


Spectral Characterization and Quantum Mechanical Studies of 5-Amino-2-(6-(2-Hydroxyethyl)-3-Oxononyl) Cyclohex-2-Enone Isolated from a Marine Algae

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Abstract: The compound C2 was synthesized and characterized with FTIR, FT-Raman, and UV-Vis, compared to experimental and simulated methods. The compound optimized with the DFT method with the WB97D/6311+G(2d,p) basis set level. In FTIR and Raman, the simulated spectrum is in good agreement with the experimental one. The wave function studies like localized orbital locator, electron localized function, average localized ionization energy, and noncovalent interaction were also studied for other theoretical evidence of the titled compound. This compound's ADMET properties are also studied to support the Swiss ADMET online tool. Molecular docking was studied with the help of Autodock/Vina software, with a binding affinity is -7.3 kcal/mol.

Keywords: DFT; IR spectra; NBO; NCI; ALIE.

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1. Introduction

Mangroves, saline-resistant ecosystems, are present in tropical and sub-tropical intertidal regions worldwide [1]. Mangrove soil offers a distinct ecological habitat for the growth of diversified microorganisms, which are used in recycling environmental nutrients and producing secondary metabolites of medicinal and industrial importance. Several factors constantly change in the ecosystem, which causes the adaption in metabolic pathways that ultimately leads to the biosynthesis of unique metabolites [2]. The microorganisms residing in the mangrove region include bacteria, fungi, actinomycetes, algae, and fungus-like protozoa. The microbial community of the mangrove ecosystem is 91% bacteria and fungi, 7% of algae, and 2% protozoa. There is extensive microbial biodiversity in this ecosystem; of this, only 1% has been explored, and 99% is not explored [3].

Actinobacteria is aerobic, branched, unicellular gram-positive bacteria with 70% GC content in soil, backwater, lake, and marine environment. It is considered a promising target for screening as it can synthesize novel compounds of therapeutic importance [4].

Actinomycetes, particularly the genus *Streptomyces* is a rich source of secondary metabolites with special attention to antibiotics [5]. *Streptomyces* possesses numerous high metabolic production rates, recycling in organic matter, and the ability to degrade chitin, lignocelluloses, etc. [6]. These bioactive molecules serve as lead molecules for drug discovery and the development new drugs [7]. Studies were reported on marine actinobacteria, but scanty reports are available concerning actinobacteria from the mangrove ecosystem.

The bioactive compounds produced from streptomycetes include enzymes, pigments, and compounds, which possess antimicrobial, anticancer, antioxidant, immunosuppressive, and other important bioactivities. Particularly antibiotics, whose biosynthesis is connected and are influenced only by primary metabolism but also by the composition of the culture medium. The bioactive compounds are released into the extracellular media, and isolation and purification involve separation techniques, i.e., solvent extraction, chemical precipitation, ion exchange chromatography, HPLC purification, etc. [8]. Further characterization involves mass spectrometry, NMR spectroscopy, and X-ray diffraction [9–13].

50 isolates of actinomycetes were mangrove soil from the area of Visakhapatnam, Andhra Pradesh and named BC 01 - BC 20 and KVR 01 - KVR 30 and maintained on the actinomycetes agar a medium. These isolates are subjected to screening for biological activity [14]. Of these, only four exhibit potent antimicrobial and antioxidant activity. BC_01 exhibits the maximum inhibition of DPPH radical scavenging activity, ferric-reducing antioxidant activity, and total antioxidant activity when compared to others [15]. The morphological, biochemical, and molecular characterization studies reveal that the potent isolate belongs to the genus *Streptomyces* and the strain's name is *Streptomyces coelicoflavus*. The bioactive fraction was isolated from the fermented broth through column chromatography. Homogeneity and purity of the isolated fractions were determined by TLC and reverse phase HPLC. Further purification procedure was subjected to the separation of three bioactive fractions named BC_C1, BC_C2, and BC_C3. The tentative chemical structure of the fraction was determined by the literature's spectral data and available data. Based on the data the compound was named as 8-(aminomethyl)-7-hydroxy-1-(1-hydroxy-4-(hydroxylmethoxy)-2,3-dimethylbutyl)-2-methyldodecahydro phenanthren-9(1H)-one [16–18].

Computational chemistry tools have emerged as a very good source of information about new compounds [19–21]. This will help people gain detailed information about several experimental materials systems [22–26].

The antibiotic compound C2 8-(aminomethyl)-7-hydroxy-1-(1-hydroxy-4-(hydroxylmethoxy)-2,3-dimethylbutyl)-2-methyldodecahydro phenanthren-9(1H)-one. This compound showed better antimicrobial efficacy against both bacteria and fungi. The minimum inhibitory concentration values for bacteria and fungi are 12.5 - 75 $\mu\text{g/ml}$ and 50 - 125 $\mu\text{g/ml}$, respectively. It exhibits the maximum inhibition of DPPH radical scavenging activity (61.92 \pm 6.95), ferric reducing antioxidant activity (56.66 \pm 2.51), and total antioxidant activity (67.83 \pm 24.79) at a concentration of 20 $\mu\text{g/ml}$. This compound exhibits potent anti-inflammatory activity compared to the standard drug Diclofenac sodium. It also possesses strong *in vitro* cytotoxic potential against various cell lines.

2. Materials and Methods

The compound C2 structure was optimized to the DFT method with WB97D/6311+G(2d,p) basis set method; with the help of the Gaussian software package [27]. The Infrared spectrum was verified from an Agilent spectrometer by KBr pellets, with a range <https://biointerfaceresearch.com/>

of 4000- 400 cm^{-1} ; FT-Raman was recorded from Bruker RFS-27 Multi RAM FT Raman spectrometer with a range of 4000 – 50 cm^{-1} . Wave function analyses like ELF, LOL, and RDG/noncovalent interaction were calculated with the help of multiwfn software [28][29]. Molecular docking analysis was done by Autodock/Vina software, and protein alignment software using for the discovery studio visualizer.

3. Results and Discussion

3.1. Structural geometry.

The titled compound C2 was optimized with WB97D/6311+G(2d,p) basic set level; [30,31] the optimized structure was shown in Figure 1. The highest bond length, bond angle, and dihedral angle are observed at 1.55, 122.88, and 179.75, corresponding to the C18-H20, C2-C3-O4, and H32-C2-C6-H35, respectively [32,33]. Some bond lengths, bond angles, and dihedral angles are discussed below; C1-C14, C1-H30, C2-H32, C3-C10, C5-H34, C7-C8, C7-H36, C8-H38, C10-H40, C14-C27, C17-H51, C18-H52, C22-H58, C1-C2-C3, C3-C2-H32, C2-C3-C10, H33-C5-H34, C2-C6-H35, C7-C6-H35, C13-C-H39, C8-C11-H48, C11-C12-C16, C15-C14-C27, C14-C15-O28, C12-C14-H50, C18-C17-H51, C20-C18-H52, C14-C1-C2-C3, C14-C1-C2-H32, H31-C1-C2-C6, H30, C1-C14-C15, C6-C2-C3-O4, C1-C2-C6-C7, C3-C2-C6-H35, C2-C3-C10-H41, C15-C5-C6-C2, H39-C9-C10-H49, H44-C12-C16-H50 and C14-C27-N29-H69 corresponding to the 1.53, 1.09, 1.10, 1.51, 1.09, 1.52, .09, .09, 1.08, 1.53, 1.09, 1.09, 1.10, 112.85, 106.14, 114.85, 107.06, 106.33, 106.36, 107.76, 109.68, 109.07, 112.84, 108.74, 112.38, 108.46, 104.79, -179.31, 62.97, -178.17, -66.93, -126.02, -177.37, 64.77, 65.08, -56.40, -177.26, 55.23 and -65.79 respectively [34].

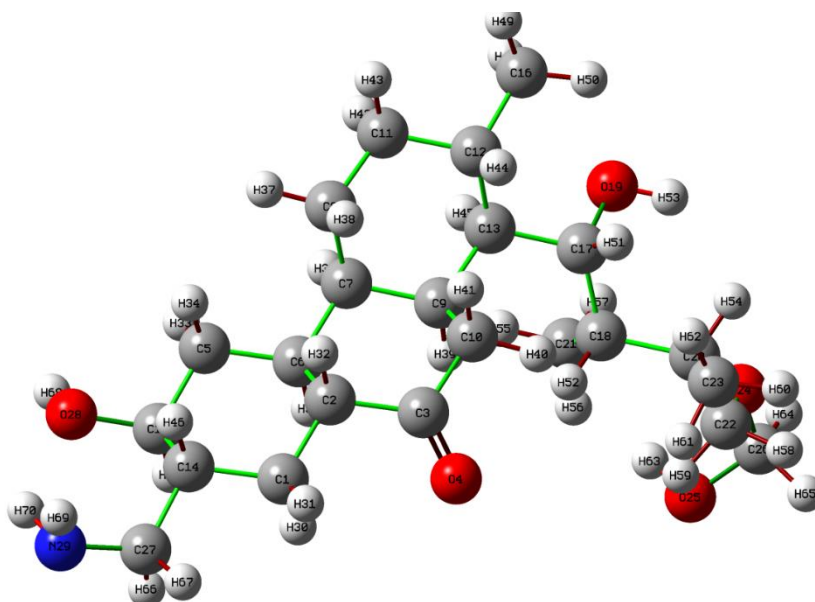


Figure 1. Optimized structure of compound C2.

3.2. Vibrational spectral analysis.

The vibrational spectra of infrared and Raman were recorded using KBr pellets [35,36]. Comparison infrared and Raman spectra are shown in Figure 2 [37]. The NH stretching vibration presented at 3364 cm^{-1} in FT-IR and FT-Raman occurs at 3422 cm^{-1} ; normally, NH vibration occurs at 3400-3300 cm^{-1} [38]. In CH, stretching vibration presented at 3014, 2924,

and 2855 cm^{-1} in infrared and Raman spectra presented at 3214, 3008, and 2921 cm^{-1} . CN stretching vibration was observed at 1653 cm^{-1} in infrared and 1686 cm^{-1} in Raman; [39] the CO stretching vibration was detected at 1246 cm^{-1} in FTIR, and Raman spectra were observed at 1413 cm^{-1} . CH_2 group vibrations are normally observed at six fundamentals: symmetric stretch, asymmetric stretch, scissoring, rocking [40,41], and wagging and twisting vibrations. Asymmetric stretching vibration is generally observed at 3000 cm^{-1} , and symmetric stretching vibration is observed at 3000 to 2900 cm^{-1} [42]. Titled compound CH_2 asymmetric stretching vibration are observed at 3014 cm^{-1} in infrared, Raman asymmetric stretching vibration was observed at 3008 cm^{-1} , and CH_2 symmetric stretching vibrations were noted at 2924 cm^{-1} in FTIR, and FT-Raman is 2921 cm^{-1} . The aromatic CH_2 vibrations of weak to medium can be observed at 1500 to 200 cm^{-1} due to scissoring, wagging, rocking, and twisting vibrations. The above-mentioned vibrations give rise to lower wavenumber and variable intensity bands [43]. In this compound, the band is 1436 and 1410 cm^{-1} for infrared, and Raman is 1413 cm^{-1} in CH_2 scissoring mode [44]. The wagging modes are presented at 1200 to 1000 cm^{-1} , with moderate to strong intensity bands. The titled compound C2, CH_2 wagging mode is presented at 1023 cm^{-1} in infrared, and CH_2 rocking mode is presented at 1318 cm^{-1} .

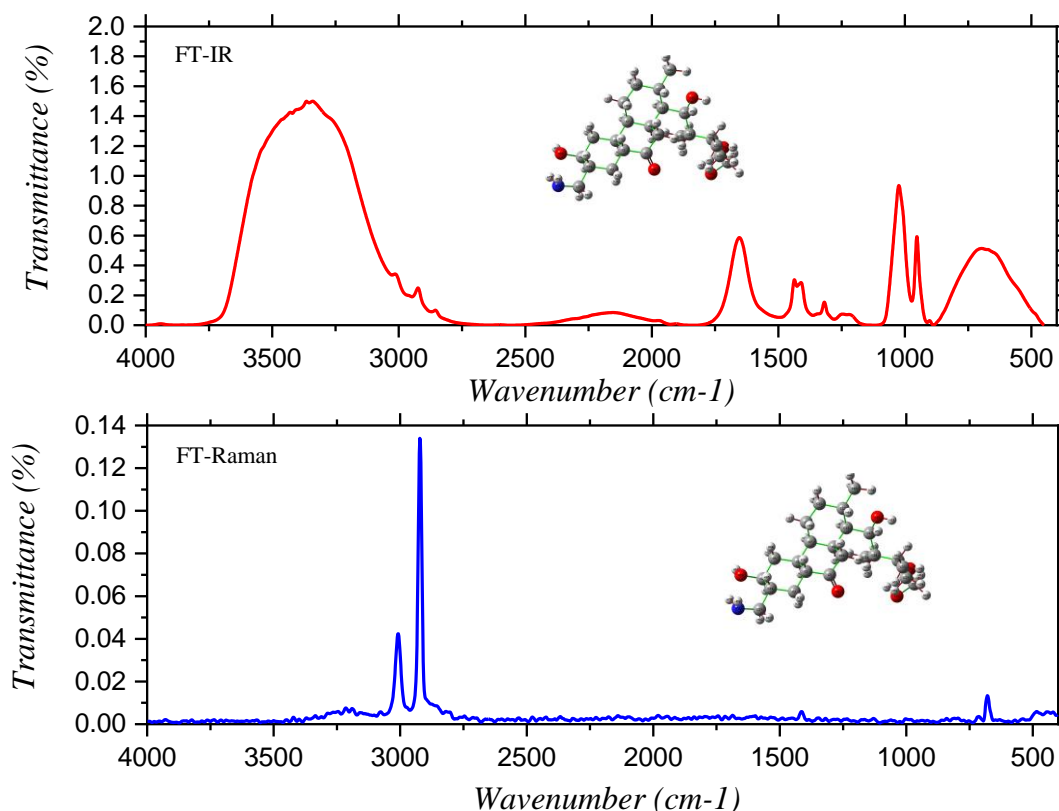


Figure 2. Comparison of Infrared and Raman spectra of the compound C2.

3.3. Molecular electrostatic potential.

The MEP is calculated from the DFT way with WB97D/6311+G(2d,p) basis set [45,46]. Calculate the nucleophilic and electrophilic attack; the molecular electrostatic potential study is very important [47,48]. The blue and red colors are represented more electron-poor and rich regions. MEP surface map is shown in Figure 3; the positive electrostatic potential is located at carbon and all hydrogen atoms, and the negative potential is located at

oxygen and nitrogen atoms [49][50]. Negative and positive electrostatic potentials are clearly indicated by the electrophilic and nucleophilic attack of the compound [51].

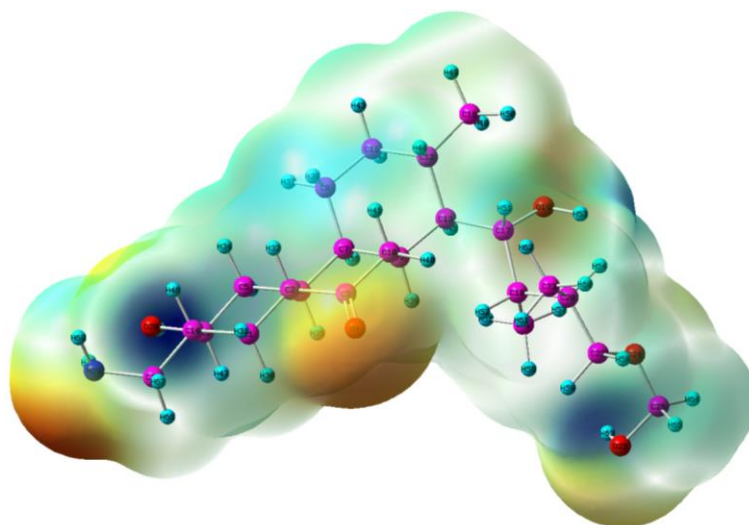


Figure 3. Molecular electrostatic potential surface map of the compound C2.

3.4. Frontier molecular orbital analysis.

Calculating the electrical and chemical properties of the FMO is the most important one [52,53]. The HOMO and LUMO represent the electron donor and acceptor [54]. The HOMO-LUMO energy surface map is shown in fig.4; C2 compound HOMO and LUMO energy are -0.36322eV and 0.18031 eV, with the energy gap, is 5.0 eV [55]. However, the hardness of the titled complex, the HOMO and LUMO energy is higher, and HOMO represents softness and LUMO energy is lower [56]. LUMO is presented at oxygen atoms and carbon atoms; HOMO is presented to an amino group, oxygen atom, and ring [57].

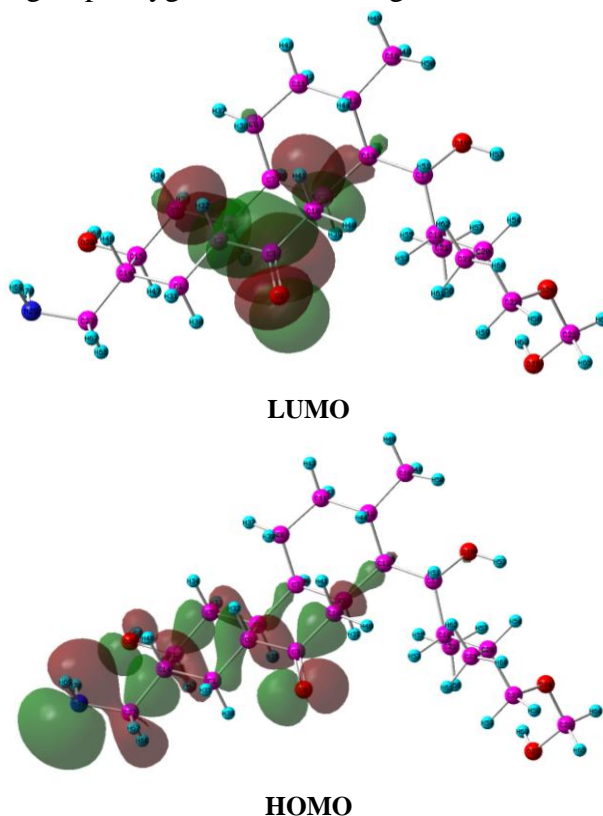


Figure 4. HOMO-LUMO surface map of the compound C2.

3.5. Natural population and bond orbital analysis.

The NPA was the distribution of electrons in various subshells; the distribution of electrons is listed in table.S3. The atom N29 has shown the highest electronegativity compared to other electronegative atoms; the value is -0.91534 [58,59]. All carbon atoms have positive and negative charges, and all hydrogen atoms have positive charges; C3 atom showed a highly positive charge compared to other atoms; the charge is 0.57247. In NPA theory, a positive atom accepts electrons and negative atoms donate electrons [60,61].

NBO analysis was determined by intermolecular and intramolecular interactions. C2 complex donor-acceptor interaction was calculated from B3LYP/631G basis set with the help of second-order perturbation theory analysis [62,63]. The donor-acceptor interactions are shown in table.S4 [64,65]. C2 highest stabilization energy is observed at LP(2) O4 to $\sigma^*(C2-C3)$ and $\sigma^*(C3-C10)$, with occupancy is 1.90407 and stabilization energy is 19.03 and 17.98 respectively; and LP(2) O24 to $\sigma^*(O25-C26)$ and LP(2) O25 to $\sigma^*(O24-C26)$, with stabilization energy is 14.75 and 14.04 and occupancy is 1.90770 and 1.93118. Furthermore, the highest stabilization energy noted at $\sigma(C23-H26)$, $\sigma(C21-H55)$ to $\sigma^*(C18-C20)$ and $\sigma(C16-H49)$, $\sigma(C9-H39)$ to $\sigma^*(C12-C13)$, with stabilization energy is 3.07, 3.62, 3.83 and 3.04 and occupancy is 1.98977, 1.98763, 1.98864 and 1.97318 respectively. The highest stabilization energy observed at $\sigma(C8-H37)$, $\sigma(C5-H33)$, and $\sigma(C1-C14)$ to $\sigma^*(C11-C12)$, $\sigma^*(C14-C15)$ and $\sigma^*(C15-O28)$, with stabilization energy, is 3.38, 3.09 and 3.21 and occupancy is 1.98067, 1.9798 and 1.97123 respectively.

3.6. Topology analysis.

Explain the topology analysis that ELF and LOL are very important [66,67]. The topology analysis is based on the ELF and LOL; the localized orbital locator and localized locator is the maximum electron pair in molecular spacing corresponding to the covalent bond [68]. The electron localized function and localized orbital locators represent the electron pair density due to the orbitals and localized orbital overlapping, respectively [69,70].

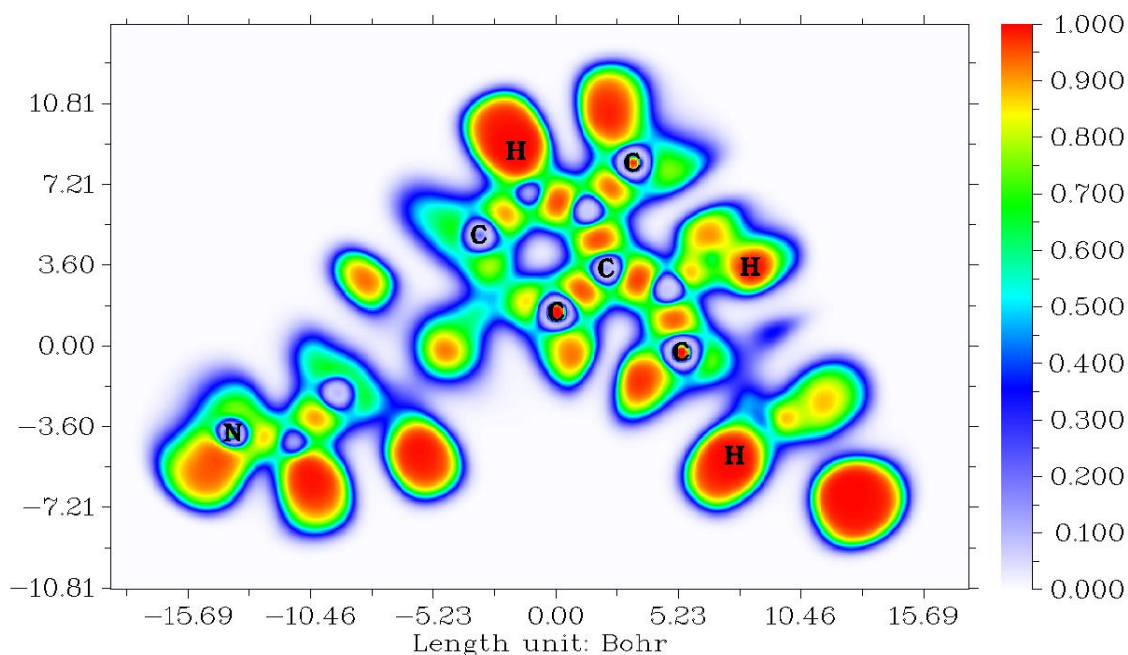


Figure 5. ELF surface map of compound C2.

Analyzing the ELF and LOL, the software used for multiwfn software [28]. The electron localized function and localized orbital locator color coding range is blue to red, and the surface map is shown in Figures 5 and 6. The ELF scale range is 0.000 to 1.000, and the LOL scale range is 0.000 to 0.800; the above color ranges are clearly indicated bonding and non-bonding orbitals [71]. The delocalized electronic region was indicated to be below the 0.5 color range, and the localized electronic region was represented above the 0.5 color range. ELF and LOL surface map, the blue and red represent bonding and non-bonding orbitals [72]. The red and blue colors indicated the maximum electrons of the bonding and non-bonding. The red color was located at hydrogen atoms; the blue color was located at nitrogen and carbon atoms in both ELF and LOL surface maps. The light blue color represents electron reduction on the inner shell and valance; this is located at the oxygen atom and surrounds the whole molecule.

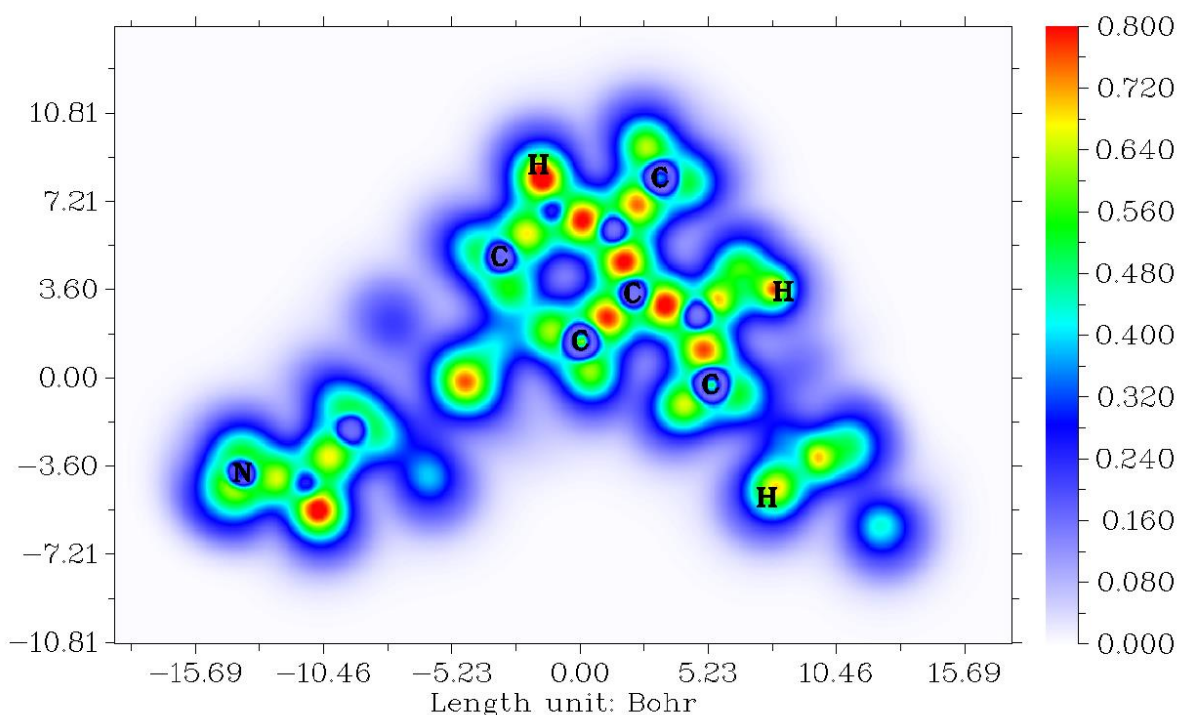


Figure 6. LOL surface map of the compound C2

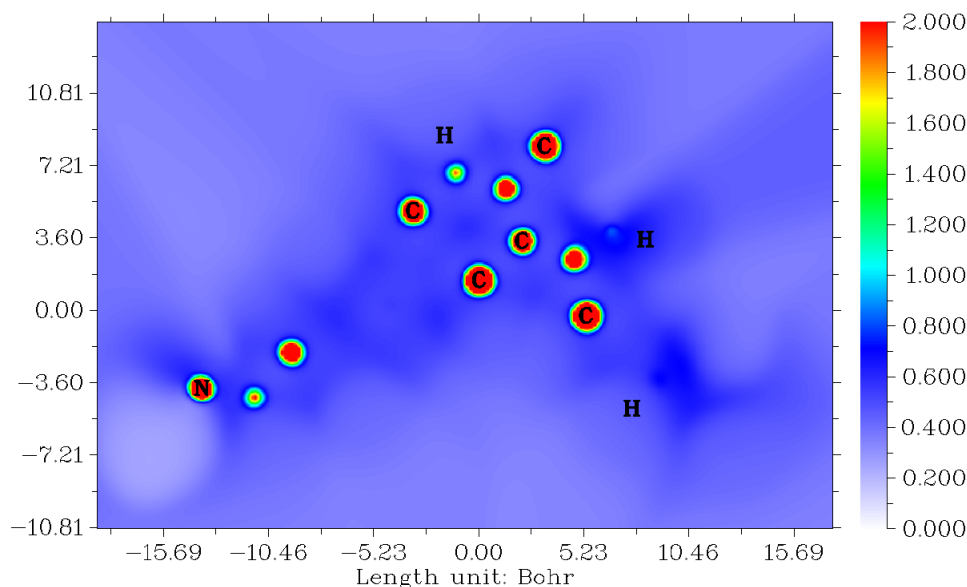


Figure 7. ALIE surface map of compound C2.

3.7. Average localized ionization energy (ALIE).

The average local energy of the compound C2 is needed to remove the electron from the point systems; [73,74] average localized ionization energy is shown in Figure 7. The lowest value of closely held electrons was preferred to the reaction site of the electrophile or radicals [75,76]. The delocalized electrons indicated a greenish-blue region, greenish blue providing different resonance structures and explaining the stability of the compound; unfortunately, no greenish-blue color was presented [77]. Stable or sigma bond indicated to blue region. Unfortunately, no blue region is presented in this compound. The multiple bonds are denoted at the red region, red color carbon, and nitrogen atoms.

3.8 Non covalent interaction.

RDG was a very useful tool for analyses of the intra- and inter-molecular interactions [73][78]. RDG function was developed by E.R. Johnson *et al.*; the gradient was plotted for RDG vs. electron density, which is shown in Figure 8 [79]. Noncovalent interaction was mainly used to predict the biological activity of the titled compound C2. The above surface map blue color indicates stronger interactive hydrogen bonds, red represents the strongest repulsion, and green represents the Van der Waals interaction of the titled compound [67].

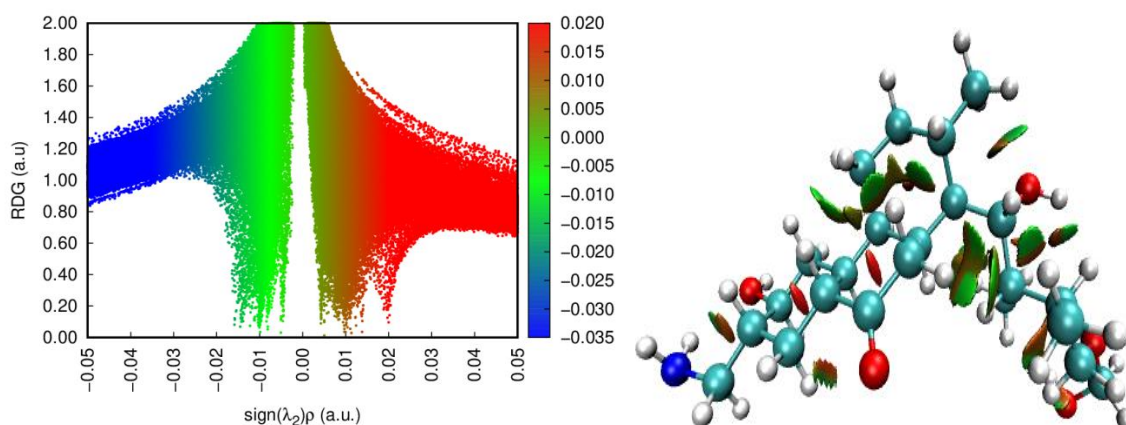


Figure 8. RDG surface map of compound C2.

3.9. Drug likeness.

Drug likeness is called ADME, the ADME was an important technique to calculate the biological property of the titled compound [80]. The Swiss online tool is used for calculating the ADME properties of this compound [81]. Biological property is calculated supporting PASS study results [82]. After the PASS study results, the most important one is the general pump inhibitor, the active probability score is 0.683, and the inactive probability score is 0.009, as listed in Table.1. All physical-chemical properties, namely rotatable bonds, heavy atoms, H-bond donor, H-bond acceptor, bioactivity, TPSA, etc., are given below.

Table 1. PASS study result of the compound C2.

Pa	Pi	Activity
0,683	0,009	General pump inhibitor
0,665	0,029	CYP3A substrate
0,647	0,033	CYP3A4 substrate
0,560	0,046	Proteasome ATPase inhibitor
0,548	0,076	Phosphatase inhibitor
0,501	0,030	Antifungal

Pa	Pi	Activity
0,474	0,019	Antibacterial
0,475	0,025	Antimetastatic
0,392	0,015	Phospholipase A1 inhibitor

3.10. Molecular docking.

Docking studies were done with Autodock/Vina free software package [83]. Earlier protein is downloaded from the RCSB-PDB data bank and prepares the protein by removing water molecules and adding polar hydrogen bonds using Discovery studio visualizer software [84,85]. The protein's active site is definite with 60A0 x 60A0 x 60A0 grid dimension [86]. The compound C2 is docked to the crystal structure of the TetR-like transcription regulator LfrR from *Mycobacterium smegmatis* (2WGB); this is an excellent docking score; [87,88] docking simulation results clearly indicated that the highest docking score is -7.3 kcal/mol and lowest docking score is -6.7 kcal/mol, shown in table.2.

Table 2. The binding affinity of the compound C2.

Mode	Binding Affinity	Distance from msd(l,p)	Best mode msd(u,p)
1	-7.3	0	0
2	-7	4.287	6.637
3	-6.8	13.489	15.255
4	-6.8	5.047	8.6
5	-6.7	2.42	3.46
6	-6.7	3.944	8.736
7	-6.7	13.541	16.054
8	-6.7	3.959	6.84
9	-6.7	25.601	28.105

Table 3. Favorable non-bond interaction of the compound C2.

Distance	Category	Type	From	From-Chem	To	To-Chem
2.81447	HB	Conventional HB	A:ARG148:HH21	H-Donor	:UNK0:O	H-Acceptor
2.72576	HB	Conventional HB	A:ARG148:HH22	H-Donor	:UNK0:O	H-Acceptor
3.05538	HB	Conventional HB	:UNK0:H	H-Donor	B:GLU160:O	H-Acceptor
2.06627	HB	Conventional HB	:UNK0:H	H-Donor	A:ASP125:OD1	H-Acceptor
4.77899	Hydrophobic	Pi-Alkyl	A:TYR107	Pi-Orbitals	:UNK0	Alkyl

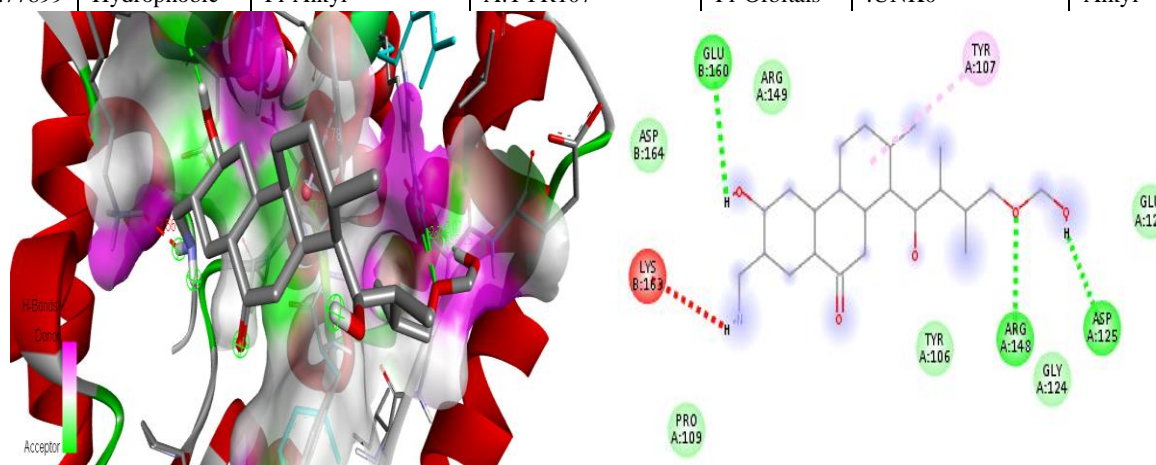


Figure 9. Interaction site of compound C2.

The favorable non-bond interactions are listed in table.5. The compound C2 interacted with protein (2WGB) has shown four conventional hydrogen bonds, namely A:ARG148:HH21 (Arginine), A:ARG148:HH22 (Arginine), B:GLU160:O (Glutamic Acid) and A:ASP125:OD1 (Aspartic Acid) with bond distance is 2.8, 2.7, 3.1 and 2.1 respectively; and one Pi-alkyl bond

such as A:TYR107 (Tyrosine), with bond distance is 4.8. The protein-ligand interaction sites are shown in Figure 9. See Table 3 for details.

4. Conclusions

Compound C2 was synthesized and characterized with a structure like infrared and Raman; after, the compound was compared to theoretical support with the support of Gaussian software. The infrared spectrum experimental and calculated values are closer. This compound is soft because the HOMO-LUMO energy gap is higher; the value is 5.0 eV. Molecular electrostatic potential study of the titled compound is most likely to nucleophile attack. An NPA and natural bond property are also calculated in this study. PASS studies, the active probability score is 0.683; this is the highest score. Drug likeness properties calculated the compound is a biologically active compound. Wave functions were also calculated in this study, namely LOL, ELF, ALIE, and RDG/noncovalent interaction. The molecular simulation studies compound C2 docked to *Mycobacterium smegmatis* (2WGB) protein and confirms the good binding affinity score; the highest binding affinity score is -7.3 kcal/mol.

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Conflicts of Interest

The authors declare no conflict of interest.

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Supplementary materials

Table S1. Optimized parameters of compound C2.

Bond	Length
(C1,C2)	1.5221
(C1,C14)	1.5314
(C1,H30)	1.0962
(C1,H31)	1.0919
(C2,C3)	1.5173
(C2,C6)	1.5428
(C2,H32)	1.0997
(C3,O4)	1.2076
(C3,C10)	1.5098
(C5,C6)	1.5261
(C5,C15)	1.5244
(C5,H33)	1.097
(C5,H34)	1.0946
(C6,C7)	1.5371
(C6,H35)	1.0998
(C7,C8)	1.5287
(C7,C9)	1.5376
(C7,H36)	1.0982
(C8,C11)	1.5244
(C8,H37)	1.0925
(C8,H38)	1.0955
(C9,C10)	1.5425
(C9,C13)	1.5429
(C9,H39)	1.0944
(C10,H40)	1.0894
(C10,H41)	1.0939
(C11,C12)	1.5333
(C11,H42)	1.0978
(C11,H43)	1.0937
(C12,C13)	1.5467
(C12,C16)	1.5278
(C12,H44)	1.0981
(C13,C17)	1.5416
(C13,H45)	1.0978
(C14,C15)	1.5266
(C14,C27)	1.5388
(C14,H46)	1.0989
(C15,O28)	1.4239
(C15,H47)	1.1016
(C16,H48)	1.0927
(C16,H49)	1.0938
(C16,H50)	1.0881
(C17,C18)	1.5465
(C17,O19)	1.4242
(C17,H51)	1.0994
(C18,C20)	1.5514
(C18,C21)	1.5302
(C18,H52)	1.0929
(O19,H53)	0.9594
(C20,C22)	1.5229
(C20,C23)	1.5312
(C20,H54)	1.098
(C21,H55)	1.0925
(C21,H56)	1.0906
(C21,H57)	1.089
(C22,O24)	1.4186
(C22,H58)	1.0983
(C22,H59)	1.0963
(C23,H60)	1.0921

Bond	Length
(C23,H61)	1.0943
(C23,H62)	1.0913
(O24,C26)	1.3923
(O25,C26)	1.4015
(O25,H63)	0.9604
(C26,H64)	1.0932
(C26,H65)	1.0956
(C27,N29)	1.4557
(C27,H66)	1.0964
(C27,H67)	1.0942
(O28,H68)	0.9591
(N29,H69)	1.0135
(N29,H70)	1.0133
(C2,C1,C14)	113.0355
(C2,C1,H30)	108.8826
(C2,C1,H31)	109.6377
(C14,C1,H30)	109.6439
(C14,C1,H31)	109.6691
(H30,C1,H31)	105.7077
(C1,C2,C3)	112.8571
(C1,C2,C6)	111.3511
(C1,C2,H32)	109.0594
(C3,C2,C6)	109.66
(C3,C2,H32)	106.1441
(C6,C2,H32)	107.497
(C2,C3,O4)	122.8863
(C2,C3,C10)	114.8512
(O4,C3,C10)	122.2369
(C6,C5,C15)	112.0039
(C6,C5,H33)	109.6332
(C6,C5,H34)	110.5504
(C15,C5,H33)	109.6657
(C15,C5,H34)	107.7847
(H33,C5,H34)	107.0648
(C2,C6,C5)	109.2965
(C2,C6,C7)	113.3754
(C2,C6,H35)	106.332
(C5,C6,C7)	113.665
(C5,C6,H35)	107.3274
(C7,C6,H35)	106.362
(C6,C7,C8)	114.9893
(C6,C7,C9)	109.9122
(C6,C7,H36)	106.678
(C8,C7,C9)	111.2501
(C8,C7,H36)	106.7874
(C9,C7,H36)	106.7448
(C7,C8,C11)	110.9321
(C7,C8,H37)	110.1069
(C7,C8,H38)	110.6977
(C11,C8,H37)	109.3838
(C11,C8,H38)	109.4307
(H37,C8,H38)	106.1696
(C7,C9,C10)	110.5651
(C7,C9,C13)	111.6914
(C7,C9,H39)	105.9694
(C10,C9,C13)	113.605
(C10,C9,H39)	106.7839
(C13,C9,H39)	107.7697
(C3,C10,C9)	110.1983
(C3,C10,H40)	108.2131
(C3,C10,H41)	108.0686
(C9,C10,H40)	112.248

Bond	Length
(C9,C10,H41)	110.3272
(H40,C10,H41)	107.6505
(C8,C11,C12)	113.9809
(C8,C11,H42)	109.172
(C8,C11,H43)	109.689
(C12,C11,H42)	108.1801
(C12,C11,H43)	109.2323
(H42,C11,H43)	106.2935
(C11,C12,C13)	109.4664
(C11,C12,C16)	109.077
(C11,C12,H44)	107.8697
(C13,C12,C16)	115.3164
(C13,C12,H44)	107.7187
(C16,C12,H44)	107.1266
(C9,C13,C12)	109.9972
(C9,C13,C17)	113.1841
(C9,C13,H45)	106.7845
(C12,C13,C17)	113.6166
(C12,C13,H45)	106.9431
(C17,C13,H45)	105.7983
(C1,C14,C15)	109.3367
(C1,C14,C27)	110.0592
(C1,C14,H46)	108.8836
(C15,C14,C27)	112.8494
(C15,C14,H46)	107.4272
(C27,C14,H46)	108.1758
(C5,C15,C14)	111.5763
(C5,C15,O28)	110.7457
(C5,C15,H47)	108.9675
(C14,C15,O28)	108.7474
(C14,C15,H47)	107.6039
(O28,C15,H47)	109.1206
(C12,C16,H48)	110.9981
(C12,C16,H49)	109.4798
(C12,C16,H50)	112.3852
(H48,C16,H49)	108.2428
(H48,C16,H50)	107.2217
(H49,C16,H50)	108.3826
(C13,C17,C18)	115.9793
(C13,C17,O19)	106.1664
(C13,C17,H51)	107.5711
(C18,C17,O19)	109.6953
(C18,C17,H51)	108.462
(O19,C17,H51)	108.7604
(C17,C18,C20)	110.0121
(C17,C18,C21)	110.6049
(C17,C18,H52)	108.4884
(C20,C18,C21)	115.3273
(C20,C18,H52)	104.7907
(C21,C18,H52)	107.2278
(C17,O19,H53)	108.8523
(C18,C20,C22)	114.163
(C18,C20,C23)	110.3219
(C18,C20,H54)	109.9103
(C22,C20,C23)	108.6829
(C22,C20,H54)	105.283
(C23,C20,H54)	108.2211
(C18,C21,H55)	110.3337
(C18,C21,H56)	110.9349
(C18,C21,H57)	111.7248
(H55,C21,H56)	107.5387
(H55,C21,H57)	107.8039

Bond	Length
(H56,C21,H57)	108.3525
(C20,C22,O24)	109.5422
(C20,C22,H58)	109.7397
(C20,C22,H59)	110.7578
(O24,C22,H58)	108.611
(O24,C22,H59)	110.5068
(H58,C22,H59)	107.6359
(C20,C23,H60)	111.2633
(C20,C23,H61)	110.5334
(C20,C23,H62)	111.9533
(H60,C23,H61)	108.0523
(H60,C23,H62)	107.4766
(H61,C23,H62)	107.378
(C22,O24,C26)	113.7814
(C26,O25,H63)	108.0926
(O24,26,O25)	113.217
(O24,C26,H64)	105.6177
(O24,C26,H65)	111.1734
(O25,C26,H64)	111.4947
(O25,C26,H65)	105.3901
(H64,C26,H65)	110.0291
(C14,C27,29)	117.5334
(C14,C27,H66)	108.4764
(C14,C27,H67)	108.3672
(N29,C27,H66)	108.2655
(N29,C27,H67)	107.734
(H66,C27,C67)	105.8706
(C15,O28,C68)	108.6693
(C27,N29,H69)	109.9954
(C27,N29,H70)	109.2752
(H69,N29,H70)	106.5085
(C14,C1,C2,C3)	-179.3184
(C14,C1,C2,C6)	-55.484
(C14,C1,C2,H32)	62.9789
(H30,C1,C2,C3)	-57.2105
(H30,C1,C2,C6)	66.624
(H30,C1,C2,H32)	-174.9132
(H31,C1,C2,C3)	57.9937
(H31,C1,C2,C6)	-178.1719
(H31,C1,C2,H32)	-59.709
(C2,C1,C14,C15)	54.7439
(C2,C1,C14,C27)	179.2319
(C2,C1,C14,H46)	-62.3523
(H30,C1,C14,C15)	-66.9365
(H30,C1,C14,C27)	57.5515
(H30,C1,C14,H46)	175.9673
(H31,C1,C14,C15)	177.4143
(H31,C1,C14,C27)	-58.0977
(H31,C1,C14,H46)	60.3181
(C1,C2,C3,O4)	-1.2629
(C1,C2,C3,C10)	176.9326
(C6,C2,C3,O4)	-126.0245
(C6,C2,C3,C10)	52.1709
(H32,C2,C3,O4)	118.1398
(H32,C2,C3,C10)	-63.6647
(C1,C2,C6,C5)	54.7003
(C1,C2,C6,C7)	-177.3739
(C1,C2,C6,H35)	-60.8577
(C3,C2,C6,C5)	-179.6708
(C3,C2,C6,C7)	-51.745
(C3,C2,C6,H35)	64.7712
(H32,C2,C6,C5)	-64.6963

Bond	Length
(H32,C2,C6,C7)	63.2294
(H32,C2,C6,H35)	179.7457
(C2,C3,C10,C9)	-55.5344
(C2,C3,C10,H40)	-178.6105
(C2,C3,C10,H41)	65.0873
(O4,C3,C10,C9)	122.6741
(O4,C3,C10,H40)	-0.402
(O4,C3,C10,H41)	-116.7043
(C15,C5,C6,C2)	-56.409
(C15,C5,C6,C7)	175.8275
(C15,C5,C6,H35)	58.5102
(H33,C5,C6,C2)	-178.3831
(H33,C5,C6,C7)	53.8534
(H33,C5,C6,H35)	-63.4639
(H34,C5,C6,C2)	63.811
(H34,C5,C6,C7)	-63.9525
(H34,C5,C6,H35)	178.7303
(C6,C5,C15,C14)	58.0612
(C6,C5,C15,O28)	179.3508
(C6,C5,C15,H47)	-60.6119
(H33,C5,C15,C14)	-179.9834
(H33,C5,C15,O28)	-58.6937
(O33,C5,C15,H47)	61.3436
(H34,C5,C15,C14)	-63.7578
(H34,C5,C15,O28)	57.5319
(H34,C5,C15,H47)	177.5692
(2,C6,C7,C8)	-71.0032
(C2,C6,C7,C9)	55.4476
(C2,C6,C7,H36)	170.8144
(C5,C6,C7,C8)	54.6218
(C5,C6,C7,C9)	-178.9275
(C5,C6,C7,H36)	-63.5607
(H35,C6,C7,C8)	172.4982
(H35,C6,C7,C9)	-61.051
(H35,C6,C7,C36)	54.3157
(C6,C7,C8,C11)	178.6327
(C6,C7,C8,H37)	-60.1398
(C6,C7,C8,H38)	56.9465
(C9,C7,C8,C11)	52.8708
(C9,C7,C8,H37)	174.0983
(C9,C7,C8,H38)	-68.8154
(H36,C7,C8,C11)	-63.2463
(H36,C7,C8,H37)	57.9813
(H36,C7,C8,H38)	175.0676
(C6,C7,C9,C10)	-57.053
(C6,C7,C9,C13)	175.3927
(C6,C7,C9,H39)	58.2979
(C8,C7,C9,C10)	71.4794
(C8,C7,C9,C13)	-56.0749
(C8,C7,C9,H39)	-173.1697
(H36,C7,C9,C10)	-172.3774
(H36,C7,C9,H13)	60.0683
(H36,C7,C9,H39)	-57.0265
(C7,C8,C11,C12)	-54.0332
(C7,C8,C11,C42)	67.0443
(C7,C8,C11,H43)	-176.8587
(H37,C8,C11,C12)	-175.6853
(H37,C8,C11,H42)	-54.6078
(H37,C8,C11,H43)	61.4893
(H38,C8,C11,C12)	68.392
(H38,C8,C11,H42)	-170.5305
(H38,C8,C11,H43)	-54.4335

Bond	Length
(C7,C9,C10,C3)	56.8271
(C7,C9,C10,H40)	177.5118
(C7,C9,C10,H41)	-62.4289
(C13,C9,C10,C3)	-176.6772
(C13,C9,C10,H40)	-55.9925
(C13,C9,C10,H41)	64.0668
(H39,C9,C10,C3)	-58.0127
(H39,C9,C10,H40)	62.672
(H39,C9,C10,H41)	-177.2688
(C7,C9,C13,C12)	57.7522
(C7,C9,C13,C17)	-173.9591
(C7,C9,C13,H45)	-57.9494
(C10,C9,C13,C12)	-68.1487
(C10,C9,C13,C17)	60.14
(C10,C9,C13,H45)	176.1497
(H39,C9,C13,C12)	173.7543
(H39,C9,C13,C17)	-57.957
(H39,C9,C13,H45)	58.0528
(C8,C11,C12,C13)	55.5943
(C8,C11,C12,C16)	-177.3826
(C8,C11,C12,H44)	-61.3495
(H42,C11,C12,C13)	-66.0334
(H42,C11,C12,C16)	60.9897
(H42,C11,C12,H44)	177.0227
(H43,C11,C12,C13)	178.6688
(H43,C11,C12,C16)	-54.3081
(H43,C11,C12,H44)	61.725
(C11,C12,C13,C9)	-56.0999
(C11,C12,C13,C17)	175.8495
(C11,C12,C13,H45)	59.5016
(C16,C12,C13,C9)	-179.5142
(C16,C12,C13,C17)	52.4353
(C16,C12,C13,H45)	-63.9127
(H44,C12,C13,C9)	60.9391
(H44,C12,C13,C17)	-67.1115
(H44,C12,C13,H45)	176.5406
(C11,C12,C16,H48)	-68.1829
(C11,C12,C16,H49)	51.2523
(C11,C12,C16,H50)	171.7469
(C13,C12,C16,H48)	55.4369
(C13,C12,C16,H49)	174.8721
(C13,C12,C16,H50)	-64.6333
(H44,C12,C16,H48)	175.3094
(H44,C12,C16,H49)	-65.2554
(H44,C12,C16,H50)	55.2391
(C9,C13,C17,C18)	38.3054
(C9,C13,C17,O19)	160.4187
(C9,C13,C17,H51)	-83.2834
(C12,C13,C17,C18)	164.6953
(C12,C13,C17,O19)	-73.1914
(C12,C13,C17,H51)	43.1066
(H45,C13,C17,C18)	-78.2876
(H45,C13,C17,O19)	43.8258
(H45,C13,C17,H51)	160.1237
(C1,C14,C15,C5)	-55.2318
(C1,C14,C15,O28)	-177.6755
(C1,C14,C15,H47)	64.2515
(C27,C14,C15,C5)	-178.0733
(C27,C14,C15,O28)	59.483
(C27,C14,C15,H47)	-58.59
(H46,C14,C15,C5)	62.7797
(H46,C14,C15,O28)	-59.664

Bond	Length
(H46,C14,C15,H47)	-177.737
(C1,C14,C27,N29)	162.2262
(C1,C14,C27,H66)	-74.6227
(C1,C14,C27,H67)	39.8805
(C15,C14,C27,N29)	-75.3363
(C15,C14,C27,H66)	47.8148
(C15,C14,C27,H67)	162.318
(H46,C14,C27,N29)	43.3759
(H46,C14,C27,H66)	166.527
(H46,C14,C27,H67)	-78.9698
(C5,C15,O28,C68)	58.998
(C14,C15,O28,C68)	-178.0574
(H47,C15,O28,C68)	-60.9478
(C13,C17,C18,C20)	-164.9521
(C13,C17,C18,C21)	66.4885
(C13,C17,C18,H52)	-50.8565
(O19,C17,C18,C20)	74.8227
(O19,C17,C18,C21)	-53.7366
(O19,C17,C18,H52)	-171.0816
(H51,C17,C18,C20)	-43.839
(H51,C17,C18,C21)	-172.3983
(C51,C17,C18,H52)	70.2567
(C13,C17,O19,H53)	-178.9114
(C18,C17,O19,H53)	-52.8828
(H51,C17,O19,H53)	65.5944
(C17,C18,C20,C22)	-159.5683
(C17,C18,C20,C23)	77.7108
(C17,C18,C20,H54)	-41.5494
(C21,C18,C20,C22)	-33.6424
(C21,C18,C20,C23)	-156.3634
(C21,C18,C20,H54)	84.3765
(H52,C18,C20,C22)	83.9941
(H52,C18,C20,C23)	-38.7268
(H52,C18,C20,H54)	-157.987
(C17,C18,C21,H55)	-54.6059
(C17,C18,C21,H56)	-173.6748
(C17,C18,C21,H57)	65.306
(C20,C18,C21,H55)	179.7726
(C20,C18,C21,H56)	60.7038
(C20,C18,C21,H57)	-60.3154
(H52,C18,C21,H55)	63.5116
(H52,C18,C21,H56)	-55.5573
(H52,C18,C21,H57)	-176.5764
(C18,C20,C22,O24)	80.3115
(C18,C20,C22,H58)	-160.5346
(C18,C20,C22,H59)	-41.834
(C23,C20,C22,H24)	-156.0793
(C23,C20,C22,H58)	-36.9254
(C23,C20,C22,H59)	81.7752
(H54,C20,C22,O24)	-40.3209
(H54,C20,C22,H58)	78.8331
(H54,C20,C22,H59)	-162.4664
(C18,C20,C23,H60)	-172.7942
(C18,C20,C23,H61)	67.1505
(C18,C20,C23,H62)	-52.509
(C22,C20,C23,H60)	61.3311
(C22,C20,C23,H61)	-58.7243
(C22,C20,C23,H62)	-178.3838
(H54,C20,C23,H60)	-52.5133
(H54,C20,C23,H61)	-172.5686
(H54,C20,C23,H62)	67.7718
(C20,C22,O24,C26)	-171.9022

Bond	Length
(H58,C22,O24,C26)	68.2501
(H59,C22,O24,C26)	-49.6067
(C22,O24,C26,O25)	71.0759
(C22,O24,C26,H64)	-166.6551
(C22,O24,C26,H65)	-47.3375
(H63,O25,C26,O24)	64.0574
(H63,O25,C26,H64)	-54.8738
(H63,O25,C26,H65)	-174.2266
(C14,C27,N29,H69)	-65.7955
(C14,C27,N29,H70)	50.794
(H66,C27,N29,H69)	170.9464
(H66,C27,N29,H70)	-72.4641
(H67,C27,N29,H69)	56.874
(H67,C27,N29,H70)	173.4634

Table S2. Frontier molecular orbital properties of the compound C2.

Property	Values
ϵ HOMO	-9.9
ϵ LUMO	-4.9
Energy gap ΔE	5.0
Ionisation energy ($I = \epsilon$ HOMO = -HOMO)	9.9
Electron Affinity ($A = \epsilon$ LUMO = -LUMO)	4.9
Global hardness ($\eta = (I-A)/2$)	2.5
Global softness ($S = 1/\eta$)	0.4
Chemical Potential ($\mu = -(I+A)/2$)	-7.4
Electronegativity ($\chi = -\mu$)	7.4
Electrophilicity index ($\omega = \mu^2/2\eta$)	8.3
Nucleophilicity index ($N = 1/\omega$)	0.1
Electronaccepting power ($\omega^+ = A^2/2(I-A)$)	0.5
Electron donating power ($\omega^- = I^2/2(I-A)$)	0.9

Table S3. Natural population analysis of the compound C2.

Atom No	Natural	Natural Population			Total
	Charge	Core	Valance	Rydberg	
C 1	-0.47452	1.99917	4.46163	0.01371	6.47452
C 2	-0.342	1.99899	4.33255	0.01046	6.342
C 3	0.57247	1.99913	3.39319	0.03521	5.42753
O 4	-0.53265	1.99977	6.52894	0.00394	8.53265
C 5	-0.49394	1.99912	4.48376	0.01106	6.49394
C 6	-0.24733	1.99908	4.2365	0.01176	6.24733
C 7	-0.25098	1.99907	4.23998	0.01193	6.25098
C 8	-0.47327	1.99917	4.46482	0.00928	6.47327
C 9	-0.25072	1.99909	4.24057	0.01106	6.25072
C 10	-0.5577	1.99908	4.5482	0.01043	6.5577
C 11	-0.46482	1.99918	4.45594	0.00969	6.46482
C 12	-0.24917	1.9991	4.23998	0.01009	6.24917
C 13	-0.26928	1.99904	4.256	0.01424	6.26928
C 14	-0.29444	1.99903	4.28373	0.01168	6.29444
C 15	0.08716	1.99891	3.90059	0.01333	5.91284
C 16	-0.68881	1.99928	4.6832	0.00633	6.68881
C 17	0.09035	1.99895	3.89583	0.01487	5.90965
C 18	-0.2816	1.99907	4.27226	0.01027	6.2816
O 19	-0.7641	1.9998	6.75815	0.00615	8.7641
C 20	-0.2706	1.99908	4.25975	0.01177	6.2706
C 21	-0.69918	1.99928	4.69186	0.00803	6.69918
C 22	-0.13736	1.99906	4.12705	0.01125	6.13736
C 23	-0.70238	1.99928	4.69718	0.00592	6.70238
O 24	-0.59716	1.99979	6.59093	0.00644	8.59716
O 25	-0.76008	1.99981	6.75443	0.00584	8.76008
C 26	0.15759	1.99934	3.82619	0.01688	5.84241
C 27	-0.27576	1.99919	4.26153	0.01504	6.27576
O 28	-0.76989	1.99979	6.76414	0.00596	8.76989
N 29	-0.91534	1.99954	5.90548	0.01032	7.91534

Atom No	Natural	Natural Population			
	Charge	Core	Valance	Rydberg	Total
H 30	0.24712	0.0000	0.75042	0.00246	0.75288
H 31	0.26403	0.0000	0.73409	0.00189	0.73597
H 32	0.26258	0.0000	0.73517	0.00225	0.73742
H 33	0.24003	0.0000	0.75778	0.00219	0.75997
H 34	0.24728	0.0000	0.75007	0.00265	0.75272
H 35	0.24788	0.0000	0.74938	0.00273	0.75212
H 36	0.25391	0.0000	0.74372	0.00237	0.74609
H 37	0.24849	0.0000	0.75005	0.00146	0.75151
H 38	0.22879	0.0000	0.76944	0.00177	0.77121
H 39	0.25256	0.0000	0.74531	0.00213	0.74744
H 40	0.26231	0.0000	0.73502	0.00267	0.73769
H 41	0.25762	0.0000	0.74092	0.00146	0.74238
H 42	0.23425	0.0000	0.76349	0.00225	0.76575
H 43	0.24769	0.0000	0.75069	0.00162	0.75231
H 44	0.23123	0.0000	0.76628	0.00249	0.76877
H 45	0.2614	0.0000	0.7357	0.00291	0.7386
H 46	0.24462	0.0000	0.75184	0.00353	0.75538
H 47	0.21776	0.0000	0.77908	0.00316	0.78224
H 48	0.24092	0.0000	0.7576	0.00148	0.75908
H 49	0.22683	0.0000	0.77168	0.00149	0.77317
H 50	0.25175	0.0000	0.74689	0.00136	0.74825
H 51	0.21785	0.0000	0.77979	0.00236	0.78215
H 52	0.24508	0.0000	0.75221	0.00271	0.75492
H 53	0.47552	0.0000	0.52293	0.00156	0.52448
H 54	0.23436	0.0000	0.76145	0.00419	0.76564
H 55	0.23183	0.0000	0.76691	0.00126	0.76817
H 56	0.23454	0.0000	0.76326	0.00219	0.76546
H 57	0.25339	0.0000	0.74417	0.00244	0.74661
H 58	0.21968	0.0000	0.77789	0.00243	0.78032
H 59	0.23128	0.0000	0.76564	0.00307	0.76872
H 60	0.24464	0.0000	0.75418	0.00118	0.75536
H 61	0.24343	0.0000	0.75512	0.00145	0.75657
H 62	0.23883	0.0000	0.76017	0.001	0.76117
H 63	0.47756	0.0000	0.52013	0.00231	0.52244
H 64	0.21396	0.0000	0.78397	0.00206	0.78604
H 65	0.20793	0.0000	0.78916	0.00292	0.79207
H 66	0.2268	0.0000	0.77078	0.00242	0.7732
H 67	0.23569	0.0000	0.76269	0.00162	0.76431
H 68	0.47389	0.0000	0.524	0.00212	0.52611
H 69	0.37511	0.0000	0.62304	0.00185	0.62489
H 70	0.40509	0.0000	0.59271	0.00221	0.59491

Table S4. Second-order perturbation theory fock matrix analysis of compound C2.

Donar	Type	ED/e	Acceptor	Type	ED/e	E(2) ^a	E(j-i) ^b	F(i,j) ^c
C 1 - C 2	σ	1.97778	C 1 - C 14	σ^*	0.01734	0.8	0.97	0.025
			C 1 - H 30	σ^*	0.01583	0.6	1.03	0.022
			C 1 - H 31	σ^*	0.01119	0.77	1.05	0.025
			C 2 - C 3	σ^*	0.06274	1.03	0.98	0.029
			C 2 - C 6	σ^*	0.03064	1.31	0.93	0.031
			C 2 - H 32	σ^*	0.02178	0.94	1.01	0.028
			C 3 - C 10	σ^*	0.05269	2.26	0.98	0.042
			C 6 - C 7	σ^*	0.02409	2.21	0.95	0.041
			C 14 - C 27	σ^*	0.04038	1.77	0.96	0.037
			C 1 - C 14	σ	1.97123	C 1 - C 2	σ^*	0.0175
C 14 - C 15	σ^*	0.03099				0.71	0.94	0.023
C 14 - C 27	σ^*	0.04038				0.62	0.95	0.022
C 14 - H 46	σ^*	0.02227				0.66	1.02	0.023
C 15 - O 28	σ^*	0.02878				3.21	0.8	0.045
C 27 - N 29	σ^*	0.00733			2.34	0.96	0.042	
			C 2 - H 32	σ^*	0.02178	2.92	0.9	0.046
C 1 - H 30	σ	1.98063	C 14 - H 46	σ^*	0.02227	2.66	0.92	0.044
C 1 - H 31	σ	1.97972	C 1 - C 2	σ^*	0.0175	0.56	0.86	0.02
C 2 - C 3	σ	1.97959	C 1 - C 2	σ^*	0.0175	1.06	1.01	0.029
C 2 - C 6	σ	1.95801	C 10 - H 40	σ^*	0.01169	1.14	1.09	0.032
			C 1 - C 2	σ^*	0.0175	1.03	0.96	0.028
			C 3 - O 4	π^*	0.01345	2.75	0.58	0.036

Donar	Type	ED/e	Acceptor	Type	ED/e	E(2) ^a	E(j-i) ^b	F(i,j) ^c
			C 5 - C 6	σ^*	0.02071	0.72	0.94	0.023
			C 5 - H 33	σ^*	0.01228	1.69	1	0.037
			C 6 - H 35	σ^*	0.02032	0.5	1	0.02
			C 7 - H 36	σ^*	0.0181	1.24	1.01	0.032
C 2 - H 32	σ	1.95484	C 1 - H 30	σ^*	0.01583	2.56	0.94	0.044
			C 3 - O 4	π^*	0.01345	5.7	0.49	0.048
			C 6 - H 35	σ^*	0.02032	2.52	0.92	0.043
C 3 - O 4	σ	1.9959	C 2 - C 3	σ^*	0.06274	0.81	1.45	0.031
			C 3 - C 10	σ^*	0.05269	0.74	1.45	0.03
C 3 - O 4	π	1.98343	C 2 - C 6	σ^*	0.03024	1.54	0.7	0.03
			C 2 - H 32	σ^*	0.02178	1.8	0.78	0.034
			C 9 - C 10	σ^*	0.02531	1.63	0.7	0.03
			C 10 - H 41	σ^*	0.01537	1.86	0.79	0.034
C 3 - C 10	σ	1.98417	C 1 - C 2	σ^*	0.0175	1.92	1.02	0.039
			C 2 - C 3	σ^*	0.06274	0.53	1.02	0.021
			C 9 - C 10	σ^*	0.02531	1.12	0.97	0.03
			C 9 - C 13	σ^*	0.02323	2.24	0.99	0.042
			C 10 - H 40	σ^*	0.01169	0.8	1.09	0.026
			C 10 - H 41	σ^*	0.01537	0.79	1.06	0.026
C 5 - C 6	σ	1.97308	C 5 - C 15	σ^*	0.02695	0.76	0.96	0.024
			C 5 - H 33	σ^*	0.01228	0.73	1.02	0.024
			C 5 - H 34	σ^*	0.01614	0.72	1.03	0.024
			C 6 - C 7	σ^*	0.02409	1	0.96	0.028
			C 6 - H 35	σ^*	0.02032	0.76	1.01	0.025
			C 7 - C 9	σ^*	0.02528	1.82	0.95	0.037
			C 15 - O 28	σ^*	0.02878	2.75	0.82	0.042
C 5 - C 15	σ	1.98275	C 14 - C 27	σ^*	0.04038	2.08	0.98	0.041
			C 15 - H 47	σ^*	0.03098	0.51	1.03	0.021
C 5 - H 33	σ	1.9798	C 14 - C 15	σ^*	0.03099	3.09	0.87	0.046
C 5 - H 34	σ	1.97969	C 15 - O 28	σ^*	0.02878	0.59	0.72	0.018
			C 15 - H 47	σ^*	0.03098	2.66	0.92	0.044
C 6 - C 7	σ	1.97304	C 7 - H 36	σ^*	0.0181	0.79	1.02	0.025
			C 8 - C 11	σ^*	0.01254	1.56	0.97	0.035
			C 9 - C 13	σ^*	0.02323	2.14	0.95	0.04
C 6 - H 35	σ	1.97261	C 2 - H 32	σ^*	0.02178	2.69	0.91	0.044
			C 5 - H 34	σ^*	0.01614	2.8	0.92	0.046
C 7 - C 8	σ	1.97855	C 8 - H 37	σ^*	0.01076	0.77	1.03	0.025
			C 8 - H 38	σ^*	0.01566	0.62	1.02	0.022
			C 9 - H 39	σ^*	0.01792	1.42	1.02	0.034
			C 11 - H 43	σ^*	0.01046	1.47	1.03	0.035
C 7 - C 9	σ	1.97148	C 7 - H 36	σ^*	0.0181	0.72	1.01	0.024
			C 8 - H 37	σ^*	0.01076	1.61	1.03	0.036
			C 9 - C 10	σ^*	0.02531	0.75	0.92	0.023
			C 9 - C 13	σ^*	0.02323	1.15	0.94	0.029
			C 9 - H 39	σ^*	0.01792	0.81	1.02	0.026
C 7 - H 36	σ	1.96989	C 8 - H 38	σ^*	0.01566	2.98	0.92	0.047
C 8 - C 11	σ	1.98315	C 11 - C 12	σ^*	0.01961	0.8	0.95	0.025
			C 11 - H 42	σ^*	0.01565	0.56	1.02	0.021
			C 11 - H 43	σ^*	0.01046	0.66	1.03	0.023
C 8 - H 37	σ	1.98067	C 11 - C 12	σ^*	0.01961	3.38	0.86	0.048
C 8 - H 38	σ	1.98103	C 11 - H 42	σ^*	0.01565	2.55	0.93	0.043
C 9 - C 10	σ	1.96467	C 10 - H 41	σ^*	0.01537	0.85	1.01	0.026
			C 13 - H 45	σ^*	0.01995	1.16	1.02	0.031
C 9 - C 13	σ	1.96919	C 13 - H 45	σ^*	0.01995	0.61	1.02	0.022
			C 17 - O 19	σ^*	0.02557	1.87	0.81	0.035
C 9 - H 39	σ	1.97318	C 10 - H 41	σ^*	0.01537	2.71	0.92	0.045
			C 12 - C 13	σ^*	0.02784	3.04	0.85	0.045
C 10 - H 40	σ	1.98071	C 9 - C 10	σ^*	0.02531	0.77	0.84	0.023
			C 10 - H 41	σ^*	0.01537	0.51	0.93	0.019
C 10 - H 41	σ	1.96404	C 3 - O 4	π^*	0.01345	1.57	1.03	0.036
			C 9 - H 39	σ^*	0.01792	2.33	0.95	0.042
C 11 - C 12	σ	1.9778	C 12 - H 44	σ^*	0.02126	0.66	1.01	0.023
			C 13 - C 17	σ^*	0.02732	2.64	0.93	0.044
			C 16 - H 50	σ^*	0.00718	1.59	1.05	0.037
C 11 - H 42	σ	1.98171	C 8 - H 38	σ^*	0.01566	2.72	0.92	0.045
			C 12 - H 44	σ^*	0.02126	2.7	0.92	0.045
C 11 - H 43	σ	1.98192	C 7 - C 8	σ^*	0.02422	3.26	0.85	0.047
			C 12 - C 13	σ^*	0.02784	3.2	0.84	0.046
C 12 - C 13	σ	1.97307	C 12 - H 44	σ^*	0.02126	0.59	1.01	0.022
			C 13 - C 17	σ^*	0.02732	0.9	0.93	0.026

Donar	Type	ED/e	Acceptor	Type	ED/e	E(2) ^a	E(j-i) ^b	F(i,j) ^c
			C 13 - H 45	σ^*	0.01995	0.85	1.01	0.026
			C 16 - H 49	σ^*	0.00785	1.19	1.02	0.031
C 12 - C 16	σ	1.9828	C 12 - H 44	σ^*	0.02126	0.74	1.01	0.024
			C 16 - H 48	σ^*	0.00814	0.58	1.03	0.022
			C 16 - H 49	σ^*	0.00785	0.53	1.02	0.021
			C 16 - H 50	σ^*	0.00718	0.75	1.05	0.025
C 12 - H 44	σ	1.97246	C 16 - H 48	σ^*	0.00814	2.62	0.94	0.044
C 13 - C 17	σ	1.97503	C 17 - H 51	σ^*	0.02842	0.75	1.03	0.025
			C 18 - C 20	σ^*	0.02904	1.45	0.94	0.033
			O 19 - H 53	σ^*	0.00953	1.69	1.01	0.037
C 13 - H 45	σ	1.96839	C 17 - H 51	σ^*	0.02842	2.26	0.91	0.041
C 14 - C 15	σ	1.97298	C 15 - H 47	σ^*	0.03098	0.63	1.02	0.023
			C 27 - H 67	σ^*	0.01206	1.02	1.05	0.029
			O 28 - H 68	σ^*	0.00782	1.73	1	0.037
C 14 - C 27	σ	1.98054	C 14 - C 15	σ^*	0.03099	0.89	0.94	0.026
			C 14 - H 46	σ^*	0.02227	0.78	1.01	0.025
			C 27 - H 67	σ^*	0.01206	0.52	1.03	0.021
C 14 - H 46	σ	1.97122	C 27 - H 66	σ^*	0.02101	2.17	0.93	0.04
C 15 - O 28	σ	1.99217	C 1 - C 14	σ^*	0.01734	1.58	1.16	0.038
C 15 - H 47	σ	1.98025	C 5 - H 34	σ^*	0.01614	2.57	0.94	0.044
			C 14 - H 46	σ^*	0.02227	2.68	0.95	0.045
C 16 - H 48	σ	1.99015	C 12 - H 44	σ^*	0.02126	2.53	0.91	0.043
C 16 - H 49	σ	1.98864	C 12 - C 13	σ^*	0.02784	3.83	0.83	0.051
C 16 - H 50	σ	1.98947	C 11 - C 12	σ^*	0.01961	2.71	0.84	0.043
			C 12 - C 16	σ^*	0.01198	0.52	0.86	0.019
C 17 - C 18	σ	1.97813	C 18 - H 52	σ^*	0.01985	0.83	1.03	0.026
			C 20 - C 22	σ^*	0.01957	1.97	0.96	0.039
			C 21 - H 56	σ^*	0.00841	1.56	1.05	0.036
C 17 - O 19	σ	1.99199	C 9 - C 13	σ^*	0.02323	1.88	1.13	0.041
			C 18 - H 52	σ^*	0.01985	1.2	1.22	0.034
C 17 - H 51	σ	1.98075	C 13 - H 45	σ^*	0.01995	2.17	0.95	0.04
			C 18 - C 21	σ^*	0.01453	2.87	0.88	0.045
C 18 - C 20	σ	1.97423	C 21 - H 55	σ^*	0.00714	1.31	1.03	0.033
			C 22 - H 58	σ^*	0.02106	1.16	1.01	0.031
			C 23 - H 60	σ^*	0.00549	1.38	1.02	0.034
C 18 - C 21	σ	1.98348	C 21 - H 55	σ^*	0.00714	0.67	1.03	0.024
			C 21 - H 56	σ^*	0.00841	0.69	1.04	0.024
			C 21 - H 57	σ^*	0.00939	0.66	1.05	0.024
C 18 - H 52	σ	1.9703	C 17 - O 19	σ^*	0.02557	3.94	0.72	0.048
			C 21 - H 57	σ^*	0.00939	2.59	0.95	0.045
O 19 - H 53	σ	1.98968	C 13 - C 17	σ^*	0.02732	2.03	1.07	0.042
C 20 - C 22	σ	1.98008	C 23 - H 62	σ^*	0.00712	1.68	1.05	0.038
			O 24 - C 26	σ^*	0.05687	1.97	0.86	0.037
C 20 - C 23	σ	1.98013	C 23 - H 61	σ^*	0.00795	0.5	1.02	0.02
			C 23 - H 62	σ^*	0.00712	0.63	1.03	0.023
C 20 - H 54	σ	1.97015	C 22 - H 59	σ^*	0.02253	2.4	0.93	0.042
			C 23 - H 61	σ^*	0.00795	2.63	0.93	0.044
C 21 - H 55	σ	1.98763	C 18 - C 20	σ^*	0.02904	3.62	0.83	0.049
			C 18 - C 21	σ^*	0.01453	0.54	0.86	0.019
C 21 - H 56	σ	1.98841	C 17 - C 18	σ^*	0.03515	3.01	0.84	0.045
C 21 - H 57	σ	1.99014	C 18 - C 21	σ^*	0.01453	0.52	0.85	0.019
			C 18 - H 52	σ^*	0.01985	2.47	0.93	0.043
C 22 - O 24	σ	1.98995	C 20 - C 23	σ^*	0.01093	1.42	1.12	0.036
			C 26 - H 64	σ^*	0.02339	0.94	1.21	0.03
C 22 - H 58	σ	1.98833	C 18 - C 20	σ^*	0.02904	3.08	0.86	0.046
			C 20 - C 22	σ^*	0.01957	0.58	0.88	0.02
C 22 - H 59	σ	1.98835	C 20 - H 54	σ^*	0.02431	2.17	0.95	0.041
C 23 - H 60	σ	1.98977	C 18 - C 20	σ^*	0.02904	3.07	0.85	0.046
C 23 - H 61	σ	1.98996	C 20 - H 54	σ^*	0.02431	2.52	0.93	0.043
C 23 - H 62	σ	1.98834	C 17 - H 51	σ^*	0.02842	0.57	0.94	0.021
			C 20 - C 22	σ^*	0.01957	2.59	0.87	0.043
O 24 - C 26	σ	1.99243	C 20 - C 22	σ^*	0.01957	1.25	1.17	0.034
O 25 - H 63	σ	1.99151	C 26 - H 65	σ^*	0.02792	1.76	1.16	0.04
C 26 - H 64	σ	1.98812	C 22 - O 24	σ^*	0.02524	3.6	0.78	0.047
C 26 - H 65	σ	1.98818	O 25 - H 63	σ^*	0.00998	3.01	0.94	0.048
C 27 - N 29	σ	1.99455	C 1 - C 14	σ^*	0.01734	1.34	1.06	0.034
C 27 - H 66	σ	1.98211	C 14 - H 46	σ^*	0.02227	2.68	0.92	0.044
			N 29 - H 69	σ^*	0.00713	2.91	0.97	0.048
C 27 - H 67	σ	1.98312	C 14 - C 15	σ^*	0.03099	3.25	0.85	0.047
			N 29 - H 70	σ^*	0.01578	2.99	1	0.049

Donar	Type	ED/e	Acceptor	Type	ED/e	E(2) ^a	E(j-i) ^b	F(i,j) ^c
O 28 - H 68	σ	1.99133	C 14 - C 15	σ^*	0.03099	2.25	1.08	0.044
N 29 - H 69	σ	1.99182	C 27 - H 66	σ^*	0.02101	2.08	1.04	0.042
N 29 - H 70	σ	1.99179	C 27 - H 67	σ^*	0.01206	2.14	1.05	0.042
LP(1)	O 4	1.97672	C 2 - C 3	σ^*	0.06274	1.91	1.07	0.041
LP(2)	O 4	1.90407	C 2 - C 3	σ^*	0.06274	19.03	0.65	0.1
LP(2)	O 4	1.90407	C 3 - C 10	σ^*	0.05269	17.98	0.65	0.097
LP(1)	O 19	1.97954	C 17 - H 51	σ^*	0.02842	1.47	1.01	0.034
LP(2)	O 19	1.96184	C 17 - C 18	σ^*	0.03515	5.46	0.64	0.053
LP(2)	O 19	1.96184	C 17 - H 51	σ^*	0.02842	4	0.73	0.048
LP(1)	O 24	1.96978	C 26 - H 65	σ^*	0.02792	1.93	1.01	0.039
LP(2)	O 24	1.9077	O 25 - C 26	σ^*	0.05372	14.75	0.52	0.079
LP(1)	O 25	1.9847	C 26 - H 64	σ^*	0.02339	1.98	1.03	0.04
LP(2)	O 25	1.93118	O 24 - C 26	σ^*	0.05687	14.04	0.56	0.079
LP(1)	O 28	1.97805	C 15 - H 47	σ^*	0.03098	2.81	0.97	0.047
LP(2)	O 28	1.98327	C 5 - C 15	σ^*	0.02695	5.94	0.71	0.058
LP(1)	N 29	1.95112	C 14 - C 27	σ^*	0.04038	9.91	0.6	0.069
LP(1)	N 29	1.95112	C 27 - H 66	σ^*	0.02101	2.65	0.68	0.038