light chain	**3bl711c**	**US20 140328862** SEQ ID NO: 538	8841
HIV 1223	Light **chain**	**3b l80kc**	**US20 140328862** SEQ ID NO: 550	8842
HTV1224	Light chain	**3bl83kc**	**US20 140328862** SEQ ID NO: 551	8843
HIV 1225	Light chain	**3b 19 lkc**	US20 140328862 SEQ ID NO: 552	8844
HEV1226	Light chain	**3b2 1kc**	**US20140328862** SEQ ID NO: 553	8845
HIV1227	Light chain	**3b27kc**	**US20 140328862** SEQ ID NO: 539	8846
HIV 1228	Light **chain**	**3b41kc**	**US20 140328862** SEQ ID NO: 540	8847
HTV1229	Light chain	**3b46kc**	**US20 140328862** SEQ ID NO: 542	8848
HIV 1230	Light chain	3b57k:	US20 140328862 SEQ ID NO: 543	8849
HEV123 1	Light chain	3b5kc	**US20140328862** SEQ ID NO: 541	8850
HIV1232	Light chain	**3b8kc**	**US20 140328862** SEQ ID NO: 544	8851
HIV1233	Light **chain**	3bncl02kc	**US20 140328862** SEQ ID NO: 554	8852

HIV1234	Light chain	3bnc104kc	US20140328862 SEQ ID NO: 555	8853
HIV1235	Light chain	3bnc105kc	US20140328862 SEQ ID NO: 556	8854
HIV1236	Light chain	3bnc107kc	US20140328862 SEQ ID NO: 557	8855
HIV1237	Light chain	3bnc108kc	US20140328862 SEQ ID NO: 558	8856
HIV1238	Light chain	3bnc117	Zhou T et al., Immunity 39 (2), 245-258 (2013), NCBI Accession # 4LSV_L(206aa)	8857
HIV1239	Light chain	3bnc117kc	US20140328862 SEQ ID NO: 559	8858
HIV1240	Light chain	3bnc134kc	US20140328862 SEQ ID NO: 560	8859
HIV1241	Light chain	3bnc142kc	US20140328862 SEQ ID NO: 561	8860
HIV1242	Light chain	3bnc151kc	US20140328862 SEQ ID NO: 562	8861
HIV1243	Light chain	3bnc153kc	US20140328862 SEQ ID NO: 563	8862
HIV1244	Light chain	3bnc156kc	US20140328862 SEQ ID NO: 564	8863
HIV1245	Light chain	3bnc158kc	US20140328862 SEQ ID NO: 565	8864
HIV1246	Light chain	3bnc159kc	US20140328862 SEQ ID NO: 566	8865
HIV1247	Light chain	3bnc15kc	US20140328862 SEQ ID NO: 567	8866
HIV1248	Light chain	3bnc176kc	US20140328862 SEQ ID NO: 568	8867
HIV1249	Light chain	3bnc193kc	US20140328862 SEQ ID NO: 569	8868
HIV1250	Light chain	3bnc196kc	US20140328862 SEQ ID NO: 570	8869
HIV1251	Light chain	3bnc31kc	US20140328862 SEQ ID NO: 571	8870
HIV1252	Light chain	3bnc42kc	US20140328862 SEQ ID NO: 572	8871
HIV1253	Light chain	3bnc53kc	US20140328862 SEQ ID NO: 573	8872
HIV1254	Light chain	3BNC55KC	US20140328862 SEQ ID NO: 545	8873
HIV1255	Light chain	3BNC60KC	US20140328862 SEQ ID NO: 546	8874
HIV1256	Light chain	3bnc62kc	US20140328862 SEQ ID NO: 574	8875
HIV1257	Light chain	3bnc65kc	US20140328862 SEQ ID NO: 575	8876
HIV1258	Light chain	3bnc66kc	US20140328862 SEQ ID NO: 576	8877
HIV1259	Light chain	3bnc75kc	US20140328862 SEQ ID NO: 577	8878
HIV1260	Light chain	3bnc79kc	US20140328862 SEQ ID NO: 578	8879
HIV1261	Light chain	3bnc81kc	US20140328862 SEQ ID NO: 579	8880
HIV1262	Light chain	3bnc84kc	US20140328862 SEQ ID NO: 580	8881
HIV1263	Light chain	3bnc87kc	US20140328862 SEQ ID NO: 581	8882
HIV1264	Light chain	3bnc89kc	US20140328862 SEQ ID NO: 582	8883
HIV1265	Light chain	3bnc91kc	US20140328862 SEQ ID NO: 583	8884
HIV1266	Light chain	3bnc95kc	US20140328862 SEQ ID NO: 584	8885
HIV1267	Light chain	412d	Huang et al., Science 317 (5846), 1930-1934 (2007), NCBI Accession # 2QAD_G (214aa)	8886
HIV1268	Light Chain	44-vrc13.01	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDJ_B (206aa)	8887
HIV1269	Light chain	45-46m2	Diskin, R., et al., Restricting HIV-1 pathways for escape using rationally designed anti-HIV-1 antibodies; J. Exp. Med. 210 (6), 1235-1249 (2013), NCBI Accession # 4JKP_L (210aa)	8888
HIV1270	Light chain	4835_F12 (PGT-124)	US20140205612 SEQ ID NO: 413	8889
HIV1271	Light chain	4838_L06 (PGT-121)	US20140205612 SEQ ID NO: 148	8890

HIV1272	Light chain	4858_P08 (PGT-123)	US20140205612 SEQ ID NO: 176	8891
HIV1273	Light chain	4869-K15 (PGT-133)	US20140205612 SEQ ID NO: 428	8892
HIV1274	Light chain	4873_E03 (PGT-121)	US20140205612 SEQ ID NO: 147	8893
HIV1275	Light chain	4876_M06 (PGT-134)	US20140205612 SEQ ID NO: 439	8894
HIV1276	Light chain	4877_D15 (PGT-122)	US20140205612 SEQ ID NO: 160	8895
HIV1277	Light chain	4964_G22 (PGT-141), 4993_K13 (PGT-141), 4995_E20 (PGT-142)	US20140205612 SEQ ID NO: 284	8896
HIV1278	Light chain	4970_K22 (PGT-144)	US20140205612 SEQ ID NO: 312	8897
HIV1279	Light chain	4980_N08 (PGT-143)	US20140205612 SEQ ID NO: 301	8898
HIV1280	Light chain	4995_P16 (PGT-145)	US20140205612 SEQ ID NO: 385	8899
HIV1281	Light chain	4e10 Fv	Finton, K.A., et al., PLoS Pathol. 9 (9), E1003639 (2013), NCBI Accession # 4LLV_B (112aa)	8900
HIV1282	Light chain	5114_A19 (PGT-128)	US20140205612 SEQ ID NO: 392	8901
HIV1283	Light chain	5120_N10 (PGT-139)	US20140205612 SEQ ID NO: 469	8902
HIV1284	Light chain	5131_A17 (PGT-132)	US20140205612 SEQ ID NO: 488	8903
HIV1285	Light chain	5136_H01 (PGT-131)	US20140205612 SEQ ID NO: 355	8904
HIV1286	Light chain	5138_G07 (PGT-138)	US20140205612 SEQ ID NO: 483	8905
HIV1287	Light chain	5141_B17 (PGT-126)	US20140205612 SEQ ID NO: 208	8906
HIV1288	Light chain	5145_B14 (PGT-127)	US20140205612 SEQ ID NO: 329	8907
HIV1289	Light chain	5147_N06 (PGT-130)	US20140205612 SEQ ID NO: 244	8908
HIV1290	Light chain	5329_C19 (PGT-136), 5366_P21 (PGT-136)	US20140205612 SEQ ID NO: 257	8909
HIV1291	Light chain	5343_B08 (PGT-135), 5344_E16 (PGT-135)	US20140205612 SEQ ID NO: 240	8910
HIV1292	Light chain	5345_I01 (PGT-137)	US20140205612 SEQ ID NO: 396	8911
HIV1293	Light chain	6808_B09 (PGT-156)	US20140205612 SEQ ID NO: 553	8912
HIV1294	Light chain	6831_A21 (PGT-151)	US20140205612 SEQ ID NO: 482	8913
HIV1295	Light chain	6843_G20 (PGT-154)	US20140205612 SEQ ID NO: 524	8914
HIV1296	Light chain	6881_N05 (PGT-158).	US20140205612 SEQ ID NO: 578	8915

HIV1297	Light chain	6889_I17 (PGT-152)	US20140205612 SEQ ID NO: 496	8916
HIV1298	Light chain	6891_F06 (PGT-153)	US20140205612 SEQ ID NO: 510	8917
HIV1299	Light chain	6892_C23 (PGT-157)	US20140205612 SEQ ID NO: 565	8918
HIV1300	Light chain	6892_D19 (PGT-155)	US20140205612 SEQ ID NO: 539	8919
HIV1301	Light chain	7H6	US20140348785 SEQ ID NO: 4	8920
HIV1302	Light chain	7N16	US20140348785 SEQ ID NO: 6	8921
HIV1303	Light chain	8anc131	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4RWY_L (213aa)	8922
HIV1304	Light chain	8ANC131KC	US20140328862 SEQ ID NO: 440	8923
HIV1305	Light chain	8anc134	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4RX4_L (213aa)	8924
HIV1306	Light chain	8ANC134KC	US20140328862 SEQ ID NO: 441	8925
HIV1307	Light chain	8ANC13KC	US20140328862 SEQ ID NO: 442	8926
HIV1308	Light chain	8ANC14KC	US20140328862 SEQ ID NO: 448	8927
HIV1309	Light chain	8ANC16KC	US20140328862 SEQ ID NO: 449	8928
HIV1310	Light chain	8anc182kc	US20140328862 SEQ ID NO: 446	8929
HIV1311	Light chain	8anc192kc	US20140328862 SEQ ID NO: 447	8930
HIV1312	Light chain	8ANC195KC	US20140328862 SEQ ID NO: 450	8931
HIV1313	Light chain	8ANC24KC	US20140328862 SEQ ID NO: 451	8932
HIV1314	Light chain	8ANC45KC	US20140328862 SEQ ID NO: 443	8933
HIV1315	Light chain	8ANC50KC	US20140328862 SEQ ID NO: 444	8934
HIV1316	Light chain	8ANC5KC	US20140328862 SEQ ID NO: 452	8935
HIV1317	Light chain	8ANC88KC	US20140328862 SEQ ID NO: 445	8936
HIV1318	Light chain	Anti-HcG	Fotinou C. et al " Structure of an Fab fragment against a C-terminal peptide of hCG at 2.0 A resolution" J. Biol. Chem. 273 (35), 22515-22518 (1998); NCBI Accession # 1SBS_L	8937
HIV1319	Light chain	B12	Zhou T et al., Structural definition of a conserved neutralization epitope on HIV-1 gp120; Nature 445 (7129), 732-737 (2007), NCBI Accession # 2NY7_L (215aa)	8938
HIV1320	Light Chain	C38-vrc16.01	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDK_L (214aa)	8939
HIV1321	Light chain	C38-vrc18.02	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDL_L (211aa)	8940
HIV1322	Light chain	CAP256-VRC26.01	WO2015128846 SEQ ID NO: 14	8941
HIV1323	Light chain	CAP256-VRC26.02	WO2015128846 SEQ ID NO: 18	8942

HIV1324	Light chain	CAP256-VRC26.03	WO2015128846 SEQ ID NO: 22	8943
HIV1325	Light chain	CAP256-VRC26.04	WO2015128846 SEQ ID NO: 26	8944
HIV1326	Light chain	CAP256-VRC26.05	WO2015128846 SEQ ID NO: 30	8945
HIV1327	Light chain	CAP256-VRC26.06	WO2015128846 SEQ ID NO: 34	8946
HIV1328	Light chain	CAP256-VRC26.07	WO2015128846 SEQ ID NO: 38	8947
HIV1329	Light chain	CAP256-VRC26.08	WO2015128846 SEQ ID NO: 42	8948
HIV1330	Light chain	CAP256-VRC26.09	WO2015128846 SEQ ID NO: 46	8949
HIV1331	Light chain	CAP256-VRC26.10	WO2015128846 SEQ ID NO: 50	8950
HIV1332	Light chain	CAP256-VRC26.11	WO2015128846 SEQ ID NO: 54	8951
HIV1333	Light chain	CAP256-VRC26.12	WO2015128846 SEQ ID NO: 58	8952
HIV1334	Light chain	CAP256-VRC26.25	WO2015128846 SEQ ID NO: 171	8953
HIV1335	Light chain	CAP256-VRC26.26	WO2015128846 SEQ ID NO: 179	8954
HIV1336	Light chain	CAP256-VRC26.27	WO2015128846 SEQ ID NO: 187	8955
HIV1337	Light chain	CAP256-VRC26-I1	WO2015128846 SEQ ID NO: 6	8956
HIV1338	Light chain	CAP256-VRC26-I2	WO2015128846 SEQ ID NO: 10	8957
HIV1339	Light chain	CAP256-VRC26-UCA.	WO2015128846 SEQ ID NO: 2	8958
HIV1340	Light chain	construct #2816, #2861	WO2015013390 SEQ ID NO: 5	8959
HIV1341	Light chain	construct #2817, #2860	WO2015013390 SEQ ID NO: 6	8960
HIV1342	Light chain	construct #2858, #2859, #2861	WO2015013390 SEQ ID NO: 7	8961
HIV1343	Light chain	Fab 2219	Stanfield, R.L., et al., J. Virol. 80 (12), 6093-6105 (2006), NCBI Accession # 2B0S_L (215aa)	8962
HIV1344	Light chain	Fab 2g12	Doores, K.J., et al., J. Virol. 84 (20), 10690-10699 (2010), NCBI Accession # 3OAU_L(212a)	8963
HIV1345	Light chain	Fab 2g12	Stanfield, R.L. et al., Crystal structure of the HIV neutralizing antibody 2G12 in complex with a bacterial oligosaccharide analog of mammalian oligomannose; Glycobiology 25 (4), 412-419 (2015), NCBI Accession # 4RBP_L (213aa)	8964
HIV1346	Light chain	Fab F425-b4e8	Bell et al., J. Mol. Biol. 375 (4), 969-978 (2008), NCBI Accession # 2QSC_L (215aa)	8965
HIV1347	Light chain	G4D	US20130195881 SEQ ID NO: 39	8966
HIV1348	Light chain	G4H	US20130195881 SEQ ID NO: 38	8967
HIV1349	Light chain	gVRC-H5(d74)/VRC-PG04LC,	WO2013090644 SEQ ID NO: 19	8968

		gVRC0H12(D74)/VRC-PG04LC		
HIV1350	Light chain	I2 (unbound) From Ch103 Lineage	Fera, D. et aL Affinity maturation in an HIV broadly neutralizing B-cell lineage through reonentalion of vanable domains; Proc. Natl. Acad. Sci. U.S.A. 111 (28), 10275-10280 (2014), NCBI Accession # 4QHN_B (213aa)	8969
HIV1351	Light chain	IGLV3-19*01	US20140348785 SEQ ID NO: 8	8970
HIV1352	Light chain	k3	WO20151117008 SEQ ID NO: 19	8971
HIV1353	Light chain	k5	WO20151 17008 SEQ ID NO: 20	8972
HIV 1354	Light chain	k53	WO20151117008 SEQ ID NO: 24	8973
HIV1355	Light chain	k59	WO20151117008 SEQ ID NO: 21	8974
HIV 1356	Light chain	k61	WO20151 17008 SEQ ID NO: 25	8975
HIV 1357	Light chain	k62	WO20151117008 SEQ ID NO: 22	8976
HIV1358	Light chain	k81	WO20151 17008 SEQ ID NO: 28	8977
HIV1359	Light chain	k31	WO20151117008 SEQ ID NO: 26	8978
HIV1360	Light chain	kl8	WO20151117008 SEQ ID NO: 23	8979
HIV 1361	Light chain	k!9	WO20151 17008 SEQ ID NO: 27	8980
HIV1362	Light chain	LSSB2066KC	US20140328862 SEQ ID NO: 501	8981
HIV1363	Light chain	LSSB2080KC	US20140328862 SEQ ID NO: 502	8982
HIV1364	Light chain	LSSB2133KC	US20140328862 SEQ ID NO: 503	8983
HIV1365	Light chain	LSSB2182KC	US20140328862 SEQ ID NO: 504	8984
HIV1366	Light chain	LSSB2339LC	US20140328862 SEQ ID NO: 467	8985
HIV1367	Light chain	LSSB2351LC	US20140328862 SEQ ID NO: 468	8986
HIV1368	Light chain	LSSB2364LC	US20140328862 SEQ ID NO: 469	8987
HIV1369	Light chain	LSSB2367LC	US20140328862 SEQ ID NO: 470	8988
HIV1370	Light chain	LSSB2490LC	US20140328862 SEQ ID NO: 471	8989
HIV1371	Light chain	LSSB2530LC	US20140328862 SEQ ID NO: 472	8990
HIV1372	Light chain	LSSB2554LC	US20140328862 SEQ ID NO: 473	8991
HIV1373	Light chain	LSSB2586LC	US20140328862 SEQ ID NO: 474	8992
HIV1374	Light chain	LSSB2612LC	US20140328862 SEQ ID NO: 475	8993
HIV1375	Light chain	LSSB2640LC	US20140328862 SEQ ID NO: 476	8994
HIV1376	Light chain	LSSB2644LC	US20140328862 SEQ ID NO: 477	8995
HIV1377	Light chain	LSSB2666LC	US20140328862 SEQ ID NO: 478	8996
HIV1378	Light chain	LSSB2680LC	US20140328862 SEQ ID NO: 479	8997
HIV1379	Light chain	LSSB2683LC	US20140328862 SEQ ID NO: 480	8998
HIV1380	Light chain	LSSB331KC	US20140328862 SEQ ID NO: 505	8999
HEV1381	Light chain	LSSB344LC	US20140328862 SEQ ID NO: 481	9000
HIV1382	Light chain	LSSNEC107LC	US20140328862 SEQ ID NO: 482	9001
HIV1383	Light chain	LSSNEC108LC	US20140328862 SEQ ID NO: 483	9002
HIV1384	Light chain	LSSNEC117LC	US20140328862 SEQ ID NO: 484	9003

HIV1385	Light chain	LSSNEC118 LC	US20140328862 SEQ ID NO: 485	9004
HIV1386	Light chain	LSSNEC122 LC	US20140328862 SEQ ID NO: 486	9005
HIV1387	Light chain	LSSNEC24L C	US20140328862 SEQ ID NO: 487	9006
HIV1388	Light chain	LSSNEC2LC	US20140328862 SEQ ID NO: 488	9007
HIV1389	Light chain	LSSNEC33L C	US20140328862 SEQ ID NO: 489	9008
HIV1390	Light chain	LSSNEC46L C	US20140328862 SEQ ID NO: 490	9009
HIV1391	Light chain	LSSNEC48L C	US20140328862 SEQ ID NO: 491	9010
HIV1392	Light chain	LSSNEC52L C	US20140328862 SEQ ID NO: 492	9011
HIV1393	Light chain	LSSNEC56L C	US20140328862 SEQ ID NO: 493	9012
HIV1394	Light chain	LSSNEC60L C	US20140328862 SEQ ID NO: 494	9013
HIV1395	Light chain	LSSNEC70L C	US20140328862 SEQ ID NO: 495	9014
HIV1396	Light chain	LSSNEC72L C	US20140328862 SEQ ID NO: 496	9015
HIV1397	Light chain	LSSNEC7LC	US20140328862 SEQ ID NO: 497	9016
HIV1398	Light chain	LSSNEC89L C	US20140328862 SEQ ID NO: 498	9017
HIV1399	Light chain	LSSNEC94L C	US20140328862 SEQ ID NO: 499	9018
HIV1400	Light chain	LSSNEC9LC	US20140328862 SEQ ID NO: 500	9019
HIV1401	Light chain	m12_Fd-aa	US7803913B2 SEQ ID NO: 7	9020
HIV1402	Light chain	m14-Fd-aa	US7803913B2 SEQ ID NO: 5	9021
HIV1403	Light chain	m16-Fd-aa	US7803913B2 SEQ ID NO: 8	9022
HIV1404	Light chain	m18_Fd-aa	US7803913B2 SEQ ID NO: 6	9023
HIV1405	Light chain	M66	Ofek, G., et al., Structural Basis for HIV-1 Neutralization by 2F5-Like Antibodies m66 and m66.6; J. Virol. 88 (5), 2426-2441 (2014), NCBI Accession # 4NRY_H (235aa)	9024
HIV1406	Light chain	M66.6	Ofek, G., et al., Structural Basis for HIV-1 Neutralization by 2F5-Like Antibodies m66 and m66.6; J. Virol. 88 (5), 2426-2441 (2014), NCBI Accession # 4NRZ_L (213aa)	9025
HIV1407	Light Chain	Mab 2158	Spurrier, B., et al., Functional Implications of the Binding Mode of a Human Conformation-Dependent V2 Monoclonal Antibody against HIV; J. Virol. 88 (8), 4100-4112 (2014), NCBI Accession # 4OAW_C (214aa)	9026
HIV1408	Light chain	MV1	US20130195881 SEQ ID NO: 40	9027
HIV1409	Light chain	Pg16 Fab	Pancera, M., et al., Nat. Struct. Mol. Biol. 20 (7), 804-813 (2013), NCBI Accession # 4DQO_L (216aa)	9028
HIV1410	Light chain	Pg9	Willis, J.R., et al., J. Clin. Invest. 125 (6), 2523-2531 (2015), NCBI Accession # 4YAQ_L (216aa)	9029
HIV1411	Light chain	Pgt121-GI Fab	Mouquet H et al., Complex-type N-glycan recognition by potent broadly neutralizing HIV antibodies; **Proc Natl** Acad Sci **U** S A. 2012	9030

			Nov 20;109(47):E3268-77, NCBI Accession # 4FQQ_A (215aa)	
HIV1412	Light chain	Pgt122	Julien, J.P., et al., PLoS Pathol. 9 (5), E1003342 (2013)", NCBI Accession # 4JY5_L (211aa)	9031
HIV1413	Light chain	Pgt123	Julien, J.P., et al., PLoS Pathol. 9 (5), E1003342 (2013)", NCBI Accession # 4JY6_A (211aa)	9032
HIV1414	Light chain	Pgt124	Garces, F., et al., Structural Evolution of Glycan Recognition by a Family of Potent HIV Antibodies; Cell 159 (1), 69-79 (2014), NCBI Accession # 4R26_L (214aa)	9033
HIV1415	Light chain	Pgt130	Doores, K.J., et al., J. Virol. 89 (2), 1105-1118 (2015), NCBI Accession # 4RNR_B (216aa)	9034
HIV1416	Light chain	Pgt135	Grover et al., Science 343 (6171), 656-661 (2014), NCBI Accession # 4NZR_L (214aa)	9035
HIV1417	Light chain	S8, S19, S20	US20110059015 SEQ ID NO: 2	9036
HIV1418	light chain	Suvizumab		9037
HIV1419	Light Chain	Vrc- Pg04	Wu, X., et al., Focused evolution of HIV-1 neutralizing antibodies revealed by structures and deep sequencing; Science 333 (6049), 1593-1602 (2011)", NCBI Accession # 3SE9_L (208aa)	9038
HIV1420	Light chain	VRC01	US8637036B2 SEQ ID NO: 2	9039
HIV1421	Light chain	VRC01 E1/I2 deletion	US2014 0322163 SEQ ID NO: 53	9040
HIV1422	Light chain	VRC01 E1/I2del F97D	US2014 0322163 SEQ ID NO: 222	9041
HIV1423	Light chain	VRC01 E1/I2del F97H	US2014 0322163 SEQ ID NO: 225	9042
HIV1424	Light chain	VRC01 E1/I2del F97K	US2014 0322163 SEQ ID NO: 223	9043
HIV1425	Light chain	VRC01 E1/I2del F97S	US2014 0322163 SEQ ID NO: 224	9044
HIV1426	Light chain	VRC01 E1/I2del V3E	US2014 0322163 SEQ ID NO: 219	9045
HIV1427	Light chain	VRC01 E1/I2del V3E, F97H	US2014 0322163 SEQ ID NO: 227	9046
HIV1428	Light chain	VRC01 E1/I2del V3E, F97S	US2014 0322163 SEQ ID NO: 226	9047
HIV1429	Light chain	VRC01 E1/I2del V3K	US2014 0322163 SEQ ID NO: 220	9048
HIV1430	Light chain	VRC01 E1/I2del V3S	US2014 0322163 SEQ ID NO: 221	9049
HIV1431	Light chain	VRC01HC/VRC03LC	WO2013090644 SEQ ID NO: 31	9050
HIV1432	Light chain	VRC01hpL02	US2014 0322163 SEQ ID NO: 50	9051
HIV1433	Light chain	VRC01hpL02 E1/I2-deletion, V3S	US2014 0322163 SEQ ID NO: 232	9052
HIV1434	Light chain	VRC01hpL03	US2014 0322163 SEQ ID NO: 228	9053

HIV1435	Light chain	VRC01hpL04	US2014 0322163 SEQ ID NO: 229	9054
HIV1436	Light chain	VRC01hpL05	US2014 0322163 SEQ ID NO: 230	9055
HIV1437	Light chain	VRC01hpL06	US2014 0322163 SEQ ID NO: 231	9056
HIV1438	Light chain	VRC01LhpL 03 E1/I2-deletion, V3S	US2014 0322163 SEQ ID NO: 233	9057
HIV1439	Light chain	VRC01LhpL 04 E1/I2-deletion, V3E	US2014 0322163 SEQ ID NO: 237	9058
HIV1440	Light chain	VRC01LhpL 04 E1/I2-deletion, V3S	US2014 0322163 SEQ ID NO: 234	9059
HIV1441	Light chain	VRC01LhpL 05 E1/I2 deletion, V3S	US2014 0322163 SEQ ID NO: 235	9060
HIV1442	Light chain	VRC01LhpL 06 E1/I2-deletion, V3S	US2014 0322163 SEQ ID NO: 236	9061
HIV1443	Light chain	VRC02	US8637036B2 SEQ ID NO: 4	9062
HIV1444	Light chain	VRC03	US8637036B2 SEQ ID NO: 28	9063
HIV1445	Light chain	VRC03HC-VRC01LC	WO2013090644 SEQ ID NO: 1	9064
HIV1446	Light chain	Vrc06b	Wu, X., et al., Maturation and Diversity of the VRC01-Antibody Lineage over 15 Years of Chronic HIV-1 Infection; Cell 161 (3), 470-485 (2015), NCBI Accession # 4XNZ_F (209aa)	9065
HIV1447	Light chain	Vrc08c	Wu, X., et al., Maturation and Diversity of the VRC01-Antibody Lineage over 15 Years of Chronic HIV-1 Infection; Cell 161 (3), 470-485 (2015), NCBI Accession # 4XNY_L (211aa)	9066
HIV1448	Light chain	Vrc23	Georgiev, I.S., et al., Delineating antibody recognition in polyclonal sera from patterns of HIV-1 isolate neutralization; Science 340 (6133), 751-756 (2013), NCBI Accession # 4J6R_L (210aa)	9067
HIV1449	Light chain	VRC-CH30	WO2013090644 SEQ ID NO: 21	9068
HIV1450	Light chain	Vrc-ch31	Zhou T et al., Immunity 39 (2), 245-258 (2013), NCBI Accession # 4LSP_L (210aa)	9069
HIV1451	Light chain	VRC-CH32	Wu X. et al, " Focused evolution of HIV-1 neutralizing antibodies revealed by structures and deep sequencing" Science 333 (6049), 1593-1602 (2011), NCBI Accession # AEM62727	9070
HIV1452	Light chain	VRC-CH33	WO2013090644 SEQ ID NO: 27	9071
HIV1453	Light chain	VRC-CH34	WO2013090644 SEQ ID NO: 29	9072
HIV1454	Light chain	VRC-PG04	Wu X. et al, " Focused evolution of HIV-1 neutralizing antibodies revealed by structures and deep sequencing" Science 333 (6049), 1593-1602 (2011), NCBI Accession # AEM62754	9073
HIV1455	Light chain	VRC-PG04b	WO2013090644 SEQ ID NO: 43	9074
HIV1456	Light chain	Vrc-pg20	Zhou T et al., Immunity 39 (2), 245-258 (2013), NCBI Accession # 4LSU_L (204aa)	9075
HIV1457	Light chain	X5	US7378093B2 SEQ ID NO: 2	9076
HIV1458	Light chain	X5	US8110192B2 SEQ ID NO: 4	9077

HIV1459	Light chain	Z13e1	Stanfield, R.L., et al, J. Mol. Biol. 414 (3), 460-476 (2011), NCBI Accession # 3Q1S_L (212aa)	9078
HIV1460	Light Chain	Z258-vrc27.01	Zhou T et al., Structural Repertoire of HIV-1-Neutralizing Antibodies Targeting the CD4 Supersite in 14 Donors; Cell 161 (6), 1280-1292 (2015), NCBI Accession # 4YDI_L (210aa)	9079
HIV1461	Light chain		NCBI Accession # 1N0X_M (215aa)	9080
HIV1462	Light chain		Okada, N., et al., Human IgM Monoclonal Antibodies Reactive with HIV-1-Infected Cells Generated Using a Trans-Chromosome Mouse; Microbiol. Immunol. 49 (5), 447-459 (2005), NCBI Accession # AAS01772.1(236aa)	9081
HIV1463	Light chain		US5804440A SEQ ID NO: 101	9082
HIV1464	Light chain		US5804440A SEQ ID NO: 102	9083
HIV1465	Light chain		US5804440A SEQ ID NO: 103	9084
HIV1466	Light chain		US5804440A SEQ ID NO: 104	9085
HIV1467	Light chain		US5804440A SEQ ID NO: 105	9086
HIV1468	Light chain		US5804440A SEQ ID NO: 107	9087
HIV1469	Light chain		US5804440A SEQ ID NO: 110	9088
HIV1470	Light chain		US5804440A SEQ ID NO: 115	9089
HIV1471	Light chain		US5804440A SEQ ID NO: 118	9090
HIV1472	Light chain		US5804440A SEQ ID NO: 121	9091
HIV1473	Light chain		US5804440A SEQ ID NO: 122	9092
HIV1474	Light chain		US5804440A SEQ ID NO: 124	9093
HIV1475	Light chain		US5804440A SEQ ID NO: 132	9094
HIV1476	Light chain		US5804440A SEQ ID NO: 147	9095
HIV1477	Light chain		US5804440A SEQ ID NO: 148	9096
HIV1478	Light chain		US5804440A SEQ ID NO: 149	9097
HIV1479	Light chain		US5804440A SEQ ID NO: 150	9098
HIV1480	Light chain		US5804440A SEQ ID NO: 151	9099
HIV1481	Light chain		US5804440A SEQ ID NO: 95	9100
HIV1482	Light chain		US5804440A SEQ ID NO: 96	9101
HIV1483	Light chain		US5804440A SEQ ID NO: 97	9102
HIV1484	Light chain		US5804440A SEQ ID NO: 98	9103
HIV1485	Light chain		WO2014063059 SEQ ID NO: 11	9104
HIV1486	Light chain		WO2014063059 SEQ ID NO: 129	9105
HIV1487	Light chain		WO2014063059 SEQ ID NO: 13	9106
HIV1488	Light chain		WO2014063059 SEQ ID NO: 15	9107
HIV1489	Light chain		WO2014063059 SEQ ID NO: 17	9108
HIV1490	Light chain		WO2014063059 SEQ ID NO: 19	9109
HIV1491	Light chain		WO2014063059 SEQ ID NO: 21	9110
HIV1492	Light chain		WO2014063059 SEQ ID NO: 23	9111
HIV1493	Light chain		WO2014063059 SEQ ID NO: 3	9112
HIV1494	Light chain		WO2014063059 SEQ ID NO: 5	9113
HIV1495	Light chain		WO2014063059 SEQ ID NO: 7	9114
HIV1496	Light chain		WO2014063059 SEQ ID NO: 9	9115

HIV1497	Light chain consensus		WO2014063059 SEQ ID NO: 1	9116
HIV1498	Light chain constant region	TNX-355, Idalizumab	US20130195881 SEQ ID NO: 2	9117
HIV1499	Light Chain Fab	Ch02	McLellan, J.S., et al., Nature 480 (7377), 336-343 (2011), NCBI Accession # 3U46_B (215aa)	9118
HIV1500	Light Chain Of Anti-HIV Fab From Human 21c Antibody	21C	Diskin, R., et al., Nat. Struct. Mol. Biol. 17 (5), 608-613 (2010), NCBI Accession # 3LMJ_L (217aa)	9119
HIV1501	Light Chain Of Anti-hiv-1 Gp120 V1v2 Antibody 830a	830a	Pan et al., J. Virol. 89 (15), 8003-8010 (2015), NCBI Accession # 4YWG_L (216aa)	9120
HIV1502	Light Chain Of Anti-hiv-1 V3 Monoclonal Antibody	Fab 2558	Gorny et al., PLoS ONE 6 (12), E27780 (2011), NCBI Accession # 3UJI_L (209aa)	9121
HIV1503	Light Chain Of Anti-hiv-1 V3 Monoclonal Antibody	Fab 4025	Gorny et al., PLoS ONE 6 (12), E27780 (2011), NCBI Accession # 3UJJ_L (213aa)	9122
HIV1504	Light chain partial	412D	Huang C. et al "Structural basis of tyrosine sulfation and VH-gene usage in antibodies that recognize the HIV type 1 coreceptor-binding site on gp120" Proc. Natl. Acad. Sci. U.S.A. 101 (9), 2706-2711 (2004), NCBI Accession # AAR88380	9123
HIV1505	Light chain partial	694/98D	Li L. et al, "A broad range of mutations in HIV-1 neutralizing human monoclonal antibodies specific for V2, V3, and the CD4 binding site", Mol. Immunol. 66 (2), 364-374 (2015); NCBI Accession # AKH36512	9124
HIV1506	Light chain variable region	0.5γ(1C10)	US8722861B2 SEQ ID NO: 2	9125
HIV1507	Light chain variable region	0.5δ (3D6)	US8722861B2 SEQ ID NO: 6	9126
HIV1508	Light chain variable region	10J4 mAb	WO2015103549 SEQ ID NO: 4	9127
HIV1509	Light chain variable region	10M6 mAb	WO2015103549 SEQ ID NO: 6	9128
HIV1510	Light chain variable region	13110 mAb	WO2015103549 SEQ ID NO: 8	9129
HIV1511	Light chain variable region	2N5mAb	WO2015103549 SEQ ID NO: 10	9130
HIV1512	Light chain variable region	35022 mAb	WO2015103549 SEQ ID NO: 2	9131
HIV1513	Light chain variable region	42F9	US8722861B2 SEQ ID NO: 8	9132
HIV1514	Light chain variable region	49G2	US8722861B2 SEQ ID NO: 10	9133
HIV1515	Light chain variable region	4O20mAb	WO2015103549 SEQ ID NO: 12	9134
HIV1516	Light chain variable region	5G2	US8722861B2 SEQ ID NO: 4	9135
HIV1517	Light chain variable region	7B9mAb	WO2015103549 SEQ ID NO: 14	9136
HIV1518	Light chain variable region	7K3mAb	WO2015103549 SEQ ID NO: 16	9137
HIV1519	Light chain variable region	B4	US7872110B2 SEQ ID NO: 4	9138

HIV1520	Light chain variable region	B4DIVKv.1	US7872110B2 SEQ ID NO: 9	9139
HIV1521	Light chain variable region	B4DIVKv.2	US7872110B2 SEQ ID NO: 10	9140
HIV1522	Light chain variable region	B4DIVKv.3	US7872110B2 SEQ ID NO: 11	9141
HIV1523	Light chain variable region	bl2 IgA2 antibody	WO2014040024 SEQ ID NO: 30	9142
HIV1524	Light chain variable region	CHμ39.1	US5773247 SEQ ID NO: 12	9143
HIV1525	Light chain variable region	CHμ5.5	US5773247 SEQ ID NO: 16	9144
HIV1526	Light chain variable region	F425-Alg8 antibody	WO2014040024 SEQ ID NO: 13	9145
HIV1527	Light chain variable region	Fab 43	US20090191216 SEQ ID NO: 9	9146
HIV1528	Light chain variable region	HGN194	US20110212106 SEQ ID NO: 46	9147
HIV1529	Light chain variable region	HJ16	US20110212106 SEQ ID NO: 14	9148
HIV1530	Light chain variable region	HK20	US20110212106 SEQ ID NO: 30	9149
HIV1531	Light chain variable region	IgA antibody	WO2014040024 SEQ ID NO: 15	9150
HIV1532	Light chain variable region	Makandal monoclonal antibody (Mmab)	US20100111990 SEQ ID NO: 3	9151
HIV1533	Light chain variable region	NM-01	US5665569 SEQ ID NO: 18	9152
HIV1534	Light chain variable region	NM-01 HuVH	US5665569 SEQ ID NO: 28	9153
HIV1535	Light chain variable region	NM-01 HuVK	US5665569 SEQ ID NO: 30	9154
HIV1536	Light chain variable region	NM-01 HuVKF	US5665569 SEQ ID NO: 32	9155
HIV1537	Light chain variable region	PGT125	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14410	9156
HIV1538	Light chain variable region	PGT126	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14411	9157
HIV1539	Light chain variable region	PGT131	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14415	9158
HIV1540	Light chain variable region	PGT136	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14417	9159
HIV1541	Light chain variable region	PGT137	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies", Nature 477 (7365), 466-470 (2011), NCBI Accession # AEN14418	9160
HIV1542	Light chain variable region	PGT141	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent	9161

			antibodies", **Nature** 477 (7365), 466-470 **(201 1), NCBI** Accession # **AEN14419**	
HIV1543	Light chain variable region	**PGT142**	**Walker** L.M. et **al** "Broad **neutralization** coverage of HIV by **multiple** highly potent antibodies", **Nature** 477 (7365), 466-470 **(201 1),** NCBI Accession # AEN14385	9162
HIV1544	Light **chain variable** region	**PGT143**	Walker L.M. et al "Broad neutralization coverage of HIV by multiple highly potent antibodies". Nature 477 (7365), 466-470 **(201 1),** NCBI Accession # **AEN1442 1**	9163
HIV1545	Light **chain** variable region	PGT144	Walker L.M. et al "Broad neutralization coverage of HIV by multiple **highly** potent antibodies", Nature 477 (7365), 466-470 **(201 1),** NCBI Accession # AEN 14422	9164
HIV1546	Light chain variable region	**PGT I51**	**Falkowska, E.** et al "Broadly Neutralizing **HIV** Antibodies Define **a** Glycan-Dependent Epitope on the Prefusion Conformation of gp4 1 on Cleaved Envelope **Trimers"** Immunity 40 **(5),** 657-668 (2014), NCBI Accession # AIC32543	9165
HIV1547	Light **chain variable** region	**PGT152**	Falkowska, E. et al "Broadly Neutralizing **HIV** Antibodies Define a **Glycan-Dependent** Epitope on the Prefusion **Conformation** of gp41 on Cleaved Envelope Trimers" Immunity 40 **(5),** 657-668 **(20 14),** NCBI Accession # AIC32544	9166
HIV1548	Light chain variable region	PGT153	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a Glycan-Dependent Epitope on the Prefusion **Conformation** of gp41 on Cleaved Envelope Trimers" **Immunity** 40 (5), 657-668 (2014), NCBI Accession # AIC32545	9167
HIV1549	Light **chain** variable region	PGT154	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a Glycan-Dependent Epitope on the Prefusion **Conformation** of gp41 on Cleaved Envelope Trimers" Immunity 40 **(5),** 657-668 (2014), NCBI Accession # AIC32529	9168
HIV1550	Light chain variable region	PGT155	Falkowska, E. et al "Broadly Neutralizing HIV Antibodies Define a Glycan-Dependent Epitope on the Prefusion Conformation **of gp41 on** Cleaved Envelope Trimers" **Immunity** 40 (5), 657-668 (2014), NCBI **Accession # AIC32547**	9169
HIV1551	Light chain **variable** region	**PGT156**	Falkowska, E. et al "Broadly Neutralizing **HIV** Antibodies Define **a** Glycan-Dependent Epitope on the Prefusion Conformation of **gp41** on Cleaved Envelope **Trimers" Immunity** 40 (5), 657-668 (2014), NCBI **Accession** # AIC32548	9170
HIV1552	Light **chain variable** region	**PGT157**	Falkowska, E. et al "Broadly Neutralizing **HIV** Antibodies Define a Glycan-Dependent Epitope on the Prefusion **Conformation** of gp41 on Cleaved Envelope **Trimers" Immunity** 40 (5), 657-668 (2014), NCBI Accession # AEC32549	9171
HIV1553	Light chain variable region	**PGT I58**	Falkowska, **E.** et al "Broadly Neutralizing **HIV** Antibodies Define **a** Glycan-Dependent Epitope on the **Prefusion** Conformation **of** gp4 1 on Cleaved Envelope Trimers" Immunity 40 (5), 657-668 (2014), NCBI Accession # AIC32550	9172
HIV1554	Light **chain variable** region	rF105	WGT993012232 SEQ ID NO: 3	9173
HIV1555	Light chain variable region	**ScFvX5-CD4**	US7378093B2 SEQ ID NO: 15	9174
HIV1556	Light **chain variable** region	**TNX-355, Idalizumab**	**US2013019588 1** SEQ ID NO: 1	9175

HIV1557	Light chain variable region	VCR14	US20150044137 SEQ ID NO: 25	9176
HIV1558	Light chain variable region	VCR14b	US20150044137 SEQ ID NO: 26	9177
HIV1559	Light chain variable region	VCR14c	US20150044137 SEQ ID NO: 27	9178
HIV1560	Light chain variable region	VCR16	US20150044137 SEQ ID NO: 33	9179
HIV1561	Light chain variable region	VCR16b	US20150044137 SEQ ID NO: 34	9180
HIV1562	Light chain variable region	VCR16c	US20150044137 SEQ ID NO: 35	9181
HIV1563	Light chain variable region	VCR16d	US20150044137 SEQ ID NO: 36	9182
HIV1564	Light chain variable region	VRC13	US20150044137 SEQ ID NO: 17	9183
HIV1565	Light chain variable region	VRC13b	US20150044137 SEQ ID NO: 18	9184
HIV1566	Light chain variable region	VRC13c	US20150044137 SEQ ID NO: 19	9185
HIV1567	Light chain variable region	VRC13d	US20150044137 SEQ ID NO: 20	9186
HIV1568	Light chain variable region	VRC13e	US20150044137 SEQ ID NO: 21	9187
HIV1569	Light chain variable region	VRC13f	US20150044137 SEQ ID NO: 22	9188
HIV1570	Light chain variable region	VRC13g	US20150044137 SEQ ID NO: 23	9189
HIV1571	Light chain variable region	VRC13h	US20150044137 SEQ ID NO: 24	9190
HIV1572	Light chain variable region	VRC15	US20150044137 SEQ ID NO: 28	9191
HIV1573	Light chain variable region		US20150004190 SEQ ID NO: 57	9192
HIV1574	Light Chain, Fab	Ch04	McLellan, J.S. et al., Structure of HIV-1 gp120 V1 V2 domain with broadly neutralizing antibody PG9; Nature 480 (7377), 336-343 (2011), NCBI Accession # 3TCL_B (215aa)	9193
HIV1575	Light Chain, Fab	N5-i5	Acharya, P., et al., Structural Definition of an Antibody-Dependent Cellular Cytotoxicity Response Implicated in Reduced Risk for HIV-1 Infection; J. Virol. 88 (21), 12895-12906 (2014), NCBI Accession # 4H8W_L (217aa)	9194
HEV1576	Light Cham, Ig Kappa Chain C Region	Nih45~46 Fab	Diskin, R., et al., Science 334 (6060), 1289-1293 (2011), NCBI Accession # 3U7Y_L (210aa)	9195
HIV1577	Light Chain, Ig Lanibda-2 Chain C region	Pgtl27	Pejchal, R., et al., Science 334 (6059), 1097-1103 (2011), NCBI Accession # 3TWC_L(211aa)	9196
HIV1578	Light Chain, Ig Kappa Chain C Region	7b2	Santra, S., et al., PLoS Pathol. 11 (8), E1005042 (2015), NCBI Accession # 4YDV_L (265aa)	9197
HIV1579	Light Chain; Fab	N60-i3	Gohain, N., et al., Cociystal Structures of Antibody N60-i3 and Antibody JR4 in Complex with gpl20 Define More Cluster A Epitopes Involved in Effective Antibody-Dependent Effector Function against HIV-1; J. Virol. 89 (17), 8840-8854 (2015), NCBI Accession # 4RFO_L (221aa)	9198

HIV1580	Light Chain; Ig Lambda-2 Chain C region	Pgt128	Pejchal, R., et al., Science 334 (6059), 1097-1103 (2011), NCBI Accession # 3TV3_L (211aa)	9199
HIV1581	Scfv	B11	US7744887B2 SEQ ID NO: 8	9200
HIV1582	Scfv		US8110192B2 SEQ ID NO: 1	9201
HIV1583	Scfv		US8110192B2 SEQ ID NO: 2	9202
HIV1584	Scfv		US8110192B2 SEQ ID NO: 3	9203
HIV1585	Scfv (SEQRES)	3b3 variant	Clark et al., Protein Sci. 18 (12), 2429-2441 (2009), NCBI Accession # 3JUY_A (256aa)	9204
HIV1586	Scfv	D5	US7744887B2 SEQ ID NO: 2	9205
HIV1587	Scfv-cd4 fusion protein		US8110192B2 SEQ ID NO: 8	9206
HIV1588		447-52d	Dhillon, A.K., et al., Acta Crystallogr. D Biol. Crystallogr. D64 (PT 7), 792-802 (2008), NCBI Accession # 3C2A_I(231aa)	9207
HIV1589		447-52d	Dhillon, A.K., et al., Acta Crystallogr. D Biol. Crystallogr. D64 (PT 7), 792-802 (2008), NCBI Accession # 3C2A_M (216aa)	9208
HIV1590		F105	Wilkinson, R.A., et al., J. Virol. 79 (20), 13060-13069 (2005), NCBI Accession # 1U6A_H (224aa)	9209
HIV1591		F105	Wilkinson, R.A., et al., J. Virol. 79 (20), 13060-13069 (2005), NCBI Accession # 1U6A_L (215aa)	9210
HIV1592		Fab 8062	Frisch, C., et al., PLoS Pathol. 6 (11), E1001182 (2010), NCBI Accession # 3MAC_H (245aa)	9211
HIV1593		Fab 8062	Frisch, C., et al., PLoS Pathol. 6 (11), E1001182 (2010), NCBI Accession # 3MAC_L (213aa)	9212
HIV1594		Fab 8066	Frisch, C., et al., PLoS Pathol. 6 (11), E1001182 (2010), NCBI Accession # 3MA9_H (245aa)	9213
HIV1595		Fab 8066	Frisch, C., et al., PLoS Pathol. 6 (11), E1001182 (2010), NCBI Accession # 3MA9_L (213aa)	9214
HIV1596		Fab'2F5 fragment	US6482928 SEQ ID NO: 6	9215
HIV1597		Fab'2F5 fragment	US6482928 SEQ ID NO: 7	9216
HIV1598		M18 Fab	Prabakaran, P., et al., J. Mol. Biol. 357 (1), 82-99 (2006), NCBI Accession # 2AJ3_D (228aa)	9217
HIV1599		M18 Fab	Prabakaran, P., et al., J. Mol. Biol. 357 (1), 82-99 (2006), NCBI Accession # 2AJ3_E (213aa)	9218
HIV1600		Pg16	Pancera, M. et al., J. Virol. 84 (16), 8098-8110 (2010), NCBI Accession # 3MME_A (238aa)	9219
HIV1601		Pg16	Pancera, M. et al., J. Virol. 84 (16), 8098-8110 (2010), NCBI Accession # 3MME_B (216aa)	9220

[ooo353] In one embodiment, the payload region of the AAV particle comprises one or more nucleic acid sequences, fragment or variants ihereofor encodes one or more polypeptides, fragments or variants thereof described in European Patent Pubhcation No. EP327000, EP478689, EP554401, EP581353 and EP711439, US Publication No. US201 10104163, US20110212106, US20130215726 and IJS20130251726, US patent No. US5266479, US5804440, US6657050, US8637036, and US9090675, and international Publication No.

WO2012154312, WO2013 163427, WO2014043386, WO2015048462, WO2015048610, WO2015048770 the contents of each of which are herein incorporated by reference in their entirety, against HIV.

Disease Specific Epitopes, Innate Defense Regulator Peptides, Cyclic Peptides

[00354] In one embodiment, the viral genomes of the AAV particles may comprise nucleic acids which have been engineered to enable expression of antibodies binding to disease-specific epitopes of proteins. Such antibodies may be used to diagnose, prevent, and/or treat the corresponding medical conditions by targeting epitopes of the protein presented by or accessible on native or non-native forms (e.g., misfolded forms of native proteins) of the target. Such epitopes may be specific to diseases involved with misfoiding of a protein due to pathologic condition and resulting in misfolded aggregates. The disease-specific proteins are considered to be toxic to neurons and to have a role in neuronal cell death and dysfunction in neurodegenerative diseases including, but not limited to. Alzheimer's disease (AD), amyotrophic lateral sclerosis (ALS), Parkinson's disease, dementia by Lewy body (DLB), and prion diseases, e.g. Creutzfeldt-Jakob disease (CJD), Gerstmann-Straussler-Scheinker syndrome (GSS), kuru, and fatal familial insomnia (FFI).

[00355] In one embodiment, the encoded disease-specific epitopes may include epitopes on SOD1 that are revealed as SODl (Superoxide dismutase [Cu-Zn]) dissociates from its homodimeric, normal state. The SOD epitopes may be selectively presented or accessible in non-native SODl forms including misfolded SODl monomer, misfolded SOD1 dimer, and the epitopes selectively presented or accessible in SODl aggregates. Such epitopes may be specific to neurodegenerative diseases including, but not limited to, amyotrophic lateral sclerosis (ALS), Alzheimer's (AD), Parkinson's (PD), and Lewy body diseases (LBD).

[00356] In one embodiment, the expressed antibodies may bind to epitopes presented by or accessible on non-native forms of SODl, such as those presented by SED ID NO: 2, 3, 5, 6, and 7 of US Patent No. IJS7977314 (the contents of which are herein incorporated by reference in its entirety), or presented by or accessible on monomelic forms of SOD1, such as those presented by SEQ ID NOs: 1 and 4 of US Patent No. US7977314, the contents of which are herein incorporated by reference in their entirety. In one embodiment, the expressed antibodies may comprise isolated peptides corresponding to such epitopes, such as those presented in SEQ ID NOs: 1-8 or SEQ ID NOs: 8-16, or epitopes presented by SEQ ID NOs: 34-63, 65-79 of US Patent No. US7977314, the contents of which are herein incorporated by reference in their entirety.

[00357] In one embodiment, the encoded disease-specific epitopes may be specific to diseases associated with prion protein (PrP); familial amyloid polyneuropathy or senile systemic amyloidosis or a disease related by the presence of misfolded transthyretine (TTR); renal accumulation of β2 microglobulin amyloid deposits or a disease related by the presence of misfolded β2 microglobulin, amyotrophic lateral sclerosis (ALS) or a disease related by the presence of misfolded SODl; Ieukemias or myelomas or a disease related by the presence of misfolded cluster of differentiation 38 (CD38); colon cancer metastasis and or a disease related by the presence of misfolded cluster of differentiation (CD44); tumors associated with tumor necrosis factor receptor (TNFR); cancers including cervical, head and neck, endometrial, lung and breast carcinomas, pleural mesotheliomas, malignant melanomas, Hodgkin lymphomas, anaplastic large cell non-Hodgkm lymphomas, or a disease related by the presence of misfolded Notch homolog 1 (NOTCH 1) e.g. acute myeloid Ieukemias and B-ceil chronic lymphoid Ieukemias; cancer in which Fas receptor (FasR) is implicated; cancers and related disorders in which misfolded epidermal growth factor (EGFR) is implicated; and/or other related diseases, disorders and conditions.

[00358] In one embodiment, the encoded disease specific epitopes may include epitopes that are revealed as the proteins misfold. In one embodiment, the expressed antibodies may bind to predicted epitopes of human PrP, such as those presented by SEQ ID NOs: 1-10 of US Patent Publication No. US201 00233 176; bovine PrP, such as those presented by SEQ ID NOs: 11-15 of US Patent Publication No. US201 00233 176, TTR, such as those presented by SEQ ID NOs: 16-22 of US Patent Publication No. US20 100233 176; beta-2 microglobulin, such as those presented by SEQ ID NOs: 23-26 of US Patent Publication No. US20100233176; SODl, such as those presented by SEQ ID NOs: 27-40 of US Patent Publication No. US20100233 S76; CD38, such as those presented by SEQ ID NOs: 41-45 of US Patent Publication No. US20I002331 76; CD44, such as those presented by 46-50 of US Patent Publication No. US20100233176; TNFR, such as those presented by 51-55 of US Patent Publication No. US20100233176; notch protein, such as those presented in SEQ ID NOs: 56-60 of US Patent Publication No. US20100233 S76; FasR, such as those presented by SEQ ID NOs: 61-65 of US Patent Publication No. US20100233176 and EGFR, such as those presented by SEQ ID NOs: 66-80 of US Patent Publication No. US20 100233 176; the contents of which are herein incorporated by reference in their entirety.

[ΘΘ359] In one embodiment, the expressed antibodies may comprise peptides corresponding to such epitopes. In one embodiment, the expressed antibodies may comprise pnon-specific pept des, such as those presented by SEQ ID NOs: 81-88 of US Patent Publication No.

US20 100233 176, the contents of which are herem incorporated by reference in their entirety, and variations thereof.

[ΘΘ360] In one embodiment, the encoded disease-specific epitopes may be specific to prion diseases, including transmissible spongiform encephalopathies (TSEs) or other prion diseases. In one embodiment, the expressed antibodies may bind to predicted epitopes of PrP, such as those presented by SEQ ID NOs: 24, 26, 28, 30, 32, 34, 36, 39-43, of US Patent Publication No. US2G150004185, the contents of which are herem incorporated by reference in their entirety. In one embodiment, the expressed antibodies may comprise prion-specific peptides or peptide fusions, such as those presented by SEQ ID NOs: 12-23, 25, 27, 29, 31, 33, 35, 37, 38, 43, and 44-48 of US Patent Publication No. US201 50004185, the contents of which are herein incorporated by reference in their entirety.

[ΘΘ36I] In one embodiment, the expressed antibodies may comprise prion peptides binding to prion specific abnormal isoform of the prion protein, such as those presented by SEQ ID NOs: 2-10 of US Patent Publication No. US20040072236, the contents of which are herein incorporated by reference in their entirety.

[00362] In one embodiment, the viral genomes of the AAV particles may comprise nucleic acids which have been engineered to express innate defense regulator (IDR) peptides. TDRs are immunomodulatory peptides that act directly on cells to effect an innate immune response. Such IDRs may be used to treat neurodegenerative diseases associated with neuroinflammation, e.g. amyotrophic lateral sclerosis (ALS), Alzheimer's disease, Friedreich's ataxia, Huntington's disease, Lewy body disease, Parkinson's disease, spinal muscular atrophy, and multiple sclerosis (MS) and other neurodegenerative diseases. In one embodiment, IDRs may be those presented by SEQ ID NOS: 1-969, and 973-1264 of International Publication No. WO2013034982, the contents of which are herein incorporated by reference in their entirety, or analogs, derivatives, anndated variations and conservative variations thereof.

[00363] In one embodiment, the viral genomes of the AAV particles may comprise nucleic acids which have been engineered to express antibodies binding to an epitope of the Tropomyosin receptor kinase (TrkC) receptor. Such antibodies may comprise a peptide, such as one presented by SEQ ID NO: 1 of US Patent No. US9200080, the contents of which are herein incorporated by reference in their entirety.

[00364] In some embodiments, the viral genomes of the AAV particles may comprise nucleic acids which have been engineered to express cyclic peptides with an amino acid sequence SNK. Non-limiting examples of other cyclic peptides include SEQ ID NO: 1-7 of US Patent No. US92I6217, the contents of which are herein incorporated by reference in their entirety. The

method of preparing the antibodies may include hyperimmune preparation method, as described in US Patent No. US9216217, the contents of which are herein incorporated by reference in their entirety.

Prions

[00365] In one embodiment, the viral genomes of the AAV particles may compose a nucleic acid sequence encoding antibodies comprising prion peptides comprising prion epitopes, and fusions and repeats thereof, such as those presented by SEQ ID NOs: 8-32, 35, and 36 of US Patent No. US9056918, the contents of which are herein incorporated by reference in their entirety.

[00366] In one embodiment, the viral genomes of the AAV particles may comprise a nucleic acid sequence encoding prion binding proteins (PrPBP). In one embodiment, the PrPBPs are cadherins, such as those presented by SEQ ID NOs: 1 and 2 of International Publication WO1997/045746, the contents of which are herein incorporated by reference in their entirety. In one embodiment, the PrPBPs are cadherins, such as those presented by SEQ ID NOs: 2 and 7-9 of international Publication No. WO2001000235, the contents of which are herein incorporated by reference in their entirety.

The nature of the polypeptides and variants

[00367] Antibodies encoded by payload regions of the viral genomes of the invention may be translated as a whole polypeptide, a plurality of polypeptides or fragments of polypeptides, which independently may be encoded by one or more nucleic acids, fragments of nucleic acids or variants of any of the aforementioned. As used herein, "polypeptide" means a polymer of amino acid residues (natural or unnatural) linked together most often by peptide bonds. The term, as used herein, refers to proteins, polypeptides, and peptides of any size, structure, or function. In some instances, the polypeptide encoded is smaller than about 50 amino acids and the polypeptide is then termed a peptide. If the polypeptide is a peptide, it will be at least about 2, 3, 4, or at least 5 amino acid residues long. Thus, polypeptides include gene products, naturally occurring polypeptides, synthetic polypeptides, homologs, orthologs, paralogs, fragments and other equivalents, variants, and analogs of the foregoing. A polypeptide may be a single molecule or may be a multi-molecular complex such as a dimer, trimer or tetramer. They may also comprise single chain or multichain polypeptides and may be associated or linked. The term polypeptide may also apply to amino acid polymers in which one or more amino acid residues are an artificial chemical analogue of a corresponding naturally occurring amino acid.

[00368] The term "polypeptide variant" refers to molecules which differ in their amino acid sequence from a native or reference sequence. The amino acid sequence variants may possess

substitutions, deletions, and/or insertions at certain positions within the amino acid sequence, as compared to a native or reference sequence. Ordinarily, variants will possess at least about 50% identity (homology) to a native or reference sequence, and preferably, they will be at least about 80%, more preferably at least about 90% identical (homologous) to a native or reference sequence.

[00369] In some embodiments "variant mimics" are provided. As used herein, the term "variant mimic" is one which contains one or more amino acids which would mimic an activated sequence. For example, glutamate may serve as a mimic for phosphoro-threonine and/or phosphoro-serine. Alternatively, variant mimics may result in deactivation or in an inactivated product containing the mimic, e.g., phenylalanine may act as an inactivating substitution for tyrosine; or alanine may act as an inactivating substitution for serine.

[00370] The term "amino acid sequence variant" refers to molecules with some differences in their amino acid sequences as compared to a native or starting sequence. The amino acid sequence variants may possess substitutions, deletions, and/or insertions at certain positions within the amino acid sequence. "Native" or "starting" sequence should not be confused with a wild type sequence. As used herein, a native or starting sequence is a relative term referring to an original molecule against which a comparison may be made. "Native" or "starting" sequences or molecules may represent the wild-type (that sequence found in nature) but do not have to be the wild-type sequence.

[00371] Ordinarily, variants will possess at least about 70% homology to a native sequence, and preferably, they will be at least about 80%, more preferably at least about 90% homologous to a native sequence. "Homology" as it applies to amino acid sequences is defined as the percentage of residues in the candidate amino acid sequence that are identical with the residues in the amino acid sequence of a second sequence after aligning the sequences and introducing gaps, if necessary-, to achieve the maximum percent homology. Methods and computer programs for the alignment are well known in the art. It is understood that homology depends on a calculation of percent identity but may differ in value due to gaps and penalties introduced in the calculation.

[00372] By "homologs" as it applies to amino acid sequences is meant the corresponding sequence of other species having substantial identity to a second sequence of a second species.

[00373] "Analogs" is meant to include polypeptide variants which differ by one or more amino acid alterations, e.g., substitutions, additions or deletions of amino acid residues that still maintain the properties of the parent polypeptide.

[00374] Sequence tags or amino acids, such as one or more lysines, can be added to the peptide sequences of the invention (e.g., at the N-terminal or C-terminal ends). Sequence tags can be used for peptide purification or localization. Lysines can be used to increase peptide solubility or to allow for biotmyiation. Alternatively, amino acid residues located at the carboxy and amino terminal regions of the amino acid sequence of a peptide or protein may optionally be deleted providing for truncated sequences. Certain amino acids (e.g., C-terminal or N-terminal residues) may alternatively be deleted depending on the use of the sequence, as for example, expression of the sequence as pari of a larger sequence which is soluble, or linked to a solid support.

[00375] "Substitutional variants" when referring to proteins are those that have at least one amino acid residue in a native or starting sequence removed and a different amino acid inserted in its place at the same position. The substitutions may be single, where only one amino acid in the molecule has been substituted, or they may be multiple, where two or more amino acids have been substituted in the same molecule.

[00376] As used herein the term "conservative ammo acid substitution" refers to the substitution of an amino acid that is normally present in the sequence with a different amino acid of similar size, charge, or polarity. Examples of conservative substitutions include the substitution of anon-polar (hydrophobic) residue such as isoleucine, valine and leucine for another non-polar residue. Likewise, examples of conservative substitutions include the substitution of one polar (hydrophilic) residue for another such as between arginine and lysine, between giutaraine and asparagine, and between glycine and serine. Additionally, the substitution of a basic residue such as lysine, arginine or histidiiie for another, or the substitution of one acidic residue such as aspartic acid or glutamic acid for another acidic residue are additional examples of conservative substitutions. Examples of non-conservative substitutions include the substitution of a non-polar (hydrophobic) amino acid residue such as isoleucine, valine, leucine, alanine, methionine for a polar (hydrophilic) residue such as cysteine, glutanime, glutamic acid or lysine and/or a polar residue for a non-polar residue.

[00377] "Insertional variants" when referring to proteins are those with one or more amino acids mseited immediately adjacent to an amino acid at a particular position in a native or starting sequence. "immediately adjacent" to an amino acid means connected to either the alpha-carboxy or alpha-amino functional group of the amino acid.

[00378] "Deletional variants" when referring to proteins, are those with one or more amino acids in the native or starting amino acid sequence removed. Ordinarily, deletional variants will have one or more amino acids deleted in a particular region of the molecule.

[00379] As used herein, the term "derivative" is used synonymously with the term "variant" and refers to a molecule that has been modified or changed in any way relative to a reference molecule or starting molecule. In some embodiments, derivatives include native or starting proteins that have been modified with an organic proteinaceous or non-proteinaceous derivatizing agent, and post-translational modifications. Covalent modifications are traditionally introduced by reacting targeted amino acid residues of the protein with an organic derivatizing agent that is capable of reacting with selected side-chains or terminal residues, or by harnessing mechanisms of post-translational modifications that function in selected recombinant host cells. The resultant covalent derivatives are useful in programs directed at identifying residues important for biological activity, for immunoassays, or for the preparation of anti-protein antibodies for immunoaffinity purification of the recombinant glycoprotein. Such modifications are within the ordinary skill in the art and are performed without undue experimentation.

[00380] Certain post-translational modifications are the result of the action of recombinant host cells on the expressed polypeptide. Glutaminyl and asparaginyl residues are frequently post-translationally deamidated to the corresponding glutamyl and aspartyl residues. Alternatively, these residues are deamidated under mildly acidic conditions. Either form of these residues may be present in the proteins used in accordance with the present invention.

[00381] Other post-translational modifications include hydroxylation of proline and lysine, phosphorylation of hydroxyl groups of seryl or threonyl residues, methylation of the alpha-amino groups of lysine, arginine, and histidine side chains (T. E. Creighton, Proteins: Structure and Molecular Properties, W.H. Freeman & Co., San Francisco, pp. 79-86 (1983)).

[00382] "Features" when referring to proteins are defined as distinct amino acid sequence-based components of a molecule. Features of the proteins of the present invention include surface manifestations, local conformational shape, folds, loops, half-loops, domains, half-domains, sites, termini or any combination thereof.

[00383] As used herein when referring to proteins the term "surface manifestation" refers to a polypeptide based component of a protein appearing on an outermost surface.

[00384] As used herein when referring to proteins the term "local conformational shape" means a polypeptide based structural manifestation of a protein which is located within a definable space of the protein.

[00385] As used herein when referring to proteins the term "fold" means the resultant conformation of an amino acid sequence upon energy minimization. A fold may occur at the secondary or tertiary level of the folding process. Examples of secondary level folds include beta sheets and alpha helices. Examples of tertiary folds include domains and regions formed due to

aggregation or separation of energetic forces. Regions formed in this way include hydrophobic and hydrophilic pockets, and the like.

[00386] As used herein the term "turn" as it relates to protein conformation means a bend which alters the direction of the backbone of a peptide or polypeptide and may involve one, two, three or more amino acid residues.

[00387] As used herein when referring to proteins the term "loop" refers to a structural feature of a peptide or polypeptide which reverses the direction of the backbone of a peptide or polypeptide and comprises four or more amino acid residues. Oiiva et al. have identified at least 5 classes of protein loops (J. Mol Biol 266 (4): 814-830; 1997).

[00388] As used herein when referring to proteins the term "half-loop" refers to a portion of an identified loop having at least half the number of amino acid residues as the loop from which it is derived. It is understood that loops may not always contain an even number of amino acid residues. Therefore, in those cases where a loop contains or is identified to comprise an odd number of amino acids, a half-loop of the odd-numbered loop will comprise the whole number portion or next whole number portion of the loop (number of amino acids of the ioop/2+/-0.5 amino acids). For example, a loop identified as a 7 amino acid loop could produce half-loops of 3 amino acids or 4 amino acids (7/2=3.5+/~Q.5 being 3 or 4).

[00389] As used herein when referring to proteins the term "domain" refers to a motif of a polypeptide having one or more identifiable structural or functional characteristics or properties (e.g.. binding capacity, serving as a site for protein-protein interactions).

[00390] As used herein when referring to proteins the term "half-domain" means portion of an identified domain having at least half the number of amino acid residues as the domain from which it is derived. It is understood that domains may not always contain an even number of amino acid residues. Therefore, in those cases where a domain contains or is identified to comprise an odd number of amino acids, a half-domain of the odd-numbered domain will comprise the whole number portion or next whole number portion of the domain (number of amino acids of the domain/2+/~0.5 amino acids). For example, a domain identified as a 7 amino acid domain could produce half-domains of 3 amino acids or 4 amino acids (7/2=3.5+/-0.5 being 3 or 4). It is also understood that sub-domains may be identified within domains or half-domains, these subdomains possessing less than all of the structural or functional properties identified in the domains or half domains from which they were derived. It is also understood that the amino acids that comprise any of the domain types herein need not be contiguous along the backbone of the polypeptide (i.e., nonadjacent amino acids may fold structurally to produce a domain, half-domain or subdomain).

[00391] As used herein when referring to proteins the terms "site" as it pertains to amino acid based embodiments is used synonymous with "amino acid residue" and "amino acid side chain". A site represents a position within a peptide or polypeptide that may be modified, manipulated, altered, derivatized or varied within the poiypeptide based molecules of the present invention.

[00392] As used herein the terms "termini or terminus" when referring to proteins refers to an extremity of a peptide or polypeptide. Such extremity is not limited only to the first or final site of the peptide or poiypeptide but may include additional amino acids in the terminal regions. The polypeptide based molecules of the present invention may be characterized as having both an N-terminus (terminated by an amino acid with a free amino group (NH2)) and a C-terminus (terminated by an amino acid with a free carboxyl group (COOH)). Proteins of the invention are in some cases made up of multiple poiypeptide chains brought together by disulfide bonds or by non-covalent forces (multimers, oligomers). These sorts of proteins will have multiple N- and C-termini. Alternatively, the termini of the polypeptides may be modified such that they begin or end, as the case may be, with a non-polypeptide based moiety such as an organic conjugate.

[00393] Once any of the features have been identified or defined as a component of a molecule of the invention, any of several manipulations and/or modifications of these features may be performed by moving, swapping, inverting, deleting, randomizing or duplicating. Furthermore, it is understood that manipulation of features may result in the same outcome as a modification to the molecules of the invention. For example, a manipulation which involves deleting a domain would result in the alteration of the length of a molecule just as modification of a nucleic acid to encode less than a full-length molecule would.

[00394] Modifications and manipulations can be accomplished by methods known in the art such as site directed mutagenesis. The resulting modified molecules may then be tested for activity using in vitro or in vivo assays such as those described herein or any other suitable screening assay known in the art.

AAV Production

[ΘΘ395] The present invention provides methods for the generation of parvoviral particles, *e.g.* AAV particles, by viral genome replication in a viral replication cell.

[00396] In accordance with the invention, the viral genome comprising a payload region encoding an antibody, an antibody-based composition or fragment thereof, will be incorporated into the AAV particle produced in the viral replication cell. Methods of making AAV particles are well known in the art and are described in e.g., United States patent Nos. US6204059, IJS5756283, IJS6258595, IJS6261551, IJS6270996, US6281010, US6365394, US6475769, US6482634, US6485966, US69430I9, US6953690, US70225I9. US7238526, US7291498 and

US7491508, US5064764, US6194191, US6566118, US8137948, or International Publication Nos. WO1996039530, WO1998010088, WO19990 14354, WO1999015685, WOI999047691, WO2000055342, WO2000075353 and WO2001023597; Methods In Molecular Biology, ed. Richard, Humana Press, NJ (1995); O'Reilly et ai., Bacuiovirus Expression Vectors, A Laboratory Manual, Oxford Univ. Press (1994); Samulski et al., *J Vir.*63:3822-8 (1989): Kajigaya et al., *Proa Nat'!. Acad. Sci. USA* 88: 4646-50 (1991); Ruffing et al., *J. Vir.* 66:6922-30 (1992): Kmibauer et al., i7r.219:37-44 (1996); Zhao et al, *Vir.*272:382-93 (2000): the contents of each of which are herein incorporated by reference in their entirety. In one embodiment, the AAV particles are made using the methods described in WO2015191508, the contents of which are herein incorporated by reference in their entirety.

[00397] Viral replication ceils commonly used for production of recombinant AAV viral vectors include but are not limited to 293 cells, COS cells, HeLa cells, KB cells, and other mammalian cell lines as described in U.S. Pat. Nos. US6156303, US5387484, US574I683, US5691I76, and US5688676; U.S. patent publication No. 2002/0081721, and International Patent Publication Nos. WO 00/47757, WO 00/24916, and WO 96/17947, the contents of each of which are herein incorporated by reference in their entireties.

[00398] In some embodiments, the present invention provides a method for producing an AAV particle having enhanced (increased, improved) transduction efficiency comprising the steps of: 1) co-transfecting competent bacterial cells with a bacmid vector and either a viral construct vector and/or AAV payload construct vector, 2) isolating the resultant viral construct expression vector and AAV payload construct expression vector and separately iransfectmg viral replication cells, 3) isolating and purifying resultant payload and viral construct particles comprising viral construct expression vector or AAV payload construct expression vector, 4) co-infecting a viral replication cell with both the AAV payload and viral construct particles comprising viral construct expression vector or AAV payload construct expression vector, and 5) harvesting and purifying the AAV particle comprising a viral genome.

[ΘΘ399] In some embodiments, the present invention provides a method for producing an AAV particle comprising the steps of I) simultaneously co-transfecting mammalian cells, such as, but not limited to I-IEK293 cells, with a payload region, a construct expressing rep and cap genes and a helper construct 2) harvesting and purifying the AAV particle comprising a viral genome.

[00400] In some embodiments, the viral genome of the AAV particle of the invention optionally encodes a selectable marker. The selectable marker may comprise a cell-surface marker, such as any protein expressed on the surface of the cell including, but not limited to receptors, CD markers, lectins, integrins, or truncated versions thereof.

[00401] In some embodiments, selectable marker reporter genes as described in International application No. WO 96/23810, Heim et al, Current Biology 2:178-1 82 (1996); Heim et al., Proc. Natl. Acad. Sci. USA (1995); or Heim et al., Science 373:663-664 (1995); WO 96/30540, the contents of each of which are incorporated herein by reference in their entireties).

II. FORMULATION AND DELIVERY

Pharmaceutical Compositions

[00402] According to the present invention the AAV particles may be prepared as pharmaceutical compositions. It will be imdersiood that such compositions necessarily comprise one or more active ingredients and, most often, a pharmaceutically acceptable excipient.

[ΘΘ403] Relative amounts of the active ingredient (e.g. AAV particle), a pharmaceutically acceptable excipient, and/or any additional ingredients in a pharmaceutical composition in accordance with the present disclosure may vary, depending upon the identity, size, and/or condition of the subject being treated and further depending upon the route by which the composition is to be administered. For example, the composition may comprise between 0.1% and 99% (w/w) of the active ingredient. By way of example, the composition may comprise between 0.1 % and 100%, e.g.. between .5 and 50%, between 1-30%, between 5-80%, at least 80% (w/w) active ingredient.

[00404] In some embodiments, the AAV particle pharmaceutical compositions described herein may comprise at least one payload. As a non-iimiting example, the pharmaceutical compositions may contain an AAV particle with 1, 2, 3, 4 or 5 payloads. In one embodiment, the pharmaceutical composition may contain a nucleic acid encoding a payload construct encoding proteins selected from antibodies and/or antibody-based compositions.

[00405] Although the descriptions of pharmaceutical compositions provided herein are principally directed to pharmaceutical compositions which are suitable for administration to humans, it will be understood by the skilled artisan that such compositions are generally suitable for administration to any other animal, e.g., to non-human animals, e.g. non-human mammals. Modification of pharmaceutical compositions suitable for administration to humans in order to render the compositions suitable for administration to various animals is well understood, and the ordinarily skilled veterinary pharmacologist can design and/or perform such modification with merely ordinary, if any, experimentation. Subjects to which administration of the pharmaceutical compositions is contemplated include, but are not limited to, humans and/or other primates: mammals, including commercially relevant mammals such as cattle, pigs, horses, sheep, cats, dogs, mice, rats, birds, including commercially relevant birds such as poultry, chickens, ducks, geese, and/or turkeys.

[00406] In some embodiments, compositions are administered to humans, human patients or subjects.

Formulations

[00407] The AAV particles of the invention can be formulated using one or more excipients to: (1) increase stability; (2) increase cell transfection or transduction; (3) permit the sustained or delayed expression of the payload; (4) alter the biodistribution (e.g., target the viral particle to specific tissues or cell types); (5) increase the translation of encoded protein; (6) alter the release profile of encoded protein and/or (7) allow for regulatable expression of the payload.

[00408] Formulations of the present invention can include, without limitation, saline, liposomes, lipid nanoparticles, polymers, peptides, proteins, cells Iransfected with viral vectors (e.g., for transfer or transplantation into a subject) and combinations thereof.

[00409] Formulations of the pharmaceutical compositions described herein may be prepared by any method known or hereafter developed in the art of pharmacology. As used herein the term "pharmaceutical composition" refers to compositions comprising at least one active ingredient and optionally one or more pharmaceutically acceptable excipients.

[00410] In general, such preparatory methods include the step of associating the active ingredient with an excipient and/or one or more other accessory ingredients. As used herein, the phrase "active ingredient" generally refers either to an AAV particle carrying a payload region encoding the polypeptides of the invention or to the antibody or antibody-based composition encoded by a viral genome of by an AAV particle as described herein.

[00411] Formulations of the AAV particles and pharmaceutical compositions described herein may be prepared by any method known or hereafter developed in the art of pharmacology. In general, such preparatory methods include the step of bringing the active ingredient into association with an excipient and/or one or more other accessor}' ingredients, and then, if necessary and/or desirable, dividing, shaping and/or packaging the product into a desired single- or multi-dose unit.

[00412] A pharmaceutical composition in accordance with the present disclosure may he prepared, packaged, and/or sold in bulk, as a single unit dose, and/or as a plurality of single unit doses. As used herein, a "unit dose" refers to a discrete amount of the pharmaceutical composition comprising a predetermined amount of the active ingredient. The amount of the active ingredient is generally equal to the dosage of the active ingredient which would be administered to a subject and/or a convenient fraction of such a dosage such as, for example, one-half or one-third of such a dosage.

[00413] In one embodiment, the AAV particles of the invention may be formulated in PBS with 0.001% of pluronic acid (F-68) at a pH of about 7.0.

[ΘΘ414] Relative amounts of the active ingredient (e.g. AAV particle), the pharmaceutically acceptable excipient, and/or any additional ingredients in a pharmaceutical composition in accordance with the present disclosure may vary, depending upon the identity, size, and/or condition of the subject being treated and further depending upon the route by which the composition is to be administered. For example, the composition may comprise between 0.1% and 99% (w/w) of the active ingredient. By way of example, the composition may comprise between 0.1 % and 100%, e.g., between 0.5 and 50%, between 1-30%, between 5-80%, at least 80% (w/w) active ingredient.

[ΘΘ415] In some embodiments, the AAV formulations described herein may contain sufficient AAV particles for expression of at least one expressed functional antibody or antibody-based composition. As anon-limiting example, the AAV particles may contain viral genomes encoding 1, 2, 3, 4 or 5 functional antibodies.

[00416] According to the present invention AAV particles may be formulated for CNS delivery. Agents that cross the brain blood barrier may be used. For example, some ceil penetrating peptides that can target molecules to the brain blood barrier endothelium may be used for formulation (e.g., Mathiipala, *Expert Opin Ther Pat.,* 2009, 19, 137-140; the content of which is incorporated herein by reference in its entirety).

Excipients and Diluents

[00417] The AAV particles of the invention can be formulated using one or more excipients or diluents to (1) increase stability, (2) increase cell transaction or transduction; (3) permit the sustained or delayed release; (4) alter the biodistribution (e.g., target the viral particle to specific tissues or ceil types); (5) increase the translation of encoded protein in vivo; (6) alter the release profile of encoded protein in vivo and/or (7) allow for regulatable expression of the polypeptides of the invention.

[00418] In some embodiments, a pharmaceutically acceptable excipient may be at least 95%, at least 96%, at least 97%, at least 98%, at least 99%, or 100% pure. In some embodiments, an excipient is approved for use for humans and for veterinary use. In some embodiments, an excipient may be approved by United States Food and Drug Administration. In some embodiments, an excipient may be of pharmaceutical grade. In some embodiments, an excipient may meet the standards of the United States Pharmacopoeia (LISP), the European Pharmacopoeia (EP), the British Pharmacopoeia, and/or the International Pharmacopoeia.

[00419] Excipients, as used herein, include, but are not limited to, any and all solvents, dispersion media, diluents, or other liquid vehicles, dispersion or suspension aids, surface active agents, isotonic agents, thickening or emulsifying agents, preservatives, and the like, as suited to the particular dosage form desired. Various excipients for formulating pharmaceutical compositions and techniques for preparing the composition are known in the art (see Remington: The Science and Practice of Pharmacy, 21st Edition, A. R. Gennaro, Lippincott, Williams & Wilkms, Baltimore, MD, 2006; incorporated herein by reference in its entirety). The use of a conventional excipient medium may be contemplated within the scope of the present disclosure, except insofar as any conventional excipient medium may be incompatible with a substance or Us derivatives, such as by producing any undesirable biological effect or otherwise interacting in a deleterious manner with any other component(s) of the pharmaceutical composition.

[ΘΘ420] Exemplary diluents include, hut are not limited to, calcium carbonate, sodium carbonate, calcium phosphate, dicalcium phosphate, calcium sulfate, calcium hydrogen phosphate, sodium phosphate lactose, sucrose, cellulose, microcrystalime cellulose, kaolin, mannito!, sorbitol, inositol , sodium chloride, dry starch, cornstarch, powdered sugar, *etc.*, and/or combinations thereof

Inactive Ingredients

[ΘΘ421] In some embodiments, AAV particle formulations may comprise at least one inactive ingredient. As used herein, the term 'inactive ingredient" refers to one or more agents that do not contribute to the activity of the active ingredient of the pharmaceutical composition included in formulations. In some embodiments, all, none or some of the inactive ingredients which may be used in the formulations of the present invention may be approved by the US Food and Drug Administration (FDA).

[Θ0422] In one embodiment, the AAV particle pharmaceutical compositions comprise at least one inactive ingredient such as, but not limited to, 1,2,6-Hexanetriol: 1,2-Dimyristoyl-Sn-Gly cero-3-(Phospho-S-(1-Gly ceroi)); 1,2-Dimyristoyl-Sn-Glycero-3-Phosphochoiine; 1,2-Dioleoyl-Sn-Glycero-3-Phosphocholine; l,2-Dipalmitoyl-Sn-Glycero-3-(Phospho-Rac-(l-Glyceroi)); l,2~Distearoyl-Sn-Glycero-3-(Phospho-Rac-(l-Giycerol»; 1,2-Distearoyl-Sn-G!ycero-3-Phosphochoiine; 1-O-Toiylbiguanide; 2-Ethyl-1,6-Hexanedioi; Acetic Acid; Acetic Acid, Glacial; Acetic Anhydride; Acetone, Acetone Sodium Bisulfite; Acetylated Lanolin Alcohols; Acetylated Monoglycerides; Acetylcysteine; Acetyitryptophan, DL-; Acrylates Copolymer; Acrylic Acid-Isooctyl Acrylate Copolymer; Acrylic Adhesive 788; Activated Charcoal; Adcote 72A103; Adhesive Tape; Adipic Acid; Aerotex Resin 3730; Alanine; Albumin Aggregated; Albumin. Colloidal; Albumin Human; Alcohol; Alcohol, Dehydrated; Alcohol,

Denatured; Alcohol., Diluted; Alfadex, Alginic Acid; Aikyl Ammonium Sulfonic Acid Beiarne; Aikyl Aryl Sodium Sulfonate; Allanloin; Ally! .Alpha.-Ionone; Almond Oil; Alpha-Terpineol; Alpha-Tocopherol; Alpha-Tocopherol Acetate, Di-; AIpha-Tocopherol, D1-; Aluminum Acetate; Aluminum Chlorliydroxy Allantoinate; Aluminum Hydroxide; Aluminum Hydroxide - Sucrose, Hydrated; Aluminum Hydroxide Gel; Aluminum Hydroxide Gei F 500; Aluminum Hydroxide Gel F 5000; Aluminum Monostearate; Aluminum Oxide; Aluminum Polyester; Aluminum Silicate; Aluminum Starch Octenylsuccinate; Aluminum Stearate; Aluminum Snbacetate; Aluminum Sulfate Anhydrous, Amerchol C; Amerchol-Cab, Aminomethylpropanol; Ammonia; Ammonia Solution; Ammonia Solution, Strong; Ammonium Acetate; Ammonium Hydroxide; Ammonium Lauryl Sulfate: Ammonium NonoxynoS-4 Sulfate; Ammonium Salt Of C-12-C-15 Linear Pnmary Aicohoi Ethoxylate; Ammonium Sulfate; Ammonyx; Aniphoteric-2; Amphoteric-9; Anethole; Anhydrous Citric Acid, Anhydrous Dextrose, Anhydrous Lactose, Anhydrous Trisodium Citrate; Aniseed Oil; Anoxid Sbn; Antifoam; Antipyrine; Apaflurane; Apricot Kernel Oil Peg-6 Esters; Aquaphor; Arginine; Arlacel; Ascorbic Acid; Ascorbyl Palmitate; Aspartic Acid, Balsam Peru, Barium Sulfate; Beeswax, Beeswax, Synthetic; Baheneth-I0, Bentonite; Benzalkonium Chloride; Benzenesuifonic Acid; Benzethonium Chloride; Benzododecinium Bromide; Benzoic Acid; Benzyl Alcohol; Benzyl Benzoate; Benzyl Chloride; Betadex; Bibapcitide; Bismuth Subgallate; Boric Acid; Brocnnat; Butane; Butyl Alcohol; Butyl Ester Of Vinyl Methyl Ether/Maleic Anhydride Copolymer (125000 Mw); Butyl Stearate; Butylated Hydroxyanisole; Butylated Hydroxytoluene, Butylene Glycol; Butyiparaben; Butyric Acid; C20-40 Pareth-24; Caffeine; Calcium; Calcium Carbonate; Calcium Chloride; Calcium Gluceptate; Calcium Hydroxide; Calcium Lactate; Calcobutroi; Caldiamide Sodium, Caloxetate Trisodium; Calteridol Calcium; Canada Balsam; Caprylic/Capric Triglyceride; Caprydio'Capric/Stearic Triglyceride; Captan; Captisol; Caramel; Carbomer 1342; Carbomer 1382; Carbomer 934; Carbomer 934p; Carbomer 940; Carbomer 941; Carbomer 980; Carbomer 981 ; Carbomer Homopolymer Type B (Allyl Pentaerythritol Crosslinked), Carbomer Homopolymer Type C (AlM Pentaerythritol Crosslinked), Carbon Dioxide; Carboxy Vinyl Copolymer; Carboxymethyleeliulose; Carboxymethylcellulose Sodium; Carboxy polymethyiene; Carrageenan; Carrageenan Salt, Castor Oil; Cedai· Leaf Oil; Cellulose; Cellulose, Macrocrystalline; Cerasynt-Se; Ceresin; Ceteareth-12; Ceteareth-15; Ceteareth-30; Cetearyl Alcohol/Ceteareth-20; Ceteaiyl Ethylhexanoate; Ceteth-10; Ceteth-2; Ceteth-20; Ceteth-23; Cetostearyl Aicohoi; Cetrimonium Chloride; Cetyl Alcohol; Cetyl Esters Wax; *Cetyl* Palmitate; Cetylpyridinium Chloride; Chlorohutanol; Chlorobutanol Hemihydrate; Chlorobutanol, Anhydrous; Chlorocresol; Chloroxylenol; Cholesterol; Choleth; Choleth-24; Citrate; Citric Acid;

Citric Acid Monohydrate; Citric Acid, Hydrous; Coeamide Ether Sulfate; Cocamine Oxide, Coco Betaine; Coco Diethanolamide; Coco Monoethanolaraide; Cocoa Butter; Coco-Glycerides; Coconut Oil; Coconut Oil, Hydrogenated; Coconut Oil/Palm Kernel Oil Glycerides, Hydrogenated; Cocoyl Caprylocaprate; ColaNitida Seed Extract; Collagen; Coloring Suspension, Corn Oil; Cottonseed Oil; Cream Base; Creatine; Creatinine; Cresol; CroscarmeUose Sodium; Crospovidone; Cupric Sulfate; Cupric Sulfate Anhydrous; Cyclometliicone; Cyclomethicone/Dimethicone Copolyoi; Cysteine; Cysteine Hydrochloride; Cysteine Hydrochloride Anhydrous; Cysteine, D1-; D&C Red No. 28; D&C Red No. 33; D&C Red No. 36; D&C Red No. 39, D&C Yellow No. 10; Dalfampridine; Daubert 1-5 Pestr (Matte) 164z; Decyl Methyl Sulfoxide; Dehydag Wax Sx; Dehydroacetic Acid; Dehymuls E; Denatonium Benzoate; Deoxychoiic Acid; Dextran; Dexiran 40; Dextrin; Dextrose; Dextrose Monohydrate, Dextrose Solution; Diatrizoic Acid; Diazolidiny] Urea; Diehlorohenzyl Alcohol; Dichlorodifluoromethane; Dichlorotetrafluoroethane; Diethanolamine; Diethyl Pyrocarbonate; Diethyl Sebacate; Diethylene Glycol Monoethyi Ether; Diethyihexyi Phthalate; Dihydroxy aluminum Ammoacetate; Diisopropanoiamine; Diisopropyl Adipate; Diisopropyl Dilinoieate, Dimethicone 350; Dimetbicone Copolyoi; Dimethicone Mdx4~42i0; Dimethicone Medical Fluid 360; Dimethyl Isosorbide; Dimethyl Sulfoxide; Diraethylaminoethyl Methacrylate - Butyl Methacrylate - Methyl Methacrylate Copolymer; Dimethyldioctadecylammonium Bentomte; Dimethylsiioxane/Methylvinylsiloxane (Copolymer; Dinoseb Ammonium Salt; Dipalmitoylphosphatidylglycerol, D1-; Dipropylene Glycol; Disodium Cocoampbodiacetate; Disodium Laureth Sulfosuccinate; Disodium Lauryi Sulfosuccinate; Disodium Sulfosaiicylate; Disofenin; Divinylbenzene Styrene Copolymer; Dmdm Hydantoin; Docosanol; Docusate Sodium, Duro-Tak 280-2516; Duro-Tak 387-2516; Duro-Tak 80-1196, Duro-Tak 87-2070; Duro-Tak 87-2194; Duro-Tak 87-2287; Duro-Tak 87-2296; Duro-Tak 87-2888; Duro-Tak 87-2979; Edetate Calcium Disodium; Edetate Disodium; Edetate Disodium Anhydrous; Edetate Sodium; Edetic Acid; Egg Phospholipids, Entsufon; Entsufon Sodium; Epilactose; Epitetracycline Hydrochloride; Essence Bouquet 9200; Ethanolamine Hydrochloride; Ethyl Acetate; Ethyl Oleate; Ethylcelluloses; Ethylene Glycol; Ethylene Vinyl Acetate Copolymer; Ethvlenediamme; Ethylenediamme Dihvdrochlonde, Ethylene-Propylene Copolymer; Ethylene-Vinyl Acetate Copolymer (28% Vinyl Acetate); Ethylene-Vinyl Acetate Copolymer (9% Vinylacetate); Ethylhexyl Hydroxystearate; Ethylparaben; Eucalyptol; Exaraetazime; Fat, Edible; Fat, Hard; Fatty Acid Esters; Fatty Acid Pentaerythnol Ester; Fatty Acids; Fatty Alcohol Citrate; Fatty Alcohols; Fd&C Blue No. 1; Fd&C Green No. 3; Fd&C Red No. 4; Fd&C Red No. 40; Fd&C Yellow No. 10 (Delisted); Fd&C Yellow No. 5; Fd&C Yellow No. 6; Ferrie Chloride;

Feme Oxide; Flavor 89-186; Flavor 89-259, Flavor Df-1 19; Flavor Df-1530; Flavor Enhancer; Flavor Fig 827118; Flavor Raspberry Pfc-8407; Flavor Rhodia Pharmaceutical No. Rf 451; Fhiorochlorohydrocarbons: Formaldehyde; Formaldehyde Solution; Fractionated Coconut Oil; Fragrance 3949-5; Fragrance 520a; Fragrance 6.007; Fragrance 91-122; Fragrance 912.8-Y; Fragrance 93498g; Fragrance Balsam Pine No. 5124; Fragrance Bouquet 10328; Fragrance Chemoderrn 6401-B; Fragrance Cheraoderra 6411; Fragrance Cream No. 73457; Fragrance Cs-28197; Fragrance Felton 066m; Fragrance Firmemch 47373; Fragrance Givaudan Ess 9090/1 c; Fragrance 11-6540; Fragrance Herbal 10396, Fragrance Nj-1085; Fragrance P O Fl-147, Fragrance Pa 52805; Fragrance Pera Derm D, Fragrance Rbd-9819; Fragrance Shaw Mudge U-7776; Fragrance Tf 044078; Fragrance Lingerer Honeysuckle K 2771; Fragrance Ongerer N5195; Fructose; Gadolinium Oxide; Galactose; Ganuna Cyclodextnn; Gelatin; Gelatin, Crosslinked; Geifoam Sponge; Gel Ian Gum (Low Acyl); Gelva 737; Gentisic Acid; Gentisic Acid Ethanolaroide; Gluceptate Sodium; Gluceptate Sodium Dihydrate; Gluconolactone; Glucuronic Acid; Glutamic Acid, D1-; Glutathione; Glycerin; Glycerol Ester Of Hydrogenated Rosin; Glyceryl Citrate; Glyceryl isostearate; Glyceryl Laurate; Glyceryl Monostearate; Glyceryl Oleate; Glyceryl Oleaxe/Propylene Glycol; Glyceryl Palmitate; Glyceryl Ricinoleate; Glyceryl Stearate; Glyceryl Stearate - Laureth-23; Glyceryl Stearate/Peg Stearate; Glyceryl Stearate/Peg-100 Stearate; Glyceryl Stearate/Peg-40 Stearate; Glyceryl Stearate-Stearamidoethyi Diethyiamine; Glyceryl Trioleate; Glycine; Glycine Hydrochloride; Glycol Distearate; Glycoi Stearate; Guanidine Hydrochloride; Guar Gum; Hair Conditioner (18n195-1m); Heptane, Hetastarch; Hexyiene Glycol; High Density Polyethylene; Histidme; Human Albumin Microspheres; Hyaluronate Sodium; Hydrocarbon; Hydrocarbon Gel. Plasucized; Flydrochloric Acid; Hydrochloric Acid, Diluted; Hydrocortisone; Hydrogel Polymer; Hydrogen Peroxide; Hydrogenated Castor Oil; Hydrogenated Palm Oil; Hydrogenated Palm/Palm Kernel Oil Peg-6 Esters; Hydrogenated Poiybutene 635-690; Hydroxide ion; Hydroxy ethyl Cellulose; Hydroxy ethyipiperazine Ethane Sulfonic Acid; Hydroxymethyl Cellulose; Hydroxy octacosanyl Hydroxy stearate; Hydroxy propyl Cellulose; Hydroxy propyl Methyicelluiose 2906; HydroxypiOpyi-Beta-cyciodextrin; Hypromellose 2208 (15000 MpaS); Hypronieliose 2910 (15000 Mpa.S); Hypromelloses; Imidurea; Iodine; lodoxamic Acid; iofetamine Hydrochloride; Irish Moss Extract; Isobutane; Isoceteth-20; Isoleucine; Isooctyl Acrylate; Isopropyi Alcohol; Tsopropyl isostearate; Isopropyi Myristate; Isopropyi Myristate - Myristyl Alcohol; Isopropyi Palmitate; Isopropyi Stearate; Isostearic Acid; isostearyi Alcohol; Isotonic Sodium Chloride Solution; Jelene, Kaolin; Kathon Cg; Katiion Cg II; Lactate; Lactic Acid; Lactic Acid. D1-; Lactic Acid. L~; Lactobionic Acid; Lactose; Lactose Monohydrate; Lactose. Hydrous; Laneth;

Lanolin; Lanolin Alcohol - Mineral Oil; Lanolin Alcohols; Lanolin Anhydrous; Lanolin Cholesterols; Lanolin Nonionic Derivatives; Lanolin, Ethoxylated; Lanolin, Hydrogenated, Lauralkonium Chloride; Lauraroine Oxide: Laurdimonium Hydrolyzed Animal Collagen; Laurefh Sulfate; Laureth-2; Laureth-23; Laureth-4; Laurie Diethanolamide; Laurie Myristic Diethanolamide, Lauroyl Sarcosine; Lauryl Lactate; Lauryi Sulfate; Lavandula Angustifolia Flowering Top; Lecithin; Lecithin Unbleached; Lecithin, Egg; Lecithin, Hydrogenated; Lecithin, Hydrogenated Soy; Lecithin, Soybean; Lemon Oil; Leucine; Levulmic Acid; Lidofenin; Light Mineral Oil; Light Mineral Oil (85 Ssu); Limonene, (+/-)-; Lipocoi Sc-15; Lysine, Lysine Acetate; Lysine Monohydrate, Magnesium Aluminum Silicate, Magnesium Aluminum Silicate Hydrate; Magnesium Chloride; Magnesium Nitrate; Magnesium Stearate: Maieic Acid; Mannitol; Maprofix; Mebrofemn: Medical Adhesive Modified S-15; Medical Antiform A-F Emulsion, Medronate Disodium; Medronic Acid, Meglumine; Menthol; Metacresol; Metaphosphoric Acid; Methanesulfonic Acid; Methionine; Methyl Alcohol; Methyl Gluceth-10; Methyl Glueeth-20; Methyl Gluceth-20 Sesquistearate; Methyl Glucose Sesquistearate; Methyl Laurate; Methyl Pyrroiidone: Methyl Salicylate, Methyl Stearate; Methyiboronic Acid; Methylcellulose (4000 Mpa.S); Methyicelluloses; Methylchloroisotbiazolinone; Methylene Blue, Methylisothiazolinone; Methylparaben: Microcrystallme Wax; Mineral Oil; Mono and Diglyceride; Monostearyl Citrate; Monothioglycerol; Multisterol Extract; Myristyl Alcohol; Myristyi Lactate; Myristyl-.Gamma.-Picolinium Chloride, N-(Carbamoyi-Methoxy Peg-40)-l,2-Distearoyl-Cephalm Sodium; N,N-Dimethylacetaraide; Niacinamide; Nioxime; Nitric Acid; Nitrogen; Nonoxynoi Iodine; Nonoxynol-15; Nonoxynol-9; Norflurane; Oatmeal; Octadecene-1/Maleic Acid Copolymer; Octanoic Acid; Octisalate; Octoxynol-1 ; Octoxynol-40; Octoxynol-9; Octyldodecarsol; Octyiphenol Polymethylene, Oleic Acid; Oleth-lO/Oieth-5; Oieth-2; Oleth-20; Oleyl Alcohol; Oleyl Oleate; Olive Oil; Oxidronate Disodium; Oxyquinoline; Palm Kernel Oil; Palniitamine Oxide; Parabens; Paraffin; Paraffin, White Soft; Parfiim Creme 45/3; Peanut Oil; Peanut Oil, Refined, Pectin; Peg 6-32 Stearate/Glycoi Stearate; Peg Vegetable Oil; Peg-100 Stearate; Peg-12 Glyceryl Laurate; Peg-120 Glyceryl Stearate; Peg-120 Methyl Glucose Dioleate; Peg-15 Cocamine; Peg-150 Distearate; Peg-2 Stearate: Peg-20 Sorbhan Isostearate; Peg-22 Methyl Ether/Dodecyl Glycol Copolymer: Peg-25 Propylene Glycol Stearate; Peg-4 Dilaurate; Peg-4 Laurate; Peg-40 Castor Oil; Peg-40 Sorbitan Diisostearate; Peg-45/Dodecyl Glycol Copolymer; Peg-5 Oleate; Peg-50 Stearate: Peg-54 Hydrogenated Castor Oil; Peg-6 isostearate; Peg-60 Castor Oil; Peg-60 Hydrogenated Castor Oil; Peg-7 Methyl Ether; Peg-75 Lanolin; Peg-8 Laurate; Peg-8 Stearate, Pegoxoi 7 Stearate; Pentadecalactone; Pentaerythritol Cocoate; Pentasodium Pentetate; Pentetate Calcium Trisodium; Pentetic Acid; Peppermint Oil;

Peril utren; Perfume 25677; Perfume Bouquet, Perfume E-1991; Perfume Gd 5604; Perfume Tana 90/42 Scba; Perfume W-l 952-1; Petrolatum; Petrolatum, White; Petroleum Distillates; Phenol; Phenol, Liquefied; Phenonip; Phenoxyethano!: Phenylalanine; Phenylethyl Alcohol; Phenylmercuric Acetate; Phenylmercuiic Nitrate; Phosphatidyl Glycerol, Egg; Phospholipid; Phospholipid, Egg; Phospholipon 90g; Phosphoric Acid; Pine Needle Oil (Pinus Sylvestris); Piperazine Hexahydrate; Plastibase-50w; Polacrilin; Polidronium Chloride; Poloxamer 124; Poloxamer 181; Poloxamer 182; Poloxamer 188; Poloxamer 237; Poloxamer 407; Poly(Bis(P~ Carboxy phenoxy)Propane Anhy dride) :Sebaci c Acid; Poly(Dimethylsik>xane/Methylvinyls^ xane/Methylhydrogensiloxane) Dimethylvinyl Or Dimethyihy droxy Or Trimethyl Endblocked; Po3y(Dl~Lactic-Co-G3ycoiic Acid), (50:50; Poly(Dl-Lactic-Co-Gly colic Acid), Ethyl Ester Terminated, (50:50; Poiyacrylic Acid (250000 Mw); Polybutene (1400 Mw); Polycarbophil; Polyester; Polyester Poly amine Copolymer; Polyester Rayon; Polyethylene Glycol 1000; Polyethylene Glycol 1450; Polyethylene Glycol 1500; Polyethylene Glycol 1540; Polyethylene Glycol 200; Polyethylene Glycol 300; Polyethylene Glycol 300-1600, Polyethylene Glycol 3350; Polyethylene Glycol 400; Polyethylene Glycol 4000; Polyethylene Glycol 540; Polyethylene Glycol 600; Polyethylene Glycol 6000; Polyethylene Glycol 8000: Polyethylene Glycol 900; Polyethylene High Density Containing Ferric Oxide Black (<1%); Polyethylene Low Density Containing Barium Sulfate (20-24%); Polyethylene T; Polyethylene Terephthaiates; Poiyglactin; Poiygly eery 1-3 Oleate; Polyglyceryl-4 Oleate; Polyhydroxyethyl Methacryiate; Polyisobutylene; Polyisobutylene (1100000 Mw); Polyisobutylene (35000 Mw); Polyisobutylene 178-236; Polyisobutylene 241-294; Polyisobutylene 35-39, Polyisobutylene Low Molecular Weight; Polyisobutylene Medium Molecular Weight; Polyisobu ty lene/Poly butene Adhesive; Polylactide; Polyols; Polyoxy ethylene - Polyoxypropylene 1800; Polyoxy ethylene Alcohols; Polyoxy ethylene Fatty Acid Esters; Polyoxy ethylene Propylene; Polyoxyi 20 Cetostearyl Ether; Polyoxyl 35 Castor Oil; Polyoxyl 40 HydiOgenated Castor Oil; Polyoxyl 40 Stearate; Polyoxyl 400 Stearate; Polyoxyl 6 And Polyoxyl 32 Palmitostearate; Polyoxyl Distearate; Polyoxyl Glyceryl Stearate; Polyoxyl Lanolin; Polyoxyl Palniitate; Polyoxyl Stearate; Polypropylene; Polypropylene Glycol; Polyquatemiuni-10; Polyquaternium-7 (70/30 Acryiamide/Dadmac; Polysiioxane; Poiysorbate 20; Polysorbate 40; Polysorbate 60; Polysorbate 65; Polysorbate 80; Polyurethane; Polyvinyl Acetate; Polyvinyl Alcohol; Polyvinyl Chloride; Polyvinyl Chloride-Poly vinyl Acetate Copolymer; Poly vmylpyridine; Poppy Seed Oil; Potash; Potassium Acetate; Potassium Alum; Potassium Bicarbonate; Potassium Bisulfite; Potassium Chloride; Potassium Citrate; Potassium Hydroxide; Potassium Metabi sulfite; Potassium Phosphate, Dibasic; Potassium Phosphate, Monobasic;

Potassium Soap: Potassium Sorbate; Povidone Aciylate Copolymer, Povidone Hydrogel; Povidone K17; Povidone K25; Povidone K29/32; Povidone K30; Povidone K90; Povidone K90f; Povidone/Eicosene Copolymer; Povidones; Ppg-12/Smdi Copolymer; Ppg-15 Stearyl Ether; Ppg-20 Methyl Glucose Ether Distearate; Ppg-26 Oleate; Product Wat; Proline; Promuigen D; Promulgen G; Propane; Propellant A-46; Propyl Gaiiate; Propylene Carbonate; Propylene Glycol: Propylene Glycol Diacetate: Propylene Glycol Dicaprylate; Propylene Glycol Monoiaurate; Propylene Glycol Monopalmitostearate; Propylene Glycol Palmitostearate; Propylene Glycol Ricinoleate; Propylene G!yeol/Diazoiidinyl Urea-'Methylparaben/Propylparben; Propylparaben; Protamine Sulfate; Protein Hydrolysate; Pvm/Ma Copolymer; Quateraium-15; Quatemium-15 Cis~Form; Quaternium-52; Ra-2397; Ra-30ll; Saccharin; Saccharin Sodium; Saccharin Sodium Anhydrous: Safilower Oil; Sd Alcohol 3a, Sd Alcohol 40, Sd Alcohol 40-2, Sd Alcohol 40b, Sepineo P 600; Serine; Sesame Oil; Shea Butter; Silastic Brand Medical Grade Tubing; Silastic Medical Adhesive,Silicone Type A; Silica, Dental; Silicon; Silicon Dioxide; Silicon Dioxide, Colloidal; Silicone; Silicone Adhesive 4102; Silicone Adhesive 4502; Silicone Adhesive Bio-Psa Q7-4201; Silicone Adhesive Bio-Psa Q7-4301; Silicone Emulsion; Sihcona/Polyester Film Strip; Simethicone; Simethicone Emulsion; Sipon Ls 20np; Soda Ash; Sodium Acetate; Sodium Acetate Anhydrous; Sodium Alkyl Sulfate: Sodium Ascorbate; Sodium Benzoate; Sodium Bicarbonate; Sodium Bisulfate; Sodium Bisulfite; Sodium Borate; Sodium Borate Decahydrate; Sodium Carbonate; Sodium Carbonate Decahydrate; Sodium Carbonate Monohydrate; Sodium Cetostearyl Sulfate; Sodium Chlorate; Sodium Chloride; Sodium Chionde Injection; Sodium Chloride Injection, Bacteriostatic; Sodium Cholesteryl Sulfate, Sodium Citrate; Sodium Cocoyl Sarcosinate; Sodium Desoxycholate; Sodium Dithionite; Sodium Dodecylbenzenesulfonate; Sodium Formaldehyde Sulfoxylate; Sodium Gluconate; Sodium Hydroxide; Sodium Hypochlorite; Sodium Iodide: Sodium Lactate; Sodium Lactate, L-; Sodium Laureth-2 Sulfate; Sodium Laureth-3 Sulfate; Sodium Laureth-5 Sulfate, Sodium Lauroyi Sarcosinate; Sodium Lauryi Sulfate; Sodium Lauryl Suifoacetate; Sodium Metabisulfite; Sodium Nitrate; Sodium Phosphate; Sodium Phosphate Dihydrate; Sodium Phosphate, Dibasic; Sodium Phosphate, Dibasic, Anhydrous; Sodium Phosphate, Dibasic, Dihydrate; Sodium Phosphate, Dibasic, Dodecahydrate; Sodium Phosphate, Dibasic, Heptahydrate; Sodium Phosphate, Monobasic; Sodium Phosphate, Monobasic, Anhydrous; Sodium Phosphate. Monobasic. Dihydrate; Sodium Phosphate, Monobasic, Monohydrate; Sodium Polyacrylate (2500000 Mw); Sodium Pyrophosphate; Sodium Pyrrolidone Carboxylate; Sodium Starch Giycolate; Sodium Succinate Hexahydrate; Sodium Sulfate; Sodium Sulfate Anhydrous: Sodium Sulfate Decahydrate; Sodium Sulfite: Sodium Sulfosuccinated

Undecycienic Monoalkylolamide; Sodium Tartrate, Sodium Thiogiycoiate; Sodium Thiomalate, Sodium Thiosuifata; Sodium Thiosulfate Anhydrous; Sodium Trirnetaphosphate, Sodium Xylenesulfonate; Somay 44; Sorbic Acid; Sorbitan; Sorbitan Isostearate; Sorbiian Monolaurate; Sorbitan Monooleate; Sorbitan Monopalmitate; Sorbitan Monostearate; Sorbitan Sesquioieate; Sorbitan Trioleate, Sorbitan Tristearaie; Sorbitol; Sorbitol Solution, Soybean Flour; Soybean Oil; Spearmint Oil; Spermaceti; Squaiane; Stabilized Oxychloro Complex; Stannous 2-Ethylliexanoate; Stannous Chloride; Stannous Chloride Anhydrous; Stannous Fluoride; Stannous Tartrate; Starch; Starch 1500, Pregelatimzed; Starch, Corn; Stearalkomum Chloride, Stearalkonium Hectorite/Propylene Carbonate; Stearamidoethyl Diethylarmne; Staareth-10, Steareth-100; Steareth-2; Steareth-20; Steareth-21; Steareth-40; Stearic Acid; Stearic Diethanoiamide; Stearoxytriniethylsilane; Steartrimonium Hydroiyzed Animal Collagen; Stearyl Alcohol; Sterile Water For Inhalation; Siyrene/Isoprena/Styrane Block Copolymer; Succimer; Succinic Acid; Sucralose; Sucrose; Sucrose Distearate; Sucrose Polyesters; Sulfacetamide Sodium; Suifobutylether .Beta.-Cyclodextrin; Sulfur Dioxide; Sulfuric Acid; Sulfurous Acid; Surfactol Qs; Tagatose, D-; Talc; Tall Oil; Tallow Glycendes; Tartaric Acid; Tartaric Acid, D1-; Tenox; Tenox-2, Tert-Butyl Alcohol; Tart-Butyl Hydroperoxide; Tert-Butylhydroquinone; Tetraki s(2-Methoxy isobutylisocy anide)Copper(l) Tetrafluoroborate; Tetrapropyl Orthosiiicate; Tetrofosmin; Theophylline; Thimerosal; Threonine; Thymol; Tin; Titanium Dioxide; Tocopherol; Tocophersolan; Total parenteral nutrition, lipid emulsion; Triacetm; Tncaprylin; Trichloromonofluoromethane; Trideceth-10; Triethanolarnme Lauryl Sulfate, Tnfluoroacetic Acid; Triglycerides, Medium Cham; Trihydroxy stearin; Trilaneth-4 Phosphate; Trilauretli-4 Phosphate, Trisodium Citrate Dihydrate; Trisodium Hedta; Triton 720; Triton X-200; Trolamine; Tromantadine; Trometliarnine (TRIS); Tryptophan; Tyloxapol; Tyrosine; Undecylenic Acid; Union 76 Amsco-Res 6038; Urea; Valine; Vegetable Oil; Vegetable Oil Glyceride, Hydrogenated; Vegetable Oil, Hydrogenated; Versetannde; Viscarin; Viscose/Cotton; Vitamin E, Wax, Emulsifying; Wecobee Fs; White Ceresin Wax; White Wax; Xanthan Gum, Zinc; Zinc Acetate; Zinc Carbonate; Zinc Chloride; and Zmc Oxide.

|00423] Pharmaceutical composition formulations of AAV particles disclosed herein may include cations or anions. In one embodiment, the formulations include metal cations such as, but not limited to, Zn2+, Ca2+, Cu2+, Mn2+, Mg+ and combinations thereof. As anon-limiting example, formulations may include polymers and complexes with a metal cation *(See e.g.,* U.S. Pat. Nos. 6,265,389 and 6,555,525, each of which is herein incorporated by reference in its entirety).

[00424] Formulations of the invention may also include one or more pharmaceutically acceptable salts. As used herein, "pharmaceutically accepiable salts" refers to derivatives of the disclosed compounds wherein the parent compound is modified by converting an existing acid or base moiety to its salt form (e.g., by reacting the free base group with a suitable organic acid). Examples of pharmaceutically acceptable salts include, but are not limited to, mineral or organic acid salts of basic residues such as amines; alkali or organic salts of acidic residues such as earboxyiie acids; and the like. Representative acid addition salts include acetate, acetic acid, adipate, alginate, ascorhate, aspartate, benzenesuifonate, benzene sulfonic acid, benzoate, bisu!fatc, borate, hutyrate, caraphorate, camphorsulfonate, citrate, cyclopentanepropionate, digluconate, dodecylsulfate, ethanesulfonate, fumarate, glucoheptonate, glycerophosphate, hemisulfate, heptonate, hexanoate, hydrobromide, hydrochloride, hydroiodide, 2-hydroxy-ethanesulfonate, laetobionate, lactate, laurate, lauryl sulfate, malate, maleate, malonate, methanes ulfonate, 2~naphtbalenesul†onate, nicou'nate, nitrate, oleate, oxalate, palmitate, pamoate, pectinate, persulfate, 3-phenylpropionate, phosphate, picrate, pivaiate, propionate, stearate, succinate, sulfate, tartrate, thiocyanate, toluenesulfonate, undecanoate, valerate salts, and the like. Representative alkali or alkaline earth metal salts include sodium, lithium, potassium, calcium, magnesium, and the like, as well as nontoxic ammonium, quaternary ammonium, and amine cations, including, but not limited to ammonium, tetraniethylammonium, tetraethylammonium, methylamine, dimethylamine, trimethylamine, tnethylamme, ethylamme, and the like. The pharmaceutically acceptable salts of the present disclosure include the conventional non-toxic salts of the parent compound formed, for example, from non-toxic inorganic or organic acids.

[00425] Solvates may be prepared by crystallization, recrystallization, or precipitation from a solution that includes organic solvents, water, or a mixture thereof Examples of suitable solvents are ethanol, water (for example, mono-, di-, and tri-hydrates), *N*-methyipyrrolidinone (NMP), dimethyl sulfoxide (BMSO), *N N* '-dimethylformamide (DMF), *N,N'*-dimethyl acetamide (DMAC), l,3-dimethyl-2-inidazolidinone (DMEIJ), I,3-dimethyl-3,4,5,6-tetrahydro-2-(lH)-pyrimidinone (DMPU), acetonitnie (ACN), propylene glycol, ethyl acetate, benzyl alcohol, 2-pyrrolidone, benzyl benzoate, and the like. When water is the solvent, the solvate is referred to as a "hydrate."

HI. ADMINISTRATION AND DOSING

Administration

[00426] The AAV particles of the present invention may be administered by any delivery route which results in a therapeutically effective outcome. These include, but are not limited to, enteral

(into the intestine), gastroenteral, epidural (into the dura mater), oral (by way of the mouth), transdermal, intracerebral (into the **cerebrum), inlracerebroventricular** (into the cerebral ventricles), epicutaneous (application onto the skin), intradermal, (into the skin itself), subcutaneous (under the skin), nasal administration (through the nose), intravenous (into a vein), intravenous **bolus,** intravenous drip, **intra-arteria!** (into an artery), **intramuscular** (into a muscle), intracardiac (into the heart), intraosseous infusion (into the bone marrow), intrathecal (into the spinal canal), intraparenchynial (into brain tissue), intraperitoneal, (infusion or injection into the peritoneum), intravesical infusion, mtravitreai, (through the eye), intracavemous injection (into a pathologic cavity) intracavitary (into the base **of the** penis), **intravagirsal** administration, intrauterine, **extra-amniotic** administration, transdermal (diffusion through the intact skin for systemic distribution), transniucosai (diffusion through a mucous membrane), transvaginal, insufflation (snorting), sublingual, **sublabial,** enema, eye drops (onto the **conjunctiva),** or **in ear** drops, auricular (in or by way of the ear), buccal (directed toward the cheek), conjunctival, cutaneous, dental (to a tooth or teeth), electro-osmosis, endoeervicai, endosinusial, endotracheal, extracorporeal, hemodialysis, infiltration, interstitial, intra-abdominal, intra-amniotic, intra-articular, **intrabiliary,** mttabronchial, mtrabursal, **intracartilagirsous (within** a cartilage), intracaudal (within the cauda equine), intracistemal (within the cisten a magna cerebellonieduiaris), intracomeal (within the cornea), dental intraeornai, mtracoronary (within the **coronary** arteries), **intracorporus** cavemosum **(within** the dilatable spaces of the corporus cavernosa **of** the penis), **intradiscal** (within a disc), intraductal (within a **duct of** a gland), intraduodenal (within the duodenum), intradural (within or beneath the dura), mtraepidernial (to the epidermis), intraesophagea! (to the esophagus), intragastric (within the stomach), **intragingival** (within the gingivae), **intraileal (within** the distal portion of the small intestine), mtralesional (within or introduced directly to a localized lesion), intraluminal (within a lumen of a tube), intralymphatic (within the lymph), intramedullary (within the marrow cavity of a bone), mtramenmgeai (withm the meninges), mtramyocardial (within the myocardium), intraocular (within the eye), mtraovarian (within the **ovary), intrapericardial** (within the pericardium), intrapleural (within the pleura), intraprostatic (within the prostate gland), infrapuinicuiary (within the lungs or its bronchi), intrasinal (within the nasal or periorbital sinuses), intraspinal (within the vertebral column), intrasynoviai (within the synovial cavity **of a joint),** intratendinous (within a tendon), mtratestieular (within the testicle), intrathecal (within the cerebrospinal fluid at any level of the cerebrospinal axis), intrathoracic (withm the thorax), intratubular (withm the tubules of an organ), intratumor (within a tumor), intratympanic (withm the aurus media), mtravascitlar (within a vessel or vessels), **intraventricular (within** a ventricle), **iontophoresis (by** means **of**

electric current where ions of soluble salts migrate into the tissues of the body), irrigation (to bathe or flush open wounds or body cavities), laryngeal (directly upon the larynx), nasogastric (through the nose and into the stomach), occlusive dressing technique (topical route administration which is then covered by a dressing which occludes the area), ophthalmic (to the external eye), oropharyngeal (directly to the mouth and pharynx), parenteral, percutaneous, periarticular, peridural, perineural, periodoniaL rectal, respiratory (within the respirator)' tract by inhaling orally or nasally for local or systemic effect), retrobulbar (behind the pons or behind the eyeball), soft tissue, subarachnoid, subconjunctival, submucosal, topical, transplacental (through or across the placenta), transtracheal (through the wall of the trachea), transtympanic (across or through the tympanic cavity), ureteral (to the ureter), urethral (to the urethra), vaginal, caudal block, diagnostic, nerve block, biliary perfusion, cardiac perfusion, photopheresis and spinal.

[ΘΘ427] In some embodiments, compositions may be administered in a way which allows them to cross the blood-brain barrier, vascular barrier, or other epithelial barrier. The AAV particles of the present invention may be administered in any suitable form, either as a liquid solution or suspension, as a solid form suitable for liquid solution or suspension in a liquid solution. The AAV particles may be formulated with any appropriate and pharmaceutically acceptable excipient.

[00428] In one embodiment, the AAV particles of the present invention may be delivered to a subject via a single route administration.

[ΘΘ429] In one embodiment, the AAV particles of the present invention may be delivered to a subject via a multi-site route of administration. A subject may be administered at 2, 3, 4, 5 or more than 5 sites.

[00430] In one embodiment, a subject may be administered the AAV particles of the present invention using a bolus infusion.

[00431] In one embodiment, a subject may be administered the AAV particles of the present invention using sustained delivery over a period of minutes, hours or days. The infusion rate may be changed depending on the subject, distribution, formulation or another delivery parameter.

[00432] In one embodiment, the AAV particles of the present invention may be delivered by intramuscular delivery route. (See, e.g., U. S. Pat. No. 6506379; the content of which is incorporated herein by reference in its entirety). Non-limiting examples of intramuscular administration include an intravenous injection or a subcutaneous injection.

[00433] In one embodiment, the AAV particles of the present invention may be delivered by oral administration. Non-limiting examples of oral administration include a digestive tract administration and a buccal administration.

[00434] In one embodiment, the AAV particles of the present invention may be delivered by intraocular delivery route. A non-limiting example of intraocular administration include an intravitreal injection.

[00435] In one embodiment, the AAV particles of the present invention may be delivered by intranasal delivery route. Non-limiting examples of intranasal delivery include administration of nasal drops or nasal sprays.

[00436] In some embodiments, the AAV panicles that may be administered to a subject by peripheral injections. Non-limiting examples of peripheral injections include intraperitoneal intramuscular, intravenous, conjunctival or joint injection. It was disclosed in the art that the peripheral administration of AAV vectors can be transported to the central nervous system, for example, to the motor neurons (e.g., U. S. Patent Publication Nos. 20100240739; and 20100130594; the content of each of which is incorporated herein by reference in their entirety).

[00437] In one embodiment, the AAV particles may be delivered by injection into the CSF pathway. Non-limiting examples of delivery to the CSF pathway include intrathecal and intracerebro ventricular administration.

[00438] In one embodiment, the AAV particles may be delivered by systemic delivery. As a non-limiting example, the systemic delivery may be by intravascular administration.

[00439] In one embodiment, the AAV particles of the present invention may be administered to a subject by intracranial delivery (See, e.g., U. S. Pat. No. 8,119,611; the content of which is incorporated herein by reference in its entirety).

[00440] In one embodiment, the AAV particles of the present invention may be administered to a subject by mtraparenchymal administration.

[00441] In one embodiment, the AAV particles of the present invention may be administered to a subject by intramuscular administration.

[00442] In one embodiment, the AAV panicles of the present invention are administered to a subject and transduce muscle of a subject. As a non-iimiting example, the AAV particles are administered by intramuscular administration.

[00443] In one embodiment, the AAV particles of the present invention may be administered to a subject by intravenous administration.

[00444] In one embodiment, the AAV particles of the present invention may be administered to a subject by subcutaneous administration.

[00445] In one embodiment, the AAV particles of the present invention may be administered to a subject by topical administration.

[00446] In one embodiment, the AAV particles may be delivered by direct injection into the brain. As a non-limiting example, the brain delivery may be by intrastriatal administration.

[00447] In one embodiment, the AAV particles may be delivered by more than one route of administration. As non-limiting examples of combination adniinistrations, AAV particles may be delivered by intrathecal and mtracerebro ventricular, or by intravenous and intraparenchymal admini strati on.

Parenteral and injectable administration

[00448] In some embodiments, pharmaceutical compositions, AAV particles of the present invention may be administered parenterally. Liquid dosage forms for oral and parenteral administration include, but are not limited to, pharmaceutically acceptable emulsions, microemulsioiis, solutions, suspensions, syrups, and/or elixirs. In addition to active ingredients, liquid dosage forms may comprise inert diluents commonly used in the art such as, for example, water or other solvents, solubiiizmg agents and emulsifiers such as ethyl alcohol, isopropyl alcohol, ethyl carbonate, ethyl acetate, benzyl alcohol, benzyl beiizoate, propylene glycol, 1,3-butylene glycol, dimethyiformamide, oils (in particular, cottonseed, groundnut, com, germ, olive, castor, and sesame oils), glycerol, tetrahydrofurfuryl alcohol, polyethylene glycols and fatty acid esters of sorbitan, and mixtures thereof. Besides inert diluents, oral compositions can include adjuvants such as wetting agents, emulsifying and suspending agents, sweetening, flavoring, and/or perfuming agents. In certain embodiments for parenteral administration, compositions are mixed with solubiiizmg agents such as CREMOPHOR ®, alcohols, oils, modified oils, glycols, poiysorbates, cyclodextrins, polymers, and/or combinations thereof. In other embodiments, surfactants are included such as hydroxypropylceliuiose.

[00449] Injectable preparations, for example, sterile injectable aqueous or oleaginous suspensions may be formulated according to the known art using suitable dispersing agents, wettmg agents, and/or suspending agents. Sterile injectable preparations may be sterile injectable solutions, suspensions, and/or emulsions in nontoxic parenterally acceptable diluents ami/or solvents, for example, as a solution in 1,3-butanediol. Among the acceptable vehicles and solvents that may be employed are water, Ringer's solution, U.S.P., and isotonic sodium chloride solution. Sterile, fixed oils are conventionally employed as a solvent or suspending medium. For this purpose, any bland fixed oil can be employed including synthetic mono- or digiycerides. Fatty acids such as oleic acid can be used in the preparation of injectabies.

[00450] Injectable formulations may be sterilized, for example, by filtration through a bacterial-retaining filter, and/or by incorporating sterilizing agents in the form of sterile solid

compositions which can be dissolved or dispersed in sterile water or other sterile injectable medium prior to use.

[ΘΘ451] In order to prolong the effect of active ingredients, it is often desirable to slow the absorption of active ingredients from subcutaneous or intramuscular injections. This may be accomplished by the use of liquid suspensions of crystalline or amorphous material with poor water solubility. The rate of absorption of active ingredients depends upon the rate of dissolution which, in turn, may depend upon crystal size and crystalline form. Alternatively, delayed absorption of a parenterally administered drug form is accomplished by dissolving or suspending the drug in an oil vehicle. Injectable depot forms are made by forming microencapsule matrices of the drug in biodegradable polymers such as polyiactide-polygiycoiide. Depending upon the ratio of drag to polymer and the nature of the particular polymer employed, the rate of drug release can be controlled. Examples of other biodegradable polymers include poly(orthoesters) and polyCanhydrides). Depot injectable formulations are prepared by entrapping the drug in liposomes or microemulsions which are compatible with body tissues.

Rectal and vaginal administration

[00452] In some embodiments, pharmaceutical compositions, AAV particles of the present invention may be administered reetally and/or vaginally. Compositions for rectal or vaginal administration are typically suppositories which can be prepared by mixing compositions with suitable non-irritating excipients such as cocoa butter, polyethylene glycol or a suppository wax which are solid at ambient temperature but liquid at body temperature and therefore melt in the rectum or vaginal cavity and release the active ingredient.

Oral administration

[00453] In some embodiments, pharmaceutical compositions, AAV particles of the present invention may be administered orally. Solid dosage forms for oral administrat on include capsules, tablets, pills, powders, and granules. In such solid dosage forms, an active ingredient is mixed with at least one inert, pharmaceutically acceptable excrpient such as sodium citrate or dicalcium phosphate and/or fillers or extenders *(e.g.* starches, lactose, sucrose, glucose, mannitol, and silicic acid), binders *{e.g.* earboxymethylceliulose, alginates, gelatin, polyviiiylpyrroiidiiione, sucrose, and acacia), humectants *(e.g.* glycerol), disintegrating agents *(e.g.* agar, calcium carbonate, potato or tapioca starch, alginic acid, certain silicates, and sodium carbonate), solution retarding agents *(e.g.* paraffin), absorption accelerators *(e.g.* quaternary ammonium compounds), wetting agents *(e.g.* cetyl alcohol and glycerol monostearate), absorbents *(e.g.* kaolin and bentonite clay), and lubricants *(e.g.* talc, calcium stearate, magnesium stearate, solid

polyethylene glycols, sodium iauryl sulfate), and mixtures thereof. in the case of capsules, tabiets and pills, the dosage form may comprise buffering agents.

Topical or transdermal administration

[00454] As described herein, pharmaceutical compositions, AAV particles of the present invention may be formulated for administration topically. The skin may be an ideal target site for deliver} as it is readily accessible. Three routes are commonly considered to deliver pharmaceutical compositions, AAV particles of the present invention to the skin: (i) topical application *(e.g.* for local/regional treatment and/or cosmetic applications); (ii) intradermal injection *(e.g.* for local/regional treatment and/or cosmetic applications), and (iii) systemic delivery *(e.g.* for treatment of dermatologie diseases that affect both cutaneous and extracutaneous regions). Pharmaceutical compositions, AAV particles of the present invention can be delivered to the skin by several different approaches known in the art.

[00455] In some embodiments, the invention provides for a variety of dressings (e.g., wound dressings) or bandages (e.g., adhesive bandages) for conveniently and/or effectively carrying out methods of the present invention. Typically dressing or bandages may comprise sufficient amounts of pharmaceutical compositions, AAV particles of the present invention described herein to allow users to perform multiple treatments.

[00456] Dosage forms for topical and/or transdermal administration may include ointments, pastes, creams, lotions, gels, powders, solutions, sprays, inhalants and/or patches. Generally, active ingredients are admixed under sterile conditions with pharmaceutically acceptable excipients and/or any needed preservatives and/or buffers. Additionally, the present invention contemplates the use of transdermal patches, which often have the added advantage of providing controlled delivery of pharmaceutical compositions, AAV particles of the present invention to the body. Such dosage forms may be prepared, for example, by dissolving and/or dispensing pharmaceutical compositions, AAV particles in the proper medium. Alternatively, or additionally, rates may be controlled by either providing rate controlling membranes and/or by dispersing pharmaceutical compositions, AAV particles in a polymer matrix and/or gel.

[00457] Formulations suitable for topical administration include, but are not limited to, liquid and/or semi liquid preparations such as liniments, lotions, oil in water and/or water in oil emulsions such as creams, ointments and/or pastes, and/or solutions and/or suspensions.

[00458] Topicaliy-administrable formulations may, for example, comprise from about 1% to about 10% (w/w) active ingredient, although the concentration of active ingredient may be as high as the solubility limit of the active ingredient in the solvent. Formulations for topical administration may further comprise one or more of the additional ingredients described herein.

Depot administration

[00459] As described herein, in some embodiments, pharmaceutical compositions, AAV particles of the present invention are formulated in depots for extended release. Generally, specific organs or tissues ("target tissues") are targeted for administration.

[00460] In some aspects of the invention, pharmaceutical compositions, AAV particles of the present invention are spatially retained within or proximal to target tissues. Provided are methods of providing pharmaceutical compositions, AAV particles, to target tissues of mammalian subjects by contacting target tissues (which comprise one or more target cells) with pharmaceutical compositions, AAV particles, under conditions such that they are substantially retained in target tissues, meaning that at least 10, 20, 30, 40, 50, 60, 70, 80, 85, 90, 95, 96, 97, 98, 99, 99.9, 99.99 or greater than 99.99% of the composition is retained in the target tissues. Advantageously, retention is determined by measuring the amount of pharmaceutical compositions, AAV particles, that enter one or more target cells. For example, at least 1%, 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99%, 99.9%, 99.99%> or greater than 99.99% of pharmaceutical compositions, AAV particles, administered to subjects are present mtracellularly at a period of time following administration. For example, intramuscular injection to mammalian subjects may be performed using aqueous compositions comprising pharmaceutical compositions, AAV particles of the present invention and one or more transfection reagents, and retention is determined by measuring the amount of pharmaceutical compositions, AAV particles, present in muscle cells.

[00461] Certain aspects of the invention are directed to methods of providing pharmaceutical compositions, AAV particles of the present invention to a target tissues of mammalian subjects, by contacting target tissues (comprising one or more target cells) with pharmaceutical compositions, AAV particles under conditions such that they are substantially retained in such target tissues. Pharmaceutical compositions, AAV particles comprise enough active ingredient such that the effect of interest is produced in at least one target ceil. In some embodiments, pharmaceutical compositions, AAV particles generally comprise one or more ceil penetration agents, although "naked" formulations (such as without ceil penetration agents or other agents) are also contemplated, with or without pharmaceutically acceptable carriers.

Pulmonary administration

[00462] In some embodiments, pharmaceutical compositions, AAV particles of the present invention may be prepared, packaged, and/or sold in formulations suitable for pulmonary administration. In some embodiments, such administration is via the buccal cavity. In some embodiments, formulations may comprise dry particles comprising active ingredients. In such

embodiments, dry particles may have a diameter in the range from about 0.5 nm to about 7 nm or from about 1 nm to about 6 nm. In some embodiments, formulations may be in the form of dry powders for administration using devices comprising dry powder reservoirs to which streams of propellant may be directed to disperse such powder. In some embodiments, self-propelling solvent/powder dispensing containers may be used. In such embodiments, active ingredients may be dissolved and/or suspended in low-boiling propeilant in sealed containers. Such powders may comprise particles wherein at least 98% of the particles by weight have diameters greater than 0.5 nm and at least 95% of the particles by number have diameters less than 7 nm. Alternatively, at least 95% of the particles by weight have a diameter greater than 1 nm and at least 90% of the particles by number have a diameter less than 6 nm. Dry powder compositions may include a solid fine powder diluent such as sugar and are conveniently provided in a unit dose form.

[ΘΘ463] Low boiling propellants generally include liquid propellents having a boiling point of below 65 °F at atmospheric pressure. Generally, propellants may constitute 50% to 99.9% (w/w) of the composition, and active ingredient may constitute 0.1% to 20% (w/w) of the composition. Propellants may further comprise additional ingredients such as liquid non-ionic and/or solid anionic surfactant and/or solid diluent (which may have particle sizes of the same order as particles comprising active ingredients).

[00464] Pharmaceutical compositions formulated for pulmonary delivery may provide active ingredients in the form of droplets of solution and/or suspension. Such formulations may he prepared, packaged, and/or sold as aqueous and/or dilute alcoholic solutions and/or suspensions, optionally sterile, comprising active ingredients, and may conveniently be administered using any nebuhzation and/or atomization device. Such formulations may further comprise one or more additional ingredients including, but not limited to, a flavoring agent such as saccharin sodium, a volatile oil, a buffering agent, a surface active agent, and/or a preservative such as methylliydroxybenzoate. Droplets provided by this route of administration may have an average diameter in the range from about 0.*1* nm to about 200 nm.

Intranasal, nasal and buccal administration

|00465] In some embodiments, pharmaceutical compositions, AAV particles of the present invention may be administered nasally and/or intranasal. In some embodiments, formulations described herein useful for pulmonary delivery may also be useful for intranasal delivery. In some embodiments, formulations for intranasal administration comprise a coarse powder comprising the active ingredient and having an average particle from about 0.2. μη to 500 μη. Such formulations are administered in the manner in which snuff is taken, *i.e.* by rapid inhalation through the nasal passage from a container of the powder held close to the nose.

[00466] Formulations suitable for nasal administration may, for example, comprise from about as little as 0.1% (w/w) and as much as 100% (w/w) of active ingredient and may comprise one or more of the additional ingredients described herein. A pharmaceutical composition may be prepared, packaged, and/or sold in a formulation suitable for buccal administration. Such formulations may, for example, be in the form of tablets **and/or** lozenges made using conventional methods, and may, for example, 0.1% to 20% (w/w) active ingredient, the balance comprising an orally dissolvable and/or degradabie composition and, optionally, one or more of the additional ingredients described herein. Alternately, formulations suitable for buccal administration may comprise powders and/or an aerosolized and/or atomized solutions and/or suspensions comprising active ingredients. Such powdered, aerosolized, and/or aerosolized formulations, when dispersed, may comprise average particle and/or droplet sizes in the range of from about 0.1 nm to about 200 nm, and may further comprise one or more of any additional ingredients described herein.

Ophthalmic or otic administration

[00467] In some embodiments, pharmaceutical compositions, AAV particles of the present invention may be prepared, packaged, and/or sold in formulations suitable for ophthalmic and/or otic administration. Such formulations may, for example, be in the form of eye and/or ear drops including, for example, a 0.1/1.0% (w/w) solution and/or suspension of the active ingredient in aqueous and/or oily liquid excipients. Such drops may further comprise buffering agents, salts, and-'or one or more other of any additional ingredients described herein. Other opbthalmieally-administrable formulations which are useful include those which comprise active ingredients in microcrysialline form and-'or in liposomal preparations. Subretinal inserts may also be used as forms of administration.

Delivery to Cells

[ΘΘ468] The present disclosure provides a method of delivering to a cell or tissue any of the above-described AAV particles, comprising contacting the cell or tissue with said AAV particle or contacting the cell or tissue with a formulation comprising said AAV particle, or contacting the cell or tissue with any of the described compositions, including pharmaceutical compositions. The method of delivering the AAV particle to a cell or tissue can be accomplished in *vitro, ex vivo,* or *in vivo.*

Delivery to Subjects

[ΘΘ469] The present disclosure additionally provides a method of delivering to a subject, including a mammalian subject, any of the above-described AAV particles comprising administering to the subject said AAV particle, or administering to the subject a formulation

comprising said AAV panicle, or administering to the subject any of the described compositions, including pharmaceutical compositions.

Dose and Regimen

[00470] The present invention provides methods of administering AAV particles in accordance with the invention to a subject in need thereof. The pharmaceutical, diagnostic, or prophylactic AAV particles and compositions of the present invention may be administered to a subject using any amount and any route of administration effective for preventing, treating, managing, or diagnosing diseases, disorders and/or conditions. The exact amount required will vary from subject to subject, depending on the species, age, and general condition of the subject, the severity of the disease, the particular composition, its mode of administration, its mode of activity, and the like. The subject may be a human, a mammal, or an animal. Compositions in accordance with the invention are typically formulated in unit dosage form for ease of administration and uniformity of dosage. It will be understood, however, that the total daily usage of the compositions of the present invention may be decided by the attending physician withm the scope of sound medical judgment. The specific therapeutically effective, prophylactic-ally effective, or appropriate diagnostic dose level for any particular individual will depend upon a variety of factors including the disorder being treated and the severity of the disorder; the activity- of the specific payioad employed; the specific composition employed; the age, body weight, general health, sex and diet of the patient; the time of administration, route of administration, and rate of excretion of the specific AAV particle employed; the duration of the treatment; drugs used in combination or coincidental with the specific AAV particle employed; and like factors well known in the medical arts.

[00471] In certain embodiments, AAV particle pharmaceutical compositions in accordance with the present invention may be administered at dosage levels sufficient to deliver from about 0.0001 mg/kg to about 100 mg/kg, from about 0.001 mg/kg to about 0.05 mg/kg, from about 0.005 mg/kg to about 0.05 mg/kg, from about 0.001 mg/kg to about 0.005 mg/kg, from about 0.05 mg/kg to about 0.5 mg/kg, from about 0.01 rag/kg to about 50 mg/kg, from about 0.1 mg/kg to about 40 mg/kg, from about 0.5 mg/kg to about 30 mg/kg, from about 0.01 mg/kg to about 10 mg/kg, from about 0.1 mg/kg to about 10 mg/kg, or from about 1 mg/kg to about 25 mg/kg, of subject body weight per day, one or more times a day, to obtain the desired therapeutic, diagnostic, or prophylactic, effect. It will be understood that the above dosing concentrations may be converted to vg or viral genomes per kg or into total viral genomes administered by one of skill in the art.

[00472] In certain embodiments, AAV particle pharmaceutical compositions in accordance with the present disclosure may be administered at about 10 to about 600 µl/site, 50 to about 500 µl/sue, 100 to about 400 µl/sue, 120 to about 300 µl/site, 140 to about 200 µl/site, about 160 µl/site. As non-iimiting examples, AAV particles may be administered at 50 µl/site and/or 150 µl/site.

[00473] The desired dosage of the AAV particles of the present invention may be delivered only once, three times a day, two times a day, once a day, every other day, every third day, every week, every two weeks, every three weeks, or every four weeks. In certain embodiments, the desired dosage may be delivered using multiple administrations *(e.g.,* two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, fourteen, or more administrations). When multiple administrations are employed, split dosing regimens such as those described herein may be used. As used herein, a "spirt dose" is the division of "single unit dose" or total daily dose into two or more doses, *e.g.* , two or more administrations of the "single unit dose". As used herein, a "single unit dose" is a dose of any therapeutic administered in one dose/at one time/single route/single point of contact, i.e., single administration event.

[00474] The desired dosage of the AAV particles of the present invention may be administered as a "pulse dose" or as a "continuous flow". As used herein, a "pulse dose" is a series of single unit doses of any therapeutic administered with a set frequency over a period of time. As used herein, a "continuous flow" is a dose of therapeutic administered continuously for a period of time in a single route/single point of contact, i.e., continuous administration event. A total daily dose, an amount given or prescribed in 24 hour period, may be administered by any of these methods, or as a combination of these methods, or by any other methods suitable for a phannaceuti cal administration.

[00475] In one embodiment, delivery of the AAV particles of the present invention to a subject provides neutralizing activity to a subject. The neutralizing activity can be for at least 1 month, 2 months, 3 months, 4 months, 5 months, 6 months, 7 months, 8 months, 9 months, 10 months, 11 months, 1 year, 13 months, 14 months, 15 months, 16 months, 17 months, 18 months, 19 months, 20 months, 20 months, 21 months, 22 months, 23 months, 2 years, 3 years, 4 years, 5 years, 6 years, 7 years, 8 years, 9 years, 10 years or more than 10 years.

[00476] In one embodiment, delivery of the AAV particles of the present invention results in minimal serious adverse events (SAEs) as a result of the delivery of the AAV particles.

[00477] In one embodiment, delivery of AAV particles to cells of the central nervous system (e.g., parenchyma) may comprise a total dose between about $1x10^6$ VG and about $1\chi 10^{16}$ VG. In some embodiments, delivery may comprise a total dose of about $1x10^6$, $2x10^6$, $3x10^5$, $4x10^6$,

5xl 0^{6}, 6xl0^{6}, 7x10^{6}, 8x10^{6}, 9xl0^{6}, lxlO^{7}, 2xl 0^{7}, 3xl0^{7}, 4x10^{7}, 5xl0^{7}, 6x10^{7}, **7x10^{7},** 8xl0^{7}, 9x10^{7}, 1x10^{8}, 2x10^{8}, 3x10^{8}, 4x10^{8}, 5x10^{8}, 6x10^{8}, 7x10^{8}, 8xl 0^{8}, 9x10^{8}, 1x10^{9}, 2x10^{9}, **3x10^{9},** 4xI 0^{9}, 5xl.0^{9}, 6x10^{9}, 7x10^{9}, 8x1.0^{9}, 9x10^{9}, **IxlO^{10},** 1.9x.lO^{10}, 2x10^{10}, 3x10^{10}, 3.73x.l 0^{10}, 4xi 0^{10}, 5xi 0^{10}, 6xl0^{10}, 7xl 0^{10}, 8xl 0^{10}, 9xl 0^{10}, lxlO^{11}, 2xl 0^{11}, 2.5xl 0^{11}, 3xlO^{11}, 4x10", 5xlO^{11}, 6xlO^{11}, 7xl O^{11}, 8xl 0^{11}, 9x10", lxlO^{12}, 2xl 0^{12}, 3x10^{12}, 4x10^{12}, 5xl 0^{12}, 6xl 0^{12}, 7xl 0^{12}, 8xl 0^{12}, 9xl 0^{12}, lxlO^{13}, **2x10",** 3x1.0^{13}, **4x10", 5x10",** 6x10^{13}, 7x10^{13}, 8x1.0^{13}, 9x10^{13}, IxlO^{34}, 2xl 0^{14}, 3x10^{34}, 4x10^{14}, 5xlO^{14}, 6x10^{14}, 7x10^{14}, 8xl0^{14}, 9x10^{s4}, lxlO^{15}, 2xl 0^{15}, 3xl0^{15}, 4x10^{15}, **5x10",** 6xl0^{15}, 7x10^{15}, 8xlG^{15}, 9x10^{35}, or IxlO^{36} VG. As a non-limiting example, the total dose is lxlO^{13} VG. As another non-limiting example, the total dose is 2.lxlO^{12} VG.

[ΘΘ478] In one embodiment, deliver}' of AAV particles to cells of the central nervous system (e.g., parenchyma) may comprise a composition concentration between about 1x10^{6} VG/mL and about lxlO^{16} VG/mL. In some embodiments, delivery may comprise a composition concentration of about IxlO^{6}, 2x10^{6}, 3xi 0^{6}, 4xi 0^{6}, 5x10^{6}, 6xl 0^{6}, 7xI 0^{6}, 8x10^{6}, 9xl 0^{6}, IxlO^{7}, 2xl 0^{7}, 3xl0^{7}, 4x10^{7}, 5xl 0^{7}, 6xl0^{7}, 7xl0^{7}, 8xl 0^{7}, 9xl0^{7}, lx10^{s}, 2x10^{8}, 3x10^{8}, 4x10^{s}, 5x10^{8}, 6x10^{s}, 7x10^{s}, 8x10^{8}, 9x10^{s}, lxlO^{9}, 2xl 0^{9}, 3xl 0^{9}, 4xl0^{9}, 5xl 0^{9}, 6xl0^{9}, 7x10^{9}, 8x10^{9}, 9x10^{9}, lxlO^{10}, 2xl0^{10}, 3χ10^{10}, 4xi 0^{10}, 5xI 0^{10}, 6xl 0^{10}, 7xl 0^{10}, 8xl 0^{10}, 9xI 0^{10}, lxlO^{11}, 2x10^{11}, 3xi 0^{11}, **4x10",** 5x1.0^{11}, 6x10^{11}, 7x1.0^{11}, 8x10^{3i}, 9xl0^{33}, 1x1.0^{12}, 2x10^{32}, **3x10",** 4xl 0^{12}, 5x10^{12}, 6x1.0^{12}, 7x10", 8xl0^{12}, 9xl 0^{12}, lxlO^{13}, 2x10°, 3xl 0^{13}, 4xl 0^{13}, 5xl 0^{13}, 6xi 0^{13}, 7x10", 8xl 0^{13}, 9xl 0^{13}, lxlO^{14}, 2xl()^{34}, 3xl ()^{14}, 4x10^{34}, 5xl 0^{14}, 6xl 0^{14}, 7x10^{34}, 8x10^{14}, 9x10^{14}, **1x10",** 2xl ()^{15}, 3x10^{35}, 4x10^{15}, 5x10", 6x10^{15}, 7xl 0^{15}, 8x10", 9x10^{15}, or lxlO^{16} VG/mL. In one embodiment, the delivery comprises a composition concentration of 1x10" VG/mL. In one embodiment, the delivery comprises a composition concentration of 2.lxlO^{12} VG/mL.

Combinations

[ΘΘ479] The AAV particles may be used in combination with one or more other therapeutic, prophylactic, research or diagnostic agents. By "in combination with," it is not intended to imply that the agents must be administered at the same time and/or formulated for delivery together, although these methods of delivery are within the scope of the present invention. Compositions can be administered concurrently with, prior to, or subsequent to, one or more other desired therapeutics or medical procedures. In general, each agent will be administered at a dose and/or on a time schedule determined for that agent. In some embodiments, the present disclosure encompasses the delivery of pharmaceutical, prophylactic, research, or diagnosti c compositions in combination with agents that may improve their bioavailability, reduce and/or modify their metabolism, inhibit their excretion, and/or modify their distribution within the body.

Measurement of Expression

[00480] Expression of payloads from viral genomes may be determined using various methods known in the art such as, but not limited to immimochemistry (e.g., IHC), ***in situ*** hybridization (TSH), enzyme-linked immunosorbent assay (EL1SA), affinity ELISA, ELISPOT, flow cytometry, immunocytoiogy, surface plasmon resonance analysis, kinetic exclusion assay, liquid chromatography-mass spectrometry (LCMS), high-performance liquid chromatography (I-IPLC), BCA assay, immtmoelectrophoresis. Western blot, SDS-PAGE, protein immunoprecipitation, and/or PCR.

Bioavailability

[00024] The AAV particles, when formulated into a composition with a delivery agent as described herein, can exhibit an increase in bioavailability as compared to a composition lacking a delivery agent as described herein. As used herein, the term "bioavailability' " refers to the systemic availability of a given amount of AAV particle or expressed payload administered to a mammal. Bioavailability can be assessed by measuring the area under the curve (AUC) or the maximum serum or plasma concentration (Cmax) of the composition following. AUC is a determination of the area under the curve plotting the serum or plasma concentration of a compound (e.g., AAV particles or expressed payloads) along the ordinate (Y-axis) against time along the abscissa (X-axis). Generally, the AUC for a particular compound can be calculated using methods known to those of ordinary skill in the art and as described in G. S. Banker, Modern Pharmaceutics, Drugs and the Pharmaceutical Sciences, v. 72, Marcel Dekker, New York, Inc., 1996, the contents of which are herein incorporated by reference in its entirety.

|00025] The Cmax value is the maximum concentration of the AAV particle or expressed payload achieved in the serum or plasma of a mammal following administration of the AAV particle to the mammal. The Cmax value of can be measured using methods known to those of ordinary skill in the art. The phrases "increasing bioavailability" or "improving the pharmacokinetics,' " as used herein mean that the systemic availability of a first AAV particle or expressed payload, measured as AUC, Cmax, or Cmin in a mammal is greater, when co-administered with a delivery' agent as described herein, than when such co-administration does not take place. In some embodiments, the bioavailability can increase by at least about 2%, at least about 5%, at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 35%, at least about 40%, at least about 45%, at least about 50%, at least about 55%, at least about 60%, at least about 65%, at least about 70%, at least about 75%, at least about 80%, at least about 85%, at least about 90%, at least about 95%, or about 100%.

Therapeutic Window

[00026] As used herein "therapeutic window" refers to the range of plasma concentrations, or the range of levels of therapeutically active substance at the site of action, with a high probability of eliciting a therapeutic effect. In some embodiments, the therapeutic window of the AAV particle as described herein can increase by at least about 2%, at least about 5%, at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 35%, at least about 40%, at least about 45%, at least about 50%, at least about 55%, at least about 60%, at least about 65%, at least about 70%, at least about 75%, at least about 80%, at least about 85%, at least about 90%, at least about 95%, or about 100%.

Volume of Distribution

[ΘΘ027] As used herein, the term "volume of distribution" refers to the fluid volume that would be required to contain the total amount of the drug in the body at the same concentration as in the blood or plasma: V_{dist} equals the amount of drug in the body/concentration of drug in blood or plasma. For example, for a 10 mg dose and a plasma concentration of 10 mg/L, the volume of distribution would be 1 liter. The volume of distribution reflects the extent to which the drug is present in the extravascular tissue. A large volume of distribution reflects the tendency of a compound to bind to the tissue components compared with plasma protein binding. In a clinical setting, V_{dist} can be used to determine a loading dose to achieve a steady state concentration. In some embodiments, the volume of distribution of the AAV particles as described herein can decrease at least about 2%, at least about 5%, at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 35%, at least about 40%, at least about 45%, at least about 50%, at least about 55%, at least about 60%, at least about 65%, at least about 70%.

Biological Effect

[ΘΘ028] In one embodiment, the biological effect of the AAV particles delivered to the animals may be categorized by analyzing the payload expression in the animals. The payload expression may be determined from analyzing a biological sample collected from a mammal administered the AAV particles of the present invention. For example, a protein expression of 50-200 pg/mi for the protein encoded by the AAV particles delivered to the mammal may be seen as a therapeutically effective amount of protein in the mammal.

IV. METHODS AND USES OF THE COMPOSITIONS OF THE INVENTION

[ΘΘ481] The present disclosure provides a method for treating a disease, disorder and/or condition in a mammalian subject, including a human subject, comprising administering to the subject any of the AAV particles described herein or administering to the subject any of the described compositions, including pharmaceutical compositions, described herein.

[00482] In one embodiment, the AAV panicles of the present invention are administered to a subject prophylacticaily.

[ΘΘ483] In one embodiment, the AAV particles of the present invention are administered to a subject having at least one of the diseases described herein.

[00484] In one embodiment, the AAV particles of the present invention are administered to a subject to treat a disease or disorder described herein. The subject may have the disease or disorder or may be at-risk to developing the disease or disorder.

[00485] In one embodiment, the AAV particles of the present invention are part of an active immunization strategy to protect against diseases and disorders. In an active immunization strategy, a vaccine or AAV particles are administered to a subject to prevent an infectious disease by activating the subject's production of antibodies that can fight off invading bacteria or viruses.

[00486] In one embodiment, the AAV particles of the present invention are part of a passive immunization strategy. In a passive immunization strategy, antibodies against a particular infectious agent are given directly to the subject.

Diseases and toxins

[ΘΘ487] Various infectious diseases may be treated with pharmaceutical compositions. AAV particles, of the present invention. As used herein, the term "infectious disease" refers to any disorders caused by organisms such as bacteria, viruses, fungi or parasites. As a non-limiting example, the infectious disease may he Acute bacterial rhinosmusitis, 14-day measles. Acne, Acrodermatitis chronica atrophicans (ACA)-(late skin manifestation of latent Lyme disease), Acute hemorrhagic conjunctivitis, Acute hemorrhagic cystitis. Acute rhmosinusitis. Adult T-cell Leukemia-Lymphoma (ATLL), African Sleeping Sickness, AIDS (Acquired Immunodeficiency Syndrome), Alveolar hydatid. Amebiasis, Amebic meningoencephalitis, Anaplasmosis, Anthrax, Arboviral or parainfectious, Ascariasis -(Roundworm infections), Aseptic meningitis. Athlete's foot (Tinea pedis), Australian tick typhus, Avian Influenza, Babesiosis, Bacillary angiomatosis, Bacterial meningitis. Bacterial vaginosis. Balanitis, Balantidiasis, Bang's disease, Barmah Forest vims infection, Bartoneilosis (Verruga peruana; Carrion's disease; Oroya fever), Bat Lyssavirus infection, Bay sore (Chiclero's ulcer), Bayiisascaris infection (Racoon roundworm infection). Beaver fever, Beef tapeworm, Bejel (endemic syphilis), Biphasic meningoencephalitis, Black Bane, Black death , Black piedra, Blackwater Fever, Blastomycosis, Blennorrhea of the newborn, Blepharitis, Boils, Bomholm disease (pleurodynia), Borreha miyamotoi Disease, Botulism, Boutonneuse fever, Brazilian purpuric fever, Break Bone fever. Brill, Bronchiolitis, Bronchitis, Brucellosis (Bang's disease), Bubonic plague, Bullous impetigo, Burkholderia mallei

(Glanders), Burkholderia pseudomaliei (Melioidosis), Buruli ulcers (also Mycoburuli ulcers), Busse, Busse-Buscbkc disease (Cryptococcosis), California group encephalitis. Campylobacteriosis, Candidiasis, Canefield fever (Canicola fever; 7-day fever; Weil's disease; leptospirosis: canefield fever), Camcoia fever, Capillanasis, Carate, Carbapenem-resistant Enterobacteriaceae (CRE), Carbuncle, Carrion's disease. Cat Scratch fever, Cave disease. Central Asian hemorrhagic fever. Central European tick, Cervical cancer, Chagas disease. Chancroid (Soft chancre), Chicago disease, Chickenpox (Varicella), Chiclero's ulcer, Chikungunya fever, Chlamydial infection. Cholera, Chrornobiastomycosts. Ciguatera, Clap, Cionorchiasis (Liver fluke infection), Clostridium Difficile Infection, ClostriDiura Pcrfringcns (Epsilon Toxin), Coccidioidomycosis fungal infection (Valley fever; desert rheumatism), Coenurosis, Colorado tick fever, Condyloma accuminata. Condyloma accuminata(Warts), Condyloma lata, Congo fever, Congo hemorrhagic fever virus. Conjunctivitis , cowpox. Crabs, Crimean, Croup, Cryptococcosis, Cryptosporidiosis (Crypto), Cutaneous Larval Migrans, Cyclosporiasis, Cystic hydatid, Cysticercosis, Cystitis, Czechoslovak tick, D68 (EV-D68), Dacryocytitis, Dandy fever. Darling's Disease, Deer fly fever. Dengue fever (1, 2, 3 and 4), Desert rheumatism, Devil's grip. Diphasic milk fever, Diphtheria, Disseminated Intravascular Coagulation, Dog tapeworm, Donovanosis, Donovanosis (Granuloma inguinale), Dracontiasis. Dracunculosis, Duke's disease. Dum Dum Disease, Durand-Nicholas-Favre disease, Dwarf tapeworm, E. Coli infection (E.Coii), Eastern equine encephalitis, Ebola Hemorrhagic Fever (Ebola virus disease EVD), Ectothrix, Ehrlichiosis (Sennetsu fever), Encephalitis, Endemic Relapsing fever. Endemic syphilis, Endophthalmitis, Endothrix, Enterobiasis (Pinworm infection), Enterotoxin - B Poisoning (Staph Food Poisoning), Enterovirus Infection, Epidemic Keratoconjunctivitis, Epidemic Relapsing fever, Epidemic typhus, Epiglottitis, Erysipelis, Erysipeloid (Erysipelothricosis), Erythema chronicum migrans, Erythema infectiosum. Erythema marginatum, Erythema multiforme. Erythema nodosum. Erythema nodosum leprosum, Erythrasma, Espundia, Eumycotic mycetoma, European blastomycosis, Exanthem subitum (Sixth disease), Eyeworm, Far Eastern tick, Faseioliasis. Fievre bouionneuse(Tick typhus), Fifth Disease (erythema infectiosum), Filatow-Dukes' Disease (Scalded Skin Syndrome; Hitter's Disease), Fish tapeworm, Fitz-Hugh-Curtis syndrome - Perihepatitis, Flinders island Spotted Fever, Flu (Influenza), Folliculitis, Four Corners Disease, Four Corners Disease (Human Pulmonary Syndrome (HPS)), Frambesia, Francis disease, Furunculosis, Gas gangrene, Gastroenteritis, Genital Herpes, Genital Warts. German measles, Gersmiann-Straussier-Schemker (GSS), Giardiasis, Gilchrist's disease, Gingivitis, Gingivostomatitis, Glanders, Glandular fever (infectious mononucleosis), Gnathostomiasis, Gonococcal Infection (Gonorrhea), Gonorrhea, Granuloma inguinale

(Donovanosis), Guinea Worm, Haemophilus Influenza disease. Hamburger disease, Hansen's disease - leprosy, Hantaan disease, Hantaan-Korean hemorrhagic fever, Hantavirus Pulmonary Syndrome , Hantavirus Pulmonary- Syndrome (HPS). Hard chancre, Hard measles, Haverhill fever - Rat bite fever, Head and Body Lice, Heartland fever, Helicobacterosis, Hemolytic Uremic Syndrome (HUS), Hepatitis A, Hepatitis B, Hepatitis C, Hepatitis D, Hepatitis E, Herpangina, Herpes- genital, Herpes labialis. Herpes- neonatal, Hydradenitis, Histoplasmosis, Histoplasmosis infection (Histoplasmosis), His-Werner disease, HIV infection, Hookworm infections, Hordeola, Hordeola (Stye), HTLV, HTLV- associated myelopathy (HAM), Human granulocytic ehrlichiosis. Human monocytic ehrlichiosis, Human Papillomavirus (HPV), Human Pulmonary- Syndrome , Hydatid cyst , Hydrophobia. Impetigo, including congenital (German Measles), Inclusion conjunctivitis, Inclusion conjunctivitis - Swimming Pool conjunctivitis- Pannus, Infantile diarrhea, Infectious Mononucleosis, Infectious myocarditis, Infectious pericarditis. Influenza, Isosporiasis, Israeli spotted fever, Japanese Encephalitis, Jock itch, Jorge Lobo disease - lobomycosis. Jungle yellow fever, Junin Argentinian hemorrhagic fever, Kala Azar, Kaposi's sarcoma, Keloidal blastomycosis, Keratoconjunctivitis , Kuru, Kyasanur forest disease, LaCrosse encephalitis, Lassa hemorrhagic fever, Legioneilosis (Legionnaires Disease), Legionnaire's pneumonia, Lemierre's Syndrome (Postanginal septicemia), Lemming fever, Leprosy , Leptospirosis (Nanukayami fever; Weil's disease), Listeriosis (Listeria), Liver fluke infection, Lobo's mycosis. Lockjaw, Loiasis, Louping Ill, Ludwig's angina, Lung fluke infection. Lung fluke infection (Paragonimiasis), Lyme disease, Lymphogranuloma venereum infection (LGV), Machupo Bolivian hemorrhagic fever, Madura foot, Mai del pinto, Malaria, Malignant pustule, Malta fever, Marburg hemorrhagic fever, Masters disease, Maternal Sepsis (Piierperal fever), Measles, Mediterannean spotted fever, Melioidosis (Whitraore's disease), Meningitis, Meningococcal Disease. M.E.RS, Milker's nodule, Mo!Iuscum contagiosum, Moniliasis, monkeypox, Mononucleosis, Mononucleosis-iike syndrome, Montezuma's Revenge, Morbilli, MRSA (methici!lin-resistant Staphylococcus aureus) infection. Mucormycosis- Zygomycosis, Multiple Organ Dysfunction Syndrome or MODS, Multiple-system atrophy (MSA), Mumps, Murine typhus, Murray Valley Encephalitis(MVE), Myeoburuli ulcers, Mycoburuli uicers- Buruli ulcers, Mycotic vulvovaginitis, Myositis, Nanukayami fever, Necrotizing fasciitis, Necrotizing fasciitis- Type 1, Necrotizing fasciitis- Type 2, Negishi, New world spotted fever, Nocardiosis, Nongonococcal urethritis, Non-Polio (Non-Polio Enterovirus). Norovirus infection, North American blastomycosis, North Asian tick typhus, Norwaik virus infection, Norwegian itch, O'Hara disease, Omsk hemorrhagic fever, Onchoeeriasis, Onychomycosis, Opisthorchiasis, Opthalmia neonatorium, Oral hairy leukoplakia, Orf, Oriental Sore, Oriental Spotted Fever,

Ornithosis (Parrot fever; Psittacosis), Oroya fever, Otitis externa, Otitis media, Pannus, Paracoccidioidomycosis, Paragonimiasis, Paralytic Shellfish Poisoning (Paralytic Shellfish Poisoning), Paronychia (Whitlow). Parotitis, PCP pneumonia, Pediculosis, Peliosis hepatica, Pelvic inflammatory Disease , Pertussis (also called Whooping cough), Phaeohyphomycosis, Pharyngoconjunctival fever, Piedra (White Piedra), PiedraiBlack Piedra), Pigbel, Pink eye conjunctivitis , Pinta, Pinworm infection, Pitted Keratolysis, Pityriasis versicolor (Tinea versicolor), Plague: Bubonic, Pleurodynia, Pneumococcal Disease, Pneumocystosis, Pneumonia, Pneumonic (Plague), Polio or Poliomyelitis, Polycystic hydatid, Pontiac fever. Pork tapeworm, Posada-Wernicke disease, Postanginal septicemia, Powassan, Progressive multifocal leukencepbaSopathy, Progressive Rubella Panencephalitis, Prostatitis, Pseudomembranous colitis, Psittacosis, Puerperal fever, Pustular Rash diseases (Small pox). Pyelonephritis, Pylephlebitis, Q-Fever, Quinsy, Quintana fever (5-day fever), Rabbit fever, Rabies, Racoon roundworm infection, Rat bite fever, Rat tapeworm, Reiter Syndrome, Relapsing fever, Respiratory syncytial virus (RSV) infection, Rheumatic fever, Rhodotorulosis, Ricin Poisoning, Rickettsialpox, Rickettsiosis , Rift Valley Fever, Ringworm, Ritter's Disease, River Blindness, Rocky Mountain spotted fever, Rose Handler's disease (Sporotrichosis), Rose rash of infants, Roseola, Ross River fever, Rotavirus infection. Roundworm infections. Rubella. Rubeola, Russian spring, Salmonellosis gastroenteritis, San Joaquin Valley fever, Sao Paulo Encephalitis, Sao Paulo fever, SARS, Scabies Infestation (Scabies) (Norwegian itch). Scalded Skin Syndrome, Scarlet fever (Scarlatina), Schistosomiasis, Scombroid, Scrub typhus, Sermetsu fever. Sepsis (Septic shock), Severe Acute Respiratory Syndrome, Severe Acute Respirator} Syndrome (SARS), Shiga Toxigenic Escherichia coli (STEC/VTEC), Shigellosis gastroenteritis (Shigella), Shinbone fever, Shingles , Shipping fever, Siberian tick typhus, Sinusitis, Sixth disease, Slapped cheek disease , Sleeping sickness. Smallpox (Variola), Snail Fever, Soft chancre. Southern tick associated rash illness, Sparganosis, Spelunker's disease. Sporadic typhus. Sporotrichosis, Spotted fever. Spring, St. Louis encephalitis, Staphylococcal Food Poisoning, Staphylococcal infection, Strep. throat, Streptococcal Disease, Streptococcal Toxic-Shock Syndrome, Strongyloiciasis, Stye, Subacute Sclerosing Panencephalitis, Subacute Sclerosing Panencephalitis (SSPE), Sudden Acute Respiratory Syndrome, Sudden Rash, Swimmer's ear. Swimmer's Itch, Swimming Pool conjunctivitis, Sylvatic yellow fever, Syphilis, Systemic inflammatory Response Syndrome (SIRS), Tabes dorsalis (tertiary syphilis), Taeniasis. Taiga encephalitis, Tanner's disease, Tapeworm infections, Temporal lobe encephalitis, Temporal lobe encephalitis, tetani (Lock jaw), Tetanus Infection, Threadworm infections. Thrush, Tick, Tick typhus, Tinea barbae. Tinea capitis. Tinea corporis. Tinea cruris, Tinea manuum, Tinea nigra,

Tinea pedis, Tinea unguium. Tinea versicolor, Torulopsosis, Torulosis, Toxic Shock Syndrome, Toxoplasmosis, transmissible spongioform (CJD), Traveler's diarrhea. Trench fever 5, TrichineUosis, Trichomoniasis, Trichomycosis axillaris, Trichuriasis, Tropical Spastic Paraparesis (TSP), Trypanosomiasis, Tuberculosis (TB), Tuberculousis, Tularemia, Typhoid Fever, Typhus fever. Ulcus mo‖e, Undulant fever. Urban yellow fever, Urethritis, Vaginitis, Vaginosis. Vancomycin Intermediate (VISA), Vancomycin Resistant (VRSA), Varicella. Venezuelan Equine encephalitis, Verruga pemana, Vibrio choierae (Cholera), Vibriosis (Vibrio), Vincent's disease or Trench mouth, Viral conjunctivitis , Viral Meningitis, Viral meningoencephalitis, Viral rash. Visceral Larval Migrans, Vomito negro, Vulvovaginitis, Warts. Waterhouse, Weil's disease, West Nile Fever, Western equine encephalitis. Whipple's disease. Whipworm infection, White Piedra, Whitlow, Whitmore's disease, Winter diarrhea, Woihynia fever, Wool sorters' disease, Yaws, Yellow Fever, Yersinosis, Yersinosis (Yersinia), Zahorsky's disease, Zika virus disease. Zoster, Zygoraycosis, John Cunningham Virus (JCV), Human immunodeficiency virus (HIV), influenza virus. Hepatitis B, Hepatitis C, Hepatitis D, Respiratory syncytial virus (RSV), Herpes simplex virus 1 and 2, Human Cytomegalovirus, Epstein-Barr virus , Varicella zoster virus, Coronaviruses , Poxviruses, Enterovirus 71, Rubella virus. Human papilloma virus, *Streptococcus pneumoniae, Streptococcus viridans., Staphylococcus aureus (S. aureus),* Methicillin-resistant *Staphylococcus aureus* (MRSA), Vancomycin-intermediate *Staphylococcus aureus* (VISA) , Vancomycin-resistant *Staphylococcus aureus* (VRSA), *Staphylococcus epidermidls (S. epidermidls), Clostridium Tetani, Bordetella pertussis, Bordetella paratussis, Mycobacterium, Francisella Tularensis, Toxoplasma gondii,* Candida (*C. albicans, C. glabrafa, C. parapsilosis, C. tropicalis, C. krusei and C. lusitaniae)* and/or any other infectious diseases, disorders or syndromes.

[00488] Various toxins may be treated with the pharmaceutical compositions, AAV particles, of the present invention. Non-limited examples of toxins include Riciii, Bacillus anthracis, Shiga toxin and Shiga-like toxin, Botulinum toxins.

[00489] Various tropical diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. Non-limited examples of tropical diseases include Chikungunya fever, Dengue fever, Chagas disease, Rabies, Malaria, Ebola virus, Marburg virus, West Nile Virus, Yellow Fever, Japanese encephalitis virus, St. Louis encephalitis virus.

[00490] Various foodbome illnesses and gastroenteritis may be treated with pharmaceutical compositions, AAV particles, of the present invention. Non-limited examples of foodbome illnesses and gastroenteritis include Rotavirus, Norwalk virus (Noro virus), Campylobacter jejuni, Clostridium difficile. Entamoeba histolytica, Helicobacter pyroli, Enterotoxin B of

Staphylococcus aureus, Hepatitis A virus (HAY), Hepatitis E, Listeria monocytogenes, Salmonella, Clostridium perfrmgens, and Salmonella

[00491] Various infectious agents may be treated with pharmaceutical compositions, AAV particles, of the present invention. Non-limited examples of infectious agents include adenoviruses, *Anaplasma phagocytoph ilium, Ascaris iumbricoides, Bacillus arithracis. Bacillus cereus, Bacteriodes* sp, Barmah Forest virus, *Bartonella bacilUforrnis, Bartonella henselae, Bartonella quintana,* beta-toxin of *Clostridium perfringens, Bordeteila pertussis. Bordetella parapertussis, Borrelia burgdorferi, Borrelia miyamotoi, Borrelia recurrentis, Borreha* sp., *Botulinum* toxin. *Brucella* sp., *Burkholderia pseudomallei,* California encephalitis virus, *Campylobacter, Candida albicans,* chikungunya virus, *Chlamydia psittaci, Chlamydia trachomatis, Clonorchis sinensis. Clostridium difficile bacteria, Clostridium tetani,* Colorado tick fever virus, *Corynebacterium diphiheriae, Corynebacterium minutissimum, Coxiella burnetii,* coxsackie A, coxsackie B, Crimean-Congo hemorrhagic fever virus, cytomegalovirus, dengue virus. Eastern Equine encephalitis vims, Ebola viruses, echovirus, *Ehrlichia chaffeensis., Ehrlichia equi., Ehrlichia* sp., *Entamoeba histolytica, Eriterobacter sp., Enterococcus feacaiis.* Enterovirus 71, Epstein-Barr virus (EBV), *Erysipelothrix rhusiopathiae, Escherichia coli,* Flavivirus, *Fusobacterium necrophorum. Gardnerella vaginalis.* Group B *streptococcus, Haemophilus aegyptius. Haemophilus ducreyi, Haemophilus influenzae,* hantavirus, *Helicobacter pylori,* Hepatitis A, Hepatitis B, Hepatitis C, Hepatitis D, Hepatitis E, herpes simplex virus 1 and 2, human herpes virus 6, human herpes Virus 8, human immunodeficiency virus 1 and 2, human T-cell leukemia viruses 1 and 11, influenza viruses (A, B, C), Jamestown Canyon virus, Japanese encephalitis antigenic, Japanese encephalitis virus, John Cunningham virus, junin virus, Kaposi's Sarcoma-associated Herpes Virus (KSHV), *Klebsiella granulomatis, Klebsiella sp.,* Kyasanur Forest Disease virus. La Crosse virus, Lassavirus, *Legionella pneumophila, Leptospira interrogans, Listeria monocytogenes,* lymphocytic choriomeningitis virus, lyssavirus, Machupovirus, Marburg virus, measles virus, MERS coronavirus (MERS-CoV), *Micrococcus sedentarius, Mobiluncus sp., Molluscipoxvirus, Moraxella caiarrhaiis, Morbilli- Rubeola virus,* Mumpsvirus, *Mycobacterium leprae. Mycobacterium tuberculosis. Mycobacterium ulcerans. Mycoplasma genita!ium, Mycoplasma* sp, Nairovirus, *Neisseria gonorrhoeae, Neisseria meningitidis, Nocardia,* Norwalk virus, norovirus, Omsk hemorrhagic fever virus, papillomavirus, parainfluenza viruses 1-3, parapoxvirus, parvovirus B19, *Peptostreptococccus* sp., *Plasmodium* sp., poiioviruses types 1, II, and III, *Proteus* sp., *Pseudomonas aeruginosa, Pseudomonas pseudomallei, Pseudomonas* sp., rabies virus, respiratory syncytial virus, ricin toxin, *Rickettsia australis, Rickettsia conori, Rickettsia honei,*

Rickettsia prowasekii, Ross River Virus, rotavirus, rubeilavirus, Saint Louis encephalitis, *Salmonella Typhi, Sarcoptes scabiei,* SARS-associated coronavirus (SARS-CoV), *Serratia sp.,* Shiga toxin and Shiga-like toxin, *Shigella* sp., Sin Nomhre Virus, Snowshoe hare virus, *Staphylococcus aureus. Staphylococcus epidermidis, Sireptobacillus moniliformis, Streptoccoccus pneumoniae, Streptococcus agalactiae. Streptococcus agalactiae, Streptococcus* group A-H, *Streptococcus pneumoniae, Streptococcus pyogenes, Treponema pallidum suhsp. Pallidum, Treponema pallidum var. carateum, Treponema pallidum var. endemicum, Tropheryma whippelh, Ureaplasma urealyticum,* Varicella-Zoster virus, variola virus, *Vibrio cholerae,* West Nile virus, yellow fever virus. *Yersinia enterocoliiica, Yersinia pestis,* and Zika virus.

[00492] Various rare diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As used herein, the term "rare disease" refers to any disease that affects a small percentage of the population. As a non-limiting example, the rare disease may be Acrocephaiosyndactylia, Acrodermatitis, Addison Disease, Adie Syndrome, Alagille Syndrome, Amylose, Amyotrophic Lateral Sclerosis, Angelman Syndrome, Angiolymphoid Hyperplasia with Eosinophilic Arnoid-Chiari Malformation, Arthritis, Juvenile Rheumatoid, Asperger Syndrome, Bardet-Biedl Syndrome, Barrett Esophagus, Beckwith-Wiedemann SyndiOme, Behcet Syndrome, Bloom Syndrome, Bowen's Disease, Brachial Plexus Neuropathies, Brown-Sequard Syndrome, Budd-Chiari Syndrome, Burkitt Lymphoma, Carcinoma 256, Walker, Caroli Disease, Charcot-Marie-Toolh Disease, Chediak-Higashi Syndrome, Chiari-Frommel Syndrome, Chondrodysplasia Punctata, Colonic Pseudo-Obstruction, Colorectal Neoplasms, Hereditary Nonpolyposis, Craniofacial Dysostosis, Creutzfeldt-Jakob Syndrome, Crohn Disease, Gushing Syndrome, Cystic Fibrosis, Dandy-Walker Syndrome. De Lange Syndrome, Dementia, Vascular, Dermatitis Herpeiiformis, DiGeorge Syndrome, Diffuse Cerebral Sclerosis of Schilder, Duane Retraction Syndrome, Dupuytren Contracture, Ebstein Anomaly, Eisenmenger Complex, Eilis-Van Creveld Syndrome, Encephalitis, Enchondroraatosis, Epidermal Necrolysis, Toxic, Facial Hemiatrophy, Factor XII Deficiency, Fanconi Anemia, Felly's Syndrome, Fibrous Dysplasia, Polyostotic, Fox-Fordyce Disease, Friedreich Ataxia, Fusobacterium, Gardner Syndrome, Gaucher Disease, Gerstmann Syndrome, Giant Lymph Node Hyperplasia, Glycogen Storage Disease Type I, Glycogen Storage Disease Type II, Glycogen Storage Disease Type IV, Glycogen Storage Disease Type V, Glycogen Storage Disease Type VII, Goldenhar Syndrome, Guiilain-Barre Syndrome, Haliermann's Syndrome, Hamartoma Syndrome, Multiple, Hartnup Disease, Hepatolenticular Degeneration, Hepatolenticular Degeneration, Hereditary Sensory and Motor Neuropathy,

Hirschsprung Disease, Histiocytic Necrotizing Lymphadenitis, Histiocytosis, Langerhans-Cell, Hodgkin Disease, Horner Syndrome, Huntington Disease, HyperaJdosteronism, Hyperhidrosis, Hyperostosis, Diffuse idiopathic Skeletal, Hypopituitarism, Inappropriate ADH Syndrome, intestinal Polyps, Isaacs Syndrome, Kartagener Syndrome, Kearns-Sayre Syndrome, Klippel-Feil Syndrome, Klippel-Trenaunay-Weber Syndrome, Kiuver-Bucy Syndrome, Korsakoff Syndrome, Lafora Disease, Lambert-Eaton Myasthenic Syndrome, Landau-Kleffner Syndrome, Langer-Giedion Syndrome, Leigh Disease, Lesch-Nyhan Syndrome, Leukodystrophy, Globoid Cell, Li-Fraumeni Syndrome. Long QT Syndrome, Machado-Joseph Disease, Mallory-Weiss Syndrome, Marek Disease, Marfan Syndrome, Meckel Diverticulum, Meige Syndrome, Melkersson-Rosenthal Syndrome, Meniere Disease, Mikulicz' Disease, Miller Fisher Syndrome, Mobius Syndrome, Moyamoya Disease, Mucocutaneous Lymph Node Syndrome, Mucopolysaccharidosis I, Mucopolysaccharidosis II, Mucopolysaccharidosis III, Mucopolysaccharidosis IV, Mucopolysaccharidosis VL Multiple Endocrine Neoplasia Type L Munchausen Syndrome by Proxy, Muscular Atrophy, Spinal, Narcolepsy, Neuroaxonai Dystrophies, Neuromyelitis Optica, Neuronal Ceroid-Lipofuscinoses, Niemann-Pick Diseases, Noonan Syndrome, Optic Atrophies, Hereditary, Osteitis Deformans, Osteochondritis, Osteochondrodysplasias, Osteolysis, Essential, Paget Disease Extra.mamma.ry, Paget's Disease, Mammary, Panniculitis, Nodular Nonsuppurative, Papilion-Lefevre Disease, Paralysis, Pelizaeus-Merzbacher Disease, Pemphigus, Benign Familial, Penile Induration, Pericarditis, Constrictive, Peroxisomal Disorders. Peutz-Jeghers Syndrome, Pick Disease of the Brain. Pierre Robin Syndrome, Pigmentation Disorders, Pityriasis Lichenoides, Polycystic Ovary Syndrome, Polyendocrinopathies, Autoimmune, Prader-Willi Syndrome, Pupil Disorders, Rett Syndrome, Reye Syndrome, Rubinstein-Taybi Syndrome, Sandhoff Disease, Sarcoma, Ewing's, Schnitzler Syndrome, Sjogren's Syndrome, Sjogren-Larsson Syndrome. Smith-Lemli -Opitz Syndrome, Spinal Muscular Atrophies of Childhood, Sturge-Weber Syndrome, Sweating, Gustatory, Takayasu Arteritis. Tangier Disease. Tay-Sachs Disease, Thromboangiitis Obliterans, Thyroiditis, Autoimmune, Tietze's Syndrome, Togaviridae Infections, Tolosa-Hunt Syndrome. Tourette Syndrome, Lveomeningoencephaiitic Syndrome, Waardenburg's Syndrome, Wegener Granulomatosis, Weil Disease, Werner Syndrome, Williams Syndrome, Wilms Tumor, Wolff-Parkinson-White Syndrome, Wolfram Syndrome, Wolraan Disease, Zellweger Syndrome, Zollinger-Ellison Syndrome, and von Willebrand Diseases.

[00493] Various autoimmune diseases and autoimmune-related diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As used herein, the term "autoimmune disease" refers to a disease in which the body produces antibodies that attack its

own tissues. As a non-limiting example, the autoimmune disease may be Acute Disseminated Encephalomyelitis (ADEM), Acute necrotizing hemorrhagic leukoencephalitis, Addison's disease. Agammaglobulinemia, Alopecia areata, Amyloidosis, Ankylosing spondylitis, Anti-GBM/Anti-TBM nephritis, Antiphosphoiipid syndrome (APS), Autoimmune angioedema, Autoimmune aplastic anemia, .Autoimmune dysautonomia, Autoimmune hepatitis, Autoimmune hyperlipidemia, Autoimmtme immunodeficiency. Autoimmune inner ear disease (ATED), Autoimmune myocarditis. Autoimmune oophoritis, Autoimmune pancreatitis, Autoimmune retinopathy. Autoimmune thrombocytopenic purpura (ATP), Autoimmtme thyroid disease, Autoimmune urticaria, Axonal & neuronal neuropathies, Balo disease, Behcet's disease, Bullous pemphigoid, Cardiomyopathy, Castleman disease. Celiac disease, Chagas disease. Chronic fatigue syndrome**, Chronic inflammatory demyehnating polyneuropathy (CXDP), Chronic recurrent multifocal ostomy elitis (CRMO), Churg-Strauss syndrome, Cicatricial pemphigoid/benign mucosal pemphigoid. Crohn's disease, Cogans syndrome, Cold agglutinin disease, Congenital heart block, Coxsackie myocarditis, CREST disease, Essential mixed cryoglobulinemia, Demyelinating neuropathies. Dermatitis herpetiformis, Dermatomyositis, Devic's disease (neuromyelitis optica), Discoid lupus, Dressier s syndrome, Endometriosis, Eosinophilic esophagitis. Eosinophilic fasciitis, Erythema nodosum. Experimental allergic encephalomyelitis, Evans syndrome, Fibromyalgia ***, Fibrosing alveolitis, Giant cell arteritis (temporal arteritis), Giant cell myocarditis. Glomerulonephritis, Goodpasture's syndrome. Granulomatosis with Poly angiitis (CPA) (formerly called Wegener's Granulomatosis), Graves' disease, Guillam-Barre syndrome, Hashimoto s encephalitis, Hashimoto's thyroiditis, Hemolytic anemia, Henoch-Schonlein purpura, Herpes gestatioms, Hypogammaglobulinemia, Idiopathic thrombocytopenic purpura (TTP), IgA nephropathy, IgG4-related sclerosing disease, Tmmunoregulatory lipoproteins. Inclusion body myositis, interstitial cystitis, Juvenile arthritis, Juvenile diabetes (Type 1 diabetes), Juvenile myositis, Kawasaki syndrome, Lambert-Eaton syndrome, Leukocytodastic vasculitis, Lichen planus, Lichen sclerosus. Ligneous conjunctivitis, Linear IgA disease (LAD), Lupus (SLE), Lyme disease, chronic, Meniere's disease. Microscopic-poly angiitis, Mixed connective tissue disease (MCTD), Mooren's ulcer, Mucha-Habermann disease. Multiple sclerosis. Myasthenia gravis. Myositis, Narcolepsy, Neuromyelitis optica (Devic's), Neutropenia, Ocular cicatricial pemphigoid, Optic neuritis, Palindromic rheumatism, PANDAS (Pediatric Autoimmune Neuropsychiatry Disorders Associated with Streptococcus), Paraneoplastic cerebellar degeneration. Paroxysmal nocturnal hemoglobinuria (PNH), Parry Romberg syndrome. Parsonnage-Turner syndrome, Pars planitis (peripheral uveitis). Pemphigus, Peripheral neuropathy. Perivenous encephalomyelitis, Pernicious anemia, POEMS syndrome,

Polyarteritis nodosa. Type I, II, & III autoimmune polyglandular syndromes. Polymyalgia rheumatica, Polymyositis, Postmyocardial infarction syndrome, Poslpericardiotomy syndrome. Progesterone dermatitis, Primary biliary cirrhosis, Primary sclerosing cholangitis, Psoriasis, Psoriatic arthritis, Idiopathic pulmonary fibrosis, Pyoderma gangrenosum, Pure red ceil aplasia, Raynauds phenomenon, Reactive Arthritis, Reflex sympathetic dystrophy, Reiter's syndrome. Relapsing polychondritis, Restless legs syndrome, Retroperitoneal fibrosis. Rheumatic fever. Rheumatoid arthritis. Sarcoidosis, Schmidt syndrome, Scleritis, Scleroderma, Sjogren's syndrome. Sperm & testicular autoimmumiy. Stiff person syndrome, Subacute bacterial endocarditis (SBE), Susac's syndrome, Sympathetic ophthalmia, Takayasu's arteritis. Temporal arteritis/Giant cell arteritis. Thrombocytopenic purpura (TTP), Tolosa-Hunt syndrome, Transverse myelitis, Ulcerative colitis, Undifferentiated connective tissue disease (UCTD), Uveitis, Vasculitis, Vesiculobullous dermatosis. Vitiligo, and Wegener's granulomatosis (now-termed Granulomatosis with Polyanghtis (GPA).

|00494] Various kidney diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As anon-limiting example, the kidney disease Abderhalden-Kaufmann-Lignac syndrome (Nephropathic Cystinosis), Abdominal Compartment Syndrome, Acute Kidney Failure/Acute Kidney Injury, Acute Lobar Nephroma, Acute Phosphate Nephropathy, Acute Tubular Necrosis, Adenine Phosphoribosyltransferase Deficiency, Adenovirus Nephritis, Alport Syndrome, Amyloidosis, ANCA Vasculitis Related to Endocarditis and Other Infections, Angiomyolipoma, Analgesic Nephropathy, Anorexia Nervosa and Kidney Disease, Angiotensin Antibodies and Focal Segmental Glomerulosclerosis, Antiphosphoiipid Syndrome, Anti-TNF-a Therapy-related Glomerulonephritis, APOL1 Mutations, Apparent Mineralocorticoid Excess Syndrome, Aristolochic Acid Nephropathy, Chinese Herbal Nephropathy, Balkan Endemic Nephropathy, Barrier Syndrome, Beeturia, β-Thalassemia Renal Disease, Bile Cast Nephropathy, BK Polyoma Virus Nephropathy in the Native Kidney, Bladder Rupture, Bladder Sphincter Dyssynergia, Bladder Tamponade, Border-Crossers' Nephropathy, Bourbon Virus and Acute Kidney Injury, Burnt Sugarcane Harvesting and Acute Renal Dysfunction, Byetta and Renal Failure, Clq Nephropathy, Cannabinoid Hyperemesis Acute Renal Failure, Cardiorenai syndrome, Carfilzomib-Induced Renal Injury, CFFIR5 nephropathy, Charcot-Mari e-Tooth Disease with Glomerulopathy, Cherry Concentrate and Acute Kidney Injury, Cholesterol Emboli, Churg-Strauss syndrome, Chyluria, Colistin Nephrotoxicity, Coliagenofibrotic Glomerulopathy, Collapsing Glomerulopathy, Collapsing Glomerulopathy Related to CMV, Congenital Nephrotic Syndrome, Conorenal syndrome (Mainzer-Saldino Syndrome or Saldino-Mainzer Disease), Contrast Nephropathy, Copper Sulpfate intoxication.

Cortical Necrosis, Crkotinib-relaied Acute Kidney Injury, Ciyoglobuinemia, Crystalglobulin-Tnduced Nephropathy, Crystal-induced Acute Kidney injury, Cystic Kidney Disease, Acquired, Cystinuria, Dasatinib-inducedNephrotic-Range Proteinuria, Dense Deposit Disease (MPGN Type 2), Dent Disease (X-linked Recessive Neplirolithiasis), Dialysis Disequilibrium Syndrome, Diabetes and Diabetic Kidney Disease, Diabetes insipidus, Dietary Supplements and Renal Failure, Drugs of Abuse and Kidney Disease. Duplicated Ureter, EAST syndrome, Ebola and the Kidney, Ectopic Kidney, Ectopic Ureter, Edema, Swelling, Erdheim-Chester Disease, Fabry's Disease, Familial Hypocaiciuric Hypercalcemia, Fanconi Syndrome, Fraser syndrome, Fibronectin Glomerulopathy, Fibrillary Glomerulonephritis and Immunotactoid Glomerulopathy, Fraley syndrome. Focal Segmental Glomerulosclerosis, Focal Sclerosis, Focal Glomerulosclerosis, Galloway Mowat syndrome, Giant Cell (Temporal) Arteritis with Kidney involvement, Gestational Hypertension, Gitelman Syndrome, Glomerular Diseases, Glomerular Tubular Reflux, Glycosuria, Goodpasture Syndrome, Hair Dye Ingestion and Acute Kidney injury, Hantavirus infection Podocytopathy, Hematuria (Blood in Urine), Hemolytic Uremic Syndrome (FIUS), Atypical Hemolytic Uremic Syndrome (aFIUS), Hemophagocytic Syndrome, Hemorrhagic Cystitis, Hemorrhagic Fever with Renal Syndrome (HFRS, Hantavirus Renal Disease, Korean Hemorrhagic Fever, Epidemic Hemorrhagic Fever, Nephropathis Epidemical Hemosiderosis related to Paroxysmal Nocturnal Hemoglobinuria and Hemolytic Anemia, Hepatic Glomerulopathy, Hepatic Veno-Occiusive Disease, Sinusoidal Obstruction Syndrome, Hepatitis C-Associated Renal Disease, Hepatorenal Syndrome, Herbal Supplements and Kidney Disease, High Blood Pressure and Kidney Disease, HIV-Associated Nephropathy (HIVAN), Horseshoe Kidney (Renal Fusion), Hturner's Ulcer, Plyperaidosteronism, Hypercalcemia, Hyperkalemia, Hypermagnesemia, Hyperaatremia, Hyperoxaluria, Hyperphosphatemia. Hypocalcemia, Hypokalemia, Hypokalemia-induced renal dysfunction, Hypokalemic Periodic Paralysis, Hypomagnesemia, Hyponatremia, Hypophosphatemia, IgA Nephropathy, IgG4 Nephropathy, Interstitial Cystitis, Painful Bladder Syndrome (Questionnaire), Interstitial Nephritis, Ivemark's syndrome, Ketamine-Associated Bladder Dysfunction, Kidney Stones, Nephrolithiasis, Kombucha Tea Toxicity, Lead Nephropathy and Lead-Related Nephrotoxicity, Leptospirosis Renal Disease, Light Chain Deposition Disease, Monoclonal Immunoglobulin Deposition Disease, Liddie Syndrome, Lightwood-Albright Syndrome, Lipoprotein Glomerulopathy, Lithium Nephrotoxicity, LMX1B Mutations Cause Hereditary FSGS, Loin Pain Hematuria, Lupus, Systemic Lupus Erythematosis, Lupus Kidney Disease, Lupus Nephritis, Lupus Nephritis with Antineutrophil Cytoplasmic Antibody Seropositivity, Lyme Disease-Associated Glomerulonephritis, Malarial Nephropathy, Malignancy-Associated Renal Disease,

Malignant Hypertension, Maiakopiakia, Meata! Stenosis, Medullary Cystic Kidney Disease, Medullary Sponge Kidney, Megaureter, Melamine Toxicity' and the Kidney, Membranoproliferative Glomerulonephritis, Membranous Nephropathy, MesoAmerican Nephropathy, Metabolic Acidosis, Metabolic Alkalosis, Metiiotrexate-reiated Renal Failure, Microscopic Polyangiitis, Milk-alkaiai syndrome. Minimal Change Disease, MDMA (Molly; Ecstacy; 3,4-Methylenedioxymethamphetamine) and Kidney Failure, Multicystic dysplastic kidney. Multiple Myeloma, Myeloproliferative Neoplasms and Glomerulopathy, Nail-patella Syndrome, Nephrocalcinosis, Nephrogenic Systemic Fibrosis, Nephroptosis (Floating Kidney, Renal Ptosis), Nephrotic Syndrome, Neurogenic Bladder, Nodular Glomerulosclerosis, Non-Gonococcal Urethritis, Nutcracker syndrome. Orofaciodigital Syndrome, Orotic Aciduria, Orthostatic Hypotension, Orthostatic Proteinuria, Osmotic Diuresis, Ovanan Hyperstimulation Syndrome, Page Kidney, Papillary Necrosis, Papillorenal Syndrome (Renal-Coloboma Syndrome, isolated Renal Hypoplasia), Parvovirus B19 and the Kidney, The Peritoneal-Renal Syndrome, Posterior Urethral Valve, Post-infectious Glomerulonephritis, Post-streptococcal Glomerulonephritis, Polyarteritis Nodosa, Polycystic Kidney Disease, Posterior Urethral Valves, Preeclampsia, Propofol infusion syndrome, Proliferative Glomerulonephritis with Monoclonal TgG Deposits (Nasr Disease), Propolis (Honeybee Resin) Related Renal Failure, Proteinuria (Protein in Urine), Pseudohyperaldosteronism, Pseudohypobicarbonatemia, Pseudohypoparathyroidism, Pulmonary-Renal Syndrome, Pyelonephritis (Kidney infection), Pyonephrosis, Radiation Nephropathy, Ranolazine and the Kidney, Refeeding syndrome, Reflux Nephropathy, Rapidly Progressive Glomerulonephritis, Renal Abscess, Peripnephric Abscess, Renal Agenesis, Renal Arcuate Vein Microthrombi-Associated Acute Kidney Injurs', Renal Artery Aneurysm, Renal Artery Stenosis, Renal Cell Cancer, Renal Cyst Renal Hypouricemia with Exercise-induced Acute Renal Failure, Renal Infarction, Renal Osteodystrophy, Renal Tubular Acidosis, Renin Secreting Tumors (Juxtaglomerular Cell Tumor), Reset Osmostat, Retrocaval Ureter, Retroperitoneal Fibrosis, Rhabdomyoiysis, Rhabdomyoiysis related to Bariatric Sugery, Rheumatoid Arthritis-Associated Renal Disease, Sarcoidosis Renal Disease, Salt Wasting, Renal and Cerebral, Schistosomiasis and Glomerular Disease, Schimke immuno-osseous dysplasia, Scleroderma Renal Crisis, Serpentine Fibuia-Polycystic Kidney Syndrome, Exner Syndrome, Sickle Cell Nephropathy, Silica Exposure and Chronic Kidney Disease, Sri Lankan Farmers' Kidney Disease. Sjogren's Syndrome and Renal Disease, Synthetic Cannabmoid Use and Acute Kidney injury, Kidney Disease Following Hematopoietic Cell Transplantation, Kidney Disease Related to Stem Cell Transplantation, Thin Basement Membrane Disease, Benign Familial Hematuria. Trigonitis, Tuberculosis, Genitourinary,

Tuberous Sclerosis, Tubular Dysgenesis, Immune Complex Tubuiointerstitial Nephritis Due to Autoantibodies to the Proximal Tubule Brush Border, Tumor Lysis Syndrome, Uremia, Uremic Optic Neuropathy, Ureteritis Cystica, Ureterocele, Urethral Caruncle, Urethral Strieture, Urinary incontinence, Urinary Tract Infection, Urinary Tract Obstruction, Vesicointestinal Fistula, Vesicoureteral Reflux, Volatile Anesthetics and Acute Kidney Injury, Von Hippei-Lindau Disease, Waldenstrom's Macroglobulinemic Glomerulonephritis, Warfarin-Related Nephropathy, Wasp Stings and Acute Kidney Injury, Wegener's Granulomatosis, Granulomatosis with Polyangiitis, West Nile Virus and Chronic Kidney Disease, and Wunderlich syndrome.

[00495] Various cardiovascular diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the cardiovascular disease may be Ischemic heart disease also known as coronary artery disease, cerebrovascular disease (Stroke), Peripheral vascular disease, Heart failure, Rheumatic heart disease, and Congenital heart disease.

[00496] Various antibody deficiencies may be treated with pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the antibody deficiencies may be X-Linked Agammaglobulinemia (XL.A), Autosomal Recessive Agammaglobulinemia (ARA), Common Variable Immune Deficiency (CVID), IgG (IgGl, IgG2, IgG3 and IgG4) Subclass Deficiency, Selective IgA Deficiency, Specific Antibody Deficiency (SAD), Transient Hypogammaglobulinemia of Infancy, Antibody Deficiency with Normal or Elevated immunoglobulins, Selective IgM Deficiency, Immunodeficiency with Thymoma (Good's Syndrome), Transcobalamin II Deficiency, Warts, Hypogammaglobulinemia, Infection, Myelokathexis (WHIM) Syndrome, Drug-Tnduced Antibody Deficiency, Kappa Chain Deficiency, Heavy Chain Deficiencies, Post-Meiotic Segregation (PMS2) Disorder, and Unspecified Hypogammaglobulinemia.

[00497] Various ocular diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the ocular disease may be thyroid eye disease (TED), Graves' disease (GD) and orbitopathy, Retina Degeneration, Cataract, optic atrophy, macular degeneration, Leber congenital amaurosis, retinal degeneration, cone-rod dystrophy, Usher syndrome, leopard syndrome, photophobia, and photoaversion.

[00498] Various neurological diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the neurological disease may be Absence of the Septum Pellucidum, Acid Lipase Disease, Acid Maltase Deficiency, Acquired Epileptiform Aphasia, Acute Disseminated Encephalomyelitis, Attention Deficit-Hyperactivity

Disorder (ADHD), Adie's Pupil. Adie's Syndrome, Adrenoleukody strophy, Agenesis of the Corpus Callosum, Agnosia, Aicardi Syndrome, Aicardi-Gouticres Syndrome Disorder, AIDS - Neurological Complications, Alexander Disease, Alpers* Disease, Alternating Hemiplegia, Alzheimer's Disease, Amyotrophic Lateral Sclerosis (ALS), Aneiiceplialy, Aneurysm, Angeiman Syndrome, Angiomatosis, Anoxia, Antiphosphohpid Syndrome, Aphasia, Apraxia, Arachnoid Cysts, Arachnoiditis, Arnold-Chiari Malformation. Arteriovenous Malformation, Asperger Syndrome, Ataxia, Ataxia Telangiectasia, Ataxias and Cerebellar or Spinocerebellar Degeneration, Atrial Fibrillation and Stroke, Attention Deficit-Hyperactrviiy Disorder, Autism Spectrum Disorder, Autonomic Dysfunction, Back Pain, Barth Syndrome, Batten Disease, Becker's Myotonia, Behcet's Disease, Bell's Palsy, Benign Essential Blepharospasm, Benign Focal Amyotrophy, Benign intracranial Hypertension, Bemhardt-Roth Syndrome, Binswanger's Disease, Blepharospasm, Bloch-Sulzberger Syndrome, Brachial Plexus Birth Injuries, Brachial Plexus injuries, Bradbury-Egg! eston Syndrome, Brain and Spinal Tumors, Brain Aneurysm, Brain Injury, Brown-Sequard Syndrome, Bulbospinal Muscular Atrophy, Cerebral Autosomal Dominant Arteriopathy with Sub-cortical Infarcts and Leukoencephalopathy (CADASIL), Canavan Disease, Carpal Tunnel Syndrome, Causalgia, Cavernomas, Cavernous Angioma, Cavernous Malformation, Central Cervical Cord Syndrome, Central Cord Syndrome. Central Pain Syndrome, Central Pontine Myelinolysis, Cephalic Disorders, Ceramidase Deficiency, Cerebellar Degeneration, Cerebellar Hypoplasia, Cerebral Aneurysms, Cerebral Arteriosclerosis, Cerebral Atrophy, Cerebral Beriberi, Cerebral Cavernous Malformation, Cerebral Gigantism, Cerebral Hypoxia, Cerebral Palsy, Cerebro-Ocuio-Facio-Skeletal Syndrome (COFS), Charcot-Mane-Tooth Disease, Chiari Malformation, Cholesterol Ester Storage Disease, Chorea, Choreoacanthocytosis, Chronic Inflammatory Dernyehnating Polyneuropathy (CIDP), Chronic Orthostatic intolerance, Chronic Pain. Cockayne Syndrome Type II. Coffin Lowry Syndrome, Colpocephaly, Coma, Complex Regional Pain Syndrome, Congenital Facial Diplegia, Congenital Myasthenia, Congenital Myopathy, Congenital Vascular Cavernous Malformations, Cortieobasal Degeneration, Cranial Arteritis, Craniosynostosis, Cree encephalitis, Creutzfeldt-Jakob Disease, Cumulative Trauma Disorders, Cushing's Syndrome, Cytomegalic Inclusion Body Disease, Cytomegalovirus Infection, Dancing Eyes-Dancing Feet Syndrome, Dandy-Walker Syndrome, Dawson Disease, De Morsier's Syndrome, Dejerine-Klumpke Palsy, Dementia, Dementia -Multi-Infarct. Dementia - Semantic, Dementia -Subcortical, Dementia With Lewy Bodies, Dentate Cerebellar Ataxia, Dentatorubral Atrophy, Dermatomyositis, Developmental Dyspraxia, Devic's Syndrome, Diabetic Neuropathy, Diffuse Sclerosis, Dravet Syndrome, Dysautonomia, Dysgraphia. Dyslexia. Dysphagia, Dyspraxia. Dyssynergia

Cerebel!aris Myocionica, Dyssynergia Cerebeliaris Progressiva, Dystonias, Early infantile Epileptic Encephalopathy, Empty Sella Syndrome, Encephalitis, Encephalitis Lethargies, Encephaloceles, Encephalopathy, Encephalopathy (familial infantile), Eneephalotrigeminal Angiomatosis, Epilepsy, Epileptic Hemiplegia, Erb's Palsy, Erb-Duchenne and Dejerine-Klumpke Palsies, Essential Tremor, Extrapontme Myeiinoiysis, Fabry Disease, Fahr's Syndrome, Fainting, Familial Dysautonomy a, Familial Hemangioma, Familial idiopathic Basal Ganglia Calcification, Familial Periodic Paralyses, Familial Spastic Paralysis, Farber's Disease, Febrile Seizures, Fibromuscuiar Dysplasia, Fisher Syndrome, Floppy Infant Syndrome, Foot Drop, Friedreich's Ataxia, Frontotemporal Dementia, Gaucher Disease, Generalized Gangliosidoses, Gerstmann's Syndrome, Gerstmann-S traussIer-Scheinker Disease, Giant Axonal Neuropathy. Giant Ceil Arteritis, Giant Cell Inclusion Disease, Globoid Cell Leukodystrophy, Glossopharyngeal Neuralgia, Glycogen Storage Disease, GuiHain-Barre Syndrome, HaHervorden-Spatz Disease, Head injury. Headache, Hemicrania Continua, Hemifacial Spasm, Hemiplegia Alterans, Hereditaiy Neuropathies, Hereditaiy Spastic Paraplegia, Heredopathia Atactica Polyneuritiformis, Herpes Zoster, I-Ierpes Zoster Oticus, Hirayama Syndrome, Holmes-Adie syndrome, Ho!oprosencephaly , HTLV-1 Associated Myelopathy, Hughes Syndrome, Huntington's Disease, Hydranencephaly, Hydrocephalus, Hydrocephalus - Normal Pressure, Hydromyeiia, Hypereortisoiism, Hypersomnia, Hypertonia, Hypotonia, Hypoxia, immune-Mediated Encephalomyelitis, Inclusion Body Myositis, Incontinentia Pigments, Infantile Hypotonia, Infantile Neuroaxonal Dystrophy, Infantile Phytanic Acid Storage Disease, Infantile Refsum Disease, Infantile Spasms, Inflammatory Myopathies, Imeneephaly, intestinal Lipodystrophy, Intracranial Cysts, Intracranial Hypertension, Isaacs' Syndrome, Joubert Syndrome, Kearns-Sayre Syndrome, Kennedy's Disease, Kinsbourae syndrome, Heine-Levin Syndrome, K!ippel-Fei! Syndrome, Kiippel-Trenaunay Syndrome (KTS), Kliiver-Bucy Syndrome, Korsakoff s Amnesic Syndrome, Krabbe Disease, Kugelberg-Welander Disease, Kuru, Lambert-Eaton Myasthenic Syndrome, Landau-Kleffner Syndrome, Lateral Femoral Cutaneous Nerve Entrapment, Lateral Medullary Syndrome, Learning Disabilities, Leigh's Disease, Lennox-Gastaut Syndrome, Lesch-Nyhan Syndrome, Leukodystrophy, Levine-Critchley Syndrome, Lewy Body Dementia, Lipid Storage Diseases, Lipoid Proteinosis, Lissencephaly, Locked-In Syndrome, Lou Gehrig's Disease, Lupus - Neurological Sequelae, Lyme Disease - Neurological Complications, Machado-Joseph Disease, Micrencephaly, Megalencephaly, Melkersson-Rosenthal Syndrome, Meningitis, Meningitis and Encephalitis, Menkes Disease, Meralgia Paresthetica, Metachroxnatic Leukodystrophy, Microcephaly, Migraine, Miller Fisher Syndrome, Mini Stroke, Mitochondrial Myopathy, Moehius Syndrome,

Monomelic Amyotrophy, Motor Neuron Diseases, Moyamoya Disease, Mucolipidoses. Mucopolysaccharidoses, Multi-Infarct Dementia, Multifocal Motor Neuropathy, Multiple Sclerosis, Multiple System Atrophy, Multiple System Atrophy with Orthostatic Hypotension, Muscular Dystrophy, Myasthenia - Congenital, Myasthenia Gravis, Myelinoclastic Diffuse Sclerosis, Myoclonic Encephalopathy of Infants, Myoclonus, Myopathy, Myopathy- Congenital, Myopathy -Thyrotoxic, Myotonia, Myotonia Congenita, Narcolepsy, Neuroacanthocytosis, Neurodegeneration with Brain Iron Accumulation, Neurofibromatosis, Neuroleptic Malignant Syndrome, Neurological Complications of AIDS, Neurological Complications of Lyme Disease, Neurological Consequences of Cytomegalovirus Infection, Neurological Manifestations of Pompe Disease, Neurological Sequelae Of Lupus, Neuromyelitis Optica, Neuromyotonia, Neuronal Ceroid Lipofuscinosis, Neuronal Migration Disorders, Neuropathy- Hereditary, Neurosarcoidosis, Neurosyphilis, Neurotoxicity, Nevus Cavemosus, Niemann-Pick Disease, O'Sullivan-McLeod Syndrome, Occipital Neuralgia, Ohtahara Syndrome, Olivopontocerebellar Atrophy, Opsoclonus Myoclonus, Orthostatic Hypotension, Overuse Syndrome, Pain -Chronic, Pantothenate Kinase-Associated Neurodegeneration, Paraneoplastic Syndromes, Paresthesia, Parkinson's Disease, Paroxysmal Choreoathetosis, Paroxysmal Hemicrania, Parry-Romberg, Pelizaeus-Merzbacher Disease, Pena Shokeir II Syndrome, Perineural Cysts, Periodic Paralyses, Peripheral Neuropathy, Periventricular Leukomalacia, Persistent Vegetative State, Pervasive Developmental Disorders, Phytanic Acid Storage Disease, Pick's Disease, Pinched Nerve, Piriformis Syndrome, Pituitary Tumors, Polymyositis, Pompe Disease, Porencephaly, Post-Polio Syndrome, Postherpetic Neuralgia, Postinfectious Encephalomyelitis, Postural Hypotension, Postural Orthostatic Tachycardia Syndrome, Postural Tachycardia Syndrome, Primary Dentatum Atrophy, Primary Lateral Sclerosis, Primary Progressive Aphasia, Prion Diseases, Progressive Hemifacial Atrophy, Progressive Locomotor Ataxia, Progressive Multifocal Leukoencephalopathy, Progressive Sclerosing Poliodystrophy, Progressive Supranuclear Palsy, Prosopagnosia, Pseudo-Torch syndrome, Pseudotoxoplasmosis syndrome, Pseudotumor Cerebri, Psychogenic Movement, Ramsay Hunt Syndrome I, Ramsay Hunt Syndrome II, Rasraussen's Encephalitis, Reflex Sympathetic Dystrophy Syndrome, Refsuni Disease, Refsuni Disease - Infantile, Repetitive Motion Disorders, Repetitive Stress Injuries, Restless Legs Syndrome, Retrovirus-Associated Myelopathy, Rett Syndrome, Reye's Syndrome, Rheumatic Encephalitis, Riley-Day Syndrome, Sacral Nerve Root Cysts. Saint Vitus Dance, Salivary Gland Disease, Sandhoff Disease, Schilder's Disease, Schizencephaly, Seiteiberger Disease, Seizure Disorder, Semantic Dementia, Septo-Optic Dysplasia, Severe Myoclonic Epilepsy of Infancy (SMEI), Shaken Baby Syndrome, Shingles, Shy-Drager Syndrome, Sjogren's Syndrome, Sleep Apnea,

Sleeping Sickness, Sotos Syndrome, Spasticity, Spina Bifida, Spinal Cord Infarction, Spinal Cord Injury, Spinal Cord Tumors, Spinal Muscular Atrophy, Spinocerebellar Atrophy, Spinocerebellar Degeneration, Steele-Richardson-Olszewski Syndrome, Stiff-Person Syndrome, Striatonigral Degeneration, Stroke, Sturge-Weber Syndrome, Subacute Sclerosing Panencephalitis, Subcortical Arteriosclerotic Encephalopathy, Short-lasting, Unilateral, Neuralgiform (SUNCT) Headache. Swallowing Disorders, Sydenham Chorea, Syncope, Syphilitic Spinal Sclerosis, Syringohydromyelia, Syringomyelia, Systemic Lupus Erythematosus, Tabes Dorsalis, Tardive Dyskinesia, Tarlov Cysts, Tay-Sachs Disease, Temporal Arteritis, Tethered Spinal Cord Syndrome, Thomsen's Myotonia, Thoracic Outlet Syndrome, Thyrotoxic Myopathy, Tic Douloureux, Todd's Paralysis, Tourette Syndrome, Transient Ischemic Attack, Transmissible Spongiform Encephalopathies, Transverse Myelitis, Traumatic Brain Injury, Tremor, Trigeminal Neuralgia, Tropical Spastic Paraparesis, Troyer Syndrome, Tuberous Sclerosis, Vascular Erectile Tumor, Vasculitis Syndromes of the Central and Peripheral Nervous Systems, Von Economo's Disease, Von Hippel-Lindau Disease (VHL), Von Recklinghausen's Disease, Wallenberg's Syndrome, Werdnig-Hoffman Disease, Wernicke-Korsakoff Syndrome, West Syndrome, Whiplash, Whipple's Disease, Williams Syndrome, Wilson Disease, **Wolman's** Disease. X-Linked Spinal and Bulbar Muscular Atrophy.

[00499] Various psychological disorders may be treated with pharmaceutical compositions, AAV particles, of the present invention. As anon-limiting example, the psychological disorders may be Aboulia, Absence epilepsy, Acute stress Disorder, Adjustment Disorders, Adverse effects of medication NOS, Age related cognitive decline, Agoraphobia, Alcohol Addiction, Alzheimer's Disease, Amnesia (also known as Amnestic Disorder), Amphetamine Addiction, Anorexia Nervosa, Anterograde amnesia. Antisocial personality disorder (also known as Sociopathy), Anxiety Disorder (Also known as Generalized Anxiety Disorder). Anxiolytic related disorders, Asperger's Syndrome (now part of Autism Spectrum Disorder), Attention Deficit Disorder (Also known as ADD), Attention Deficit Hyperactivity Disorder (Also known as ADHD), Autism Spectrum Disorder (also known as Autism), Autophagia, Avoidant Personality Disorder, Barbiturate related disorders, Benzodiazepine related disorders, Bereavement, Bibliomania, Binge Eating Disorder, Bipolar disorder (also known as Manic Depression, includes Bipolar I and Bipolar ïï), Body Dysmorphic Disorder, Borderline intellectual functioning. Borderline Personality Disorder, Breathing-Related Sleep Disorder, Brief Psychotic Disorder, Bruxism, Bulimia Nervosa, Caffeine Addiction, Cannabis Addiction, Catatonic disorder. Catatonic schizophrenia, Childhood amnesia. Childhood Disintegrative Disorder (now part of Autism Spectrum Disorder), Childhood Onset Fluency Disorder (formerly

known as Stuttering), Orcadian Rhythm Disorders, Claustrophobia, Cocaine related disorders. Communication disorder, Conduct Disorder, Conversion Disorder, Cotard delusion. Cyclothymia (also known as Cyclothymic Disorder), Delerium, Delusional Disorder, dementia , Dependent Personality Disorder (also known as Asthenic Personality Disorder), Depersonalization disorder (now known as Depersonalization/Derealization Disorder), Depression (also known as Major Depressive Disorder), Depressive personality disorder, Derealization disorder (now known as Depersonalization/Derealization Disorder), Dermotillomania, Desynchronosis, Developmental coordination disorder, Diogenes Syndrome, Disorder of written expression, Dispareunia, Dissocial Personality Disorder, Dissociative Amnesia. Dissociative Fugue, Dissociative identity Disorder (formerly known as Multiple Personality Disorder), Down syndrome, Dyslexia, Dyspareunia, Dysthymia (now known as Persistent Depressive Disorder), Eating disorder NOS, Ekbom's Syndrome (Delusional Parasitosis), Emotionally unstable personality disorder. Encopresis, Enuresis (bedwetting), Erotomania, Exhibitionistic Disorder, Expressive language disorder. Factitious Disorder, Female Sexual Disorders, Fetishistic Disorder, Folie à deux, Fregoli delusion, Frotteuristic Disorder, Fugue State, Ganser syndrome, Gambling Addiction, Gender Dysphoria (formerly known as Gender Identity Disorder). Generalized Anxiety Disorder, General adaptation syndrome. Grandiose delusions, Hallucinogen Addiction, Haltiose personality disorder, Histrionic Personality Disorder, Primary hypersomnia, Huntington's Disease, Hypoactive sexual desire disorder, Hypochondriasis, Hypomania, Hyperkinetic syndrome, Hypersomnia, Hysteria, impulse control disorder, Impulse control disorder NOS, Inhalant Addiction, insomnia, Intellectual Development Disorder, Intermittent Explosive Disorder, Joubert syndrome, Kleptomania, Korsakoff's syndrome. Lacunar amnesia. Language Disorder, Learning Disorders, Major Depression (also known as Major Depressive Disorder), major depressive disorder, Male Sexual Disorders, Malingering, Mathematics disorder, Medication-related disorder, Melancholia, Mental Retardation (now known as intellectual Development Disorder), Misophonia, Morbid jealousy. Multiple Personality Disorder (now known as Dissociative Identity Disorder). Munchausen Syndrome, Munchausen by Proxy, Narcissistic Personality Disorder, Narcolepsy, Neglect of child, Neurocognitive Disorder (formerly known as Dementia), Neuroieptic-reiated disorder, Nightmare Disorder, Non Rapid Eye Movement, Obsessive-Compulsive Disorder, Obsessive-Compulsive Personality Disorder (also known as Anankastic Personality Disorder), Oneirophrenia, Onychophagy, Opioid Addiction, Oppositional Defiant Disorder, Orthorexia (ON), Pain disorder. Panic attacks. Panic Disorder, Paranoid Personality Disorder, Parkinson's Disease, Partner relational problem, Passive-aggressive personality disorder, Pathological

gambling, Pedophilic Disorder, Perfectionism, Persecutory delusion, Persistent Depressive Disorder (also known as Dysthymia), Personality change due to a general medical condiiion, Personality disorder, Pervasive developmental disorder (FDD), Pheneyc!idine related disorder, Phobic disorder, Phonological disorder, Physical abuse, Pica, Polysubstance related disorder, Postpartum Depression, Post-traumatic embiitermeni disorder (PTED), Post Traumatic Stress Disorder, Premature ejaculation, Premenstrual Dysphoric Disorder, Psychogenic amnesia, Psychological factor affecting medical condition, Psychoneurotic personality disorder, Psychotic disorder, not otherwise specified, Pyromania, Reactive Attachment Disorder, Reading disorder, Recurrent brief depression, Relational disorder, REM Sleep Behavior Disorder, Restless Leg Syndrome, Retrograde amnesia, Retts Disorder (now part of Autism Spectrum Disorder), Rumination syndrome, Sadistic personality disorder, Schizoaffective Disorder, Schizoid Personality Disorder, Schizophrenia, Schizophreniform disorder, Schizotypal Personality Disorder, Seasonal Affective Disorder, Sedative, Hypnotic, or Anxiolytic Addiction, Selective Mutism, Self-defeating personality disorder, Separation Anxiety Disorder, Sexual Disorders Female, Sexual Disorders Male, Sexual Addiction, Sexual Masochism Disorder, Sexuai Sadism Disorder, Shared Psychotic Disorder, Sleep Arousal Disorders, Sleep Paralysis, Sleep Terror Disorder (now part of Nightmare Disorder, Social Anxiety Disorder, Somatization Disorder, Specific Phobias, Stendhal syndrome, Stereotypic movement disorder, Stimulant Addiction, Stuttering (now known as Childhood Onset Fluency Disorder), Substance related disorder, Tardive dyskinesia, Tobacco Addiction, Tourettes Syndrome, Transient tic disorder, Transient global amnesia, Transvesiic Disorder, Trichotillomania, Undifferentiated Somatoform Disorder, Vaginismus, and Voyeuristic Disorder.

[00500] Various lung diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the lung diseases may be Asbestosis, Asthma, Bronchiectasis, Bronchitis, Chronic Cough, Chronic Obstructive Pulmonary Disease (COPD), Croup, Cystic Fibrosis, Hantavirus, Idiopathic Pulmonary Fibrosis, Pertussis, Pleurisy, Pneumonia, Pulmonary Embolism, Pulmonary Hypertension, Sarcoidosis, Sleep Apnea, Spirometry, Sudden Infant Death Syndrome (SIDS), Tuberculosis, Alagille Syndrome, Autoimmune Hepatitis, Biliary Atresia, Cirrhosis, ERCP (Endoscopic Retrograde Cholangiopancreatography), and Hemochromatosis. Nonalcoholic Steatohepatitis, Porphyria, Primary Biliary Cirrhosis, Primary Sclerosing Cholangitis.

[ΘΘ501] Various bone diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As anon-limiting example, the bone diseases may be osteoporosis, neurofibromatosis, osteogenesis imperfecta (01), rickets, osteosarcoma,

achondroplasia, fracture, osteomyelitis, Ewing tumour of bone, osteomalacia, hip dysplasia, Paget disease of bone, marble bone disease, osteochondroma, bone cancer, bone disease, osteochondrosis, osteoma, fibrous dysplasia, cleidocranial dysostosis, osteoclastoma, bone cyst, metabolic bone disease, meiorheostosis, callus, Caffey syndrome, and mandibulofacial dysostosis.

[00502] Various blood diseases may be treated with pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the blood diseases may be Anemia and CKD (for health care professionals), Aplastic Anemia and Myelodysplasia Syndromes, Deep Vein Thrombosis, Hemochromatosis, Hemophilia, Henoch-Schonlein Purpura, idiopathic Thrombocytopenic Purpura, Iron-Deficiency Anemia, Pernicious Anemia, Pulmonary Embolism, Sickle Cell Anemia, Sickle Cell Trait and Other Hemoglobinopathies, Thalasseniia, Thrombotic Thrombocytopenic Purpura, and Von Willebrand Disease.

[00503] Various diseases associated with TNF-alpha may be treated with the pharmaceutical compositions, AAV particles, of the present invention. As anon-limiting example, the disease may be respiratory disorder; asthma; allergic and nonaliergic asthma; asthma due to infection; asthma due to infection with respiratory syncytial virus (RSV); chronic obstructive pulmonary disease (COPD); a condition involving airway inflammation; eosinophilia; fibrosis and excess mucus production; cystic fibrosis; pulmonary fibrosis; an atopic disorder; atopic dermatitis; urticaria, eczema; allergic rhinitis; allergic enterogastritis; an inflammatory and/or autoimmune condition of the skin; an inflammatory and/or autoimmune condition of gastrointestinal organs; inflammatory bowel diseases (1BD); ulcerative colitis; Crohn's disease; an inflammatory and/or autoimmune condition of the liver; liver cirrhosis; liver fibrosis; liver fibrosis caused by hepatitis B and/or C virus, scleroderma; tumors or cancers; hepatocellular carcinoma, gliobiastoma; lymphoma; Hodgkin's lymphoma; a viral infection; a bacterial infection; a parasitic infection; HTLV-1 infection; suppression of expression of protective type 1 immune responses, and suppression of expression of a protective type 1 immune response during vaccination, rheumatoid arthritis, osteoarthritis, juvenile chronic arthritis, septic arthritis, Lyme arthritis, psoriatic arthritis, reactive arthritis, spondyloarthropathy, systemic lupus erythematosus, Crohn's disease, ulcerative colitis, inflammatory bowel disease, insulin dependent diabetes mellitus, thyroiditis, asthma, allergic diseases, psoriasis, dermatitis scleroderma, graft versus host disease, organ transplant rejection, acute or chronic immune disease associated with organ transplantation, sarcoidosis, atherosclerosis, disseminated intravascular coagulation, Kawasaki's disease, Grave's disease, nephrotic syndrome, chronic fatigue syndrome, Wegener's granulomatosis, Henoch-Schoenlein purpurea, microscopic vasculitis of the kidneys, chronic

active hepatitis, uveitis, septic shock, toxic shock syndrome, sepsis syndrome, cachexia, infectious diseases, parasitic diseases, acquired immunodeficiency syndrome, acute transverse myelitis, Huntington's chorea, Parkinson's disease, Alzheimer's disease, stroke, primary biliary cirrhosis, hemolytic anemia, malignancies, heart failure, myocardial infarction, Addison's disease, sporadic, polyglandular deficiency type I and polyglandular deficiency type II, Schmidt's syndrome, adult (acute) respirator}' distress syndrome, alopecia, alopecia greata, seronegative arthropathy, arthropathy, Reiter's disease, psoriatic arthropathy, ulcerative colitic arthropathy, enteropathic synovitis, chlamydia, yersinia and salmonella associated arthropathy, spondyloarthropathy, atheromatous disease/arteriosclerosis, atopic allergy, autoimmune bullous disease, pemphigus vulgaris, pemphigus foHaceus, pemphigoid, linear IgA disease, autoimmune haemolytic anaemia, Coombs positive haemolytic anaemia, acquired pernicious anaemia, juvenile pernicious anaemia, myalgic encephalitis/Royal Free Disease, chronic mucocutaneous candidiasis, giant cell arteritis, primary sclerosing hepatitis, cryptogenic autoimmune hepatitis, Acquired inmiunodeficiency Disease Syndrome, Acquired immunodeficiency Related Diseases, hepatitis B, hepatitis C, common varied immunodeficiency (common variable hypogaramaglobulinaemia), dilated cardiomyopathy, female infertility, ovarian failure, premature ovarian failure, fibrotic lung disease, cryptogenic fibrosing alveolitis, post¬ inflammatory interstitial lung disease, interstitial pneumonitis, connective tissue disease associated interstitial lung disease, mixed connective tissue disease associated lung disease, systemic sclerosis associated interstitial lung disease, rheumatoid arthritis associated interstitial lung disease, systemic lupus erythematosus associated lung disease, dermatomyositis/poly myositis associated lung disease, Sjogren's disease associated lung disease, ankylosing spondylitis associated lung disease, vasculitis diffuse lung disease, baemosiderosis associated lung disease, drug-induced interstitial lung disease, fibrosis, radiation fibrosis, bronchiolitis obliterans, chronic eosinophilic pneumonia, lymphocytic infiltrative lung disease, postinfectious interstitial lung disease, gouty arthritis, autoimmune hepatitis, type-1 autoimmune hepatitis (classical autoimmune or lupoid hepatitis), type-2 autoimmune hepatitis (anti-LKM antibody hepatitis), autoimmune mediated hypoglycaemia, type B insulin resistance with acanthosis nigricans, hypoparathyroidism, acute immune disease associated with organ transplantation, chronic immune disease associated with organ transplantation, osteoarthrosis, primary sclerosing cholangitis, psoriasis type I, psoriasis type 2, idiopathic leucopaenia, autoimmune neutropaenia, renal disease NOS, giomerulonephntides, microscopic vasculitis of the kidneys, Lyme disease, discoid lupus erythematosus, male infertility idiopathic or NOS, sperm autoimmunity, multiple sclerosis (all subtypes), sympathetic ophthalmia, pulmonary

hypertension secondary to connective tissue disease, Goodpasture's syndrome, pulmonary manifestation of polyarteritis nodosa, acute rheumatic fever, rheumatoid spondylitis, Still's disease, systemic sclerosis, Sjorgren's syndrome, Takayasu's disease/arteritis, autoimmune thrombocytopaenia, idiopathic thrombocytopaenia, autoimmune thyroid disease, hyperthyroidism, goitrous autoimmune hypothyroidism (Hashimoto's disease), atrophic autoimmune hypothyroidism, primary myxoedema, phacogenic uveitis, primary vasculitis, vitiligo acute liver disease, chronic liver diseases, alcoholic cirrhosis, alcohol-induced liver injury, choleostasis, idiosyncratic liver disease, drug-Induced hepatitis, non-alcoholic steatohepatitis, allergy and asthma, group B streptococci (GBS) infection, mental disorders (e.g., depression and schizophrenia), Th2 'Type and Thl Type mediated diseases, acute and chronic pain (different forms of pain), and cancers such as lung, breast, stomach, bladder, colon, pancreas, ovarian, prostate and rectal cancer and hematopoietic malignancies (leukemia and lymphoma) ahetalipoproteinemia, acrocyanosis, acute and chronic parasitic or infectious processes, acute leukemia, acute lymphoblastic leukemia (ALL), acute myeloid leukemia (AML), acute or chronic bacterial infection, acute pancreatitis, acute renal failure, adenocarcinomas, aerial ectopic beats, AIDS dementia complex, alcohol-induced hepatitis, allergic conjunctivitis, allergic contact dermatitis, allergic rhinitis, allograft rejection, alpha-] -antitrypsin deficiency, amyotrophic lateral sclerosis, anemia, angina pectoris, anterior horn ceil degeneration, anti-CD3 therapy, antiphosphoiipid syndrome, anti-receptor hypersensitivity reactions, aortic and peripheral aneurysms, aortic dissection, arterial hypertension, arteriosclerosis, arteriovenous fistula, ataxia, atrial fibrillation (sustained or paroxysmal), atrial flutter, atrioventricular block, B ceil lymphoma, bone graft rejection, bone marrow transplant (BMT) rejection, bundle branch block, Rurkitt's lymphoma, burns, cardiac arrhythmias, cardiac stun syndrome, cardiac tumors, cardiomyopathy, cardiopulmonary bypass inflammation response, cartilage transplant rejection, cerebellar cortical degenerations, cerebellar disorders, chaotic or multifocal atrial tachycardia, chemotherapy associated disorders, chronic myelocytic leukemia (CML), chronic alcoholism, chronic inflaminatory pathologies, chronic lymphocytic leukemia (CLL), chronic obstructive pulmonary disease (COPD), chronic salicylate intoxication, colorectal carcinoma, congestive heart failure, conjunctivitis, contact dermatitis, corpulmonale, coronary artery disease, Creutzfeldt-Jakob disease, culture negative sepsis, cystic fibrosis, cytokine therapy associated disorders, dementia pugilistica, demyelinating diseases, dengue hemorrhagic fever, dermatitis, dermatologic conditions, diabetes, diabetes mellitus, diabetic arteriosclerotic disease, Diffuse Lewy body disease, dilated congestive cardiomyopathy, disorders of the basal ganglia, Down's Syndrome in middle age, drug-induced movement.

disorders induced by drugs which block CNS dopamine receptors, drug sensitivity, eczema, encephalomyelitis, endocarditis, endocrinopathy, epiglottitis, Epstein-Barr virus infection, erythromelalgia, extrapyramidal and cerebellar disorders, familial hemophagocytie lymphohistiocytosis, fetal thymus implant rejection, Friedreich's ataxia, functional peripheral arterial disorders, fungal sepsis, gas gangrene, gastric ulcer, glomerular nephritis, graft rejection of any organ or tissue, gram negative sepsis, gram positive sepsis, granulomas due to intracellular organisms, hairy cell leukemia, Hallervorden-Spatz disease, Hashimoto's thyroiditis, hay fever, heart transplant rejection, hemochromatosis, hemodialysis, hemolytic uremic syndrome/thrombolytic thrombocytopenic purpura, hemorrhage, hepatitis (A), His bundle arrhythmias, HIV infection/HIV neuropathy. Hodgkin's disease, hyperkinetic movement disorders, hypersensitivity reactions, hypersensitivity pneumonitis, hypertension, hypokinetic movement disorders, hypothalamic-pituitary-adrenal axis evaluation, idiopathic Addison's disease, idiopathic pulmonary fibrosis, antibody mediated cytotoxicity, asthenia, infantile spinal muscular atrophy, inflammation of the aorta, influenza a, ionizing radiation exposure, iridocyclitis/uveitis/optic neuritis, ischemia-reperfusion injury, ischemic stroke, juvenile rheumatoid arthritis (JRA), juvenile spinal muscular atrophy, Kaposi's sarcoma, kidney transplant rejection, legionella, leishmaniasis, leprosy, lesions of the corticospinal system, lipedema, liver transplant rejection, lymphedema, malaria, malignant lymphoma, malignant histiocytosis, malignant melanoma, meningitis, meningococcemia, metabolic/idiopathic, migraine headache, mitochondrial multi-system disorder, mixed connective tissue disease, monoclonal gammopathy, multiple myeloma, multiple systems degenerations (Menzei, Dejerine-Thomas, Shy-Drager, and Machado-Joseph), myasthenia gravis, mycobacterium avium intraceliulare, mycobacterium tuberculosis, myelodysplastic syndrome, myocardial infarction, myocardial ischemic disorders, nasopharyngeal carcinoma, neonatal chronic lung disease, nephritis, nephrosis, neurodegenerative diseases, neurogenic I muscular atrophies, neutropenic fever, non-Hodgkins lymphoma, occlusion of the abdominal aorta and its branches, occlusive arterial disorders, OKT3® therapy, orchitis/epidydimitis, orchitis/vasectomy reversal procedures, organomegaly, osteoporosis, pancreas transplant rejection, pancreatic carcinoma, paraneoplastic syndrome/hypercalcemia of malignancy, parathyroid transplant rejection, pelvic inflammatory disease, perennial rhinitis, pericardial disease, peripheral atherosclerotic disease, peripheral vascular disorders, peritonitis, pernicious anemia, Pneumocystis carinii pneumonia, pneumonia. POEMS syndrome (polyneuropathy, organomegaly, endocrinopathy, monoclonal gammopathy, and skin changes syndrome), post perfusion syndrome, post pump syndrome, post-Mi cardiotomy syndrome, preeclampsia, progressive supranucleo palsy, primary pulmonary

hypertension, radiation therapy, Raynaud's phenomenon and disease, Raynaud's disease, Refsum's disease, regular narrow QRS tachycardia, renovascular hypertension, reperfusion injuty, restrictive cardiomyopathy, sarcomas, scleroderma, senile chorea, senile dementia of Lewy body type, seronegative arthropathies, shock, sickle ceil anemia, skin allograft rejection, skin changes syndrome, small bowel transplant rejection, solid tumors, specific arrhythmias, spinal ataxia, spinocerebellar degenerations, streptococcal myositis, structural lesions of the cerebellum, subacute sclerosing panencephalitis, syncope, syphilis of the cardiovascular system, systemic anaphylaxis, systemic inflammatory response syndrome, systemic onset juvenile rheumatoid arthritis, T-cell or FAB ALL, telangiectasia, thromboangitis obliterans, thrombocytopenia, toxicity, transplants, trauma/hemorrhage, type III hypersensitivity reactions, type IV hypersensitivity, unstable angina, uremia, urosepsis, umcaria, valvular heart diseases, varicose veins, vasculitis, venous diseases, venous thrombosis, ventricular fibrillation, viral and fungal infections, viral encephalitis/aseptic meningitis, viral-associated hemophagocytic syndrome, Wernicke-Korsakoff syndrome, Wilson's disease, xenograft rejection of any organ or tissue, acute coronary syndromes, acute idiopathic polyneuritis, acute inflammatory demyelinating polyradiculoneuropathy, acute ischemia, adult Still's disease, alopecia greata, anaphylaxis, anti-phospholipid antibody syndrome, aplastic anemia, arteriosclerosis, atopic eczema, atopic dermatitis, autoimmune dermatitis, autoimmune disorder associated with streptococcus infection, autoimmune enteropathy, autoimmune hearing loss, autoimmune lymphoproliferative syndrome (ALPS), autoimmune myocarditis, autoimmune premature ovarian failure, blepharitis, bronchiectasis, bullous pemphigoid, cardiovascular disease, catastrophic antiphospholipid syndrome, celiac disease, cervical spondylosis, chronic ischemia, cicatricial pemphigoid, clinically isolated syndrome (CIS) with risk for multiple sclerosis, conjunctivitis, childhood onset psychiatric disorder, chronic obstructive pulmonary disease (COPD), dacryocystitis, dermatomyositis, diabetic retinopathy, diabetes meilitus, disk herniation, disk prolapse, drug induced immune hemolytic anemia, endocarditis, endometriosis, endophthalmitis, episcleritis, erythema multiforme, erythema multiforme major, gestational pemphigoid, Guillain-Barré syndrome (GBS), hay fever, Hughes syndrome, idiopathic Parkinson's disease, idiopathic interstitial pneumonia, IgE-mediated allergy, immune hemolytic anemia, inclusion body myositis, infectious ocular inflammatory disease, inflammatory demyelinating disease, inflammatory heart disease, inflammatory kidney disease, IPF/UIP, iritis, keratitis, keratojuncuvitis sicca, Kussmaui disease or Kussmaul-Meier disease, Landry's paralysis, Langerhan's cell histiocytosis, livedo reticularis, macular degeneration, microscopic polyangiitis, morbus bechterev, motor neuron disorders, mucous membrane pemphigoid,

multiple organ failure, myasthenia gravis, myeiodysplastic syndrome, myocarditis, nerve root disorders, neuropathy, non-A non-B hepatitis, optic neuritis, osteolysis, ovarian cancer, paraarticular JRA, peripheral artery occlusive disease (PAOD), peripheral vascular disease (PYB), peripheral artery disease (PAD), phlebitis, polyarteritis nodosa (or periarteritis nodosa), polychondritis, polymyalgia rheumatica, poliosis, polyarticular JRA, polyendocrine deficiency syndrome, polymyositis, polymyalgia rheumatica (PMR), post-pump syndrome, primary Parkinsonism, prostate and rectal cancer and hematopoietic malignancies (leukemia and lymphoma), prostatitis, pure red cell aplasia, primary adrenal insufficiency, recurrent neuromyelitis optica, restenosis, rheumatic heart disease, sapho (synovitis, acne, pustulosis, hyperostosis, and osteitis), scleroderma, secondary amyloidosis, shock lung, scieritis, sciatica, secondary adrenal insufficiency, silicone associated connective tissue disease, Sneddon-Wilkinson dermatosis, spondylitis ankyiosans, Stevens-Johnson syndrome (SJS), systemic inflammatory response syndrome, temporal arteritis, toxoplasmic retinitis, toxic epidermal necrolysis, transverse myelitis, TRAPS (tumor necrosis factor receptor associated periodic syndrome), type 1 allergic react on, type II diabetes, urticaria, usual interstitial pneumonia (UIP), vasculitis, vernal conjunctivitis, viral retinitis, Vogt-Koyanagi-Harada syndrome (VKH syndrome), wet macular degeneration, wound healing, yersinia or salmonella associated arthropathy.

[00504] Various receptor for advanced glycation endproducts (RAGE) diseases may be treated with the pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the disease may be Amytropic Lateral Sclerosis, Brachial Plexus Injury, Brain Injury, including traumatic brain injury, Cerebral Palsy, Friedrich's Ataxia, Guiiiain Barre, Leukodystrophies, Multiple Sclerosis, Post Polio, Spina Bifida, Spinal Cord Injury, Spinal Muscle Atrophy, Spinal Tumors, Stroke, Transverse Myelitits, dementia, senile dementia, mild cognitive impairment, Alzheimer-related dementia, Huntington's chorea, tardive dyskinesia, hyperkinesias, manias, Morbus Parkinson, steel-Richard syndrome, Down's syndrome, myasthenia gravis, nerve trauma, vascular amyloidosis, cerebral hemorrhage I with amyloidosis, brain inflammation, Friedrich's ataxia, acute confusion disorder, amyotrophic lateral sclerosis, glaucoma, Alzheimer's disease, diabetic nephropathv, sepsis, rheumatoid arthritis and related inflammatory diseases.

[00505] Various neurite degenerative diseases may be treated with the pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the disease may be multiple sclerosis, Parkinson's disease, Alzheimer's disease, Tay-Sachs disease, Niemann-Pick disease, Gaucher's disease, Hurler's syndrome, Huntington's disease, amyotrophic

lateral sclerosis, idiopathic inflammatory demyeiinating diseases, vitamin B12 deficiency, central pontine myelinolysis, tabes dorsalis, transverse myelitis, Devic's disease, progressive multifocal ieukoencephalopathy, optic neuritis, traumatic injury to the CNS, an ischemic cerebral stroke, glaucoma, diabetic retinopathy, age-dependent macular degeneration, and a leukodystrophy.

[00506] Various neurological diseases may be treated with the pharmaceutical compositions, AAV particles, of the present invention. As a non-limiting example, the disease may be Amvotrophic Lateral Sclerosis, Brachial Plexus Injury, Bram injury, including traumatic brain injury, Cerebral Palsy, Guillain Barre, Leukodystrophies, Multiple Sclerosis, Post Polio, Spina Bifida, Spinal Cord Injury, Spinal Muscle Atrophy, Spinal Tumors, Stroke, Transverse Myelitis; dementia, senile dementia, mild cognitive impairment, Alzheimer-related dementia, Huntington's chorea, tardive dyskinesia, hyperkinesias, manias, Morbus Parkinson, steel-Richard syndrome, Down's syndrome, myasthenia gravis, nerve trauma, vascular amyloidosis, cerebral hemorrhage with amyloidosis, bram inflammation, acute confusion disorder, amyotrophic lateral sclerosis, glaucoma and Alzheimer's disease.

[00507] Various cancers may be treated with pharmaceutical compositions, AAV particles, of the present invention. As used herein, the term "cancer" refers to any of various malignant neoplasms characterized by the proliferation of anaplastic ceils that tend to invade surrounding tissue and metastasize to new body sites and also refers to the pathological condition characterized by such malignant neoplastic growths. Cancers may be tumors or hematological malignancies, and include but are not limited to, all types of lymphomas/1eukemias, carcinomas and sarcomas, such as those cancers or tumors found in the anus, bladder, bile duct, bone, brain, breast, cervix, colon/rectum, endometrium, esophagus, eye, gallbladder, head and neck, liver, kidney, larynx, lung, mediastinum (chest), mouth, ovaries, pancreas, penís, prostate, skin, small intestine, stomach, spinal marrow, tail bone, testicles, thyroid and uterus.

[00508] Types of carcinomas which may be treated with the AAV particles of the present invention include, but are not limited to, papillorna'carcmoma, choriocarcinoma, endodermal sinus tumor, teratoma, adenonia-adenoearcinoma, melanoma, fibroma, lipoma, leiomyoma, rhabdomyoma, mesothelioma, angioma, osteoma, chondroma, glioma, lynrphomaleukemia, squamous ceil carcinoma, small cell carcinoma, large cell undifferentiated carcinomas, basal cell carcinoma and sinonasal undifferentiated carcinoma.

[00509] Types of sarcomas which may be treated with the AAV particles of the present invention include, but are not limited to, soft tissue sarcoma such as alveolar soft part sarcoma, angiosarcoma, dermatofibrosarcoma, desmoid tumor, desmoplastic small round ceil tumor,

extraskeletai chondrosarcoma, extraskeletai osteosarcoma, fibrosarcoma, hemangiopericytoma, hemangiosarcoma, Kaposi's sarcoma, leiomyosarcoma, liposarcoma, lymphangiosarcoma, lymphosarcoma, malignant fibrous histiocytoma, neurofibrosarcoma, rhabdomyosarcoma, synovial sarcoma, and Asian's tumor, Ewing's sarcoma (primitive neuroectodermal tumor), malignant hemangioendothelioma, malignant sdiwannoma, osteosarcoma, and chondrosarcoma.

[00510] As a non-limiting example, the cancer which may be treated may be Acute granulocytic leukemia. Acute lymphocytic leukemia. Acute myelogenous leukemia, Adenocarcinoma, Adenosarcoma, Adrenal cancer. Adrenocortical carcinoma.. Anal cancer, Anaplastic astrocytoma, Angiosarcoma, Appendix cancer, Astrocytoma, Basal cell carcinoma, B-Cell lymphoma), Bile duct cancer, Bladder cancer. Bone cancer, Bowel cancer, Brain cancer. Brain stem glioma, Brain tumor, Breast cancer. Carcinoid tumors. Cervical cancer, Cholangiocarcinoraa, Chondrosarcoma, Chronic lymphocytic leukemia, Chronic myelogenous leukemia. Colon cancer, Colorectal cancer, Craniopharyngioma, Cutaneous lymphoma, Cutaneous melanoma, Diffuse astrocytoma, Ductal carcinoma in situ. Endometrial cancer, Ependymoma, Epithelioid sarcoma. Esophageal cancer, Ewing sarcoma, Exirahepatic bile duct cancer, Eye cancer, Fallopian tube cancer. Fibrosarcoma, Gallbladder cancer, Gastric cancer, Gastrointestinal cancer, Gastrointestinal carcinoid cancer, Gastrointestinal stromal tumors, General, Germ cell tumor, Glioblastoma multiforme, Glioma, Hairy cell leukemia. Head and neck cancer, Hemangioendothelioma, Hodgkin lymphoma, Hodgkin's disease, Hodgkin's lymphoma, Hypopharyngeal cancer, Infiltrating ductal carcinoma. Infiltrating lobular carcinoma, Inflammatory breast cancer, Intestinal Cancer, Intrahepatic bile duct cancer, invasive / infiltrating breast cancer, islet cell cancer, Jaw cancer, Kaposi sarcoma. Kidney cancer. Laryngeal cancer, Leiomyosarcoma, Leptomeningeal metastases, Leukemia, Lip cancer, Liposarcoma, Liver cancer, Lobular carcinoma in situ, Low-grade astrocytoma, Lung cancer. Lymph node cancer. Lymphoma, Male breast cancer, Medullary carcinoma, Medulloblastoma, Melanoma, Meningioma, Merkel cell carcinoma, Mesenchymal chondrosarcoma, Mesenchymous, Mesothelioma, Metastatic breast cancer. Metastatic melanoma, Metastatic squamous neck cancer, Mixed gliomas. Mouth cancer, Mucinous carcinoma, Mucosal melanoma, Multiple myeloma. Nasal cavity cancer. Nasopharyngeal cancer. Neck cancer. Neuroblastoma, Neuroendocrine tumors, Non-Hodgkin lymphoma, Non-Hodgkin's lymphoma, Non-small cell lung cancer. Oat cell cancer. Ocular cancer. Ocular melanoma, Oligodendroglioma, Oral cancer, Oral cavity cancer, Oropharyngeal cancer, Osteogenic sarcoma. Osteosarcoma, Ovarian cancer. Ovarian epithelial cancer. Ovarian germ cell tumor. Ovarian primary peritoneal carcinoma, Ovarian sex cord stromal tumor, Paget's disease,

Pancreatic cancer, Papillary carcinoma, Paranasal sinus cancer, Parathyroid cancer, Pelvic cancer, Penile cancer. Peripheral nerve cancer, Peritoneal cancer. Pharyngeal cancer, Pheochromocytoma, Pilocytic astrocytoma, Pineal region tumor, Pineoblastoma, Pituitary gland cancer, Primary central nervous system lymphoma, Prostate cancer, Rectal cancer. Renal cell cancer. Renal pelvis cancer. Rhabdomyosarcoma, Salivary gland cancer. Sarcoma, Sarcoma, bone. Sarcoma, soft tissue, Sarcoma, uterine. Sinus cancer. Skin cancer, Small cell lung cancer. Small intestine cancer, Soft tissue sarcoma, Spinal cancer, Spinal column cancer, Spinal cord cancer, Spinal tumor. Squamous cell carcinoma, Stomach cancer, Synovial sarcoma, T-cell lymphoma), Testicular cancer, Throat cancer. Thymoma/thymic carcinoma. Thyroid cancer, Tongue cancer. Tonsil cancer, Transitional cell cancer. Transitional cell cancer. Transitional cell cancer, Triple-negative breast cancer, Tubal cancer, Tubular carcinoma, Ureteral cancer, Ureteral cancer. Urethral cancer, Uterine adenocarcinoma. Uterine cancer, Uterine sarcoma. Vaginal cancer, and Vulvar cancer.

[00511] The AAV particles or pharmaceutical compositions of the present invention useful in preventing or treating HIV and AIDS may alternatively, or in combination, encode an antibody that targets a different infectious agent (e.g., an infectious agent that is not HIV-1 or 2). Non-limiting examples of other target antigens include any of the following, including fragments or variants thereof, adenoviruses, *Anaplasma phagocytophilium, Ascaris lumbricoides, Bacillus anthracis, Bacillus cereus, Bacteriodes* sp. Barman Forest virus, *Bartonella bacilliformis, Bartonella henselae, Bartonella quintana,* beta-toxin of *Clostridium perfringens, Bordetella pertussis. Bordetella parapertussis, Borrelia burgdorferi Borrelia miyamotoi, Borrelia recurrentis, Borrelia* sp., *Botulinum* toxin, *Brucella* sp., *Bwkho!deria pseudomallei,* California encephalitis virus, *Campylobacter, Candida albicans,* chikungunya virus, *Chlamydia psittaci, Chlamydia trachomatis, Clonorchis sinensis, Clostridium difficile bacteria, Clostridium tetani,* Colorado tick fever virus, *Corynebacterium diphtlieriae, Corynebacterium minutissimum, Coxiella burnetii,* coxsackie A, coxsackie B, Crimean-Congo hemorrhagic fever virus, cytomegalovirus, dengue virus. Eastern Equine encephalitis virus, Ebola viruses, echovirus. *Ehrlichia chqffeensis.. Ehrlichia equl, Ehrlichia* sp., *Entamoeba histolytica, Enterobacter sp., Enterococcus feacalis,* Enterovirus 71, Epstein-Barr virus (EBV), *Erysipelothrix rhusiopathiae, Escherichia coli,* Fiavivirus, *Fusobacterium necrophorum, Gardnerella vaginalis.* Group B *streptococcus, Haemophilus aegypthis, Haemophilus ducreyi, Haemophilus influenzae,* hantavirus, *Helicobacter pylori,* Hepatitis A, Hepatitis B, Hepatitis C, Hepatitis D, Hepatitis E, herpes simplex virus 1 and 2,, human herpes virus 6, human herpes Virus 8, human immunodeficiency virus 1 and 2, human T-cell leukemia viruses I and II, influenza viruses (A,

B, C), Jamestown Canyon virus, Japanese encephalitis antigenic, Japanese encephalitis virus, John Cunnmham vinisjuninvirus, Kaposi's Sarcoma-associated Herpes Virus (KSHV), *Klebsiella gramdomatis, Klebsiella sp.,* Kyasanur Forest Disease virus, La Crosse virus, Lassavirus, *Legionella pneumophila, Leptospira interrogans, Listeria monocytogenes,* lymphocytic choriomeningitis virus, lyssavirus, Machupovirus, Marburg virus, measles virus, MERS coronavirus (MERS-CoV), *Micrococcus sedenlarius, Mobiluncus sp., Molluscipoxvirus, Moraxella catarrhalis, Morbilli- Rubeola virus,* Munipsvirus, *Mycobacterium leprae. Mycobacterium tuberculosis, Mycobacterium ulcerans. Mycoplasma genitalium, Mycoplasma* sp, Nairovirus,, *Neisseria gonorrhoeae. Neisseria meningitidis, Nocardia,* Norwaik virus, norovims, Omsk hemorrhagic fever virus, papilloma virus, parainfluenza viruses 1-3, parapoxvirus, parvovirus B19, *Feptostreptococccus* sp., *Plasmodium* sp., polioviruses types I, II, and III, *Proteus* sp., *Pseudomonas aeruginosa, Pseudomonas pseudomallei. Pseudomonas* sp., rabies virus, respiratory syncytial virus, ricin toxin, *Rickettsia australis. Rickettsia conori, Rickettsia honei, Rickettsia prowazekii,* Ross River Virus, rotavirus, rubellavirus. Saint Louis encephalitis. *Salmonella Typhi, Sarcoptes scabiei,* SARS-associated coronavirus (SARS-CoV), *Serratia sp.,* Shiga toxin and Shiga-like toxin, *Shigella* sp.. Sin Norabre Virus, Snowshoe hare virus. *Staphylococcus aureus, Staphylococcus epidermidis, Slreptobacillus moniliformis, Streptoecoccus pneumoniae, Streptococcus agalactiae. Streptococcus agalactiae. Streptococcus* group A-H, *Streptococcus pneumoniae, Streptococcus pyogenes, Treponema pallidum subsp. Pallidum, Treponema pallidum var. carateum, Treponema pallidum var. endemicum, Tropheryma whippelii, Lfreaplasma urealyticum,* Varicella-Zoster virus, variola virus. *Vibrio cholerae,* West Nile virus, yellow fever virus. *Yersinia enteroc-olitica. Yersinia pestis,* Zika virus.

Diagnostic applications

[ΘΘ512] The AAV particles of the present invention may be used for diagnostic purposes or as diagnostic tools for any of the aforementioned diseases or disorders. As a non-limiting example, the AAV particles of the present invention or the antibodies encoded within the viral genome therein may be used as a biomarker for disease diagnosis. As a second non-limiting example, the AAV particles of the present invention or the antibodies encoded within the viral genome therein may be used for diagnostic imaging purposes, e.g., MRI, PET, CT or ultrasound.

Preventative applications

[00513] The AAV particles of the present invention or the antibodies encoded by the viral genome therein may be used to prevent disease or stabilize the progression of disease. In one embodiment, the AAV particies of the present invention are used to as a prophylactic to prevent a disease or disorder in the future. In one embodiment, the AAV particles of the present

invention are used to halt further progression of a disease or disorder. As a non-iimiting example, the AAV particles of the invention may be used in a manner similar to that of a vaccine.

Research applications

[00514] The AAV particles of the present invention or the antibodies encoded by the viral genome therein may also be used as research tools. The AAV particles of the invention may be used as in any research experiment, e.g., *in vivo* or *in vitro* experiments. In a non-limiting example, the AAV particles of the invention may be used in cultured cells. The cultured cells may be derived from any origin known to one with skill in the art, and may be as non-limiting examples, derived from a stable cell line, an animal model or a human patient or control subject. In a non-limiting example, the AAV particles of the invention may be used in *in vivo* experiments in animal models (i.e., mouse, rat, rabbit, dog, cat, non-human primate, guinea pig, ferret, c-eJegans, drosophila, zebrafish, or any other animal used for research purposes, known in the art). In another non-iimiting example, the AAV particles of the invention may be used in human research experiments or human clinical trials.

Combination applications

[00515] The AAV particles of the invention may be used as a combination therapy with any other therapeutic molecule known in the art. The therapeutic molecule may be approved by the US Food and Drug Administration or may be in clinical trial or at the preclinical research stage. The therapeutic molecule may utilize any therapeiuic modality known in the art, with non-iimiting examples including gene silencing or interference (i.e., miRNA, siRNA, RNAi, shRNA), gene editing (i.e., TALEN, CRISPR/Cas9 systems, zinc finger nucleases), and gene, protein or enzyme replacement.

Therapeutic applications: Infectious Diseases

[00516] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and-'or treat infectious disease. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Tables 21-42 (SEQ ID NO: 4138-9220).

[00517] The methods, components and compositions of the present invention may be used to diagnose, prevent, treat and/or manage infectious diseases. Infectious diseases, also known as transmissible diseases or communicable diseases, are caused by invasion and multiplication of agents in the body. Infection agents are species typically not present within the body and may be, but are not limited to, viruses, bacteria, prions, nematodes, fungus, parasites or arthropods. Additionally, an infection or symptoms associated with an infection may be caused by one or

more toxins produced by such agents. Humans, and other mammals, react to infections with an innate immune system response, often involving an inflammation. The illnesses and symptoms involved with infections vary according to the infectious agent. Many infections may be subclinical without presenting any definite or observable symptoms, whereas some infections cause severe symptoms, require hospitalization or may be life-threatening. Some infections are localized, whereas some may overcome the body through blood circulation or lymphatic vessels. Some infections have long-term effects on wellbeing of infected individuals.

[00518] Infectious agents may be transmitted to humans via different routes. For example, infection agents may be transmitted by direct contact with an infected human, an infected animal, or an infected surface. Infections may be transmitted by direct contact with bodily fluids of an infected human or an animal, e.g. blood, saliva, sweat, tears, mucus, female ejaculate, semen, vomit or urine. For example, infection may be transmitted by a fecal-oral route, referring to an infected person shedding the virus in fecal particles which then enters to person's mouth causing infection. The fecal-oral route is especially common transmission route in environments with poor sanitation and hygiene. Non-limiting examples of agents transmitted by the fecal-oral route include bacteria, e.g. shigella, *Salmonella typhii* and *Vibrio Cholerae,* virus, e.g. norovirus, rotavirus, enteroviruses, and hepatitis A, fungi, e.g. *Entamadeba histolytica,* parasites, tape worms, transmitted by contaminated food or beverage, leading to food poisoning or gastroenteritis. Infections may be transmitted by a respiratory route, referring to agents that are spread through the air. Typical examples include agents spread as small droplets of liquid or as aerosols, e.g. respiratory droplets expelled from the mouth and nose while coughing and sneezing. Typical examples of respiratory transmitted diseases include the convmon cold mostly implicated to rhinoviruses, influenza caused by influenza viruses, respiratory tract infections caused by e.g. respiratory syncytial virus (RSV). infections may be transmitted by a sexual transmission route. Examples of common sexually transmitted infections include e.g. human immunodeficiency virus (HIV) causing acquired immune deficiency syndrome (AIDS), chlamydia caused by *Neisseria gonorrhoeae* bacteria, fungal infection Candidiasis caused by Candida yeast, and Herpes Simplex disease caused by herpes simplex virus. Infections may be transmitted by an oral transmission route, e.g. by kissing or sharing a drinking glass. A common infection transmitted by oral transmission is an infectious mononucleosis caused by Epstein-Barr virus. Infections may be transmitted by a vertical transmission, also known as "mother-to-child transmission," from mother to an embryo, fetus or infant during pregnancy or childbirth. Examples of infection agents that may be transmitted vertically include HIV, chlamydia, rubella. *Toxoplasma gondii,* and herpes simplex virus. Infections may be transmitted by an iatrogenic

route, referring to a transmission by medical procedures such as injection (contaminated reused needles and syringes), or transplantation of infected material, blood transfusions, or infection occurring during surgery. For example. methicillin -resistant *Staphylococcus aureus* (MRSA). which may cause several severe infections, may be transmitted via iatrogenic route during surgery. infections may also be transmitted by vector-bome transmission, where a vector may be an organism transferring the infection agents from one host to another. Such vectors may be triatomine bugs, e.g. trypaiiosomes, parasites, animals, arthropods including e.g. mosquiios, flies, lice, flees, tick and mites or humans. Non-limiting examples of mosquito-borne infections include Dengue fever, West Nile virus related infections, Yellow fever and Chikungunya fever. Non-limiting examples of parasite-borne diseases include malaria, Human African trypanosomiasis and Lyme disease. Non-limiting examples of diseases spread by humans or mammals include HIV, Ebola hemorrhagic fever and Marburg fever.

[00519] Traditionally infectious diseases are treated with medications and/or good supportive care. Medical prevention, treatment and/or management of bacterial infections may include administration of antibiotics. Antibiotics may inhibit the colonization of bacteria or kill the bacteria. Antibiotics include e.g. penicillins, cephalosporins, macrolides, fluoroquinolones, sulfonamides, tetracyclines, and aminoglycosides. Antibiotics may be specific to a certain bacteria or act against broad spectrum of bacteria. Some types of bacteria are especially susceptible to antibiotics, whereas some bacteria are more resistant. Development of bacterial strain mutations that are resistant to antibiotics is an increasing concern. Methicillin-resistant Staphylococcus aureus (MRSA), vancomycin-resistant Enterococcus (VRE), multi-drug-resistant Mycobacterium tuberculosis (MDR-TB) and *Klebsiella pneumoniae carbapenemase*- producing bacteria (KPC) are examples of bacteria that are resistant to most general antibiotics. Due to the emerging resistance, unnecessary administration and overdosing of antibiotics should be avoided. Medical prevention, treatment and/or management of viral infections may include administration of antiviral medications. Antiviral medications may be specific to a certain bacteria or act against a broad spectrum of viruses. Currently antiviral medications are available for e.g. HIV, influenza, hepatitis B and C. Medical prevention, treatment and/or management of viral infections may include administration of antifungal medication. .Antifungal medication kills or prevents the growth of fungi. Types of antifungal medications include e.g. imidazoles, triazoles and thiazoies, allylamines, and echinocandins. Development of antifungal medication capable of targeting fungal cells without affecting human cells is a challenge due to the similarities of human and fungal cell on the molecular level. Typically, medical treatment is combined with good supportive care, which includes provision of fluids, bed rest, medication to

relieve pain and lower fever, supportive alternative medicine such as vitamins, antioxidants and other supplements important for welibeing of patients.

[ΘΘ520] Antibody therapies for infectious diseases have also been developed. Examples of commercial therapeutic antibodies include raxibacumab (deveioped by Cambridge Antibody Technology and Human Genome Sciences) which is an antibody for the prophylaxis and treatment of inhaled anthrax, SHlGAMAB ™ (deveioped by Bellus Health Inc.) is a monoclonal antibody for treatment of Shiga toxin induced hemolytic uremic syndrome, and actoxumab and bez!otoxumab (developed by Medarex Inc. and the University of Massachusetts Medical School) are commercial human monoclonal antibodies targeting C. difficile toxin A and toxin B, respectively.

[00521] Infectious diseases and/or infection related diseases, disorders, and/or conditions that may be treated by methods, components and compositions of the present invention include, but are not limited to, 14-day measles, 5-day fever, acne, acquired immunodeficiency syndrome (AIDS), acrodermatitis chronica atrophicans (ACA), acute hemorrhagic conjunctivitis, acute hemorrhagic cystitis, acute rhinosinusitis, adult T-ce!l ieukemia-lymphoma (ATLL), African sleeping sickness, alveolar hydatid, amebiasis, amebic meningoencephalitis, anaplasmosis, anthrax, arbovirai, ascariasis, aseptic meningitis. Athlete's foot, Australian tick typhus, avian influenza, babesiosis, bacillary angiomatosis, bacterial meningitis, bacterial vaginosis, balanitis, balantidiasis. Bang's disease, Barmah Forest virus, bartoneliosis, hat lyssavirus. Bay sore, Baylisascaris, beaver fever, beef tapeworm, bejel, biphasic meningoencephalitis, black bane, black death, black piedra, Blackwaier fever, blastomycosis, blennorrhea of the newborn, blepharitis, boils, Bomholm disease, borrelia miyamotoi disease, botulism, boutonneuse fever, Brazilian purpuric fever, break bone fever, brill, bronchiolitis, bronchitis, brucellosis, bubonic, bubonic plague, bullous impetigo, burkholderia mallei, burkholderia pseudomallei, burly ulcers mycoburuli ulcers, Busse-Buschke disease, California group encephalitis, campyiobacteriosis, candidiasis, canefieid fever, canicoia fever, capillar!asis, carate, carhapenem-resistant enterobacteriaceae (CRE), Carrion's disease, cat scratch fever, cave disease, central Asian hemorrhagic fever. Central European tick, cervical cancer, Chagas disease, cancroid, Chicago disease, chickenpox, Chiciero's ulcer, chikunguny a fever, chlamydial, cholera, chromoblastomy cosis, ciguatera, clap, clonorchiasis, Clostridium difficile, Clostridium perfringens, coccidioidomycosis fungal, coenurosis, Colorado tick fever, condyloma accuminata, condyloma lata, Congo fever, Congo hemorrhagic fever virus, conjunctivitis, cowpox, crabs, Crimean disease, croup, crypto, cryptococcosis, cryptosporidiosis, cutaneous larval migrans, cyclosporiasis, cystic hydatid, cysticercosis, cystitis, Czechoslovak tick, d68 (EV~d68),

dacryocystitis, dandy fever, darling's disease, deer fly fever, dengue fever types 1, 2, 3, and 4, desert rheumatism, devil's grip, diphasic milk fever, diphtheria, disseminated intravascular coagulation, dog tapeworm, donovanosis, dracontiasis, dracuncuiosis, duke's disease, dum dum disease, Durand-Nicholas-Favre disease, dwarf tapeworm, E. coli, eastern equine encephalitis, Ebola hemorrhagic fever, Ebola virus disease (EVD), ectothrix, ehrlichiosis, encephalitis, endemic relapsing fever, endemic syphilis, endophthalmitis, endothrix. enterobiasis, enterotoxin - B poisoning (staph food poisoning), enterovirus, epidemic keratoconjunctivitis, epidemic relapsing fever, epidemic typhus, epiglottitis, epsilon toxin, erysipelis, erysipeloid, erysipelothricosis, erythema chronicura migrans, erythema infectiosum, erythema marginatum, erythema multiforme, erythema nodosum, erythema nodosum leprosura, erythrasma, espundia, eumycotic mycetoma, European blastomycosis, exanthem subitum, eyeworm, Far-Eastern tick, fascioiiasis, fievre boutormeuse, fifth disease, Filatow-Dukes' disease, fish tapeworm, Fitz-Hugh-Curtis syndrome - perihepatitis, flinders island spotted fever, flu, folliculitis, four comers disease, frambesia, francis disease, furunculosis, gas gangrene, gastroenteritis, genital herpes, genital warts, German measles, Gerstmann-Straussler-Scheinker (GSS), giardiasis, Gilchrist's disease, gingivitis, gingivostomatitis, glanders, glandular fever, gnathostomiasis, gonococcal, gonorrhea, granuloma inguinale, guinea worm, Haemophilus influenza disease, hamburger disease, Hansen's disease, Hantaan disease, Hantaan-Korean hemorrhagic fever, haniavims pulmonary syndrome (HPS), hard chancre, hard measles, Haverhill fever, head and body lice, heartland fever, helicobacterosis, hemolytic uremic syndrome (HUS), hepatitis A, hepatitis B, hepatitis C, hepatitis D, hepatitis E, herpaiigina, herpes- genital, herpes labialis, herpes- neonatal, hidradenitis, histoplasmosis, histoplasmosis, his-wemer disease, hiv, hookworm s, hordeola, HTLV- associated myelopathy (HAM), human granulocytic ehrlichiosis, human monocytic ehrlichiosis, human papiilomarivus (HPV), human pulmonary syndrome, human pulmonary syndrome (HPS), human T-cell lymphotropic virus (HTLV), hydatid cyst, hydrophobia, impetigo, including congenital, inclusion conjunctivitis, infantile diarrhea, infectious mononucleosis, infectious myocarditis, infectious pericarditis, influenza, isosporiasis, Israeli spotted fever, Japanese encephalitis, jock itch, jorge lobo disease, jungle yellow fever, Junin Argentinian hemorrhagic fever, kala azar, Kaposi's sarcoma, keloidal blastomycosis, keratoconjunctivitis, kuru, Kyasarmr forest disease, lacrosse encephalitis, lassa hemorrhagic fever, legionellosis, legionnaires disease, legionnaire's pneumonia, Lemierre's syndrome, lemming fever, leprosy, ieptospirosis, listeria, listeriosis, liver fluke, iobo's mycosis, lockjaw, lockjaw, loiasis, louping ill, Ludwig's angina, lung fluke, Lyme disease, lymphogranuloma venereum (LGV), Machupo Bolivian hemorrhagic fever. Madura foot, mal del pinto, malaria,

malignant pustule, Malta fever, Marburg hemorrhagic fever, masters disease, maternal sepsis, measles, Mediterranean spotted fever, melioidosis, meningitis, meningococcal disease, Middle East Respiratory Syndrome (MERS), methiciilin-resistant staphylococcus aureus (MRSA), milker's nodule, molluscum contagiosity, moniliasis, monkeypox, mononucleosis, mononucleosis-like syndrome, Montezuma's revenge, morbilli, mucormycosis, multiple organ dysfunction syndrome (MODS), multiple-system atrophy (MSA), mumps, murine typhus, Murray Valley encephalitis (MVE), mycoburuli ulcers, mycotic vulvovaginitis, myositis, Nanukayami fever, necrotizing fasciitis, necrotizing fasciitis- type 1, necrotizing fasciitis- type 2, negishi, new world spotted fever, nocardiosis, nongonococcal urethritis, non-polio enterovirus, norovirus. North American blastomycosis. North Asian tick typhus. Norwalk virus, Norwegian itch, O'hara disease, Omsk hemorrhagic fever, onchoceriasis, onychomycosis, opisthorchiasis, opthalmia neonatorium, oral hairy leukoplakia, orf, oriental sore, oriental spotted fever, ornithosis, Oroya fever, otitis externa, otitis media, pannus, paracoccidioidomycosis, paragonimiasis, parainfectious, paralytic shellfish poisoning, paronychia, parotitis, parrot fever, pediculosis, peiiosis hepatica, peivic inflammatory disease, pertussis, phaeohvphomycosis, pharingoconjunctival fever, piedra, pighel, pink eye conjunctivitis, pinta, pinworm, pitted keratolysis, pityriasis versicolor, plague, pleurodynia, pneumococcal disease, Pneumocystis pneumonia, pneumocystosis, pneumonia, polio, poliomyelitis, polycystic hydatid, Pontiac fever, pork tapeworm, Posada-Wernicke disease, postanginal septicemia, Powassan, progressive multifocal leukencephaiopathy (PML), progressive rubella panencephalitis, prostatitis, pseudomembranous colitis, psittacosis, puerperal fever, pustular rash diseases, pyelonephritis, pylephlebitis, q-fever, quinsy, quintana fever, rabbit fever, rabies, racoon roundworm, rat bite fever, rat tapeworm, Reiter syndrome, relapsing fever, respiratory syncytial virus (RSV), rheumatic fever, rhodotorulosis. ricin poisoning, rickettsialpox, rickettsiosis, Rift valley fever, ringworm, Ritter's disease, river blindness, rocky mountain spotted fever, rose handler's disease, rose rash of infants, roseola, Ross river fever, rotavirus, roundworms, rubella, rubeola, Russian spring, salmonellosis gastroenteritis, San Joaquin valley fever, Sao Paulo encephalitis. Sao Paulo fever, scabies infestation, scalded skin syndrome, scalded skin syndrome, scarlatina, scarlet fever, schistosomiasis, scombroid, scrub typhus, sennetsu fever, sepsis, septic shock, severe acute respiratory syndrome, severe acute respiratory syndrome (SARS), shiga toxigenic Escherichia coli, shigella, shigellosis gastroenteritis, shinbone fever, shingles, shipping fever, Siberian tick typhus, sinusitis, sixth disease, slapped cheek disease, sleeping sickness, small pox, smallpox, snail fever, soft chancre, southern tick associated rash illness, sparganosis, Spe! tinker's disease, sporadic typhus, sporotrichosis, spotted fever, spring, St. Louis encephalitis,

staphylococcal food poisoning, staphylococcal, strep. throat, streptococcal disease, strepiococcal toxic-shock syndrome, strongyloiciasis, stye, subacute sclerosing panencephalitis (SSAPE), sudden acute respiratory syndrome, sudden rash, swimmer's ear, swimmer's itch, swimming pool conjunctivitis, sylvatic yellow fever, syphilis, systemic inflammatory response syndrome (SIRS), tabes dorsalis, taeniasis, taiga encephalitis, tanner's disease, tapeworm s, temporal lobe encephalitis, tertiary syphilis, tetani, tetanus, threadworm s, thrush, tick, tick typhus, tinea barbae, tinea capitis, tinea corporis, tinea cruris, tinea manuum, tinea nigra, Tinea pedis, tinea unguium, tinea versicolor, torulopsosis, torulosis, toxic shock syndrome, toxoplasmosis, transmissible spongioforrn, traveler's diarrhea, trench fever 5, trichinellosis, trichomoniasis, trichomycosis axillaris, trichuriasis, tropical spastic paraparesis (TSP), trypanosomiasis, tuberculosis (TB), tularemia, typhoid fever, typhus fever, ulcus molle, undulant fever, urban yellow fever, urethritis, vaginitis, vaginosis, valley fever, vancomycin intermediate (VISA), vancomycin resistant (VRSA), varbuncle, varicella, variola, varrion's disease, Venezuelan equine encephalitis. Verruga peruana, vibrio, vibrio cholerae, vibriosis, vmcent's disease or trench mouth, viral conjunctivitis, viral meningitis, viral meningoencephalitis, viral rash, visceral larval migrans, vomito negro, vulvovaginitis, warts, Waterhouse, Weil's disease. West Nile fever. Western equine encephalitis, Whipple's disease, whipworm, white piedra. whitlow, Whitmore's disease, whooping cough, winter diarrhea, wolhynia fever, wool sorters' disease, yaws, yellow fever, yersinosis, zahorsky's disease, zika virus disease, zoster, zygomycosis, acute bacterial rhmosini!sitis, lobomycosts, and/or any other infectious diseases, disorders or conditions.

John Cunningham Virus (JCV)

[00522] John Cunningham Virus is a common human poiyomavirus. The transmission route of JCV is unknown. The virus is suspected to be spread by contaminated water and may be obtained through tonsils or by the gastrointestinal tract. 70-90 % of humans are estimated to be infected by the virus, and for normal healthy individuals the infection is asymptomatic. However, for patients with weakened immune system. JCV may lead to Progressive multifocal leukoencephalopathy (PML). PML is a condition characterized by multifocal progressive damage or inflammation of the white matter of the brain. The symptoms include clumsiness, progressive weakness and changes in visual, speech and personality. PLM has a mortality rate of 30-50 % and patients who survive the disease are left with severe neurological disabilities. PML occurs in patients with a severe immunodeficiency, most commonly in patients with HIV/AIDS. As many as 5 % of HIV/AIDS patients are affected by PML. Individuals with other autoimmune conditions such as multiple sclerosis, rheumatoid arthritis, and systemic lupus erythematosus are also at risk, as well as individuals going through immunosuppressive therapy for cancer, e.g.

lymphoma orHodgkin's disease, or organ transplant. PML associated with immunosuppressive therapy is an increasing concern. For example, commercial antibody nataHzumab (TYSABRI®, developed by Biogen Idee) for treatment of multiple sclerosis increases susceptibility to PML. Other drugs associated with increased risk of PML include Rituxiniab (RiTUXAN ®, developed by IDEC Pharmaceuticals), Efalizumab (RAPT1VA® developed by Genentech and XOMA) and Mycophenolate mofetil (CELLCEPT®, developed by Genentech).

[00523] JCV is a nonenveioped, T=7 icosahedral virus with a closed circular, double-stranded DNA genome. The major capsid component is the viral protein VPl is made of 72 pentamers formed by VPl monomers linked through the C terminal end. VPl starts the infection by binding to the receptor target cells. After initial infection, typically occurring in childhood or adolescence, the virus stays quiescent in the kidneys and the lymphoid organs. In healthy individuals, the virus may replicate in kidney without causing any symptoms. However, in patients with weakened immune system, JCV may cross the blood-brain barrier into the central nervous system causing PML.

[00524] As of today, there is no known cure for PML. Current therapies focus on reversing the immune deficiency to slow down or stop the progress of the disease. There remains a need for therapies neutralizing JVC for prevention, management and treatment of JCV infection and PML Goidmann et al. demonstrated that neutralizing activity with JCV VPl protein in sera of a rabbit (see Goidmann C. et al., 1999, *J Virol.;* 73(5): 4465-4469). Therapies based on neutralizing JCV antibodies could be applied for treatment, management and/or prevention of PML. Recently, immunological approaches have been under investigation and neutralizing antibodies binding to JC virus, especially targeting the VPl protein, have been developed e.g. as described in US Patent Publication US2015/0191530, US2015/0056188 and US2015/0050271, the contents of each of which are incorporated herein by reference in their entirety. Such antibodies may cause reduction of JCV replication, proliferation or mfeetivify. Antibodies may bind to a conformational epitope of JCV VPl protein or to the sialic acid binding pocket of VPl protein of JCV.

[00525] In some embodiments, methods of the present invention may be used to prevent, manage and/or treat JCV infection and/or PML.

[00526] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat JCV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 26 (SEQ ID NO: 6802-6865).

Influenza virus

[00527] Influenza -viruses cause a common respiratory infection called influenza (flu). Influenza viruses are categorized into three main groups, virus A, B and C. Influenza viruses are negative-sense, single-stranded, segmented RNA viruses. Influenza A contains two proteins on the surface of the viral envelope: hemagglutinin (H), which is a protein responsible for red blood cell agglutination and neuraminidase (N), which is an enzyme cleaving the glycosidic bonds of neuraminic acid. Influenza A mutates at a faster rate than types B and C. Several serotypes of H and subtypes of N have been identified. Influenza Type B, similarly to Type A, contains H and N protein. Type C influenza virus is a single stranded RNA virus with glycoprotein called hemagglutinin-esterase fusion. Influenza strains vary according to geographical presentation.

[ΘΘ528] Influenza in general is a highly contagious disease and may be transmitted by the respirator}' route. influenza symptoms include e.g. high fever, runny nose, headache, sore throat, muscle pain, cough and occasionally nausea and vomiting. Influenza may lead to other complications such as pneumonia or sinus infections. influenza may be dangerous to young children, the elderly, pregnant women and individuals with chrome medical conditions or weakened immune system. According to Centers for Disease Control and Prevention (CDC), the estimated annual number of flu-associated deaths in the United States ranges between 3000 and 49, 000, depending on the severity of the seasonal variations.

[ΘΘ529] Influenza may be treated with good supportive care and antiviral medication. Antiviral medications include newaminidase inhibitors, e.g. ose!tamivir and zanamivir and M2 protein inhibitors. However, some strains of influenza appear to be resistant to these antiviral medications. Seasonal vaccinations to influenza are very efficient in prevention of the disease and are recommended annually.

[00530] There remains a need for prevention and treatment therapies for influenza, especially for those providing long lasting and broad neutralization. Therapeutic antibodies against influenza viruses have been developed. In general, antibody responses to different subtypes and serotypes of influenza A, B and C are unique. Some therapeutic antibodies are specific to an antibody type, whereas some have a broad coverage. Navrvumab (developed by Celltrion, Inc.) taught in US Patent application US20140234336, firivumab (developed by Celltrion, Inc.) taught in US Patent application US20 130004505 and diridavumah (developed by Jansen Biotech, Crucell and Johnson&Johnson) taught in international Patent application WO/2008/028946 are examples of therapeutically antibodies against influenza A hemagglutinin HA.

[00531] AAV particles and methods of using the AAV particles described in the present invention may he used to prevent, manage and/or treat influenza. As a non-iimiting example, the

AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 21 (SEQ ID NO: 4138-5222).

Hepatitis

[00532] Hepatitis is an inflammation of the liver. Hepatitis may be caused by an infection of hepatitis viruses A, B, C, D or E. In some cases, hepatitis may be asymptotic. A typical symptom of hepatitis is jaundice, characterized by yellowing of the skin, mucous membrane and conjunctiva. Other symptoms include loss of appetite, diarrhea, nausea and fever. Hepatitis may lead to a liver failure. Acute form of hepatitis is healed within six months of infection. The inflammation may also progress to a chronic hepatitis, which may lead to liver complications such as fibrosis, cirrhosis or hepatocellular carcinoma. There is no specific treatment for hepatitis. Typically, acute hepatitis is treated with good supportive care, including good nutritional balance, fluid and rest. Chronic hepatitis may be treated with antiviral drugs. Hepatitis may be prevented by vaccinations.

[00533] Hepatitis A (HAY) virus belongs to the family of *Picornaviridae.* HAV is encapsidated in an icosahedrai structure formed by 60 copies of three viral structural proteins (VP1 , VP2 and VP3), (see e.g. Kim et al. 2004. *Virology.*;318(2):598-607, and references therein). HAV is spread by the fecal-oral-route. Typical transmission is through contaminated food or drink or in contact with an infected individual. Improperly cooked shellfish is a common source of HAV. Hepatitis A is more abundant in developing countries with poor sanitary conditions. According to the World Health Organization (WHO), an estimated 1.4 million people are infected by HAV every year.

[00534] Vaccines for prevention of HAV infection exists and are recommended to be administered to children under 1 year of age by CDC. As of today, there is no specific treatment for HAV infection. The treatment includes supportive therapy and may last for weeks or even months. There remains a need for treatment therapies for HAV. Antibodies for prevention and/or treatment of HAV have been developed. For example, US Patent IJS763476, International Publication WO201 1114353 and Kim et al in *Virology.* 2004 Jan 20;318{2):598~607, the contents of each of which are incorporated herein by reference in their entirety, teach neutralizing antibodies targeting HAV antigens.

[00535] Hepatitis B (HBV) belongs to the family of *Orthohepadnaviridae.* HBV comprises a 3.2 kb-partialiy double-stranded circular DNA genome. HBV virus may be transmitted via the sexual transmission route, vertical transmission at birth, iatrogenic route (e.g. blood transfusions, contaminated reused needles and synnges). as well as via exposure to certain body fluids of an infected individual. According to the WHO. an estimated 240 million people are chronically

infected with hepatitis B annually, and more than 780 000 people die to associated complications.

[00536] HBV may be prevented by vaccination. The WHO recommends vaccination for all infants, as well as for adults living in increased risk of the infection. HBV infection may be treated with antiviral medications, e.g. tenofovir and entecavir. The medication does not cure the disease but suppresses the replication of the virus. individuals with chronic hepatitis B infection are administered antiviral medications for life. There remains a need for therapies providing long lasting management and/or cure for HBV infection. Antibodies for prevention and/or treatment of HBV infection are described e.g. in US Patent publication US20120308580 and International publication WO20131 65972, the contents of each of which are herein incorporated by their reference in their entirety.

[00537] Hepatitis C (HCV) belongs to the family of *Flaviviridae.* HCV is a positive-sense single-stranded RNA virus with an open reading frame with 9600 nucleotide bases. HCV is most commonly transmitted by the sexual transmission route or iatrogenic route. Hepatitis C may be transmitted also via the vertical route, though uncommon. According to WHO, 130-150 million people have a chronic HCV infection and approximately half a million people die from complications associated with HCV annually.

[00538] As of today, there is no vaccine for HCV infection. Traditional treatment of hepatitis C is based on antiviral medication therapy with e.g. ribavirin and interferon. More recently, direct antiviral agents (DAA) have been developed to treat hepatitis C infections. However, there remains a need for efficient prevention and treatment therapies for HCV infection.

[00539] Hepatitis D (HDV) is a small spherical enveloped RNA virus belonging to the genus of deltaviruses. HDV infection may only replicate in the presence of a HBV virus and therefore HDV infection has a dependency on HBV. HDV virus may be transmitted as coinfection with HBV or be superimposed on chronic HBV or HBV carrier state. HDV may be transmitted similarly to HBV, e.g. via the sexual transmission route, vertical transmission at birth, iatrogenic route, as well as via exposure to certain body fluids of an infected individual. Treatment and vaccination against HBV may be applied against HDV, and there remains a need for therapies to cure both infections.

[00540] Hepatitis E (HEV) is a linear, monopartite, single-stranded RNA virus belonging to the family *of Hepeviridae.* HEV may be transmitted via the fecal-oral route due to contaminated food or beverage, the iatrogenic route (e.g. blood transfusions, contaminated reused needles and syringes) or the vertical transmission route during pregnancy. Contaminated drinking water is the most common source of infection. Improperly cooked shellfish are a common source of HEV.

The disease is present worldwide but is more abundant in East and South Asia, and especially in environments with poor sanitation and hygiene. According to WHO, an estimated 20 million HEV infections occur annually leading to 56 600 death associated with HEV complications.

[00541] There is no specific treatment for HEV. The disease is typically cured with good supportive care. As of today, vaccinations against HEV are not globally available, though development in the field has been done. There remains a need for prevention and treatment therapies for HEV infection. Antibodies for prevention and treatment of HEV have been developed. For example, neutralizing antibodies targeting HEV have been taught in US Patent US 7148323, Tang et al. 2011, *Proc. Natl. Acad. Sci.* U.S.A. 108 (25), 10266-10271 and Gu et al. 2015, *Cell Res.* 25 (5), 604-620, the contents of each of which are incorporated herein by reference in their entirety.

[00542] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by HAV, HBV, HCV, HDV and/or HEV.

[00543] AAV particles and methods of using the AAV panicles described in the present invention may be used to prevent, manage and/or treat HAV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 6 (SEQ ID NO: 3197-3237).

[00544] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HBV. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described In Table 23 (SEQ ID NO: 6311-6627).

[00545] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HDV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 23 (SEQ ID NO: 6311-6627).

[00546] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HEV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 6 (SEQ ID NO: 3197-3237).

Respiratory syncytial virus (RSV)

[00547] Respiratoiy syncytial virus (RSV) is a smgie-stranded RNA virus belonging to the family of *Paramyxoviridae.* The RSV RNA is contained in a nucleocapsid made of 11 proteins and covered with a lipid envelope (see. e.g. Piedimonte, 2015, *Cleve Clin J Med.* 82{\1 Suppl

1): S13-8, and references therein). RSV attaches to the epithelial cells of the host airway cells with the surface glycoproteins G and F and merges the viral envelope to the membranes of adjacent cells. G and F glycoproteins are the principal antigens exposed to the host immune system.

[00548] Respiratory syncytial virus (RSV) causes infections of the respiratory tract including the lungs and breathing passages. RSV is transmitted through the respiratory transmission route, in direct contact with nasal or oral secretions of infected individuals, or indirectly e.g. by touching a contaminated surface. The symptoms include a runny nose, decrease of appetite, coughing, sneezing, fever and wheezing. The infection may progress into a pneumonia or bronchiolitis. Additionally, RSV infection may have a role in triggering asthma attacks and in the inception of asthma for individuals with a family history of asthma. In healthy adults, RSV infection is typically mild and does not require hospitalization. However, the infection may be dangerous for young children and infants, and for individuals with a weakened immune system. According to the CDC, almost all children under 3 years of age will acquire an RSV infection and up to 2 % of cases require hospitalization. RSV infection the most common cause for bronchiolitis and pneumonia in children younger than 1-year-old.

[00549] As of today, there is no specific medical treatment for RSV infection on the market and typically the infection is treated with good supportive care. There remains a need for prevention and treatment therapies for RSV infections and associated complications. Antibodies for treatment and prevention of RSV infection have been developed. For example, palivizumab (developed by Medlmmune) taught in US Patent US 8153133, the contents of which are incorporated herein by reference in their entirety, is a nearly human monoclonal antibody targeting the RSV F glycoprotein. Palivizumab is used for passive immunity for infants at risk for severe infection, including children with hemodynamicaliy significant congenital heart defects, profound immunodeficiency and pulmonary or neuromuscular pathologies impairing airway clearance.

[00550] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by RSV.

[00551] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat RSV. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 22 (SEQ ID NO: 5223-6310).

Herpes simplex virus I and 2

[00552] Herpes simplex viruses 1 and 2 (HSV1 and HSV2), also known as human herpes virus 1 and 2 (HHV-1 and HHV-2), belong to the family of *Herpesviridae.* Herpesviruses in general, consist of an icosahedral capsid surrounded by a membrane envelope. The capsid contains the viral double stranded DNA. The capsid is surrounded by an amorphous tegument of 30 viral proteins. The virion is enveloped by lipids with multiple viral glycoproteins and cellular proteins (see, e.g. McAllister and Schieiss, 2014, *Expert Rev Vaccines:,* 13(11): 1349-1360, and references therein).

[00553] HSV1 and HSV2 cause an infection known as herpes, which is characterized by blisters in the skin, or mucous membranes of the mouth, lips, also known as "cold sores'", or genitals. Typically, the symptoms are mild or asymptomatic. However, HSV1 and HSV2 are neurotropic and neuromvasive viruses persisting in the body by becoming latent, and sustain in the cell bodies of neurons. The infection is lifelong with outbreaks, or sporadic episodes of viral reactivation, when the virus in the nerve ceils become active causing new blistering. The infection may be dangerous to individuals with weakened immune system. Neonatal herpes of infants may be fatal. Occasionally HSVl infections may lead to encephalitis or keratitis. HSVl and HSV2 are transmitted by contact with an infected area during reactivations of the virus. HSVl is mainly transmitted by oral-to-oral contact, skin contact or the sexual transmission route. HSVl may also be transmitted vertically during birth. HSV2 is transmitted via the sexual transmission route and is one of the most common sexually transmitted infections. According to the WHO, an estimated 67 % of world's population aged under 50 years has an HSV-1 infection. An estimated 11% of world's population aged 15-49 years has an HSV2 infection.

[00554] As of today, there is no vaccination for prevention of HSV1 and HSV2 infections on the market. HSVl and HSV2 infections may be treated with antiviral medications, such as acyclovir, famciclovir and valacyclovir. Antiviral medications do not cure the infection, but reduce the seventy and frequency of symptoms. There remains a therapy for prevention and cure for these infections. Antibodies for prevention, treatment and management of HSVl and HSV2, targeting the viral glycoproteins, have been developed, as described e.g. in US Patent US8431118, US Patent US5646041, Haynes US Patent Publication US2014/0302062, the contents of each of which are incorporated herein by reference in their entirety.

[00555] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by HSVl and HSV2.

[00556] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HSV. As a non-limiting example, the

AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 24 (SEQ ID NO: 6628-6736).

Human Cytomegalovirus

[00557] Human Cytomegalovirus (HCMV) also known as human herpesvirus 5 (HHV-5) belongs to the family of *Herpesviridae,* a sub-family of *Beiaherpesvirmae* . HCMV is a double-stranded DNA enveloped virus composed of anucleocapsid surrounded by structured tegument layer and bounded by a trilaminate membrane envelope.

[00558] In most occasions, an initial HCMV infection is asymptomatic, or associated with mild symptoms e.g. sore throat fatigue, flu-like symptoms, and fever. After initial infections, HCMV virus resides in mononuclear ceils without detectable symptoms. HCMV infection may be dangerous to individuals with weakened immune system. HCMV may be transmitted by contact with certain body fluids of an infected individuals (e.g. saliva, urine, semen). HCMV may be transmitted vertically, especially if acquired during pregnancy, leading to a congenital HCMV infection. According to CDC, about 1 in 150 children are born with congenital CMV infection. In about 20 % of cases, congenital HCMV infection may lead to premature birth, birth defects or developmental disabilities, e.g. liver, lung, spleen problems, small head size, small body size or seizures.

[00559] As of today, there is no specific treatment or prevention therapy for HCMV infection. In severe cases of congenital HCMV infection, infants may be treated with an antiviral drug, ganciclovir, to prevent hearing loss and developmental outcomes. However, the drug has serious side effects. There remains a need for prevention therapy and improved therapies for treatment and cure of HCMV infection. Antibodies neutralizing HCMV have been developed. Such antibodies are taught e.g. in international Patent Publication WO20 10007463, US Patent IJS9149524, US8492529 and US8202518, the contents of each of which are mcorporated herein by reference in their entirety.

[00560] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by HCMV.

[00561] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HCMV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 24 (SEQ ID NO: 6628-6736).

Epstein-Barr virus

[00562] Epstein-Barr virus (EBV), also known as human herpesvirus 4 (HHV-4) belongs to the family of *Herpesviridae.* EBV is a double-stranded DNA virus composed of a protein

nucleocapsid surrounded by a tegument layer and bounded by an envelope containing lipids and surface projection of glycoproteins. EBV may enter B cells and epithelial ceils.

[00563] EBV infection causes glandular fever known as infectious mononucleosis, also known as the kissing disease. Typical symptoms include e.g. sore throat, fever swollen lymph nodes in the neck, enlarged spleen, swollen liver, rash and fatigue. Additionally, EBV infection is associated with certain cancers, e.g. central nervous system lymphomas, Hodgkiirs lymphoma, Burkitt's lymphoma, Guiliain-Barre syndrome, multiple sclerosis, and higher susceptibility to certain autoimmune diseases. The virus is transmitted via contact with certain bodily fluids of an infected individual, especially through saliva. The infection affects majority of population. According to CDC, 90 % of adult population have antibodies demonstrating current or past EBV infection.

[00564] As of today, there is no specific therapy for prevention or treatment of EBV infection on the market. Typically, EBV infection is treated with good supportive care. Antibodies for prevention, management and treatment of EBV infection and associated diseases have been developed, e.g. by Wang and Fogg in US Patent publication US20 1500641 74 and Fang et al. in *Inter virology* 50 (4), 254-263 (2007), the contents of each of which are incorporated herein by reference in their entirety.

[00565] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by EBV.

[00566] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat EBV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 31 (SEQ ID NO: 6898-6911).

Varicella zoster virus

[00567] Varicella zoster virus (VZV), also known as human herpes virus 3 (HHV-3) and chickenpox virus, belongs to the family of *Herpes viridae.* VZV is a linear duplex DNA molecule containing two segments **(L** and S) joined covalently. At least five clades of the virus have been identified.

[00568] VZV causes varicella, also known as chickenpox, which is an infection characterized by blister-like rash, itching, fatigue and fever. Chickenpox may be dangerous for babies, adults and individuals with weakened immune system. After primary phase of the infection, VZV resides in the nerves, including cranial nerve ganglia, dorsal root ganglia and autonomic ganglia, and may eventually lead to shingles, which is a viral disease characterized with a painful skin rash, blistering and occasionally nerve pain. Additionally, VZV has been associated with other

complications, e.g. neurological conditions, inflammation of arteries, myelitis, Ramsay Bunt syndrome, Mollaret's meningitis. VZV is transmitted by direct contact or by the respiratory route. VZV is highly contagious. According to CDC, before VZV vaccination, about 4 million people would be affected by chickenpox in the US annually, with more than 10, 000 hospitalized.

[00569] VZV infection may be prevented by a vaccination, which is recommended by CDC to all children and unvaccinated adults. Chickenpox may be treated with antiviral medications, e.g. acyclovir, valacyclovir and famciclovir, or with other symptom relieving medications and therapies. However, the present antiviral medications may have undesirable side effects. There remains a need for improved therapies to treat VZV infection, and its reactivation stages. Antibodies targeting VZV have been developed, e.g. as described in US Patent US5506132, and US Patent application US20J00074906, the contents of which are herein incorporated by their reference in their entirety.

[00570] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by VZV.

[00571] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat VZV. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 31 (SEQ ID NO: 6898-691 1).

Coronavimses

[00572] Coronavimses are a diverse group of enveloped viruses belonging to the family of *Coronaviridae*. Coronavimses contain an envelope, a helical capsid, and a singie-stranded, positive-sense RNA genome. Coronavimses have a characteristic structure with viral spike-shaped glycoprotein populating the stirface of the virus and causing an appearance resembling the solar corona. Coronavimses are a common cause of mammalian and avian infections causing upper respiratory tract, gastrointestinal and central nervous system diseases.

[00573] Human coronavirus 229E, OC-43, NL63, and HKUi are a cause a behind typical, short term "common cold" and affect individuals ail over the world. Typical symptoms of the infections include coughing, sneezing, fatigue and fever. Occasionally the viruses can cause lower-respiratory tract illnesses, such as pneumonia. The viruses are spread by direct contact or by the respiratory route. The infections may be dangerous to the elderly and individuals with weakened immune system. There is no specific treatment or prevention therapy for these coronavirus infections.

[00574] Severe Acute Respiratory Syndrome coronavirus (SARS-CoV) causes a viral respiratory illness. Typical symptoms of the infection include a high fever, headache, body aches, dry coughing and eventually pneumonia SARS-CoV was identified in 2003 in an outbreak starting from Asia. SARS-CoV is transmitted by direct contact with an infected individual or by the respiratory route. According to the WHO, during the 2003 oiitbreak of SARS-CoV, 8098 people worldwide were infected with symptoms and out of them, 774 died. As of today, there is no specific treatment or prevention therapy for SARS on the market. Antiviral medication and steroids may be prescribed to certain patients. Antibodies targeting SARS-CoV have been developed, e.g. as described in US Patent US7728110 and US Patent publication US20150159001, the contents of each of which are herein incorporated by their reference in their entirety.

[00575] Middle East Respiratory syndrome coronavirus (MERS-CoV) causes an acute severe respiratory infection affecting the lungs and breathing tubes. MERS-CoV was identified in 2012. Typical symptoms include fever, cough and shortness of breath, eventually pneumonia and additionally gastrointestinal symptoms. MERS-CoV is highly dangerous to humans. According to the WHO, 36 % of the infections are fatal. MERS-CoV is a zoonotic virus transmitted to humans from animals, e.g. bats and camels, or from human to human. Camels are suggested to be a reservoir for MERS-CoV. Majority of MERS-CoV infection have occurred in the Arabian Peninsula, and especially in Saudi Arabia. As of today, no specific treatment of prevention therapy for MERS-CoV infection is available on the market. Antibodies targeting MERS-CoV have been developed, e.g. as described in international publication WO2015057942, the contents of which are herein incorporated by their reference in their entirety.

[00576] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by SARS-CoV, MERS-CoV and/or other coronaviruses.

[00577] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat coronaviruses. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 25 (SEQ ID NO: 6737-6801).

Poxviruses

[00578] Poxviruses affecting humans include orthopoxvirus, parapoxvirus, yatapoxvirus and moiiusipoxvirus. Poxviruses are typically brick-shaped, enveloped, single, liner or double-stranded viruses with DNA genome. Typically, poxvirus infections cause lesions, skin nodules, or disseminated rash. Poxviruses may be transmitted by direct contact with contaminated

humans, animals or materials. Diseases caused by poxviruses include e.g. smallpox, monkeypox, molluscum conagiosum, vaccinia virus and orf virus infection.

[00579] Smallpox virus infection is highly fatal, and though it does not occur in nature anymore, smallpox virus is considered to be a potential chemical or biological warfare agent. The threat of terrorism has created a need for efficient and impro ved methods for treatment and/or prevention of smallpox infection. The traditional vaccination for smallpox, also applicable against monkeypox, has a rare but severe side effect due to vaccinia virus, which is the active constituent of the vaccine that eradicated smallpox. Vaccinia Immune Globulin (VIG) is the only licensed therapeutic treatment for smallpox, hut is highly variable and available in limited quantities. Antibodies against smallpox have been developed, as described e.g. in US Patent US8623370 and US Patent publication US20140186370, the contents of each of which are herein incorporated by their reference in their entirety.

[00580] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by smallpox virus and/or other poxviruses.

[00581] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat poxvirus. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 27 (SEQ ID NO: 6866-6875).

Enterovirus 71

[00582] Enterovirus 71 (EV71) belongs to the family of *Picornmnridae.* Enterovirus 71 is a single-stranded RNA positive sense virus. The virus has approximately 7411 nucleotides. The RNA genome is enclosed in an icosahedral capsid of structural proteins VPI-VP4. (see, e.g. Tan et al., 2014, *J Biomed Sci;* 21(1): 140, and references therein).

[00583] EV71 infections typically cause hand, foot and mouth (HFMD), which is characterized by fever, mouth ulcers, and vesicles on the palms of the hands and feet. Additionally, EV71 causes severe neurological manifestations, including poliomyelitis-like acute flaccid paralysis, brainstem encephalitis in infants and children. These neurological manifestations may be fatal, or cause permanent neurological consequences, such as delayed neurodevelopment or reduced cognitive function in children. EV71 is transmitted through direct contact with certain bodily fluids, such as saliva, or the respirator)' route, or the fecai-to-moutb route. Outbreaks of EV71 have been reported by WHO in the US, Europe, and more frequently in Asia-Pacific region in the past 30 year.

[00584] As of today, no specific treatment or prevention therapy for EY71 is on the market. Antiviral drugs, e.g. pleconaris and other capsid-function inhibitors (see, e.g. Tan et al. J Riorned Sci. 2014; 21(1): 140), may be prescribed against EV71 infections, though their effectiveness is not well established. There remains a need for prevention and treatment therapies for EV71 infection. Antibodies neutralizing EY71 have been developed. Non-limiting examples include the anti-EV71 antibody MAB979 (developed by Merck Millipore) and those taught by Carderosa et al. in International Patent Publication WO2015092668, the contents of which are incorporated herein by reference in their entirety.

[00585] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by EV71.

[00586] AAV panicles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat EV71. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 28 (SEQ ID NO: 6876-6891).

Rubella virus

[00587] Rubella virus belongs to the family of *Togaviridae*. Rubella virus is a positive sense, single-stranded RNA virus with spike-like, hemagglutinin containing surface projections. The virus core is enveloped by glycosylated E1 and E2. proteins.

[00588] Rubella, also known as German measles or three-day measles, is a viral infection typically characterized by a rash, low fever, nausea, swollen lymph glands behind the ears and the neck, and mild conjunctivitis. At later stage, the infection may develop arthritis and pain in the joints. Typical symptoms of rubella infection are mild and affect children and young adults. Rubella virus is transmitted by the respiratory route and the virus replicates in the nasopharyngeal mucosa and local lymph nodes. However, when an infection is acquired during pregnancy, the virus is transmitted through vertical route with 90% chance and may cause fetal death or congenita! defects known as congenital rubella syndrome (CRS). infants with CRS may have hearing impairments, eye and heart defects, diabetes mellitus, thyroid dysfunction and/or autism. According to the WHO, about 10,000 infants with CRS are born every year, majority occurring in countries with low vaccine coverage.

[00589] As of today, there is no specific treatment for rubella. Rubella may be prevented with vaccination, and rubella has been part of the vaccination program for the past 40 years. However, the infection still persists and an increasing concern related to the life-time of vaccine efficiency exists. There remains a need for long lasting prevention therapy, as well as treatment for rubella virus infection. Antibodies against rubella have been described e.g. in US Patent

US20100143376, the contents of each of which are herein incorporated by reference in their entirety.

[00590] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by rubella.

[00591] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Rubella. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 29 (SEQ ID NO: 6892-6895).

Human papilloma virus

[00592] Human papilloma virus (HPV) is a non-enveloped double-stranded DNA virus belonging to the family of *Papillomaviridae.* Over 170 types of HPV have been identified.

[00593] HPV infections may be asymptomatic, or cause infection related to warts (e.g. plantar, flat or anogeiiital warts), oral infections such as papillomas or multifocal epithelial hyperplasia. The infection may be undetected, and clears from the body to low levels within two years. Infections caused by human papillomavirus (HPV) have been associated with certain cancers of stratified epithelial tissues, e.g. cervical, anal, vaginal, vulvar and penile cancers, lung and throat cancers. Especially HPV16 and HPV18 are known to be carcinogenic. According to the WHO, persistent genital HPV infection may cause cervical cancer which is the second most common cancer in women worldwide. In developing countries, cervical cancer counts for 13 % of all female cancers, and survivor rate worldwide is approximately 50%. HPV is very common. CDC estimates that every one in four individuals in the US has an HPV infection. Most commonly HPV is transmitted by the sexual route, but also the vertical transmission route, or by direct contact to infected blood, or objects may occur.

[00594] Cancers caused by HPV may be prevented by vaccines developed against certain HPV types. The vaccines are available worldwide and are recommended by CDC for all preteen aged children. As of today, there are no specific treatment for HPV infection. There remains a need for prevention and treatment therapy affecting a broad range of HPV infections. Antibodies for HPV have been developed, e.g. as described in International publication WO2015096269, the contents of each of which are herein incorporated by reference in their entirety.

[00595] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by HPV.

[00596] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HPV. As a non-limiting example, the

AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 30 (SEQ ID NO: 6896 and 6897).

Pseudomonas Aeruginosa

[00597] *Pseudomonas Aeruginosa (P. Aeruginosa)* is a common Gram-negative, aerobic, rod-shaped bacterium belonging to the family of *Pseudomonodaceae. P. Aeruginosa* is found in soil, water, skin, Oora, and in most man-made environments around the world. *P. Aeruginosa* is considered as an opportunistic pathogen taking advantage of a weakened immune system.

[00598] *P. Aeruginosa* may cause a variety of mild infections, such as, urinary tract infections, respiratory system infections, dermatitis, soft tissue infections, bacteremia, bone and joint infections, gastrointestinal infections, blood infections, ear infections, skin rash, eye infections and a variety of systemic infections. *P. Aeruginosa* is transmitted through water, contaminated hands, materials or objects. In general, *P. Aeruginosa* infections in healthy individuals are very mild or asymptomatic. However, the infections expose a significant risk for individuals with weakened immunity, such as patients with other underlying illnesses or complications, and especially when in a hospital environment. For example, patients with cystic fibrosis have a susceptibility towards loss of king function due to respiratory tract infection with the bacterium. Patients attached to breathing machines, patients with catheters, or with surgery wounds or burn wounds are potentially at risk for serious and life-threatening infections. *P. Aeruginosa* infection may lead to a fatal sepsis. According to CDC, approximately 51, 000 health-care associated infection occur in the US every year, leading to approximately 400 deaths.

[00599] As of today, there are no prevention therapies for *P. Aeruginosa* infection on the market. Some strains of P. *Aeruginosa* may be treated with antibiotics, e.g. gentamicin, tobramycin, colistin, and amikacin. However, an increasing number of strains of P. *Aeruginosa*, especially those affecting hospitalized patients, are resistant to antibiotics and no specific treatment therapy exists. There remains a need for improved treatment and prevention therapies against P. *Aeruginosa* infections. Antibodies against P. *Aeruginosa* have been developed, such as, panobacumab (developed by Kenta Biotech Inc.), which is an antibacterial antibody against *P. Aeruginosa.*

[00600] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *P. Aeruginosa.*

[00601] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *P. Aeruginosa.* As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 32 (SEQ ID NO: 6912-7196).

Streptococcus bacteria

[00602] *Streptococcus* is a germs of gram-positive bacteria belonging to the family of *Streptococcaceae.* Species of *Streptococcus* are divided into alpha- and beta-hemolytic species. Alpha-hemolytic species cause oxidation of iron in hemoglobin molecules within the red blood cells. Alpha-hemoiytic streptococci include e.g. *Streptococcus pneumoniae* and *Streptococcus viridans.* Beta-hemolytic species cause complete rupture of the red blood cells and include e.g. Laneefield groups A and B, also known as 'group A strep' and 'group B strep'. *Streptococcus* genus includes overall more than 50 species. *Streptococcus* bacteria cause a variety of infections in humans, including dental caries, pneumonia, endocarditis, meningitis, respiratory tract infections, urinary tract infections, neonatal meningitis, pharyngitis and/or sepsis.

[00603] *Streptococcus pneumoniae* is a common bacterium causing, i.e. pneumonia, meningitis, bronchitis, acute sinusitis, conjunctivitis, osteomyelitis, endocarditis and/or septic arthritis. The bacteria is transmitted by direct contact or via the respiratory route. The bacteria resides in the nasopharynx of healthy carriers and proceeds into an infection under certain Circumstances. The infection may be prevented by vaccines, e.g. conjugate vaccine or polysaccharide vaccines. The infection may be treated with antibiotics, e.g. broad-spectrum cephalosporin, and vancomycin, but there is a concern over increasing resistant towards antibodies. According to CDC, *Streptococcus pneumoniae* is currently resistant to one or more antibiotics in 30 % of infections. *Streptococcus pneumoniae* is resistant to e.g. penicillins. There remains a need for improved, non-antibiotic, therapies for treatment of *Streptococcus pneumoniae* and other *Streptococcus* infections. Antibodies for *Streptococcus* have been developed, as described e.g. in US Patent IJS5686070 and US Patent publication US20070003561, the contents of each of which are herein incorporated by reference in their entirety.

[00604] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *Streptococcus pneumoniae* and other *Streptococcus* bacteria.

[00605] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *Streptococcus pneumoniae.* As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 33 (SEQ ID NO: 6900-6902. 6905-6907, 6911, 7197-7229).

Staphylococcus bacteria

[00606] *Staphylococcus* is a genus of gram-positive bacteria belonging to the family of *Staphylococcaceae.* The germs includes overall approximately 40 species. Most species of the genus are harmless and reside in the skin and mucous membranes of humans. *Staphylococcus* bacteria may also be found in the soil. The bacteria may cause diseases either through toxin production or penetration. Staphylococcal toxins are a common cause of food poisoning. Staphylococcus bacteria may cause a variety of diseases, e.g. localized or diffuse skin infection, gastroenteritis, ear infections, septic arthritis, osteomyelitis, sinusitis, infective endocarditis and/or toxic shock syndrome.

[00607] *Staphylococcus aureus (S. aureus)* is typically residing in human nose asymptomatically. In certain circumstances, *S. aureus* infections may affect many tissues and organs. Individuals with chronic conditions, e.g. diabetes, cancer, vascular disease, eczema and lung disease, have an increased susceptibility towards *S. aureus* infections. *S. aureus* may cause skin infections, such as, pimples, impetigo, atopic dermatitis, cellulitis folliculitis. More serious forms of infections include pneumonia, meningitis, osteomyelitis and endocarditis. *S. aureus* may also cause food poisoning. In severe cases, *S. aureus* infection may enter the blood stream causing bacteremia and/or sepsis. As of today, there is no medical therapy for prevention of the infection. Some strains of *S. aureus* may be treated with antibiotics. However, increasing resistance towards antibiotics is a concern. Currently several antibiotic resistant forms of *S. aureus* exist, including, but not limited to, Methiciilin-resistant *Staphylococcus aureus* (MRSA), Vancomycin-raterraediaie *Staphylococcus aureus* (VISA) and Vancoraycin-resistant *Staphylococcus aureus* (VRSA). The drug resistant forms of S. aureus are more frequent in hospital environments.

[00608] *Staphylococcus epidermidis (S. epidermidis)* resides in the normal human skin flora and may cause an infection to individuals with weakened immune system, and to individuals who have catheters, prostheses or surgical implants. *S. epidermidis* has an ability to colonize on plastic materials or devices placed within the body. The infection may be treated with some antibiotics, but they do *not* remove the infection and can only be used to manage such infections. Many *S. epidermis* strains are resistant to antibiotics, such as penicillin, methiciilin and/or amoxicillin, and increasing resistance to antibiotics in a concern.

[00609] There remains a need for prevention and/or improved treatment therapies against *Staphylococcal* infections. Antibodies targeting *Staphylococcal* bacteria have been developed. As an example, pagadaximab (developed by Medlmmune and AstraZeneca) is a monoclonal antibody for prevention of staphylococcal sepsis and may be administered to infants with low birth weight.

[00610] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *Staphylococcus* bacteria

[00611] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *Staphylococcal* infections. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 34 (SEQ ID NO: 7230-7478).

Clostridium Tetani

[00612] *Clostridium Tetani (C. Tetani)* is a rod-shaped, anaerobic, Gram-positive bacteria belonging to the family of *Clostridiaceae.* A matured bacterium develops a terminal spore, which is resistant to heat and common antiseptics. *C. tetani* produces tetanospasrain toxin. *C. tetani* is found as spores in soil and in the gastrointestinal tract of animals.

[00613] *C. tetani* infection spreads the tetanosparain toxin to the body, causing tetanus, also known as lock jaw. Tetanus is a dangerous disease characterized by painful tightening of the muscles. The disease may lead to locking of the jaw and neck, leading to inability to open mouth or swallow. The tightening may affect the whole body. In severe cases, the infection may lead to breathing difficulties, pneumonia, or pulmonary embolism. Even more serious is an infection acquired during pregnancy, leading to almost always fatal neonatal tetanus of an infant. The bacteria is typically transmitted through broken skin by direct contact with contaminated soil or objects, or saliva or feces of a contaminated animal. Especially susceptible are individuals with bums, puncture wounds, crush injuries or injuries with dead tissue, individuals having animal bites or scratches. Tetanus is fairly uncommon in developed countries. However, the WHO reported an estimated 50, 000 neonatal tetanus deaths in year 2008. A program form elimination of tetanus was started in 1989 by the WHO.

[00614] Tetanus may be prevented efficiently by a four vaccine combination, DTaP, Tdap. DT, and Td, given to children and adults. For adequate immunity, the primary vaccine is administered during childhood, a booster dose during adolescence and every 10 years thereafter during adulthood. *C. tetani* infection may be treated with antibiotics, wound care and with human tetanus immune globulin (an antitoxin). Despite the existing treatment methods, approximately 10 % of tetanus infections lead to death, according to CDC. There remains a need for longer lasting vaccine as well as improved treatment therapies against *C. tetani* infections. Antibodies against *C. Tetani* have been developed, as described e.g. by Larrick, J.W. et al., 1992, *Immunol. Rev.* 130, 69-85, and de Kruif, J. et al, 2009, *J. Mol Biol.* 387 (3), 548-558, the contents of each of which are herein incorporated by reference in their entirety.

[00615] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *C. Tetoni.*

[ΘΘ616] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *C. Tetani.* As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 35 (SEQ ID NO: 7479-7535).

Bordeteila

[00617] *Bordeteila* is a genus of Gram-negati ve, *coccobacilli* belonging to the family of *Alcahigenaceae.* The structure of the bacteria consists of an outer membrane with lipopolysaccharides and phospholipids forming a capsule. *Bordeteila* bacteria affecting humans include, but are not limited to, *B. pertussis, B. parapertussis* and *B. bronchiseptica. B. pertussis* resides in the upper air pathways, mostly the trachea and the bronehii, of humans. *B. parapertussis* resides in the upper air pathways of mammals. The bacteria release toxins that cause damage and swelling of the respirator}' pathways.

[00618] Pertussis, also known as whooping coiigh, is a highly contagious infection of the respiratory track caused most commonly by *B. pertussis,* and occasionally by *B. parapertussis.* Typical symptoms of the infection include severe coughing and difficulty to breathe accompanied by a runny nose, apnea and fever. Additional complications for infants include pneumonia, convulsions, apnea, and encephalopathy. The bacteria are transmitted through the respiratory tract route. The disease is especially dangerous for infants. According to CDC, about 30,000 infections were reported in the US in 2014. CDC reports 277 deaths occurring from 2000 through 2014, out of which 241 where infants less than 3 months of age.

[00619] Pertussis may be treated with antibiotics, such as, erythromycin, claritlvromycin or azithromycin. However, an increasing resistance to antibiotics is a concern. Pertussis caused *8. pertussis* may be prevented by vaccination, e.g. by DTaP combination vaccine, which is recommended routinely for infants by CDC and WHO. Despite the widespread vaccination, the disease has insisted. The protection provided by the traditional vaccination is estimated to be 3-6 years. There remains a need for prevention therapies providing a longer lasting immunity, as well as for improved, non-antibiotic, treatments. Antibodies for prevention and-'or treatment of pertussis have been developed, as described e.g. in International publication WO2014160098, and Hussein, A.H. et al., 2007. *Infect. Immun.* 75 (11), 5476-5482, the contents of each of which are herein incorporated by reference in their entirety.

[00620] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *B. pertussis, B. parapertussis* and/or other *Bordetella* bacteria.

[00621] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *Bordetella* infection. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 36 (SEQ ID NO: 7536-7560}.

Mycobacterium

[00622] *Mycobacterium* is a genus of nonmottle and aerobic bacteria, belonging to its own family *of Mycobacteriaceae. Mycobacteria* have an outer membrane, and a hydrophobic and waxy cell wail with mycolic acid/mycolates. The cell wall is neither truly Gram-positive nor -- negative. In general, the infections are difficult to treat and the bacteri a is naturally resistant to many antibiotics, e.g. penicillin, due to the cell wall. Mycobacteria includes species, such as, but not limited to, *M. tuberculosis, Nontuberculous mycobacteria* (NTM), *M. leprae, M. bovis, M. africanum, andM. microti.*

[00623] *M. tuberculosis* is a genetically diverse bacterium and most common and dangerous of the mycobacteria family species. *M. tuberculosis* causes tuberculosis (TB) which is an infection mainly affecting the lungs. Typical early symptoms include cough, fever, night sweat, and weight loss. The disease may be mild for a period of time and therefore early diagnosis is difficult. Eventually the symptoms get more severe and coughing sputum and blood may occur. TB may be transmitted by the respirator/ tract. TB affects ail ages of the population, but is most dangerous to children, and individuals with weakened immune systems, e.g. HIV patients. According to the WHO, TB is referred to as a top infectious disease killer worldwide. WHO reports an estimated 9.6 million infections of TB in 2014. out of which 1.5 million cases were fatal. The disease is globally spread, but it is most abundant in the South-East Asia and Western Pacific Regions.

[00624] TB may be prevented by vaccinations, i.e. Bacille Clamette-Guerin vaccine. The vaccine is provided for children and adults exposed to environments with high risk of infection. However, the vaccine is not always efficient against TB, e.g. due to the diversity of strains geographically. TB may be treated with a 6 to 9 month course of combinational antimicrobial drug therapy. Antimicrobial drugs effective against TB include e.g. isoniazid, rifampin, ethambutol, and pyrazinamide. However, an increasing resistance towards the medication is a concern. Certain strains of existing TB are identified as multi-drug resistant TB strains, which do not respond to therapy with e.g. isoniazid. rifampicin, or other common drugs. WHO reports an

estimated 480 000 muhidrug-resistant TB infections in 2014. There remains a need for prevention therapies protecting against broad spectrum of strains, as well as for improved treatment of *M. tuberculosis* and/or other *mycobacteria.* Antibodies against mycobacteria have been developed as described e.g. in US Patent publications US20130309237, US20130309237, IJS20060229438, the contents of each of which are herein incorporated by reference in their entirety.

[00625] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *M. tuberculosis* and/or other *mycobacteria.*

[00626] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *myobacterium* related diseases. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 37 (SEQ ID NO: 7561-7576).

Francisella Tularensis

[00627] *Francisella Tularensis (F. tularensis)* is a facultative intracellular Gram-negative, rod-shaped bacterium belonging to the family of *Francisellaceae. F. tularensis* resides in invertebrates, birds, reptiles, fish, and mammals, including humans. It is one of the most infectious and pathogenic bacteria known (see, e.g. Pechous et al, 2009, *Microbiol Mol Biol Rev.:* 73(4): 684-711).

[00628] *F. Tularensis* causes infection called Tularemia. Severity of tularemia varies from mild to fatal. *F. Tularensis* may be transmitted to a human by direct skin or eye contact, by the respiratory roiite or by consumption of contaminated food or drink. Most commonly, the infection is acquired while handling infected animals. Most common form of tularemia is ulceroglanduiar tularemia, characterized by skin ulcers on the site of infection accompanied by swelling or regional lymph glands. Ulceroglanduiar tularemia is typically acquire by a tick, or deer fly bite. Pneumonic tularemia is an infection of the respiratory tract characterized by a cough, chest pain, and difficulty of breathing. Pneumonic tularemia is transmitted through the respiratory route and may be fatal if not treated. Oropharyngeal tularemia is transmitted by contaminated food or beverage and causes a sore throat, mouth ulcers, tonsillitis and swelling of lymph glands in the neck. Other forms of tularemia include glandular, oculoglandular (affecting the eyes) and typhoidal (combination of the general symptoms). *F. Tularensis* is considered to be a potential biological and chemical warfare agent.

[00629] As of today, there is not preventive therapy for tularemia infection on the market. Some vaccines have been under development (see, e.g. Pechous et al., Microbiol Mol Biol Rev.

2009 Dec; 73(4): 684-71 1). Tularemia may be treated with antibiotics, such as, streptomycin, gentamicin, doxycyctine, and ciprofloxacin. However, increasing resistance against antibiotics is a concern. There remains a need for improved prevention and treatment therapies for *F. Tularensis* infections. Antibodies against *F. Tularensis* have been developed, e.g. as described by Rynkievvicz, M.J. et al., 2012, *Biochemistry,* 51 (28), 5684-5694 and Lu, Z., et al., 2013, *Immunology,* 140 (3). 374-389, the contents of each of which are herein incorporated by reference in their entirety.

[00630] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *F. Tularensis.*

[ΘΘ631] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *F. Tularensis* related infections. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 38 (SEQ ID NO: 7577-7592).

Toxoplasma gondii

[00632] *Toxoplasma gondii* is a parasitic protozoan infecting warm-blooded animals, including humans. Domestic cats and other felines are the most desired hosts for toxoplasma gondii, as they are the only hosts where the protozoan is capable of sexual reproduction. According to CDC, more than 60 million people in the US may be infected by *Toxoplasma gondii.*

[00633] *Toxoplasma gondii* causes toxoplasmosis, which is typically asymptomatic in healthy individuals and is controlled by the natural immune system. The infection may be obtained from undercooked, contaminated food, especially pork, lamb and venison, from food contaminated by utensils, or contaminated hands, occasionally from contaminated drinking water, or by the fecal-to-oral route from cat feces. *Toxoplasma gondii* may also be transmitted by vertical route, especially when the protozoan is acquired during pregnancy. Children infected during or just prior to pregnancy may have eye problems, or brain damage at birth, or may develop symptoms later in their lives. Toxoplasmosis may be dangerous to individuals with a weakened immune system, such as patients with AIDS, undergoing certain chemotherapies or having organ transplants.

[00634] Toxoplasmosis may be treated with certain medications such as antibiotics called sulfadiazine and pyrimethamine, which is an anti-parasite medication used for e.g. malaria. However, resistance to both of the medications is an increasing concern. There remains a need for improved treatment methods as well as prevention therapies against *Toxoplasma gondii* infection. Antibodies targeting *Toxoplasma gondii* have been developed, as described by e.g.

Graiiie, M. et **al.,** 2005, *J. Mol Biol.* 354 (2), 447-458, the contents of which are herein incorporated by reference in their entirety.

[00635] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *Toxoplasma gondii.*

[00636] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *Toxoplasma gondii* related infections. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 40 (SEQ ID NO: 7617 and 7618).

Candida yeast

[00637] Typically, species of yeast are commensals and endosymbionts of human hosts, but may cause an infection under certain circumstances. *C. albicans* is a yeast belonging to the family of Saccharoraycetaceae. *C. albicans* causes infection of the mouth characterized by white patches on the tongue, mouth and throat. The infection of the mouth is most typical with new¬ born babies, the elderly and individuals with weakened immune system, e.g. HIV/AIDS patients. Optionally, the infection may affect the nails, leading o brittle and defected nails. Optionally, the infection may cause an infection of the vagina, leading to genital burning or uncomfortable discharge. Typically, Candida albicans infections are mild and localized. However, the infection may be severe or fatal for individuals with underlying health problems and left untreated. Invasive candidiasis refers to an infection spreading to many parts of the body, including the heart, brain, eyes, bones and/or joints. Candidemia refers to an infection where Candida yeast is present in the blood stream. Severe forms of *C. albicans* infections affect individuals in health care environments, e.g. patients with central venous catheter, patients treated at an intensive care unit, patients undergoing antibiotic treatments, treatments for kidney failure, recovering from a surgery, and patients with chronic diseases, e.g. diabetes and/or HIV/AIDS. *C. albicans* is typically transmitted from mother to an infant during childbirth and it remains as a species of human's normal microflora. It may also be transmitted through the sexual transmission route. Other species of Candida yeast family include, e.g., *C. glabrata, C. parapsilosis, C. tropicalis, C. krusei and C. lusilaniae.*

[00638] *C. albicans* infection may be treated with antifungal drugs, e.g. nystatin, clotrimazole, amphotericin B oral suspension) or systemic oral azoles (e.g. fluconazole, itraconazole, or posaconazole). Despite the medical therapy available, some forms of *C. albicans* infections are dangerous, or life-threatening. There remains a need for improved prevention, and/or treatment

therapies against *C. albican* infections, for example by antibody therapies. Efungumab (developed by NeuTee Pharrna) is an antibody for treatment of invasive *C. albicans* infection.

[00639] In some embodiment, methods of the present invention may be used to prevent and/or treat *C. albican* infections.

[00640] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *C. albican* related infections. As a non-iimiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 41 (SEQ ID NO: 7619).

Human Immunodeficiency Virus (HIV)

[00641] Human immunodeficiency virus (HIV) is a roughly spherical enveloped ENA vims belonging to the family of Retroviridae. HIV is composed of two positive single-stranded RNA copies. The viral core contains a viral capsule protein, p24, which surrounds the two single stranded RNAs and the enzymes for HIV replication. The viral envelope consists of two lipid layers, the outer layer glycoprotein 120 (gp 120) and the transmembrane glycoprotein 41 (gp41). Gpl20 attached to the host cell whereas gp41 has a role in the cell fusion process. For replication, the virus needs a host cell and the RNA first transcribes into DNA by enzyme reverse transcriptase. HIV infects the CD4 lymphocyte (T cell) leading to depletion of CD4÷ T cells and loss of CD4+ T-eeli function, as infected cell loses its function and converts to a HIV-replicating cell. (see, e.g. Okoye and Picker, 2013, Immunol Rev.; 254(1): 54-64, and references therein). Additionally, HIV infection leads to B lymphocyte (B cell) hyper-activation and dysfunction (see, e.g. Moir and Fauci, 2009, Nat Rev Immunol.; 9(4): 235-245, and references therein). The virus may be transmitted through sexual transmission route, vertical transmission route, iatrogenic (medical procedure) route, or in direct contact with certain body fluids with high concentration of HIV, including e.g. blood, breast milk, semen, vaginal and rectal secretions. Two types of HIV (HTV-1 and HIV-2) have been identified. HIV-1 has higher mfectivity and has spread around the globe whereas HIV-2 is more localized to West Africa. According to CDC. there is about 36.9 million people in the world with HIV/AIDS with about 2 million cases arising every year. The infection is most abundant in Sub-Saharan Africa.

[00642] In acute HIV infection stage, within 2-4 weeks after infection, infected patients experience flu-like illness. In the second stage the infection is asymptomatic and the HIV replication is at low level. The second stage may last for years or decades, especially when treated with HiV medication. Eventually, HIV causes acquired immune deficiency syndrome (AIDS), which is a clinical condition characterized by severe immunosuppression attacking the CD4 cells, making individuals susceptible to life-threatening malignancies and infections.

Complications associated with HIV/AIDS include common bacterial and viral infections, parasite infections, certain cancers (e.g. Kaposi's sarcoma, Non-Hodgkin's lymphoma, and angiosarcoma), progressive multifocal leukoencephalopathy (PML) and wasting.

[00643] As of today, there is no prevention therapy or cure for HIV/AIDS. However, with antiretro viral (ART) therapy, the disease may be managed for a long period of time. ART therapy comprises of five classes of drugs used in different combinations to treat HIV. The drugs target the different phases of the retrovirus life-cycle. However, there remains a need for improved therapies for prevention, management and/or treatment of HIV/AIDS.

[00644] Antibodies for treatment and prevention oi' HIV infection have been developed. For example, commercial antibody ibalizumah (developed by Tainted Biologies Inc.) is anon-inimnnosuppressive monoclonal antibody binding to CD4, *Anaplasma phagocytophilium* inhibiting the viral entry process. As another example, suvizumab (developed by Kaktsuden, Chemo-Sero Therapeutic Research Institute) is a humanized antibody targeting the HTV-l envelope glycoprotein GP120. As a non-limiting example, any of the antibodies in Table 42, variants or fragments thereof may be used in the treatment and/or prevention of HIV.

[00645] Antibodies neutralizing HIV-1 and HIV-l strains have been identified, but as of today, the researchers have not been able to develop a vaccination for HIV. HIV has a capability to evolve with unusually high somatic mutation and recombination rate. So far, conventional vaccines have not succeeded in eliciting analogues of the broadly neutralizing antibodies. An alternative approach suggested involves using adeno-associated vectored gene delivery for expression of antibodies from muscle tissue (e.g. Baiasz et al, 2012, Nature Letter, 481, 81-84, Balasz et al, 2014, Nat Med.; 20(3): 296-300, Saunders et al, 2015, J Virol., 89(16):8334-45, and US Patent publication US200302 19733, the contents of which are herein incorporated by reference in their entireiy). The studies have demonstrated efficient and long lasting protection from HIV infection by e.g. intravenous or mucosal surface transmission.

[00646] AAV Particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat HIV infection and AIDS. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 42 (SEQ ID NO: 7620-9220).

Therapeutic applications: Toxins

[00647] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat infectious disease. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Tables 14-17 (SEQ ID NO: 3549-3914).

[00648] Toxins are a group of substances that are highly poisonous and dangerous to humans. Toxins are infectious agents in form of bacteria, viruses, fungi, proteins, and other chemical and/or biological substances. Toxins may lead to fatal conditions. Toxins are produced by nature, and may be produced synthetically. Exposure to toxins may be unintentional and occur when in contact with toxic plants, or contaminated food, water, livestock or animals. Due to the life-threatening impact of toxins, they are considered to be potential biological and/or chemical warfare agents that may be applied as weapons of mass destruction in war field. They also impose a threat to be used as means for terrorist attacks.

Ricin

[00649] Is a naturally occurring carbohydrate-binding lectin protein produced by castor oil plant growing in Eastern Africa, India, Southeastern Mediterranean basin area, and in tropical regions. Ricin may also be manufactured from the waste products when processing castor beans. Ricin has a globular structure with two toxin chains, chain A and chain B, which both need to be present for the cytotoxic affect. Ricin kills cells by inhibiting protein synthesis. Chain B penetrates to the cell whereas the disulfide bond joining chain A to chain B lectin has an affinity to bind to cell surface carbohydrates, (see, e.g. Friedman and Rasooly, 2013, *Toxins (Basel): 5(4):* 743-775). Ricin is highly toxic to humans with median lethal dose (LD_{50}) of 22 micrograms per kilogram of body weight. The exposure to Ricin may be by inhaling, ingestion or by injection. The symptoms are dependent of the method of exposure. When inhaled, ricing causes severe inflammation of the lungs, causing would has symptoms including coughing, difficulty breathing, muscle ache and nausea. When ingested, ricin induces internal bleeding of the stomach and intestines leading to pain, vomiting and bloody diarrhea, and eventual failure of the kidneys, liver and spleen. When injected, ricin induces failure of the muscles and lymph nodes, and eventually failure of the liver, kidney and spleen. There is no known treatment for Ricin poisoning.

[00650] Unintentional poisoning by Ricin is uncommon. However, Ricin is a potential biological and chemical warfare agent creating a need for treatment and prevention therapies for ricin poisoning. Antibodies targeting ricin have been developed, as described e.g. in International publication WO2015100409, the contents of which are herein incorporated by reference in their entirety.

[00651] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by ricin.

[00652] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Ricin related infections and/or conditions.

As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 14 (SEQ ID NO: 3.549-3568).

Bacillus anthracis

[00653] *Bacillieus anthracis* is a Gram-positive, rod-shaped bacterium causing anthrax disease (see. e.g. Spencer. 2003, *J Clin Pathol.*;*56(3):* 182-187, and references therein). Most animals, especially herbivores, are susceptible to infection of *Bacillieus anthracis.* Anthrax may be infected via respiratory exposure, skin contact or eating contaminated food, in most cases meat. inhaled anthrax causes flu-like symptoms, pneumonia and severe respiratory collapse. Gastrointestinal anthrax causes severe diarrhea, acute inflammation of the intestinal tract, and vomiting of blood. Skin exposure to the bacteria will lead to boii-like skin lesions forming an ulcer with black center. Typically, infection to humans occurs by eating contaminated meat or while handling infected animals or their product such as skin, wool or meat *Bacillieus anthracis* is a potential biological warfare agent. In 2001, weeks following the September 11 ten-oust attacks, letters containing *Bacillieus anthracis* were mailed to news media offices and two U.S. Senators resulting in death of five people and infected many more.

[00654] Anthrax may be treated with antibiotics, such as penicillin and amoxicillin, and may be prevented by vaccines, developed both for humans and animals. However, due to increased threat of biological warfare and terrorism, improved methods of treatment are in demand. Anthrax may also be treated by antibody therapy. For example. Raxibacumab (developed by-Cambridge Antibody Technology and Human Genome Sciences) is an antibody for the prophylaxis and treatment of inhaled anthrax.

[00655] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *Bacillieus anthracis.*

[00656] AAV panicles and methods of using the AAV particies described in the present invention may be used to prevent, manage and/or treat *Bacillieus anthracis* related infections and/or conditions. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 15 (SEQ ID NO: 3569-3813).

Shiga toxin and Shiga-like toxin

[00657] Shiga toxin, including two major types Stx1 and Stx2, is a toxin produced by *Shigella dysenteriae,* a rod-shaped bacteria belonging to bacterial genus *Shigella.* Shiga toxin inhibits protein synthesis within cells. The toxin enters cell via a marcopinosome and inhibits the protein synthesis by cleaving a specific nucleobase RNA of the 60S subunit of ribosome. Shiga-like

toxins 1 and 2 are structurally similar to Six l and Stx2 and are produced by enterohemorrhagi c strains of *Escherichia coli* (EHECj strains. (see, e.g. Friedman and Rasooly, Toxins (Basel). 2013 Apr; 5(4): 743-775). EHEC type 0 157 is the most common pathogen causing *E. Coli* outbreaks in the US. Stx2 is considered to be orders of magnitude more toxic that Stxl. The severity of Shiga toxin foodborne illnesses range from mild diarrhea to a iife-threatening complication known as hemolytic uremic syndrome (HUS). HUS is a disease associated with hemolytic anemia, acute kidney failure and low platelet count. Cattle is the major source or infection to humans, but the disease may be spread by birds or pigs. Shiga infection is typically obtained from contaminated food or drink, such as meat, unpasteurized milk, or contaminated water, or by contact with cattle. Shiga toxin and Shiga-Iike toxins considered to be potential chemical and biological warfare agents.

[00658] As of today, there is no prevention therapy or specific treatment for Shiga and Shiga-Iike toxins. Recent developments have been made in antibody therapy of Shiga toxin induced HUS. For example, SHIGAMAB™ (developed by Belius Health inc.) is a monoclonal antibody for treatment of Shiga toxin induced HUS.

[00659] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *Shigella dysenteriae*.

[00660] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *Shigella dysenteriae* related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 17 (SEQ ID NO: 3844-3914).

Botulinum toxins

[00661] Botulinum toxins are neurotoxins produced by *Clostridium bacteria* and they cause a disease called botulism which is characterized by weakness, problems in vision, tiredness, and problems with speech, followed by weakness of the arms, chest muscles and legs. Botulism may be fatal. There are seven different botulinum neurotoxins with a four-domain structure varying in antigenic properties and interactions with intracellular targets. L-ehain enters the cytosoi, cleaves the synaptosomal protein and blocks neurotransmitter release resulting in peripheral neuromuscular blockade and flaccid paralysis in humans. (see, e.g. Friedman and Rasooly, Toxins (Basel). 2013 Apr; 5(4): 743-775) Botulinum neurotoxins are highly dangerous to humans, serotype A having a median lethal dose (LD_{50}) of 0.8 micrograms for a human of 70 kg weight. The bacteria is common in soil and water and may produce the botulinum toxins when exposed to low oxygen levels and certain temperatures. Outbreaks of foodborne botulism occur

occasionally. Most susceptible to contamination by botulinum are baked products, fresh mussels, canned fruit and vegetables. Infant botulism occurs when the toxins are produced and released by bacteria in the infant's intestines. Botulism may also occur in wounds where the bacteria in the absence of oxygen produces and releases the toxins. Wound botulism is most common in cases where contaminated needles are used for injection. Botulinum toxins are potential biological and chemical warfare agents.

[00662] As of today, there is no prevention therapy for botulism. Botulism may be treated with antitoxins that block the circulation of toxins in the blood and prevent worsening of the disease. However, the antitoxins are expensive and not easily available. In cases of wound botulism, the area infected may be removed surgically. Additionally, good supportive care therapy is applied. There remains a need for therapies to prevent and treat botulism. Antibodies targeting botulinum toxins are developed, as described e.g. in US Patent publication US20I30058962, and International publication WO2015100409, the contents of each of which are herein incorporated by reference in their entirety.

[00663] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by botulinum toxins.

[00664] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat botulinum toxin related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 16 (SEQ ID NO: 3814-3843).

Therapeutic applications: Neglected Tropical Diseases (NTDs)

[00665] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat infectious disease. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Tables 10-13 (SEQ ID NO: 3327-3548).

[00666] Neglected Tropical diseases (NTDs) are a diverse category of communicable diseases present in tropical and subtropical environments. NTDs affect more thai one billion people in about 150 countries. NTDs are a significant public health problem costing the involved developing economies billions of dollars annually. The diseases affect mostly the populations with inadequate sanitation, and those in contact with infectious vectors, domestic animals and livestock. In May 2013, the 66th WHO Assembly announced resolution WHA66.12 to integrate measures and plan investments to improve the wei!being of populations affected by NTDs. NTDs include Buruli ulcer, Chagas disease, Dengue and Chikungunya, Dracunculiasis (guinea-

worm disease). Echinococcosis, Endemic treponematoses (Yaws), Foodbome trematodiases, Human African trypanosomiasis (sleeping sickness), Leishmaniasis, Leprosy (Hansen disease), Lymphatic filariasis, Onchocerciasis (river blindness), Rabies, Schistosomiasis, Soil-transmitted helminthiases, Taeniasis/Cysticercosis and Trachoma.

Chikungunya virus

[00667] Chikungunya virus is an arbovirus belonging to the *Togoviridae* family. The genome is a single-strand RNA molecule encoding four non-structural and three structural glycoproteins (€, El, E2) (see, e.g. Caghoti ***etal.,*** 2013, *New Microbiol* .;36(3):2\1-27, and references therein). Chikungunya fever is a mosquito-borne disease caused by chikungunya virus. The symptoms include a fever lasting 2-7 days, rash and flu-like symptoms accompanied by a joint pain that may last for weeks, months or even years. The disease may be dangerous for the elderly and individuals with chronic medical problems. Chikungunya virus is spread by *Aedes albopictus and Aedes aegypti.* Outbreaks of chikungunya fever have occurred in Africa, Asia. Europe and Indian and Pacific Oceans, and more recently in islands in the Caribbean. As an example, according to the WHO, an outbreak of 1.9 million cases in India, Indonesia, Maldives, Myanmar and Thailand since 2005 has been reported. More recently, as of April 2015 more than million cases have reported in Caribbean Islands. Latin American countries and the United States.

[00668] As of today, there is no specific treatment or vaccination for chikungunya fever. The disease is typically treated with supportive care therapy, as well as anti-inflammatory drugs and medicines to relieve the symptoms. Research and development on vaccinations has been done but none has been approved for commercial use so far. Antibodies for detection and treatment of Chikungunya have been developed. E.g. fully human antibodies binding to an epitope located in an antigenic site of the chikungunya virus E1 and E2 envelope proteins were in US Patent Publication US20130189279, the contents of each of which are incorporated herein by reference in their entirety.

[00669] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by chikungunya virus.

[00670] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat chikunguyna virus related infections and/or conditions. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 13 (SEQ ID NO: 3543-3548).

Dengue virus

[00671] Dengue virus belongs to the family *of Flaviviridae,* genus of F!avivirus. It is an enveloped, positive strand RNA virus containing two integral membrane proteins envelope (E) and premembrane (prM). Dengue virus is closely related to e.g. Yellow fever. West Nile virus and St. Louis and Japanese encephalitis viruses. There are five serotypes of the virus that can cause dengue fever, which is a mosqtiito-borne tropical disease. Neutralizing antibodies target the protein E as it binds to the cellular receptors and mediates the viral entry into cells. infection with a serotype may produce a lifelong immunity to that serotype but no long-term mummify against other serotypes, (see e.g., Wahala and de Silva, 2011, *Viruses.;* 3(12): 2374-2395, and references therein). In fact, an infection by a second serotype may lead to a more severe form of disease, due to the complexity of the antibody respond and possible antibody dependent enhancement (ADE), which hypothesizes that weakly neutralizing antibodies from the first infection bind to the second serotype and enhance the infection. The symptoms of dengue fever are similar to flu, including fever, headache, muscle and joint pain and skin rash. The disease may also manifest as a potentially lethal complication called severe dengue, also known as dengue hemorrhagic fever. The disease may be dangerous to individuals with chronic diseases, such as diabetes or asthma, or children and the elderly. Dengue virus is spread by several mosquito species, out of which *Aeaes aegypti* is the most common. Dengue may also be transmitted via infected blood or organ donation or by the vertical transmission route. According to the WHO, the estimated niimber of dengue infections annually could be as high as 390 million.

|00672] As of today, there is no specific treatment or prevention therapy for dengue fever. Antibodies targeting dengue virus have been developed. As an example, antibodies neutralizing four serotypes of dengue virus have been in US Patent publication US20150225474, US2015Q218255 and in US Patent US 9073981, the contents of each of which are incorporated herein by reference in their entirety.

[00673] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by dengue virus.

|00674] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Dengue virus reiated infections **and/or** conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 10 (SEQ ID NO: 3327-3449).

Trypanosoma cruzi

[00675] *Trypanosoma Cruzi* (*T. cruzi)* is a species of parasitic eugienoid protozoan. *T. cruzi* causes Chagas disease, also known as American trypanosomiasis, which is a tropical parasitic disease. The symptoms of Chagas disease at the early stage include fever, swollen lymph nodes, headaches or local swelling at the site of bite. The chronic phase of Chagas starts after 8-12 weeks, which may be symptomless, or include enlargement of the ventricles of the heart, which may result in heart failure, or to an enlarged esophagus or enlarged colon. The severity of Chagas disease varies from almost unnoticeable to fatal. Chagas disease is spread by an insect vector triatomine bug. These bugs get infected with *T. cruzi* by feeding on the blood of an infected human or animals, and they spread it further by bites and ingestion of blood. The triatomine bug is also known as a "kissing bug" referring to its tendency to feed on people's faces. *T. cruzi* may also be transmitted through blood transfusions or through breast milk. Chagas disease is present mainly in 12 Latin American countries, but has also spread to other continents. According The WHO, over 10 000 people die every year from Chagas disease, and 25 million people are in the risk of infection.

[00676] As of today, there is no specific prevention or treatment therapy for Chagas disease. The traditional therapies for Chagas have been involved with attempts to kill the parasite and treatment of the symptoms. For example, azole and nitro-derivative drugs have been used, but have not been successful in removal of the parasite fully. Other mechanisms to treat the disease have been under research. After infection in mammals, the parasite incorporates a charged carbohydrate (sialic acid) to survive to the chronic phase of the disease. To do so, the parasite scavenged sialic acid it from the host's sialogiycoconjugates, through a transglycosylation reaction catalyzed by an enzyme calied trans-sialidase. The trans-siaiidase has been identified as a potential target for drug development. Buschiazzo et al. have reported an antibody inhibiting the T. cruzi trans-siahdase enzyme providing an antibody therapy mechanism for Chagas disease (see, Buschiazzo et al., 2012, *PLoS Pathol.* 8 (1), E1002474, and references therein).

[00677] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by Chagas disease.

[00678] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Chagas disease. As a non-iirniting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 12 (SEQ ID NO: 3541 and 3542).

Rabies virus

[00679] Lyssaviruses are a genus of RNA viruses belonging to the family of *Rhabdoviridae.* Rabies virus is a neurotropic virus with cylindrical morphology. After infection, rabies virus

enters the peripheral nervous system, and further to central nervous system by retrograde axonal transport. Rabies virus and Australian bat lyssavirus cause rabies. Rabies affects humans and warm-blooded animals. The early stage symptoms include flu-like signs, but later the disease manifests as paralysis, anxiety, insomnia, abnormal behavior, hallucinations. Humans and animals infected may also experience hydrophobia, "fear of water", which is considered a characteristic symptom of the disease. Eventually the disease affects the central nervous system and brain, causing death. Humans are typically infected by being bitten, scratched or licked by an animal with the disease. Most commonly the infection is by dogs. Whereas efficient vaccination programs for animals have been able to reduce or even eliminate rabies in developing countries, the disease still affects poor population mainly in Africa and Asia. According to the WHO, post-bite treatment and vaccination is provided for 15 million people annually.

[00680] Rabies is a vaccine-preventable disease and especially systematic vaccination of dogs has been a cost-effective strategy for prevention of rabies. Post-exposure prophylaxis (PEP), the treatment of bite victims immediately after the exposure, includes local treatment of the wound, rabies vaccination and administration of rabies immunoglobulin. Though efficient vaccines for rabies have been developed, there remains a need for treatment/or management of rabies to prevent death after rabies virus has entered the central nervous system (see, e.g., Hicks et al, 2012, *Clin Exp Immunol;* 169(3): 199-204, and references therein). The genome of rabies virus codes for five viral proteins. Out of the five, G protein, which is an external surface glycoprotein, forms protrusions that cover the outer surface of the virion envelope and is known to induce neutralizing antibodies. Also, nucleoprotem (N) molecules and the phospho-protein (NS) participate in immune responses. G protein has been the target of antibody developments. For example, therapeutic antibodies against rabies virus are taught in US Patents US70713 19, US6890532, and US Patent 9005624, the contents of each of which are incorporated herein by-reference in their entirety.

[00681] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by rabies virus.

[00682] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat rabies virus related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 11 (SEQ ID NO: 3450-3540).

Therapeutic applications: Tropical Diseases (TPs) and Vector-Borne Diseases

[00683] AAV particles and methods of using the AAV panicles described in the present invention may be used to prevent, manage and/or treat infectious disease. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Tables 10-13 (SEQ ID NO: 3327-3548).

Plasmodium falciparum

[00684] *Plasmodium falciparum (P. falciparum)* is a protozoan parasite belonging to *Plasmodium* parasite family. *P. falciparum* is the main cause of malaria and responsible for nearly all death cases in malaria. *P. falciparum* is released to the human bloodstream through mosquito saliva. The parasite has a high rate of replication and capability to alter. P, falciparum, among other *Plasmodium* parasites, cause malaria, which is a mosquito borne tropical disease. The early stage symptoms include fever, headache, chilis and vomiting. If not treated at the early-stage, malaria can progress to a life-threatening condition involving multiple organs, resulting in skin yellowing, seizures and coma. In children, malaria may cause severe anemia, respirator}; distress in relation to metabolic acidosis, and/or cerebral malaria. The disease is especially dangerous for young children, pregnant women and individuals without immunity to the disease, such as travelers from non-malaria areas. An infection may develop a partial immunity, allowing the following infections to be asymptomatic. According to the WHO, about half of world's population are at risk of malaria. Sub-Saharan Africa carries the highest density of malaria. In 2015, 88% of malaria cases and 90% of malaria deaths was in Sub-Saharan Africa. Malaria is spread by female *Anopheles* mosquitos and caused by 5 different parasite species, out of which *Plasmodium falciparum* is the most prevalent and responsible for the severe cases of malaria.

[00685] Despite tremendous efforts, there is no commercial vaccination for malaria. Traditional treatment for malaria consists of antimalarial medicine tlierapies. such as arteraisinin-based combination therapies, which consists of artemisinin combined with antimalarial drugs such as amodiaquine, lumefaiitrine, mefloquine and siilfadoxine/pynmethainiiie. However, drug resistance has been a serious challenge in malaria treatment. Currently resistance is common for all antimalarial medications apart from artemisinin combination therapy. The cost of artemisinin treatment is high and there remains a need for prevention therapies and improved treatment against malaria.

[00686] Due to the polymorphic nature and high replication rate *of P. Falciparum,* tolerance to malaria is achieved only after years of repeated infections. Antibodies for prevention and treatment of malaria have been developed. For example, antibodies against *P. falciparum* are taught in US Patent US781 1569, in US Patent publication US20150197562 and in international Patent publication WO2014087007, the contents of each of which are incorporated herein by

reference in their entirety. A need for mechanism to deliver constant, effective concentration of malaria antibody for a long period is still in need. Studies by Deal et al. demonstrate results on vectored iramunoprophylaxis delivery of malaria antibodies to mice (see. Deal et al. FNAS, 2014, 111(34), 12528-12532).

[00687] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *P.falciparum.*

[00688] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *P.falciparum* related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 18 (SEQ ID NO: 3915-3971).

Ebola virus

[00689] Genus of Ebola virus includes five viruses. Zaire, Reston, Sudan, Tai Forest and Bundibygyo Ebola viruses, is a negative-sense RNA virus belonging to the family *offiloviridae.* The West Africa outbreak has been associated with Zaire Ebola virus. Hie genome of Ebolavrrus encodes seven genes. The glycoprotein GP gene encodes two distinct gene products: sGP which is a dimeric and secreted glycoprotein and less abundant GP, which is a trirneric-virion attached, membrane embedded envelope glycoprotein and responsible for the virus attachment, fusion and entry during infection. Ebola virus disease is a hemorrhagic fever disease caused. The early symptoms include fever, sore throat, muscular pain, followed by a diarrhea and rash. Eventually the disease will affect the liver and kidney function, and cause internal bleeding. The disease is highly fatal, as about 50% infected individuals die. The Ebola virus is transmitted by direct contact with the blood and body fluids and tissues of an infected person or an animal, most commonly a chimpanzee, gorilla, fruit bat, monkey, forest antelope and porcupine. The disease is also transmitted when handling dead bodies of infected animals or humans. Also, sexual transmittance of the disease has been suggested. The WHO has reported more than 28 000 infections and 11 000 deaths in Ebola virus disease outbreak in West Africa (2014-present). mainly affecting Guinea, Sierra Leone and Liberia.

[00690] As of today, there is no licensed treatment or prevention therapy proven to neutralize the virus. Typically. Ebola virus disease is treated with a good supportive care. A variety of blood, immunological and drug therapies are under investigation, as well as preventive vaccines undergoing evaluations. However, a demand for effective therapies for treatment and prevention of Ebola virus disease remain.

[00691] Viral surface ofGP has been identified as a target for neutralizing antibodies. Antibodies targeting GP of Ebola virus have been taught, e.g. in International Patent publication WO2015127136 and Olal, D., *etal,* 2012. *J Virol* 86 (5), 2809-2816, the contents of each of which are incorporated herein by reference in their entirety.

[00692] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by Ebola virus.

[00693] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Ebola related infections and/or conditions. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 19 (SEQ ID NO: 3972-4024).

Marburg virus

[00694] Marburg virus belongs to *thefiloviridae* family of viruses with coiled, toroid or branched structures with seven proteins. The structure of Marburg vims is similar to Ebola virus, however, the involved antigens are different. The fiioviruses express a single glycoprotein on their surface. The glycoprotein is responsible for the infection, as it is involved in the attachment and entry of the viruses causing infection. Marburg virus disease is a hemorrhaging fever disease caused by Marburg virus. It is highly fatal disease and related to Ebola virus diseases. The early symptoms of the disease include severe headache and malaise. Severe hemorrhagic manifestations in later stages include bleeding from multiple sites. The Marburg virus is transmitted by direct contact with the blood and body fluids and tissues of infected persons or animals, most commonly fruit bats and monkeys. The disease is also transmitted when handling dead the bodies of infected animals or humans. Marburg virus disease is uncommon, but outbreaks typically have a high rate of fatality. According to the WHO, the death rate was as high 80 % in outbreaks of 1998-2000 in Democratic Republic of Congo and 2005 in Angola.

[00695] As of today, there is no preventive or treatment therapy for Marburg virus disease. The current treatment methods include good supportive treatment. The surface glycoprotein has been a target for development of antibodies for Marburg disease vaccines and treatments. For example. International Patent publication WO2015127140, and US Patent publication US20140356354, the contents of which are incorporated herein by reference in their entirety, teach therapeutic antibodies that recognize glycoprotein of fiioviruses for different strains of Marburg, as well as Ebola.

[00696] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by Marburg virus.

[00697] AAV particles and methods of using the AAV panicles described in the present invention may be used to prevent, manage and/or treat Marburg related infections arid/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Tables 3-42. (SEQ ID NO: 2948-9220).

Wesi Nile Virus

[00698] West Nile virus (WNV) is a positive-stranded RNA of the flavivirus genome and member of the Japanese encephalitis serocornpiex of fiaviviruses, (see Throsby, *M.,* J. Viroi. 80 (14), 6982-6992 (2006)). Two lineages of the virus have been identified. The genome of the virus encodes a single polyprotein producing three structural proteins, capsid C, precursor membrane prM and envelope E as well as seven nonstructural proteins. WNV causes mosquito-borne infections with a variety of manifestations. Tough about 80 % of WNV infections are symptomless and not harmful, in certain cases, the disease may lead to fatal neurological diseases. Infection of MNV may lead to a West Nile fever, wiiich causes flu-like symptoms accompanied by high fever, headache, chills, excessive sweating, fatigue, weakness, swollen lymph nodes, and joint pains. Infection by MNV may also occur as cutaneous manifestations, including rashes that may include punctate erythematous, macular and popular eruptions. West Nile infections may also affect the central nervous system resulting in West Nile neuromvasive diseases, including meningitis, encephalitis, meningoencephalitis and po!iomyelitis-like syndrome. These neuroinvasive forms of NWV infections occur in only about 1 % of infections, but they may be life-threatening. WNV is commonly found in Africa, Europe, the Middle East, North America and West Asia. WNV is typically transmitted to humans and other mammals by mosquitos and is maintained in nature in a cycle involving transmission between birds and mosquitoes. WNV is carried by different types of mosquitos, dependent on geographical distribution. Transmission to humans may also occur from birds, horses or other humans.

[00699] As of today, there is no specific treatment or prevention therapy for MNV infections. Current methods of treatment include good supportive care. Due to severity of some of the manifestations, there remains a need for such therapies. Envelope E has been a target of most antibody related studies. Antibodies targeting M and the first non-structural protein have also been investigated. As an example, Thorsby et al, 2006, *J. Virol.* 80 (14), 6982-6992, the contents of which are incorporated herein by reference in their entirety, teaches antibodies binding to E and prM proteins. US Patents US8663950 and US7527973, the contents of each of which are incorporated herein by reference in their entirety, teach antibodies binding to E protein of WNV.

[00700] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by West Nile virus.

[ΘΘ701] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat West Nile vims related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 20 (SEQ ID NO: 3333, 3359, 3393, 3418, 3442, 4025-4137).

Yellow Fever virus

[00702] Yellow fever virus is an enveloped RNA virus belonging to the Flavivirus family. Yellow fever, also known as Yellow Jack, Yellow Plague or Bronze John, is a mosquito-borne viral hemorrhagic disease. In most cases, the symptoms include fever, headache, chills, loss of appetite, nausea, and muscle pain. In some occasions, the disease progresses to a second stage which includes fever accompanied by abdominal pains, liver damage resulting in jaundice, kidney problems and/or bleeding. The disease is spread primarily by *Aedes* and *Haemogogus type* mosquiios. The disease is most typical in tropical environments. Accordrng to the WHO, there are 200 000 annual cases of yellow fever resulting in 30 000 deaths mainly in Africa and Latin America. 90 % of cases occur in Africa.

[00703] Preventive live-attenuated vaccines for yellow fever are available. However, concern related to post-vaccine adverse events has decreased the popularity of the vaccines. The vaccination is not recommended to infants younger than 9 months, pregnant women and individuals with an immune deficiency. As of today, there is no specific treatment for yellow fever. Current methods for treatment involve with supportive care to treat dehydration, respiratory failure and fever. There is a need for improved prevention and treatment therapies against yellow fever virus.

[00704] Envelope E glycoprotein of yellow fever virus has been identified as a potential target for antibody therapies. Neutralizing antibodies for yellow fever virus have been reported by Thibodeaux, B.A. et al,201 2, *Antiviral Res.* 94 (l), 1-8 and Daffis, S. et al., 2005, *Virology,* 337 (2), 262-2.72, the contents of each of which are incorporated herein by reference in their entirety.

[00705] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by yellow fever virus.

[00706] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat yellow fever virus related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention

comprise a nucleic acid sequence encoding at least one of the sequences described in Table 20 (SEQ ID NO: 3333, 3359, 3393, 3418, 3442, 4025-4137).

Japanese encephalitis virus

[00707] Japanese encephalitis virus is an enveloped positive sense single-stranded RNA virus belonging to Fiavivirus family and closely related to St. Lotus encephalitis and West Nile virus. The virus causes Japanese encephalitis, also known as Japanese **B** encephalitis. In majority of cases, the disease is symptomless. However, in less than 1 % of infections, the disease leads to a life-threatening encephalitis. The early stage symptoms include fever, headache and malaise. As the disease progresses into an acute encephalitis, the symptoms include neck rigidity, cachexia, hemiparesis, convulsions and fever, accompanied by lifelong neurological problems such as deafness, and/or mental retardation. The disease is transmitted to humans via mosquitos of the *Culex* species. The virus exists in a transmission cycle between mosquitos, pigs, and water birds. The disease affects 24 countries in the South-East Asia and Western Pacific. According to the WHO, an estimated 68 000 clinical cases are reported annually, with case-fatality rate as high as 30 %. Major outbreaks of the disease occur every 2-15 years.

[00708] The disease may be prevented by a vaccination, most common vaccination being a live attenuated vaccine. In general, the vaccines initially show high effectiveness, but the protection decreases over time. As of today, there is no specific treatment for the disease. Current treatment therapies include good supportive care. There remains a need for longer lasting, improved prevention therapies, and treatment for Japanese encephalitis virus infections.

[00709] Antibodies for treatment of Japanese encephalitis have been developed. For example, Hsieh et al. teach antibodies that target cellular receptors and interrupts their function in fiavivirus infections in US Patent publication US20080292644, the contents of which are incorporated herein by reference in their entirely.

[00710] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by Japanese encephalitis virus.

[00711] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Japanese encephalitis vims related infections and/or conditions. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 20 (SEQ ID NO: 3333, 3359, 3393, 3418, 3442, 4025-4137).

St. Louis encephalitis virus

[00712] St. Louis encephalitis virus is a positive-stranded RNA virus and member of the Fiavivirus family and closely related to Japanese encephalitis virus. St. Louis encephalitis is a

mosquito-borne disease caused by the virus. In majority of cases, the disease is symptomless. However, in less than 1 % of the cases, the disease may lead to encephalitis, which may he life-threatening, especially for the elderly. The early stage symptoms include fever, headache, dizziness, malaise and nausea. If the disease progresses to the central nervous system, symptoms include stiff neck, confusion, disorientation, dizziness, tremor and unsteadiness, and in severe cases coma or even death. St. Louis encephalitis virus is transmitted to humans through Culex mosquitos. The virus exists in a transmission cycle between mosquitos and birds. The disease mainly affects the USA, especially eastern and central states. The disease has also spread to Canada and Mexico.

[00713] As of today, there is no vaccine or specific treatment for St. Louis encephalitis. Current treatment therapies include good supportive care. There is a demand for preventive and treatment therapies for the disease. Neutralizing antibodies for St. Louis encephalitis virus have been reported in Thibodeaux, B.A., et ai, 2012, *Antiviral Res*. 94 (1), 1-8 and Daffis, S. *el al*., 2005, *Virology* 337 (2), 262-272, the contents of which are incorporated herein by reference in their entirety.

[00714] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by St. Louis encephalitis virus.

[00715] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat St. Louis encephalitis virus related infections and/or conditions. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 20 (SEQ ID NO: 3333, 3359, 3393, 3418, 3442, 4025-4137).

Therapeutic applications: Foodborne illness and Gastroenteritis

[00716] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat infectious disease. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Tables 3-9 (SEQ ID NO: 2948-3326).

[00717] Foodborne illnesses, also known as food poisoning, are a common and costly public health problem. The illnesses are typically transmitted by the fecahoral-route. The transmission to humans is by consuming contaminated food or beverage. More than 250 different foodborne diseases, mostly infections caused by viruses, bacteria, parasites or fungus, are identified by the CDC. CDC estimates that approximately 48 million individuals are affected by foodborne illnesses annually in the United States. Gastroenteritis is an inflammation of the gastrointestinal tract involving stomach and small intestine. Gastroenteritis is also caused by an infection caused

by viruses, bacteria, parasites or fungus. The transmission to humans is by person-to-person contact, or by consuming contaminated food or beverage. Foodborne illnesses and gastroenteritis have similar symptoms including diarrhea, vomiting, abdominal pain, dehydration. In some cases, the diseases may require hospitalization or be fatal. Both illnesses are best prevented by proper hand hygiene, proper hygiene while preparing food, treatments to kill bacteria such as pasteurizing, cooking or heating food, and proper methods to store food.

Rotavirus

[00718] Rotavirus is a double-stranded RNA virus belonging to the family of *Reoviridae.* The rotavirus genome consists of 10 segments coding for a single protein, and segment 11 coding for two proteins. The virions are non-enveloped, triple-layered and icosahedral in structure (see. e.g. Aiyegbo *et ah,* 2013, *Flos One* 8, 61101, and references therein). The virus is spread by the fecal-oral-route. Rotavirus is very common especially among infants and young children and spreads easily. Almost all children worldwide are infected with rotavirus by the age of 5, and the disease leads to death of half a million children annually. Rotavirus causes rotavirus gastroenteritis with symptoms including nausea, vomiting, diarrhea and fever. Rotavirus is associated with dehydration. The disease is milder in adults and more severe in young children, infants and the elderly. Though infection does not provide full immunity to the virus, the first infection is typically the most severe in symptoms.

[00719] As of today, there is no specific treatment rotavirus infections. Present treatment includes good supportive care including drinking of fluids to prevent dehydration. In severe cases, the rotavirus gastroenteritis requires hospital care e.g. treatment with intravenous fluids. Vaccines for prevention of the disease have been developed and CDC recommends rotavirus vaccination for infants as part of the routine vaccinations. There remains a need for medical treatment therapies for the infection. Development has been done in the field of antibodies. E.g. Aiyegbo et ai., in Flos One 8, 61101 (2.013, teach antibodies targeting the intermediate capsid layer of VP6 of the triple-layered particle and Frenken et al. teach anti-rotavirus antibodies in US Patent US8105592, the contents of which are incorporated herein by reference in their entirety.

[00720] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by rotavirus.

[00721] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat rotavirus related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 8 (SEQ ID NO: 3286-3310).

Norwalk virus/Norovirus

[00722] Norwalk virus, also known as winter vomiting bug, is the only member of genus norovirus belonging to the family of *Caliciviridae*. Norwalk vims is a single-stranded RNA with three open-reading frames that encode a polyprotem precursor to non-structural proteins, and two polypeptides of different sizes (see e.g. Jiang et ai., 1993, Virology; 195(1):51-61, and references therein). Norwalk virus is spread by the fecal-oral-route. Norwalk virus is extremely contagious and can be transmitted through contaminated food or drink, touching contaminated surfaces or objects or from a contact with an infected individual. The Norwalk virus causes an inflammation of stomach and/or intestines. The symptoms associated with the infection include stomach pain, nausea, vomiting and diarrhea. The disease can be dangerous, especially for your children or young adults. According to CDC, every year 19-21 million infections occur leading to 570-800 deaths in the US.

[00723] As of today, there is no vaccine or specific treatment for Norwalk virus associated gastroenteritis. Antibodies for prevention and treatment of Norwalk virus have been developed. For example, international Patent publication WO2014126921 and WO2014183052, the contents of each of which are incorporated herein by reference in their entirety, teach neutralizing antibodies binding to the polypeptides of Norwalk vims.

[00724] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by Norvvalk virus.

[00725] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat Norwalk virus related infections and/or conditions. As a non-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 7 (SEQ ID NO: 3238-3285).

Campylobacter jejuni

[00726] *Campylobacter jejuni (C. jejuni)* is an oxidase-posiiive, caialase-positive, nonfermentative Gram-negative bacteria with a helical shape. The *C. jejuni* inhabits in the intestinal tract of animals (e.g. poultry, cattle, pigs, sheep, ostriches and shellfish), and in pets (e.g. cats and dogs). The bacteria may be transmitted to humans foodbome, e.g. when eating contaminated food or drink, such as unpasteurized milk. According to the WHO, Campylobacter is the most common cause of gastroenteritis worldwide. *C. jejuni* causes campylohacteriosis infection. The typical symptoms include diarrhea with blood in the feces, abdominal pain, fever, headache, nausea and/or vomiting. The infection may be dangerous to young children, the elderly and individuals with immunodeficiency and is most abundant with malnourished

children. *C. jejuni* infections have been associated with severe long-term complications such as Guillain-Barre Syndrome, inflammatory bowel disease and reactive arthritis (see, e.g., Plaits-Mills and Kosek, 2014, *Ciirr Opin Infect Dis.;* 27(5): 444-450, and references therein).

[00727] Typically, *C. jejuni* infection does not require specific treatment in addition to good supportive care. In more severe cases, in humans and in poultry, the infection has been treated with antibiotics such as fluoroquinoles and macrolides. However, spread of antibiotic-resistant strains is an increasing concern. The treatment with antibiotics is recommended in cases where the bacteria has invaded the intestinal mucosa cell and damaged the tissues, or to eliminate the carrier state. There remains a need for prevention therapies, as well as improved, non-antibiotic, therapies for treatment of the infection. Antibodies targeting *C. jejuni* have been taught e.g. in International Patent publication WO2014063253, the contents of which are incorporated herein by reference in their entirety.

[00728] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *C. Jejuni.*

[00729] AAV particles and methods of using the AAV panicles described in the present invention may be used to prevent, manage and/or treat *C. Jejuni* related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 4 (SEQ ID NO: 3089-3098).

Clostridium difficile

[00730] *Clostridium difficile* bacteria *(C. difficile)* is a Gram- positive, anaerobic spore-forming bacteria belonging to the genus of *Clostridium. C. difficile* inhabits in the soil. *C. difficile* produces toxins, most commonly enterotoxin A and cytotoxin B. Toxins A and B both have a C-terminal receptor-binding domain containing repeating sequences, a central hydrophobic domain and N-termmai glucosyltranferase domain. The toxins bind to the intestinal epithelial cells leading to glucosylation of target Rho GTPases, disruption of the cytoskeleton and cell death. *C. difficile* toxins A and B are a common cause *C. difficile* associated diarrhea and *Clostridium difficile* colitis, which is an inflammation of the large intestine. Typical symptoms of the colitis include flu-like symptoms, bloating, diarrhea, and/or abdominal pain. The disease may lead to dehydration, kidney failure, bowel perforation, toxic megacolon resulting in colon rupture. The elderly and individuals with a weakened immunity are more susceptible to severe and recurring infections which can be life-threatening. *C. difficile* is transmitted by the fecal-oral-route. Due to the ability to form heat-resistant spores, the bacteria is not killed by alcohol-based cleansers or routine surface cleaning. The bacteria may be cultured

on almost any surface and survives in clinical environments, such as hospitals. *C. difficile* is one of the most common and severe healthcare-associated infections. According to CDC, an estimated about half a million infections occur in the United States annually. In 201 1, 29, 000 deaths related to *C. difficile* were reported.

[00731] Currently *C. difficile* infections are treated with antibiotics such as vancomycin and metronidazole. However, increasing an antibiotic-resistance to the bacteria is a concern. Especially in cases of recurring infections, antibiotic treatments have an incomplete response and they disrupt the normal colonic flora. There remains a need for prevention and improved treatment therapies for the infection. Antibodies targeting *C. difficile* have been developed. For example, actoxumab and bezlotoxumab (developed by Medarex Inc. and the University of Massachusetts Medical School) are human monoclonal antibodies targeting *C. difficile* toxin A and toxin B, respectively. The antibodies may be administered as a combination for the prevention of recurring *C. difficile* infection.

|00732] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *C. difficile.*

[00733] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *C. difficile* related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 3 (SEQ ID NO: 2948-3088).

Entamoeba histolytica

[00734] *Entamoeba histolytica (E. histolytica)* is an anaerobic one-celled parasite protozoan belonging to the genus of *Entamoeba.* The active stage of the protozoan exists only in the host and in fresh feces. Cysts survive outside the host in water, soil and food in moist conditions. *E. histolytica* causes an infection called amebiasis, also known as ameobiasis or entamoebiasis. In majority of cases, amebiasis is symptomless. In 10-20 % of individuals infected have symptoms that include loose feces, stomach pain and cramping. The severe more form of amebiasis called amebic dysentery is associated with stomach pain, blood stools and fever. in rare cases, E. *histolytica* invades the liver, forms an abscess and may spread to other parts of the body, such as the lungs or brain. The transmission to humans is mostly via the fecal-oral-route. The disease is typically caused by ingestion of mature cysts in contaminated food, water or via hands. The disease may also be transmitted in close person-to-person contact, e.g. sexual contact. *E. histolytica* infections are most common in tropical areas and especially in poor sanitary

conditions. It is estimated that 50 million cases of amebiasis occur annually, leading to 100, 000 deaths.

[00735] As of today, there are no preventive vaccines for *E. histolytica* infections, though cellular immunity is important for the prevention of liver invasive amebiasis. Amebiasis is typically treated with amebicides, which are medicines *targeting E. histolytica* at specific pails of the body, e.g. the intestine tissue or liver. Optionally, the treatment may involve one or more antibiotics, as well as steroids. However, increasing antibiotic-resistance of *E. histolytica* is a concern. There remains a need for prevention therapy as well as for improved treatments. Antibodies targeting *E. histolytica* are taught in, e.g., 2009, *Infect Immun.*; 77(1): 549-556, and Tachibana et al., 1999, *Clin Diagn Lab Immunol.*;6(3):383~7, the contents of which are incorporated herein by reference in their entire!}7.

[0736] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *E. histolytica.*

|00737] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *E. histolytica* related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 9 (SEQ **ID** NO: 3311-3326).

IIelicobacter pyroli

[0738] *Helicobacter pyroli (II. pyroli)* is a Gram-negative, spiral-shaped microaerophilic bacterium. *H. pyroli* infection is typically asymptomatic and is suggested to be transmitted through the fecal-oral route or oral-oral route. According to CDC, two-thirds of the world's population is infected with H. pyroli. The infection may cause chronic active, chronic, persistent, and atrophic gastritis, duodenal and gastric ulcers and is associated with cancer. CDC reposts 25 million Americans suffering from an ulcer during their lifetime. Typical symptoms associated with ulcer are gnawing or burning pain in the epigastrium, especially between meais. Additional symptoms include nausea, vomiting, loss of appetite, internal bleeding leading to anemia and fatigue.

[00739] Typical treatment for *H. pyroli* infection involves antibiotics. increasing antibiotic resistance and patient noncompliance are major challenges associated with the antibiotic treatment. There remains a need for improved, non-antibiotic, treatment and prevention therapies targeting *H. Pyroli.* Antibodies targeting *H. pyroli* infection have been developed. For example, Boren et al. teach antibodies targeting the BAbA antigen expressed by *H. pyroli* in US patent US8025880, the contents of which are incorporated herein by reference in their entirety.

[00740] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by *H. pyroli.*

[ΘΘ741] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat *H. pyroli* related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 5 (SEQ ID NO: 3099-3196).

Enterotoxin B

[00742] Enterotoxin B is a toxin produced by certain strains of Gram-positive bacteria *Staphylococcus aureus* and is a common cause for food poisoning. *Staphylococcus* species thrive and produce toxins in unrefrigerated meats, dairy, and bakery products. The symptoms associated with enterotoxin B infection are severe diarrhea, nausea and intestinal cramping. The toxin may remain active in the human body after the bacteria has been killed. Enterotoxin B is a so-called superantigen. Superantigens are toxins that may activate T cells by forming a budge between a MHC II on antigen presenting cells (APCs) and the T cell receptors (TCR). Due to binding of enterotoxin B, the T cells release large amount of cytokines leading to an inflammation and gastroenteritis. Though enterotoxin B infection is typically not life threatening, enterotoxin B has been identified as a potential chemical and biological warfare agent.

[00743] As of today, there is no specific prevention or treatment for enterotoxin B infection. Antibodies that neutralize enterotoxin B have been investigated, e.g. as described in US Patent. US8895704.

[00744] In some embodiment, methods of the present invention may be used to prevent, manage and/or treat infections and complications caused by enterotoxin B.

[00745] AAV particles and methods of using the AAV particles described in the present invention may be used to prevent, manage and/or treat enterotoxin B related infections and/or conditions. As anon-limiting example, the AAV particles of the present invention comprise a nucleic acid sequence encoding at least one of the sequences described in Table 5 (SEQ ID NO: 3099-3196).

V. KITS AND DEVICES

Kits

[00746] In one embodiment, the invention provides a variety of kits for conveniently and/or effectively carrying out methods of the present invention. Typically, kits will comprise sufficient amounts and/or numbers of components to allow a user to perform multiple treatments of a subject(s) and/or to perform multiple experiments.

[00747] Any of the AAV particles of the present invention may be comprised in a kit. In some embodiments, kits may further include reagents and/or instructions for creating and/or synthesizing compounds and/or compositions of the present invention. In some embodiments, kits may also include one or more buffers. In some embodiments, kits of the invention may include components for making protein or nucleic acid arrays or libraries and thus, may include, for example, solid supports.

|00748] In some embodiments, kit components may be packaged either in aqueous media or in !yophihzed form. The container means of the kits will generally include at least one vial, test tube, flask, bottle, syringe or other container means, into which a component may be placed, and preferably, suitably aliquoted. Where there is more than one kit component, (labeling reagent and label may be packaged together), kits may also generally contain second, third or other additional containers into which additional components may be separately placed. In some embodiments, kits may also comprise second container means for containing sterile, pharmaceutically acceptable buffers and/or other diluents. In some embodiments, various combinations of components may be comprised in one or more vial. Kits of the present invention may also typically include means for containing compounds and/or compositions of the present invention, e.g., proteins, nucleic acids, and any other reagent containers in close confinement for commercial sale. Such containers may include injection or blow-molded plastic containers into which desired vials are retained.

[00749] In some embodiments, kit components are provided in one and/or more liquid solutions. in some embodiments, liquid solutions are aqueous solutions, with sterile aqueous solutions being particularly preferred. in some embodiments, kit components may be provided as dried powder(s). When reagents and/or components are provided as dry powders, such powders may be reconstituted by the addition of suitable volumes of solvent. In some embodiments, it is envisioned that solvents may also be provided in another container means. In some embodiments, labeling dyes are provided as dried powders. In some embodiments, it is contemplated that 10, 20, 30, 40, 50, 60. 70, 80, 90, 100, 120. 120, 130, 140, 150. 160, 170, 180, 190, 200, 300, 400, 500, 600, 700, 800, 900, 1000 micrograms or at least or at most those amounts of dried dye are provided in kits of the invention. In such embodiments, dye may then be resuspended in any suitable solvent, such as DMSO.

[ΘΘ750] In some embodiments, kits may include instructions for employing kit components as well the use of any other reagent not included in the kit. instructions may include variations that may be implemented.

Devices

[00751] In one embodiment, the AAV particles may delivered to a subject using a device to deliver the AAV particles and a head fixation assembly. The head fixation assembly may be, but is not limited to, any of the head fixation assemblies sold by MRI interventions. As a non-limiting example, the head fixation assembly may be any of the assemblies described in US Patent Nos. 8,099,150, 8,548,569 and 9,031,636 and international Patent Publication Nos. **WO201** 108495 and **WO2014014585,** the contents of each of which are incorporated by reference in their entireties. A head fixation assembly may be used in combination with an MRI compatible drill such as, but not limited to, the MRi compatible drills described in International Patent Publication No. WO2013181008 and US Patent Publication No. US20130325012, the contents of which are herein incorporated by reference in its entirety.

[00752] In one embodiment, the AAV particles may be delivered using a method, system and/or computer program for positioning apparatus to a target point on a subject to deliver the AAV particles. As anon-limiting example, the method, system and/or computer program may be the methods, systems and/or computer programs described in US Patent No. 8, 340,743, the contents of which are herein incorporated by reference in its entirety. The method may include: determining a target point in the body and a reference point, wherein the target point and the reference point define a planned trajectory line (PTL) extending through each; determining a visualization plane, wherein the PTL intersects the visualization plane at a sighting point; mounting the guide device relative to the body to move with respect to the PTL, wherein the guide device does not intersect the visualization plane; determining a point of intersection (GPP) between the guide axis and the visualization plane; and aligning the GPP with the sighting point in the visualization plane.

[00753] In one embodiment, the AAV particles may be delivered using an MRT-guided device. Non-limiting examples of MRI-guided devices are described in US Patent Nos. 9,055,884, 9,042,958, 8,886,288, 8,768,433, 8,396,532, 8,369,930, 8,374,677 and 8,175,677 and US Patent Application No. IJS20 140024927 the contents of each of which are herein incorporated by reference in their entireties. As a non-limiting example, the MRI-guided device may be able to provide data in real time such as those described in US Patent Nos. 8,886,288 and 8,768,433, the contents of each of which is herein incorporated by reference in its entirety. As another non-limiting example, the MRI-guided device or system may be used with a targeting cannula such as the systems described in US Patent Nos. 8,175.677 and 8,374,677, the contents of each of which are herein incorporated by reference in their entireties. As yet another non-limiting example, the MRI-guided device includes a trajectory guide frame for guiding an interventional device as described, for example, in US Patent No. 9,055,884 and US Patent Application No.

US20140024927, the contents of each of which are herein incorporated by reference in their entireties.

[00754] In one embodiment, the AAV particles may be delivered using an MRI-compatible tip assembly. Non-limiting examples of MRI-compatible tip assemblies are described in US Patent Publication No. US20140275980, the contents of which is herein incorporated by reference in its entirety.

[00755] In one embodiment, the AAV particles may be delivered using a cannula which is MRI-compatible. Non-hmiting examples of MRI-compatible cannulas include those taught in International Patent Publication No. WO2011130107, the contents of which are herein incorporated by reference in its entirety.

[00756] In one embodiment, the AAV particles may be delivered using a catheter which is MRI-compatible. Non-limiting examples of MRI-compatihle catheters include those taught in International Patent Publication No. WO2012116265, US Patent Publication No. 8,825,133 and US Patent Publication No. LiS20140024909, the contents of each of which are herein incorporated by reference in their entireties.

[00757] In one embodiment, the AAV particles may be delivered using a device with an elongated tubular body and a diaphragm as described in US Patent Publication Nos. US20140276582 and US20140276614, the contents of each of which are herein incorporated by reference in their entireties.

[00758] In one embodiment, the AAV particles may be delivered using an MRI compatible localization and/or guidance system such as, but not limited to, those described in US Patent Publication Nos. US20150223905 and US20150230871, the contents of each of which are herein incorporated by reference in their entireties. As anon-limiting example, the MRI compatible localization and/or guidance systems may comprise a mount adapted for fixation to a patient, a targeting cannula with a lumen configured to attach to the mount so as to be able to controilably translate in at least three dimensions, and an elongate probe configured to snugly advance via slide and retract in the targeting cannula lumen, the elongate probe comprising at least one of a stimulation or recording electrode.

[00759] In one embodiment, the AAV particles may be delivered to a subject using a trajectory frame as described in US Patent Publication Nos. **US20150031982** and IJS20I40066750 and international Patent Publication Nos. WO2015057807 and WO2014039481, the contents of each of which are herein incorporated by reference in their entireties.

[00760] In one embodiment, the AAV particles may be delivered to a subject using a gene gun.

VI. DEFINITIONS

[00761] At various places in the present specification, subsiituents of compounds of the present disciosiire are disclosed in groups or in ranges. It is specifically intended that the present disclosure include each and every individual subcombination of the members of such groups and ranges.

[00762] *About:* As used herein, the term "about" means +/- 10% of the recited value.

[00763] *Adeno-associated virus:* The term "adeno-associated virus" or "AAV" as used herein refers to members of the dependovirus genus comprising any particle, sequence, gene, protein, or component derived therefrom.

[00764] *AAV Particle;* As used herein, an "AAV particle" is a virus which comprises a viral genome with at least one payload region and at least one 1TR region. AAV vectors of the present disclosure may be produced recombinantly and may be based on adeno-associated virus (AAV) parent or reference sequences. AAV particle may be derived from any serotype, described herein or known in the art, including combinations of serotypes (i.e., "pseudolyped" AAV) or from various genomes (e.g., single stranded or self-complementary). In addition, the AAV particle may be replication defective and/or targeted.

[00765] *Activity:* As used herein, the term "activity" refers to the condition in which things are happening or being done. Compositions of the invention may have activity and this activity may involve one or more biological events.

[00766] *Administered in combination:* As used herein, the term '"administered in combination" or "combined administration" means that two or more agents are administered to a subject at the same time or within an interval such that there may be an overlap of an effect of each agent on the patient. In some embodiments, they are administered within about 60, 30, 15, 10, 5, or 1 minute of one another. In some embodiments, the administrations of the agents are spaced sufficiently closely together such that a combinatorial *(e.g.,* a synergistic) effect is achieved.

[00767] *Amelioration:* As used herein, the term "amelioration" or '"ameliorating" refers to a lessening of seventy of at least one indicator of a condition or disease. For example, in the context of neurodegeneration disorder, amelioration includes the reduction of neuron loss.

|00768] *Animal:* As used herein, the term "animal" refers to any member of the animal kingdom. In some embodiments, "animal" refers to humans at any stage of development. in some embodiments, "animal" refers to non-human animals at any stage of development. In certain embodiments, the non-human animal is a mammal *(e.g.,* a rodent, a mouse, a rat, a rabbit, a monkey, a dog, a cat, a sheep, cattle, a primate, or a pig). In some embodiments, animals include, but are not limited to, mammals, birds, reptiles, amphibians, fish, and worms. in some embodiments, the animal is a transgenic animal, genetically -engineered animal, or a clone.

[00769] *Antibody:* As used herein, the term "antibody" is referred to in the broadest sense and specifically covers various embodiments including, but not limited to monoclonal antibodies, polyclonal antibodies, multispecific antibodies (e.g. bispecific antibodies formed from at least two intact antibodies), and antibody fragments (e.g., diabodies) so long as they exhibit a desired biological activity (e.g., "functional"). Antibodies are primarily amino-acid based molecules but may also comprise one or more modifications (including, but not limited to the addition of sugar moieties, fluorescent moieties, chemical tags, etc.). Non-limiting examples of antibodies or fragments thereof include V_H and V_L domains, scFvs, Fab, Fab', F(ab')2, Fv fragment, diabodies, linear antibodies, single chain antibody molecules, multispecifie antibodies, bispecific antibodies, intrabodies, monoclonal antibodies, polyclonal antibodies, humanized antibodies, codon-optimized antibodies, tandem scFv antibodies, bispecific T-cell engagers, mAb2 antibodies, chimeric antigen receptors (CAR), tetravalent bispecific antibodies, bi©synthetic antibodies, native antibodies, miniaturized antibodies, unihodies, maxibodies, antibodies to senescent cells, antibodies to conformers, antibodies to disease specific epitopes or antibodies to innate defense molecules.

[00770] *Antibody-based composition:* As used herein, "antibody-based" or "antibody-derived" compositions are monomelic or muiti-meric polypeptides which comprise at least one amino-acid region derived from a known or parental antibody sequence and at least one ammo acid region derived from a non-antibody sequence, e.g., mammalian protein.

[00771] *Approximately:* As used herein, the term "approximately" or "about," as applied to one or more values of interest, refers to a value that is similar to a stated reference value. In certain embodiments, the term "approximately" or "about" refers to a range of values that fall within 25%, 20%, 19%, 18%, 17%, 16%, 15%, 14%, 13%, 12%, 11%, 10%, 9%, 8%, 7%, 6%, 5%, 4%, 3%, 2%, 1%, or less in either direction (greater than or less than) of the stated reference value unless otherwise stated or otherwise evident from the context (except where such number would exceed 100% of a possible value).

[00772] *Associated with:* As used herein, the terms "associated with," "conjugated," "linked," "attached," and "tethered," when used with respect to two or more moieties, means that the moieties are physically associated or connected with one another, either directly or via one or more additional moieties that serves as a linking agent, to form a structure that is sufficientiy stable so that the moieties remain physically associated under the conditions in which the structure is used, e.g., physiological conditions. An "association" need not be strictly through direct covalent chemical bonding. It may also suggest ionic or hydrogen bonding or a

hybridization based connectivity sufficiently stable such that the "associated" entities remain physically associated.

[00773] *Bifimctional:* As used herein, the term 'Afunctional' ` refers to any substance, molecule or moiety which is capable of or maintains at least two functions. The functions may effect the same outcome or a different outcome. The structure that produces the function may be the same or different.

|00774] *Biocompatible:* As used herein, the term "biocompatible" means compatible with living cells, tissues, organs or systems posing little to no risk of injury, toxicity or rejection by the immune system.

[ΘΘ775] *Biodegradable:* As used herein, the term "biodegradable" means capable of being broken down into innocuous products by the action of living things.

[Θ0776] *Biologically active:* As used herein, the phrase "biologically active" refers to a characteristic of any substance that has activity in a biological system and/or organism. For instance, a substance that, when administered to an organism, has a biological effect on that organism, is considered to be biologically active. In particular embodiments, an AAV particle of the present invention may be considered biologically active if even a portion of the encoded payioad is biologically active or mimics an activity considered biologically relevant.

[00777] *Capsid:* As used herein, the term "capsid" refers to the protein shell of a virus particle.

[00778] *Chimeric antigen receptor (CAR):* As used herein, the term "chimeric antigen receptor" or "CAR" refers to an artificial chimeric protein comprising at least one antigen specific targeting region {ASTR), a transmembrane domain and an intracellular signaling domain, wherein the antigen specific targeting region comprises a full-length antibody or a fragment thereof. As a non-limiting example the ASTR of a CAR may be any of the antibodies listed in Tables 3-42, antibody-based compositions or fragments thereof. Any molecule that is capable of binding a target antigen with high affinity can be used in the ASTR of a CAR. The CAR may optionally have an extracellular spacer domain and/or a co-stimulatory domain. A CAR may also be used to generate a cytotoxic cell carrying the CAR.

|00779] *Complementary and. substantially complementary:* As used herein, the term "complementary" refers to the ability of polynucleotides to form base pairs with one another. Base pairs are typically formed by hydrogen bonds between nucleotide units in antiparallel polynucleotide strands. Complementary polynucleotide strands can form base pair in the Watson-Crick manner (e.g., A to T, A to U, C to G), or in any other manner that allows for the formation of duplexes. As persons skilled in the art are aware, when using RNA as opposed to DNA, uracil rather than thymine is the base that is considered to be complementary to adenosine.

However, when a U is denoted in the context of the present invention, the ability to substitute a T is implied, unless otherwise stated. Perfect complementarity or 100% complementarity refers to the situation in which each nucleotide unit of one polynucleotide strand can form hydrogen bond with a nucleotide unit of a second polynucleotide strand. Less than perfect complementarity refers to the situation in which some, but not ail nucleotide units of two strands can form hydrogen bond with each other. For example, for two 20-mers, if only two base pairs on each strand can form hydrogen bond with each other, the polynucleotide strands exhibit 10% complementarity. In the same example, if 18 base pairs on each strand can form hydrogen bonds with each other, the polynucleotide strands exhibit 90% complementarity. As used herein, the term "substantially complementary" means that the siRNA has a sequence (e.g., in the antisense strand) which is sufficient to bind the desired target mRNA, and to trigger the RNA silencing of the target mRNA.

[00780] *Compound:* Compounds of the present disclosure include all of the isotopes of the atoms occurring in the intermediate or final compounds. "Isotopes" refers to atoms having the same atomic number but different mass numbers resulting from a different number of neutrons in the nuclei. For example, isotopes of hydrogen include tritium and deuterium.

[00781] The compounds and salts of the present disclosure can be prepared in combination with solvent or water molecules to form solvates and hydrates by routine methods.

[00782] *Comprehensive Positional Evolution (CPE™):* As used herein, the term "comprehensive positional evolution" refers to an antibody evolution technology that allows for mapping of the effects of amino acid changes at every position along an antibody variable domain's sequence. Tins comprehensive mutagenesis technology can be used to enhance one or more antibody properties or characteristics.

[00783] *Comprehensive Protein Synthesis (CPS™):* As used herein, the term "comprehensive protein synthesis" refers to a combinatorial protein synthesis technology that can be used to optimize antibody properties or characteristics by combining the best properties into a new, high-performance antibody.

[00784] *Conditionally active:* As used herein, the term "conditionally active" refers to a mutant or variant of a wild-type polypeptide, wherein the mutant or variant is more or less active at physiological conditions than the parent polypeptide. Further, the conditionally active polypeptide may have increased or decreased activity at aberrant conditions as compared to the parent polypeptide. A conditionally active polypeptide may be reversibly or irreversibly inactivated at normal physiological conditions or aberrant conditions.

[00785] *Conserved:* As used herein, the term "conserved" refers to nucleotides or amino acid residues of a polynucleotide sequence or polypeptide sequence, respectively, that are those that occur unaltered in the same position of two or more sequences being compared. Nucleotides or amino acids that are relatively conserved are those that are conserved amongst more related sequences than nucleotides or amino acids appearing elsewhere in the sequences.

[00786] In some embodiments, two or more sequences are said to be "completely conserved" if they are 100% identical to one another. In some embodiments, two or more sequences are said to be "highly conserved" if they are at least 70% identical, at least 80% identical, at least 90% identical, or at least 95% identical to one another. In some embodiments, two or more sequences are said to be "highly conserved" if they are about 70% identical, about 80% identical, about 90% identical, about 95%, about 98%, or about 99% identical to one another. In some embodiments, two or more sequences are said to be "conserved" if they are at least 30% identical, at least 40% identical, at least 50% identical, at least 60% identical, at least 70% identical, at least 80% identical, at least 90% identical, or at least 95% identical to one another. In some embodiments, two or more sequences are said to be "conserved" if they are about 30% identical, about 40% identical, about 50% identical, about 60% identical, about 70% identical, about 80% identical, about 90% identical, about 95% identical, about 98% identical, or about 99% identical to one another. Conservation of sequence may apply to the entire length of a polynucleotide or polypeptide or may apply to a portion, region or feature thereof.

[00787] *Control Elements:* As used herein, "control elements", "regulatory control elements" or "regulatory sequences" refers to promoter regions, polyadenylation signals, transcription termination sequences, upstream regulatory domains, origins of replication, internal ribosome entry sites ("IRES"), enhancers, and the like, which provide for the replication, transcription and translation of a coding sequence in a recipient cell. Not all of these control elements need ahvays be present as long as the selected coding sequence is capable of being replicated, transcribed and/or translated in an appropriate host cell.

[00788] *Controlled Release:* As used herein, the term "controlled release" refers to a pharmaceirtical composition or compound release profile that conforms to a particular pattern of release to effect a therapeutic outcome.

[00789] *Cytostatic:* As used herein, "cytostatic" refers to inhibiting, reducing, suppressing the growth, division, or multiplication of a cell (e.g., a mammalian cell (eg., a human cell)), bacterium, vims, fungus, protozoan, parasite, prion, or a combination thereof.

[00790] *Cytotoxic:* As used herein, "cytotoxic" refers to killing or causing injurious, toxic, or deadly effect on a cell *{e.g.,* a mammalian cell *{e.g.,* a human cell)), bacterium, virus, fungus, protozoan, parasite, prion, or a combination thereof.

[00791] *Delivery:* As used herein, "delivery" refers to the act or manner of delivering an AAV particle, a compound, substance, entity, moiety, cargo or payload.

[00792] *Delivery Agent:* As used herein, "deliver)' agent" refers to any substance which facilitates, at least in part, the *in vivo* deliver}' of an AAV particle to targeted ceils.

[00793] *Destabilized:* As used herein, the term "destabie," "destabilize," or "destabilizing region" means a region or molecule that is less stable than a starting, wild-type or native form of the same region or molecule.

[00794] *Detectable label:* As used herein, "detectable label" refers to one or more markers, signals, or moieties which are attached, incorporated or associated with anotber entity that is readily detected by methods known in the art including radiography, fluorescence, ehennliimmeseence, enzymatic activity, absorbance and the like. Detectable labels include radioisotopes, fluorophores, chromophores, enzymes, dyes, metal ions, ligands such as biotin, avidin, streptavidin and haptens, quantum dots, and the like. Detectable labels may be located at any position in the peptides or proteins disclosed herein. They may be within the amino acids, the peptides, or proteins, or located at the N- or C- termini.

[00795] *Digest:* As used herein, the term "digest" means to break apart into smaller pieces or components. When referring to polypeptides or proteins, digestion results in the production of peptides.

[00796] *Distal:* As used herein, the term "distal" means situated away from the center or away from a point or region of interest.

[00797] *Dosing regimen: As* used herein, a "dosing regimen" is a schedule of administration or physician determined regimen of treatment, prophylaxis, or palliative care.

[00798] *Encapsulate:* As used herein, the term "encapsulate" means to enclose, surround or encase.

[00799] *Engineered:* As used herein, embodiments of the invention are "engineered" when they are designed to have a feature or property, whether structural or chemical, that varies from a starting point, wild type or native molecule.

[00800] *Effective Amount:* As used herein, the term "effective amount" of an agent is that amount sufficient to effect beneficial or desired results, for example, clinical results, and, as such, an "effective amount" depends upon the context in which it is being applied. For example, in the context of administering an agent that treats cancer, an effective amount of an agent is, for

example, an amount sufficient to achieve treatment, as defined herein, of cancer, as compared to the response obtained without administration of the agent.

[00801] *Epitope:* As used herein, an "epitope" refers to a surface or region on a molecule that is capable of interacting with a biomolecule. For example, a protein may contain one or more amino acids, e.g., an epitope, which interacts with an antibody, e.g., a biomolecule. In some embodiments, when referring to a protein or protein module, an epitope may comprise a linear stretch of amino acids or a three-dimensional structure formed by folded amino acid chains.

[00802] EvoMap™: As used herein, an EvoMap™ refers to a map of a polypeptide, wherein detailed informatics are presented about the effects of single amino acid mutations within the length of the polypeptide and their influence on the properties and characteristics of that polypeptide.

[00803] *Expression:* As used herein, "expression" of a nucleic acid sequence refers to one or more of the following events: (1) production of an RNA template from a DNA sequence *(e.g.*, by transcription); (2) processing of an RNA transcript *(e.g.*, by splicing, editing, 5' cap formation, and/or 3' end processing); (3) translation of an RNA into a polypeptide or protein; and (4) post-translational modification of a polypeptide or protein.

[00804] *Feature:* As used herein, a "feature" refers to a characteristic, a property, or a distinctive element.

[00805] *Formulation:* As used herein, a "formulation" includes at least one AAV particle and a delivery agent.

[00806] *Fragment:* A "fragment," as used herein, refers to a portion. For example, fragments of proteins may comprise polypeptides obtained by digesting full-length protein isolated from cultured cells.

[00807] *Functional:* As used herein, a "functional" biological molecule is a biological molecule in a form in which it exhibits a property and/or activity by which it is characterized.

[00808] *Gene expression:* The term "gene expression" refers to the process by which a nucleic acid sequence undergoes successful transcription and in most instances translation to produce a protein or peptide. For clarity, when reference is made to measurement of "gene expression", this should be understood to mean that measurements may be of the nucleic acid product of transcription, e.g., RNA or mRNA or of the amino acid product of translation, e.g., polypeptides or peptides. Methods of measuring the amount or levels of RNA, mRNA, polypeptides and peptides are well known in the art.

[00809] *Homology:* As used herein, the term "homology" refers to the overall relatedness between polymeric molecules, *e.g.* between polynucleotide molecules *(e.g.* DNA molecules

and/or RNA molecules) and/or between polypeptide molecules. In some embodiments, polymeric molecules are considered to be "homologous" to one another if their sequences are at least 25%, 30%. 35%, 40%, 45%. 50%, 55%, 60%, 65%. 70%, 75%, 80%. 85%, 90%, 95%, or 99% identical or similar. The term "homologous" necessarily refers to a comparison between at least two sequences (polynucleotide or polypeptide sequences). In accordance with the invention, two polynucleotide sequences are considered to be homologous if the polypeptides they encode are at least about 50%, 60%, 70%, 80%, 90%, 95%, or even 99% for at least one stretch of at least about 20 amino acids. In some embodiments, homologous polynucleotide sequences are characterized by the ability to encode a stretch of at least 4-5 uniquely specified amino acids. For polynucleotide sequences less than 60 nucleotides in length, homology is determined by the ability to encode a stretch of at least 4-5 uniquely specified amino acids. In accordance with the invention, two protein sequences are considered to be homologous if the proteins are at least about 50%, 60%, 70%, 80%, or 90% identical for at least one stretch of at least about 20 amino acids.

[00810] *Heterologous Region:* As used herein the term "heterologous region" refers to a region which would not be considered a homologous region.

[00S11] *Homologous Region:* As used herein the term "homologous region" refers to a region which is similar in position, structure, evolution origin, character, form or function.

[00812] *Identity.* As used herein, the term "identity" refers to the overall relatedness between polymeric molecules, *e.g.,* between polynucleotide molecules *(e.g.* DNA molecules and/or RNA molecules) and'or between polypeptide molecules. Calculation of the percent identity of two polynucleotide sequences, for example, can be performed by aligning the two sequences for optimal comparison purposes *(e.g.,* gaps can be introduced in one or both of a first and a second nucleic acid sequences for optimal alignment and non-identical sequences can be disregarded for comparison purposes). In certain embodiments, the length of a sequence aligned for comparison purposes is at least 30%, at least 40%, at least 50%, at least 60%, at least 70%, at least 80%, at least 90%, at least 95%, or 100% of the length of the reference sequence. The nucleotides at corresponding nucleotide positions are then compared. When a position in the first sequence is occupied by the same nucleotide as the corresponding position in the second sequence, then the molecules are identical at that position. The percent identity between the two sequences is a function of the number of identical positions shared by the sequences, taking into account the number of gaps, and the length of each gap, which needs to be introduced for optimal alignment of the two sequences. The comparison of sequences and determination of percent identity between two sequences can be accomplished using a mathematical algorithm. For example, the

percent identity between two nucleotide sequences can be determined using methods such as those described in Computational Molecular Biology, Lesk, A. M., ed., Oxford University Press, New York, 1988; Biocomputing: informatics and Genome Projects. Smith. D. W., ed., Academic Press, New York, 1993; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; Computer Analysis of Sequence Data, Part I. Griffin, A. M., and Griffin, H. G., eds., Humana Press, New Jersey, 1994; and Sequence Analysis Primer, Gribskov, M. and Devereux, I , eds., M Stockton Press, New York, 1991; each of which is incorporated herein by reference. For example, the percent identity between two nucleotide sequences can be determined using the algorithm of Meyers and Miller (CABIOS, 1989. 4:11-17), which has been incorporated into the ALIGN program (version 2.0) using a PAM120 weight residue table, a gap length penalty of 12 and a gap penalty of 4. The percent identity between two nucleotide sequences can. alternatively, be determined using the GAP program in the GCG software package using an NWSgapdna.CMP matrix. Methods commonly employed to determine percent identity between sequences include, but are not limited to those disclosed in Canllo, H., and Lipman, D., SIAM J Applied Math.. 48:1073 (1988); incorporated herein by reference. Techniques for determining identity are codified in publicly available computer programs. Exemplary computer software to determine homology between two sequences include, but are not limited to, GCG program package, Devereux, j., *et al.*, *Nucleic Acids Research,* 12(1), 387 (1984)). BLASTP, BLASTN, and FASTA Aitschui, S. F. *et al.*, *J. Mo!ec. Biol,* 215, 403 (1990)).

|00813] *Inhibit expression of a gene:* As used herein, the phrase 'inhibit expression of a gene" means to cause a reduction in the amount of an expression product of the gene. The expression product can be an RNA transcribed from the gene *(e.g.*, an mRNA) or a polypeptide translated from an mRNA transcribed from the gene. Typically, a reduction in the level of an mRNA results in a reduction in the level of a polypeptide translated therefrom. The level of expression may be determined using standard techniques for measuring mRNA or protein.

[00814] *In vitro:* As used herein, the term *"in vitro"* refers to events that occur in an artificial environment, *e.g.*, in a test tube or reaction vessel, in cell culture, in a Petri dish, *etc.*, rather thai withm an organism *(e.g.*, animal, plant, or microbe).

[00815] *In vivo:* As used herein, the term *"in vivo"* refers to events that occur within an organism *(e.g.*, animal, plant, or microbe or cell or tissue thereof).

[ΘΘ816] *Isolated:* As used herein, the term "isolated" refers to a substance or entity that has been separated from at least some of the components with which it was associated (whether in nature or in an experimental setting). Isolated substances may have varying levels of purity in

reference to the substances from which they have been associated. isolated substances and/or entities may be separated from at least about 10%, about 20%, about 30%, about 40%, about 50%, about 60%, about 70%, about 80%, about 90%, or more of the other components with which they were initially associated. In some embodiments, isolated agents are more than about 80%, about 85%, about 90%, about 91%. about 92%, about 93%, about 94%, about 95%. about 96%, about 97%, about 98%. about 99%, or more than about 99% pure. As used herein, a substance is "pure" if it is substantially free of other components.

[00817] *Substantially isolated:* By "substantially isolated" is meant that a substance is substantially separated from the environment in which it was formed or detected. Partial separation can include, for example. a composition enriched in the substance or AAV particles of the present disclosure. Substantial separation can include compositions containing at least about 50%, at least about 60%, at least about 70%, at least about 80%. at least about 90%, at least about 95%. at least about 97%, or at least about 99% by weight of the compound of the present disclosure, or salt thereof. Methods for isolating compounds and their salts are routine in the art.

[00818] *Linker:* As used herein "linker" refers to a molecule or group of molecules which connects two molecules, such as a **VH** chain and **VL** chain or an antibody. A linker may be a nucleic acid sequence connecting two nucleic acid sequences encoding two different polypeptides. The linker may or may not be translated. The linker may be a cleavabie linker.

[00819] *MicroRNA (miRNA) binding site:* As used herein, a microRNA (miRNA) binding site represents a nucleotide location or region of a nucleic acid transcript to which at least the "seed" region of a miRNA binds.

[00820] *Modified:* As used herein "modified" refers to a changed state or structure of a molecule of the invention. Molecules may be modified in many ways including chemically, structurally, and functionally.

[00821] *Naturally Occurring:* As used herein, "naturally occurring" or "wild-type" means existing in nature without artificial aid, or involvement of the hand of man.

[00822] *Non-human vertebrate:* As used herein, a "non-human vertebrate" includes all vertebrates except *Homo sapiens,* including wild and domesticated species. Examples of non-hitman vertebrates include, but are not limited to, mammals, such as alpaca, banteng. bison, camel, cat, cattle, deer, dog, donkey, gayal, goat, guinea pig, horse, llama, mule, pig, rabbit, reindeer, sheep water buffalo, and yak.

[00823] *Off-target:* As used herein, "off target" refers to any unintended effect on any one or more target, gene, or cellular transcript.

[00824] *Open reading frame:* As used herein, "open reading frame" or "ORF" refers to a sequence which does not contain a stop codon in a given reading frame.

[00825] *Operably linked:* As used herein, the phrase "operably linked" refers to a functional connection between two or more molecules, constructs, transcripts, entities, moieties or the like.

[00826] *Particle:* As used herein, a "particle" is a virus comprised of at least two components, a protein capsid and a polynucleotide sequence enclosed within the capsid.

[00827] *Patient:* As used herein, "patient" refers to a subject who may seek or be in need of treatment, requires treatment, is receiving treatment, will receive treatment, or a subject who is under care by a trained professional for a particular disease or condition.

[00828] *Payload:* As used herein, "payload" or "payload region" refers to one or more polynucleotides or polynucleotide regions encoded by or within a viral genome or an expression product of such polynucleotide or polynucleotide region, e.g., a transgene, a polynucleotide encoding a polypeptide or multi-polypeptide or a modulatory nucleic acid or regulatory nucleic acid.

[00829] *Payload construct:* As used herein, "payload construct" is one or more polynucleotide regions encoding or comprising a payload that is flanked on one or both sides by an inverted terminal repeat (ITR) sequence. The payload construct is a template that is replicated in a viral production cell to produce a viral genome.

[00830] *Payload construct vector:* As used herein, "payload construct vector" is a vector encoding or comprising a payload construct, and regulatory regions for replication and expression in bacterial cells.

[00831] *Payload construct expression vector:* As used herein, a "payload construct expression vector" is a vector encoding or comprising a payload construct and which further comprises one or more polynucleotide regions encoding or comprising components for viral expression in a viral replication cell.

[00832] *Peptide:* As used herein, "peptide" is less than or equal to 50 amino acids long, e.g., about 5, 10, 15, 20, 25, 30, 35, 40, 45, or 50 amino acids long.

[00833] *Pharmaceutically acceptable:* The phrase "pharmaceutically acceptable" is employed herein to refer to those compounds, materials, compositions, **and/or** dosage forms which are, within the scope of sound medical judgment, suitable for use in contact with the tissues of human beings and animals without excessive toxicity, irritation, allergic response, or other problem or complication, commensurate with a reasonable benefit/risk ratio.

[00834] *Pharmaceutically acceptable excipients:* The phrase "pharmaceutically acceptable excipient," as used herein, refers any ingredient other than the compounds described herein (for

example, a vehicle capable of suspending or dissolving the active compound) and having the properties of being substantially nontoxic and noninflammatory in a patient. Excipients may include, for example: antiadherents. antioxidants, binders, coatings, compression aids, disintegrates, dyes (colors), emollients, emulsifiers, fillers (diluents), film formers or coatings, flavors, fragrances, giidants (flow enhancers), lubricants, preservatives, printing inks, sorbenis, suspensing or dispersing agents, sweeteners, and waters of hydration. Exemplary excipients include, but are not limited to: butyiated hydroxytoluene (BHT), calcium carbonate, calcium phosphate (dibasic), calcium stearaie, croscarmeiiose, crossiinked polyvinyl pyrroiidone, citric acid, crospovidone, cysteine, etbylcelhilose, gelatin, hydroxypropyl cellulose, hydroxypropyl metbyiceilulose, lactose, magnesium stearate, maltitoi, mannitol, methionine, methyl cellulose, methyl paraben, microcrystalline cellulose, polyethylene glycol, polyvinyl pyrroiidone, povidone, pregelatinized starch, propyl paraben, retinyl palmitate, shellac, silicon dioxide, sodium carboxyraethyl cellulose, sodium citrate, sodium starch giycoiate, sorbitol, starch (com), stearic acid, sucrose, talc, titanium dioxide, vitamin A, vitamin E, vitamin C, and xyhtol.

[00835] *Pharmaceutically acceptable salts:* The present disclosure also includes pharmaceutically acceptable salts of the compounds described herein. As used herein, "pharmaceutically acceptable salts" refers to derivatives of the disclosed compounds wherein the parent compound is modified by converting an existing acid or base moiety to its salt form (e.g., by reacting the free base group with a suitable organic acid). Examples of pharmaceutically acceptable salts include, but are not limited to, mineral or organic acid salts of basic residues such as amines; alkali or organic salts of acidic residues such as carboxylic acids; and the like. Representative acid addition salts include acetate, acetic acid, adipate, alginate, ascorbate, aspartate, henzenesulfonate, benzene sulfonic acid, benzoate, bisulfate, borate, butyrate, campborate, camphorsulfonate, citrate, cyclopentanepropionate, digiuconate, dodecyisulfate. ethanesuifonate, fumarate, giucoheptonate, glycerophosphate, hennsulfate, lieptonate, hexanoate, hydrobromide, hydrochloride, hydroiodide, 2-hydroxy-ethanesuifonate, lactobionate, lactate, laurate, lauryi sulfate, raalate, maleate, malonate, methanesulfonate, 2-naphthalenesulfonate, nicotinate, nitrate, oieate, oxalate, palmitate, pamoate, pectinate, persuifate, 3-phenylpropionate, phosphate, picrate, pivalate, propionate, stearate, succinate, sulfate, tartrate, thiocyanate, toiuenesuifonate. undecanoate, valerate salts, and the like. Representative alkali or alkaline earth metal salts include sodium, lithium, potassium, calcium, magnesium, and the like, as well as nontoxic ammonium, quaternary ammonium, and amine cations, including, but not limited to ammonium, tetramethylammonium, tetraethylammonium, methylamine, dimethylamine, trimethylamine, triethyiamine, ethylarnine. and the like. The pharmaceutically acceptable salts

of the present disclosure include the conventional non-toxic salts of the parent compound formed, for example, from non-toxic inorganic or organic acids. The pharmaceutically acceptable salts of the present disclosure can be synthesized from the parent compound which contains a basic or acidic moiety by conventional chemical methods. Generally, such salts can be prepared by reacting the free acid or base forms of these compounds with a stoichiometric amount of the appropriate base or acid in water or in an organic solvent, or in a mixture of the two; generally, nonaqueous media like ether, ethyl acetate, ethanol, isopropanoi, or acetonitnle are preferred. Lists of suitable salts are found in *Remington's Pharmaceutical Sciences,* 17th ed., Mack Publishing Company, Easton, Pa., 198.5, p. 1418, *Pharmaceutical Salts: Properties, Selection, and Use,* P.H. Stahl and C.G. Werrnuth (eds.), Wiley-VCH, 2008, and Berge et al., *Journal of Pharmaceutical. Science,* 66, 1-19 (1977), each of which is incorporated herein by reference in its entirety.

[00836] *Pharmaceutically acceptable solvate:* The term "pharmaceutically acceptable solvate," as used herein, means a compound of the invention wherein molecules of a suitable solvent are incorporated in the crystal lattice. A suitable solvent is physiologically tolerable at the dosage administered. For example, solvates may be prepared by crystallization, recrystallization, or precipitation from a solution that includes organic solvents, water, or a mixture thereof. Examples of suitable solvents are ethanol, water (for example, mono-, di-, and tri-hydrates), *N*-methylpyrrolidinone (NMP), dimethyl sulfoxide (DMSO), *N,N'*-dimethylformamide (DMF), *N,N'*-diraethylacetami de (DMAC), 1,3-diraethyl-2-imidazolidinone (DMEU), 1,3-diraethyl-3,4,5,6-tetrahydro-2-(lH)-pyriinidinone (DMPU), acetonitnle (ACN), propylene glycol, ethyl acetate, benzyl alcohol, 2-pyrrolidone, benzyl benzoate, and the like. When water is the solvent, the solvate is referred to as a "hydrate."

[00837] *Pharmacokinetic:* As used herein, "pharmacokinetic" refers to any one or more properties of a molecule or compound as it relates to the determination of the fate of substances administered to a living organism. Pharmacokinetics is divided into several areas including the extent and rate of absorption, distribution, metabolism and excretion. This is commonly referred to as ADME where: (A) Absorption is the process of a substance entering the blood circulation; (D) Distribution is the dispersion or dissemination of substances throughout the fluids and tissues of the body; (M) Metabolism (or Biotransformation) is the irreversible transformation of parent compounds into daughter metabolites; and (E) Excretion (or Elimination) refers to the elimination of the substances from the body. In rare cases, some drugs irreversibly accumulate in body tissue.

[00838] *Physicochemical:* As used herein, "physicochemical" means of or relating to a physical and/or chemical property.

[00839] *Preventing:* As used herein, the term "preventing" refers to partially or completely delaying onset of an infection, disease, disorder and/or condition; partially or completely delaying onset of one or more symptoms, features, or clinical manifestations of a particular infection, disease, disorder, and/or condition; partially or completely delaying onset of one or more symptoms, features, or manifestations of a particular infection, disease, disorder, and/or condition, partially or completely delaying progression from an infection, a particular disease, disorder and/or condition; and/or decreasing the risk of developing pathology associated with the infection, the disease, disorder, and/or condition.

[00840] *Proliferate:* As used herein, the term "proliferate" means to grow, expand or increase or cause to grow, expand or increase rapidly. "Proliferative" means having the ability to proliferate. "Anti-proliferative" means having properties counter to or inapposite to proliferative properties.

[00841] *Prophylactic:* As used herein, "prophylactic" refers to a therapeutic or course of action used to prevent the spread of disease.

[00842] *Prophylaxis:* As used herein, a "prophylaxis" refers to a measure taken to maintain health and prevent the spread of disease.

[00843] *Protein of interest:* As used herein, the terms "proteins of interest" or "desired proteins" include those provided herein and fragments, mutants, variants, and alterations thereof

[00844] *Proximal:* As used herein, the term "proximal" means situated nearer to the center or to a point or region of interest.

[00845] *Purified:* As used herein, "purify," "purified," "purification" means to make substantially pure or clear from unwanted components, material defilement, admixture or imperfection. "Purified" refers to the state of being pure. "Purification" refers to the process of making pure.

[00846] *Region:* As used herein, the term "region" refers to a zone or general area. In some embodiments, when referring to a protein or protein module, a region may comprise a linear sequence of amino acids along the protein or protein module or may comprise a three-dimensional area, an epitope and/or a cluster of epitopes. In some embodiments, regions comprise terminal regions. As used herein, the term "terminal region" refers to regions located at the ends or termini of a given agent. When referring to proteins, terminal regions may comprise N- and/or C-termini. N-termini refer to the end of a protein comprising an amino acid with a free amino group. C-termini refer to the end of a protein comprising an amino acid with a free

carboxyl group. N- and/or C-terminal regions may there for comprise the N- and/or C-termini as well as surrounding amino acids. In some embodiments. N- and/or C-terminal regions comprise from about 3 amino acid to about 30 amino acids, from about 5 amino acids to about 40 amino acids, from about 10 amino acids to about 50 amino acids, from about 2.0 amino acids to about 100 amino acids and/or at least 100 ammo acids. In some embodiments, N-ierminal regions may comprise any length of amino acids that includes the N-terminus, but does not include the C-termmus. In some embodiments, C-terminal regions may comprise any length of amino acids, which include the Oterminus, but do not comprise the N-terminus.

[00847] In some embodiments, when referring to a polynucleotide, a region may comprise a linear sequence of nucleic acids along the polynucleotide or may comprise a three-dimensional area, secondary structure, or tertiary structure. In some embodiments, regions comprise terminal regions. As used herein, the term "terminal region" refers to regions located at the ends or termini of a given agent. When referring to polynucleotides, terminal regions may comprise 5' and 3' termini. 5' termini refer to the end of a polynucleotide comprising a nucleic acid with a free phosphate group. 3` termini refer to the end of a polynucleotide comprising a nucleic acid with a free hydroxy] group. 5' and 3` regions may there for comprise the 5" and 3' termini as well as surrounding nucleic acids. in some embodiments, 5" and 3' terminal regions comprise from about 9 nucleic acids to about 90 nucleic acids, from about 15 nucleic acids to about 120 nucleic acids, from about 30 nucleic acids to about 150 nucleic acids, from about 60 nucleic acids to about 300 nucleic acids and/or at least. 300 nucleic acids. In some embodiments, 5' regions may comprise any length of nucleic acids that includes the 5` terminus, but does not include the 3` terminus. In some embodiments, 3' regions may comprise any length of nucleic acids, which include the 3` terminus, but does not. comprise the 5' terminus.

[00848] *RNA or RNA molecule;* As used herein, the term "RNA" or "RNA molecule" or "ribonucleic acid molecule" refers to a polymer of ribonucleotides; the term "DMA" or "DNA molecule" or "deoxyribonucleic acid molecule" refers to a polymer of deoxy ribonucleotides. DNA and RNA can be synthesized naturally, e.g., by DNA replication and transcription of DNA. respectively; or be chemically synthesized. DNA and RNA can be smgie-stranded (i.e., ssRNA or ssDNA, respectively) or multi-stranded (e.g., double stranded, i.e., dsRNA and dsDNA, respectively). The terra "rnRNA" or "messenger RNA", as used herein, refers to a single stranded RNA that encodes the amino acid sequence of one or more polypeptide chains.

[ΘΘ849] *Sample:* As used herein, the term "sample" or "biological sample" refers to a subset of its tissues, cells or component parts (e.g. body fluids, including but not limited to blood, mucus, lymphatic fluid, synovial fluid, cerebrospinal fluid, saliva, amniotic fluid, amniotic cord blood,

urine, vaginal fluid and semen). A sample further may include a homogenate, lysaie or extract prepared from a whole organism or a subset of its tissues, ceils or component parts, or a traction or portion thereof, including but not limited to, for example, plasma, serum, spinal fluid, lymph fluid, the external sections of the skin, respíratory, intestinal, and genitourinary tracts, tears, saliva, milk, blood cells, tumors, organs. A sample further refers to a medium, such as a nutrient broth or gel, which may contain cellular components, such as proteins or nucleic acid molecule.

[00850] *Self-complementary viral particle:* As used herein, a "self-complementary viral particle" is a particle comprised of at least two components, a protein capsid and a polynucleotide sequence encoding a self-complementary genome enclosed within the capsid.

[00851] *Signal Sequences:* As used herein, the phrase "signal sequences" refers to a sequence which can direct the transport or localization of a protein.

[00852] *Single unit dose:* As used herein, a "single unit dose" is a dose of any therapeutic administered in one dose/at one time/single route/single point of contact, i.e., single administration event. In some embodiments, a single unít dose is provided as a discrete dosage form (e.g., a tablet, capsule, patch, loaded syringe, vial, etc.).

[00853] *Similarity.* As used herein, the term "similarity" refers to the overall relatedness between polymeric molecules, *e.g.* between polynucleotide molecules *(e.g.* **DNA** molecules and/or RNA molecules) and/or between polypeptide molecules. Calculation of percent similarity of polymeric molecules to one another can be performed in the same manner as a calculation of percent identity, except that calculation of percent similarity takes into account conservative substitutions as is understood in the art.

[00854] *Split dose:* As used herein, a "split dose" is the division of single unit dose or total daily dose into two or more doses.

[00855] *Stable:* As used herein "stable" refers to a compound that is sufficiently robust to survive isolation to a useful degree of purity from a reaction mixture, and preferably capable of formulation into an efficacious therapeutic agent.

[00856] *Stabilized:* As used herein, the term "stabilize", "stabilized," "stabilized region" means to make or become stable.

[00857] *Subject:* As used herein, the term "subject" or "patient" refers to any organism to which a composition in accordance with the invention may be administered, *e.g.,* for experimental, diagnostic, prophylactic, and/or therapeutic purposes. Typical subjects include animals *(e.g.,* mammals such as mice, rats, rabbits, non-human primates, and humans) and/or plants.

[00858] *Substantially:* As used herein, the term "substantially" refers to the qualitative condition of exhibiting total or near-total extent or degree of a characteristic or property of interest. One of ordinary skill in the biological arts will understand that biological and chemical phenomena rarely, if ever, go to completion and'or proceed to completeness or achieve or avoid an absolute result. The term "substantially" is therefore used herein to capture the potential lack of completeness inherent in many biological and chemical phenomena.

[00859] *Substantially equal:* As used herein as it relates to time differences between doses, the term means plus/minus 2%.

[00860] *Substantially simultaneously:* As used herein and as it relates to plurality of doses, the term means within 2 seconds.

[00861] *Suffering from:* An individual who is "suffering from" a disease, disorder, and/or condition has been diagnosed with or displays one or more symptoms of a disease, disorder, and/or condition.

[00862] *Susceptible to:* An individual who is "susceptible to" a disease, disorder, and/or condition has not been diagnosed with and/or may not exhibit symptoms of the disease, disorder, and/or condition but harbors a propensity to develop a disease or its symptoms. In some embodiments, an individual who is susceptible to a disease, disorder, and/or condition (for example, cancer) may be characterized by one or more of the following: (1) a genetic mutation associated with development of the disease, disorder, and/or condition; (2) a genetic polymorphism associated with development of the disease, disorder, and/or condition; (3) increased and/or decreased expression and/or activity of a protein and/or nucleic acid associated with the disease, disorder, and/or condition; (4) habits and/or lifestyles associated with development of the disease, disorder, and/or condition; (5) a family history of the disease, disorder, and/or condition; and (6) exposure to and/or infection with a microbe associated with development of the disease, disorder, and/or condition. In some embodiments, an individual who is susceptible to a disease, disorder, and/or condition will develop the disease, disorder, and-'or condition. In some embodiments, an individual who is susceptible to a disease, disorder, and'or condition will not develop the disease, disorder, and/or condition.

[00863] *Sustained release:* As used herein, the term "sustained release" refers to a pharmaceutical composition or compound release profile that conforms to a release rate over a specific period of time.

[00864] *Synthetic:* The term "synthetic" means produced, prepared, and-'or manufactured by the hand of man. Synthesis of polynucleotides or polypeptides or other molecules of the present invention may be chemical or enzymatic.

[00865] *Targeting:* As used herein, '-targeting" means the process of design and selection of nucleic acid sequence that will hybridize to a target nucleic acid and induce a desired effect.

[00866] *Targeted Cells:* As used herein, "targeted cells" refers to any one or more cells of interest. The cells may be found *in vitro, in vivo, in situ* or in the tissue or organ of an organism. The organism may be an animal, preferably a mammal, more preferably a human and most preferably a patient.

[00867] *Therapeutic Agent:* The term "therapeutic agent" refers to any agent that, when administered to a subject, has a therapeutic, diagnostic, and'or prophylactic effect and/or elicits a desired biological and/or pharmacological effect.

[00868] *Therapeutically effective amount:* As used herein, the term "therapeutically effective amount" means an amount of an agent to be delivered *(e.g.,* nucleic acid, drug, therapeutic agent, diagnostic agent, prophylactic agent, *etc.)* that is sufficient, when administered to a subject suffering from or susceptible to an infection, disease, disorder, and/or condition, to treat, improve symptoms of, diagnose, prevent, and/or delay the onset of the infection, disease, disorder, and/or condition. In some embodiments, a therapeutically effective amount is provided in a single dose. In some embodiments, a therapeutically effective amount is administered in a dosage regimen comprising a plurality of doses. Those skilled in the art will appreciate that in some embodiments, a unit dosage form may be considered to comprise a therapeutically effective amount of a particular agent or entity if it comprises an amount that is effective when administered as part of such a dosage regimen.

|00869] *Therapeutically effective outcome:* As used herein, the term "therapeutically effective outcome" means an outcome that is sufficient in a subject suffering from or susceptible to an infection, disease, disorder, and/or condition, to treat, improve symptoms of, diagnose, prevent, and/or delay the onset of the infection, disease, disorder, and/or condition.

[00870] *Total daily dose:* As used herein, a "total daily dose" is an amount given or prescribed in 24 hr period. It may be administered as a single unit dose.

[00871] *Transfection:* As used herein, the term "transfection" refers to methods to introduce exogenous nucleic acids into a cell. Methods of transfection include, but are not limited to, chemical methods, physical treatments and canonic lipids or mixtures.

[00872] *Treating:* As used herein, the term "treating" refers to partially or completely alleviating, ameliorating, improving, relieving, delaying onset of, inhibiting progression of, reducing severity of, and/or reducing incidence of one or more symptoms or features of a particular infection, disease, disorder, and/or condition. For example, "treating" cancer may refer to inhibiting survival, growth, and/or spread of a tumor. Treatment may be administered to

a subject who does not exhibit signs of a disease, disorder, and/or condition and/or to a subject who exhibits only early signs of a disease, disorder, and/or condition for the purpose of decreasing the risk of developing pathology associated with the disease, disorder, and/or condition.

[00873] *Unmodified:* As used herein, "unmodified" refers to any substance, compound or molecule prior to being changed in any way. Unmodified may, but does not always, refer to the wild type or native form of a biomoiecule. Molecules may undergo a series of modifications whereby each modified molecule may serve as the "unmodified" starting molecule for a subsequent modif cation.

[00874] *Vector.* As used herein, a "vector" is any molecule or moiety which transports, transduces or otherwise acts as a earner of a heterologous molecule. Vectors of the present invention may he produced recorabinantly and may be based on and/or may comprise adeno-associated virus (AAV) parent or reference sequence. Such parent or reference AAV sequences may serve as an original, second, third or subsequent sequence for engineering vectors. In non-limiting examples, such parent or reference AAV sequences may comprise any one or more of the following sequences: a polynucleotide sequence encoding a polypeptide or multi-polypeptide, which sequence may be wild-type or modified from wild-type and which sequence may encode full-length or partial sequence of a protein, protein domain, or one or more subunits of a protein; a polynucleotide comprising a modulatory or regulatory nucleic acid which sequence may be wild-type or modified from wild-type; and a transgene that may or may not be modified from wild-type sequence . These AAV sequences may serve as either the "donor" sequence of one or more codons (at the nucleic acid level) or amino acids (at the polypeptide level) or "acceptor" sequences of one or more codons (at the nucleic acid level) or amino acids (at the polypeptide level).

[00875] *Viral genome:* As used herein, a "viral genome" or "vector genome" is a polynucleotide comprising at least one inverted terminal repeat (ITR) and at least one encoded payioad. A viral genome encodes at least one copy of the payload.

|00876] Described herein are compositions, methods, processes, kits and devices for the design, preparation, manufacture and/or formulation of AAV particles. In some embodiments, payioads, such as but not limited to AAV polynucleotides, may be encoded by payioad constructs or contained within piasmids or vectors or recombinant adeno-associated viruses (AAVs).

[00877] The details of one or more embodiments of the invention are set forth in the accompanying description below. Although any materials and methods similar or equivalent to

those described herein can be used in the practice or testing of the present invention, the preferred materials and methods are now described. Other features, objects and advantages of the invention will be apparent from the description. In the description, the singular forms also include the plural unless the context clearly dictates otherwise. Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this invention belongs. In the case of conflict, the present description will control.

[00878] The present invention is further illustrated by the following non-limiting examples.

VII. EXAMPLES

EXAMPLE 1. Production and Purification of AAV particles

[ΘΘ879] AAV particles described herein may be produced using methods known in the art, such as, for example, triple transfection or baculovirus mediated virus production. Any suitable permissive or packaging cell known in the art. may be employed to produce the vectors. Mammalian cells are often preferred. Also preferred are trans-complementing packaging cell lines that provide functions deleted from a replication-defective helper virus, e.g., 293 cells or other E1a trails-complementing cells.

[ΘΘ880] The gene cassette may contain some or all of the parvovirus (e.g., AAV) cap and rep genes. Preferably, however, some or all of the cap and rep functions are provided in trans by introducing a packaging vector(s) encoding the capsid and/or Rep proteins into the cell. Most preferably, the gene cassette does not encode the capsid or Rep proteins. Alternatively, a packaging cell line is used that is stably transformed to express the cap and/or rep genes

[00881] Recombinant AAV virus particles are, in some cases, produced and purified from culture supematants according to the procedure as described in US20160032254, the contents of which are incorporated by reference. Production may also involve methods known in the art including those using 293T cell, sf9 insect cells, triple transfection or any suitable production method.

[Θ0882] In some cases, 293 cells are transfected with CaP04 with plasrmds required for production of AAV, i.e., AAV2 rep, an adenoviral helper construct and a ITR flanked transgene cassette. The AAV2 rep piasmid also contains the cap sequence of the particular virus being studied. Twenty-four hours after transfection, which occurs in serum containing DMEM, the medium is replaced with fresh medium with or without serum. Three (3) days after transfection, a sample is taken from the culture medium of the 293 adherent cells. Subsequently cells are scraped and transferred into a receptacle. After centrifugation to remove cellular pellet, a second sample is taken from the supernatant after scraping. Next ceil lysis is achieved by three

consecutive freeze-ihaw cycles (-80C. to 37(1). Cellular debris is removed and sample 3 is taken from the medium. The samples are quantified for AAV particles by DNase resistant genome titration by Taqman.TM. PGR. The total production yield from such a transfection is equal to the particle concentration from sample 3.

[00883] AAV vector titers are measured according to genome copy number (genome particles per milliliter). Genome particle concentrations are based on Taqman.RTM. PGR of the vector DNA as previously reported (Clark et al. (1999) Hum. Gene Ther., 10:1031-1039; Veldwijk et al. (2002) Mol. Ther., 6:272-278).

EXAMPLE 2. Tissue specific **expression**

[00884] To evaluate the expression of various encoded antibody payloads in tissues, a series of AAV particles carrying the encoded antibody sequences driven by a panel of ubiquitous and tissue-specific promoters are made. These particles are administered to the specific tissue, e.g.. intramuscularly, via an appropriate route, e.g., a single injection in the gastrocnemius muscle and expression is monitored to determine the relative expression potential of the payload as well as of each promoter in this target tissue. Measurement of antibody production is performed using standard techniques, for example by ELISA.

[00885] In some cases, the cytomegalovirus immediate early promoter (CMV), chimeric chicken-beta-actin (CAG), and ubiquitin C (UBC), CBA, H1 promoters provide robust expression.

EXAMPLE 3, Generation of antibodies

Antibody production by hyhridoma technology

[00886] Host animals (e.g. mice, rabbits, goats, and llamas) are immunized by an injection with an antigenic protein to elicit lymphocytes that specifically bind to the antigen. Lymphocytes are collected and fused with immortalized cell lines to generate hybridomas. Hybridomas are cultured in a suitable culture medium that is enriched with appropriate selection agents to promote growth.

[00887] Antibodies produced by the cultured hybridomas are subjected to analysis to determine binding specificity of the antibodies for the target antigen. Once antibodies with desirable characteristics are identiiiecl corresponding hybridomas are subcioned through limiting dilution procedures and grown by standard methods. Antibodies produced by these cells are isolated and purified using standard immunoglobulin purification procedures.

Recombinant antibody production

[00888] Recombinant antibodies are produced using heavy and light chain variable region cDNA sequences selected from hybridomas or from other sources. Sequences encoding antibody

vanable domains expressed by hybridomas are determined by extracting RNA molecules from antibody-producing hybridoma cells and producing cDNA by reverse transcriptase polymerase chain reaction (PGR). PCR is used to amplify cDNA using primers specific for heavy and light chain sequences. PCR products are then subcloned into plasmids for sequence analysis. Antibodies are produced by insertion of resulting variable domain sequences into expression vectors.

Recombinant antibodies are also produced using phage display technology. Target antigens are screened, in vitro, using phage display libraries having millions to billions of phage particles expressing unique single chain variable fragments (scFvs) on their viral coat. Precipitated phage particles are analyzed and sequences encoding expressed scFvs are determined. Sequences encoding antibody variable domains and/or CDRs are inserted into expression vectors for antibody production.

Recombinant antibodies are further produced using yeast surface display technology, wherein antibody variable domain sequences are expressed on the cell surface of *Saccharomyces cerevisiae.* Recombinant antibodies are developed by displaying the antibody fragment of interest as a fusion to e.g. Aga2p protein on the surface of the yeast, where the protein interacts with proteins and small molecules in a solution. scFvs with affinity towards desired receptors are isolated from the yeast surface using magnetic separation and flow cytometry. Several cycles of yeast surface display and isolation will be done to attain scFvs with desired properties through directed evolution.

EXAMPLE 4. Optimization of the encoded Antibody

[00889] To design an optimal framework for the expression of an antibody, the heavy and light chains of several antibodies separated by an F2A self-processing peptide sequence are cloned into a mammalian expression vector under the control of the CMV promoter. 293T cells or any suitable cell line transfected with these vectors exhibit secretion of human IgG into the culture supernatant that is then detected by EL!SA.

[ΘΘ890] To increase expression, the antibody chains and/or the processing peptide are codon optimized for mammalian expression. In some instances, a furin cleavage site at the N-terminus is inserted for better processing.

[00891] To improve secretion of the antibody, the endogenous signal sequences are replaced with a sequence which may or may not be codon optimized, derived from any gene. In some cases, the human growth hormone signal sequence is used. Any of the heavy, light or both chains may be driven by any signal sequence, whether the same or different. Antibody expression is confirmed using standard immunohistochermcal techniques, including EL1SA.

EXAMPLE 5. Vectored Antibodies

[00892] Viral genomes are designed for AAV delivery of antibodies to cells. The viral genome comprises a payioad region and at least one inverted terminal repeat (1TR) region. The payioad region may optionally encode regulatory- elements e.g., a promoter region, an intronic region, or a polyadenyiatiott sequence. The payioad region comprises a sequence encoding one or more polypeptides selected from the group consisting of those listed in Table 3. An exemplary payioad region comprises a sequence encoding an antibody heavy chain, a region encoding an antibody light chain and a region encoding a linker connecting the heavy and light chain sequences or polypeptides before further processing. A promoter is selected to target the desired tissue or for desired regulation of expression, or both. The promoter may be selected from human EFla, CMV, CBA, and its derivative CAG, GUSB, UBC, or any other promoter known to one with skill in the art, or combinations thereof. The 5' and 3' ITRs may or may not be of the same serotype as the capsid of the AAV particle.

|00893] Payioad regions may optionally encode a linker between light and heavy antibody chain sequences or polypeptides. Sequence encoding linkers are derived from an internal ribosome entry site (IRES; SEQ ID NO: 899), foot and mouth disease virus 2A (F2A; SEQ ID NO: 900), porcine teschovirus-l virus 2A (P2A; SEQ ID NO: 901), a form cleavage site (F; SEQ ID NO: 902), or a 5xG4S (SEQ ID NO: 9221 encoded by SEQ ID NO: 903) linker sequence. In various payioad regions, the order of heavy and light chains is alternated with respect to 5' to 3' direction. Payloads are further designed to encode protein signal sequences (to aid in protein processing, localization, and/or secretion) as well as an untranslated poly A tail

[00894] Each viral genome is then incorporated into an AAV cloning vector to create payioad expression vectors.

[00895] The payioad expression vectors are expressed in e.g. Expi 293 cells. The supernatants are collected and expressed antibodies are purified using protein A/G beads. Supernatants are diluted with a loading buffer and applied to a column prepared with A/G beads. Unbound proteins are washed tlirough with loading buffer. Elution buffer is added to the column, fractions collected, and fractions containing proteins of interest are identified with absorption spectroscopy technique, pooled together, and neutralized. Western blotting techniques are used to identify payioad regions producing the antibody proteins of interest. Purified antibodies are then tested for their affinity to their specific target by e.g. ELISA essay technique and antibodies with the highest affinity are identified and selected.

[00896] Finally, the rAAVs are produced using, for example, HEK293T cells. The cells are transfeeted simultaneously with the viral genome of the present invention, a viral genome encoding helper proteins and a viral genome encoding replication and capsid proteins.

EXAMPLE 6. **In** Vivo **Expression** and **efficacy of antibody payloads**

[00897] To determine the efficacy or comparative expression of encoded antibodies, dose-dependent expression is determined at a series of time points. Samples from mice treated with AAV particles encoding antibodies or luciferase at various levels are examined for expression using standard techniques such as nucleic acid analyses for RNA levels, protein analyses for antibody levels and compared to the expression of the luciferase control.

EXAMPLE 7. Treatment of infections disease

[00898] AAV particles of the current invention encoding an antibody are administered to a patient who has been diagnosed with an infectious disease, disorder or condition. The purpose of the treatment may be aimed to manage the disease, prevent or slow the progression of the disease, treat the symptoms associated with the disease and/or cure the disease.

[00899] The AAV particles may be administered through an intramuscular injection to the skeletal muscle. The administration may include one or more injections over a period of time. The level and distribution of AAV particles and antibody expression is monitored by standard diagnostic techniques known in the art. Such diagnostic techniques include e.g. (e.g. from blood, urine, or saliva), cerebrospinal fluid (CSF) testing, or any other testing useful for monitoring antibody levels in the body.

[00900] Additionally, the progression of the disease and the health of the patient is monitored by standard diagnostic techniques known in the art. Such techniques may include diagnostic imaging (e.g. X-ray, MRA scans. Ultrasound scans, PET scans. Nuclear scans, mammography), biopsy, laboratory tests (e.g. from blood, mine, or saliva), cerebrospinal fluid (CSF) testing, vital signs, clinical tests (cognitive, motor or reflex tests) and other relevant techniques. Treatment with the AAV particles may results in cure of the non-infectious disease, slowing down or stabilizing the progression of the disease, or have no effect on the progression of the disease. Additionally, the treatment may reduce severity of one or more symptoms associated with the disease, eliminate one or more symptoms associated with the disease or have no effect on the symptoms.

EXAMPLE 8. Treatment of HIV or AIDS

[00901] AAV particles of the current invention encoding an antibody are administered to a patient who has been diagnosed with HIV or AIDS. The purpose of the treatment may be aimed

to manage the disease, prevent or slow the progression of the disease, treat the symptoms associated with the disease and/or cure the disease.

[00902] The AAV particles may be administered through an intramuscular injection to the skeletal muscle. The administration may include one or more injections over a period of time. The level and distribution of AAV particles and antibody expression is monitored by standard diagnostic techniques known in the art. Such diagnostic techniques include e.g. (e.g. from blood, urine, or saliva), cerebrospinal fluid (CSF) testing, or any other testing useful for monitoring antibody levels in the body.

[00903] Additionally, the progression of the disease and the health of the patient is monitored by standard diagnostic techniques known in the art. Such techniques may include diagnostic imaging (e.g. X-ray, MRA scans, Ultrasound scans, PET scans, Nuclear scans, mammography), biopsy, laboratory tests (e.g. from blood, urine, or saliva), cerebrospinal fluid (CSF) testing, vital signs, clinical tests (cognitive, motor or reflex tests) and other relevant techniques. Treatment with the AAV particles may results in cure of the non-infectious disease, slowing down or stabilizing the progression of the disease, or have no effect on the progression of the disease. Additionally, the treatment may reduce severity of one or more symptoms associated with the disease, eliminate one or more symptoms associated with the disease or have no effect on the symptoms.

VIII, EQUIVALENTS AND SCOPE

[ΘΘ904] Those skilled in the art will recognize, or be able to ascertain using no more than routine experimentation, many equivalents to the specific embodiments in accordance with the invention described herein. The scope of the present invention is not intended to be limited to the above Description, but rather is as set forth in the appended claims.

[00905] In the claims, articles such as "a," "an," and "the" may mean one or more than one unless indicated to the contrary or otherwise evident from the context. Claims or descriptions that include "or" between one or more members of a group are considered satisfied if one, more than one, or all of the group members are present in, employed in, or otherwise relevant to a given product or process unless indicated to the contrary or otherwise evident from the context. The invention includes embodiments in which exactly one member of the group is present in, employed in, or otherwise relevant to a given product or process. The invention includes embodiments in which more than one, or the entire group members are present in, employed in, or otherwise relevant to a given product or process.

[00906] It is also noted that the term "comprising" is intended to be open and permits but does not require the inclusion of additional elements or steps. When the term "comprising" is used herein, the term "consisting of" is thus also encompassed and disclosed.

[00907] Where ranges are given, endpoints are included. Furthermore, it is to be understood that unless otherwise indicated or otherwise evident from the context and understanding of one of ordinary skill in the art, values that are expressed as ranges can assume any specific value or subrange within the stated ranges in different embodiments of the invention, to the tenth of the unit of the lower limit of the range, unless the context clearly dictates otherwise.

[00908] In addition, it is to be understood that any particular embodiment of the present invention that falls within the prior art may be explicitly excluded from any one or more of the claims. Since such embodiments are deemed to be known to one of ordinary skill in the art, they may be excluded even if the exclusion is not set forth explicitly herein. Any particular embodiment of the compositions of the invention (e.g., any antibiotic, therapeutic or active ingredient; any method of production; any method of use; etc.) can be excluded from any one or more claims, for any reason, whether or not related to the existence of prior art.

[00909] It is to be understood that the words which have been used are words of description rather than limitation, and that changes may be made within the purview of the appended claims without departing from the true scope and spirit of the invention in its broader aspects.

[00910] While the present invention has been described at some length and with some particularity with respect to the several described embodiments, it is not intended that it should be limited to any such particulars or embodiments or any particular embodiment, but it is to be construed with reference to the appended claims so as to provide the broadest possible interpretation of such claims in view of the prior art and, therefore, to effectively encompass the intended scope of the invention.

CLAIMS

1. An AAV particle comprising **a capsid** and a viral genome, said viral genome comprising at least one inverted terminal repeat **(ITR)** region and a **payload** region, said payload region comprising a **regulatory** sequence **operably** linked to at least a first nucleic acid segment, said first nucleic acid segment encoding one or more polypeptides selected from the group consisting of any member given in Tables 3-42 and fragments thereof.
2. The AAV particle of claim 1, wherein the capsid is selected from the group of serotypes consisting of Table 1.
3. The AAV particle **of** claim 2, wherein the **regulatory** sequence comprises a promoter.
4. The AAV particle of claim 3, wherein the promoter is selected from the group consisting **of human** elongation factor 1**a-subunit** (EFla), cytomegalovirus (CMV) immediate-early enhancer and/or promoter, chicken β**-actin** (CBA) and its derivative CAG, β glucuronidase **(GUSB),** or **ubiquitin** C **(UBC).** Tissue-specific expression elements can be used to restrict expression **to** certain cell types such as, but not limited to, muscle specific promoters, B cell promoters, monocyte promoters, leukocyte promoters, macrophage promoters, pancreatic acinar cell promoters, endothelials ceil promoters, lung tissue promoters, astrocyte promoters, or nervous system promoters **which** can be used to restrict expression to neurons, astrocytes, or oligodendrocytes.
5. The AAV particle of claim 1, wherein the viral genome is single stranded.
6. The AAV particle of claim 1, wherein the viral genome is **self-complementary.**
7. The AAV particle of claim 1, wherein **at** least one region of the viral genome is codon-optimized.
8. The AAV particle of claim 7, wherein the first nucleic acid segment is **codon-optimized.**
9. The AAV particle of any **of claims** 1-8, wherein the first nucleic acid segment encodes one or more polypeptides selected from the group consisting of an antibody heavy chain, an antibody light chain, a linker, and combinations thereof.
10. The AAV particle **of claim** 9, wherein any of the polypeptides encoded by first nucleic acid segment **of the** payload region is humanized.
11. The AAV particle of claim 9, wherein the linker is selected from one or more of the members of the group given in Table 2.

12. The AAV particle of claim 9, wherein the first nucleic acid segment encodes from 5' to 3', an antibody heavy chain, a linker, and an antibody light chain.
13. The AAV particle of claim 9, wherein the first nucleic acid segment encodes from 5' to 3', an antibody light chain, a linker, and an antibody heavy chain.
14. The AAV particle of claim 9, wherein the first nucleic acid segment encodes one or more antibody heavy chains.
15. The AAV particle of claim 14, wherein the first nucleic acid segment encodes one or more antibody heavy chains selected from those listed in Tables 3-42.
16. The AAV particle of claim 9, wherein the first nucleic acid segment encodes one or more antibody light chains.
17. The AAV particle of claim 16, wherein the first nucleic acid segment encodes one or more antibody light chains selected from those listed in Tables 3-42.
18. The AAV particle of claim 9, wherein the first nucleic acid segment encodes one or more antibody heavy chains and one or more antibody light chains and, optionally one or more linkers.
19. The AAV particle of any of claims 9-18, wherein said linker is selected from the group consisting of Table 2 and combinations thereof.
20. The AAV particle of claim 1, wherein the first nucleic acid segment encodes an antibody, having at least 95% identity to any of the sequences selected from the group consisting of SEQ ID NO: 2948-9220 (Tables 3-42).
21. An AAV particle comprising a capsid and a viral genome, said viral genome comprising at least one inverted terminal repeat **(ITR)** region and a payload region comprising a regulatory' sequence operably linked to at least a first nucleic acid segment, said first nucleic acid segment encoding a bispecific antibody derived from any of the sequences listed in Tables 3-42 or portions or fragments thereof.
22. The AAV particle of claim 21, wherein the bispecific antibody comprises a light and a heavy chain selected from two different starting antibodies selected from the group consisting of SEQ ID NO: 2948-9220 (Tables 3-42).
23. A method of producing a functional antibody in a subject in need thereof, comprising administering to said subject the AAV particle of any of claims 1-22.
24. The method of claim 23, wherein the level or amount of the functional antibody in the target cell or tissue after administration to the subject is from about .001 ug/mL to 100 mg/mL.

25. The method of claim 23, wherein the functional antibody is encoded by a single first nucleic acid segment of a viral genome within said AAV particle.
26. The method of claim 23, wherein the functional antibody is encoded by two different viral genomes, said two different viral genomes packaged in separate capsids.
27. A pharmaceutical composition comprising an AAV particle of any of the preceding claims in a pharmaceutically acceptable excipient.
28. The pharmaceutical composition of claim 27, wherein the pharmaceutically acceptable excipient is saline.
29. The pharmaceutical composition of claim 27, wherein the pharmaceutically acceptable excipient is 0.001% pluronic in saline.
30. A method of expressing an antibody in a cell or tissue comprising administering the AAV particle of any of claims i-29 via a delivery route selected from the group consisting of enteral (into the intestine), gastroenteral, epidural (into the dura mater), oral (by way of the mouth), transdermal, intracerebral (into the cerebrum), intracerebroventricular (into the cerebral ventricles), epicutaneous (application onto the skin), intradermal, (into the skin itself), subcutaneous (under the skin), nasal administration (through the nose), intravenous (into a vein), intravenous bolus, intravenous drip, intra-arteriai (into an artery), intramuscular (into a muscle), intracardiac (into the heart), intraosseous infusion (into the bone marrow), intrathecal (into the spinal canal), intraparenchymal (into brain tissue), intraperitoneal, (infusion or injection into the peritoneum), intravesical infusion, mtravitreal, (through the eye), intracavernous injection (into a pathologic cavity) intracavitary (into the base of the penis), intravaginal administration, intrauterine, extra-amniotic administration, transdermal (diffusion through the intact skin for systemic distribution), transmucosai (diffusion through a mucous membrane), transvaginal, insufflation (snorting), sublingual, subiabial, enema, eye drops (onto the conjunctiva), or in ear drops, auricular (in or by way of the ear), buccal (directed toward the cheek), conjunctival, cutaneous, dental (to a tooth or teeth), electro-osmosis, endocervical, endosmusiai, endotracheal, extracorporeal, hemodialysis, infiltration, interstitial, intra-abdominal, intra-amniotic, intra-articular, intrabiliary, intrabronchial, intrabursal, intracartilaginous (within a cartilage), **intracaudal** (within the cauda equine), intracisternal (within the cisterna magna cerebellomedularis), mtracomeal (withm the cornea), dental intracornal, mtracoronary (within the coronary arteries), mtracorporus cavernosum (within the dilatable spaces of the corporus cavernosa of the penis).

intradiscal (within a disc), intraductal (within a duct of a gland), **intraduodenal** (within **the** duodenum), intradural (within or beneath the **dura), intraepidermal** (to the epidermis), **intraesophageal (to the** esophagus), intragastric (within the stomach), intragingival (within the gingivae), mtraileai (withm the distal portion of the small intestine), intraiesional (within or introduced directly to a localized lesion), intraluminal (within a lumen of a tube), **intralymphatic** (within the lymph), intramedullary (within the marrow **cavity of** a bone), **intrameningeal** (within the meninges), **intrarnyocardial** (within the myocardium), intraocular **(within** the eye), **intraovarian** (within the ovary), **intraperi cardial** (within the pericardium), intrapleural (within the pleura), **intraprostatic** (within the prostate gland), **intrapulmonary** (within the lungs or its bronchi), **intrasinal** (within the nasal or periorbital sinuses), intraspinal (within the vertebral column), **intrasynovial** (within the synovial cavity **of a joint),** **mtratendinous** (within a tendon), **intratesticular** (within the testicle), intrathecal (within the cerebrospinal fluid at any level of the cerebrospinal axis), intrathoracic (within **the** thorax), **intratubular** (within the tubules **of** an organ), **intratumor** (within a tumor), **intratympanic** (within **the** aurus media), intravascular (within a vessel or vessels), intraventricular (within a ventricle), iontophoresis (by means of **electric** current where ions of soluble salts migrate into the tissues of the body), irrigation (to bathe or flush open wounds or body cavities), laryngeal (directly upon the larynx), nasogastric (through the nose and into the stomach), occlusive dressing technique (topical route administration which is then covered by a dressing which occludes the area), ophthalmic (to the external eye), oropharyngeal (directly to the mouth and pharynx), parenteral, percutaneous, periarticular, peridural, perineural, periodontal, rectal, respiratory (within **the respiratory** tract by inhaling orally or nasally for local or systemic effect), retrobulbar (behind the pons or behind the eyeball), soft tissue, subarachnoid, subconjunctival, submucosal, topical, transplacental (through or across **the** placenta), transtracheal (through the wall of the trachea), transtympanic (across or through the tympanic **cavity),** ureteral (to the ureter), urethral (to the urethra), vaginal, caudal block, diagnostic, nerve block, biliary perfusion, cardiac perfusion, **photopheresis** and spinal.

31. The method of claim 30, wherein the delivery route is intramuscular.

32. The method of claim 31, wherein the intramuscular administration is to at least one limb.

33. The method of claim 30, wherein the deliver}' route is intravascular.

34. The method of claim 30, wherein the delivery route is intrathecal.
35. The method of claim 30, wherein the deliver); route is intracerebroventricuiar.
36. The method of claim 30, wherein the deliver} route is intraparenchymal.
37. The method of claim 30, wherein the AAV particle is encapsulated in a nanoparticle.
38. The method of claim 30, wherein the AAV particle is delivered by a device.
39. The method of claim 38, wherein the device is a gene gun.
40. A method of preventing a disease or disorder in a subject comprising administering to said subject the pharmaceutical composition of any of claims 27-29.
41. The method of claim 40, wherein the administration is at a prophylactically effective dose.
42. The method of claim 41, wherein the dose is from about 1 ug/mL to about 500 ug/mL of expressed polypeptide or 1x10e4 to lx!0el6 VG/mL from the pharmaceutical composition.
43. The method of claim 42, wherein the pharmaceutical composition is administered once.
44. The method of claim 42, wherein the pharmaceutical composition is administered more than once.
45. The method of claim 42, wherein the pharmaceutical composition is administered daily, weekly, monthly or yearly.
46. The method of claim 42, wherein the pharmaceutical composition is co-administered as part of a combination therapy.
47. A method of treating a disease or disorder in a subject in need thereof comprising administering to said subject the pharmaceutical composition of any of claims 27-29.
48. The method of claim 47, wherein said disease or disorder is selected from the group consisting of diseases caused by John Cunningham Virus (JCV), influenza, hepatitis A, hepatitis B, hepatitis D, hepatitis E, respiratory syncytial virus (RSV), herpes simplex virus 1, herpes simplex virus 2, hyman cytomegalovirus, Epstein-Barr virus, Varicella zoster virus, Coronavirus, Poxvirus, Enterovirus 71, rubella virus, human papilloma vims, *Pseudomonas Aeruginosa,* Streptococcus bacteria, Stapylococcus bacteria, *Clostridium Telani,* Bordetella, Mycobacterium, *Francisella Tularensis, Toxoplasma gondii,* Candida yeast, ricin, *bacillus anthracis,* shiga toxin, shiga-like toxin, botulinum toxins, chikungunya virus, dengue virus, trvpasnosoma cruzi, rabies virus, *Plasmodium falciparum,* ebola virus, Marburg virus. West Nile virus, Yellow Fever virus, Japanese encephalitis virus, St. Louis encephalitis virus, rotavirus,

Norwalk virus, *Campylobacter jejuni, Clostridium difficile, Entamoeba histolytica, Helicobacter pyroli,* and Enterotoxin B.

49. The AAV particle of claim 1, wherein the viral genome comprises 2 ITR regions.

50. The AAV particle of claim 1, wherein the at least one ITR region is derived from the same parental serotype as the capsid.

51. The AAV particle of claim 1, wherein the at least one ITR region is derived from a different serotype as the capsid.

52. The AAV particle of claim 1, wherein the at least one ITR region is derived from AAV2.

53. The AAV particle of claim 1, wherein the at least one ITR region is 100-150 nucleotides in length.

54. The AAV particle of claim 1, wherein the at least one ITR region is 102 nucleotides in length.

55. The AAV particle of claim 1, wherein the at least one ITR region is 140-142 nucleotides in length.

56. The AAV particle of claim I, wherein the at least one ITR region is 140 nucleotides in length.

57. The AAV particie of claim 1, wherein the at least one ITR region is 141 nucleotides in length.

58. The AAV particle of claim 1, wherein the at least one ITR region is 142 nucleotides in length.

59. The AAV particle of claim 1, wherein the viral genome further comprises an intron or staffer sequence.

60. A method of producing an antibody in a subject comprising administering the AAV particle of claim 1 to said subject, with the proviso that the antibody is not a virus neutralizing antibody.

61. A method of producing an antibody in a subject comprising administering the AAV particle of claim I to said subject, with the proviso that the antibody is not an HIV or HCV virus neutralizing antibody.

62. The AAV particie of claim I, wherein the payload region of the viral genome comprises a second nucleic acid segment, said second nucleic acid segment encoding an aptamer, siRNA, saRNA, ribozyme, microRNA, mRNA or combination thereof.

63. The AAV particle of claim 62, wherein the second nucleic acid segment encodes an siRNA and said siRNA is designed to target the mRNA that encodes the target of the antibody encoded by the first nucleic acid segment.

64. The AAV particle of claim 62, wherein the second nucleic acid segment encodes a microRNA and said microRNA is selected to target the mRNA that encodes the target of the antibody encoded by the first nucleic acid segment.

65. The AAV particle of claim 62, wherein the second nucleic acid segment encodes an mRNA and said mRNA encodes one or more peptides inhibitors of the same target of the antibody encoded by the first nucleic acid segment.

66. The AAV particle of claim 1 or 62, wherein the payload region of the viral genome comprises a third nucleic acid segment.

67. The AAV particle of claim 66, wherein the third nucleic acid segment encodes a nuclear export signal.

68. The AAV particle of claim 66, wherein the third nucleic acid segment encodes a polynucleotide or polypeptide which acts as a regulator of expression of the viral genome in which it is encoded.

69. The AAV particle of claim 66, wherein the third nucleic acid segment encodes a polynucleotide or polypeptide which acts as a regulator of expression of the payload region of the viral genome in which it is encoded.

70. The AAV particle of claim 66, wherein the third nucleic acid segment encodes a polynucleotide or polypeptide which acts as a regulator of expression of the first nucleic acid segment of the payload region of the viral genome in which it is encoded.

FIG. 1

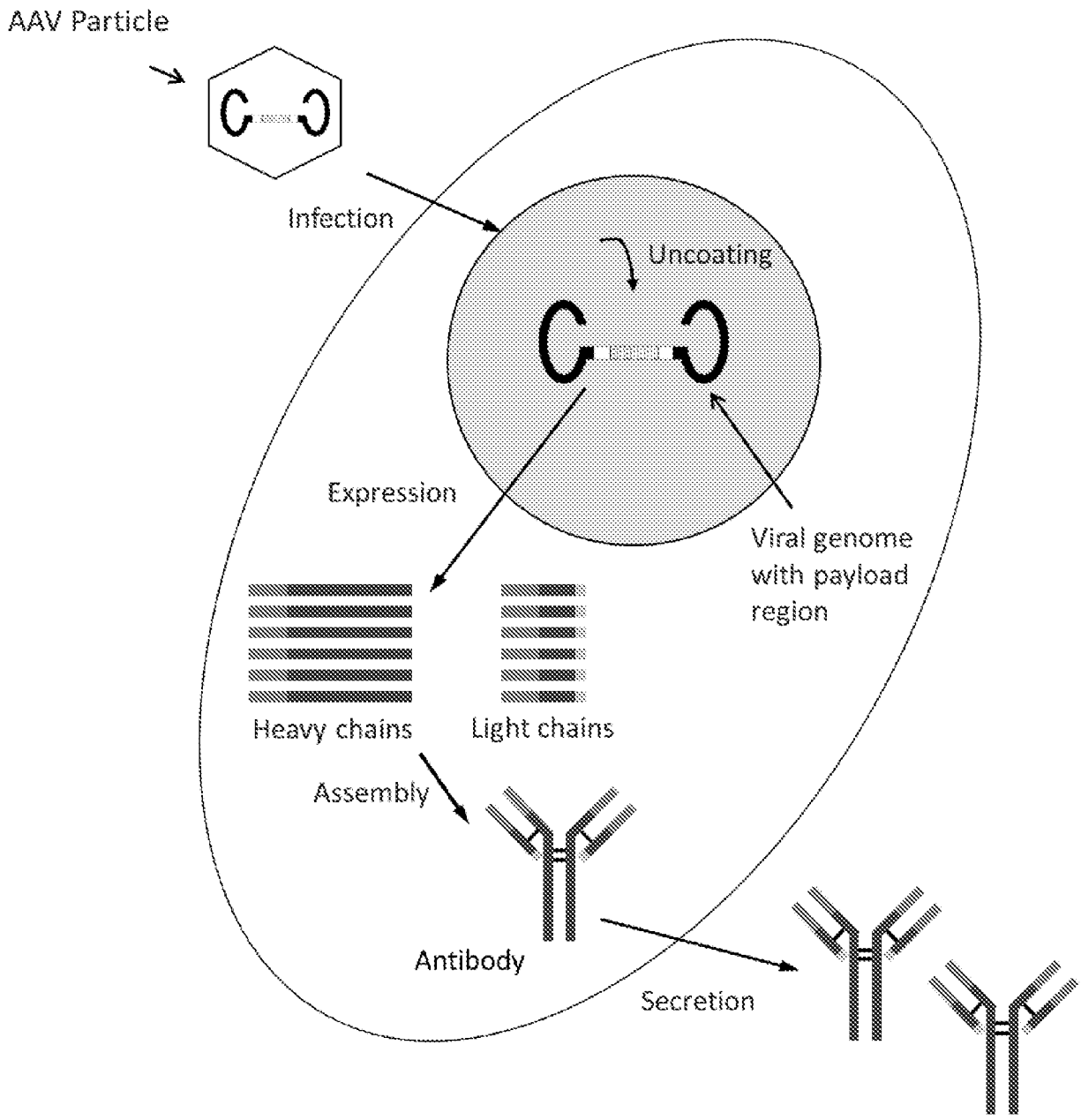

AAV Particle
Infection
Uncoating
Expression
Viral genome with payload region
Heavy chains
Light chains
Assembly
Antibody
Secretion

FIG. 2

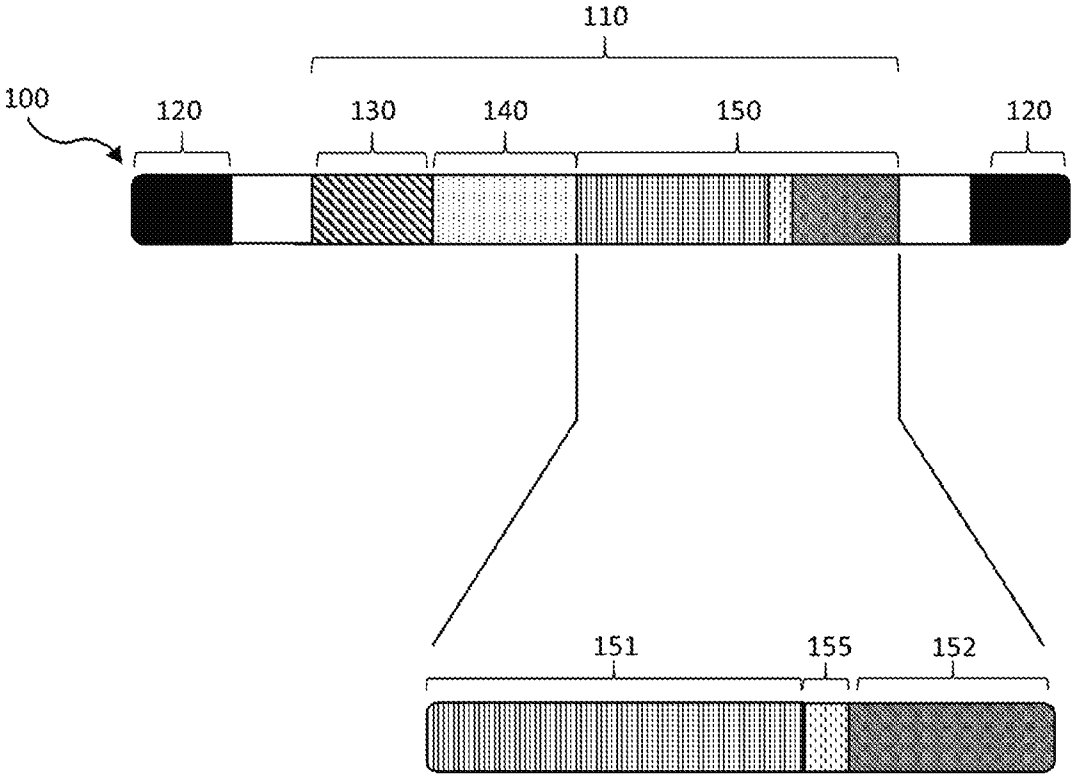
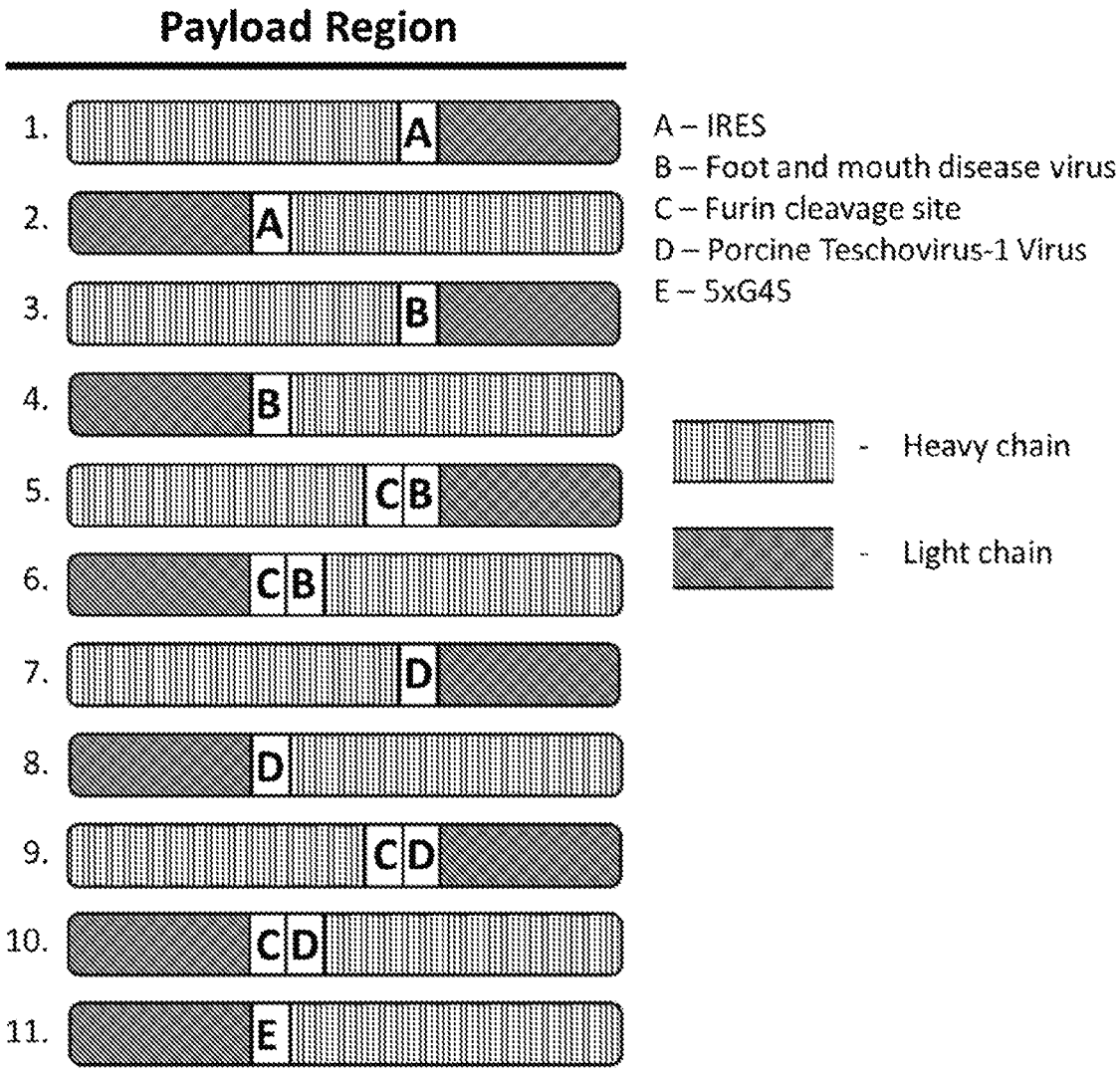


FIG. 3


Payload Region
1.
A
2.
A
3.
B
4.
B
5.
C B
6.
C B
7.
D
8.
D
9.
C D
10.
C D
11.
E
A – IRES
B – Foot and mouth disease virus
C – Furin cleavage site
D – Porcine Teschovirus-1 Virus
E – 5xG4S
- Heavy chain
- Light chain