

THE EVOLUTION OF PLANT PHYSIOLOGY

*Edited by
Alan R. Hemsley and Imogen Poole*



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List of contributors

Pieter Baas, Nationaal Herbarium Nederland, Universiteit Leiden Branch, PO Box 9514, 2300 RA Leiden, The Netherlands

Hendrik Bargel, Institut für Botanik, Zellescher Weg 22, 01062 Dresden, Germany

Wilhelm Barthlott, Botanisches Institut, Abteilung Systematik und Biodiversität, Meckenheimer Allee 170, 53115 Bonn, Germany

David J Beerling, Department of Animal and Plant Sciences, University of Sheffield, Sheffield S10 2TN, UK

Pim F van Bergen, Organic Geochemistry, Earth Sciences, Utrecht University, PO Box 80021, 3508 TA Utrecht, The Netherlands

Peter C Bilkey, AgResearch International, 7841 East Oakbrook Circle, Madison, WI 53717, USA

Peter Blokker, Vrije Universiteit, Analytical Chemistry and Applied Spectroscopy, Faculty of Sciences, De Boelelaan 1083, 1081 HV Amsterdam, The Netherlands

William J Bond, Department of Botany, University of Cape Town, Private Bag, Rondebosch, 7700 South Africa

Adrianus C Borstlap, Transport Physiology Research Group, Department of Plant Sciences, Utrecht University, Sorbonnelaan 16, NL-3584 CA Utrecht, The Netherlands

Tim Brodribb, Parque Nacional Santa Rosa, Costa Rica

William G Chaloner, Department of Geology, Royal Holloway, University of London, Egham Hill, Egham, Surrey TW20 0EX, UK

Mark W Chase, Molecular Systematics Section, Jodrell Laboratory, Royal Botanic Gardens, Kew, Richmond TW9 3DS, UK

Jerry D Cohen, Department of Horticultural Science, University of Minnesota, Saint Paul, MN 55108, USA

Margaret E Collinson, Department of Geology, Royal Holloway University of London, Egham, Surrey TW20 0EX, UK

Martha E Cook, Department of Biological Sciences, Illinois State University, Normal, IL, USA

Todd J Cooke, Department of Cell Biology and Molecular Genetics, University of Maryland, College Park, MD 20742, USA

- Stephen D Davis, Pepperdine University, Natural Science Division, Malibu, CA 90263-4321, USA
- Michael E Day, Department of Forest Ecosystem Science, University of Maine, 5755 Nutting Hall, Orono, Maine, USA
- Steven Dessein, Laboratory of Plant Systematics, Institute of Botany and Microbiology, K.U.Leuven, Kasteelpark Arenberg 31, B-3001 Leuven, Belgium
- Joost van Dongen, Max Planck Institute of Molecular Plant Physiology, Am Mühlenberg 1, 14476 Golm, Germany
- Dianne Edwards, School of Earth, Ocean and Planetary Sciences, Cardiff University, PO Box 914, Cardiff CF1 3YE, UK
- Frank W Ewers, Michigan State University, Department of Plant Biology, East Lansing, MI 48824, USA
- Richard D Firn, Department of Biology, University of York, York YO1 5DD, UK
- Madeline M Fisher, Wisconsin Alumni Research Foundation, University of Wisconsin, Madison, WI 53706, USA
- James M Graham, Department of Botany, University of Wisconsin, Madison, WI 53706, USA
- Linda E Graham, Department of Botany, University of Wisconsin, Madison, WI 53706, USA
- Howard Griffiths, Department of Plant Sciences, Downing Street, University of Cambridge, Cambridge CB2 3EA, UK
- John M Hackney, Department of Botany, University of Wisconsin–Madison, WI 53706, USA
- David T Hanson, Molecular Plant Physiology, Research School of Biological Sciences, National University, Canberra, ACT 2601, Australia
- Robert S Hill, Centre for Evolutionary Biology and Biodiversity, South Australian Museum, Adelaide, South Australia 5000; Department of Environmental Biology, Adelaide University, South Australia 5005
- Martin Ingrouille, School of Biological and Chemical Sciences, Birkbeck University of London, Malet Street, London WC1E 7HX, UK
- Richard Jagels, Department of Forest Ecosystem Science, University of Maine, 5755 Nutting Hall, Orono, Maine, USA
- Steven Jansen, Laboratory of Plant Systematics, Institute of Botany and Microbiology, K.U.Leuven, Kasteelpark Arenberg 31, B-3001 Leuven, Belgium
- Philip John, School of Plant Sciences, The University of Reading, Reading RG6 6AS, UK
- Clive G Jones, Institute of Ecosystem Studies, Box AB, Millbrook NY 12545-0129 Millbrook, USA
- Kerstin Koch, Botanisches Institut, Abteilung Systematik und Biodiversität, Meckenheimer Allee 170, 53115 Bonn, Germany

- Robin B Kodner, Department of Organismal and Evolutionary Biology, Harvard University, Cambridge, MA 02138, USA
- Pieter J C Kuiper, Department of Plant Biology, University of Groningen, The Netherlands
- Cécile M H Lapré, Freelance Research Consultancy, Haren, The Netherlands
- Tracy Lawson, Department of Biological Sciences, University of Essex, Wivenhoe Park, Colchester, Essex CO4 3SQ, UK
- Jan W de Leeuw, Organic Geochemistry, Earth Sciences, Utrecht University, PO Box 80021, 3508 TA Utrecht; Marine Biogeochemistry and Toxicology, Royal NIOZ, PO Box 59, AB Den Burg, Texel, The Netherlands
- Ben A LePage, Department of Earth and Environmental Science, University of Pennsylvania, 240 S. 33rd St, Philadelphia, PA 1910-6316, USA
- Kate Maxwell, Department of Plant Sciences, Downing Street, University of Cambridge, Cambridge CB2 3EA, UK
- Guy F Midgley, Ecology and Conservation, Kirstenbosch Research Centre, National Botanical Institute, Private Bag X7 Claremont, 7735 South Africa
- James I L Morison, Department of Biological Sciences, University of Essex, Wivenhoe Park, Colchester, Essex CO4 3SQ, UK
- Christoph Neinhuis, Institut für Botanik, Zellescher Weg 22, 01062 Dresden, Germany
- John Obst, UDSA Forest Products Laboratory, Madison, WI 53706, USA
- Colin P Osborne, Department of Animal and Plant Sciences, University of Sheffield, Sheffield S10 2TN, UK
- Christopher N Page, Honorary Associate, Royal Botanic Garden, Edinburgh. Correspondence: Cornwall Geological Museum, Penzance TR18 2QR, UK
- Norman W Pammenter, School of Life and Environmental Sciences, University of Natal, Durban, 4041 South Africa
- DorothyBelle Poli, Department of Cell Biology and Molecular Genetics, University of Maryland, College Park, MD 20742, USA
- John A Raven, Division of Environmental and Applied Biology, Biological Sciences Institute, School of Life Sciences, University of Dundee, Dundee DD1 4HN, UK
- Elizabeth A Reynolds, School of Plant Sciences, The University of Reading, Reading RG6 6AS, UK
- David Richardson, Department of Plant Sciences, Downing Street, University of Cambridge, Cambridge CB2 3EA, UK
- Elmar Robbrecht, National Botanic Garden of Belgium, Domein van Bouchout, B-1860 Meise, Belgium
- Wendy Robe, Department of Plant Sciences, Downing Street, University of Cambridge, Cambridge CB2 3EA, UK
- Nick Rowe, Botanique et bioinformatique de l'architecture des plantes, UMR 5120, TA 40/PS 2 Boulevard de la Lironde, 34398 Montpellier, cedex 5, France

-
- Lukas Schreiber, Botanisches Institut, Abteilung Ökophysiologie, Kirschallee 1, 53115 Bonn, Germany
- Jaap S Sinninghe Damsté, Organic Geochemistry, Earth Sciences, Utrecht University, PO Box 80021, 3508 TA Utrecht; Marine Biogeochemistry and Toxicology, Royal NIOZ, PO Box 59, AB Den Burg, Texel, The Netherlands
- Erik Smets, Laboratory of Plant Systematics, Institute of Botany and Microbiology, K.U.Leuven, Kasteelpark Arenberg 31, B-3001 Leuven, Belgium
- Thomas Speck, Plant Biomechanics Group, Botanical Garden of the Albert-Ludwigs-Universität, Schänzlestrasse 1, D-79104 Freiburg, Germany
- David R Vann, Department of Earth and Environmental Science, University of Pennsylvania, 240 S. 33rd St, Philadelphia, PA 1910-6316, USA
- Toshihiro Watanabe, Graduate School of Agriculture, Hokkaido University, Sapporo, 0608589, Japan
- Charles H Wellman, Department of Animal and Plant Sciences, University of Sheffield, Alfred Denny Building, Western Bank, Sheffield S10 2TN, UK
- Elisabeth A Wheeler, North Carolina State University, Department of Wood & Paper Science, Box 8005, Raleigh, NC 27695-8005, USA
- Lee W Wilcox, Department of Botany, University of Wisconsin, Madison, WI 53706, USA
- Christopher J Williams, Department of Earth and Environmental Science, University of Pennsylvania, 240 S. 33rd St, Philadelphia, PA 1910-6316, USA

Preface

Despite its extensive history as a field of study, plant physiology has rarely been considered by palaeobotanists in the context of the fossil record. Similarly, those involved with modern physiology have rarely considered that the fossil record might have anything to offer with respect to a modern view of plants and their responses to environmental change. This is of no great surprise since few fossils are amenable to the traditional methods used in modern physiology and fossils are, quite rightly, viewed as being deficient in useful characters when compared with living specimens. However, over the past few years, the emerging field of palaeophytophysiology (the study of the physiology of living plant ancestors and their extinct relatives) has begun to redress this imbalance and the wealth of physiological information hidden within the palaeobotanical realm is finally being unearthed. It was with these thoughts in mind that the symposium, sharing the same title as this book, was organized jointly between the Linnean Society of London (Palaeobotany Specialist Group) and the Royal Botanic Gardens, Kew with sponsorship from the Annals of Botany Company. Its aim was to bring together researchers from a range of disciplines, each with their own perspective on the overlap between an interest in plant physiology and the botanical fossil record. At this unique and somewhat unusual event we were able to begin considering the mechanisms, responses, effects and subsequent repercussions of plant physiology through geological time.

The synthesis of such previously disparate disciplines has required the development of new techniques and interpretative frameworks. These have brought about an understanding of palaeophytophysiology in its widest context and have provided exciting ideas for physiologists, palaeobotanists and climate modellers alike. Cutting edge developments in this novel field provide the basis of this book drawing on subjects as distant as animal evolution, biochemistry, computer modelling, phylogenetic analyses, organic geochemistry and plant ecology to provide greater insights into the evolution of plant physiology in its widest context.

The origins of plant physiology

We begin with a focus on the physiology of early land plants with reference to the problems faced by bryophytes and embryophytes; their photosynthetic limitations and the mechanistic means of overcoming associated physiological limitations. The necessary advances in spore wall physiology, involving crucial adaptive responses to the new harsh subaerial environment, which ensured a successful invasion of the land, are discussed.

Evolution of plant physiology from the molecular level

Any consideration of physiological evolution must include reference to the associated biochemistry. This section delves deeper into our understanding of how and why selected

molecules and molecular structures have played an important role in palaeophytophysiology. These chapters provide an introduction to this area with focus on specific biomolecules such as auxin, aquaporins, ethylene and phenolics and their resulting influence on the plants themselves concurrent with evidence from the fossil record. Biomacromolecules with protective and supportive roles are considered.

Evolution of anatomical physiology

Physiological adaptation to environmental variables cannot improve without associated advances in morphology and anatomy. Evolutionary development of the leaf and its associated anatomy is an obvious example but without an improved hydraulic system the functioning of the leaves would undoubtedly fail. This section focuses on the development of the megaphyll leaf, the stomata (a crucial advancement for photosynthesis and controlling water loss through transpiration) and the plan of hydraulic delivery of water throughout the plant. This section also considers physiology with respect to reproduction and its phylogenetic utility.

Evolution of environmental and ecosystem physiology

Evolutionary adaptation is inevitably a response to environmental change. Throughout the course of geological time, the environments in which plants grew have been changing, often radically and irreversibly. Therefore it is only right to include a section on their adaptations to environments. Such adaptations include responses to factors as far reaching as the unique polar regime, specific elements present within the soil and large-scale relationships between physiology, environment and species distribution.

This broad, but readable collection of contributions from leading specialists in systematics, plant physiology, palaeobotany and bio/geochemistry provides an essential resource base for both the newcomer and the established researcher in this new field. The contributions are individual, thought provoking and sometimes even provocative. In some cases authors disagree, but we view this as inevitable in a newly emerging field. Already, new terminology and conceptual frameworks are accruing; clearly the idea of 'trade-offs' among past physiological requirements permeates this book. Our personal interest and enthusiasm for this research area is only dampened somewhat by the realization that previous publications, and the chapters that make up this volume, represent only a small body of work and that this currently constrains the intellectual walls against which we push. We are confident, however, that increasing interest, inspiring curiosity-driven research, and the obvious relevance of palaeophytophysiology to all aspects of palaeoecology and environmental change, coupled with the development of newly emerging techniques, will promote rigorous evaluation and notable expansion of this field. Regardless of the ultimate conclusions, palaeophytophysiology certainly merits further investigation and we are confident that this volume will act as a seed for the pursuit and dispersal of additional, more specialized and comprehensive texts in the not too distant future.

Finally, we would like to thank all those who helped make the symposium and this publication a reality, including the independent reviewers for their time and effort spent on each chapter. Special mention goes to John Marsden and the staff at the Linnean Society

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Alan R Hemsley
Imogen Poole

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Turning the land green: inferring photosynthetic physiology and diffusive limitations in early bryophytes

Howard Griffiths, Kate Maxwell, David Richardson and
Wendy Robe

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Introduction

The tremendous interest in the form and function of the earliest land plants mirrors the enormous effect such plants had on the early climate, increasing the drawdown of CO₂ directly through photosynthesis and indirectly via weathering (Berner, 1998), both likely to lower global temperatures and encourage additional diversification on land (Algeo *et al.*, 2001). Despite recent developments in our understanding of land plant evolution, the physiological ecologist could feel somewhat marginalized, particularly if wary of engaging in *post hoc* speculations and reconstructions. There seems little doubt that bryophytes were key early players, since the fossil record has revealed some exquisite examples of early morphological details (Edwards *et al.*, 1995, 1998; Niklas, 1997). For many specimens, spore structure is consistent with bryophytes occurring throughout the Silurian,

particularly when associated with fossilized axes containing non-tracheophyte conducting elements in the Late Silurian/Early Devonian (Edwards, 1998, 2000). There has been considerable debate regarding the phyletic origins of bryophytes (Kenrick and Crane, 1997; Niklas, 1997; Renzaglia *et al.*, 2000; Kenrick, 2000). Despite the widespread use of *rbcL* as a phylogenetic marker (Chase *et al.*, 1993; Lewis *et al.*, 1997; Qiu and Palmer, 1999), analysis of the coevolution of Rubisco kinetic properties and variations in CO₂ concentrating mechanisms (CCM) have, in equivalent terms, received less attention (but see Badger and Andrews, 1987; Badger *et al.*, 1998; Raven *et al.*, 1998; Raven, 2000). Given that Rubisco is arguably the most abundant and important protein on Earth, such deficiencies need to be redressed.

Additionally, there have been few attempts to examine constraints to Rubisco carboxylation, mesophyll conductance and light utilization in extant representatives of early land-plant life-forms, such as the bryophytes (but see Green and Snelgar, 1982; Proctor *et al.*, 1992; Green and Lange, 1994; Deltoro *et al.*, 1998; Green *et al.*, 1998; Csintalan *et al.*, 1999; Zotz *et al.*, 2000; Proctor, 2001). This is despite many theoretical approaches (Raven, 1977, 1995; Edwards *et al.*, 1998), which have also called for additional measurements of water relations and photosynthetic characteristics of bryophytes. Accordingly, it is the aim of this chapter to redress the imbalance in some of these approaches and to consider why terrestrial land plants did *not* adopt the more widely used biophysical CCMs found in most algae and then in the Anthocerotae (Smith and Griffiths, 1996a,b), only to develop biochemical CCMs such as the C₄ pathway and crassulacean acid metabolism (CAM) much later in plant evolution.

Phylogeny of bryophytes

There has been considerable debate in the literature on whether the bryophytes (mosses, liverworts and hornworts) represent a monophyletic or paraphyletic group, relative to the tracheophytes (Kenrick and Crane, 1997; Kenrick, 2000). Based on a suite of morphological and molecular characteristics, Renzaglia *et al.* (2000) propose that the hornworts were the earliest divergent clade. Advanced features associated with reproductive and sporophyte development, as well as stomata and conducting tissues, arose in parallel and were not considered homologous across the bryophytes (Renzaglia *et al.*, 2000; Ligrone *et al.*, 2000). Other evidence suggests that hornworts represent the basal topology of the land-plant phylogenetic tree: from Rubisco large subunit (chloroplast *rbcL*) and small subunit rDNA sequences (Nickrent *et al.*, 2000), as well as mitochondrial *nad5* (Beckert *et al.*, 1999) and from an analysis of marchantioid liverwort radiation (Wheeler, 2000). Alternatively, liverworts have been suggested to be the earliest land plants, with hornworts monophyletic with mosses and closer to the tracheophytes, as identified by mitochondrial DNA markers and *rbcL* sequences (Lewis *et al.*, 1977; Qiu *et al.*, 1998; Qiu and Palmer, 1999).

Three alternative strategies were proposed by Kenrick (2000), based on a combination of traditional and molecular phylogenies, but ultimately he suggested that mosses were the immediate progenitor of higher plants in a progression including extinct protracheophytes. In conclusion, it seems that the three bryophyte groups are sufficiently similar for all authorities to support the notion strongly that they gave rise to the tracheophytes in a paraphyletic fashion, although the precise relationships are still to be resolved. Even the fossil record is not helpful here, since the hepatic characteristics which might be expected to be associated with spores and sporangia in the mid-Ordovician (460 million years (Ma))

are not well represented even in the Late Silurian, as compared to the megafossil record of protracheophytes, which occurs more clearly in the Early Devonian (Edwards, 2000; Kenrick, 2000). However, we note the warning given by Schuster (1981) in interpreting the phylogenetic progression, whereby many of the bryophyte lines were ultimately unsuccessful in colonizing land, and many of the more advanced liverworts families diversified in moist microclimates beneath angiosperms some 300 Ma later.

Rubisco: a discriminating marker for photosynthetic metabolism

Excursions in the stable carbon isotope record have long been used to infer changes in mass balance of $^{13}\text{C}:^{12}\text{C}$, which represent changes in the partitioning between geosphere and biosphere (Schidlowski, 2001). At present, the source air is progressively being depleted in CO_2 as we return the equivalent of some 60 years of net C_3 photosynthesis to the atmosphere by means of fossil fuel combustion (Hall and Rao, 1994). Carbon isotopes can also be used to distinguish photosynthetic pathways, such that a low discrimination (more enriched ^{13}C signal) is associated with terrestrial C_4 and CAM pathway and aquatic CCMs (Farquhar *et al.*, 1989; Griffiths *et al.*, 1999). In Figure 1.1 we collate data for a variety of such photosynthetic pathways, including terrestrial bromeliads (Figure 1.1A), which show the traditional bimodal distribution of carbon isotope discrimination (a measure which corrects the measured $\delta^{13}\text{C}$ of organic material for source CO_2 contribution to provide a positive value of biological discrimination). Thus CAM and C_4 plants show a lower discrimination because of the biochemical CCM and the action of PEP carboxylase, which suppress the inherent discrimination of Rubisco; in C_3 plants, this potentially high value of Rubisco fractionation is tempered by the diffusive limitation imposed by stomata, such that lower values of discrimination can be used to infer high water-use efficiency under comparable growth conditions

The biophysical CCM in algae and cyanobacteria is normally associated with low values of carbon isotope discrimination (Beardall *et al.*, 1982; Máguas *et al.*, 1995) and the values for lichens are included in Figure 1.1B to show the effect of assimilating CO_2 when high rates of respiratory CO_2 (from the associated fungal partner, the mycobiont) are presented to the photobiont. There are lessons here for bryophytes, which normally grow appressed to the soil substrate and hence would be likely to receive a respiratory CO_2 bonus (Raven, 2000; Raven and Edwards, 2001). Despite this, an analysis of the carbon isotope discrimination in a number of bryophyte species also shows a bimodal pattern much more closely allied to the C_4 /CAM range (Figure 1.1C), because of the operation of a biophysical CCM in some hornworts (Smith and Griffiths, 1996a,b, 2000). Therefore, carbon isotopes provide one means to distinguish the occurrence of a CCM in bryophytes.

In addition, as shown below, we can also characterize the expression and activity of a CCM by measuring carbon isotope discrimination instantaneously during photosynthesis. Together with other measures, such as CO_2 compensation point, accumulation of an internal pool of dissolved inorganic carbon (DIC) and high carboxylation efficiency, it is possible to diagnose the operation of a CCM (Smith and Griffiths, 1996a,b). However, it should be noted that these are mostly indirect measures, and care should be taken in using CO_2 compensation points as a primary means of identifying CCM activity (Badger *et al.*, 1998; Raven *et al.*, 2000).

Why, then, does Rubisco need this type of photosynthetic turbocharger? The answer lies in the kinetic deficiencies of this extraordinary enzyme: often characterized as slow

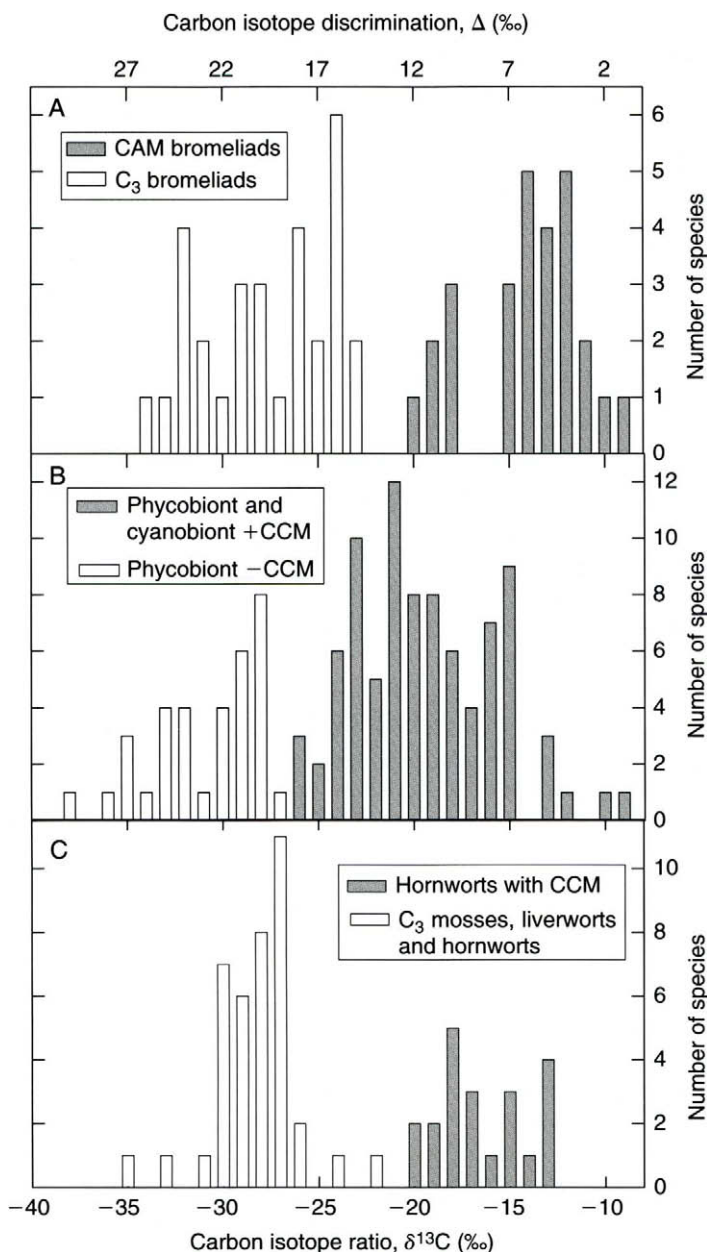


Figure 1.1 Carbon isotope ratio ($\delta^{13}\text{C}$) and discrimination (Δ) in organic material of contrasting plant groups: (A) bromeliads from Trinidad, showing distribution of C_3 and CAM photosynthetic pathways (data redrawn from Griffiths and Smith, 1983); (B) lichens, showing C_3 -like discrimination associated with non-pyrenoidal algal photobionts and the more C_4 -like signal associated with biophysical CO_2 concentrating mechanism (CCM) in pyrenoid (phycobiont) and carboxysome (cyanobacterial) of photobionts (data redrawn from Máguas *et al.*, 1995; Smith and Griffiths, 1996b); (C) bryophytes, showing C_3 -like signal in most mosses and liverworts, with the CCM in Anthocerotae possessing a chloroplast pyrenoid (data redrawn from Proctor *et al.*, 1992; Smith and Griffiths, 1996b).

and inefficient, it has also been suggested to have evolved as a 'qwerty' enzyme, whereby, like the first typewriter keyboard which was designed to *slow* the typist, carboxylation efficiency is deliberately inefficient and prevents the total drawdown of CO₂ in the biosphere (Nisbet and Sleep, 2001). Additionally, having evolved prior to oxygenation of the atmosphere, this enzyme has a potentially fatal predilection for O₂ (as well as CO₂), leading to the production of phosphoglycolate, which can either be excreted (some algae: Raven *et al.*, 2000) or metabolized in photorespiration. Such competition between gaseous substrates would intuitively be over-run by the current molar ratios of CO₂:O₂ in the atmosphere. Indeed, Rubisco affinity for CO₂ is low (K_m , or $K_{0.5}$, the substrate concentration required to half-saturate the enzyme), around 10–20 μmol CO₂ at 25°C, which is close to the CO₂ concentration in the cell cytosol. Thus, the enzyme should only ever be able to operate at half maximum velocity under normal conditions of diffusive CO₂ supply. However, the relative solubility of O₂ is lower than CO₂ at moderate temperatures, with O₂ now only some 25-fold higher in concentration. Another saving grace is that the affinity for O₂ is also relatively lower, around twice the dissolved O₂ concentration at 25°C, so that under these conditions Rubisco operates with the ratio of carboxylase: oxygenase rates (v_c/v_o) at around 2.5.

Rubisco is also catalytically slow, having a turnover rate (k_{cat}) of some 3 s⁻¹, as compared to 30 000 s⁻¹ for carbonic anhydrase, an enzyme associated with the interconversion of bicarbonate and CO₂ in solution as a substrate for Rubisco, as well as central to many types of CCM activity. There is considerable variation in the kinetic properties of Rubisco, with one key indicator being the specificity factor (τ or S_{rel}), which reflects the selectivity for CO₂ over O₂. In general terms, cyanobacteria and other primitive (L2) forms of the enzyme have low specificity, at 25°C, while there seems to have been a progressive improvement from chlorophyte algae through to higher plants, and then, most surprisingly, to certain thermophilic red algae (Badger and Andrews, 1997; Badger *et al.*, 1998; Raven, 2000). While this is usually reflected in the K_m for CO₂, there have been certain studies using site-directed mutagenesis which have led to dramatic increases in the K_m for O₂ (Zhu *et al.*, 1998; Schlitter and Wildner, 2000). There is also usually a 'trade-off' between specificity and catalytic turnover, such that an enzyme with a high affinity for CO₂ tends to show a much lower k_{cat} . At any event, we now urgently need a more detailed survey of Rubisco specificity for the three bryophyte groups if we are to map the kinetic properties on to the wealth of phylogenetic information available to date.

What are the implications for Rubisco in bryophytes and the colonization of land? The elevated CO₂ likely to have been prevalent at 460 my would have suppressed oxygenase activity and photorespiration (Raven, 2000; Sage, 2002). Secondly, appressed to the soil surface, the respiratory bonus from today's organically enriched substrates may help to offset any oxygenase activity, particularly if thallus surfaces limit diffusive uptake of CO₂ (Raven, 2000; Raven and Edwards, 2001). In this regard, it is important to note that CO₂ diffuses 10 000 times more slowly in water than in air and that surficial films of water on bryophyte thalli can impose a significant limitation to CO₂ uptake. Therefore, an increasing degree of ventilation and internalization of air spaces seen in liverwort thalli represent a progression towards restricting external diffusion limitations at the thallus surface. Finally, the occurrence of a CCM in hornworts suggests that at some stage during the colonization of the land, the energetic differences associated with powering active transport for the CCM may have been replaced by those associated with recycling carbon skeletons during photorespiration. We evaluate the relative costs of these limitations experimentally below.

Life on land: caught in a compromising situation

An important point has recently been raised regarding the pattern of land plant evolution (Sage, 2002): while we accept that land plants evolved from a group within the Charophyceae, it is perhaps no coincidence that this has been the only group to evolve a high efficiency photorespiratory pathway to dispose of oxygenase products for aerial organs (see also Raven *et al.*, 2000; Raven, 2000). When moving onto land, two irreconcilable problems arose for plants: the compromise between water loss and desiccation and the need to dispose of glycolate as an oxygenase waste product in an aerial environment. This, Sage (2002) suggests, provides compelling evidence for the origins of land plants *via* the only group to evolve an effective photorespiratory pathway associated with the development of the peroxisome. The subsequent exploitation of the aerial environment has left us with a green world rather than red, yellow or brown should one of the other classes of algae have come to dominate. However, if hornworts were basal to the phylogeny of land plants, what happened to the genetic capacity to express a CCM, and why was it not more widely adopted?

In addition to direct effects of O₂, we have also alluded to other possible problems for a thalloid life form – the compromise between diffusion limitation and minimizing the thickness of water films through which CO₂ must diffuse (*via* internalization of air spaces); the difficulties of occupying a high UV world and need to develop mechanisms to control and dissipate excess photon energy; the need for additional C reserves to be allocated for lignin and structural support rather than simply for reproduction; and finally, the need to accommodate air spaces and operation of stomata (Raven, 1995, 2000). In addition, another problem might have been high temperatures (Algeo *et al.*, 2001), which in themselves promote the rate of oxygenase activity, spurred on by the double disadvantage of O₂ being increasingly soluble at high temperatures relative to CO₂, and the disproportionate shift in Rubisco S_{rel} in favour of O₂ at high temperatures. A final point to consider is whether Rubisco activase would have been present, and if so, operational under these conditions. Recent evidence has suggested that at temperatures close to 40°C, the activase does not activate Rubisco as effectively (Crafts-Brandner and Salvucci, 2000), rather acting to protect the enzyme (Rokka *et al.*, 2001).

Why is there no biophysical CCM in terrestrial plants other than hornworts?

Two features are required to allow the development of a CCM: the first is a compartment within which CO₂ may be concentrated. For C₄ plants, this is the bundle sheath; for CAM, in chlorenchyma throughout the entire leaf or cladode, as stomata close in the light; for a CCM, in cyanobacteria and phycobiont algae, some means to concentrate Rubisco and generate elevated CO₂, respectively, via the carboxysome and the pyrenoid. Secondly, some mechanism to concentrate CO₂, whether via biochemistry (C₄, CAM) or a biophysical CCM (cyanobacteria, algae and hornworts). The distribution of pyrenoids and association with single chloroplasts in algal cells has been reviewed in the context of the activity of the CCM and changes in Rubisco kinetics (Badger *et al.*, 1998). The association between pyrenoids, Rubisco and other Calvin cycle enzymes had been long established (Vaughn *et al.*, 1990; McKay and Gibbs, 1991). Earlier studies had suggested that the concentration of Rubisco and Rubisco activase in the pyrenoid reflected the evolutionary

progression from uniplastidic to more advanced multiplastidic systems, with Rubisco distributed throughout the stroma (McKay and Gibbs, 1991). It now seems clear that the pyrenoid is consistently associated with CCM activity (Badger *et al.*, 1993, 1998; Palmqvist, 1993; Máguas *et al.*, 1995; Smith and Griffiths, 1996a,b). Indeed, there may be specialized thylakoid lamellae which are inserted through the pyrenoid which are enriched in Photosystem I (PSI). Such observations have led to suggestions that spatial separation of O₂ evolving PSII is one advantage (Pronino and Semenenko, 1992), and that cyclic electron flux may contribute to the ATP requirements of the CCM (Badger *et al.*, 1998). Whether this also indirectly generates the required pH environment for a specific intrathylakoid carbonic anhydrase enzyme still requires validation (Raven, 1997). Alternatively, pseudocyclic ATP generation has been associated with pyrenoid function (Sültemeyer *et al.*, 1993). Most recently, changes in allocation of Rubisco to the pyrenoid have been associated with diurnal changes in dinoflagellate photosynthesis (Nassoury *et al.*, 2001). In conclusion, theoretical models of eukaryote CCM activity strongly support the role of the pyrenoid in facilitating CCM activity, together with a central role for carbonic anhydrase (Badger *et al.*, 1998; Thoms *et al.*, 2001).

However, the evidence that the loss of the pyrenoid (and hence CCM) in the hornworts is associated with the multiplastidic condition is compelling, and is mirrored in a similar progression in unicellular algae (Smith and Griffiths, 1996b; Badger *et al.*, 1998). Despite analyses of the morphological correlates (Brown and Lemon, 1990), one key feature requiring more detailed analysis is the molecular control of chloroplast differentiation, so as to determine why genetic expression of the pyrenoid was lost, seemingly irrevocably. It seems likely that the development of the multiplastidic cell would bring about a dramatic increase in mesophyll conductance, allowing smaller chloroplasts to be appressed close to air spaces in increasingly well-ventilated photosynthetic thalli, thereby dramatically reducing the liquid- and lipid-phase diffusion limitation in the aerial environment. Perhaps it is this 'trade-off' which allowed the energetic disadvantage of photorespiration to be offset by higher values of C_c , the concentration of CO₂ at Rubisco. In a uniplastidic cell, the retention of the pyrenoid +CCM would be essential since Rubisco is so tightly packed in the pyrenoid that the low mesophyll conductance without a CCM would cause a major drawdown of CO₂ intracellularly (analogous to CAM plants during Phase IV of gas exchange: Maxwell *et al.*, 1997).

Comparative physiology of bryophyte photosynthesis

First, we consider the general characteristics of the CCM in hornworts, as compared to gas exchange and carbon isotope discrimination characteristics for other liverworts (Table 1.1). Rates of net CO₂ assimilation are similar for contrasting thalloid life forms when expressed on a chlorophyll basis (and also on a weight and area basis: Smith and Griffiths, 1996a,b, 2000). However, the higher carboxylation conductance of hornworts, as inferred from the $K_{0.5}$ and the lower CO₂ compensation point (Γ), is related to the magnitude of the dissolved inorganic carbon (DIC) pool accumulated (Table 1.1). Alternatively, for bryophytes with a varying degree of thallus ventilation, the gas exchange characteristics were uniformly more 'C₃-like', with higher $K_{0.5}$ values and higher Γ , suggesting that the efficiency of CO₂ acquisition is lower. The values for *Conocephalum conicum* (L.) Underw., with generally a low chlorophyll content, show a slight DIC pool, which is probably associated with alkalization of the chloroplast stroma during the light-dark

Table 1.1 Net CO₂ assimilation, gas exchange and isotope discrimination characteristics for hornworts and liverworts

| <i>Species</i> | <i>Net CO₂ assimilation, A</i> (nmol CO ₂ mg ⁻¹ chl s ⁻¹) | <i>Half saturation constant for CO₂, K_{0.5}</i> (μmol mol ⁻¹) | <i>CO₂ compensation point, Γ</i> (μmol mol ⁻¹) | <i>Dissolved inorganic carbon (DIC) pool</i> (nmol CO ₂ mg ⁻¹ chl) | <i>Instantaneous carbon isotope discrimination, Δ</i> (‰) |
|------------------------------|---|---|---|--|---|
| <i>Anthoceros crispulus</i> | 4.1 | 167 | 26 | 17.6 | 12.2 |
| <i>Phaeoceros laevis</i> | 5.1 | 110 | 25 | | 12.4 |
| <i>Marchantia</i> spp. | 4.9 | 185 | | 0 | |
| <i>Marchantia polymorpha</i> | 4.4 | | 54 | | 26.6 |
| <i>Pellia endivifolia</i> | 4.3 | 310 | 55 | 0 | 31.4 |
| <i>Pellia epiphylla</i> | 4.8 | 225 | 69 | 0 | 27.6 |
| <i>Conocephalum conicum</i> | 2.0 | 287 | 49 | 5.5 | 24.8 |

Data from Smith and Griffiths (1996b, 2000).

transient measurements (Smith and Griffiths, 1996a,b; Badger *et al.*, 1998). Finally, the on-line measurements of carbon isotope discrimination, which show the extent of discrimination directly during photosynthetic gas exchange (Griffiths *et al.*, 1999), are low in the two hornworts (*Phaeoceros laevis* L. and *Anthoceros crispulus* L.), close to the range normally associated with C₄-plant organic material (see Figure 1.1; Smith and Griffiths, 1996b), while the values for other liverworts are clearly within the C₃ range (Table 1.1).

Are there any advantages for maintaining a CCM in bryophytes? We have recently conducted a comparative study of bryophyte photosynthesis and light use, presented in Figure 1.2 (unpublished data). The CO₂ response of photosynthesis for the two non-ventilated thalli (*Phaeoceros* and *Pellia endivifolia* (Dicks) Dum.) are clearly distinguished: thus the diffusion limitation imposed in *Pellia* results in a linear A/Ca response, with low rates of maximum CO₂ uptake (Figure 1.2A). The effect of the CCM is to lower the CO₂ compensation point and achieve maximum rates of net CO₂ uptake similar to the well-ventilated *Marchantia*. In terms of carbon gain, therefore, the CCM helps to overcome diffusive limitations in hornworts, but rates of C gain are then only comparable to the bryophytes which have reduced the diffusive limitation within the thallus by increasing 'internal' air spaces. In today's climate, the relative costs of photorespiration equate to those of driving the biophysical CCM, similar to C₃ and C₄ plants at 18–20°C (Raven, 1985; Raven *et al.*, 1998); in the environment under which similar life forms might have evolved on land, the perhaps 10-fold higher atmospheric CO₂ concentration (Berner, 1998; Sage, 2002) would have largely suppressed photorespiration, and so the energetic 'cost' of the CCM, needed to overcome diffusive limitations both internal (single chloroplast) and external (non-ventilated thallus), did not provide a selective advantage.

However, a comparison of photosynthetic light use, measured as Photosystem II fluorescence, reveals that the CCM-based photosynthetic system does allow electron transport

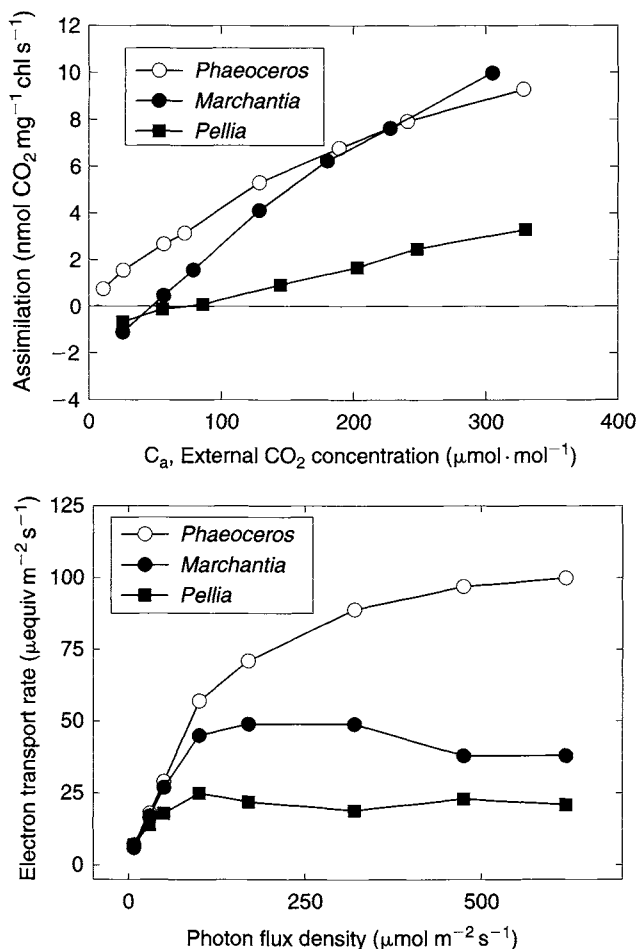


Figure 1.2 Photosynthetic CO₂ and light (electron transport) responses for hornworts and liverworts with a varying degree of thallus ventilation. CO₂ and PSII fluorescence were measured in a modified leaf cuvette (LD2, Hansatech, Kings Lynn, UK) coupled to an ADC 225 mk III infrared gas analyser (ADC, Hoddesdon, UK) in a closed system, and a PAM 101 Fluorometer (H. Walz, Effeltrich, Germany). Each response is the mean of measurements on two replicate thallus samples maintained under photon flux density of 50 μmol m⁻² s⁻¹, with CO₂ responses measured under 150 μmol m⁻² s⁻¹.

to be sustained under high photon fluxes up to full sunlight (Figure 1.2), as compared to both *Marchantia* and *Pellia*. Electron transport is saturated around a photon flux of 100 μmol m⁻² s⁻¹ for the latter, as opposed to around 600 μmol m⁻² s⁻¹ for *Phaeoceros* (Figure 1.2). Given that stomata do occur in the sporophyte generation of hornworts, the capacity to cope with high light seems to represent a lost opportunity for adopting more exposed habitats during colonization of the land, and supports observations that hornworts do thrive in relatively exposed conditions today. Perhaps the limitations imposed by the pyrenoid and the uniplastidic condition, with the dominance of chloroplast division over cell division processes (Brown and Lemon, 1990), inhibits the development of more complicated life forms.

Conclusions

A definitive phylogenetic tree would help to clarify the paraphyletic development of the bryophytes in the context of the protracheophytes and help to resolve how the dramatic change in reproductive life cycle was accomplished during the progression towards vascular plants. However, it would not necessarily resolve the occurrence of the pyrenoid in hornworts, since in the genus *Megaceros* there is the gradual loss of the pyrenoid associated with the development of the multiplastidic condition, which seemingly represents the derived condition (Burr, 1970; Brown and Lemon, 1990; Vaughn *et al.*, 1992; Badger *et al.*, 1998). However, we may make some physiological contribution towards the debate regarding phylogeny: if hornworts are to be considered basal (Beckert *et al.*, 1999; Wheeler, 2000; Nickrent *et al.*, 2000), and by inference closest to the *Coleochaete*-like ancestor, why was the genetic basis for expressing a pyrenoid lost? That the pyrenoid-containing members of the family are more primitive and uniplastidic (Vaughn *et al.*, 1992; Badger *et al.*, 1998), as well as possessing stomata in the sporophyte, seems to make this family the sister group for other bryophytes and hence all embryophytes (Beckert *et al.*, 1999; Nickrent *et al.*, 2000; Renzaglia *et al.*, 2000). Then again, the conclusions that liverworts are basal, using *rbcL* sequence of many bryophytes (Lewis *et al.*, 1997; Qiu *et al.*, 1998; Qiu and Palmer, 1999) also seems compelling, and so ultimately we must adopt a compromise position (Kenrick, 2000) to account for the lack of agreement between molecular phylogeny and the bryophyte fossil appearance in terms of poor bryophyte preservation and changing geological conditions, rather than rapid diversification. Importantly, we now need to conduct a search at the molecular level for genes encoding the pyrenoid in other bryophytes and early tracheophytes (*cf.* Pfannschmidt *et al.*, 1999).

As far as Rubisco functioning, a study has now been completed in the variation of catalytic properties in a range of hornworts (Hanson *et al.*, 2002). We urgently need a comparative study on the range of bryophytes to clarify the interrelationships between the CCM and the changes in Rubisco specificity which may have occurred in terrestrial vascular plants. Since lower S_{rel} values are associated with C_4 plants which have perhaps maintained Rubisco under elevated CO_2 for only 10 my, it would be intriguing to determine whether any significant variations have developed between *Anthoceros* and *Megaceros*, respectively with and without the pyrenoid, or across the mosses and liverworts. At any event, when bryophytes first colonized the land some 460 my ago, the advantage of elevated CO_2 at that time would have been offset by the higher ambient temperatures, likely to have caused v_c/v_o to have decreased from 5.7 to 2.2 for an increase from 15 to 35°C (at current CO_2 concentrations) and also reduced activation by any Rubisco activase (Crafts-Brandner and Salvucci, 2000; Rokka *et al.*, 2001).

The increasing ventilation of thalli, in a progression seen in extant liverworts as well as in the fossil record through increasing stomatal densities (Osbourne *et al.*, 2001) would undoubtedly increase internal conductance to CO_2 (g_i), but at the cost of higher water loss. When we performed measurements of photosynthesis and carbon isotope discrimination on a wetted thallus, it was noticeable that the on-line discrimination signal decreased by 4‰ consistent with higher diffusion limitation, although net CO_2 uptake rate was barely affected (Smith and Griffiths, 1996b). Our data suggest that the CCM can operate in a non-wetted thallus and help to overcome diffusion limitation and *Anthoceros* is often observed growing in quite exposed conditions in arable fields (MCF Proctor, personal communication) and so wetting does not seem to be a prerequisite for CCM activity. We need additional information on the occurrence, location and activity of

carbonic anhydrase throughout the hornworts (whether external, periplasmic, cytosolic, stromal or intrathylakoidal) and their interrelationship with CCM operation (Raven, 1997; Badger *et al.*, 1998; Thoms *et al.*, 2001).

Gas exchange characteristics of extant bryophytes provide some insight into the likely benefits of a CCM for hornworts as compared to the non-ventilated *Pellia* (see Figure 1.2). However, despite a higher carboxylation conductance (i.e. low $K_{0.5}$ for CO_2), and low compensation point, maximum rates of carbon gain are similar to the ventilated liverwort thalli (see Table 1.1, Figure 1.2). Indeed, we may infer that the expected energetic costs of the CCM are substantial (in contrast to the theoretical predictions of Raven, 1985), given the higher rates of electron transport needed to support this rate of CO_2 assimilation in *Anthoceros* (see Figure 1.2).

Ultimately, without a better understanding of the genes and regulatory processes leading to the expression (or suppression) of the pyrenoid or multiplastidic cells, we cannot make any more detailed inferences on the selective processes likely to have shaped the earliest terrestrial bryophytes. Perhaps we may uncover molecular evidence to show how and when the pyrenoid was lost from *Coleochaete* and hornworts and with it the potential to express a CCM. One thing seems certain – by becoming multiplastidic and internalizing airspaces, mesophyll conductance would have been dramatically increased. As long as oxygenase products could be detoxified via photorespiration, the higher light intensities available to drive electron transport (and not the CCM) could then have led to additional carbon reserves for creating structural material and conducting tissues for competing in the developing land-plant canopy. Despite the early opportunities and genetic basis for developing a CCM in early land plants, it appears that 400 to 450 my later, when the C_4 and CAM pathways became widespread, that potential had been irrevocably lost.

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2

Physiological evolution of lower embryophytes: adaptations to the terrestrial environment

John A Raven and Dianne Edwards

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Introduction

The physiology of embryophytes differs from that of their algal ancestors in a number of ways. Most relate to the differences in water relations of organisms which live on land, i.e. extant embryophytes, except the small number of species which have returned to live in water and those which live permanently in water, i.e. the great majority of species of algae.

This chapter sets out to examine the differences in physiology between embryophytes and their algal ancestors, with particular emphasis on their water relations. The embryophytes have very significant variations in water relations and the chapter considers their evolution within the embryophytes as well as the evolution of embryophyte water relations from those of their algal ancestors. The chapter also considers the relationship of the likely evolution of embryophyte water relations to cladistic analyses of embryophyte phylogeny and to the fossil record. A final point concerns the history of our understanding of this subject area and the possible constraints on achieving earlier syntheses.

The ancestors of embryophytes

The origins of the physiology of plants, i.e. embryophytes, must be sought in the extant algae which are most closely related to their algal ancestors. These algae are the Charophyceae *sensu lato*, and it is to the comparative electron microscopic studies of Pickett-Heaps (reviewed by Pickett-Heaps, 1975) and of Stewart, Mattox, Floyd and O'Kelly (reviewed by Stewart and Mattox, 1975) that the relationship of the Charophyceae to embryophytes was firmly established. Data on the occurrence of different enzymes catalysing the oxidation of glycolate, the hydrolysis of urea and the dismutation of superoxide radical anions supported the ultrastructural evidence (Bekheet and Syrett, 1977; Syrett and Al-Houty, 1984; De Jesus *et al.*, 1989). These findings have been supported by molecular phylogenetic studies (Nickrent *et al.*, 2000) and multifactorial cladistic analyses based on non-molecular studies (van den Hoek *et al.*, 1995; Kenrick and Crane, 1997; Graham and Wilcox, 2000a). The closest living relative of the embryophytes in the Charophyceae *sensu lato* has been widely held to be the small discoid alga *Coleochaete* Bréb. (van den Hoek *et al.*, 1995; Kenrick and Crane, 1997; Graham and Wilcox, 2000a; but see Karol *et al.*, 2001) which lives epilithically or epiphytically in fresh waters. The vegetative phase of *Coleochaete* is haploid and the only diploid phase in the life cycle is the zygote. Although earlier suggestions that there were mitotic divisions within the zygote before meiosis occurred have not been substantiated, one aspect of the life cycle of *Coleochaete* resembles that of the lower embryophytes. This is matrotrophy, i.e. nutrition of the zygote by the haploid vegetative phase using 'transfer cell'-like wall invaginations (Graham and Wilcox, 2000b). This similarity of *Coleochaete* to embryophytes is not shared by some charophyceans such as *Stichococcus* Nägeli and *Klebsormidium* Silva Mattox *et* Blackwell which are, however, sometimes found in terrestrial habitats (Graham, 1993). The analysis by Karol *et al.* (2001) of the phylogeny of the Charophyceae *sensu lato* in relation to the Embryophyta uses the sequences of one nuclear, one mitochondrial and two plastid genes in a range of analytical methods including Bayesian inference. Karol *et al.* (2001) conclude that the Charales are the sister clade to the embryophytes. Their findings suggest that the ancestors of embryophytes were vegetatively somewhat more complex than *Coleochaete*. However, Raven (1977) points out that extant members of the Charales, with most of the volume of the organism occupied by giant cells with volumes up to 10 mm³, are not good mechanical or physiological prototypes for early embryophytes (Raven, 1977, 1986).

The embryophytes are characterized by an alternation of gametophyte and sporophyte phases, with the gametophyte phase occupying a decreasing fraction of the biomass in the life cycle in proceeding from the bryophyte grade of evolution through the pteridophyte grade to the spermatophyte grade. In parallel to this the frequency of desiccation tolerant species decreases and the frequency of species with internal conduction pathways for water increases

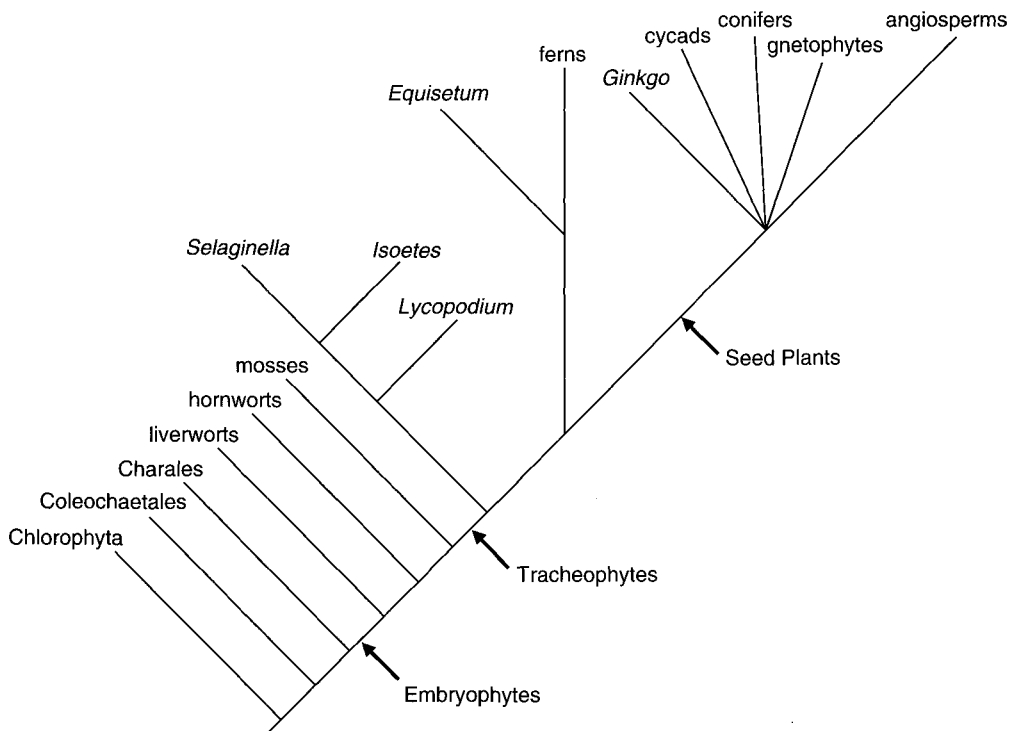


Figure 2.1 A phylogeny of the major groups of green plants based on several recent synthetic studies (summarized in Oliver *et al.* (2000), Karol *et al.* (2001)).

(Raven, 1977, 2002). Cladistic phylogeny (Kenrick and Crane, 1997; Qiu *et al.*, 1998; Qiu and Lee, 2000; Renzaglia *et al.*, 2000; Pryer *et al.*, 2001) and the fossil record (Chaloner, 1970; Gensel and Andrews, 1984; Edwards, 1993, 1996, 1998, 2000, 2003; Edwards *et al.*, 1996, 1998; Kenrick and Crane, 1997; Wellman and Gray, 2000; Raven and Edwards, 2001; Raven, 2002) are used as a framework for considering the evolution and the physiology of embryophytes (Figures 2.1 and 2.2).

Water, carbon dioxide and energetics of land plants

The physiology of the embryophytes is a very large subject and our considerations will be limited to resource acquisition and related mechanical matters. In particular, the acquisition of photons and inorganic carbon for photosynthesis from the atmospheric environment necessitates the loss of water vapour to the atmosphere. The use of light energy in photosynthesis necessarily involves dissipation of at least 73% of the absorbed photosynthetically active radiation other than in the reduced organic products of photosynthesis. The influx of CO₂ from the bulk atmosphere to Rubisco in chloroplasts requires a high-conductance diffusion pathway in the gas phase to a wet cell wall in which the CO₂ dissolves prior to diffusion in solution to Rubisco. The combination of an energy source for evaporation of water, the presence of a high conductance pathway for gas diffusion to the atmosphere and the general lack of saturation with water vapour of the atmosphere results in the loss of water from the photosynthetic structures to the atmosphere (Raven, 1977; Jones, 1992).

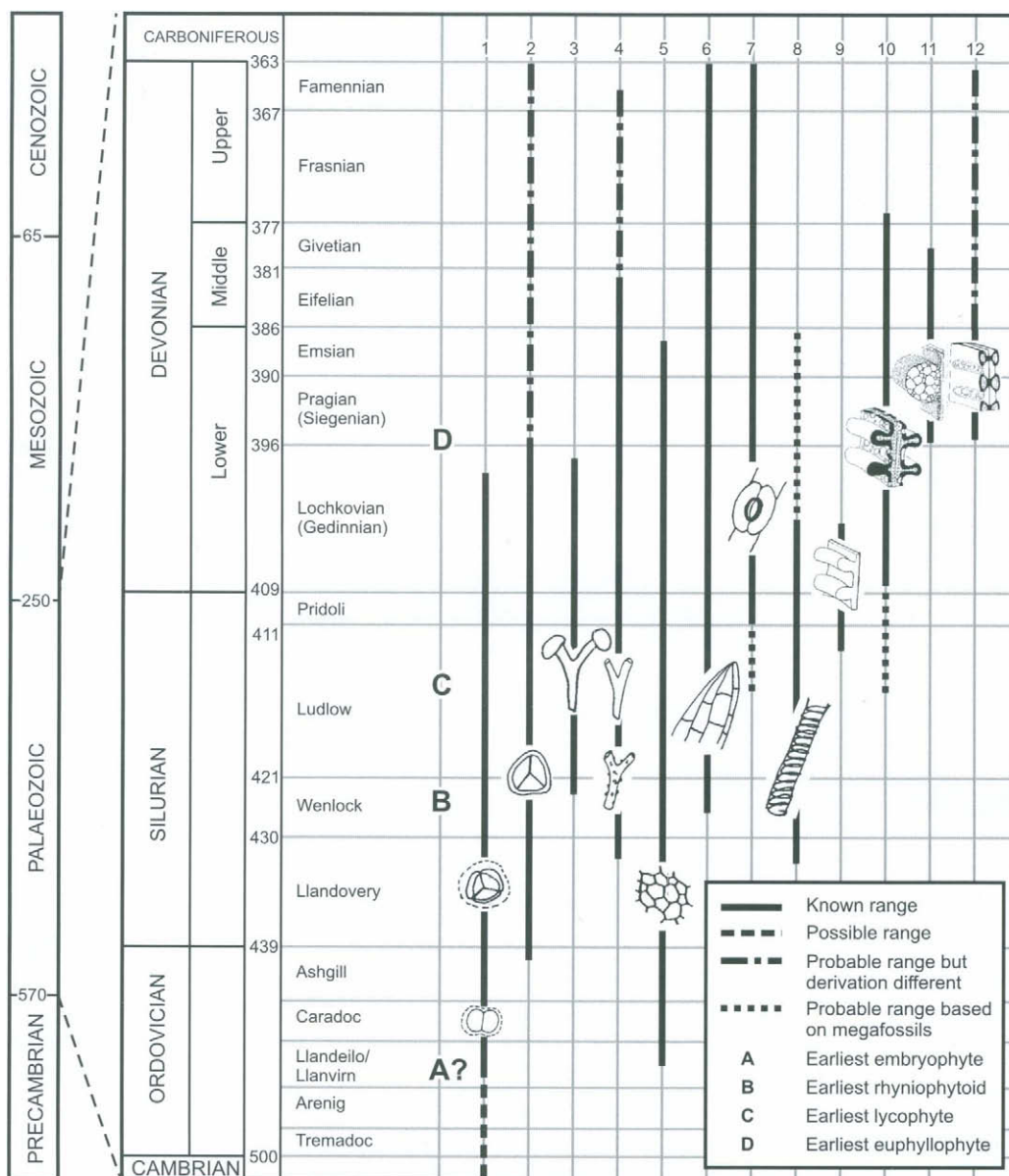


Figure 2.2 Stratigraphic ranges of fossils relevant to the early history of land plants. 1 = cryptospores (dyads and tetrads) thought to derive from bryophytes; 2 = trilete spores (monads) produced by tracheophytes; 3 = *Cooksonia* s.l.; 4 = axial fragments of ?tracheophytes; 5 = *Nematothallus* cuticles; 6 = sporangial and ?tracheophyte cuticles; 7 = stomata; 8 = unevenly thickened tubes of nematophyte affinity resembling tracheids; 9 = cooksonioid tracheids; 10 = G-type tracheid (zosterophylls and lycophytes); 11 = S-type tracheid (Rhyniopsida); 12 = P-type tracheid (typical early euphyllphyte tracheid).

An important aspect of the argument is the physics underlying the partitioning of energy dissipation from the photosynthetic apparatus. In addition to dissipation as the latent heat of evaporation of water, radiation, conduction and convection can also be involved in energy dissipation (Jones, 1992; Denny, 1993). Conduction between the photosynthetic

structures and the environment is much less in photosynthetic structures in air than in water as a result of the much lower thermal conductivity of air than of water as modulated by differences in boundary layer thickness and volume-based specific heat (Denny, 1993). This means that the temperature of the photosynthetic structures of the aquatic ancestors of the terrestrial embryophytes was much closer to that of surrounding water than the photosynthetic structures of the embryophytes are to the surrounding air. The temperature of the photosynthetic structures in air is also dependent on the extent of energy dissipation in its evaporating water and the temperature of the structures in air and in water is also determined by and determines, energy dissipation as long-wave radiation according to the Stefan-Boltzman law (Jones, 1992; Denny, 1993).

The outcome of these various considerations on the energy balance and gas phase conductances is that photosynthetic organs on land with C₃ physiology (diffusive supply of CO₂ to Rubisco) lose hundreds of grams of water vapour during the photosynthesis equivalent to the gain of one gram dry matter. The exact value of the water cost of growth depends on the radiation environment, wind speed, air temperature and relative humidity and the adaptive, acclimatory and regulatory characteristics of the photosynthetic organism (Jones, 1992). The CO₂ content of the atmosphere is another factor which alters the water cost of photosynthetic growth; more CO₂ in the atmosphere can permit lower water costs (Edwards *et al.*, 1998; Woodward, 1998; Konrad *et al.*, 2000). The CO₂ content of the atmosphere varies annually by about 5 μmol mol⁻¹ out of a current total of some 360 μmol mol⁻¹ as a result of the excess of terrestrial photosynthesis, mainly in the northern hemisphere, over terrestrial respiration in the summer and the excess of terrestrial respiration over photosynthesis in the winter (Keeling *et al.*, 1995; Berner and Berner, 1996). The present trend of year-on-year increases in atmospheric CO₂ as a result of changed land use, combustion of fossil fuels and cement manufacture could double atmospheric CO₂ over the next century (Berner and Berner, 1996). Clearly these anthropogenic effects are not relevant to the changes in atmospheric CO₂ prior to the evolution of man and rates of change of atmospheric CO₂ known from the Pleistocene ice record (Petit *et al.*, 1999) are lower than the current rate of increase. More relevant to the evolution of an embryophytic land flora is the suggested high CO₂ in the Ordovician, Silurian and Lower Devonian (Berner and Kothavala, 2001; *cf.* Boucot and Gray, 2001), as modelled for photosynthesis by Lower Devonian organisms at the pteridophyte grade of organization (Konrad *et al.*, 2000).

Another atmospheric factor which influences the water cost of growth is O₂ (which shows much less interannual variation in relative terms, than does CO₂: Keeling and Shertz, 1992; Keeling *et al.*, 1995; Berner and Berner, 1996). Higher O₂ mol fractions in the atmosphere, combined with a CO₂ mol fraction which is limiting for photosynthesis, limits gross photosynthesis via the oxygenase activity of Rubisco relative to an atmosphere with less O₂ (Raven *et al.*, 1994). The high CO₂ in the Ordovician, Silurian and Lower Devonian atmosphere (Berner and Kothavala, 2001), together with the evidence that the O₂ level was similar to the extant values (Berner, 2001), shows that O₂ was probably not a significant restriction on photosynthesis by the earliest embryophytes.

These considerations show that the water lost per unit dry matter increase is, other things being equal, lower in the atmosphere found in the Ordovician, Silurian and Lower Devonian than in the present atmosphere. A factor that would decrease the water cost of dry matter gain, especially at low CO₂ partial pressures, is the occurrence of CO₂ concentration mechanisms (Surif and Raven, 1990). Most green algae, including *Coleochaete* and many other members of the Charophyceae *sensu lato*, as well as some hornworts

(see Figure 2.3), have pyrenoids which are invariably correlated with the presence of a CO₂ concentrating mechanism (CCM) (van den Hoek *et al.*, 1995; Smith and Griffiths, 1996a,b, 2000; Badger *et al.*, 1998; Graham and Wilcox, 2000a). These mechanisms decrease the external inorganic C concentration required to half-saturate photosynthesis and usually lead to whole-cell photosynthesis which is O₂-insensitive and which has a greater CO₂ affinity than does Rubisco from the same organism *in vitro* (Badger *et al.*, 1998; Raven, 2000). In addition to their occurrence in many charophycean algae, pyrenoids and the associated CCM are also found in a number of species of hornwort (Smith and Griffiths, 1996a,b, 2000). These data could be construed as indicating that the common ancestor of *Coleochaete* and the embryophytes had pyrenoids. Regardless of whether the liverworts (see Figure 2.1; Qiu *et al.*, 1998; Qiu and Lee, 2000) or the hornworts (Nickrent *et al.*, 2000; Renzaglia *et al.*, 2000) are the taxon of embryophytes which is most closely related to the algal ancestors of embryophytes, losses of pyrenoids would be required to explain the pyrenoid-less state of almost all embryophytes on the basis of the occurrence of pyrenoids in the algal ancestor of embryophytes. However, it is likely that pyrenoids are polyphyletic (Raven, 1997) and evolved in response to low CO₂ concentrations and so are unlikely to have occurred before the low CO₂ in the Upper Devonian and Carboniferous. Accordingly, it is not likely that the earliest embryophytes had pyrenoids and so they would not have had higher CO₂ affinity and higher water-use efficiencies than plants with C₃ physiology. In any case, C₃ plants growing in the high CO₂ environment of the Ordovician, Silurian and Lower Devonian would, as indicated above, already have relatively low water costs of growth (Konrad *et al.*, 2000).

Desiccation tolerance, desiccation intolerance, poikilohydry and homoiohydry

Poikilohydry of algae and early-evolving embryophytes

The inevitability of water loss in terrestrial photolithotrophs during growth, combined with variability of water supply, leads to potentially lethal water loss in organisms which combine two characteristics: poikilohydry and desiccation intolerance. Poikilohydric plants have little or no capacity to restrict water loss when the rate of evaporative water loss exceeds the rate of liquid water supply. Desiccation-intolerant plants have a lethal lower limit of water content corresponding to a cell or tissue water potential which is still relatively high. Before attempting to quantify the 'little or no capacity to restrict water loss' and 'relatively high water potential', we point out that the 'transmigrant' charophyceans were poikilohydric.

Whether these 'transmigrant' charophyceans were desiccation-tolerant is not clear, but it is very likely that they were (Oliver *et al.*, 2000). Some extant charophyceans (e.g. members of the Zygnematales) have desiccation-tolerant zygotes. As for the embryophytes, Oliver *et al.* (2000) (see also Tuba *et al.*, 1999) have performed a parsimony analysis of the occurrence and evolution of vegetative desiccation tolerance. A very significant fraction of the taxa in extant liverworts, hornworts and mosses are desiccation-tolerant and Oliver *et al.* (2000) suggest that these organisms are ancestrally desiccation-tolerant. Oliver *et al.* (2000) suggest that vegetative desiccation tolerance was lost early in the evolution of the vascular plant sporophyte; the loss was permitted by the evolution of homoiohydricity and was evolutionarily favoured by the metabolic costs of desiccation tolerance which exceed those of homoiohydricity. However, while there are data sets with lower specific growth rates for desiccation-tolerant algae (e.g. *Porphyra* C.A. Agardh sp.) than for less desiccation-tolerant algae (e.g. *Enteromorpha* Link in Nees, nom. con sp. and *Ulva lactuca* L.) (Nielsen and

Sand-Jensen, 1990; Abe *et al.*, 2001), phylogenetic bias must be considered (Raven, 1999). More research is needed to determine if the metabolic costs of desiccation tolerance exceed those of homoiohydry (Raven, 1999). There is also the correlation of vegetative desiccation tolerance with plants of relatively small stature (Raven, 1999). Propagules (spores and ultimately seeds) are frequently desiccation tolerant even in vascular embryophytes which are vegetatively intolerant of desiccation (Raven, 1999; Oliver *et al.*, 2000). The occurrence of reproductive desiccation tolerance could form the basis for the polyphyletic (re)evolution of vegetative desiccation tolerance in the lycopsid *Selaginella* Pal. and in ferns and at least eight times among the angiosperms (Figures 1 and 2 of Oliver *et al.*, 2000; Pryer *et al.*, 2001; see also Gaff, 1981, 1997).

Tuba *et al.* (1999) reviewed the responses of desiccation-tolerant and desiccation-intolerant terrestrial photosynthetic organisms to CO₂ levels higher than the present level and concluded that neither theoretical nor observational results suggest that elevated CO₂ will lead to any substantial shift in the balance of advantage between the two groups of plants. The theoretical aspects of the study should apply to the early embryophytes in their high CO₂ environment. Tuba *et al.* (1999) could find data on CO₂ enrichment only for one vascular plant species (*Xerophyta scabrida*) so that all of the data for desiccation-tolerant plants comes from poikilohydric plants (bryophytes) and lichens. Tuba *et al.* (1999) noted that *Polytrichum formosum* Hedw. is the moss gametophyte which comes closest in its response to increased CO₂ to non-desiccation-tolerant homoiohydric plants. Some *Polytrichum* spp. do approach the homoiohydric condition (Raven, 2002). Desiccation tolerance in photosynthetic organisms has recently been related to the expression of certain proteins (late embryogenesis expressed proteins, dehydrins, rehydrins: Close, 1997; Campbell and Close, 1997; Li *et al.*, 1998; Farnsworth, 2000; Velten and Oliver, 2001; *cf.* Thomson *et al.*, 1998). Although Close (1997) refers to unpublished immunological evidence of dehydrin protein in green algae there seem to be no sequence data confirming the presence of dehydrins in green and, especially, charophycean algae. However, the presence of dehydrins in cyanobacteria, based on sequence evidence (Close, 1997), as well as the extensive sequence evidence for dehydrins in seed plants and a lycopod (Close, 1997) is consistent with the universal occurrence of dehydrins in photosynthetic organisms, although lateral gene transfer cannot be ruled out. However, it is likely that other genes are significant in permitting desiccation tolerance.

Thus far our considerations of poikilohydry and desiccation tolerance as putatively ancestral states in embryophytes, i.e. present in their algal ancestors, has not dealt with whether the poikilohydric and the desiccation-tolerant states are qualitatively or quantitatively different from their alternate states, i.e. homoiohydry and desiccation intolerance. Desiccation intolerance has already been considered; it is the (apparently) obligatory state for the vegetative phase of all homoiohydric plants more than 1 m in height, and for the great majority of homoiohydric plants of smaller stature (Oliver *et al.*, 2000; Raven, 2002). Homoiohydry is the capacity to maintain vegetative tissue hydration over a range of external conditions of water availability in the soil and evaporative demand by the atmosphere. The restriction on water loss rate when evaporative demand exceeds the soil supply of water requires a waxy cuticle over above-ground parts of the plant combined with stomatal closure. The occurrence of photosynthesis and transpiration when evaporative demand does not exceed the soil supply of water requires open stomata, an intercellular gas space system supplying atmospheric CO₂ to photosynthetic cells and an endohydric conducting system (true xylem in all extant homoiohydric plants: Walter and Stadelmann, 1968; Raven, 1977, 1984, 1993, 1996, 2002; Jarvis, 1998).

Dealing first with poikilohydry, is there a qualitative difference between poikilohydric and homoiohydric plants? Slowing down the rate of water loss from plants in a habitat with intermittent water supply can to some extent replace homoiohydric in maintaining cell and tissue hydration. This is because the likelihood of the organisms remaining above a given water potential by the time another rain event occurs is increased. The requirement here is a lower conductance for water vapour between the evaporating surface and the bulk atmosphere, so that a given concentration difference of water vapour between the evaporating surface and the bulk atmosphere yields a smaller flux of water vapour. The geometry which lowers the conductance for water vapour occurs at the cell and tissue level (Green and Lange, 1994) and at the canopy level (Rice *et al.*, 2001) in bryophytes. Of course, a lower gas phase conductance for water vapour involves a lower gas phase conductance for CO₂, so that there is a lower rate of CO₂ fixation in photosynthesis on a unit biomass basis. Because there is a biochemical conductance involved in photosynthesis but not in transpiration, a given decrement in gas-phase conductance gives a greater decrement in water loss than in photosynthesis. This effect is partially offset by the higher tissue temperatures which occur if there is the same biomass-based rate of absorption of photosynthetically active radiation but less dissipation of this radiation as the latent heat of evaporation of water. Nevertheless, slowing the rate of dehydration as a result of a decrease in the gas-phase conductance also yields a greater dry matter gain per unit water lost (Table 2.1).

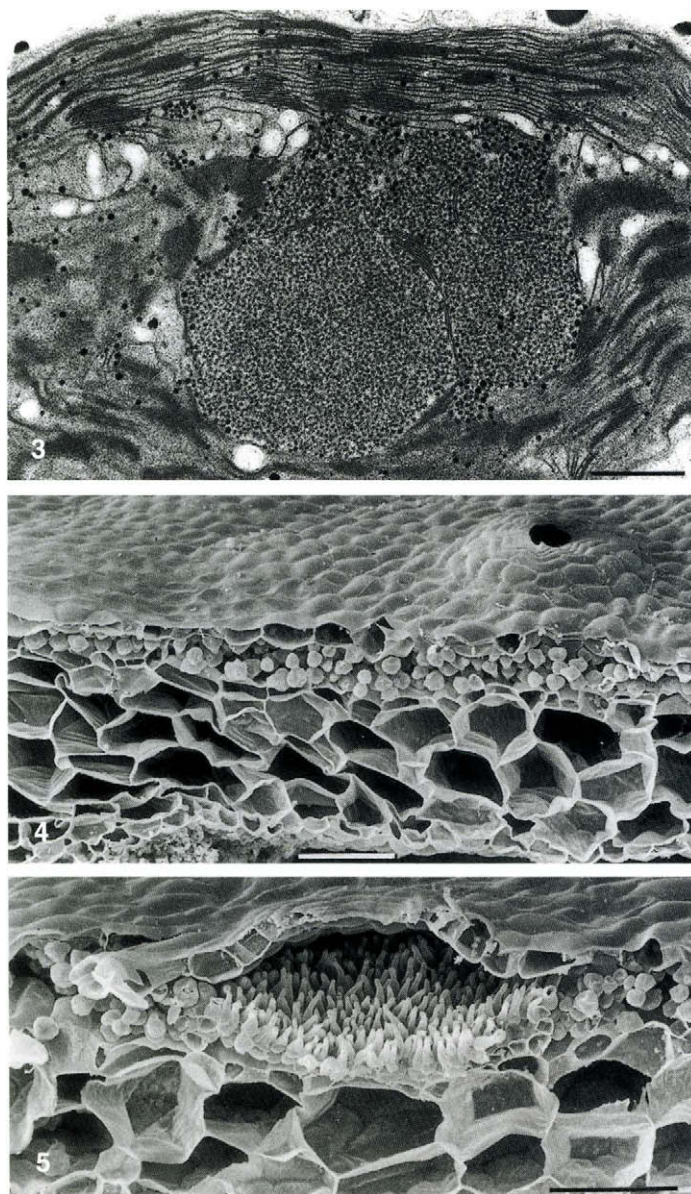
Such a low gas-phase conductance means that the achieved photosynthetic rate is lower than would be the case if there was a larger conductance. A low gas-phase conductance means that a given biochemical capacity for photosynthesis is not used to the extent that is possible with a higher gas-phase conductance. In the low gas-phase conductance case the rate of photosynthesis on a tissue nitrogen basis is lower than in the case of high gas-phase conductance. If there is a lower biochemical capacity for photosynthesis in parallel with a lower gas-phase conductance then the photosynthetic rate per unit nitrogen is the same in the high and the low gas-phase conductance cases, but the dry matter gain per unit water lost is also the same in the two cases (Table 2.1). A low gas-phase conductance reduces the use of the latent heat of evaporation of water in minimizing the increase in temperature of the tissue relative to the atmosphere (Table 2.1); changes in surface reflectance as a function of water content can also have an influence here (Hamerlynck *et al.*, 2000).

Conclusions from this analysis (see Table 2.1) show that there are cost-benefit 'trade-offs' for poikilohydric terrestrial plants in relation to variations in gas-phase and liquid-phase conductance in the absence of any specific regulatory changes in the conductances as a response to changed tissue water status, soil water availability, or the driving force for water loss to the atmosphere. Presumably different strategies (see Table 2.1) are more appropriate for particular habitats with (for example) different mean intervals between rainfall events, although it is not easy to find data which test this suggestion. Overall, poikilohydry, defined as the occurrence of little or no capacity to restrict water loss when the rate of evaporative water loss exceeds the rate of liquid water supply, is qualitatively different from homoiohydric.

However, there are examples of essentially poikilohydric plants which show some homoiohydric structural features. Examples are the endohydric, ventilated, cuticularized bryophyte gametophytes of many marchantiaceous liverworts and polytrichaceous mosses (Green and Lange, 1994; Raven, 2002). Despite the occurrence of intercellular gas spaces in the marchantiaceous liverworts, resembling those in homoiohydric sporophytes of vascular plants, there is no evidence of regulation of gas-phase conductance via changed pore

Table 2.1 The influence on poikilohydric plants of variation in gas-phase conductance and in biochemical conductance on maximum growth rate, dry weight gain per unit water lost, rate of dry weight gain per unit tissue nitrogen and the capacity to retain tissue hydration in environments of episodic rainfall. Based on Raven (1977, 1984, 1993, 2002) and Konrad *et al.* (2000), and discussion in the text of this chapter

| <i>Gas-phase conductance</i> | <i>Aqueous phase and biochemical conductance</i> | <i>Potential specific growth rate</i> | <i>Dry weight gain per unit water lost</i> | <i>Rate of dry weight gain per unit tissue nitrogen</i> | <i>Capacity to maintain hydration in environments with episodic rainfall</i> | <i>Excess of tissue temperature over air temperature in high light</i> |
|------------------------------|--|---------------------------------------|--|---|--|--|
| High | High | High | Moderate | Moderate | Low | Low |
| High | Low | Moderate | Low | High | Low | Low |
| Low | High | Moderate | High | Low | High | High |
| Low | Low | Low | Moderate | Moderate | High | High |



Figures 2.3–2.5 Figure 2.3 TEM (transmission electron micrograph) of part of cell of hornwort, *Dendroceros javanicus* Colenso, showing pyrenoid and part of chloroplast. Scale bar = 1 μm . Figures 2.4 and 2.5. SEMs (scanning electron micrographs) of fractured gametophyte of *Conocephalum conicum* (L.) Dum. after critical point drying. Scale bars = 100 μm . Figure 2.4 Note pore and photosynthetic tissue immediately below upper epidermis. Figure 2.5 Section through pore showing photosynthetic lamellae.

aperture (Figures 2.4 and 2.5) as a function of the driving force for water loss to the atmosphere or soil water availability or even tissue water status in the pre-emptive manner shown by stomata (Raven, 2002). The ventilatory system of the leaves of polytrichaceous mosses, i.e. photosynthetic lamellae on the upper leaf surface, is structurally less like the

Table 2.2 Variations in desiccation tolerance mechanisms in embryophytes (based on Oliver *et al.*, 2000; see van der Willigen *et al.*, 2001)

| Characteristic | Grade of organization of desiccation-tolerant embryophyte | | |
|---|--|----------------|----------------|
| | Bryophytes | Pteridophytes | Angiosperms |
| Tolerance of rapid (less than an hour) dehydration | Yes | No | No |
| Constitutive or induced tolerance of dehydration | Usually constitutive (<i>Funaria hygrometrica</i> is an exception) | Always induced | Always induced |
| Involvement of dehydrins or dehydrins-like proteins | Yes | Yes | Yes |
| Involvement of rehydrins | Yes | Yes | No |
| Involvement of ABA desiccation tolerances | Not in those with constitutive desiccation tolerance; involved in <i>Funaria</i> | Yes | Yes |

intercellular gas spaces of the marchantiaceous liverworts (Figures 2.4 and 2.5) and homoiohydric vascular plant sporophytes. However, it appears that the regulation of gas-phase conductance by movements of leaf components, and of leaves relative to stems, have more resemblance to the pre-emptive functioning of stomata than does behaviour of marchantiaceous pores (Raven, 2002; see also Tuba *et al.*, 1998). These regulatory functions of changes in gas-phase conductance in gametophytes of polytrichaceous mosses could be construed as showing some, as yet poorly quantified, approach to the homoiohydric state.

Desiccation tolerance and intolerance

As to desiccation tolerance, again there is evidence of a continuum of tolerance and intolerance of desiccation in vegetative tissues (Table 2.2). Oliver *et al.* (2000) review the evolution of desiccation tolerance in land plants and argue that the ancestral desiccation tolerance mechanism in embryophytes is the constitutive mechanism shown by algae and bryophytes and by lichens. These organisms are tolerant of rapid water loss (which happens as a consequence of their poikilohydric condition), although less rapid water loss (several hours rather than one hour) correlates with more rapid recovery upon re-addition of water in the much-investigated moss *Tortula ruralis* (Hedw.) Gaertn *et al.* (Oliver *et al.*, 2000). The organisms which can tolerate dehydration only if water loss is gradual (several hours to days) are typically the desiccation-tolerant homoiohydric sporophytes of vascular plants (Oliver *et al.*, 2000). The most detailed work has involved flowering plants, where it seems that there is an induction of a cellular protection mechanism during drying, which presumably accounts for intolerance of rapid drying (Oliver *et al.*, 2000). Relatively slow drying is permitted by the homoiohydric nature of these organisms. Oliver *et al.* (2000) conclude that the vegetative desiccation tolerance of vascular plants involved recruitment to the vegetative phase of the desiccation tolerance mechanism found in the

propagules, which in turn had evolved from the vegetative and reproductive desiccation-tolerance mechanism in algae and bryophytes. In the flowering plants the majority are homoiochlorophyllous (i.e. are able to tolerate vegetative desiccation in the green state). A few monocotyledonous flowering plants from the almost soil-less habitats of tropical inselbergs are tolerant of desiccation by a mechanism that involves loss of chloroplast fine structure with the need for three days of rehydration to re-establish full photosynthetic function (Oliver *et al.*, 2000). There is some evidence that vascular plants at the pteridophyte grade of organization have a desiccation-tolerance mechanism which is closer to the ancestral form than that of flowering plants. Thus, while pteridophytes, like angiosperms but unlike most bryophytes (an exception is *Funaria hygrometrica* Hedw.), show an induction of a cellular protection system upon dehydration, they resemble bryophytes in possessing rehydrins (Oliver *et al.*, 2000) (see Table 2.2).

Clearly the bryophyte mechanism(s) of vegetative desiccation tolerance are more relevant to the origin and early evolution of an embryophytic land flora. However, most of the quantitative data (tissue water potentials) or the extent of survivable dehydration in desiccation-tolerant and in desiccation-intolerant land plants comes from vascular plants. Raven (2002) suggests that desiccation-intolerant plants can tolerate water potentials not lower than -22 MPa, although the plants cannot grow at such low water potentials. By contrast, desiccation-tolerant plants can tolerate water losses equivalent to a cell water potential of -200 MPa or lower, and some of them (algae, bryophytes) can photosynthesize, and even grow, at water potentials as low as -30 MPa (Raven, 2002).

Evolution of homoiohydric

We have seen that the charophycean ancestors of the embryophytes, and the earliest embryophytes, were certainly poikilohydric and almost certainly desiccation tolerant. In considering how homoiohydric evolved, i.e. the order in which the homoiohydric characteristics occurred, what is physiologically plausible must be considered in relation to phylogeny of the embryophytes and to the fossil record.

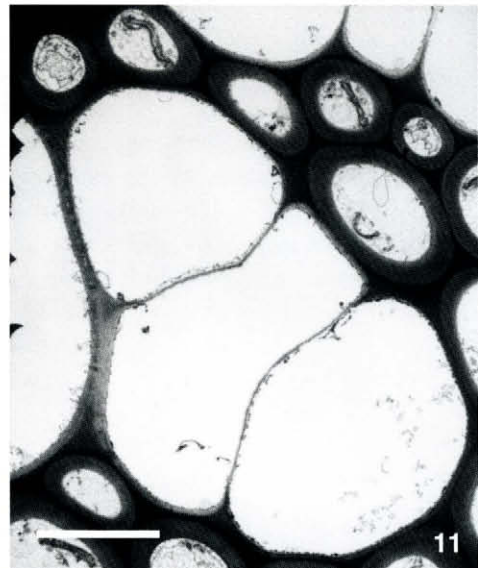
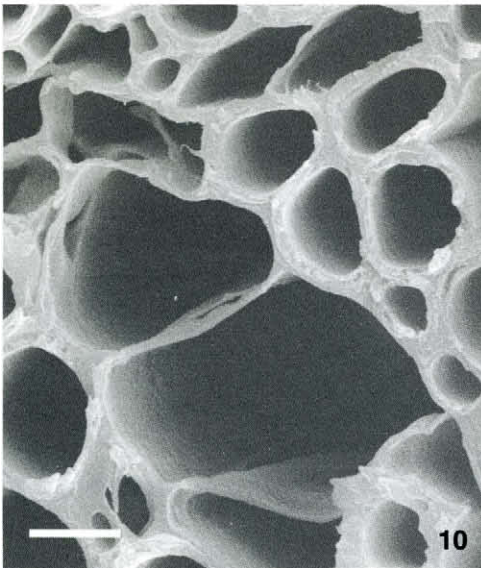
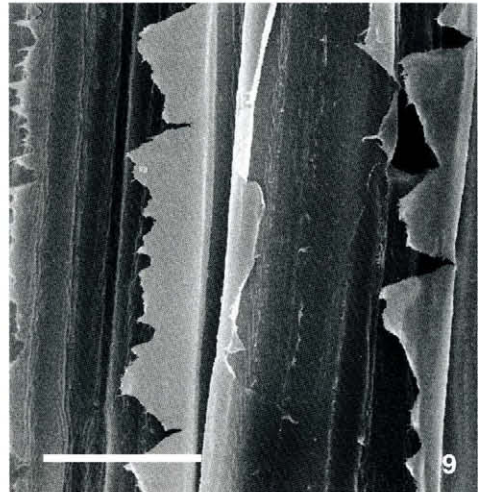
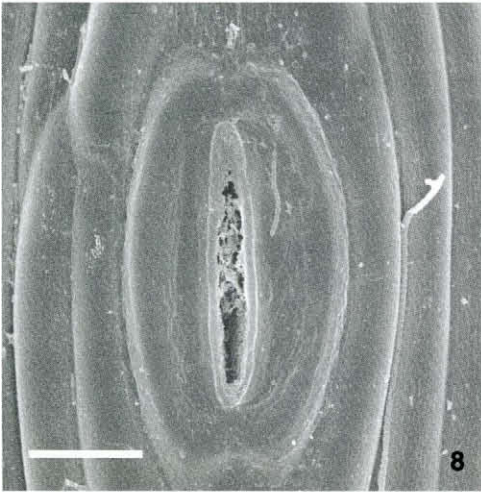
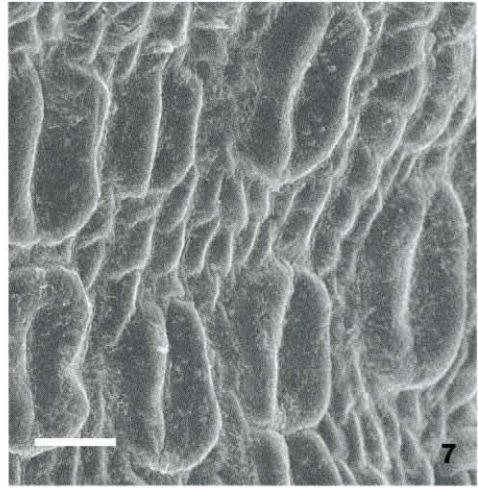
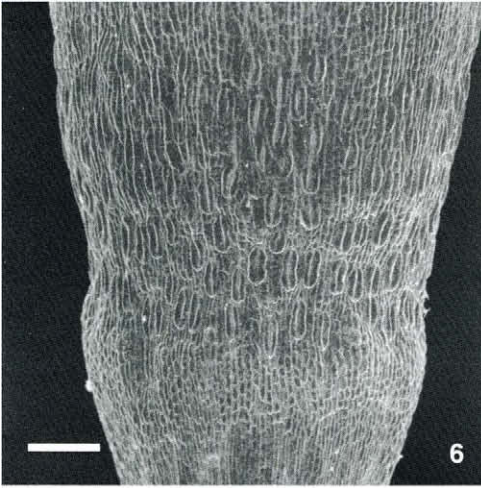
The consideration of the origins of differentiated endohydric conducting tissues and stomata comes to different conclusions as to polyphyletic origins or losses and depends on the assumptions made about the phylogeny of the lower embryophytes. One interpretation has the hornworts as the extant embryophytes which are most closely related to ancestral embryophytes (Kenrick and Crane, 1997; Nickrent *et al.*, 2000; Renzaglia *et al.*, 2000; Karol *et al.*, 2001; see Figure 2.1), while another interpretation has liverworts as the extant organisms closest to the ancestral embryophytes (Kenrick and Crane, 1997; Qiu *et al.*, 1998; Qiu and Lee, 2000). The liverworts have some extant representatives with endohydric conducting systems in their gametophytes (see Figures 2.4 and 2.5), and others with an intercellular gas space system and pores in their gametophytes, together with a cuticle in some gametophytes (Raven, 1996; Kenrick and Crane, 1997; Ligrone *et al.*, 2000; Raven, 2002). This assumption of liverworts as the ancestral embryophyte taxon (at least as far as extant taxa are concerned) permits monophyly of stomata (and intercellular gas spaces) in sporophytes without loss of these structures at the higher taxon (superclass or subdivision) level. By contrast, assuming that the hornworts are the ancestral embryophytes requires either that stomata are polyphyletic, or that stomata were lost in the whole super-class or subdivision of liverworts (Kenrick and Crane, 1997; Raven, 2002). Hornworts lack differentiated endohydric-conducting tissues in both the gametophyte or the sporophyte phases.

Regardless of whether the hornworts or the liverworts were closest extant relatives of the ancestral embryophytes, it is clear that the stomata (Figures 2.6–2.9), with intercellular gas spaces and cuticle, preceded the vascular plant endohydric conducting system and may have preceded the occurrence of a differentiated endohydric conducting system. This would presumably involve a sporophyte of thalloid morphology, similar to the structure and function of the thalloid gametophytes of extant marchantiaceous liverworts. Such organisms are endohydric but, with only a few hundred μm vertically of aqueous phase transport pathway from rhizoids to transpiring surface, the transpiratory flux can be supported by movement through undifferentiated cell walls of parenchyma cells (Raven, 1993, 2002). The occurrence of cuticle and intercellular gas spaces, with stomata, would permit increased photosynthesis on a thal-
lus projected area basis and maintain hydration even when evaporative demand with open stomata exceeds water supply from the soil (Raven, 1984, 1993, 1994a,b, 1998, 2002).

The alternative of endohydric water conduction (Ligrone *et al.*, 2000) with water-repellent cuticle but with no intercellular gas spaces or stomata has an extant analogue in the gametophytes of polytrichaceous mosses. Although there is no intercellular gas space system, these gametophytes are ‘ventilated’ via photosynthetic lamellae on the upper leaf surface and movements of lamellae and leaves can limit water loss at the expense of the photosynthetic rate (Green and Lange, 1994; Raven, 2002). However, these regulatory responses do not make the gametophytes homoiohydric. The sporophyte phase of these mosses has an endohydric conducting system as well as intercellular gas spaces, stomata (see Figures 2.8–2.11) and cuticle (Raven, 2002). While this apparently gives these sporophytes the attributes of homoiohydry, it must be remembered that these sporophytes are parasitic on the poikilohydric gametophyte, thereby constraining the extent to which the hydration state of the sporophyte can be maintained when soil water supply is low relative to the evaporative demand of the atmosphere (Raven, 1993, 2002).

While both of these hypotheses as to the evolution of homoiohydry seem evolutionarily plausible in terms of ecophysiology, the ‘stomata before differentiated endohydric conducting system’ seems more plausible in terms of phylogeny, at least if the endohydric system is equated with xylem *sensu stricto* (Raven, 1993, 2002).

Turning to the endohydric conducting system in the early embryophytes, including the conducting system in the gametophytes in some Lower Devonian polysporangiophytes, the plants were of relatively low stature with what are now known to be relatively low conductivity xylem ($\text{m}^3 \text{H}_2\text{O m}^{-2}$ transverse section area of conducting tissue $\text{s}^{-1} \text{Pa}^{-1} \text{m} = \text{m}^2 \text{H}_2\text{O Pa}^{-1} \text{s}^{-1}$) present in relatively small amounts in terms of the fraction of the axis cross-sectional area which is occupied by the conducting tissue (Raven, 1977, 1984). A low conductivity is a function of the small radius of the tracheids in the conducting tissue; there seems to be no evidence as to the length of the tracheids. The radius impacts on conductivity as defined above via the Hagen-Poiseuille equation (Jones, 1992; Nobel, 1999) in which the conductivity is directly proportional to the square of the conduit radius. The length of the tracheids relates to conductivity via the very low conductivity of the end walls relative to the lumen of the tracheids; the more frequent the cross walls, the lower the conductivity. There is a complex ‘trade-off’ between conductivity and prevention of cavitation and embolism; the former requires a large radius conduit and a large distance between cross-walls if a high conductivity is to occur, while a small radius and a small length of conduits means that a given cavitation or embolism event has less impact on overall conductivity. This latter correlation has a further relationship to overall conductivity: with more conducting units in a given cross-sectional area of xylem when the individual elements are small, the decreased effect of the failure of an individual element on the overall



conductivity of the xylem might be offset by a greater number of elements in a given cross-sectional area or volume. The impact of element size on the likelihood of a given decrement in conductivity in response to a certain set of environmental conditions essentially depends on whether cavitation or embolism is a function of the volume of an element, or of the number of elements, or the total volume of the segment of xylem.

The discussion of the factors involved in the evaluation of the endohydric conducting system, and specifically of xylem, has taken place in the context of the cohesion-tension hypothesis (Dixon, 1914). This hypothesis seems to account for most of the data which are currently available (Stuedle, 2001), although there is still dispute (Canny *et al.*, 2001; Cochard *et al.*, 2001; Richter, 2001).

As to the fossil record, the crucial evolutionary events apparently occurred in the Silurian (see Figure 2.2). Cuticles occur throughout the Silurian; although the earlier ones did not have any obvious relation to embryophytes (see Figures 2.12 and 2.13), the younger cuticles include those with clear relation to embryophytes (Edwards, 1993, 1996, 1998, 2000; Edwards *et al.*, 1996, 1998). Alas, the earliest embryophyte cuticles in the record yield no direct evidence on their chemistry or permeability (Edwards *et al.*, 1996). Fossil evidence for embryophyte cuticles only preceded the endohydric conducting system (Edwards and Wellman, 1996); later came stomata (see Figures 2.14 and 2.17) and xylem, with these two structures appearing in the record at a very similar time in the Upper Silurian (Edwards, 1993, 1996, 1998, 2000; Edwards *et al.*, 1996, 1998).

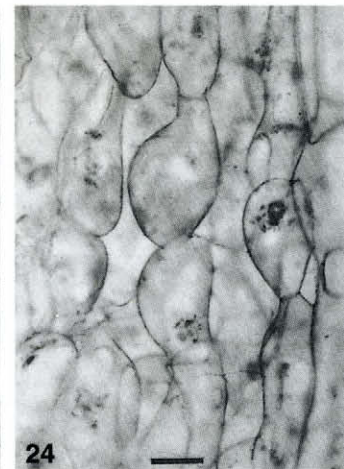
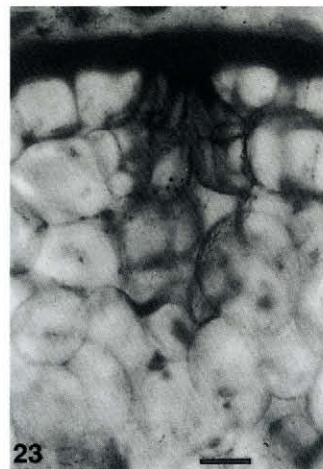
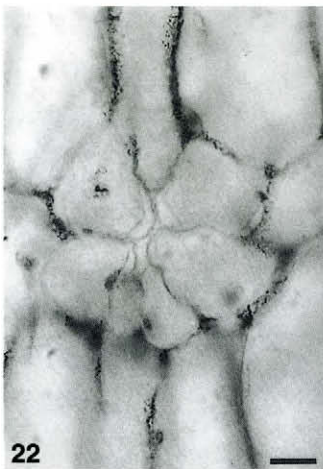
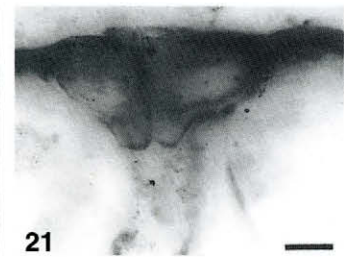
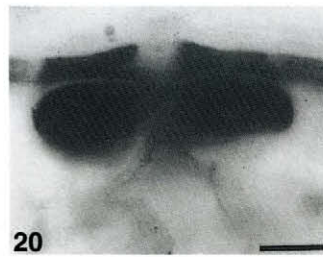
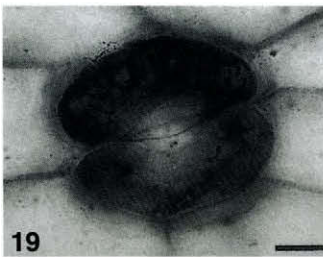
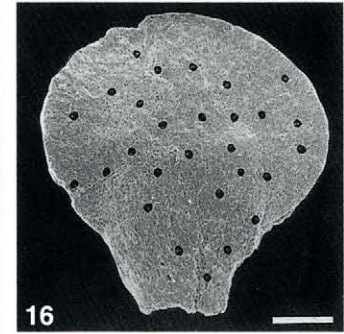
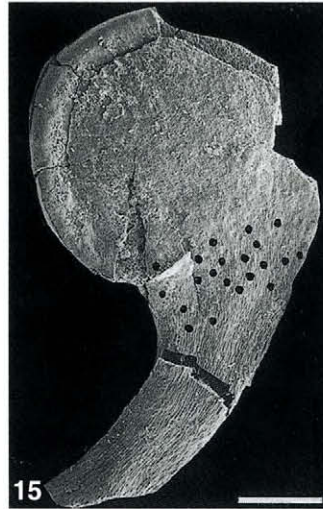
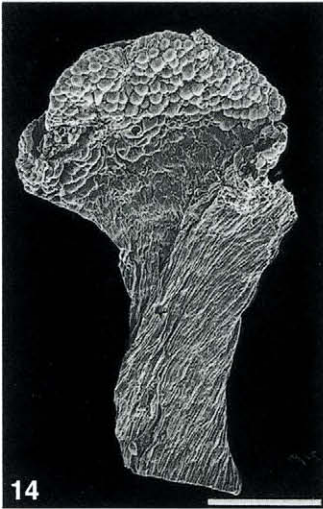
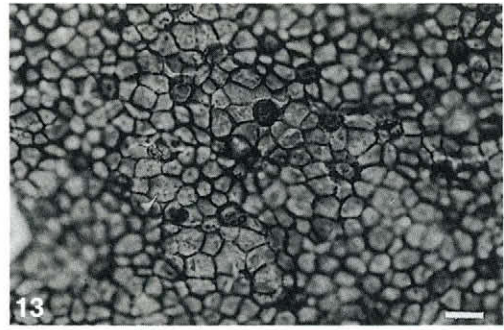
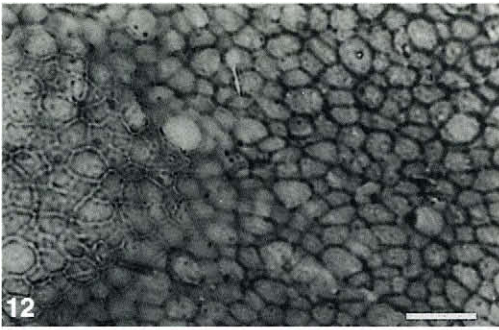
History of physiological interpretations of early embryophytes

Introduction

Here we consider the timing of some major discoveries in the palaeobotany of lower embryophytes and of major discoveries in plant physiology, and especially plant water relations. The three case histories considered in more detail are those of the role of transpiration and the endodermis in nutrient supply to the shoot of endohydric plants, the mechanism of water movement up the xylem during transpiration, and the role of stomata in determining the rate of photosynthesis and the water cost of photosynthesis in relation to the functioning of stomata.

Dawson, between 1850 and 1890, discovered and described Devonian vascular plants (e.g. *Psilophyton*) from the Maritime Provinces of Canada. The Rhynie Chert in Aberdeenshire, Scotland was discovered in 1912, and Kidston and Lang published on the Rhynie Chert embryophytes, with exquisitely preserved (permineralized) cellular structure, between 1917 and 1921. This work clearly shows xylem (or what in some cases are now interpreted to be analogous to endohydric water conducting systems: Edwards, 1993), stomata, intercellular gas spaces and cuticle (see Figures 2.19–2.33). Subsequent work extended the geographical and stratigraphical range of early embryophytes with good anatomical, cell-level preservation (Edwards, 1993). Scanning electron microscopy has

Figures 2.6–2.11 Figures 2.6 and 2.7 SEMs of stomata on seta of *Polytrichum commune* Hedw. Scale bars = 200 and 35 μm . Figure 2.8 SEM of surface view of stoma of hornwort, *Phaeoceros laevis* (L.) Prosk. Scale bar = 20 μm . Figure 2.9 SEM of longitudinal fracture through imperforate hydroids in central strand of *Dawsonia superba* Grev. Scale bar = 5 μm . Figure 2.10 SEM of transverse fracture through hydroids and surrounding stereids in *Dawsonia superba* Grev. Scale bar = 5 μm . Figure 2.11 TEM section through similar tissue. Scale bar = 5 μm .



been very important in investigating the anatomy of early embryophytes, and has been particularly significant in examining coalified fragmentary polysporangiophyte (Figures 2.14–2.16) and possible bryophyte remains (Edwards, 1993, 1996, 2000).

Discoveries in plant photosynthesis, water relations and mineral nutrition include the discovery of photosynthesis by van Helmont and measurements of the water use efficiency of growth (Woodward, 1699) in the late seventeenth century. In the eighteenth century plant plumbing was investigated and root pressure discovered by Hales (1727), and the role of O₂ in photosynthesis and respiration was discovered. The nineteenth century saw advances in understanding of nutrition (Liebig's Law of the Minimum; Liebig, 1840), and by early in the twentieth century the physical basis of gas diffusion through stomata and of the cohesion-tension hypothesis of xylem water movement in transpiration had been formalized (Brown and Escombe, 1900, 1905; Dixon, 1914) as had limitation of rate (Blackman, 1905), as well as extent (Liebig, 1840), of physiological processes by external factors. Later in the twentieth century great advances were made in quantifying energetic as well as kinetic aspects of transpiratory water loss (the Penman–Monteith equation: Jones, 1992) and formalizing concepts of homoiohydry and poikilohydry in relation to desiccation tolerance and desiccation intolerance (Walter and Stadelmann, 1968).

Transpiration rate and endodermal function in regulating nutrient supply to the shoot

Turning now to the interpretation of the physiology of the early land embryophytes in relation to their anatomy, Haberlandt set an excellent precedent in the late nineteenth century by attempting to interpret the functional anatomy of extant plants. However, later attempts to interpret the evolution of land plants by Church (1919) made little use of fossil data

Figures 2.12–2.24 Figure 2.12 LM (Light micrograph) of isolated cuticle of *Nematohallus*. Area out of focus results from the typical undulating nature of the cuticle. Lochkovian: north of Brown Clee Hill, Shropshire. DE1. Scale bar = 20 μm. Figure 2.13 LM of papillate cuticle of *Cosmochlaina maculata*. Lochkovian: M50 motorway near exit 3 (Newent, Gloucestershire). NMW85.20G.6. Scale bar = 20 μm. Figure 2.14 SEM of holotype of *Hollandophyton collicula* with terminal bivalved sporangium terminating stomatiferous axis. Pridoli: Ludford Corner, Shropshire. NMW96.11G.7. Scale bar = 500 μm. Figure 2.14 was published in *Special Papers in Palaeontology*, 67, 2002, 233–249 and is reproduced here with kind permission of The Palaeontological Association. Figure 2.15 SEM of bivalved sporangium, *Sporathylacium salopense*, with oblique band of stomata (dots). Lochkovian: north of Brown Clee Hill, Shropshire. NMW94.60G.11. Scale bar = 500 μm. Figure 2.16 SEM of unidentified sporangium with scattered stomata (dots). Lochkovian: north of Brown Clee Hill, Shropshire. NMW96.5G.7. Scale bar = 500 μm. Figures 2.15 and 2.16 were published in *Plant Cuticles*, 1996, 1–31 and are reproduced here with kind permission of BIOS Scientific Publishers Ltd, Oxford. Figure 2.17 SEM of stoma from *Hollandophyton* (Figure 2.14): the earliest stomata illustrated in a fertile taxon. Scale bar = 20 μm. Figure 2.18 SEM of stoma from Figure 2.15 Scale bar = 20 μm. Figures 2.19–2.24 LMs of silicified sections of axes of *Aglaophyton major*. Pragian: Rhynie Chert, Rhynie, Aberdeenshire. Scale bars = 20 μm (Figures 2.19–2.22) and 50 μm (Figures 2.23 and 2.24). Figure 2.19 Paradermal section through stoma. Note dark staining of guard cells compared with adjacent epidermal cells. P1603. Figure 2.20 TS guard cells with slightly separating cuticle and ledges. P1407. Figure 2.21 TS polar regions of stoma with extensions of the inner periclinal walls. P1611. Figure 2.22 TLS hypodermal cells below stoma: note thickenings surrounding channel to substomatal cavity. P1603. Figure 2.23 TS aerial axis just below a stoma to margin of the substomatal cavity. P1826. Figure 2.24 LS aerial axis showing parenchyma adjacent to substomatal cavity. Figures 2.19–2.24 were published in *Journal of Experimental Botany*, 49, special issue, 1998, 255–298 and are reproduced here with kind permission of Oxford University Press.

and did not take fully into account what was known about the physics of stomatal function from the work of Brown and Escombe (1900, 1905) (see Edwards *et al.*, 1996; Raven, 2002). Thus, Church (1919) held that stomata were not involved in photosynthetic CO₂ uptake, but were involved in the transpirational flux of nutrients from the soil solution to the shoot. This suggestion is not entirely without foundation in evidence from the fossil record as far as transpiration and nutrient uptake are concerned. One line of evidence comes from the distribution of stomata on the sporophytes of Upper Silurian and Lower Devonian embryophytes, where the stomata are often concentrated around sporangia (Figures 2.15, 2.16 and 2.18), a presumed site of a high requirement for soil-derived nutrients such as N, P, K, Ca, Mg, S and Fe (Edwards *et al.*, 1996, 1998; Raven, 2002). A second line of evidence comes from the fact that, to date, presence of an endodermis in below-ground, nutrient-absorbing structures in embryophytes before the Carboniferous has not been demonstrated (Raven, 1984; Raven and Edwards, 2001), although it must be acknowledged that we have next to no anatomical information on such structures through this time interval. The endodermis is widely held to be a means of limiting entry of soil solutes which are present in the soil solution in concentrations in excess of plant demand, granted the likely transpiratory water use in growth equivalent to 1 g gain in dry matter and the elemental content per g dry matter. The endodermis is also held to be involved in limiting leakage of nutrients from the stelar tissues in those cases where energized transport of nutrients from the soil solution to the stele compensates for lower nutrient concentrations in the soil solution than would satisfy plant elemental requirement (per g dry matter) and the transpiration occurring during production of 1 g dry matter (Raven, 1984; Raven and Edwards, 2001). However, among extant endohydric plants an anatomically evident endodermis is absent from the gametophytes of endohydric mosses and leafy liverworts, and the sporophytes of *Lycopodium* spp. among vascular plants. Accordingly, the absence of an anatomically evident endodermis may not prevent the accumulation in, or exclusion from, the xylem of components of the soil solution (Raven and Edwards, 2001). Church (1919) must not, of course, be castigated for not commenting on these functions of the endodermis (or its anatomically indistinguishable analogue) in relation to nutrient acquisition since, although what is now known as root pressure was investigated by Hales in the eighteenth century and the barrier role of the endodermis had been shown by Lavisson (1910), the function of the root system in the accumulation and exclusion of soil solutes was only fully characterized late in the twentieth century (Raven and Edwards, 2001).

The endodermis can also be related to Bower's suggestions (Bower, 1921, 1930) on the role of elaboration of the anatomy of primary stele tissue with increasing size of diameter of plant axial structure, both ontogenetically and phylogenetically as shown in the fossil record. Bower suggested that the increased complexity of the shape of the primary xylem in transverse section with increasing cross-sectional area of the axial plant structure was related to maintaining a near constant area of exchange of resources between xylem and the surrounding ground tissue. However, the nature of the exchanges was not specified and the restriction on radial solute fluxes may occur in the endodermis or a functional analogue thereof (Raven and Edwards, 2001). Nevertheless, Bower was prescient in attempting further functional interpretations of the anatomy and morphology of fossil plants.

Mechanism of endohydric water movement

In the context of the perception of the mechanism of endohydric water fluxes in interpreting the structure of the endohydric conduits in fossils of early embryophytes, the publication of

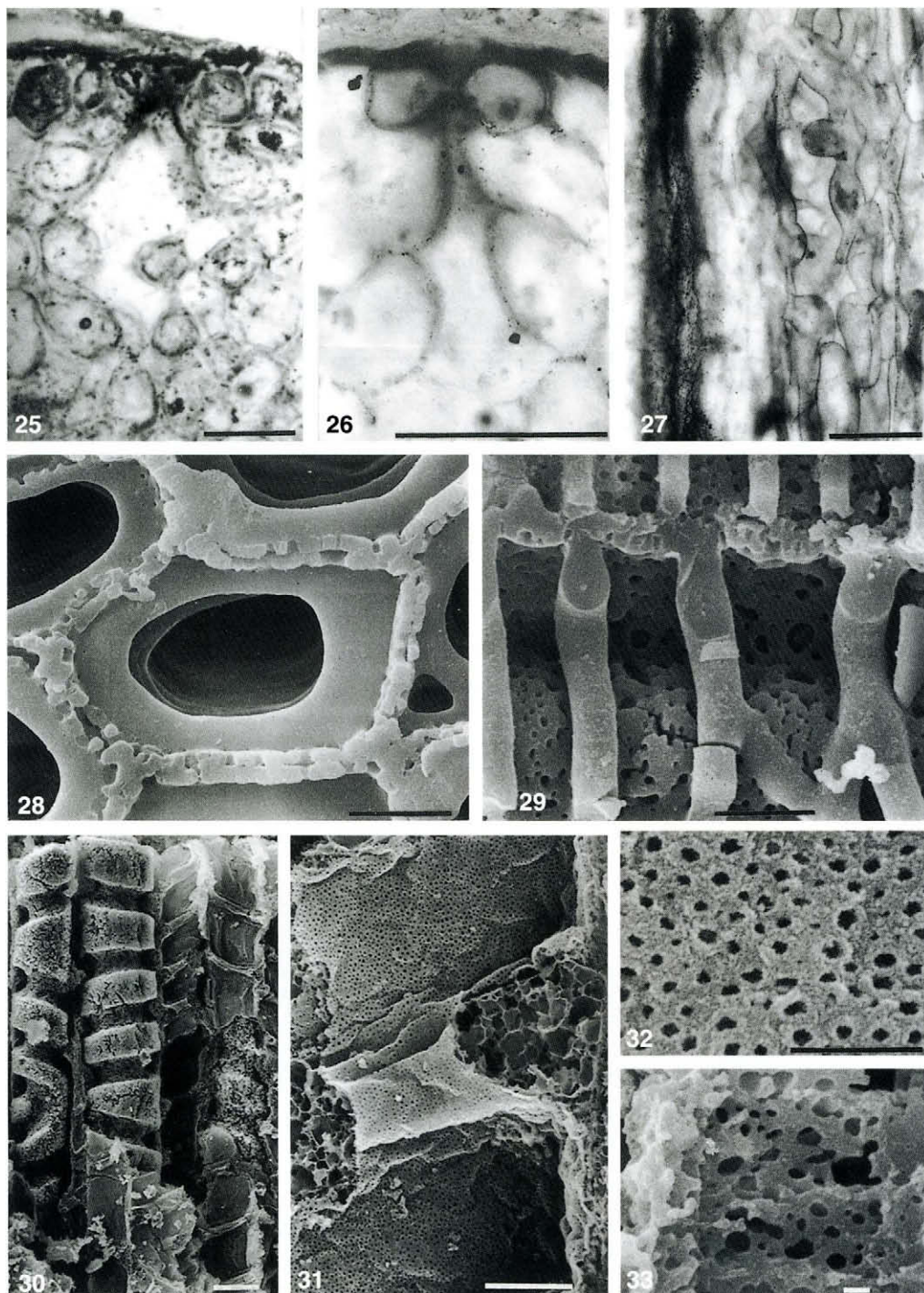
the book by Dixon (1914; see Dixon and Joly, 1894, 1895) on the cohesion-tension hypothesis just predated the publication of anatomical details of the Rhynie Chert plants between 1917 and 1921 by Kidston and Lang. Integration of the cohesion-tension hypothesis into the structural data from early embryophytes (Figures 2.27–2.32) took some time (Raven, 1977). Indeed, in their ultrastructural detail, the walls of tracheids have no counterparts in extant tracheophytes (e.g. S-types: Figures 2.30, 2.31 and 2.33) and some of the features of xylem, e.g. the presence of lignin, are not generally directly demonstrable in early fossil embryophytes (e.g. by pyrolysis/mass spectrometry of fossils with organic matter preserved: Ewbank *et al.*, 1997). Lignin is an important component of the cell wall of xylem conduits, providing rigidity and resisting the tendency to implode with cell contents under tension (Wainwright, 1970). Further work is needed on the functioning of the endohydric conducting system (hydrome) in, for example, extant polytrichaceous mosses to determine the tolerance of non-lignified but possibly polyphenolic endohydric conducting systems to increasingly negative pressures of the solution they contain.

Role of stomata in determining the rate of photosynthesis and the water cost of photosynthesis

Returning to the functioning of stomata, this time in the context of the possible photosynthetic rates and ratios of carbon assimilation to transpiratory water loss, post-Church (1919) analyses include the very important perception of Chaloner (see McElwain and Chaloner, 1995) about the significance of stomatal density measurements on early vascular plant fossils in the context of the potential rate of photosynthesis. Such concepts have now been integrated with the knowledge that the CO₂ mole fraction in the atmosphere has varied greatly over the Phanerozoic (McElwain and Chaloner, 1995); for a recent model for the CO₂ content of the atmosphere over the last 500 million years see Berner and Kothavala (2001).

However, these modelling studies relying on stomatal densities as proxies for atmospheric CO₂ (McElwain and Chaloner, 1995) and on the role of tracheophytes, via increased weathering, in lowering the CO₂ level in the Upper Palaeozoic (Berner and Kothavala, 2001) have been challenged by Boucot and Gray (2001). The main criticisms by Boucot and Gray (2001) include the mismatch between data from climatically sensitive sediments and the climate models and the lack of consideration of pretracheophytic land plants and of aquatic primary production in the consideration of the effect of biota on atmospheric CO₂ levels.

Despite the critique of Boucot and Gray (2001), we consider it likely that the CO₂ content of the atmosphere was 10–20 times higher than the present value at the time at which the earliest stomata are known from the fossil record and it is clear that the stomatal density of these Upper Silurian and Lower Devonian plants was much lower than that found today or in the Carboniferous when the CO₂ level was little greater than that found today (McElwain and Chaloner, 1995; Edwards *et al.*, 1998; Berner and Kothavala, 2001; Lake *et al.*, 2001; Raven, 2002). The possible photosynthetic rates and water costs of growth of early homoiohydric plants have been suggested by Raven (1977) who assumed that the atmospheric CO₂ content was the same as that found today. Konrad *et al.* (2000) give much more plausible estimates of gas exchange in Lower Devonian homoiohydric plants. The structural adaptations in the vicinity of stomata, including the deep seated substomatal cavity accessed by a narrow subporal canal illustrated here in certain Rhynie plants (see Figures 2.18–2.27), combined with low stomatal frequencies, infer high water-use efficiency (Edwards *et al.*, 1998). Further modelling of stomatal function has been carried out by Beerling *et al.* (2001a) who have related the evolution of the euphyllophyte leaf



Figures 2.25–2.33 Figures 2.25 and 2.26 Guard cells and substomatal cavities in TS axes of *Aglaophyton major* (P1980) and *Rhynia gwynne-vaughanii* (P2219). Pragian: Rhynie Chert, Rhynie, Aberdeenshire. Scale bars = 100 μm . Figure 2.27 LS axis of *R. gwynne-vaughanii* with epidermis to left and aerating tissue at margin of substomatal cavity on right. P2238. Pragian: Rhynie Chert, Rhynie, Aberdeenshire. Scale bar = 100 μm . Figures 2.25–2.27 were published in *Journal of*

(Kenrick and Crane, 1997; Pryer *et al.*, 2001) in the context of decreasing atmospheric CO₂ content of the atmosphere during the Devonian and thermal balance of the photosynthetic structures. These suggestions have not been universally accepted (Hedrich and Steinmeyer, 2001; Tanner, 2001; *cf.* Beerling *et al.*, 2001b).

Conclusions

The physiological changes which occurred in the evolution from algal ancestors to the different grades of organization of embryophytes has been determined from the physiology of extant plants in relation to their phylogeny as determined by cladistic analysis and from the order in which anatomical features appeared in the fossil record. The fossil record of embryophytes also tells us about organisms with characteristics which are not found today. Examples are gametophytes with the anatomical characteristics of homoiohydry and homoiohydric sporophytes of polysporangiophytes which lacked true xylem as their endohydric conduits in the Rhynie Chert (Lower Devonian). The fossil record is not helpful in telling us about such important properties as desiccation tolerance or intolerance, except by applying an empirical correlation from extant plants that no embryophyte more than 1 m in height is desiccation tolerant in the vegetative phase. Overall conclusions from consideration of the lines of evidence indicated above are that the earliest embryophytes were desiccation tolerant and poikilohydric. Achieving homoiohydry could have followed the cuticle plus gas spaces plus stomata before endohydric conducting system, or vice versa.

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Experimental Botany, 49, special issue, 1998, 255–298 and are reproduced here with kind permission of Oxford University Press. Figures 2.28 and 2.29 SEMs of coalified examples of G-type tracheids. Lochkovian: north of Brown Clee Hill, Shropshire. NMW99.20G.1. Scale bars = 5 µm. In these tracheids that characterize zosterophylls and basal lycophytes, annular or helical secondary thickenings are connected by perforated sheets of resilient material, believed to be deposited on the intervening compound middle lamella. Figure 2.28 Transverse fracture of xylem showing pitting in adjacent cell walls and a simple annular thickening. Note that the pits are only rarely coincident on adjacent walls. Figure 2.29 Longitudinal fracture of parts of tracheids. Note variation in size and shape of the pits. Figures 2.30–2.32 Xylem in *Sennicaulis hippocrepiformis*, preserved in pyrite, composed of S-type tracheids. These S-type tracheids occur in rhyniophytes such as *R. gwynne-vaughanii*. They are composed of a resilient spongy material and have broad predominantly helical thickenings. The lumen of the tracheid is lined by a layer (with perforations of nanometre dimensions) which also covers the secondary thickenings. Lower Old Red Sandstone: Mill Bay West, South Wales. Figures 2.30–2.32 were published in *Palaeontology*, 34, 1991, 751–766 and are reproduced here with kind permission of The Palaeontological Association. Figure 2.30 Longitudinally fractured xylem. Note residues of coalified material on left, pyrite casts of lumina on right. NMW90.42G.3. Scale bar = 20 µm. Figure 2.31 Fragment showing part of a thickening and its spongy matrix with overlying perforate layer. NMW90.42G.3. Scale bar = 5 µm. Figures 2.32 and 2.33 illustrate relative size of perforations in S and G-type tracheids. Figure 2.32 NMW90.42G.3. Scale bar = 1 µm. Figure 2.33 (from Figure 2.29) NMW99.20G.1. Scale bar = 1 µm. Figures 2.28, 2.29 and 2.33 were published in *Philosophical Transactions of the Royal Society of London, B*, 355, 2000, 733–755 and are reproduced here with kind permission of The Royal Society.

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3

Origin, function and development of the spore wall in early land plants

Charles H Wellman

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Introduction

The embryophytes (i.e. land plants) are one of the major kingdoms of eukaryote life. There is overwhelming evidence indicating that they are monophyletic, with extant charophycean green algae their sister group (e.g. Graham, 1993; Mishler *et al.*, 1994; Kenrick and Crane, 1997). All sexually reproducing land plants produce either spores, or their more derived homologues pollen, as part of their lifecycle. The resistant sporopollenin wall that encloses spores and pollen (spore exospore or pollen exine) is considered to be a synapomorphy of the embryophytes (e.g. Graham, 1993). It seems quite plausible that the invasion of the terrestrial subaerial habitat was not possible until a sporopollenin spore wall had evolved. Virtually all land plants possess spore/pollen walls composed of sporopollenin, except very rarely where this character has been secondarily lost (e.g. in certain aquatic angiosperms). This chapter is concerned with the role of the sporopollenin spore wall in the origin and early diversification of land plants (i.e. mid Ordovician-end Devonian). The aim is to explore and review our current understanding of: (1) the origin of the spore wall; (2) the function(s) of

the spore wall in early land plants (including evolution of the spore wall in response to changes in the habit and external environment of the plants); and (3) mode of spore wall formation in early land plants, including recent developments in the study of the genetic basis for spore/pollen wall formation. Referencing is not extensive, but is intended to provide easy access into the literature: it includes major texts and review papers, along with some more specific papers that are either directly relevant to discussions or are of particular interest.

Origin of the spore wall

The embryophytes can be considered as a grade of organization that evolved as an adaptive response to their invasion of the terrestrial environment, i.e. the origin of land plants is intimately related to the invasion of the land, with the unique features of the land plants (autapomorphies) evolving as an adaptive response that enabled transmigration from an aquatic to subaerial habitat. In this respect it is intriguing to consider the origin and function of the spore wall. Almost intuitively we tend to equate acquisition of a sporopollenin spore wall with the origin of land plants/invasion of the land. Hence sporopollenin spore/pollen walls are considered a synapomorphy of the embryophytes and are not present in their putative green algal precursors.

When plants made the transition from an aqueous to subaerial environment their reproductive propagules experienced a dramatic change in environment from aqueous (aqueous green algal precursors) to subaerial (earliest land plants). The aqueous environment may be considered as one of relative comfort and a medium in which propagule movement is relatively straightforward. The subaerial environment, on the other hand, is one that is extremely hostile and in which propagule transportation is problematic. Thus the process of propagule transport was translocated from: (1) the relative safety encountered in an aqueous medium, into the harsh subaerial environment, where propagules were exposed to increased solar UV-B radiation levels, desiccation and likelihood of mechanical damage; and (2) an aqueous environment in which movement is relatively easy, into a subaerial environment where transportation must essentially be subaerial (but with free water provided during some stage in the lifecycle). In this scenario the sporopollenin wall is viewed as an adaptation ideally suited to overcome these problems and protect the propagules in their harsh new environment. Of course, evolution of the spore is fundamentally related to evolution of the embryophyte life cycle (i.e. alternation of generations, e.g. Kenrick, 1994).

Sporopollenin production, however, is almost certainly preadaptive. It occurs in a variety of diverse algal groups (usually enclosing resting cysts or reproductive structures), including the charophyceans (the sister group to embryophytes) where it is located in an inner layer of the wall enclosing the zygote (Graham, 1993). In the vast majority, if not all, of these cases, the function of the sporopollenin coating is usually considered to be protective, i.e. to protect the enclosed resting cyst or reproductive structure from mechanical damage and/or microbial and fungal attack, and if exposed to the subaerial environment, desiccation and/or solar UV-B radiation (see p. 46). When considering mutations that enabled green algal ancestors to invade the subaerial environment and survive as the earliest land plants, Knoll and Bambach (2000, p. 8) note that 'principal among these mutations must have been one involving a simple change in the timing of gene expression for sporopollenin biosynthesis from just after zygote formation to after meiosis and spore formation... such a mutation, arguably lethal in water, would have provided protection against desiccation and harmful radiation at a critically vulnerable phase of the lifecycle'.

However, the extent to which the sporopollenin of spore/pollen walls of embryophytes and the resting cysts/reproductive structures of the diverse algal groups are similar is unclear. Studies of sporopollenin chemistry are difficult to assess. Sporopollenin is one of the toughest and most chemically resistant biological polymers known (hence its utility as a protective coating). These properties, however, render it difficult to analyse and the chemical composition and structure of sporopollenin are poorly understood as are the biochemical pathways leading to its synthesis. Recent reports suggest that the sporopollenin of different algal groups varies in chemical composition and structure (see, for example, Graham and Gray, 2001). In fact the sporopollenin produced by different embryophyte groups also appears to vary in chemical composition and structure (Hemsley *et al.*, 1995). However, it is unwise to use such evidence to prove/disprove homologies because: (1) analysis of such resistant material is fraught with difficulties; and (2) chemical composition/structure may simply have evolved in response to relatively minor biochemical changes, possibly related to changing utility and/or environment. It is generally considered that spore/pollen walls are homologous to the zygote walls of charophycean algae – we simply have a heterochronic switch from enclosing the zygote to the products of meiosis (e.g. Blackmore and Barnes, 1987; Graham, 1993; Hemsley, 1994). However, this needs to be tested at the biochemical and molecular level. Similarities in the mode of sporopollenin deposition in embryophytes and certain green algae (see p. 47) support interpretation of these structures as homologous. It is anticipated that ongoing research in the field of molecular genetics may also aid identification of homology by identifying homologous genes involved in sporopollenin synthesis and/or regulation of its deposition.

Function of the spore wall

In the diverse algal groups possessing sporopollenin walls that enclose resting cysts/reproductive structures, the primary function of the sporopollenin is most likely one of protection of the contents. In aquatic algae this involves protection from mechanical damage and fungal/microbial attack. There are almost certainly multiple secondary functions. These include aids to buoyancy (e.g. extensions to the walls of resting cysts) etc.

In the charophycean green algae, the sporopollenin enclosed zygote is generally confined to an aquatic habitat, but may be exposed to the subaerial environment, e.g. when a pond dries out. Graham and Gray (2001) recently reviewed the function of the sporopollenin zygote wall of charophycean green algae. They suggest that, in charophyceans, the primary function of the sporopollenin wall is to protect the zygote from mechanical damage and fungal/microbial attack during seasonal dormancy (and possibly also transport). Interestingly, they suggest that desiccation resistance is almost certainly a secondary function (if one at all). Graham and Gray (2001) note that ‘there is no definitive evidence that sporopollenin (or other wall modifications) of algal resting cells prevents cellular water loss over extended periods of time, although it is possible that it functions as a short-term barrier to diffusion, as compact sporopollenin is regarded as essentially impermeable to water. Rather, sporopollenin in cell walls of charophyceans (and other aquatic algae) may provide protection from varied environmental extremes, where resistance to degradation of cytoplasmic contents is essential, such as during obligate periods of zygote dormancy’. They conclude that ‘desiccation protection in aerial distribution, if any, is probably a secondary function of sporopollenin in charophycean algae’. This interpretation is based on the fact that of the algal groups that produce propagules with a sporopollenin component, few are

demonstrably dispersed subaerially. In the majority of those that are subaerially dispersed, birds and insects are the dispersal vector and the sporopollenin is likely to protect against enzymatic digestion in the digestive tracts. In any case, the presence of a sporopollenin wall might even impede rehydration during germination.

The arguments developed are interesting regarding the function of the sporopollenin spore/pollen wall of embryophytes, particularly in that these are usually regarded as homologous to the sporopollenin walls present in various green algal groups. One of the major problems faced by land plants during the transition from an aquatic to subaerial environment was that of protection and dispersal of propagules outside of the aqueous environment (see p. 44). To what extent are the functions of the sporopollenin spore/pollen wall similar to those in green algal precursors and to what extent are they exaptations? Graham and Gray (2001) suggest that the primary function of the sporopollenin spore/pollen wall in embryophytes is similar to that in extant charophycean green algae, i.e. to provide mechanical protection to the spore protoplasm during seasonal dormancy prior to germination and to resist fungal and microbial attack. They consider that protection from desiccation is likely to be a secondary function (possibly as is protection from solar UV-B radiation). However, it is clear that sporopollenin does provide at least short-term protection of the spore protoplasm from desiccation. It is also clear that sporopollenin plays a critical role in protecting spore contents from solar UV-B radiation (summarized in Rozema *et al.*, 2001a,b).

As the embryophytes have evolved and diversified the spore/pollen wall morphology evolved as it acquired additional functions (i.e. exaptations) (e.g. Chaloner, 1976; Traverse, 1988). These include: (1) dispersal (by wind, water or animal vectors, including protection when passing through their guts); (2) protection from herbivores and detritivores; (3) lodging; and (4) germination control. A brief examination of spore/pollen walls in fossil and extant plants reveals prodigious morphological diversity (structure and ornament). This almost certainly reflects the multiple functions of the wall (both primary functions and exaptations) and these of course are intimately related to the evolving mode of life of plants and changes in the external environment. Ultimately, however, such diverse morphological adaptations function to facilitate sexual reproduction.

Spore wall development

So how do embryophytes actually build the sporopollenin spore/pollen wall? Clearly a very flexible system is involved that allows diverse morphology to evolve in response to changing lifestyles and environments. In the following section current knowledge of spore wall formation is summarized, concentrating on the information relevant to early land plant spores.

Basic mechanisms of spore wall formation

Ultrastructural studies carried out across the plant kingdom (extant and fossil) have shed light on spore wall ontogeny and the structures/processes operative during development. Blackmore and Barnes (1987) recognize four basic modes of sporopollenin deposition in spore walls. These are:

1. Accumulation on white-line-centred-lamellae (WLCL)
2. Deposition from surrounding cells of the sporangium onto previously existing layers
3. Accumulation within primexine (a pre-patterned cell surface glycocalyx)
4. Centripetal accumulation onto pre-existing layers.

Blackmore *et al.* (2000) subsequently considered these four principal modes of spore wall formation in greater detail and subdivided them. This was largely in order that they could be utilized as characters in a cladistic analysis of extant free-sporing vascular plants, but also to reflect an increasing understanding of these modes of spore wall formation.

Regarding mode of formation (1), WLCL have long been prominent in studies of spore wall formation. They appear to be involved in sporopollenin deposition in various green algae and the vast majority of, if not all, land plants. They have also been identified in fossil spores, including those over 400 million years old (Wellman *et al.*, 1998a). Deposition on a system of WLCL is considered to be the most primitive mode of sporopollenin wall formation. It occurs in many algae and is plesiomorphic in all embryophytes. WLCL appear to form at the plasma membrane with sporopollenin polymerizing out on either side of the white line. In *Chlorella* Beijerinck they link up to form a single continuous layer (Atkinson *et al.*, 1972). In bryophytes we see the acquisition of numerous superimposed lamellae (this appears to be an early transformation in the evolution of land plants). They can also occur in other forms (as extensions, overlapping etc.) and although walls thus formed are usually rather simple, complex ornamentation can be built up in this fashion. In their recent cladistic analysis, character 8 of Blackmore *et al.* (2000) concerned exine deposition involving WLCL formed on the plasma membrane (notably initial exine deposition). In mature spore/pollen walls parts formed by the accumulation of WLCL may appear lamellate or homogeneous. The latter occurs when compression and/or sporopollenin deposition conceals early sub-structures. Hence it is desirable to study ontogenetic sequences, not just the mature wall.

Mode of formation (2) (deposition from surrounding cells of the sporangium onto previously existing layers) was subdivided by Blackmore *et al.* (2000). Their character 10 concerned tapetal contribution to the exine. However, they distinguished exine formation involving a tapetum-derived component incorporated onto surfaces formed by the microspore (and usually based on WLCL) from epispore, perispore and paraexospore formation. Epispore (their character 13) is a layer within the sporoderm of heterosporous pteridophytes that is formed during the latest stages of exospore development, is composed of sporopollenin and is largely, if not exclusively, of tapetal origin. Perispore (their character 14) consists of material derived from the degenerating tapetum that condenses in one or more continuous layers over the surface of the spores. Taylor (2000) notes that sporopollenin derived from a tapetum may produce mature walls that are homogeneous and/or structures known as globules, orbicules or Ubisch bodies. Blackmore *et al.* (2000) also distinguish paraexospores (*sensu* Lugardon, 1976) (their character 11). A paraexospore is an outer layer of the exospore that is separated from the inner component by a large discontinuity (gap). Both layers are structurally similar and formed by WLCL. Clearly mode of formation (2), in its various guises, has major input from the diploid sporophyte (i.e. the tapetum). Blackmore and Barnes (1987) suggested that a tapetal contribution to the sporoderm was absent in basal land plants but acquired within both mosses and vascular plants. They considered this mode of deposition largely in terms of perispore formation (present in mosses and all vascular plants except *Lycopodium* L.). However, Blackmore *et al.* (2000) noted that a tapetal contribution to the sporoderm takes many forms (including addition to layers previously formed from WLCL or the formation of an epispore).

Mode of formation (3), accumulation within primexine (a pre-patterned cell surface glycolyx), will not be considered further. This mode is confined to the pollen of seed plants and is thus of little relevance to our consideration of spore wall formation in early land plants.

Mode of formation (4) involves centripetal accumulation onto pre-existing layers. Blackmore *et al.* (2000) covered this in their character 12. They note that spore wall

formation may involve accumulation below an earlier formed layer, either by accumulation by WLCL or deposition of granular or amorphous sporopollenin.

Blackmore *et al.* (2000) note that during sporogenesis a wide variety of mature forms is generated by a limited number of ontogenetic processes. They also note that the phylogenetic pattern of sporoderm synthesis is not simply a sequence of gains of additional wall components as there also appear to have been losses of developmental processes. However, later stages of development of spores are not always contingent upon the successful completion of earlier ones, so the deletion of an entire phase may not always prove fatal (Blackmore and Crane, 1988). Blackmore and Crane (1988) noted that sporoderm ontogeny is complex, but includes non-terminal additions or deletions of developmental steps without concomitant loss of reproductive viability. Blackmore *et al.* (2000) consider 'in our view this reflects the particular aspect of spore ontogeny that is unique among all plant cell walls, namely development through an interplay of haploid, gametophytic and diploid, sporophytic processes'.

Substructural organization of spore walls

A number of models for the substructure of spore/pollen walls have been proposed. Two recent models are briefly discussed below.

Rowley (e.g. 1995, 1996) suggests that all embryophyte spore/pollen walls are composed of similar substructures. He interprets these substructures as like wire-wound springs (termed tufts) that are always of similar size and configuration. Rowley suggests that the tufts form an early substructure and act as receptors for sporopollenin (i.e. sporopollenin polymerizes out on these tufts). Furthermore, he suggests that they can open and close and thus act as migration routes for various materials, including those utilized during spore wall construction. Rowley also suggests that the tufts combine actually to form WLCL. He suggests that the tufts orientate perpendicular to the white line and hence the sporopollenin polymerizes out on either side of the white line. He considers that the WLCL can also open and close and thus act as a migration route for various materials. Because they form early, and are subsequently coated in sporopollenin, tufts are difficult to study (they can only be viewed in early maturation or degraded walls). Rowley (1995) provides reconstructions of these substructures alongside photographs of immature and degraded spore/pollen walls in which they can purportedly be seen.

Another exciting development is the discovery that self-assembly is operative during spore wall formation in the rather thick walls of certain lycopsid megaspores (fossil and extant) (e.g. Hemsley *et al.*, 1994, 2000; Gabarayeva, 2000). It will be interesting to see how widespread this is in the plant kingdom, or if it is confined to the abnormally thick walls of megaspores. Interestingly, Scott (1994) has suggested that self-assembly of molecules (lipids and so on) is responsible for the formation of WLCL. I see absolutely no reason why self-assembly should not be involved in actual wall construction. However, I am convinced that this is ultimately under genetic control, in that the amount, composition and delivery of materials is genetically controlled, as is ornament morphology through various template systems.

Spore wall formation in extant plants

Extant plants most relevant to the study of early land plants essentially belong to the more 'primitive' free-sporing plants (i.e. non-seed plants): 'bryophytes' (liverworts, hornworts, mosses) and 'pteridophytes' (lycopsids, sphenopsids, ferns). Spore wall formation has been

studied in representatives of all of these groups and includes studies of the isospores of homosporous forms and the microspores and megaspores of heterosporous forms. For reviews of spore wall formation in these plant groups the reader is referred to the following: bryophytes (Brown and Lemmon, 1988, 1990, 1991); pteridophytes (Lugardon, 1990; Tryon and Lugardon, 1991; Blackmore *et al.*, 2000). A brief summary of this information follows.

In most liverworts, immediately following meiosis, a spore special wall is deposited directly outside of the plasma membrane. This spore special wall appears to predict exospore ornamentation. More rarely, however, ornament is determined by exine precursors produced by the sporocyte (rather than individual haploid spores). In liverworts exospore development proceeds in a centripetal fashion. WLCL are formed external to the spore cytoplasm. These take on various forms and orientations (see Brown and Lemmon, 1990, Figure 8). Sometimes they are organized in multilaminar bands consisting of numerous parallel WLCL. The entire wall essentially comprises sporopollenin deposited on WLCL and the lamellate structure is clearly discernible at maturity. There is apparently no tapetal input and thus no extra-exospore layers. Interestingly, the spores of *Sphaerocarpos* Boehmer often remain united in tetrads and the permanent tetrads may be enclosed within an envelope (Gray, 1985, 1991). These spores are similar to the envelope-enclosed cryptospore permanent tetrads produced by the earliest land plants and it has been suggested that such cryptospores derive from bryophyte-like plants closely related to the liverworts. Some workers have suggested that the cryptospore envelopes are tapetally derived (Gray, 1991; Edwards *et al.*, 1999). However, Renzaglia and Vaughn (2000) have demonstrated that in extant *Sphaerocarpos* it derives from the spore mother cell.

Spore wall formation in hornworts is little studied and poorly understood. A spore special wall is formed after meiosis and acts as a primexine in which the exospore is deposited. Interestingly, however, the exospore appears to develop without WLCL and the mature exospore is homogeneous.

In mosses, spore wall formation appears to occur in the absence of a spore special wall. Three types of spore wall are recognized among the mosses: Sphagnidae-type; Andreaeidae-type and Bryopsida-type (Brown and Lemmon, 1990).

Sphagnidae-type is illustrated by *Sphagnum* L. *Sphagnum* has a complex spore wall consisting of five layers: the endospore, lamellate inner exospore (A-layer), homogeneous outer exospore (B-layer), a unique translucent layer and the perine. The A-layer forms first and consists of 20–30 alternating light/dark layers that appear to derive from the spore, forming by sporopollenin accumulation on WLCL. The homogeneous B-layer accumulates above the A-layer. Overlying the exospore is the unique translucent layer, that consists of unconsolidated exospore lamellae in a matrix of unknown composition. Above this is the perine.

Andreaeidae-type is illustrated by *Andreaea* Hedwig, which has a spongy exospore that develops in the absence of WLCL. Development begins with the accumulation of homogeneous globules outside of the plasma membrane. These build up into an irregular layer with many interstitial spaces.

The Bryopsida-type of spore wall is homogeneous throughout, except for an inconspicuous foundation layer. The foundation layer develops first via sporopollenin accumulation on WLCL. In most cases a thick homogeneous exospore layer is deposited outside of the foundation layer in centrifugal fashion. It is likely that the outer homogeneous layer is in part of extrasporal origin, possibly deposited from the tapetum-like lining of the spore sac which may be secretory. Additionally, in some cases homogeneous material, presumably deriving from the spore cytoplasm, is deposited on the inside of the foundation layer (hence

the foundation layer is not always located at the base of the exospore, but is sometimes located within the exospore). The perine is finally deposited on the exospore. Ornament may be elaborated in the exospore below, or perine may provide the sole ornament. The perine is believed to be of extrasporal origin. The perine is not acetolysis resistant.

Lycopodium clavatum L. is a good example of a homosporous lycopsid in which spore wall development is well understood (e.g. Uehara and Kurita, 1991). Here the plasma membrane folds shortly after meiosis indicating ornament pattern. Short WLCL begin to form on the plasma membrane and are elaborated in a centripetal direction, forming the bulk of the exospore. After the outer lamellar layer is formed, an inner granular layer is deposited, but only in the proximal region. There is no perispore (or other form of extra-exospore layer). Spore wall development in all homosporous lycopsids appears to be similar to that in *L. clavatum*, except that a very thin perispore, consisting of one or two layers and forming after exospore completion, is present in some taxa (Lugardon, 1990; Tryon and Lugardon, 1991).

In the heterosporous lycopsids, spore wall structure and development differ between: (i) different taxa; (ii) the microspores and megaspores in the same taxon. Both *Selaginella* Palisot de Beauvois and *Isoetes* L. have been studied in detail and I will consider the microspores and megaspores of both.

The microspores of *Selaginella* have a bilayered exospore. The inner layer forms first. It is narrow and consists of imbricate lamellae, formed on WLCL, that are initiated in a centripetal direction. The outer layer forms second, after the inner layer is completed, and consists of amorphous sporopollenin that is deposited onto it. Extra-exospore layers vary in the microspores of different species of *Selaginella*. It may consist of a perispore, a paraexospore (*sensu* Lugardon, 1976) or be absent. In *S. selaginoides* (L.) Link (Tryon and Lugardon, 1991) a paraexospore is present. This is ontogenetically and chemically related to the exospore. It begins to form before the exospore, consisting of a granulate accumulation on the inner surface of the special wall surrounding the tetraspores. Lamellae are formed within the granulate material as the exospore begins to develop below the paraexospore. The exospore and the paraexospore are completed at the same time, as similar amorphous sporopollenin is deposited forming the outer exospore layer and also accumulates on the lamellae of the paraexospore. In some species of *Selaginella* a perispore is present, that is usually thin and firmly attached to the exospore surface.

Selaginella megaspores have a bilayered exospore. Two layers of similar thickness are recognized early in sporogenesis. The inner layer is lamellate and the outer layer consists of small and poorly delimited elements. During exospore development the inner layer does not thicken and the lamellae form a compact basal layer. The outer layer, however, increases in thickness dramatically, due to self-assembly (Hemsley *et al.*, 1994, 2000; Gabarayeva, 2000). The outer layer has a very characteristic appearance. In most species silica is deposited in the voids of the outer layer and on the exospore surface. There are no extra-exospore layers. The endospore forms between the exospore and the plasma membrane during the final stages of sporogenesis.

Uehara *et al.* (1991) describe microspore development in *Isoetes japonica* A. Br. The exospore is bilayered, with a large gap between the two layers. The outer layer forms first. Immediately after meiosis WLCL develop on the plasma membrane. These form an undulating plate consisting of two to three long and irregularly fused lamellae. The inner layer forms second. This forms by the centripetal accumulation of WLCL that develop on the plasma membrane. It comprises 12–14 lamellae (although more may be added later). At this point the lamellae of the outer exine layer are thickened by the addition of homogeneous

sporopollenin. Lastly the perispore forms, from electron dense material that is presumably tapetally derived, and the endospore forms, between the exospore and the plasma membrane. Lugardon (1990) and Tryon and Lugardon (1991) regard the exospore outer layer of *Isoetes* as a paraexospore, because it is initiated prior to the exospore, but consists of sporopollenin similar to the exospore (i.e. the inner exospore layer of Uehara *et al.*) and both are completed at the same time.

The megaspores of *Isoetes* are similar in wall structure and development to those of *Selaginella*. It is bilayered, with the outer layer initiated before the inner layer. Large amounts of silica are deposited within and above the outer layer prior to exospore completion. During the late stages of sporogenesis the endospore is deposited between the exospore and plasma membrane.

Spores in extant sphenopsids are highly distinctive in terms of morphology and development and are unique among the plant kingdom. They are most likely highly derived. Uehara and Kurita (1989) provide a detailed description of spore wall development in *Equisetum arvense* L. They note that the spore wall consists of four layers: endospore, exospore, middle layer and pseudoelaters. The exospore forms after meiosis and consists of two distinct layers (inner exospore and outer exospore). The inner exospore is first formed by the accumulation of plate-like structures that are deposited on the plasma membrane. As this layer accumulates it becomes thick and homogeneous. The outer exospore then forms. It accumulates as granular material on the inner exospore and becomes thick and homogeneous. The inner and outer exospore are similar in thickness. After the exospore is completed, the middle layer is deposited. Initially the middle layer consists of a membranous structure, which subsequently thickens up to about 0.2–0.3 μm thick. It forms in the gap between the spore and the plasmodial plasma membrane, completely surrounding the spore, but only in contact in the region of the aperture. Next the pseudoelaters form. Initially the pseudoelaters appear as membranous structures on the surface of the plasmodial plasma membrane, spirally coiled around the middle layer. The pseudoelaters are subdivided into two distinct layers. The inner layer initially comprises longitudinal microfibrils, that surround the spore in spiral fashion, but is homogeneous at maturity. The outer layer is formed by discharge of granules from vesicles in the plasmodial cytoplasm and is homogeneous. The pseudoelaters are joined to the spore, via the middle layer, at the aperture. The final part of the spore wall to form is the endospore which is deposited on the inside of the exospore. Only the exospore and middle layer are acetolysis-resistant.

In homosporous ferns the active plasmodial tapetum surrounds the spore tetrads and exospore formation commences. The exospore develops in a centrifugal fashion. It is two-layered: an inner substructure and an outer thick layer of amorphous sporopollenin. The inner substructure consists of between one and twelve partly fused sheets. These form by sporopollenin accumulation of WLCL and develop in a centrifugal manner. The number of sheets appears to be greater in more primitive ferns (up to twelve sheets) than in more derived ferns (often reduced to a single sheet). The outer thick layer of homogeneous material contains very small cavities and thin radial canals or fissures. The perispore forms after the exospore is completed, through the condensation of particulate material and is deposited from the tapetum as it decays. The perispore is always present, but is extremely variable in morphology. Where the perispore is multilayered, the inner layer is first formed with other layers successively deposited on pre-existing layers.

Spore wall development is similar in all of the heterosporous ferns and is similar in both microspores and megaspores (although the spore wall is clearly much thicker in the latter). Spore wall development is similar to that in advanced homosporous ferns, except

that the plasmodial tapetum penetrates the tetrad at an early stage enveloping each of the spores, and the perispore is replaced by an episore (*sensu* Lugardon and Husson, 1982). The exospore consists of a single substructural sheet (similar to that in more derived homosporous ferns) covered by a thick outer layer of amorphous sporopollenin. In both microspores and megaspores the outer layer consists of an episore. This is initiated between the tapetum and the incomplete exospore, and consists of amorphous sporopollenin deriving from the tapetum that is deposited simultaneously on the episore and exospore until both are completed. Thus the completed episore is partly fused with the exospore (especially in the distal and equatorial areas).

Spore wall formation in early land plants

The wealth of data obtained from ultrastructural studies of spore wall formation in extant plants has prompted the extension of similar investigations into the fossil record. Such research has been extremely profitable (see, for example, the series of investigations reported in Kurmann and Doyle, 1994). However, analysis of fossil material poses its own set of distinct problems, most important of which involves the identification of preservational artefacts. However, it has been demonstrated that well-preserved fossil spores can preserve exquisite ultrastructure (e.g. Wellman, 2001).

At present, studies of spore wall ultrastructure in early land plants is limited. Dispersed spores of latest Ordovician-earliest Silurian age have been reported by Taylor (1995a,b, 1996, 1997, 2000). *In situ* spores, from exceptionally preserved floras of latest Silurian-earliest Devonian age, have provided a wealth of information on spores that can, in many cases, be related to their parent plants (Rogerson *et al.*, 1993; Edwards *et al.*, 1995a,b, 1996, 1999; Wellman, 1999; Wellman *et al.*, 1998a,b; Habgood, 2000). Dispersed Middle Devonian spores have been described by Wellman (2001, 2002). There are a number of other reports of wall ultrastructure from a variety of Devonian *in situ* and dispersed spores (Pettitt, 1966; Fletcher, 1976; Gensel, 1979; Taylor *et al.*, 1980; Taylor and Brauer, 1983; Gensel and White, 1983; Cichan *et al.*, 1984; Hemsley, 1989; Meyer-Melikyan and Telnova, 1989; Telnova, 1993; Foster and Balme, 1994; Taylor and Scheckler, 1996).

Taylor's work on wall ultrastructure in dispersed spores of latest Ordovician-earliest Silurian age has shed light on the mode of spore wall formation in the earliest land plants, in addition to providing evidence for their affinities. Spores of this age derive from the 'bryophyte-like' plants that are believed to have comprised the land flora for the first 40 or so million years of its existence.

Taylor reports a variety of types of spore wall ultrastructure. Some have entirely homogeneous walls, and he notes that these can be produced in several ways, with the final product leaving little evidence for mode of formation. Hence little can be discerned regarding wall development in spores whose wall is homogeneous at maturity. However, many of Taylor's spores have walls that are, at least in part, lamellate. He reports a wide range of lamella morphology, with many similarities to those produced by extant plants.

Taylor notes that the walls of certain dyads are most similar to those of extant sphaerocarpacean liverworts. However, development of the envelope that often encloses early land plant spores (including monads and polyads) is problematic. Taylor considers that, in extant liverworts, the tapetum (if present at all) does not contribute to sporopollenin deposition and hence does not form a perispore. He also notes that the sporangium does not contribute to the sporopollenin wall of extant charophycean green algae, as the enclosed zygote represents the entire sporophyte generation. Thus the presence of an envelope, most

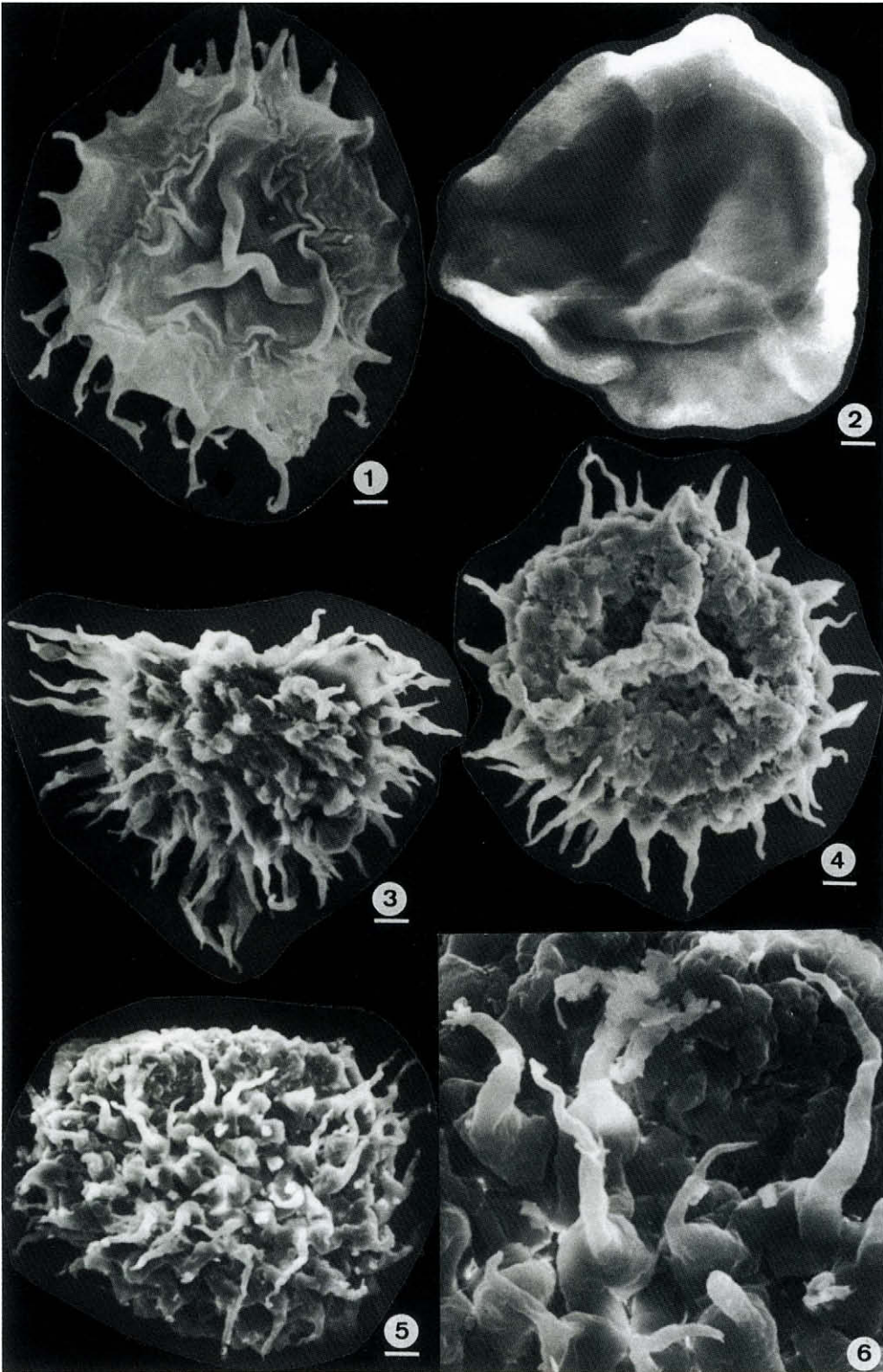
easily explained as a perispore, is at odds with liverwort affinities. Taylor, however, considers that the absence of a perispore in extant liverworts does not preclude its presence in an ancestor of that lineage. It is interesting to note that Renzaglia and Vaughn (2000) consider that ?homologous envelopes in extant *Sphaerocarpos* derive from the spore mother cell.

Taylor (2000) goes on to propose developmental scenarios for the walls of a number of early land plant spore morphologies. He concludes that 'it is difficult to envision a reasonable developmental pathway for producing these complex cryptospore walls that does not involve tapetal deposition of sporopollenin and/or the envelope (perispore)'. He also notes that these early land plants 'had already evolved the basic structural units (and the developmental patterns to produce these units) that make up the sporoderm of most modern plants'.

Wall ultrastructure has been described in a number of different *in situ* spores from exceptionally preserved floras of latest Silurian-earliest Devonian age from the Anglo-Welsh Basin (see Figures 3.8–3.9). These include trilete spores (Rogerson *et al.*, 1993; Edwards *et al.*, 1995a,b, 1996; Wellman, 1999), hilate cryptospores (Wellman *et al.*, 1998a), permanent dyads (Wellman *et al.*, 1998b; Habgood, 2000) and permanent tetrads (Edwards *et al.*, 1999; Habgood, 2000). These spores derive from a mixture of vascular rhyniophytes and plants of uncertain status (including among them rhyniophytoids). In many of these the walls are entirely homogeneous. Others are at least in part lamellate, exhibiting a diverse array of different lamella morphologies, including those typical of WLCL (Wellman *et al.*, 1998a) (see Figure 3.8). Some spore morphotypes possess outer wall layers that are globular and interpreted as being tapetally derived (e.g. Wellman *et al.*, 1998a) (see Figures 3.7 and 3.9). Many, if not all, of the structures encountered can be recognized among extant plants (principally bryophytes and pteridophytes) and include evidence of sporopollenin deposition from both the haploid spore and diploid tapetum.

Wellman (2001, 2002), has commenced work documenting wall ultrastructure in dispersed spores of Middle Devonian age from the Middle Old Red Sandstone of Scotland. The spores are exceptionally well preserved and preserve exquisite wall ultrastructure (Figures 3.1–3.7, 3.10–3.14). Some of the examined spores have morphology far more complex than that in the previously discussed simple spores of Ordovician-Early Devonian age (e.g. *Ancyrospora* Richardson and *Samarisporites* Richardson are multilayered, acamerate and zonate; *Rhabdosporites* Richardson is two-layered and camerate). These spores preserve a number of different lamella morphologies, including WLCL (Figures 3.10 and 3.12) and clear evidence for extra-exosporal layers that are tapetally derived (Figure 3.7). Despite the complexities of the spore walls, Wellman was able to link preserved structures to those present in extant free-sporing plants and hence interpret spore wall formation in terms of developmental processes observed in extant plants. Some of the spore types could clearly be related to those of extant plants. For example, it was demonstrated that spore wall formation in *Ancyrospora* was similar to that in extant lycopsids and hence the parent plants (hitherto unknown) were almost certainly lycopsid (Wellman, 2002). Furthermore, it was demonstrated that complex structures, such as inner bodies, zona, camera and grapnel-tipped processes, were easily constructed using the simple mechanisms of spore wall formation identified in extant free-sporing plants.

In conclusion, I consider that virtually all of the structure we see in the spore walls of primitive extant plants (bryophytes and pteridophytes) and in early land plants fossil spores can be accounted for by a small number of relatively simple modes of formation involving principally lamellae (of various morphologies) and tapetally-derived material. Clearly we have a very flexible system that is capable of producing immense morphological diversity that can satisfy the rapidly evolving and diverse functions of spore walls.



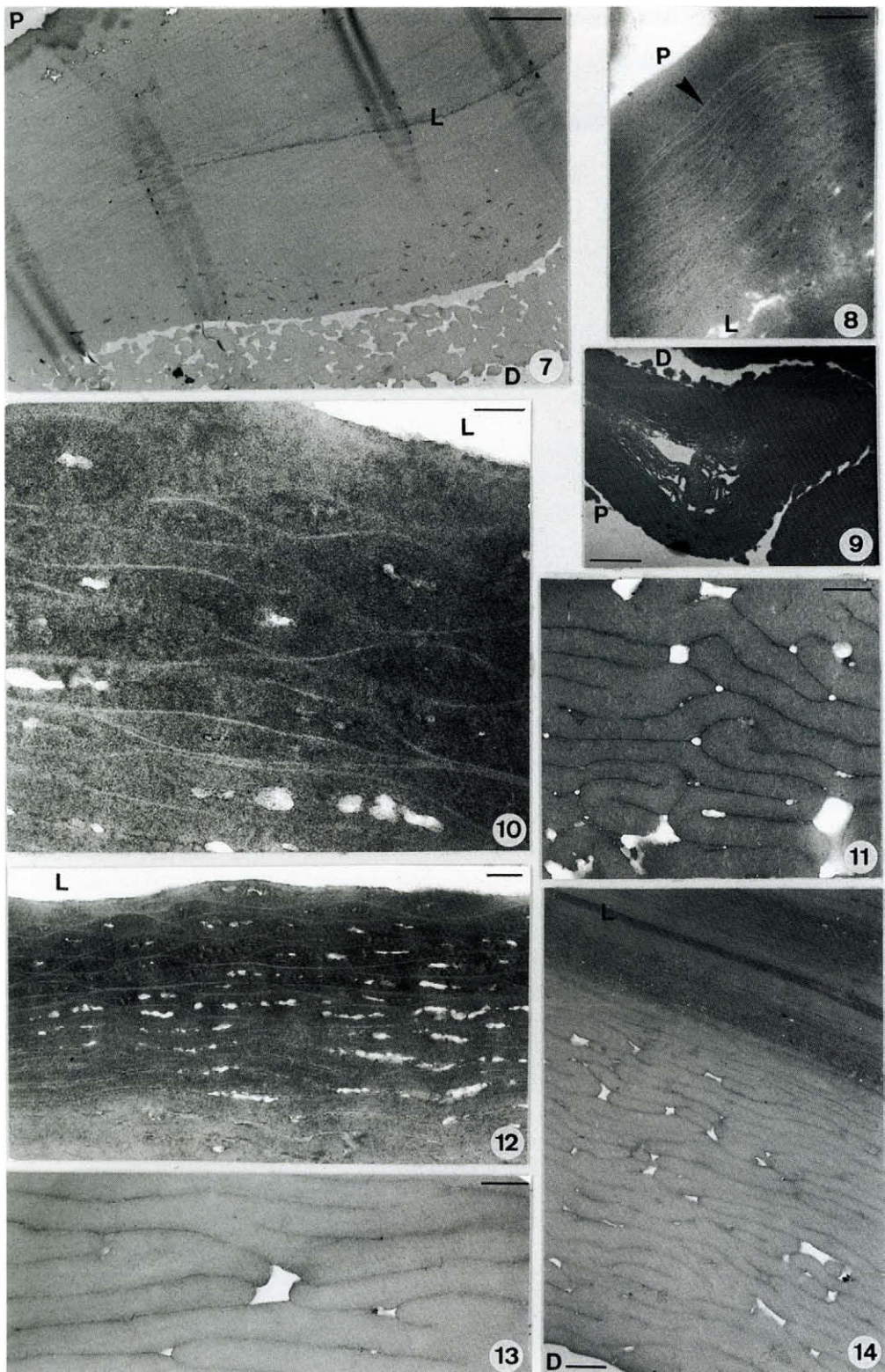
Molecular genetics of spore wall development

Research into the molecular genetics of sporogenesis is in its infancy, but understanding of this subject is progressing rapidly as a number of active research groups explore this fascinating topic. In the following section I have attempted to summarize current understanding of the molecular genetics of spore/pollen wall development. This review draws heavily on the most recent reviews of the subject (e.g. Davies *et al.*, 1992; Mascarenhas, 1989, 1990, 1992; McCormick, 1991, 1993; Scott *et al.*, 1991a), but also discusses important recent findings (e.g. Paxson-Sowers *et al.*, 2001). It is important to note that to date research has been confined to certain 'model' angiosperms (e.g. *Arabidopsis thaliana* (L.) Heynh., *Brassica napus* L.). Towards the end of this section I will discuss the potential relevance of this work on angiosperms with respect to more primitive plants (e.g. charophycean green algae, 'bryophytes' and 'pteridophytes') that are of more relevance to our understanding of spore wall development in early land plants.

Regarding genetic control of spore/pollen wall development, a couple of aspects have long been known. First, it is clear that in building the sporopollenin wall there is input from both the haploid spore and the diploid sporophyte. Secondly, control of wall patterning appears usually to be associated with the diploid pre-meiotic sporocyte. There are two lines of evidence for the latter: (1) ontogenetic studies often reveal pre-patterned templates forming in the sporocyte (e.g. Brown *et al.*, 1986); and (2) following abnormal meiosis, cytoplasmic fragments lacking a nucleus produce normal spore walls. However, for more detailed studies of the molecular genetics of spore/pollen wall development, we currently rely on research conducted on various model angiosperms.

The production of the angiosperm male gametophyte is a complex developmental process involving a tightly controlled series of cytological and biochemical changes. These are coordinated with the expression of anther-specific genes. Currently relatively little is known concerning the genes involved, although it is considered to involve a large number of both sporophytic and gametophytic genes. Evidence for the involvement of sporophytic genes includes numerous male-sterile mutations that disrupt microsporogenesis. The recessive nature of most such mutations suggests that expression of the involved genes occurs in diploid cells and not in the haploid developing pollen grain. Additionally, some male-sterile mutants exhibit altered tapetal metabolism. Evidence for the involvement of gametophytic

Figures 3.1–3.6 SEM images of early land plant fossil spores from the Eifelian (Middle Devonian) of Scotland (see Wellman, 2001, 2002). Figure 3.1 *Ancyrospora ancyrea* (Eisenack) Richardson, 1962: a bilayered, acamerate, zonate, trilete spore of probable lycopsid affinity. Scale bar = 10 μm . Note the ornament comprising long processes with grapnel-tipped endings. The function of the grapnel-tips is uncertain, but possible facilitated attachment to arthropods that possibly acted as a dispersal vector (Wellman, 2002). Figure 3.2 *Rhabdosporites langii* (Eisenack) Richardson, 1960: a bilayered, camerate, trilete spore of probable aneurophytalean (progymnosperm) affinity. Scale bar = 10 μm . The spore has a thick-walled, rigid inner body enclosed within a very loose, thin-walled outer layer, that is attached to the inner body only over the proximal surface. This 'bladder' possibly increases buoyancy. Figures 3.3–3.6 *Acinosporites macrospinosus* Richardson, 1965: a trilete spore, with a distinct apical prominence and a dense distal ornament of long and prominent spines, possibly of lycopsid affinity (based on wall ultrastructure evidence). Scale bars = 10 μm (Figure 3.3), 10 μm (Figure 3.4), 10 μm (Figure 3.5) and 3 μm (Figure 3.6). Figure 3.3 is preserved in lateral compression. Note that the apical prominence is missing (cf. Figure 3.5). Figure 3.4 is preserved in polar compression. Note the nature of the apical prominence and associated trilete mark. Figure 3.5 is preserved in lateral compression, clearly displaying the apical prominence. Figure 3.6 is an enlarged portion of the specimen in Figure 3.5 illustrating the nature of the ornament.



genes includes proof that transcription and translation from the haploid genome occurs during pollen development based on: (1) studies of several dimeric enzymes; and (2) utilizing pollen-expressed and pollen-specific clones to demonstrate activation of specific genes after meiosis and at specific periods during spore/pollen development (see Macarenhas 1989, 1992).

Male gametogenesis may be divided into three distinct phases. First, 'sporogenesis' consists of a series of archesporial cell mitotic divisions which give rise to the tapetum and sporogenous cells, the latter undergoing meiosis to produce microsporogenous cells. Secondly, 'microspore development' involves the development of the free uninucleate microspores. Thirdly, 'pollen maturation' encompasses microspore mitosis and the formation of the mature bi- or trinucleate gametophyte.

It is during the first and second phases that the pollen wall develops. In the majority of angiosperms, meiosis is synchronous, forming tetrads of microspores enclosed within a thick callose wall. This callose wall is digested and the microspores liberated, following the release of callase (β -(1,3)-glucanase) secreted into the locule from the tapetum. Pollen wall development commences while the microspores are still united in a tetrad, but following liberation there is further elaboration of the wall and the deposition of reserves such as lipids. The 'microspore development phase' essentially terminates when microspore mitosis commences.

It is generally believed that the protein components of intine are derived from gametophyte gene expression while those of the exine are produced by the sporophyte tapetal layer. The sporopollenin component of the pollen wall is considered to be polymerized from a

Figures 3.7–3.14 TEM images of wall ultrastructure in early land plant fossil spores. For all images: L = lumen; P = proximal surface; D = distal surface. Figure 3.7 *Rhabdosporites langii*: a bilayered, camerate, trilete spore from the Eifelian (Middle Devonian) of Scotland. Scale bar = 2 μ m. The spore has a thick-walled inner body, that has lamellate ultrastructure only discernible at higher magnification. The outer layer is globular and almost certainly formed by the accumulation of tapetally-derived globules. The outer/inner layers are attached over the proximal surface, where the outer layer is thin and compressed, but separated over the distal surface (i.e. camerate), where the outer layer is thick and uncompressed. However, due to compression of the fossil, the camera is reduced. Figure 3.8 *Laevolancis divellomedia* Type C (*sensu* Wellman *et al.*, 1998a): a laevigate hilate cryptospore from the Lochkovian (Early Devonian) of the Welsh Borderland (see Wellman *et al.*, 1998a). Scale bar = 0.25 μ m. The spore is bilayered, with a thick inner layer comprising structures typical of WLCL and a thinner outer layer, interpreted as tapetally derived. The arrowhead marks the junction between the two layers. Figure 3.9 *Laevolancis divellomedia* Type B (*sensu* Wellman *et al.*, 1998a): a laevigate hilate cryptospore from the Pridoli (Late Silurian) of the Welsh Borderland (see Wellman *et al.*, 1998a). Scale bar = 1 μ m. The spore wall is bilayered with a lamellate inner layer, consisting of relatively thick and laterally continuous lamellae and a homogeneous outer layer. Globules of extra-exospore material, similar to the material forming the outer layer of the exospore, adhere to the surface and are interpreted as tapetally derived. Figures 3.10–3.12 *Acinosporites macrospinosus*: a trilete spore with a dense distal ornament of long and prominent spines, from the Eifelian (Middle Devonian) of Scotland. Scale bars = 0.1 μ m (Figure 3.10) and 0.25 μ m (Figures 3.11–3.12). In the inner part of the spore wall laterally impersistent and overlapping WLCL are conspicuous (Figures 3.10, 3.12). Towards the outside of the wall these become less conspicuous (see Figure 3.12), presumably as sporopollenin has accreted onto them, forming a series of lamellae that thicken towards the outside of the wall, and are often folded (Figure 3.11). Figures 3.13–3.14 *Ancyrospora ancyrea*: a bilayered, acamerate, zonate, trilete spore from the Eifelian (Middle Devonian) of Scotland (see Wellman, 2002). Scale bars = 0.25 μ m (Figure 3.13) and 0.5 μ m (Figure 3.14). The spore has an electron dense inner body, consisting of narrow, parallel and continuous lamellae. The outer layer consists of thicker lamellae, that become increasingly thicker towards the outside of the wall. These lamellae are laterally continuous and bifurcate.

lipid-like monomer derived from caretenoids and/or caretenoid esters. The sporopollenin almost certainly derives from both the microspore and the tapetum, although the bulk usually derives from the latter. The species-specific sculpture patterns of the exine are believed to be genetically determined by the sporophyte.

The number of genes expressed in developing microspores and pollen grains (incorporating sporogenesis and gametogenesis) is considered to be very large. It has been estimated that during the lifecycle of pollen at least 15 000 different genes may be transcribed. However, substantial overlap occurs between genes active in the male gametophyte and in the sporophyte and studies of dimeric enzymes suggest that the vast majority of genes expressed in the male gametophyte are also expressed in the sporophyte, suggesting that many of these genes are simple housekeeping genes etc. Very few (usually less than 10%) are pollen specific (Mascarenhas, 1989).

Two distinct sets of genes have been identified with respect to the timing of their activity during pollen development (e.g. Mascarenhas, 1989, 1990). The so called 'early genes' are active soon after meiosis is completed. The mRNAs reach their maximum accumulation by late pollen interphase and then decrease substantially by anthesis. It is generally surmised that 'early genes' might encode cytoskeletal proteins and proteins needed for wall synthesis or starch deposition. The so-called 'late genes' are activated after microspore mitosis. The mRNAs increase in content up to maturity, suggesting a major function during germination and early tube growth (and possibly also during the later parts of pollen maturation).

Far more is known about the 'late genes' than 'early genes', which is unfortunate as the latter are almost certainly those involved in spore wall formation. This disparity occurs because the majority of studies have been based on cDNA libraries constructed from mature pollen or mature anther RNA, and pollen-specific cDNAs thus isolated are representative of these 'late genes'. We do not at present have any estimates for the numbers of 'early genes' or of their similarity to genes expressed in later development.

However, workers such as Scott *et al.* (1991a) have illustrated a strong allometric relationship between bud length/anther length and stages of anther development. They have demonstrated how this relationship can be utilized in the isolation and characterization of anther-specific cDNAs expressed at defined stages of anther development, including early stages (i.e. 'early genes'). Examples of identified 'early genes' are actin, alcohol dehydrogenase, β -galactosidase and those specifying the anther-specific tobacco transcripts TA25 and TA29 (see for example: Stinson and Mascarenhas, 1985; Albin *et al.*, 1990, 1991; Koltunow *et al.*, 1990; Smith *et al.*, 1990; Scott *et al.*, 1991b; Theerakulpisut *et al.*, 1991; Foster *et al.*, 1992; Roberts *et al.*, 1993).

For some of the identified 'early genes' possible functions have been suggested, including, in some cases, involvement in spore wall formation. Scott *et al.* (1991b) isolated I3 cDNA from immature anthers of *Brassica napus* that exhibits microspore-specific expression. They speculated that the protein encoded by the gene represented by I3 may be involved in early cell surface structure of developing pollen, or even in directing the development of the intine. Foster *et al.* (1992) report on the isolation of a *Brassica napus* mRNA which is expressed during the 'microspore development phase' in both the developing microspore and the tapetum. The predicted protein of the E2 cDNA shows high homology with phospholipid transfer proteins (PLTPs), and Foster *et al.* speculated that the E2 PLTP isoenzyme might be involved in sporopollenin production.

More recently Paxson-Sowders *et al.* (2001) identified a gene (DEX1) they considered important in exine pattern formation. They worked on the *dex1* mutation of *Arabidopsis* which disrupts exine formation and patterning. Pollen wall development in *dex1* plants is

similar to that in wild-type plants until the early tetrad stage. Thereafter, in *dex1* plants, primexine deposition is delayed and reduced, rippling of the plasma membrane and spacer production is absent and the produced sporopollenin is randomly deposited on the plasma membrane, but is not anchored to the microspore, and forms large aggregates on both the developing microspore and the locule walls. Following isolation and molecular characterization of DEX1 it was demonstrated that it encodes a novel plant protein, that was predicted to be membrane associated and to contain several potential calcium-binding domains. Factors controlling exine pattern are poorly understood and structures such as the primexine matrix, plasma membrane, endoplasmic reticulum and microtubules have been implicated. Interestingly, this work suggests that the plasma membrane does have an integral role in exine pattern formation. The fact that sporopollenin was synthesized and deposited, but failed to anchor to the surface of the microspore, suggests that DEX1 may function as the nucleation point for sporopollenin deposition. The authors speculated that DEX1 may either be a component of the primexine matrix or endoplasmic reticulum and involved in the assembly of primexine precursors. Recent discoveries of a number of other mutants in which exine development is disrupted, in *Arabidopsis* (e.g. Taylor *et al.*, 1998) and other plants such as *Zea mays* L. (Loukides *et al.*, 1995), are encouraging regarding the progress of similar research in the near future.

It is clear from the brief review outlined above that studies of the molecular genetics of spore/pollen wall formation are in their infancy. As more work is undertaken and techniques improve we can expect a dramatic, and hopefully rapid, increase in our understanding of these issues. Initially such work will be confined to angiosperms, but it is hoped that studies on other plant groups (gymnosperms, 'pteridophytes' and 'bryophytes') will follow. Only then will we be able to assess similarities/differences between the molecular genetics of spore/pollen wall formation in angiosperms and more primitive plant groups and ascertain how conserved the molecular mechanism and involved genes actually are. Also fruitful, and of direct relevance to early land plants, will be comparisons with the charophycean green algae, the sister group to embryophytes (land plants). Of interest will be the molecular genetics of the development of the sporopollenin layer surrounding the zygote, which will potentially shed light on the evolution of sporopollenin walled reproductive propagules in the earliest land plants.

Conclusions

It is concluded that the sporopollenin wall surrounding charophycean zygotes and embryophyte spores/pollen grains is homologous, and that the spore/pollen wall is an embryophyte synapomorphy that evolved as an adaptive response to the invasion of the land. It is considered that the primary function of the spore/pollen wall involves protection in the harsh subaerial environment. However, spore/pollen walls have taken on multiple secondary functions (exaptations), particularly as they evolved in response to changes in the habit and external environment of early land plants. Regarding spore wall development, ultrastructural studies have demonstrated that structural elements present in the spore walls of extant plants can be recognized in the fossil spores of early land plants, suggesting that spore wall development was similar in extant and ancient land plants. It is clear that spore wall construction is a simple and flexible process allowing rapid evolution of complex structure and morphology in response to evolving mode of life and external environment. Finally, studies of molecular genetics of spore wall formation are in their infancy, but have the potential to

solve many unanswered questions regarding spore wall homologies and developmental processes, particularly when such studies are commenced on more 'primitive' land plants (bryophytes and pteridophytes) and their extant sister group (charophycean green algae).

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4

The evolution of plant biochemistry and the implications for physiology

Richard D Firn and Clive G Jones

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Introduction

Plant physiology and plant biochemistry are seen by some as brother and sister but by others as distant cousins. All would agree that there is a relationship but few would agree how close it is. The fact that books have been written on plant biochemistry with very little discussion of the physiology of plants shows that families can lose touch with each other. The aim of this chapter is to show that a functional understanding of plants can be gained by integrating physiological and biochemical knowledge. Our attempts to bring about integration start with an evolutionary perspective.

When introductory biochemistry is taught, there is a tendency initially to concentrate on the central metabolic pathways of 'the cell'. The default cell is usually mammalian. Plant cells are usually treated as unusual, but only in so far as they have walls and chloroplasts. This emphasis on the commonality of the basic metabolism of organisms is helpful to those learning the subject but it does tend to limit conceptual approaches to biochemistry. The reality is that most cells are biochemical specialists. Evolution has selected for biochemical traits of cells that are appropriate for their particular cellular environment in a manner analogous to the selection of organisms that are more suited to their environment. This specialization is found in both unicellular and multicellular organisms. However, in multicellular organisms, higher order coordination and control – physiological processes – can have a marked effect on the environment of specialist cells. To provide this coordination, specialist cells have evolved to play a central role in controlling many plant physiological processes (e.g. gas exchange is regulated by the guard cells, abscission depends on the abscission layer, seed reserve mobilization in grasses is dependent on the aleurone cells, etc.). Consequently there will be very clear links between physiological and biochemical processes. In this chapter we will argue that the selection pressures operating on the evolution of metabolism will have given rise to certain metabolic traits and, because of the intimate links between biochemistry and physiology, those traits have helped shape physiological processes. The intimate connection between the physiology and biochemistry of plants is well established in the case of some aspects of physiology, especially where there are dramatic morphological and anatomical adaptations which make the links clear – photosynthesis, germination, stomatal functioning, for example – and the links between the physiology and biochemistry of these processes will not be considered here. Instead, some more general principles concerning the evolution of metabolism will be discussed. These simple principles will then be used to consider the links between metabolism and physiology in cases where the anatomical and morphological signposts are less clear.

Molecular evolution, biochemical evolution and metabolic evolution – hierarchical terms

The term *molecular evolution* is currently used to refer to the way in which the evolution of protein structure relates to protein function, frequently with emphasis on tracing the lineage of base sequences in specific genes. Those working on molecular evolution generally focus on the role of one protein (or family of proteins). *Biochemical evolution* overlaps with molecular evolution but can extend to consider more than one enzyme in a pathway and the control of that pathway. *Metabolic evolution* could be considered to operate at the next higher level, determining the way in which the whole of metabolism evolves, with extensions or deletions of metabolic steps being the result of descent via modification in lineages subjected

to natural selection. It is *metabolic evolution* that determines the scope of metabolism in an organism, with biochemical evolution and molecular evolution determining the degree to which operation of the enzymes and their controlling elements effect organismal fitness. It is *metabolic evolution* on which we focus, because it is at this level that links between physiology and biochemistry are most apparent.

Metabolic evolution – what determines whether a new enzyme is retained?

New enzymes usually arise as a result of gene duplication and subsequent mutation of one gene copy such that the mutated enzyme has an altered substrate tolerance and can act on a new substrate to produce a new product (Petsko *et al.*, 1993). Whether that new metabolite enhances the fitness of the producer depends entirely on whether the benefits of possession of the new product exceed the costs of producing it. In turn, the benefits depend upon the properties that the new metabolite brings to the organism. We identify three major classes of property that natural selection appears to have favoured during *metabolic evolution*.

Biomolecular activity – the evolution of ‘secondary metabolism’

It has been well established from screening trials of collections of synthetic or naturally occurring molecules that the probability of any individual chemical possessing potent biological activity is very low (Jones and Firn, 1991; Firn and Jones, 1996). We proposed that this fact must have been a great constraint on the evolution of pathways leading to molecules that benefit their maker by possessing biological activity. However, the relevance of this fact to the understanding of the evolution of secondary metabolism has been challenged (Berenbaum and Zangerl, 1996). Berenbaum and Zangerl argued that the analogy between humans screening for a useful biological activity and organisms evolving chemical diversity in order to gain fitness by making biologically active chemicals was inappropriate – the nub of their argument was that vagueness in the definition of the term ‘biological activity’ led to a false analogy. However, by defining the term *biomolecular activity* quite precisely (as the ability of a molecule to interact with a biologically functional molecule such that its biological function is significantly changed), Firn and Jones (2000) countered this objection and provided evidence that the low probability of any chemical possessing potent *biomolecular activity* is a predictable and well understood consequence of ligand–protein interactions. Consequently, they reiterated a refined argument stating that the low probability of specific ligand–protein binding has been a significant evolutionary constraint on the production of biologically active molecules by organisms. Humans and other organisms may adopt different means of trying to reduce the impact of this fundamental constraint but they have both had to evolve ways of doing so. The methodology that humans have used when seeking to exploit the biological activity of chemicals (for example as pesticides or pharmaceuticals) has been the development and use of screening trials. These trials provide useful data about the frequency of any particular biological activity occurring in random collections of chemicals. Because such trials have been used to screen synthetic and naturally-occurring chemicals, and because they have been used against a wide range of biological targets, the information available from such trials is very extensive.

Do screening trials reveal any other relevant information? Indeed they do. They show that the type of biomolecular activity possessed by any individual chemical is unpredictable. Humans synthesizing new chemicals to test for biological activity are often surprised that a chemical made with the intention of producing one type of biological activity actually turns out to possess a very different, equally valuable activity. The discovery of Viagra is a recent well known example of human serendipity, but previous examples abound (Roberts, 1989). The herbicides paraquat and diquat emerged from an observation that a surfactant used in an experimental formulation was surprisingly phytotoxic, which led to some diquateryny dyestuff intermediates being examined in a herbicide screen with the result that the bipyridylum herbicides were discovered. Another example would be that the discovery of the pyrimidine fungicides started with attempts to make insecticides. In other words, the structure of any individual molecule is only partly predictive as to whether a molecule will possess biomolecular activity and is poorly predictive as to the type of biomolecular activity. The successful organism, like the successful company, exploits fortuitous events. Consequently, it is reasonable to predict that as a result of evolutionary adaptation, a pathway in an organism initially leading to one form of biomolecular activity can eventually lead to a quite different form of biomolecular activity. Hence we have two constraints that must be taken into account when considering the evolution of chemical diversity in plants – any new molecule has a very low probability of possessing biomolecular activity and the type of biomolecular activity cannot be reliably predicted from the biomolecular activity (if any) of the chemical(s) from which it derived.

Molecules retained because of their physicochemical properties

The terms 'primary metabolism' and 'secondary metabolism' are very commonly used, however, there are large groups of chemicals in organisms that sit between these two traditional groupings. Lipids, many pigments such as the carotenoids, many polysaccharides and some anthocyanins fall into this category. For example, consider lipids. All cellular organisms need lipids but they do not need them all – as a group lipids are essential but individually they are not. The paradox can be resolved by recognizing that the properties that have been selected for are physicochemical traits – lipophilicity, light absorption, structural properties, etc. These properties depend on the molecular properties shared by large numbers of structurally similar chemicals. Minor changes to part of the molecular structure might predictably make little difference to these properties – this is a marked contrast to the impossibility of predicting how such changes might alter the biomolecular activity of the same molecule. (Once again the analogy with chemicals made by humans is appropriate. A chemist who has synthesized a novel chemical can predict with some confidence its lipophilicity or its spectral properties but cannot reliably make any precise predictions about the biomolecular activity of the molecule.) Thus an organism that gains fitness by making a chemical that protects it against harmful irradiation, might as a result of extending the pathway leading to that pigment, produce another chemical with predictably similar useful properties. Slightly different extensions of the pathway leading from that point might have similar chances of creating new molecules with similar properties to the common precursor hence it is predictable that different species will produce a rather different mix of such chemicals – their overall requirement for a certain mix of chemicals with an average physicochemical property can be met in many different ways. The diversity of molecules selected for their physicochemical properties is thus predictable but the contribution that any one metabolite will make to the overall requirements is unpredictable. Under these

circumstances, it would seem that the advantage given to the producer of a new chemical depends greatly on the existing overall mix of existing similar chemicals.

The diversity of molecules selected for their contribution to the overall physicochemical requirements carries with it a chemical diversity that can be exploited as a route to molecules possessing useful biomolecular activity. Thus it is predictable that compounds with useful biomolecular activity will sometimes arise from pathways usually considered to operate principally for other purposes. Thus the phenylpropanoid pathway can generate chemicals that have a role in absorbing visible or non-visible light but this pathway may also lead to compounds that can enhance the fitness of the producer because they possess useful biomolecular activity (Winkel-Shirley, 2001). Likewise, the isoprenoid pathway can give rise to photoprotecting pigments such as the xanthophylls (Taiz and Zeiger, 1991) or carotenoid pigments in flowers or fruits (Harborne, 1988), yet also gives rise to the plant hormones such as the gibberellins or abscisic acid (Davies, 1995).

Primary metabolism – canalized metabolism, each step depending on other pre-existing metabolic capabilities

For the purpose of this discussion we define ‘primary metabolism’ as metabolic pathways where there is a great interdependence of the individual steps and where the individual molecules made by an enzyme serve only to feed into another enzyme – the complete pathway is greater than the sum of the parts. A significant difference between ‘primary metabolism’ and the previous two categories of metabolism, is that in primary metabolism selection is less environmentally contingent and the evolved homeostatic mechanisms of the organism and of the cell make the selection pressure more constant. At some stage early in the evolutionary history of primary metabolism the incorporation of a new metabolite into the developing primary metabolism could only have occurred if the new molecule fitted usefully into the specific scheme of existing pathways. The actual advantage to the producer of this new metabolite would arise solely from the ability of the new molecule to be acted upon by an existing enzyme(s) to produce another molecule(s) that had properties that enhanced fitness. The most extreme outcome would be the production of a metabolic cycle where there is no single ‘end point’ on which selection can act but the overall net efficiency of the cycle is subject to intense selection. However, in such cycles (e.g. Calvin cycle, the photorespiratory carbon oxidation cycle, the C₄ cycle, etc.), selection that has fitted existing capabilities to local circumstance makes it much harder to introduce a new, compatible transformative capacity. Consequently, the pathway becomes severely canalized – the optimization of the coordinated processes increasingly reduces the opportunities to evolve radically different methods of achieving the same goal. (A dramatic example of canalization would be the genetic code – it is only one possible code of many that could have been used but once organisms evolved with a workable system on which selection could improve, the die was cast.) The powerful selection against new chemical diversity is in contrast to the previous property classes where there is a tolerance of chemical diversity (physicochemical properties) or selection for chemical diversity (biomolecular activity).

Selection for different molecular properties has consequences for metabolic evolution

Although it has been biochemical dogma that enzymes are highly substrate specific, this dogma has largely arisen from studies of enzymes involved in primary metabolism where

canalization is severe. The biochemical properties evident in a highly evolved and specialized metabolism tell us about the outcome of the selection processes operating on that sphere of metabolism and provide little reliable guidance as to the properties of individual new enzymes arising as a result of mutation. We would argue that a mutant enzyme would have a high probability of possessing a broad substrate tolerance initially. The reasoning is that a mutant enzyme that cannot access an existing substrate because it has narrow substrate specificity cannot produce a new product. Hence the enzyme will not be selected for in any circumstance in which new products would confer evolutionary advantage. In contrast, a mutant with a broad substrate tolerance has a greater initial probability of producing new substances with useful properties, irrespective of the type of property considered. However, it is clear that the selection pressures that would operate after the new product(s) have been generated would differ greatly depending on the type of property brought to the cell. For example, gaining fitness by producing biomolecular activity requires chemical diversity to be generated and maintained, hence there might be little selection pressure narrowing substrate tolerance (Jones and Firn, 1991). Indeed, enzymes capable of acting on more than one substrate might bring benefits. So if a new enzyme produces a new lipid that adds to the physicochemical properties of the cell, as long as the enzyme does not generate a chemical with adverse cellular properties, there may be little selection to narrow the initial broad substrate tolerance. Where there will be strong selection for narrow substrate tolerance is in the final property class – participation in primary metabolism. Here an enzyme accessing a common, important intermediate would be expected to have a negative effect on the overall metabolic and homeostatic mechanisms due to substrate competition and the possible generation of compounds that act allosterically to hinder rather than aid metabolic control. These simple concepts predict that metabolic evolution will have produced rather different metabolic traits in different pathways and even at different stages within a pathway.

Biochemical evolution and physiology

Why should the ideas outlined above be of interest to physiologists or biochemists? The fact that these issues have rarely been addressed previously suggests that both physiologists and biochemists have been comfortable working without such general principles for most of the 150 years that these disciplines have existed. In order to help promote interest in building the evolutionary frameworks underpinning the links between plant biochemistry and physiology, we will use two topics as examples of how simple ideas about metabolic evolution could guide experiments:

1. the physiology of chemical interactions between plants and other organisms
2. the physiology of intraplant signalling (plant hormones).

The former area is chosen because it is one where an alternative evolutionary model – based on ecological rather than metabolic considerations – was well developed and widely accepted. That model is now challenged by our metabolic evolutionary model. The second area – the evolution of plant hormonal control – is chosen because plant hormones have been shown to play a role in many very important, basic physiological processes.

The interaction of plants with other organisms

The human experience

Human experience has been both a useful guide and a distraction in understanding the role of plant chemicals in the coevolution of plants with other organisms. Humans are themselves massively influenced by plant chemistry, although most people go about their lives unaware of this fact because plant chemicals are so embedded in most cultures. Few readers will be reading this sentence without some plant chemicals being active in their bodies – it will be the rare reader who starts this chapter (and an even rarer reader who finishes this chapter) without having taken one of the following during the previous few hours – coffee, tea, tobacco, chocolate, a recreational or prescription drug, a tasty wine, a flavoursome beer, a fruit drink or a piece of confectionery. If a reader cannot concentrate on this chapter because their mind would prefer to think of the meal that awaits them, it will not be the expectation of a plate of starch or a piece of protein that excites their mind but the thought of the pasta sauce, the exciting curry, the sharp onions, garlic or interesting green salad. Plant secondary chemicals feed the mind while plant storage products feed the body. The human craving for particular plant chemicals has been so powerful that it has driven colonial expansion in the past and today many national economies are dependent on such chemicals. The power that a few plant species exert over humans by making just a few strange chemicals is quite remarkable and yet traditionally we have considered such chemicals ‘secondary products’! If we delve a little deeper into the human experience with such chemicals we find that humans value such chemicals for different reasons:

1. such chemicals attract or repel humans by acting on sense organs – smell and taste mechanisms have been honed by evolution in many animals just to identify potential food sources and avoid intoxication based on their chemistries. Had there been no chemical diversity in the plant or microbial derived material would we have such fine senses?
2. the ability of these chemicals to influence mental processes (behaviour, well-being, etc) means that we can change our perception of the world
3. less commonly in humans, these chemicals can have a physiological effect, acting on some metabolic pathways in a positive or negative manner.

Each of these apparently different modes of action shares a common feature in that the effect of the chemical at a cellular level is caused by the chemicals binding to a specific receptor. Here the rules of ligand/receptor interactions apply. However, there is one crucial difference in the taste/smell receptors that differentiates them from the neurological and physiological receptors. In the case of the taste/smell receptors, the receptors have evolved to detect ligands and that means that there is potential for coevolution of the ligand/receptor interaction. A pollinator that is attracted to a food source by a plant-derived odour gains fitness if it is a mutant that has an odour binding protein that better matches the structure of the odorous chemical – the evolution of the detection system becomes ‘locked on’ to the chemistry of the producer. In contrast, neurological and physiological receptors are usually ‘fortuitous receptors’ – the receptor proteins have roles unrelated to their fortuitous ability to bind the plant-derived chemical and selection for their primary role would be paramount. If the plant-derived chemical has a serious adverse effect on the organism by interrupting its normal function there will be massive selection pressure to select individuals that make a mutant protein which functions in its original role but which cannot bind

the plant-derived chemical at the concentration that it occurs – the evolution of detection is ‘locked off’ (this is why insecticide-resistant pests and antibiotic-resistant microbes are an inevitable consequence of human attempts to use chemicals to reduce the fitness of competing organisms).

The human experience of plant and microbe-derived chemical diversity has thus been very important in human affairs and this experience can inform us about ligand/protein interactions but it can tell us little about the role of plant chemicals in the organisms that make them. However, this human experience does tell us a final, very important lesson. The great majority of chemicals made by plants or microbes have no direct impact on humans in any observable way. Even in plants grown and consumed by humans, the majority of plant chemicals made by those plants are unsensed by humans, these chemicals have no unambiguous physiological or neurological effect and there must be many such chemicals that are as yet uncharacterized by humans (who have tended to concentrate on the compounds that occur in large amounts or which are physiologically or neurologically active).

Plant/microbe and plant/insect interactions

Being (with a few exceptions) primary producers, plants are subject to attack by other organisms that seek access to the resources captured by the plant, resources that are now in a form usable by attackers. The evolution of plant physical and life cycle strategies which can reduce the rate, frequency or effectiveness of attack has obviously been an important feature of plant evolution (Rauscher, 1992) but such strategies will not be considered further, rather we shall concentrate on chemical defences. However, it is worth noting that there is an intimate link between physical defences and chemical ones. As in the case of physical defences used by humans, the chemistry of the material used to construct the physical form is crucial to its effectiveness. Likewise in plants, the cuticle, some trichomes and cell walls have chemistries that make the physical structure more suited to its purpose. The evolution of the chemistries of such structures is a topic which the following discussion might inform but it will not be considered further. We will concentrate on low molecular weight chemicals which are made by plants to gain fitness by acting directly on the interacting organism.

Because many organisms interacting with plants use volatile phytochemicals in order to locate the plant, plants will have evolved in response to the selection pressure that is associated with these volatile-mediated interactions (the clearest example of such selection comes from human selective breeding where human preferences for certain scents and flavours has resulted in plants (for example roses, carnations, apples, etc.) with extreme characteristics being bred and widely propagated throughout the globe by humans). The attraction of insects to a few plant volatiles can result in increased fitness for the plant in the case of pollination or in attraction of the insect natural enemies of plant-feeding insects (Harborne, 1988; Vet, 1999). However, a role can only be assigned (as an attractant or repellent) to a very few plant volatiles in the complex mixture made by any single plant species. There has certainly been a tendency to focus attention on the very small fraction of plant chemistry that possesses clear biological activity when building an evolutionary framework explaining all plant ‘secondary’ chemical diversity. Being impressed by the potent biological activity of some secondary chemicals, it was argued that these chemicals are used by the plant to increase fitness by negatively or positively affecting the metabolism or behaviour of other organisms (Fraenkel, 1959; Ehrlich and Raven, 1964; Harborne, 1988). To account for the tens of thousands of secondary chemicals for which there was no convincing evidence that the production enhanced the fitness of the producer, various

explanations were offered. For example, given the diversity of other plants, animals and microbes that come into contact with a single plant species over evolutionary time and throughout the contemporary geographical range, it was argued that in order to confer specificity to the chemical interactions between a plant and these other organisms, a very large number of biologically active compounds would be expected to be found in any single plant species. This idea, supplemented with an idea of evolutionary relics, led to the view that 'every natural product has, or had, a purpose in the evolutionary strategy of the taxon concerned' (Swain, 1975). This model ignores the fact that the probability of any compound possessing any biomolecular activity is very low (see earlier). To generate a new chemical with useful biomolecular activity, repeatedly at each stage of a linear pathway, is extremely unlikely. Furthermore, there are countering selection pressures which would further reduce the chances of such linear pathways evolving. Consider a scenario, where by such extreme chance, a plant species has evolved a four-step synthesis of a compound that reduces the fitness of a herbivorous insect. Once the end product of the pathway has lost its effectiveness (which it will do rapidly judging by >100 years experience with evolution of resistance to many synthetic pesticides and antibiotics), the chances of the plant gaining fitness by evolving another novel new biologically active chemical from the now-redundant chemical is much lower than the chance of gaining fitness by reducing the costs by eliminating the now redundant four-step pathway. The longer the pathway the greater the problem becomes because the number of genes in which a mutation can give cost savings is greatly in excess of the number of genes that can give rise to a new useful product. Furthermore, in each gene, there will be a greater chance of destroying enzyme activity via mutation than changing it usefully. The alternative model to explain secondary product chemical diversity (The Screening Hypothesis; Jones and Firn, 1991), was based on well-defined general principles of ligand/protein interactions instead of being based on limited and maybe selective ecological evidence. The Screening Hypothesis proposes that organisms that gain fitness by making compounds with potent biomolecular activity are selected for because of the overall metabolic traits they possess that enhance the generation and retention of chemical diversity. Some of the chemicals made by this metabolic capacity will possess fitness enhancing properties but many (the great majority?) chemicals made will possess no properties that contribute to the current fitness of the maker. Many concepts that have driven previous studies should now be questioned:

1. *Presence indicates purpose?* Not necessarily. The presence of a chemical in a plant, even if the chemical is shown to possess interesting biomolecular properties in some assay, is insufficient evidence that selection has operated to promote the synthesis of this specific property. The safer deduction is that selection has operated to retain the overall metabolic capacity which gave rise to this and other molecules, one or more of which must possess properties that enhance fitness, provided that the overall fitness benefits outweigh the overall metabolic costs.
2. *Quantity indicates purpose?* Not necessarily. It is known from many structure-activity studies that the relative biomolecular activity of several structurally related molecules can vary by many orders of magnitude (Firn and Jones, 1996) hence the amount of chemical made by a plant provides little evidence of role.
3. *Pathways indicate purpose?* Not necessarily. Because the biomolecular properties of a chemical are unpredictable, a pathway at one stage in evolution contributing to plant defence against insect herbivores could subsequently contribute to another property (such as reducing the fitness of a pathogen).

4. *Compound X has been shown to defend plant A hence surely a similar chemical in plant B plays a similar role?* Not necessarily. The different evolutionary experience of different species and the flexibility of the metabolic traits used to make molecules possessing biomolecular activity means that similar compounds could serve different roles in different species and the different compounds could serve the same role in different species.
5. *'A well-known physiological function of the anthocyanin pigments and flavonol copigments is the recruitment of pollinators and seed dispersers....'* (Winkel-Shirley, 2001). *'Glucosinolates are biologically active secondary metabolites....'* (Kliebenstein *et al.*, 2001). Both these statements come from papers published recently and they are inaccurate in that they generalize for a pathway and imply that all the compounds made from a particular pathway play a particular role. It has not been shown that all flavonoids play a role in attracting insects. It has not been shown that all glucosinolates are biologically active (and it would mean little unless that activity was shown to be of benefit to the producer). It might be true to say that some members of these chemical groups play a particular role but there needs to be the recognition that most chemicals made by these pathways have not been shown to play any role.

The Screening Hypothesis clearly places great demands on those studying the role of chemicals in plants (and microbes). The 'rules' which have shaped metabolism in plants are operating at the molecular level and the outcome of these rules is usually studied at higher levels of organization. The rules do not predict outcomes. As in a game of chess, the few simple rules cannot predict the outcome. The operation of any rules simply opens up multiple opportunities and it is the players that ultimately shape the game.

The evolution of regulatory systems for secondary metabolism

As in the case of the immune system (another strategy evolved to counter the low probability of any antibody possessing the correct properties to enable it to bind at low concentrations to a specific hapten), an ability to induce the chemical defence only when needed provides a very great cost saving. To achieve cost saving by having inducible defences three extra elements are needed:

1. a sensing system(s) that can detect the conditions that are an accurate indicator of a need for defence
2. a regulatory step in the chemical production capacity (by either regulating transcription, translation or by allosteric means)
3. a linkage mechanism between the detection system and the regulatory system.

Evidence is accumulating that plants possess all three of the above abilities and evolution may have provided multiple ways of linking the three elements. Damage or invasion detection mechanisms involving the detection of low molecular weight compounds or proteins, either arising from the attacker or as a result of the attacker creating low molecular weight compounds as they break into the plant, have now been found in many plants (Karban and Baldwin, 1997; McDowell and Dangel, 2000). One linkage between the detection system and the regulation of gene expression involves transcription factors (Tamagnone *et al.*, 1998), however, we shall not consider the detailed mechanisms involved in such responses.

Rather, we will consider the evolutionary strategy that has given rise to inducibility. How could an organism best gain the cost-saving benefits of inducibility yet retain the flexibility to generate and retain chemical diversity? Might not a well-defined regulatory system operating on a pathway begin to canalize an area of metabolism in a manner which ultimately constrains the production and retention of new chemical diversity?

A speculative scenario for the evolution of the control of pathways leading to compounds retained because they possess biomolecular activity

Given the advantages that inducibility confers (reducing costs increases chances that benefits outweigh costs), it is expected that inducibility should have evolved at an early stage in the evolutionary history of any secondary product pathway. Once the inducibility had evolved at that position in a pathway, as long as the mechanisms provided adequate control of the amount of active compound made at a later stage along the pathway, the selection pressure to regulate the pathway at a later stage might be small. Thus regulation of flux through a pathway could be achieved by regulation at an entry point or early stage of a metabolic sequence. An immediate implication of this logic is that inducibility becomes quite a poor predictor of the role of any chemical made as a consequence of a pathway being induced by a particular stimulus. Just as the inducibility of the immune system tells one little about the role of each type of antibody, maybe the inducibility of secondary chemicals after a biotic challenge indicates a mechanism of response and not a role for each chemical made. However, a further complexity is introduced because of the predicted multifunctionality of secondary metabolic pathways. Because a pathway may serve different roles at different stages of evolution, or in different organisms, how can natural selection result in a sensing/induction system that adapts to the new role that a pathway may best serve? Consider two extreme scenarios:

1. At one extreme, a particular pathway could evolve with associated regulatory processes finely tuned to deal with one specific type of challenge only – an insect herbivore defence system using products of an alkaloid pathway, for example. This would result in excellent cost savings when it first evolved, but the chances of evolving novel anti-insect compounds (i.e. compounds that are sufficiently different in their mode of action from a now redundant one for which the insect now has evolved to resist) would be greatly reduced if there was a reliance on this one pathway. If the insect damage sensing system is uniquely linked to this alkaloid pathway then to evolve an extension of a non-alkaloid pathway brings a requirement for a whole new regulatory system for that pathway in order to gain cost savings.
2. At the other extreme, regulatory systems for several pathways could be evolved which could respond to one or more of several different challenges (insects, fungal, physical damage, etc.). Such a strategy would increase the chances of evolving new compounds with biomolecular activity able to serve any purpose. Thus a pathway evolved because it enhanced the fitness of a plant by making compounds that reduced the fitness of insects could at any time produce a compound that enhanced plant fitness by reducing the fitness of an invading pathogen (see the earlier analogous human experience of using whatever biological activity one finds despite the original purpose). The multifunctionality of the pathways maybe dictates a multifunctional sensing system.

These extremes are not mutually exclusive and a plant may have different pathways which fall anywhere along the spectrum of the extremes. However, in recent years there has been

a growing awareness of 'cross talk' between signalling and response mechanisms in plants (Felton *et al.*, 1999a,b; Feys and Parker, 2000). This is precisely what would be predicted by one of the extreme scenarios just discussed where the multifunctionality of a pathway leads to a flexible response system which has 'cross talk' built into it. The possibility that cross talk is an inevitable consequence of the metabolic traits of such pathways has many consequences for those studying the role of products made by such pathways. Inducibility becomes a very poor predictor of the role of a chemical because inducibility has possibly been evolved as a general means of cost-saving to be applied in a non-specific manner because the underlying metabolism needs to retain the ability to generate chemical diversity.

Signalling molecules within plants

The concept of specific chemicals acting as the controllers of developmental and functional processes in plants has dominated the thinking of plant physiologists for many decades. At the centre of this thinking are the well established roles for the five major groups of plant growth substances (auxin, ethylene, abscissic acid, cytokinins and the gibberellins (Davies, 1995)). However, there are numerous reports in the literature of a very wide range of other chemicals (nearly always secondary plant metabolites) purportedly playing a regulatory role (Gross, 1975). It is often suggested that these secondary metabolites either replace or supplement the five major types of plant growth substances in particular circumstances (Gross, 1975). During the 35 years (1928–1963) that the major hormone groups were being discovered, a large number of plant extracts were tested for biological activity in plant-based bioassays, and many reports of new endogenous regulators appeared during that period. Possibly because the discovery of each of the major groups of plant hormone was unusual in some respect, with the active compounds first being isolated from unlikely sources or in a study that did not establish an unambiguous role (Table 4. 1), close scrutiny was not always given to other claims that new endogenous regulators had been discovered in certain plants. The result is that many substances or groups of substances have been ascribed roles as endogenous regulators in plants. The most commonly discussed examples are the polyamines (Evans and Malmberg, 1989; Bagni and Torrigiani, 1992), oligosaccharines (Albersheim, 1985), acetyl choline (Tretyn *et al.*, 1990) and the jasmonates (Pathier *et al.*, 1992). However, claims were also made for a much larger number of compounds of more much limited taxonomic distribution. Gross (1975) reviewed over 100 such substances that were considered to play a role as endogenous regulators, these included representatives of aliphatic and aromatic carboxylic acids, phenols, alkaloids, terpenes and S- and N-heterocyclic compounds. However, a satisfactory evolutionary explanation to explain why different species use different chemicals to control the same basic physiological process does not yet exist. Why should plant X use compound A to control flowering when plant Y uses compound B? Are the links between physiology and biochemistry in plants really anarchic? An answer to this question might be formed by considering the links between secondary metabolism and plant 'hormones'.

The link between secondary metabolism and hormonal control

Are plant hormones 'secondary metabolites'?

It is known that chemical communication is important in some simple organisms as a way of coordinating sexual reproduction. For example, in the water mold *Achlya*, the terpenoid compounds oogoniol or antheridiol are used to coordinate the sexual reproduction of

Table 4.1 The major groups of plant growth substance and their discovery

| <i>Hormone group</i> | <i>Biosynthetic pathway or precursor</i> | <i>Discovery</i> |
|-----------------------------------|--|---|
| Indolyl-3-acetic acid (IAA) auxin | Tryptophan (?) | First isolated in the 1930s during a search for auxin activity from human urine, <i>Rhizopus</i> and <i>Saccharomyces</i> cultures. At that time plants were thought to contain a cyclopentane auxin (<i>auxin a</i> – now known not to exist). IAA identified in corn seed extracts in 1946 and widely reported in other plant extracts subsequently. |
| Gibberellins | Isoprenoid | First isolated from fungal cultures by Japanese phytopathologists investigating a disease of rice (1926). Several related compounds were subsequently found in fungal cultures. Gibberellins were not isolated from plants until the 1950s, some decades after some of their effects on plant growth had been described. |
| Abscissic acid | Isoprenoid | Isolated and characterized from abscising cotton bolls and dormant tree buds in 1963. No longer thought to play an important role in abscission or bud dormancy but good evidence for a role in controlling stomatal aperture and seed dormancy. |
| Cytokinin | Isoprenoid and purine | Searching for a substance capable of promoting cell division in cell and tissue cultures, coconut milk, malt extract, yeast extract and autoclaved herring sperm DNA were found to be active. A purine was isolated from the latter source by the mid-1950s. A decade elapsed before a related compound was isolated from maize endosperm. |
| Ethylene | Methionine | As a constituent of town gas, ethylene was known to have a potent effect on plants since 1906 and this compound was isolated from ripening apples in 1935. Some physiologists did not accept ethylene as a true plant growth substance until well into the 1960s, despite its widespread occurrence and high biological activity. |

All the major hormones were either first isolated from unusual sources or were discovered as a result of the study of a physiological process in which the hormone now plays a disputed role.

colonies and in some simple fungi, trisporic acid (C15 isoprenoid) is used as a sporulation coordinator (Gooday and Adams, 1993). Thus the roots of plant hormonal control may lie in simple organisms communicating between cells of the same species in *different* individuals. With the evolution of true multicellular organisms, the need for coordinated development and functioning would have extended the scope for chemical signalling and using chemicals to coordinate the functioning or fate of cells within the *same* individual would be a small step. The extent and way in which such chemical signalling would have evolved would have depended on the nature of the specialized functions that appeared in various types of organism over evolutionary time. It can be postulated that in all early multicellular organisms, secondary metabolites, already selected for the capacity to generate compounds with potent biomolecular activity were put to a new role. Evidence for such a scenario can be found in the multiple roles played by members of the isoprenoid pathway. Individual isoprenoids function as plant growth substances, plant defence compounds, fungal sexual coordinators, animal hormones and animal olfactory attractants and repellents. Of the plant growth substances, three are derived fully or partly from the isoprenoid pathway – the gibberellins, abscissic acid and cytokinins (see Table 4.1). The other two plant hormones, auxin and ethylene, are derived from amino-acid precursors and amino acids also serve as the precursors for some chemical regulators in animals and play a part in plant defences, possibly because amino acids have commonly been used as precursors in ‘inventive biosynthesis’ (Wong, 1981). It is interesting that the exact route leading to the biosynthesis of IAA has been debated for nearly half a century, with the expectation that the biosynthesis of such an important molecule would be finely controlled. Traditional biochemical investigations could not provide definitive evidence for such a pathway, but unexpected phenotypes resulting from the manipulation of cytochrome P450 genes suggests that, in some species, there may be an intimate link between the synthesis of indole glucosinolates and the synthesis of IAA (Feldmann, 2001). IAA could be considered to be a ‘secondary product’ with a primary role. If the evolutionary recruitment of a ‘secondary metabolite’ to serve a role as an endogenous coordinator or regulator in plants has occurred, it is likely that the event will have brought to the emerging ‘hormonal control’ a series of metabolic traits that have evolved to serve a quite different purpose – the generation and retention of chemical diversity. If there is not a duplication of all the enzymes involved in the pathway leading to the plant hormone, the plant hormonal control will inevitably be somewhat compromised by the metabolic features of a pathway which is even more multifunctional in that it now includes a role as an internal signalling molecule. Could such multifunctionality explain some previously puzzling aspects of plant hormone biosynthesis? Maybe this multifunctionality explains why so many stimuli (from insects and fungi to many forms of physical stimuli such as various wavelengths of light, low and high temperatures, too little or too much water, etc.) can change the hormone content of plants (Davies, 1995).

Gibberellin synthesis – generating diversity?

Over 100 different gibberellins have been isolated and characterized – some are found in many species but others have a much more limited taxonomic distribution. A single plant species usually contains several gibberellins. The great majority of gibberellins do not possess high biological activity (or possess activity only because they are converted to other more active gibberellins) (Davies, 1995). The fact that the great majority of gibberellins possess low activity is itself consistent with the Screening Hypothesis, but is harder for other models of secondary metabolite evolution to explain. But why do plants (and some

fungi) make so many gibberellins? We would argue that this group of plant hormones are showing their evolutionary origin as secondary products. Even when there is a need for a gibberellin to act as an essential regulator in the plant, the metabolic traits that generate chemical diversity are retained (or at least are not selected against). The particular trait of relevance to gibberellin biosynthesis is the proposal that some enzymes involved in secondary metabolite biosynthesis may possess a relatively broad substrate specificity, leading to matrix grid transformations. In such matrix grid biosynthetic routes, a few enzymes could add or transform substituents to a carbon skeleton and the order in which they are added or transformed is not fixed. Hence a number of intermediates can be generated. As long as the possession of at least one of the compounds made results in a net gain in plant fitness, the chemical diversity represented in all the intermediates can be retained. Evidence in support of this prediction can be found in the study of the gibberellin 20-oxidase which can convert two precursors into at least 11 products because it is multifunctional (Lange *et al.*, 1994). The fact that some parts of the biosynthetic pathway leading to the active gibberellin, GA₁, can operate as a matrix (Taiz and Zeiger, 1991) would be evidence that this type of mechanism does operate in plants even in a pathway used to make compounds which are central to controlling plant growth and development and that some enzymes involved in the pathway are following rules that would allow the extension and retention of chemical diversity. A similar complex metabolic grid may also exist in the biosynthetic pathways leading to brassinosteroids (Wang and Chory, 2000).

Plant hormone degradation – another role?

For many decades it has been argued by some that enzymes involved in metabolizing plant hormones (to give either breakdown products or conjugates) may play an important role in plant hormone homeostasis (Bandurski *et al.*, 1992). The control of hormone concentration by controlling the breakdown, rather than the synthesis, seemed counter intuitive to those schooled in hormonal control in mammals. However, if the biosynthetic machinery leading to plant hormone production carries with it some of the flexibility (and maybe the inducibility) of a metabolism evolved for multifunctionality, and where precise control of the amount of product may not be something that has been highly selected for, the evolution of other means of controlling the amount of any hormone by degradation perhaps deserves attention. However, an alternative explanation could be that, in some cells (maybe cells that are insensitive themselves to hormones), an ability to generate and retain chemical diversity has resulted in a production of hormones and it is in these cells that selection has resulted in a fairly crude method (degradation) of hormone concentration regulation so as to avoid disturbing more carefully regulated hormone levels in other cells. However, if these cells retain a capacity to make hormones and genes can be induced in them, possibly the extra burst of hormone synthesis could give rise to the hormone level changes that are sometimes associated with insect or fungal attack. This may be yet another opportunity for cross talk?

Summary

The aim of this chapter has not been to inform readers how plants work or indeed how plants have evolved. Rather it has been to try to engage readers in at least considering whether there might be an appropriate evolutionary framework for metabolism that could help us investigate and eventually understand many physiological processes. We have indulged in

speculation in order to provoke readers and some may find the lack of certainty unsatisfactory. Readers provoked into rejecting the evolutionary model that has been advanced should feel free to make such a rejection ... as long as they have a better model with which to replace it. After any discovery, the question 'why do plants do that?' needs some evolutionary explanation. Maybe any plant physiology or plant biochemistry textbook without a section on evolution should be regarded as seriously incomplete.

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5

Did auxin play a crucial role in the evolution of novel body plans during the Late Silurian–Early Devonian radiation of land plants?

Todd J Cooke, DorothyBelle Poli and Jerry D Cohen

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Introduction

The overall objective of this chapter is to identify plausible developmental mechanisms that might have contributed to the rapid diversification of early land plant lineages during the Late Silurian to Early Devonian Periods (Kenrick and Crane, 1997). An even more rapid diversification of bilateral animal lineages seems to have occurred during the well-known

Cambrian radiation (Gould, 1989; Raff, 1996; Conway Morris, 2000). Given that the lineages that ultimately gave rise to animals and to plants are most likely to have diverged as ancient lineages of unicellular flagellates (Baldauf *et al.*, 2000), the origins of early land plants occurred via different macroevolutionary processes than those operating in ancient animals (Meyerowitz, 2002). Nonetheless, as an archetypal example of organismal radiation, our current understanding of the Cambrian radiation can be used as a conceptual framework for considering the analogous radiation of early land plants.

Brief overview of Cambrian radiation of bilateral animals

Rapid diversification of animal phyla

Although well-preserved fossils of the soft-bodied Ediacaran fauna are widely distributed in Late Precambrian rocks, it has been very problematical to trace the evolution of the simple features of Ediacaran fossils into the more complex body plans of Cambrian metazoa (Knoll and Carroll, 1999; Conway Morris, 2000). Therefore, palaeontologists have focused on the Cambrian radiation that resulted in the rapid (or 'explosive' with respect to geological time) appearance of the crown phyla of bilateral animals, including molluscs, arthropods, echinoderms and chordates, around 550–530 million years ago (Ma) (Raff, 1996; Erwin *et al.*, 1997; Valentine *et al.*, 1999; Conway Morris, 2000). In marked contrast to the fossil record, molecular clock estimates predict that initial divergences of these lineages occurred much earlier in the Precambrian era (Bromham *et al.*, 1998; Heckman *et al.*, 2001). Even if a long Precambrian fuse was needed to ignite the Cambrian explosion (Fortey, 2001), it is still undeniable that earliest bilateral animals diversified into the crown metazoan lineages during a rather narrow slice of deep geological time, with the result that all extant animal phyla appeared before the end of the Cambrian. Some authors attribute the Cambrian radiation to the occurrence of large-scale environmental perturbations resulting in new adaptive landscapes at the Precambrian-Cambrian boundary (Knoll and Carroll, 1999), while others emphasize the possible role of new ecological selection pressures associated with the rise of filter feeding and macroscopic predation (Conway Morris, 2000). Of particular relevance to this chapter are the putative developmental mechanisms discussed below that may be responsible for generating new phylum-specific body plans in the presence of those selective pressures.

Characteristic body plan of each phylum

Traditionally, animal biologists have recognized different phyla on the basis of their fundamental *Baupläne* (or body plans in English), which encapsulate the nature and organization of tissues and organs within the animal body. For example, the arthropods comprise a diverse assemblage of bilateral organisms, including horseshoe crabs, spiders, millipedes, crustaceans and insects. Yet the arthropod phylum as a whole exhibits a common body plan with such unifying features as a segmented exoskeleton, jointed legs, ventral nerve cord and dorsal heart with an open circulatory system. Interestingly, the characteristic body plans of all extant animal phyla were already expressed in their marine ancestors by the end of the Cambrian radiation (Raff, 1996). While certain phyla underwent major structural innovations accompanying their post-Cambrian invasion of the land environment, these terrestrial adaptations occurred within the conserved patterns of their pre-existing body plans.

Early establishment of body plan

For two related reasons, until the recent innovations in molecular analyses, phylogenetic schemes for animal phyla were typically based on the comparative morphology of embryos and larvae. (1) the basic body plan of each phylum is first expressed at these early stages of animal development (Gilbert, 2000). Ever since von Baer (1828), it has been recognized that the general features of each phylum appear in embryonic and/or early postembryonic development as opposed to the more specialized features of individual classes that develop at later stages. Indeed, it is the common embryonic or larval features that disclose the close evolutionary relationships within those phyla like molluscs or arthropods with divergent adult forms. Furthermore, the common body plan of vertebrate embryos makes them also appear remarkably similar even though the specialized features of the adults are distinctly different. (2) the structures developing in the early, so-called phylotypic stage are more evolutionarily conserved than those formed at later stages (Raff, 1996). For example, in insects, the germ band or early larval stages have a segmented worm-like appearance that resembles the body plans of other taxa, such as the onychophorans, related to the arthropods (Ballard *et al.*, 1992). As another intriguing example, living echinoderms develop free-swimming bilateral larvae that undergo a complicated metamorphosis to become adults exhibiting 5-parted radial symmetry. Peterson and Davidson (2000) have hypothesized that ancestral stem-groups of bilateral animals developed simple larva similar to those characteristic of extant echinoderms. These larvae are then envisioned to have served as the structural platform for the activity of 'set-aside cells' required to generate the complex body plans of different crown bilateral animals.

Altered expression of embryonic genes resulting in new body plans

The basic approach in evolutionary developmental biology ('evo-devo') is to study the genetic regulation of embryonic and larval development of extant organisms belonging to various lineages in order to elucidate the developmental mechanisms responsible for the origin and/or diversification of those lineages (Raff, 1996; Gilbert, 2000; Arthur, 2002). This research is based on the reasonable, albeit untestable, assumption that genetic regulation of embryonic development is extraordinarily stable over vast geological time. The justification for this assumption is that any disruption in embryonic regulation should result in an avalanche of disruptive, and inevitably fatal, consequences for postembryonic development (Raff, 1996). Similar arguments have been also put forth to explain the failure of any new metazoan body plans to evolve following the Cambrian radiation (Gilbert, 2000).

The paradigmatic case of genetic regulation of metazoan body plan involves a conserved group of homeobox (*Hox*) genes (Erwin *et al.*, 1997; Gellon and McGinnis, 1998; Valentine *et al.*, 1999; Carroll *et al.*, 2001). *Hox* genes play crucial regulatory roles in various aspects of metazoan axis specification, such as external segment identity in arthropods and internal body segmentation in vertebrates. Molecular analyses of *Hox* gene diversity have shown that these genes were gradually duplicated during the ancient evolution of non-bilateral animals, with the result that the stem group of bilateral animals appear to have evolved a fully-fledged cluster consisting of a minimum of seven *Hox* genes (de Rosa *et al.*, 1999; Peterson and Davidson, 2000). Some workers have proposed that the *Hox* genes were responsible for encoding the original anterior-posterior axis in the common ancestor of all animals (e.g. Slack *et al.*, 1993), while others argue that the *Hox* cluster was recruited for pattern formation of the pre-existing anterior-posterior axis during the evolution of the stem group of bilateral animals (e.g. Peterson and Davidson, 2000). In either

event, it appears that the evolution of different body plans in the lineages of bilateral animals did not depend on the evolution of new *Hox* clusters or other developmental master genes. Instead, the diversification of bilateral animals required various innovations in regulatory networks controlling the expression of these genes (Gellon and McGinnis, 1998; Knoll and Carroll, 1999; Carroll *et al.*, 2001). The molecular palette of developmental genes was already present in primordial bilateral animals before these genes were rewired to sculpt novel bilateral body plans.

In summary, it seems reasonable to conclude from this brief description that the Cambrian radiation of bilateral animals can be viewed as having arisen from the rapid emergence of new expression patterns of pre-existing developmental regulatory genes. These altered expression patterns resulted in a wide range of new body plans for bilateral animals. These new animals were presumably subjected in turn to ecological selection pressures that favoured those animals with body plans best adapted for surviving in Cambrian oceans.

Silurian–Devonian radiation of land plants

Using the conceptual framework derived from our consideration of the Cambrian radiation of bilateral animals, we intend here to discuss plausible developmental mechanisms underlying the evolution of novel plant body plans and hence the macroevolution of different plant lineages in the Late Silurian to Middle Devonian periods. The hormone auxin appears to regulate the organizational features comprising body plans of contemporary plants; therefore, particular attention is granted to the hypothesis that evolutionary changes in auxin action were causally involved in the generation of different body plans during the radiation of early vascular plants.

Did early land plants diverge in a rapid evolutionary radiation?

Recent work has provided cogent molecular, biochemical and cellular evidence that ancient charophycean green algae (also called charophytes), whose living descendants include the orders Zygnematales, Coleochaetales and Charales, represent the primordial group giving rise to land plants (Graham, 1993; Graham *et al.*, 2000; Karol *et al.*, 2001). Living charophytes, sometimes derided as ‘pond scum’, have haplobiontic life cycles, with a dominant haploid gametophyte and a diploid phase solely consisting of the zygote that undergoes meiosis to produce four haploid cells. These algae exhibit a diverse range of growth forms, including unicells, tip-growing filaments, margin-expanding discs and complex shoot-like axes consisting of alternating large multinucleate internodal cells and multicellular nodes bearing lateral branches. This diversity of growth forms in the charophytes is thought to reflect the more permissive nature of aquatic habitats (Niklas, 2000).

According to molecular clock estimates, ancient charophycean green algae are predicted to have invaded the land during the Precambrian around 600 Ma (Heckman *et al.*, 2001). Dating from the Middle Ordovician around 470 Ma, the oldest microfossils, which are considered to have originated from genuine land plants, are obligate spore tetrads with sporopollenin-impregnated walls, imperforate cuticles and narrow tubes (Gray, 1985; Edwards and Wellman, 2001; Graham and Gray, 2001). Indeed, the ability of these first land plants successfully to colonize terrestrial environments is often attributed to the desiccation-resistant coverings observed on these microfossils. This microfossil evidence suggests that the earliest land plants are likely to have exhibited a bryophyte-grade of structural organization,

at least with respect to spore morphology (Gray, 1985; Edwards and Wellman, 2001; Graham and Gray, 2001). Available molecular sequence information is also consistent with the perspective that the three extant bryophyte lineages diverged earlier than the monophyletic lineage giving rise to the vascular plants (Qiu *et al.*, 1998; Nickrent *et al.*, 2000; Karol *et al.*, 2001).

It is sometimes postulated that the first land plants had quickly evolved the embryo, via the intercalation of mitotic divisions of the zygote prior to the occurrence of sporic meiosis (Graham and Wilcox, 2000). However, the lack of any confirming fossil evidence makes it conceivable that land plants did not develop embryos until long after the successful colonization of terrestrial habitats (Niklas, 1997). The earliest mesofossils with possible bryophyte affinities have been identified as miniature branching axes in Lower Devonian rocks (Edwards *et al.*, 1995; Edwards, 2000; Edwards and Axe, 2000). However, very few macrofossils, including *Sporogonites*, *Tortilicaulis* and *Pallaviciniites* from Devonian strata, appear to exhibit the characteristics of dorsiventral thalli and/or monosporangiate sporophytes that disclose their likely affinities to extant bryophytes (Taylor and Taylor, 1993). Thoughtful discussions about the inadequacy of the fossil record for elucidating the major events in bryophyte evolution can be found in many sources (e.g. Kenrick and Crane, 1997; Niklas, 1997; Edwards, 2000; Kenrick, 2000).

It is irrefutable that a considerable time interval exists between the first putative land plant spores in mid-Ordovician strata and the first protracheophyte *Cooksonia* in Upper Silurian rocks. In marked contrast to the bryophyte enigma, the fossil record for vascular plants provides compelling evidence that the appearance of *Cooksonia* was followed by the rapid radiation of numerous vascular plant lineages including those lineages that evolved into extant lycophytes, ferns and horsetails (Taylor and Taylor, 1993; Kenrick and Crane, 1997; Niklas, 1997). This radiation of vascular plants is dated to have occurred in the Late Silurian (424–409 Ma) according to cladistic analysis (Kenrick and Crane, 1997: Figure 7.15) or in the Early to Middle Devonian (409–381 Ma) according to macrofossil stratigraphy (Taylor and Taylor, 1993; Kenrick and Crane, 1997). The rapid diversification of early vascular plant lineages culminated with the origin of the progymnosperms (i.e. the progenitor lineage for seed plants) in the Middle Devonian, as indicated by the fossil *Svalbardia* (Taylor and Taylor, 1993; Berry and Fairon-Demaret, 2001).

The fossil record indicates that the Late Silurian to Middle Devonian was indeed the most innovative interval in the morphological diversification of land plants. During this interval, land plants appear to have evolved many features often assumed to serve as crucial terrestrial adaptations, including multicellular primary and secondary meristems, vascular tissues, vegetative organs (roots, stems, and leaves), and sporangium-bearing organs (sporophylls and sporangiophores) (Graham, 1993; Taylor and Taylor, 1993; Kenrick and Crane, 1997). Of particular interest to this chapter, it is often presumed that the first sporophyte consisted of a spherical embryo that was directly modified to form a simple sporangium embedded in the archegonium (Niklas, 1997; Graham and Wilcox, 2000). Its evolutionary elaboration into the complex vascular plant sporophyte with elevated sporangium-bearing axes must have been a critical adaptation for flourishing in terrestrial environments, because this plant body is well-constructed to produce numerous meiospores for effective aerial dispersal.

In parallel to the Cambrian radiation of bilateral animals, the rapid diversification of vascular plants starting in the Late Silurian hints at the possibility that the ancestral stem group for all vascular plants had just previously experienced a limited number of genetic innovations, which permitted and/or facilitated the rapid origin of new body plans. Contrary

to the selection pressures operating on the first bilateral animals in Cambrian oceans, it is reasonable to propose that during the Silurian–Devonian radiation of land plants, natural selection acted to favour those body plans that enabled these plants to survive in terrestrial environments. The following sections will address the developmental mechanisms responsible for generating the body plans of land plants.

Are the characteristic body plans of land plants established during embryonic development?

Although not many plant biologists make explicit reference to the concept of body plan, its occasional appearance in the botanical literature has led to some confusion because it has been applied to different levels of plant organization. For instance, those plant biologists focusing on algal life forms have referred to different types of cellular organization, such as unicellular, siphonous, colonial and multicellular, as representing the basic body plans of photosynthetic eukaryotes (e.g. Niklas, 2000). The present chapter adopts an alternative perspective from the discipline of plant morphology, where the body plan concept, when used, is generally restricted to vascular plant sporophytes (e.g. Troll, 1943; Groff and Kaplan, 1988). The multicellular sporophyte of each land plant division can be said to exhibit a characteristic body plan that is based on such features as meristem organization, vascular tissue arrangement, positional relationships among vegetative organs and positional relationships of reproductive structures on vegetative organs (Bold *et al.*, 1987; Gifford and Foster, 1989). In this section, the sporophytes from representative divisions of extant land plants are illustrated in order to evaluate whether or not the embryo or the young post-embryonic plant expresses the fundamental body plan of each division.

As an example of the bryophyte grade of plant organization, the mature sporophyte of liverwort *Marchantia polymorpha* L. consists of three parts: absorptive foot, elongated seta and spore-producing capsule (Figure 5.1A–D) (Smith, 1995; Bold *et al.*, 1987; Crum, 2001). This sporophyte is initiated as a zygote that develops into a small spherical embryo inside the archegonium (Figure 5.1B). Then the spherical embryo differentiates into three regions in which different cell shapes reveal the ultimate fate of each region (Figure 5.1C). The small basal cells destined to form the foot are generally oriented perpendicular to the future growth axis. The presumptive seta is composed of enlarged isodiametrical cells, whereas the immature cells of the capsule are greatly elongated and oriented parallel to the future axis. Although each region is capable of a limited number of additional cell divisions, the embryo does not develop any localized region of cell division recognizable as a genuine meristem. Instead, the embryo expands and differentiates into the mature sporophyte illustrated in Figure 5.1D, which displays the typical foot and capsule found in almost all but a few semi-aquatic liverworts. The *Marchantia* seta is similar to other liverwort setae in that it elongates via diffuse growth over its entire length, but it is considerably shorter. This reduced length is attributed to *Marchantia* sporophyte developing suspended from an elevated archegoniophore, which does not arise on most other liverwort thalli. Because liverwort sporophytes never develop an apical meristem capable of producing additional organs, they do not exhibit the modular organization of reiterated units that is characteristic of vascular plant sporophytes.

In essence, the liverwort sporophyte can thus be said to exhibit a tripartite body plan that is first expressed in the young embryo. Hornworts and mosses, the other two bryophyte divisions, also exhibit tripartite body plans; however, the cellular activities associated with axial growth are distinctly different among bryophyte divisions (Smith, 1955; Wardlaw, 1955;

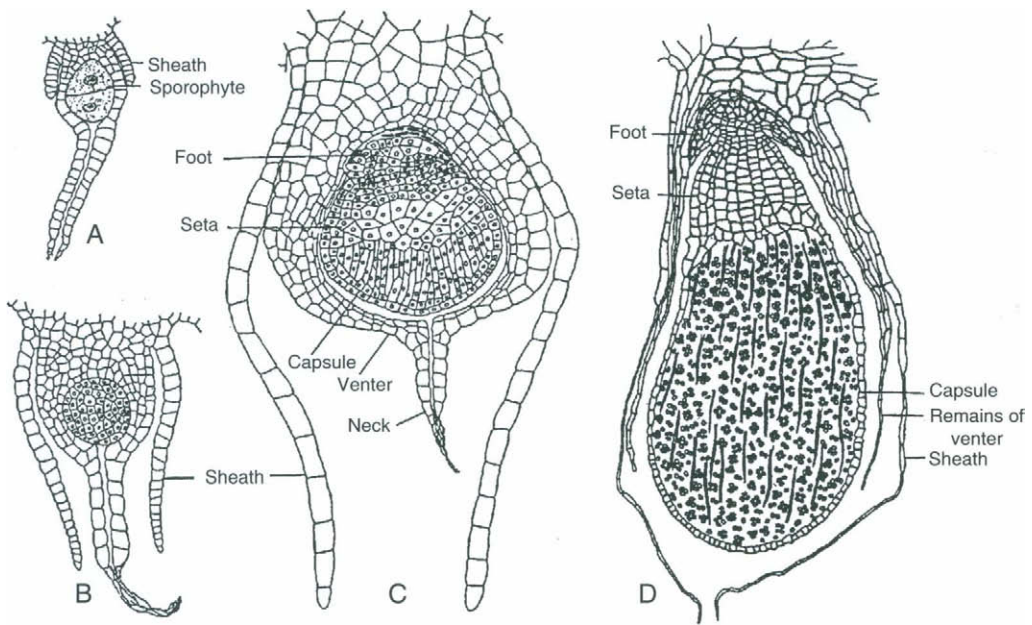


Figure 5.1 The development of the embryo and sporophyte of the liverwort *Marchantia polymorpha* L. (A) Two-celled embryo inside the archegonium. (B) Young spherical embryo lacking obvious cellular differentiation. (C) Expanded spherical embryo exhibiting tripartite organization. (D) Nearly mature sporophyte with well-defined foot, seta and capsule. Redrawn from Smith (1955) with permission from McGraw-Hill Companies.

Bold *et al.*, 1987; Crum, 2001). The embryo of the hornworts is also composed of a basal tier destined to become the foot and an apical tier representing the future apicalmost cells of the capsule (Campbell, 1918; Smith, 1955; Wardlaw, 1955; Crum, 2001). The intermediate tier develops into a narrow band of dividing cells called an intercalary meristem that remains positioned just above the foot. The intercalary meristem undergoes unifacial divisions on its capsule side, with the result that these new cells compose almost the entire linear capsule that rises above the gametophyte. Thus, the three-tiered hornwort embryo is directly enlarged into a tripartite body plan. In most mosses, it is difficult to recognize three cellular tiers in the first stages of embryonic development (Smith, 1955; Wardlaw, 1955; Lal and Bhandari, 1968; Bold *et al.*, 1987; Crum, 2001). A transient apical cell differentiates into the upper hemisphere and then this apical cell and its derivatives act to generate a spindle-shaped embryo. This embryo exhibits three well-defined regions that are destined to develop into the foot, seta and capsule of the mature moss sporophyte. In marked contrast to the diffuse growth of the liverwort seta, a unifacial intercalary meristem arises in the moss seta just beneath the immature capsule and its activity generates most of the cells composing the elongating seta.

The fern *Pityrogramma triangularis* (Kaulf.) Maxon serves here as a representative example of the pteridophyte grade of plant organization (Figure 5.2). The fern zygote within the archegonium undergoes a series of segmentation divisions resulting in the formation of a globular-shaped embryo (Campbell, 1918; Smith, 1955; Wardlaw, 1955; Gifford and Foster, 1989). This embryo is often said to be composed of four quadrants representing the shoot apex, first leaf, first root and foot (Figure 5.2A–D). The shoot apex is solely responsible

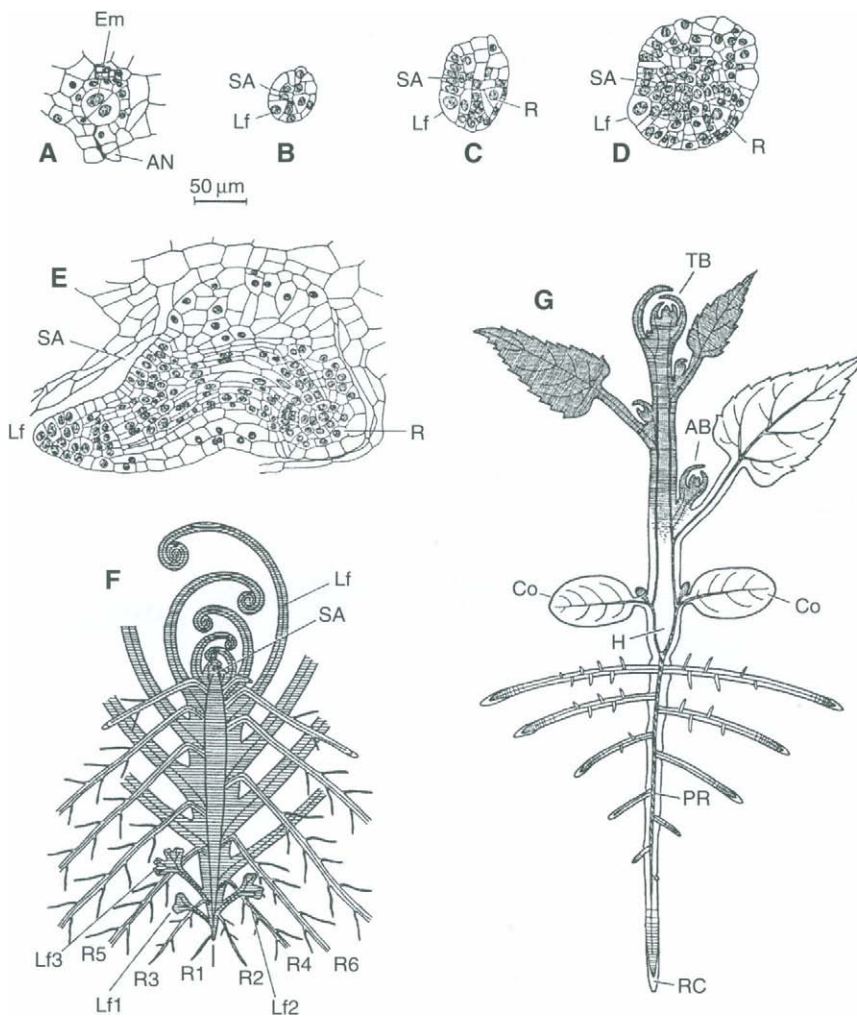


Figure 5.2 Embryonic development of the fern *Pityrogramma triangularis* (Kaulf.) Maxon (A–E) and the body plans of the ferns (F) and the dicots (G). (A). Two-celled embryo inside the archegonium. (B) Young spherical embryo exhibiting characteristic apical cells for shoot apex and first leaf in apical hemisphere. (C–E) Successive stages of the four-quadrant embryo composed of shoot apex, first leaf, first root and foot (unlabelled). The first root arises in a lateral position near the base of the first leaf. Panels (A–E) were redrawn with permission from unpublished work of W. Hagemann. (F) Diagrammatic illustration of typical fern body plan, which shows that the positional relationships first expressed in the unipolar embryo are reiterated in postembryonic development. (G) Diagrammatic illustration of the model dicot body plan, which shows that the positional relationships first expressed in the bipolar embryo are reiterated in postembryonic development. This situation does not hold true for all dicots. Panels (F) and (G) were redrawn from Troll (1959) with permission of Georg Thieme Verlag. Abbreviations: Em, embryo; AN, archegonial neck; SA, shoot apex; Lf, leaf; R, root; TB, terminal bud; AB, axillary bud; Co, cotyledon; PR, primary root; RC, root cap.

for generating the growth axis of the postembryonic plant; hence, the fern embryo is referred to as being unipolar (Groff and Kaplan, 1988). The first root arises near the base of first leaf (Figure 5.2E). All postembryonic roots are also observed to originate near leaf bases and thus, fern roots are consistently lateral with respect to the longitudinal axis of

the growing plant (Figure 5.2F), which has been termed the homorhizic condition (Troll, 1943; Groff and Kaplan, 1988). The horsetails (*Equisetum*), which are thought to comprise a monophyletic group with the ferns (Pryer *et al.*, 2001), exhibit similar organographic arrangements, with the first root subtending the first leaf and all postembryonic roots arising at the nodes (Gifford and Foster, 1989).

The relative positions of embryonic organs illustrated in Figure 5.2 foreshadow the postembryonic body plan in almost all ferns (Eames, 1936; Smith, 1955; Bierhorst, 1971; Gifford and Foster, 1989). In even those ferns exhibiting unusual habits, the body plan is irrevocably fixed by its embryonic organization. In marked contrast to the subterranean root systems of dicot trees, the tree ferns from such genera as *Dicksonia* and *Cyathea* develop buttress-like coverings of interwoven roots that arise at the bases of lower leaves and grow down the outside of the trunk and into the ground (Smith, 1955; Gifford and Foster, 1989). Additional roots continue to form at the bases of higher tree fern leaves, even though these roots are destined to remain short and never penetrate the soil. Mature plants of the aquatic floating fern *Salvinia* do not develop roots and this rootless feature can be traced back to its embryo, where the non-growing sector subjacent to the first leaf is often interpreted as a vestigial first root (Eames, 1936; Bierhorst, 1971). Even the problematic whisk ferns (Psilotales), which are apparently closely related to eusporangiate ferns according to molecular analyses (Pryer *et al.*, 2001), exhibit an embryonic organization that correlates with its mature morphology, at least with respect to the absence of any roots. In *Tmesipteris tannensis* Bernh. and *Psilotum nudum* (L.) Pal. Beauv., the embryos display a two-parted organization consisting of a distal shoot apex and a proximal foot with no evidence at all for leaf or root quadrants (Holloway, 1921, 1939; Bierhorst, 1971). The embryonic shoot apex develops into a branched plagiotropic rhizome lacking roots, which may be an adaptive response to the epiphytic habit frequently adopted by whisk ferns. (Eventually, the rhizome produces aerial axes bearing reduced dorsiventral structures called enations which are actually homologous to genuine leaves (for resolution of this controversial issue, see Kaplan, 2001).) A rare exception to the general rule about the embryonic encapsulation of fern body plan is seen in certain *Ophioglossum* species where roots develop subterminal buds capable of growing into new shoots (Peterson, 1970); nevertheless, the roots of the daughter shoots also emerge at the base of their fronds, thereby replicating the positional relationships of the original shoot.

The other major pteridophyte group, the lycophytes, also manifests unipolar embryonic organization, which is largely reproduced in the body plans of adult plants (Groff and Kaplan, 1988; Gifford and Foster, 1989). However, postembryonic roots in the lycophytes do not exhibit the same positional relationships as the ephemeral first root. For example, the embryos of different *Selaginella* species exhibit wide variation in the position of the first root relative to other embryonic organs, which are not reflected in the organographic arrangements of their postembryonic plant bodies (Bower, 1935; Gifford and Foster, 1989). Subsequent roots in many *Selaginella* species originate from leafless axes called rhizophores that form at stem bifurcations. The distal ends of these rhizophores bear typical roots with root caps. Some workers suggest that rhizophores are considered as true roots (e.g. Gifford and Foster, 1989), while others view them as unique root-bearing structures (e.g. Lu and Jernstedt, 1996). A similar uncoupling of embryonic root position and postembryonic organization is observed in other lycophytes. In most *Lycopodium* species, postembryonic roots originate close to the shoot tip and traverse down the cortex before emerging into the soil (Gifford and Foster, 1989). The postembryonic roots of *Isoetes* and its extinct relatives are borne on different specialized structures called rhizomorphs of uncertain homology

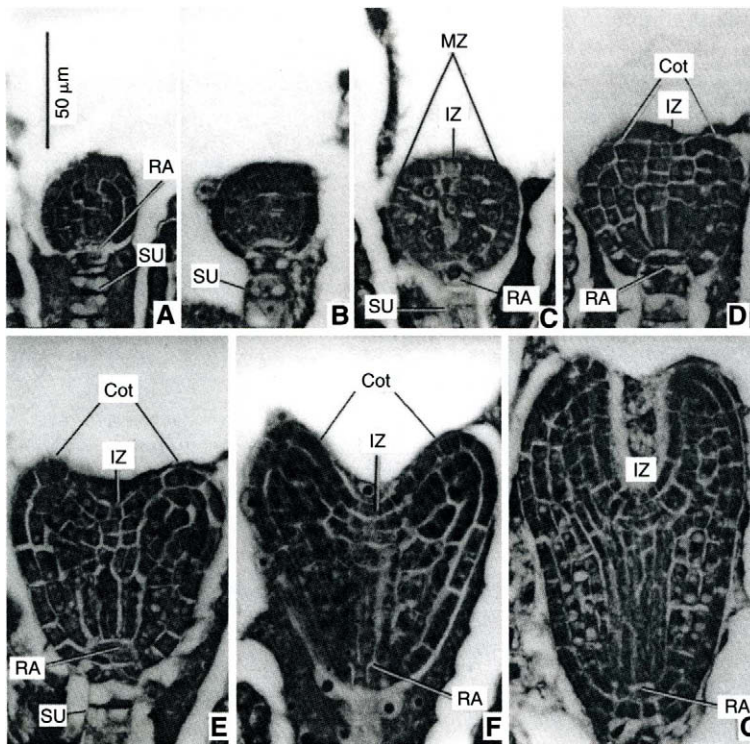


Figure 5.3 Embryonic development of the dicot *Capsella bursa-pastoris* (L.) Medic. (A) and (B) Young embryos exhibiting well-defined suspensor and embryo proper. The incipient root apical meristem can be recognized by the periclinal cell divisions delimiting the future root cap. (C) Mid-globular embryo displaying incipient shoot apical meristem with a lighter staining central (or initial) zone and a darker staining lateral (or morphogenetic) zone. (D–F) Early to late heart embryos showing cotyledon emergence from the lateral zones. (G) Early torpedo stage. Reprinted from Kaplan and Cooke (1997) with permission from American Society of Plant Biologists. Abbreviations: RA, root apex; SU, suspensor; MZ, morphogenetic zone; IZ, initial zone; Cot, cotyledon.

(Paolillo, 1963, 1982). In conclusion, it appears that the fundamental pteridophyte body plan is fully manifested in either developing embryos (ferns and horsetails) or young post-embryonic plants (lycophytes) and then it is reiterated throughout the life of the plant.

Characteristically, seed plant embryos exhibit bipolar (or allorhizic) organization with the embryonic shoot and root apices arising at opposite poles (Troll, 1943; Groff and Kaplan, 1988). In many, but not all, embryos, these incipient meristems act to perpetuate the bipolar organization, with the result that the primary plant body exhibits a central axis with opposite shoot and root systems, as is diagrammatically illustrated in Figure 5.2G. For example, in the dicot *Capsella bursa-pastoris* (L.) Medic. (Figure 5.3), the incipient root and shoot apical meristems arise during the globular stage of embryonic development (Kaplan and Cooke, 1997). These meristems can first be recognized by the cellular activities that accompany the formation of their distal or lateral structures. In particular, the origin of the root apical meristem is disclosed by periclinal divisions that delimit the distal root cap from the more proximal root body (Figure 5.3A–D). This first root will then generate the root system of the mature plant. The embryonic shoot apical meristem is revealed by the presence of cytohistological zonation at the shoot pole of the late globular embryo (Figure 5.3C–G).

The dark-staining lateral regions of this incipient meristem are clearly distinguishable from the light-staining central region. The lateral regions give rise to two cotyledons, which are homologous to the first leaf of pteridophyte embryos. The central region is destined to become the epicotylar shoot apical meristem that is ultimately responsible for generating the entire shoot system. Thus, the bipolar organization of the *Capsella* embryo establishes the positional relationships that are expressed throughout postembryonic development.

Although Figure 5.2G was proposed as the model for the bipolar condition in dicots (Troll, 1943), it is clear that the body plans of individual seed plants may represent either direct reiterations or modified arrangements of the original embryonic organization. The dicots in particular exhibit the greatest variation in the relationship between embryonic organization and mature body plan (Groff and Kaplan, 1988). Although many dicots maintain the bipolar embryonic organization throughout postembryonic development, the embryonic organization of numerous other species is subsequently modified by: (1) the shoots bearing lateral roots (e.g. many vines like *Vitis vinifera* L. and *Hedera helix* L.); (2) the roots bearing new shoots (e.g. saprophytic plants like *Monotropa uniflora* L.); or (3) both shoot-borne roots and root-borne shoots (e.g. many perennial herbs and certain trees) (Groff and Kaplan, 1988). It can be argued that root-borne shoots are simply mimicking the original bipolar axis, with the new shoot apex at one pole and the bud-producing root at the other. However, the origin of lateral roots from the shoot does not reflect the embryonic organization but rather represents a postembryonic modification of the bipolar body plan.

The bipolar organization of gymnosperm embryos is characteristically maintained during postembryonic growth, with very few reported examples of root-borne shoots or vice versa (Groff and Kaplan, 1988). By contrast, resolving the nature of positional relationships in monocot embryos represents an extraordinary challenge that can only cursorily be addressed here. The traditional perspective is that the single cotyledon occupies the terminal position in the developing monocot embryo (Souèges, 1931; Gifford and Foster, 1989), which suggests that the shoot apex should be viewed as a lateral structure. However, the weight of morphological evidence argues that the monocot shoot apex does indeed arise in the terminal position (Haccius, 1952, 1960; Swamy and Laksmanan, 1962), but it is subsequently displaced into what appears to be a lateral position by the pronounced growth of its overarching cotyledon (for further discussion, see Gifford and Foster, 1989). As an additional complication, the orientation of the first root relative to the central axis shows considerable variability in monocot embryos. The first root of many monocots is positioned at the opposite embryonic pole, thereby displaying the allorhizic organization of the typical bipolar embryo (Troll, 1943). However, the first roots of certain monocots such as *Zea mays* L. (Randolph, 1936) arise at the opposite pole but much later in embryo development, which resembles the timing of the first root initiation in unipolar fern embryos, while the first roots of still other monocots such as *Aponogeton madagascariensis* Mirbel (Yamashita, 1976) are reported to originate as genuine lateral structures. Nevertheless, the embryonic first root of almost all monocots is short lived or else poorly developed so that their postembryonic root system is almost entirely derived from shoot-borne roots. Therefore, although the monocot embryo is generally interpreted to have bipolar organization, the postembryonic monocot body plan is regarded as being secondarily homorhizic, due to the origin of subsequent roots as shoot-borne organs (Troll, 1943).

This section has established the following generalizations: embryonic pattern is amplified to form the postembryonic body in bryophytes, or it is reiterated to generate the postembryonic body in vascular plants. The most noteworthy exception to these generalizations

is that in some vascular plants, the positional relationships of all roots except the embryonic root are established in the young postembryonic body.

What developmental mechanisms act to generate plant body plans?

It must be appreciated that the botanical research on this question is far less advanced than comparable research in animal development. However, the initial evidence reviewed below suggests that the hormone auxin (indole-3-acetic acid) acts as a critical developmental mechanism for generating the body plans of land plants.

As discussed in a previous section, the current theory for explaining the developmental mechanisms underlying the Cambrian radiation is that the genes responsible for regulating embryonic development in basal animals experienced repeated duplication and altered transcriptional regulation, with the result that these genes were able to specify more complex body plans. A similar working hypothesis is now being adopted by the emerging discipline of plant evolutionary developmental biology (for an excellent overview, see Cronk, 2001). For instance, plant homeobox genes can be classified into at least three subfamilies: *KNOTTED1*-like (*KNOX*) genes affecting meristematic cell fates; homeodomain-leucine zipper (*HD-ZIP*) genes regulating later developmental and physiological processes; and *GLABROUS2*-like genes specifying epidermal cell fates (Chan *et al.*, 1998; Williams, 1998; Bharathan *et al.*, 1999). The archetypal example of the first subfamily is the so-called *SHOOTMERISTEMLESS* gene (Barton and Poethig, 1993) that is expressed in the central region of the globular embryo of *Arabidopsis thaliana* (L.) Heynh. (Long *et al.*, 1996). The identical stage in the related *Capsella* embryo is illustrated in Figure 5.3C. Although the name of this gene implies that it is essential for the origin of the shoot apical meristem, an alternative interpretation is that it maintains the proliferative activity of the embryonic central region in order to generate the epicotylar shoot apical meristem (Kaplan and Cooke, 1997). The recent report of two *KNOX*-like genes in the moss *Physcomitrella patens* (Hedw.) B.S.G. indicates that this subfamily started diverging early in the evolution of land plants (Champagne and Ashton, 2001). The ability of *Physcomitrella* to perform homologous recombination should allow these investigators to determine the precise roles of homeobox genes in mosses (Theissen *et al.*, 2001). However, research to date has not revealed any direct relationship between evolutionary changes in homeobox genes and new body plans in plants. Since the primeval protist lineages evolving into animals and plants had probably diverged as single-celled eukaryotes, it should come as no surprise that plants might depend on different genetic mechanisms to organize their body plans (Meyerowitz, 2002).

On the other hand, it is quite intriguing that the molecular diversity of the MADS-box family of transcription factors correlates with the morphological complexity of land plants (Theissen *et al.*, 2000; Vergara-Silva *et al.*, 2000; Cronk, 2001). Seven MADS-box genes have been isolated from the moss *Physcomitrella patens* (Krogan and Ashton, 2000; Henschel *et al.*, 2002) in contrast to 15 genes from the fern *Ceratopteris richardii* Brogn. and to even larger numbers from several angiosperm species. Of particular importance to angiosperm reproductive development are the MADS-box genes that act as homeotic selector genes for controlling floral organ identity. Indeed, the ABC model provides molecular validation for the classical morphological concept of serial homology, i.e. all lateral dorsoventral structures from cotyledons to carpels are leaf homologues (Goethe, 1790), because the triple mutant in all three functions exhibits foliage leaf-like structures in place of floral organs (Coen and Meyerowitz, 1991). The molecular evolution literature expresses considerable optimism that the MADS-box gene diversity underlies the evolution of plant body

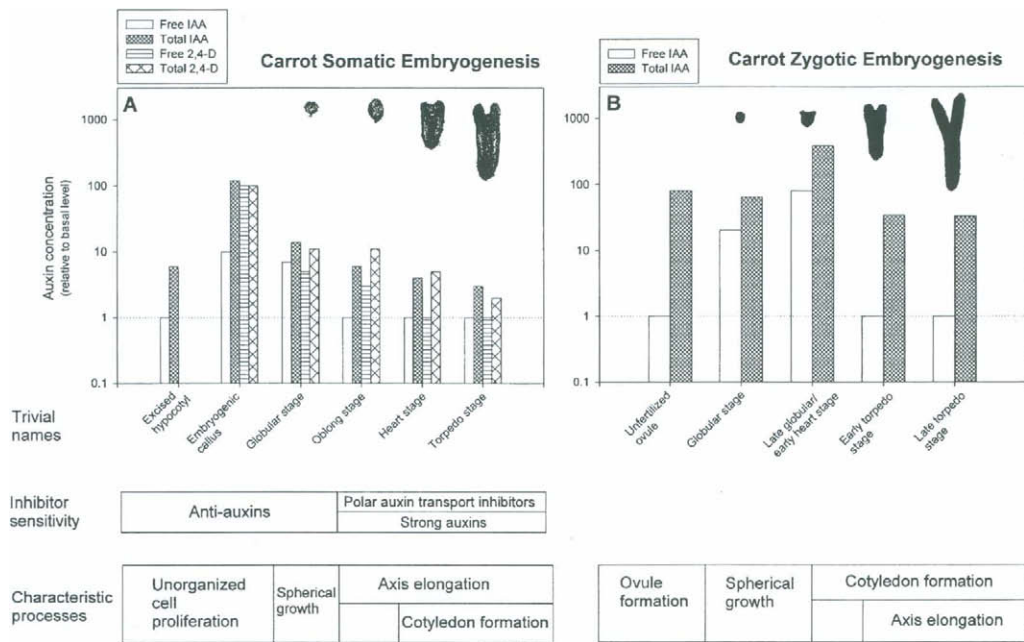


Figure 5.4 Auxin regulation of the somatic (A) and zygotic (B) embryogenesis of the carrot *Daucus carota* L. (for details, see Schiavone and Cooke, 1987; Michalczuk *et al.*, 1992a; Ribnicky *et al.*, 2002). Trivial names refer to the shapes of each embryonic stage; characteristic processes refer to the morphogenetic processes occurring at each stage; and inhibitor sensitivity refers to the auxin agonists or antagonists active at each stage. For the purpose of normalizing the data for free auxin concentrations between embryo types, all data are expressed relative to basal levels, which are 7 ng IAA/g FW and 14 ng 2,4-D/g FW in oblong to torpedo stages in somatic embryos and 26 ng IAA/g FW in unfertilized ovules and torpedo stages in zygotic embryos. Redrawn from Ljung *et al.* (2002) with permission from Kluwer Academic Publishers. Abbreviations: IAA, indole-3-acetic acid = endogenous auxin; 2,4-D,2,4-dichlorophenoxyacetic acid = synthetic auxin.

plans in general (for enthusiastic advocacy, see Vergara-Silva *et al.*, 2000). We wish to point out to the contrary that no compelling evidence yet exists to support the notion that MADS-box genes play even minor roles in establishing the fundamental tripartite, unipolar or bipolar organization of any division of land plants, including the angiosperms.

In our opinion, auxin is causally involved in the establishment of plant body plans, at least in the seed plants (Figure 5.4). The progression through both somatic and zygotic embryogenesis of the carrot *Daucus carota* L. appears to require the sequential activation of two different auxin biosynthetic pathways (Michalczuk *et al.*, 1992a,b; Ribnicky *et al.*, 1996, 2002). During the initial stages of embryo growth, a tryptophan-dependent pathway for auxin biosynthesis produces high levels of free (i.e. active) auxin, which are apparently critical for mediating the rapid cell divisions needed to generate the globular embryo. Then the embryo switches to a tryptophan-independent pathway for auxin biosynthesis that appears capable of exercising greater homeostatic control over the free auxin levels. The action of this pathway results in much lower free auxin levels that may be a necessary precondition for establishing auxin gradients regulating the polarized growth of older embryos. The results from inhibitor experiments are entirely consistent with these concepts: both synthetic auxins and polar auxin transport inhibitors are able to block or alter the polarized growth, but not the initial isodiametric expansion, of all angiosperm embryos examined

(Schiavone and Cooke, 1987; Liu *et al.*, 1993; Fischer *et al.*, 1997; Hadfi *et al.*, 1998). It remains critical to characterize auxin levels and biosynthetic pathways in other embryos than those of carrots. Auxin levels are reported to increase between the initial embryonic and subsequent postembryonic development in the wheat (*Triticum aestivum* L.) caryopsis (Fischer-Iglesias *et al.*, 2001). It is clear that these results are not directly comparable to those obtained from carrot embryo work because wheat embryos undergo prolonged postembryonic growth while still retained inside the caryopsis.

Molecular research is beginning to reveal the molecular basis for auxin action in *Arabidopsis* embryos (Souter and Lindsey, 2000; Hamann, 2001). For instance, a homozygous null mutation in the putative auxin receptor gene (*ABP1*) results in embryo development being blocked at the early globular stage (Chen *et al.*, 2001). Of even greater importance are the observations on *gnom* mutant embryos, which become enlarged spherical structures unable to initiate a polarized growth axis. The molecular basis for the mutant phenotype is that the embryos fail to localize the auxin efflux carrier PIN1 in the proper position for carrying out polar auxin transport (Steinmann *et al.*, 1999).

Moreover, auxin acts as a critical regulator of postembryonic body plan of seed plants. In particular, localized synthesis and/or polarized transport are thought to establish auxin gradients that appear absolutely critical for the positioning of new leaf primordia on the shoot apex (Meicenheimer, 1981; Reinhardt *et al.*, 2000) and of new lateral root primordia along developing roots (Reed *et al.*, 1998; Casimiro *et al.*, 2001). Finally, it is well documented that auxin exercises predominant control over many aspects of vascular tissue development, including the induction of primary vascular tissues (Roberts *et al.*, 1988; Aloni, 1995); the positioning of primary vascular bundles (Sachs, 1991; Berleth *et al.*, 2000); and the activity of vascular cambia (Uggla *et al.*, 1996, 1998). All this evidence taken together demonstrates that auxin acts as a very important regulator of the body plans of seed plants.

Much less research has been devoted to the auxin regulation of developmental processes in bryophytes and pteridophytes (for review, see Cooke *et al.*, 2002). In fact, to the best of our knowledge, no published work has directly studied auxin action in body plan organization in these plants. Nonetheless, three arguments can be advanced in support of the prediction that auxin must also help to regulate the body plans of non-seed plants. (1) it seems quite plausible that the seed plants would not have evolved *de novo* development mechanisms for generating body plans but rather would have modified pre-existing mechanisms already operating in the common ancestor of non-seed and seed plants. (2) we have noted elsewhere that the nature of metabolic regulation of auxin levels appears to have evolved in concert with the increasing morphological complexity in the land plant lineage (Sztejn *et al.*, 2000). (3) certain auxin-mediated processes, including tropisms, apical dominance and axis elongation, are widespread among all land plants, including bryophytes (Cooke *et al.*, 2002). Among the positional relationships subject to auxin regulation in non-seed plants are: rhizoid initiation in bryophyte gametophytes (Kaul *et al.*, 1962; Kumra and Chopra, 1987; Nyman and Cutter, 1981; Chopra and Vashistha, 1990), root initiation in pteridophyte sporophytes (Wardlaw, 1957; Partanen and Partanen, 1963; Wochok and Sussex, 1975) and vascular differentiation in fern sporophytes (Ma and Steeves, 1992). It is also worthwhile here to mention the provocative modelling work of Stein (1993) who attempted to relate predicted patterns of auxin concentration in the shoot apex to the observed arrangements of primary vascular tissues in a wide range of fossil plants. Although this model is based on several assumptions about auxin biosynthesis, movement and accumulation that have never been evaluated with real meristems, it does result in a close correspondence between predicted hormone distributions and underlying stelar patterns for many fossil

plants, which suggests that altered patterns of auxin action may have been involved in the diversification of stelar patterns throughout the land plant lineage.

Did major changes in auxin regulation occur prior to the Silurian–Devonian radiation?

The approach adopted in evolutionary development biology to address such questions involves: (1) the use of the fossil record and/or molecular phylogenies to identify those groups that diverged from the stem group prior to the radiation; and (2) the characterization of the developmental mechanisms in extant organisms from those early diverging lineages (Raff, 1996; Arthur, 2002). The ultimate goal is to predict those developmental mechanisms operating in the stem group that may have contributed to body plan diversification during the radiation. The critical assumption discussed earlier is that the developmental mechanisms for generating embryonic body plans are assumed to remain extraordinarily stable over geological time. This assumption has been validated in non-bilateral animals by the direct relationship between their homeobox gene diversity and their phylogenetic relationships (Peterson and Davidson, 2000), but we have no evidence to evaluate its validity for plants.

A comparable approach toward the Silurian–Devonian radiation of vascular plants would involve an investigation of the genes responsible for regulating auxin action in the three extant bryophyte lineages, which diverged from the vascular plant stem group prior to this radiation, according to morphological and molecular phylogenies (Kenrick and Crane, 1997; Qiu *et al.*, 1998; Nickrent *et al.*, 2000; Renzaglia *et al.*, 2000; Karol *et al.*, 2001). It must be acknowledged that this effort is hampered by several limitations. For instance, it is probably premature to attempt a comparative study of auxin regulatory genes in land plants because few non-seed plants, with the notable exception of *Physcomitrella patens*, are readily amenable to molecular manipulation. Moreover, the phylogenetic relationships among the bryophyte lineages remain unresolved, although the research cited above does frequently, but not always, place the mosses as the sister group to the vascular plants. Lastly, since macrofossils exhibiting the morphological characteristics of modern bryophytes (e.g. monosporangiate sporophytes and heteromorphic gametophyte-dominant generations) appear relatively late in the fossil record, one must always keep in mind the possibility that one or more modern bryophyte lineages may be derived from isomorphic, polysporangiate ancestors (Kenrick and Crane, 1997).

What can be accomplished here is that we can discuss the distribution of auxin regulatory processes in charophytes and bryophytes (Table 5.1) in order to speculate about developmental mechanisms that may have been operating in the stem group before the diversification of vascular plants in the Late Silurian–Early Devonian radiation. The tryptophan-independent pathway has definitively been identified as the predominant auxin biosynthetic pathway in the liverwort *Marchantia polymorpha* L., the moss *Polytrichum ohioense* Ren. & Card., and several vascular plants, which indicates that the capacity for tightly regulating auxin biosynthesis may be ubiquitous in the land plant lineage (Sztein *et al.*, 2000). Because the charophyte *Nitella* spp. maintains auxin levels that are comparable with those measured in bryophytes, it is likely that the tryptophan-independent pathway is also operative in this group. In so far as charophytes and liverworts carry out metabolic inter-conversions between active free and inactive conjugated forms of auxin at slow rates, it appears that the free auxin levels are maintained in these groups via the balance between biosynthetic and degradative reactions. However, hornworts and mosses share the ability for rapid auxin conjugation with vascular plants, which provides these lineages the potential

Table 5.1 Phylogenetic distribution of critical processes involved in auxin action in green plants. The occurrence of polar auxin transport (PAT) was assessed by the direct measurement of PAT in agar-block experiments and/or by the sensitivity of auxin efflux to PAT inhibitors.

| <i>Process</i> | <i>Charophytes</i> | <i>Liverworts</i> | <i>Hornworts</i> | <i>Mosses</i> | <i>Vascular plants</i> |
|---|---------------------------|---------------------------|-------------------------|-------------------------|-------------------------|
| Tryptophan-independent pathway | Predicted | Yes | n.d. | Yes | Yes |
| Auxin conjugation rates | Very slow | Slow | Intermediate to rapid | Intermediate to rapid | Rapid to very rapid |
| Predicted mechanism for regulating auxin levels | Biosynthesis/ degradation | Biosynthesis/ degradation | Conjugation/ hydrolysis | Conjugation/ hydrolysis | Conjugation/ hydrolysis |
| PAT in gametophyte rhizoids | Yes? | n.d. | n.d. | Yes | n.d. |
| PAT in gametophyte thalli | No | Yes | n.d. | Yes | n.d. |
| PAT in young sporophytes | n.s. | No | No | Yes | Yes |

n.d., no data; n.s., non-existent structure. For references, see text.

for even more precise regulation of auxin levels (Sztein *et al.*, 1995, 1999, 2000). One significant difference is that almost all vascular plants produce two specific conjugates, IAA-aspartate (or -glutamate) and IAA-1-*O*-glucose, that are not accumulated in bryophytes.

Membrane proteins capable of mediating the transmembrane auxin transport also evolved before the origin of the land plants, as evidenced by their activity in the thalli of the charophyte *Chara vulgaris* L. (Dibb-Fuller and Morris, 1992). However, there is contradictory evidence about whether charophytes carry out the more sophisticated process of intercellular auxin transport, as characterized by basipetal polarity and inhibitor sensitivity. The standard inhibitors of polar auxin transport do not affect auxin efflux from intact thalli (Dibb-Fuller and Morris, 1992), but these inhibitors are reported to have significant effects on decapitated thalli with growing rhizoids (Klambt *et al.*, 1992). Intercellular auxin transport with strong basipetal polarity and inhibitor sensitivity has been measured in liverwort thalli (Maravolo, 1976; Gaal *et al.*, 1982); moss protonemata (Rose *et al.*, 1983; Geier *et al.*, 1990), and moss rhizoids (Rose and Bopp, 1983). The sporophytes of hornworts and liverworts do not exhibit polar auxin transport (Thomas, 1980; DB. Poli, unpublished observations). By contrast, young setae of the moss *Polytrichum* maintain significant fluxes of basipetal polar transport that are even higher than those measured in corn coleoptiles (DB. Poli, unpublished observations). If mosses are the actual sister group of the vascular plants, then this observation suggests that their common ancestor evolved the auxin-dependent mechanism for generating polarized axes that is still being utilized in the sporophytes of extant members of both groups. Finally, bryophytes exhibit many of the auxin-mediated responses, such as apical dominance, phototropism and axis elongation also found in vascular plants (Cooke *et al.*, 2002). In summary, the range of auxin-regulated processes reported in extant bryophytes lends some credibility to the hypothesis that auxin was intimately involved in establishing the body plans of ancestral Silurian plants prior to the diversification of vascular plant lineages.

Conclusions

Using the conceptual framework derived from the study of the Cambrian radiation of bilateral animals, this chapter attempted to address four questions concerning the evolution of the early land plants. (1) the evidence assembled indicates that the early land plants did undergo a rapid evolutionary radiation during the Late Silurian to Early Devonian periods. (2) the characteristic body plans of different divisions of extant land plants are established during embryonic and early postembryonic development, which means that the regulatory mechanisms operating in embryonic development are also critical for generating these body plans. (3) the research evidence surveyed in this chapter establishes that the hormone auxin serves as a primary mechanism for regulating embryo and postembryonic development, at least in vascular plants. (4) judging from our current knowledge of auxin action in early-divergent bryophyte lineages, it appears that major changes in auxin action occurred in the earliest land plants prior to the Late Silurian. Therefore, the evidence available to date lends support to the appealing perspective that genetic changes in auxin action in early land plants were instrumental in the subsequent diversification of body plans in vascular plants. It follows that increased knowledge of the auxin regulation of embryonic mechanisms in extant plants should help to elucidate the developmental events that generated novel body plans during the Silurian–Devonian radiation of new vascular plant lineages.

Our enthusiasm for these interpretations is dampened somewhat by the realization that due to the limited number of relevant papers, we are skating over the conceptual equivalent of thin ice. A rigorous evaluation of the question posed in the title requires that much greater research effort be devoted toward the characterization of: (1) the phylogenetic relationships among the lineages of both extant bryophytes and putative bryophyte fossils; (2) the genetic regulation of auxin action in embryo development; and (3) the developmental mechanisms operating in charophytes, bryophytes and pteridophytes. Regardless of the ultimate answer, that question will be worth considering if it inspires further research on those three subjects.

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6

Aquaporins: structure, function and phylogenetic analysis

Joost T van Dongen and Adrianus C Borstlap

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Introduction

The colonization of land by plants – one of the most important events in the history of planet Earth – coincided with the diversification of basal embryophytes, which led to the emergence of mosses, horsetails, ferns and seed plants (Kenrick and Crane, 1997). The first embryophytes evolved as far back as 500 million years (Ma) ago from charophycean green algae, probably on the margins of drying pools. The earliest land plants inhabited damp places and were poikilohydric, that is their degree of hydration depended on atmospheric humidity. Large central vacuoles are typical of plant cells and must have already been present in the algal ancestors. In the earliest land plants vacuoles may have served, for the first time, as a reserve water supply stabilizing the hydration of the cytoplasm during short periods of drought stress (Wiebe, 1978).

The transition of poikilohydric into homoiohydric organisms has been a major step in the evolution of plants and allowed them to invade drier habitats (Walter and Stadelmann, 1968). Homoiohydric plants can maintain a constant internal water balance independent of atmospheric humidity. Their aerial parts are characterized by internal gas spaces and, more importantly, by a cuticle and closable stomata by which transpiration can be regulated.

The emergence of homoiohydric plants also required the innovation of a root system, or some similar underground structure, to exploit soil water resources (Raven and Edwards, 2001). In addition, an endohydric water-conducting system was needed to supply water to the aerial parts at rates sufficient to support the transpiratory loss of water during photosynthesis. Early land plants had an unexpected diversity of conducting cells, which evolved by parallel evolution (Ligrone *et al.*, 2000). Among these, the precursors of the xylem tracheary elements were the most successful and became the water-conducting system of all vascular plants (Friedman and Cook, 2000). At the same time the phloem evolved as a food conducting system to supply the heterotrophic roots with the products of photosynthesis.

The tracheal system in the xylem provides a low-resistance, apoplasmic pathway for the upward flow of water. But the passage of water through plant tissues also involves symplasmic and transcellular pathways and, therefore, transmembrane water fluxes (Steudle and Peterson, 1998). The sieve elements in the phloem constitute a symplasmic route by which organic nutrients are translocated. Being hydraulically operated, the functioning of this food-conducting system depends on the influx of water into the sieve elements at the loading site and the efflux of water at the site of unloading. Influx and efflux of water across the plasmalemma and tonoplast is also at the basis of turgor-regulated movements in plants, such as the opening and closure of stomata and the diurnal movements of leaf pulvini. Thigmotropic, turgor-dependent movements may be very fast as, for example, in the pulvini of the sensitive plant (*Mimosa pudica* Mimosa pudica L.), the traps of insectivorous plants (*Dionaea* *Dionaea* Ellis, *Utricularia* *Utricularia* L.), and the gynostemium of trigger plants (*Stylidium*). These rapid movements (15–30 ms in *Stylidium* *Stylidium* Sw. ex Willd) are thought to result from a very fast efflux of both solutes and water from motor cells (Hill and Findlay, 1981).

Transport of water across cell membranes

In the 18th century, Stephen Hales was the first to measure the transpiration of plants and the force of upward sap pressure in stems (Hales, 1727). Since then, the study of water flows and water relations in plants has been a major theme in plant physiology, though the mechanism by which water crosses plant cell membranes has received relatively little attention. The basic structure of cell membranes is a fluid lipid bilayer, which mainly consists of phospholipids and sterols. The intrinsic (or integral) membrane proteins are 'floating' in this 'sea' of lipid molecules, whereas extrinsic membrane proteins are located on both surfaces of the membrane (Singer and Nicolson, 1972). The lipid bilayer is a real barrier for water movements, impeding the diffusion of water by at least a factor of 3000.

Water can cross a membrane by diffusion or in response to an osmotic pressure gradient. The rate at which water permeates a membrane may accordingly be defined by the diffusion permeability coefficient (P_d) or by the osmotic permeability coefficient (P_f). Interestingly, the relationship between P_f and P_d depends on the nature of the membrane. For an 'oil' membrane, such as a phospholipid bilayer, the concentration of water within the membrane is so low that interactions between water molecules may be neglected. As a consequence, P_f and P_d for such membranes are equal. This may be different for porous membranes. If the movement of water through the membrane is restricted to the pores, and if the pores are so narrow that the transport of water molecules occurs in a single file, the remarkable relationship $P_f/P_d = N$ can be derived, where N is the number of water molecules in the single file that fills the pore (Finkelstein, 1987).

The permeability of lipid membranes for water has been the subject of intensive investigations (Finkelstein, 1987). Depending on parameters such as lipid chain length, degree of chain saturation and sterol content, P_f (or P_d), values for artificial lipid bilayers have been found to span almost three orders of magnitude, ranging from 2×10^{-5} to $1 \times 10^{-2} \text{ cm s}^{-1}$. As these values encompass nearly the entire range of values reported for cell membranes it seemed that the water permeability of a membrane was fully attributable to its bilayer structure. Besides the magnitude of P_f , however, some additional issues remained that required an explanation. First, the P_f and P_d values for cell membranes were often found to be widely different. In careful experiments with red cell membranes, for instance, P_d amounted to $0.37 \times 10^{-2} \text{ cm s}^{-1}$, whereas the value of P_f was nearly 6-fold higher and determined at $2.15 \times 10^{-2} \text{ cm s}^{-1}$ (Moura *et al.*, 1984). Secondly, unlike the water flux through lipid bilayers, that through cell membranes could be largely inhibited by sulphhydryl reagents such as *p*-chloromercuribenzenesulphonate (pCMBS), indicating that a proteinaceous channel is involved. Finally, the water permeability of epithelial cells in the urinary bladder of frog and toad and the mammalian collecting kidney tubules can increase up to 50-fold in response to vasopressin (the 'antidiuretic hormone'). This hormonal regulation is difficult to explain if water transport occurred exclusively through the bilayer. Thus around 1985 evidence had accumulated that the membranes of animal cells could contain pores which transported water and little else (Finkelstein, 1987). Similar conclusions were reached from studies of water transport into characean cells (Wayne and Tazawa, 1990). The nature of these pores, however, remained a mystery.

Discovery of aquaporins and the MIP-family

The first aquaporin was serendipitously discovered in Peter Agre's laboratory during efforts to purify the 32-kDa subunit of the red cell Rh blood group antigen. A novel 28-kDa protein was found, which displayed some similarity with certain channel-forming integral proteins, hence its name CHIP28. Further studies showed that the protein was also abundantly present in the epithelia of proximal tubules and thin descending limbs of rat kidneys. Renal epithelia are the site of intense water transport. In humans they are responsible for the absorption of 180 litres of fluid each day. Its presence in the kidney therefore sparked the idea that the novel 28-kDa polypeptide may be the long sought water channel (Agre *et al.*, 1998; Borgnia *et al.*, 1999). When the gene coding for CHIP28 was cloned this hypothesis could be tested by expression in oocytes of the African claw frog, *Xenopus laevis*. After injection of cRNA of CHIP28 into the oocytes, the permeability of oocyte membrane for water increased dramatically (from $\sim 0.2 \times 10^{-2}$ to $\sim 2 \times 10^{-2} \text{ cm s}^{-1}$), which caused them to explode rapidly when transferred to a hypotonic medium (Preston *et al.*, 1992). Thus CHIP28 was identified as the first water-transporting protein and aptly renamed as aquaporin 1 (AQP1).

AQP1 belongs to the family of major intrinsic proteins (MIPs), one of the 46 families of the class of α -helix-type channels (Saier, 2000). The family has been named after the major intrinsic protein in the cell membrane of lens fibre cells, where it comprises >60% of the membrane proteins (Gorin *et al.*, 1984). The existence of the MIP family first transpired when striking similarities were noticed in the amino acid sequences of the MIP from eye lens fibre cells (now known as AQP0) and two other membrane proteins from quite disparate cells (Baker and Saier, 1990). One of these was GlpF, the glycerol facilitator in the cell membrane of *Escherichia coli*. The other was Nodulin 26, a plant-encoded abundant

protein isolated from the peribacteroid membrane of soybean nodules. At the time CHIP28 joined the family and was identified as the first aquaporin, some additional plant members of the MIP family were already known. These included the turgor-responsive gene *clone 7a/TRG-31* from the pea (Guerrero *et al.*, 1990; Guerrero and Crossland, 1993); *TobRB7*, which was identified in a cDNA library from tobacco roots (Yamamoto *et al.*, 1990); and α -TIP, an abundant protein from the tonoplast of protein-storing vacuoles in seeds of the common bean (Johnson *et al.*, 1990). The gene that codes for another tonoplast localized MIP-like protein in *Arabidopsis thaliana* (L.) Heynh, called γ -TIP, was also cloned in Maarten Chrispeels' laboratory, and subsequently characterized as the first plant aquaporin (Höfte *et al.*, 1992; Maurel *et al.*, 1993). The discovery of the PIPs, plant aquaporins that localize to the plasma membrane, followed soon after (Kammerloher *et al.*, 1994).

Structure and function of MIPs

Most MIPs are composed of 250–290 amino acids. All have six hydrophobic, putative transmembrane helices, which are connected by loops (Figure 6.1). The two halves of the protein exhibit substantial sequence similarity to one another. The B and E loop contain the asparagine–proline–alanine (NPA) motif which is highly characteristic for aquaporins. Because of the uneven number of transmembrane helices in each repeat, the two domains are obversely oriented in the membrane.

The functionality of MIPs is generally assessed after expression in *Xenopus* oocytes. These studies have shown that most MIPs can be classified either as aquaporins, which selectively facilitate the transport of water across the membrane, or as aquaglyceroporins that transport water as well as glycerol. Recently atomic models were published for the prototypes of the two functional classes, AQP1 and GlpF (Murata *et al.*, 2000; Fu *et al.*, 2000).

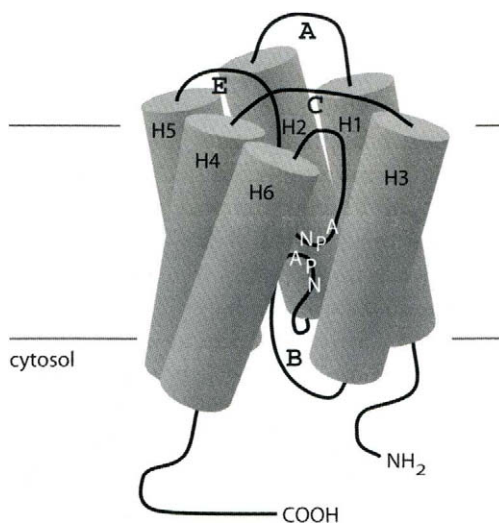


Figure 6.1 Topology of a major intrinsic protein. Six transmembrane helices (H1–6) are linked by loops (A–E). Loops B and E contain the highly conserved asparagine–proline–alanine (NPA) motif and bend inward into the protein. The NH₂- and COOH-termini are localized in the cytosol, whereas the opposite part of the protein is orientated towards the apoplast or the vacuolar lumen. Note that loop D between helices 4 and 5 is not visible in this figure.

Although AQP1 and GlpF are less than 30% identical, their gross structural features are very similar. Both proteins are right-handed helix bundles, organized in two symmetrical domains. The aquaporin has a narrow central constriction and wider external openings. The aqueous pathway is largely lined with hydrophobic residues that permit rapid water transport. The narrowest part of the pore, 0.3 nm (3 Å) wide, is in the centre of the membrane, just behind the region where loops B and E interact with each other. Four hydrophobic amino acid residues in helices 1, 2, 4, and 5, respectively, together with the asparagines of the NPA-motifs line the inside of the constriction pore (Murata *et al.*, 2000). The unit permeability of AQP1 has been determined at $\sim 10^9$ water molecule per second, remarkably close to the value predicted by theory (Sansom and Law, 2001).

In GlpF the selectivity filter consists of a 2.8 nm (28 Å) long, 0.34–0.38 nm (3.4–3.8 Å) wide amphipathic channel. The alkyl backbone of the glycerol molecule is wedged against the hydrophobic side of the channel, whereas the three hydroxyl groups of glycerol form successive hydrogen bonds with side groups of the hydrophilic side of the pore (Fu *et al.*, 2000).

MIPs of bacteria, fungi and animals

The genomes of the archaeobacteria *Archaeoglobus fulgidus* and *Methanobacterium thermoautotrophicum* contain a single gene that codes for a MIP (Figure 6.2), but MIPs are lacking in *Methanococcus janaschii* and other sequenced archaea. Bacteria and fungi may possess MIPs of the aquaporin or the aquaglyceroporin type (Hohmann *et al.*, 2000). In *E. coli* the aquaporin AqpZ is involved in cellular hydration, whereas GlpF has a role in the utilization of glycerol. In the genome of bakers' yeast (*Saccharomyces cerevisiae*) three genes are related to aquaglyceroporins and three others to aquaporins. Curiously, most laboratory strains have no functional aquaporins because of mutations in the coding genes. Glycerol is an important osmolyte in many fungi, including yeast. The yeast aquaglyceroporin Fps1 is involved in the export of glycerol when the cells are exposed to a hypotonic medium (Luyten *et al.*, 1995).

The worm *Caenorhabditis elegans* has eight MIP genes (Kuwahara *et al.*, 1998). Only a few MIPs are known from insects. One is BIB (big brain) of *Drosophila*, which is a homologue of the mammalian AQP4. Another insect MIP is AQP_{cic} which was identified in the filter chamber of a sap-sucking cicada (*Cicadella viridis*) and, recently, the presence of an aquaporin in the Malpighian tubules of the bug *Rhodnius prolixus* was demonstrated (Borgnia *et al.*, 1999; Echevarría *et al.*, 2001).

In mammals 11 different MIPs have been identified. The mammalian aquaporins (AQP0, AQP1, AQP2, AQP4, AQP5, AQP8) play key roles in transmembrane water fluxes in many tissues (reviewed by Deen and Van Os, 1998; Borgnia *et al.*, 1999). Kidney, airways, eye and brain have tissues with complex expression patterns involving multiple aquaporins and aquaglyceroporins. The kidney, for instance, harbours at least four aquaporins and two aquaglyceroporins. AQP0 seems to be expressed exclusively in lens fibre cells, whereas AQP1 is expressed widely in various epithelial and capillary endothelia, including proximal tubule and thin descending limb in kidney. In kidney collecting ducts AQP2 mediates the vasopressin-induced increase in water permeability, which is required for the concentration of urine. AQP4 is the main aquaporin in brain. AQP5 resides in lung and salivary and lachrymal glands and presumably regulates airways humidification and the release of saliva and tears. AQP8 is found in various tissues, including those of testis, pancreas and liver. AQP6 is the only known intracellular aquaporin in animal cells. It has

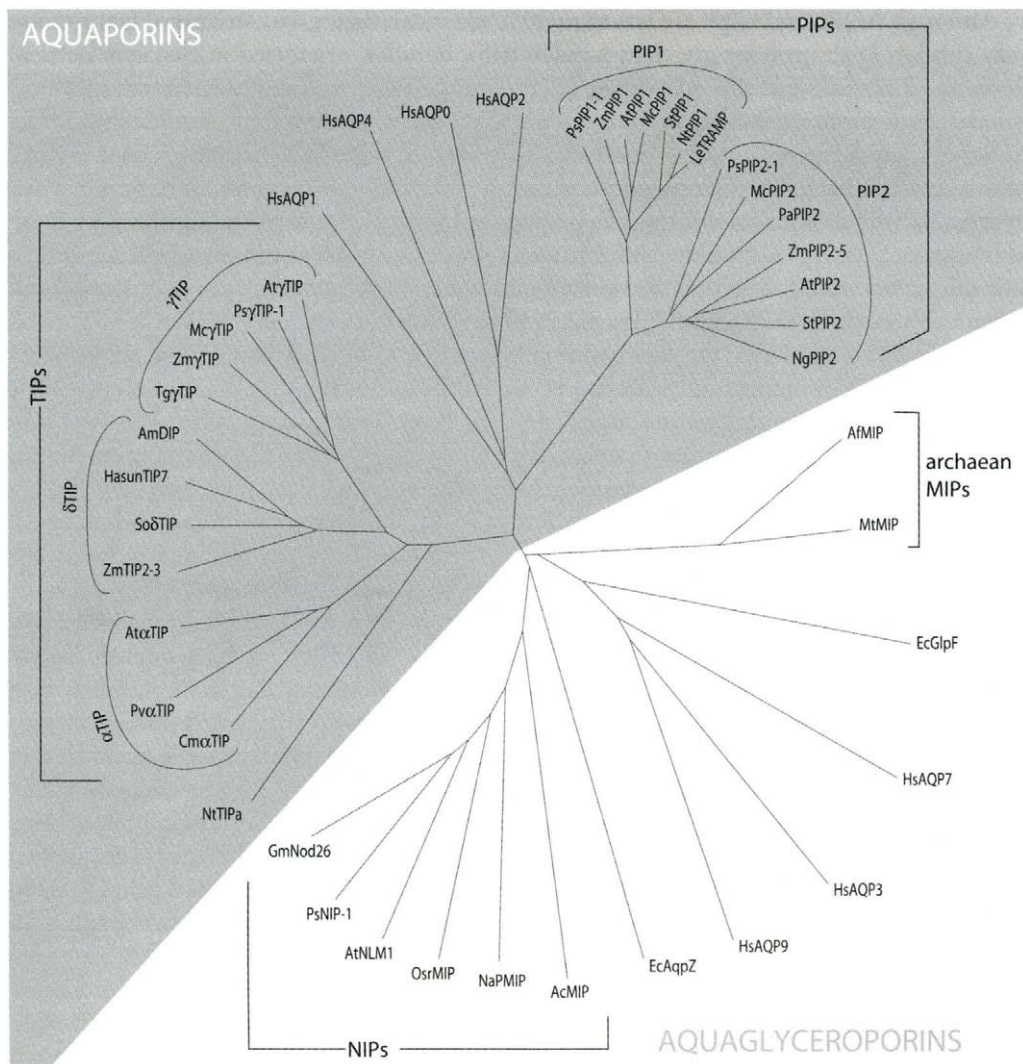


Figure 6.2 Phylogenetic analysis of some selected members of the family of major intrinsic proteins. Included are the MIPs from archaeobacteria, the two MIPs from *E. coli*, several human MIPs and representatives of three subfamilies of plants MIPs (PIPs, plasmamembrane intrinsic proteins; TIPs, tonoplast intrinsic proteins; NIPs, Nodulin 26-like MIPs). Protein sequences were aligned using ClustalW at DDBJ (<http://www.ddbj.nig.ac.jp/>) and the tree was drawn with TreeView 1.6.5 (Page, 1996). Amino acid sequences of the following proteins were used as input. Plant PIP1-members: *Pisum sativum* PsPIP1-1 (P25794), *Zea mays* ZmPIP1 (Q9XF59), *Arabidopsis thaliana* AtPIP1 (Q39196), *Mesembryanthemum crystallinum* McPIP1 (Q40266), *Solanum tuberosum* StPIP1 (Q9XGF4), *Nicotiana tabacum* NtPIP1 (O24662), *Lycopersicon esculentum* LeTRAMP (Q08451). Plant PIP2-members: *Pisum sativum* PsPIP2-1 (Q9XGG8), *Zea mays* ZmPIP2-5 (Q9XF58), *Arabidopsis thaliana* AtPIP2 (P43287), *Mesembryanthemum crystallinum* McPIP2 (O24050), *Solanum tuberosum* StPIP2 (Q9XGF3), *Nicotiana glauca* NgPIP2 (Q9FPZ7), *Picea abies* PaPIP2 (O49921). Plant α TIP-members: *Phaseolus vulgaris* Pv α TIP (P23958), *Arabidopsis thaliana* At α TIP (P26587), *Cucurbita maxima* Cm α TIP (Q39646). Plant γ TIP-members: *Pisum sativum* Ps γ TIP (Q9XGG6), *Tulipa gesneriana* Tg γ TIP (Q41610), *Arabidopsis thaliana* At γ TIP (P25818), *Zea mays* Zm γ TIP (O64964), *Mesembryanthemum crystallinum* Mc γ TIP (O24048). Plant δ TIP-members: *Spinacia oleracea* So δ TIP (Q9SM46), *Helianthus annuus* HasunTIP7 (Q39958), *Antirrhinum majus* AmDIP (P33560), *Zea*

low water permeability but, interestingly, is also permeable for anions (Yasui *et al.*, 1999). Of the aquaglyceroporins AQP3 is abundantly expressed in kidney collecting duct and at lower levels at multiple sites including airways, whereas AQP10 seems to be specific for the intestine (Ishibashi *et al.*, 2002). AQP9 from the liver is an extraordinary aquaglyceroporin as it is also permeable for various other neutral solutes (Tsukaguchi *et al.*, 1998). The aquaglyceroporin AQP7, isolated from testis, is identical to AQPap from adipocytes, which is involved in the glycerol export during lipolysis (Kishida *et al.*, 2000, 2001). The physiological role of the other aquaglyceroporins is not so clear.

Plant MIPs

Genes coding for MIPs are particularly abundant in plant genomes. In *Arabidopsis* as many as 35 MIP-genes have been recorded, which is 3-fold more than in any other sequenced organism and hints at the importance of hydraulics in a wide range of plant processes. Based on sequence homology, members of the MIP-family in higher plants fall into three major subgroups: the plasma membrane intrinsic proteins (PIPs), the tonoplast intrinsic proteins (TIPs), and the Nodulin 26-like MIPs (NLMs or NIPs). A fourth subfamily, the short, basic intrinsic proteins (SIPs), was recently revealed by genomic analysis, but nothing is known about their function, intracellular localization or expression patterns. *Arabidopsis* possesses 13 PIPs, 10 TIPs, 9 NIPs and 3 SIPs (Johanson *et al.*, 2001).

The PIP subfamily can be further divided into the highly homologous but clearly distinct subgroups PIP1 and PIP2, which differ significantly in the N- and C-terminus (Kammerloher *et al.*, 1994). Inasmuch as PIP2-members expressed in *Xenopus* oocytes induce relatively high permeabilities for water, but not for small neutral solutes like glycerol and urea, they may be regarded as orthodox aquaporins. The function of PIP1-members is less clear because, after expression in oocytes, they show a much lower aquaporin activity than PIP2-members or no activity at all (Kammerloher *et al.*, 1994; van Dongen, 2001). On the basis of circumstantial evidence it has been suggested that some PIP1-members (TRG-31 from pea and the ripening-associated membrane protein from tomato, TRAMP) might function as solute transporters (Jones and Mullet, 1995; Chen *et al.*, 2001). Another PIP1-member from maize expressed in oocytes slightly increased the transport of boric acid (Dordas *et al.*, 2000).

On the basis of a phylogenetic analysis of TIPs in *Arabidopsis* and maize, five subgroups are currently distinguished (Chaumont *et al.*, 2001; Johanson *et al.*, 2001). Best known are the α -, γ -, and δ -TIPs which, according to a newly proposed nomenclature are designated as the TIP3-, TIP1- and TIP2-group, respectively (Johanson *et al.*, 2001). Representatives of the α -, γ -, and δ -TIPs have been shown to function as aquaporins (reviewed by van Dongen, 2001). Different vacuolar compartments in the plant cell may

mays ZmTIP2-3 (O81216), *Nicotiana tabacum* Nt δ TIPa (Q9XG70). Members of the plant NIP-subfamily: *Pisum sativum* PsNLM (Q9XGG7), *Glycine max* GmNod26 (P08995), *Arabidopsis thaliana* AtNLM1 (O48595), *Oryza sativa* OsMIP (Q40746), *Nicotiana glauca* NaPMIP (P49173), *Adiantum capillus-veneris* AcMIP (Q9FXW2). Human aquaporins: *Homo sapiens* HsAQP0 (P30301), HsAQP1 (P29972), HsAQP2 (P41181), HsAQP4 (P55087). Human aquaglyceroporins: *Homo sapiens* HsAQP3 (Q92482), HsAQP7 (O14520), HsAQP9 (O43315). MIPs from *Escherichia coli* EcAqpZ (P48838), EcGlpF (P11244). Archaeal MIPs: *Archaeoglobus fulgidus* AfMIP (O28846), *Methanobacterium thermoautotrophicum* MtMIP (O26206).

have different complements of TIP-subtypes. The α -TIPs are found in the membrane of protein storage vacuoles in seeds, whereas δ -TIPs localize to the membrane of vacuoles containing vegetative storage proteins and pigments. The γ -TIPs are typical for the membrane of lytic vacuoles, but may be also present in the tonoplast of storage vacuoles together with α - and/or δ -TIPs (Jauh *et al.*, 1999).

NIPs probably represent the most ancient MIPs in the plant kingdom, as their closest homologues are the MIPs from the archaeobacteria (see Figure 6.2). The best-studied NIP is Nodulin 26, the major intrinsic protein of the peribacteroid membrane of the soybean nodule (Dean *et al.*, 1999, and references therein). Nodulin 26 as well as LIMP2 from the legume *Lotus japonicus* (Regel) K. Larsen (Guenther and Roberts, 2000) are exclusively expressed in nodules. A pea homologue (possibly orthologue) of Nodulin 26 is specifically expressed in the coat (integument) of developing seeds (van Dongen, 2001). It may be noted that the peribacteroid membrane in nodules and the plasma membrane of seed coat parenchyma cells have a common function. Both membranes have to accommodate the export of a variety of nutrients from the cytosol of the plant cell to feed the symbiotic bacteroids or the plant embryo.

It has been suggested repeatedly that the Nodulin 26 channel might be involved in the release of nutrients from the host cell, but up to now only two transport functions have been revealed. When expressed in oocytes, Nodulin 26 appears to function as a channel for water and glycerol (Dean *et al.*, 1999), whereas purified Nodulin 26 reconstituted in lipid vesicles functions as an ion channel (Lee *et al.*, 1995).

Though once thought to be unique for the legume nodule, NIPs have also been identified in non-legumes. The NIPs from *Arabidopsis* were reported to be expressed exclusively (*AtNLM1* and *AtNLM5*) or predominantly (*AtNLM2* and *AtNLM4*) in roots (Weig and Jakob, 2000 and references therein). A gene coding for a NIP was also cloned from the pollen of *Nicotiana glauca* Link and Otto and another one from rice anthers (Liu *et al.*, 1994), the latter being expressed in shoots but not in roots. Interestingly, the only MIP-gene cloned so far from a non-seed plant, the fern *Adiantum capillus-veneris* L., belongs to the NIP subfamily (see Figure 6.2). As compared with other MIP genes, those coding for non-nodule NIPs seem to be expressed at low levels. Because the peribacteroid membrane is derived from the plasma membrane of the host cell, it may be expected that the non-nodule NIPs are localized in the latter membrane, but experimental evidence for this is lacking.

Classification of the NIP-subfamily into the aquaporin cluster (Heymann and Engel, 1999) has now to be revised because the five NIP-members whose functionality has been assessed all appeared to transport water as well as glycerol (reviewed by van Dongen, 2001). This seems to imply that members of the NIP-subfamily are the aquaglyceroporins of plants. As for mammalian aquaglyceroporins, the physiological significance of their plant counterparts is largely unknown. Glycerol is generally not regarded as an important osmolyte in plants, although the possibility that it might be so in certain specialized cells cannot be excluded.

That NIPs are aquaglyceroporins is underscored by sequence alignments with other glycerol transporters. The constriction pore (selectivity filter) in GlpF is lined by three residues: Trp⁴⁸, Phe²⁰⁰ and Arg²⁰⁶, which are crucial for the selective permeation of glycerol by this transporter (Fu *et al.*, 2000; Unger, 2000). Alignment of several glycerol-transporting MIPs showed that a tryptophan residue similar to Trp⁴⁸ in GlpF is conserved in Fsp1, the glycerol facilitator from *Saccharomyces cerevisiae*, and in several members of the NIP-subfamily, whereas in the mammalian aquaglyceroporins (AQP3, AQP7 and AQP9) it is replaced by a phenylalanine residue (van Dongen, 2001).

Plant aquaporins may play a role in a wide variety of processes, from water uptake during cell elongation to the water fluxes through plant tissues that are linked to the transpiration stream. Most likely they are also involved in other hydraulic processes such as the translocation of organic nutrients through sieve elements and turgor dependent movements of stomata, pulvini and the traps of insectivorous plants. The vacuolar aquaporin γ TIP seems to play a crucial role in the large and rapid turgor variations in pulvinar motor cells of the sensitive plant *Mimosa pudica* (Fleurat-Lessard *et al.*, 1997).

As for other multigene families the expression of various MIPs in plants probably runs the gamut from general and overlapping to specific and localized. Expression of certain plant aquaporins is regulated by blue light, gibberellic acid or water deprivation. Specific aquaporins are involved in the inhibition of self-pollination, nematodal infection and the interaction between the plant parasite *Cuscuta* L. with its host (reviewed by Maurel, 1997; Werner *et al.*, 2001).

Phylogenetic analysis

Phylogenetic analysis of MIP sequences reveals two main clusters (see Figure 6.2), which have been designated as the AQP and GLP cluster (Heymann and Engel, 1999). The AQP cluster includes the plant aquaporins of plasma membrane and tonoplast and the animal aquaporins. The GLP cluster includes the aquaglyceroporins of bacteria, yeast, animals and plants (NIPs), but also AqpZ, the aquaporin from *E. coli*, and the two MIPs from archaeobacteria.

One interpretation of this analysis could be that the common ancestor of archaea, bacteria and eukaryotes contained a single gene that coded for a MIP. This gene may be thought to have persisted in extant archaeobacteria, while a gene duplication in the common ancestor of bacteria and eukaryotes gave rise to aquaporins and aquaglyceroporins. This scenario would provide a simple explanation for the presence of the two functional MIP-types in all major clades, except the archaea. However, it seems now generally accepted that the precursor of the eukaryotic cell originated from an archaean line and that this event was preceded by an earlier diversification into the bacteria and archaeans (Doolittle, 1999; Woese, 2000). This would mean that the bacterial aquaglyceroporins (such as GlpF) and those of eukaryotes (the mammalian AQP3, AQP7, AQP9 and AQP10 and the NIP-members in plants) have evolved by parallel evolution.

Nearly all known sequences of plant MIPs are from angiosperms. Members of the PIP1 and PIP2 subtypes as well as of the α , γ , and δ subtypes of TIPs are present in both monocotyledons and dicotyledons. This indicates that the various subtypes of plant aquaporins were already invented before the two angiosperm classes diverged. Gymnosperm MIPs are represented in the databases by a few PIP1- and PIP2-members. The only MIP-gene cloned so far from a non-seed plant, the fern *Adiantum capillus-veneris*, belongs to the NIP subfamily (see Figure 6.2). Among the 10954 sequences in an EST library of the moss *Physcomitrella patens* (Hedw.) Bruch Schimp. (<http://www.moss.leeds.ac.uk>) we found only a single fragment of a MIP gene. This fragment (GenBank acc. no. AW476973) is most similar to the corresponding fragment of a gymnosperm PIP2-member.

It seems likely that the plant aquaporins of the plasma membrane (PIPs) and the tonoplast (TIPs) are derived from a common ancestor. Tonoplast aquaporins probably arose after the diversification of fungi and green plants, because the vacuole membrane in fungi, at least that in yeast, seems to lack aquaporins. The various types of TIPs possibly evolved concurrently with the emergence of distinct types of plant vacuoles. Because of the lack of

information of MIPs in green algae, bryophytes, lycopods, horsetails and ferns we have no idea when the various types of PIPs and TIPs have originated during the evolution of plants.

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7

Evolutionary origin of the ethylene biosynthesis pathway in angiosperms¹

Elizabeth A Reynolds and Philip John

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Introduction

In angiosperms, ethylene is an important regulator of seed germination, the response to wounding, leaf and flower abscission, senescence and the ripening of climacteric fruits. Its biosynthesis follows a well-characterized pathway from methionine which gives rise to the immediate precursor of ethylene, 1-aminocyclopropane-1-carboxylic acid (ACC). In non-angiosperms, much less is known of how ethylene is generated. However, it is known that in non-seed plants, ACC is not involved (Osborne, 1989). In this chapter we describe a feasible scenario for the evolutionary origin of the angiosperm ethylene biosynthesis pathway, refining previously published preliminary accounts (John, 1997; John *et al.*, 1997, 1999) and presenting new molecular evidence relating to the origin of ACC oxidase.

¹Dedicated to the memory of Dr Andy G Prescott.

Evolution of the angiosperm ethylene biosynthesis pathway

Early responses to stress conditions

In angiosperms the ethylene biosynthesis pathway is as shown in Figure 7.1. Responsiveness to ethylene appears to be present throughout the plant kingdom. All species of algae, ferns and gymnosperms that have been tested detect and respond to ethylene (Edwards and Miller, 1972; Cookson and Osborne, 1978; Abeles *et al.*, 1992; Kong and Yeung, 1994; Kwa *et al.*, 1995; Chernys and Kende, 1996; Ingemarsson and Bollmark, 1997). Presumably evolution of ethylene responsiveness was driven by the natural connection between ethylene generation and plant stress. When cells are damaged, oxidative breakdown of cell constituents, particularly membrane fatty acids (Abeles *et al.*, 1992) generates ethylene, albeit in amounts that are orders of magnitude lower than those generated physiologically. Thus the gas constitutes a potentially useful signal of stress. An ability to detect (and then respond to) ethylene would have allowed plants to develop stress responses. Representatives of all major groups of land plants possess a high-affinity ethylene-binding activity (Bleeker, 1999). This activity was also found in the cyanobacterium, *Synechocystis*, which neither makes ethylene, nor responds to it. The protein responsible may function simply as a copper-binding protein. Thus we have evidence that the ability to detect ethylene arose early in the evolution of land plants (see *a* in Figure 7.2).

As land plants evolved, they acquired increasingly complex physiological and biochemical responses to the stresses to which they were exposed. Among the biochemical responses were betaine synthesis, to reduce water potential during drought, and lignification, to limit predator and pathogen attack. The synthesis of both betaine and lignin depends upon the availability of *S*-adenosylmethionine (SAM). This is an abundant metabolite in all organisms, playing a key role in transmethylation reactions. It is derived from methionine by the action of SAM synthetase (see *b* in Figure 7.2). As a present-day example of how stress stimulates the biosynthesis of SAM, Mayne *et al.* (1996) showed that drought conditioning of *Pinus banksiana* Lamb. seedlings, which enhanced betaine synthesis 3-fold, increased SAM synthetase activity 2-fold and mRNA abundance 6-fold.

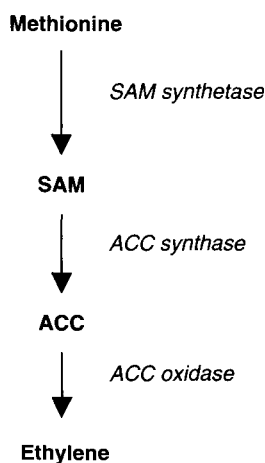


Figure 7.1 The biosynthetic pathway for ethylene as characterized in angiosperms.

The role of ACC

SAM is also the precursor of ACC, which arises by the action of ACC synthase on SAM. ACC is detectable in representatives of all major groups of land plants (J. Hodson, unpublished data). ACC has been shown to be present in the ferns *Regnellidium diphyllum* Lindman (Chernys and Kende, 1996; Osborne *et al.*, 1996) and *Marsilea quadrifolia* L., and notably, in *M. quadrifolia* it is generated by an ACC synthase with many of the properties of the angiosperm ACC synthase (Chernys and Kende, 1996). However, ACC is not used as the precursor of ethylene in these ferns (Cookson and Osborne, 1978; Osborne *et al.*, 1996; Chernys and Kende, 1996). Thus we propose that ACC was accumulated in these non-seed plants without being used as a precursor of ethylene. Naturally occurring compounds, such as ACC, that contain a cyclopropane ring can have insecticidal, antimicrobial and neurochemical properties (Salaün and Baird, 1995), with ACC itself known to have neurochemical properties (Zapata *et al.*, 1996). These properties would have made ACC useful in the phytochemical armoury of the plant. Therefore, it is proposed that before ACC became the precursor of ethylene, it accumulated as a response to predator or pathogen attack. Like the other biochemical responses to stress, ACC accumulation would have resulted from enhanced rates of SAM generation (see *b* in Figure 7.2).

The acquisition of ACC oxidase

The key step in the evolution of the complete angiosperm ethylene biosynthetic pathway is the acquisition of the final stage: the ability to generate ethylene from ACC by the action of the enzyme ACC oxidase (Figure 7.3). ACC readily penetrates plant tissues and, therefore, in principle, a simple test for the presence of ACC oxidase is available by measuring the ethylene generated when plant tissues are presented with ACC. On the basis of

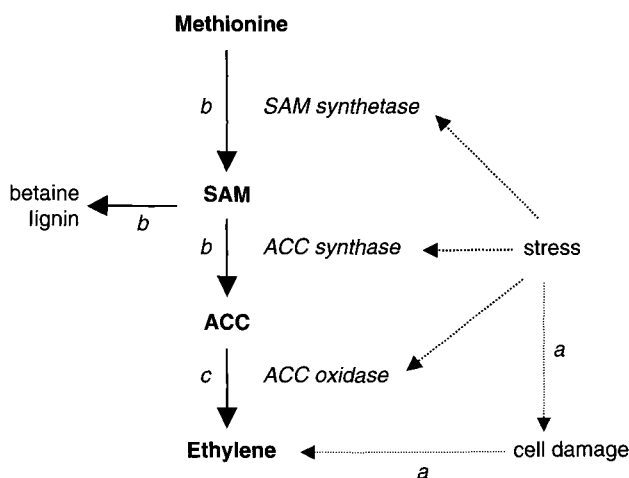


Figure 7.2 The evolutionary sequence leading to the pathway for ethylene biosynthesis in angiosperms. *a*, in the earliest land plants, cell damage arising from stress leads to the formation of ethylene by chemical breakdown of cell constituents; *b*, in ferns and allied non-seed plants, stress leads to the increased expression of S-adenosylmethionine (SAM) synthetase for lignin and betaine synthesis; and to enhanced ACC synthase activity and the accumulation of ACC; *c*, with acquisition of ACC oxidase in a group of seed plants ancestral to the Gnetales, Pinopsida and angiosperms, ethylene is produced from ACC.

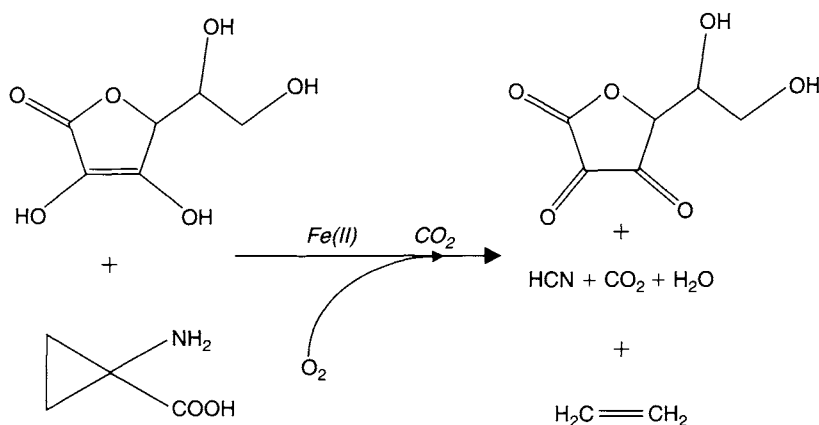


Figure 7.3 The reaction catalysed by ACC oxidase. 1-aminocyclopropane-1-carboxylate is oxidized to ethylene, and ascorbate is oxidized to dehydroascorbate. Fe(II) and CO_2 act as cofactors.

Table 7.1 Distribution among gymnosperms of ACC oxidase activity measured *in vitro* (from data in Reynolds and John, 2000)

| | | Leaf | Germinated seed |
|--------------|---|------|-----------------|
| (angiosperms | <i>Cucumis</i> | Yes | Yes) |
| Pinopsida | <i>Pinus</i> , <i>Pseudotsuga</i> × <i>Cupressocyparis</i> | No | Yes |
| Gnetales | <i>Ephedra</i> | No | Yes |
| Ginkgoales | <i>Ginkgo</i> | No | No |
| Cycadales | <i>Cycas</i> , <i>Dioon</i> , <i>Zamia</i> | No | No |

such tests, Osborne (1989) proposed that ACC was used by seed plants, but not by ferns and mosses. After it became possible for ACC oxidase to be assayed *in vitro* (Verweridis and John, 1991), Reynolds and John (2000) demonstrated that among the extant seed plants, only angiosperms showed ACC oxidase activity in leaf material; activity was absent from gymnosperm leaf extracts. Moreover, among representatives of the four extant gymnosperm groups, ACC oxidase activity was present in seedlings of the Pinopsida and Gnetales, but absent from seedlings of the Ginkgoales and Cycadales (Table 7.1). As a representative of gymnosperm ACC oxidases, the enzyme from *Pinus nigra* var. *nigra* Arnold was shown to resemble biochemically the angiosperm enzyme, including a requirement for ascorbate, CO_2 and Fe(II) (Reynolds and John, 2000). The simplest explanation of these findings was that ACC oxidase arose in a common ancestor of the three groups: angiosperms, Gnetales and Pinopsida, with the Ginkgoales and Cycadales basal to this proposed origin (Reynolds and John, 2000).

Our finding that ACC oxidase was strongly represented in germinated seeds of certain gymnosperms, but absent from leaf tissues of all gymnosperms tested (Table 7.1), led us to suggest (Reynolds and John, 2000) that the earliest role played by ACC oxidase may have been associated with germination, and only later in the evolution of angiosperms did it expand its repertoire of functions to leaves, and to the distinctively angiosperm organs of flowers and fruits. However, this suggestion must remain tentative until more is known of the relationship between ethylene production and germination (Petruzzelli *et al.*, 2000;

Matilla, 2000). Even in the relatively well-studied angiosperms 'the role of ethylene in germination remains controversial. Some authors hold that (it) is a consequence of the germination process, while others contend that ethylene production is a requirement for germination' (Matilla, 2000). Among the gymnosperms, nothing is known of the role of ethylene in germination.

Chemistry was on the side of ACC becoming the ethylene precursor in more ways than one. The cyclopropane ring of ACC is readily broken, with the release of ethylene, by oxidative reactions, such as the Fenton reaction in which hydrogen peroxide formed by the action of Fe(II) on ascorbate acts as the oxidizing agent (McRae *et al.*, 1983). ACC oxidase uses Fe(II) and CO₂ as cofactors and ascorbate as a cosubstrate to catalyse an accelerated and controllable conversion of ACC to ethylene. Additionally, one of the other products generated from ACC by ACC oxidase is cyanide, which may have helped the plant defend itself against attack (Grossman, 1996).

In the preceding we have identified the acquisition of ACC oxidase among the gymnosperm group as being the critical step in the evolutionary development of the biosynthetic pathway of ethylene biosynthesis present in angiosperms (see *c* in Figure 7.2). The production of ethylene from ACC meant that ethylene production now became a signal that was related to the *response* of the plant to the stress, as opposed to (or in addition to) ethylene production as a signal of plant stress *per se*. In angiosperms all three enzymes in the pathway from methionine to ethylene increase activity in response to stress (see Figure 7.2). Similarly in the non-angiosperms, in which ACC is not a precursor of ethylene, the activities of SAM synthetase (Mayne *et al.*, 1996) and ACC synthase (Chernys and Kende, 1996) increase in response to stress. The acquisition of ACC oxidase can thus be viewed as taking ACC from the relative obscurity of a secondary metabolite involved in stress responses, to being a key intermediate in the biosynthesis of a major plant growth regulator.

Two characteristic features of the angiosperm ACC oxidase assayed *in vitro* are: (1) a low specific activity of the purified enzyme compared with other 2-ODDs (John, 1994); and (2) a non-linear reaction rate due to progressive inactivation during catalysis (Smith *et al.*, 1994). It is not known whether these are features of the enzyme activity *in vivo*; they may be attributable to the *in vitro* conditions used for assay. However, it has been noted that in ripening fruits the enzyme is present in large amounts relative to the amount of ethylene formed: in apple, ACC oxidase has been estimated to constitute 5% of the total soluble protein (Abeles *et al.*, 1992). Enzymes involved in secondary metabolism commonly show high abundance where the enzyme has acquired an important function (Pichersky and Gang, 2000). The rationale for this high abundance is that the molecular changes required to increase expression of a protein are simpler and more feasible than those required to increase the rate of substrate turnover or enhance stability (Pichersky and Gang, 2000). Thus we suggest that ACC oxidase, as the 'newest recruit' to the ethylene biosynthesis pathway, compensates for its low catalytic efficiency by a high level of expression.

How ACC oxidase originated

ACC oxidase as a 2-oxoacid-dependent dioxygenase

Having identified the acquisition of ACC oxidase as the critical step in the evolutionary development of the ethylene biosynthetic pathway of angiosperms, we now turn to the evolutionary origin of the enzyme itself. What was the ancestral enzyme? What molecular gymnastics did it undergo to result in ACC oxidase activity?

By similarity of primary sequence and function, ACC oxidase belongs to the non-heme Fe(II), 2-oxoacid dependent dioxygenase (2-ODD) family of enzymes (Prescott and John, 1996). It is to these enzymes that we must turn when considering the possible evolutionary ancestor of ACC oxidase. The 2-ODDs are a large and important class of enzymes that catalyse a wide variety of oxidative reactions, including hydroxylation, desaturation and epoxidation (Prescott, 2000). Known 2-ODDs in plants are involved in a variety of pathways. Commonly, more than one 2-ODD is found in a biosynthetic pathway, notably those involving alkaloids, gibberellins and flavonoids. In *Arabidopsis* Heynh. it is estimated (Prescott, 2000) that there are about 100 genes that code for 2-ODD proteins, but it is only for a minority that any function can be ascribed.

The presence in specific plant groups of 2-ODDs catalysing particular reactions can be assumed on the basis of the presence of particular biochemical products. Flavonoids are present in mosses and higher orders; gibberellins in ferns and higher orders; and anthocyanins in gymnosperms and angiosperms (Koes *et al.*, 1994). On the basis of this, it can be inferred that, as a first approximation, the evolutionary sequence in which 2-ODDs have appeared is: (1) flavonoid enzymes, (2) gibberellin enzymes, (3) anthocyanin enzymes.

Molecular changes

Protein sequence considerations have been used to identify the gross molecular changes required in an evolutionary transition from a 2-ODD ancestor to a functional ACC oxidase. Multiple alignment of all available ACC oxidase sequences revealed that only 30% of the residues are functionally conserved in all sequences examined (Reynolds, 2001). Of the 95 conserved residues, 36 are also conserved in related 2-ODDs (Reynolds, 2001). Thus one can define ACC oxidase at a primary sequence level. However, these conserved sequences are more usefully interpreted in a functional context. In its enzymatic action, ACC oxidase differs from other 2-ODDs not only in its primary substrate specificity, but also in using ascorbate as a cosubstrate (rather than 2-oxoglutarate) and in using CO₂ as an essential cofactor (see Figure 7.3). To account for these unique functional features of ACC oxidase, the protein alignments must be viewed in terms of protein structure. The 3-D structure of ACC oxidase is unknown, but a structure for the microbial 2-ODD, isopenicillin *N* synthase (IPNS) is available (Roach *et al.*, 1995). There is sufficient sequence homology between IPNS and ACC oxidase for the IPNS structure to provide a basis for structural features of the ACC oxidase protein to be recognized (Roach *et al.*, 1995).

A summary of secondary structure predictions for ACC oxidase is shown in Figure 7.4. The most extensive region of conservation in ACC oxidase is in the centre of the protein from Pro¹⁶⁹ to Pro²⁵⁵ (Figure 7.4). This region also shares most similarity with other 2-ODDs. Based on homology with IPNS, this domain probably constitutes the jellyroll motif (Stuart, 1993) of eight β -sheets, containing the active site of the protein. Conserved residues within this catalytic domain will provide the binding sites for Fe(II), oxygen, ACC and probably ascorbate (Roach *et al.*, 1995).

The region of ACC oxidase that diverges most from other 2-ODDs is the C-terminus (see Figure 7.4). This region can therefore be considered diagnostic for ACC oxidases compared with other 2-ODDs. The C-terminal region beginning at Tyr²⁸¹ contains 14 conserved residues, none of which is conserved in related 2-ODDs. Secondary structure prediction indicates that the C-terminal part of the protein forms a domain consisting of two α -helices connected to the body of the protein by a loop section. This loop section is a poorly conserved region of ACC oxidase which can be up to 16 residues in length.

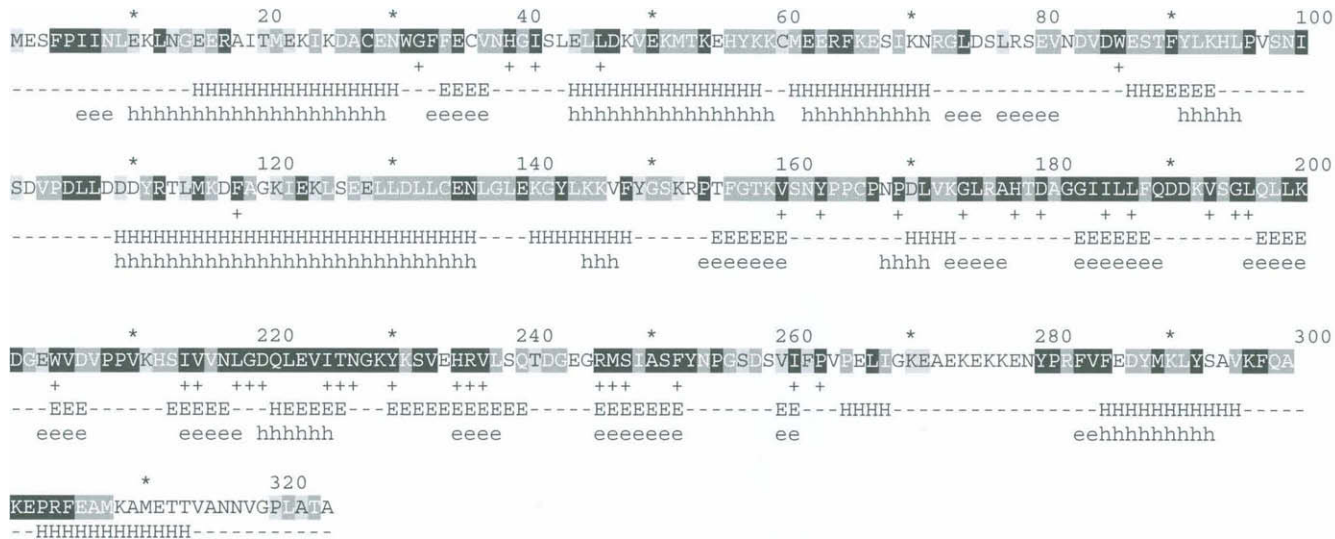


Figure 7.4 Summary of analysis of all available ACC oxidase protein sequences. The protein sequence of ACC oxidase from *Arabidopsis thaliana* L. is shown (Accession number AAC97998) shaded according to the multiple alignment of 65 ACC oxidase protein sequences. Residues shaded black are functionally conserved in all sequences, those shaded dark grey are functionally conserved in >80% of the sequences and light grey >60%. The + beneath the protein sequence indicates residues that are conserved in all 2-ODDs. The secondary structure prediction is given, where H/h = helix and E/e = sheet, - indicates no predicted structure. The line in upper case is a summary of predictions made using SOPM and SSPro and the line in lower case is the structure inferred by alignment with the related microbial protein of known structure, isopenicillin N-synthase (IPNS). (From Reynolds, 2001.)

The high degree of sequence conservation in the C-terminus region of ACC oxidase suggests that it is important functionally. ACC oxidase is unique among 2-ODDs in its requirement for CO₂ as a cofactor. The C-terminus region of ACC oxidase is rich in positively charged residues, such as arginine and lysine, which are potential binding sites for CO₂. Site-directed mutagenesis of ACC oxidase has provided evidence for the location of the CO₂ binding site at a conserved arginine residue in the C-terminus region: Arg³⁰⁰ in ACC oxidase from kiwi fruit (Lay *et al.*, 1996); Arg²⁹⁹ in ACC oxidase from apple fruit (Kadyrzhanova *et al.*, 1999). Thus we propose that acquisition of the C-terminal domain, by an ancestral 2-ODD was crucial in the step change in catalytic function to ACC oxidase, and it coincided with the requirement for CO₂ as a cofactor.

In the evolutionary acquisition of ACC oxidase activity, the ability to bind and turn over the novel substrate ACC was acquired, but also, we assume that ascorbate took the place of 2-oxoglutarate as cosubstrate. This latter assumption is strengthened by the finding (Iturriagagoitia-Bueno *et al.*, 1997) that 2-oxoglutarate, and other 2-oxo acids, inhibit ACC oxidase activity competitively with respect to ascorbate. Fe(II) chelation was excluded as an explanation of the action of 2-oxoglutarate, and it was concluded that ascorbate and 2-oxoglutarate were competing for the same binding site (Iturriagagoitia-Bueno *et al.*, 1997).

The enzyme ancestral to ACC oxidase

Prescott (2000) has created a phylogenetic tree of 64 2-ODDs from *Arabidopsis* (Figure 7.5). In this phylogeny, the three *Arabidopsis* ACC oxidases appear as terminal branches of a subgroup of five proteins, the other members of which are uncharacterized. The proteins of known function most closely related to ACC oxidases are those of the flavonoid pathway. They, and the ACC oxidases, appear to diverge from a common origin (Figure 7.5). On this basis we suggest that ACC oxidases evolved from an ancestral 2-ODD that was involved in flavonoid biosynthesis.

Flavonoids and ACC do not appear to have much in common chemically and further information is needed before our suggestion can be substantiated. However, in support of our suggestion, we note that 2-ODDs can show a relatively loose substrate specificity (Prescott and John, 1996; Prescott, 2000). This substrate flexibility is shown by the 2-ODDs in alkaloid, antibiotic and gibberellin synthetic pathways catalysing more than one reaction; ACC oxidase itself can convert *D*-valine to *iso*-butanal (Gibson *et al.*, 1998). We envisage that the substrate flexibility of 2-ODDs played a part in the early stages of the evolutionary development of the ACC oxidase.

When did the ethylene biosynthesis pathway arise?

Palaeoclimatic considerations

Although seed plants may have first appeared during the Devonian period, it was during the relatively dry Jurassic period that they diversified extensively and began to dominate (Willis and McElwain, 2002). Ethylene diffuses 10⁴ times faster in air than in water (Musgrave *et al.*, 1972). Thus, in general, ethylene would have been a more useful signalling compound after the end of the relatively wet Devonian and Carboniferous periods, as suggested by Osborne *et al.* (1996). It is pertinent to note that the submerged aquatic angiosperm *Potamogeton pectinatus* L. is incapable of producing ethylene because it lacks

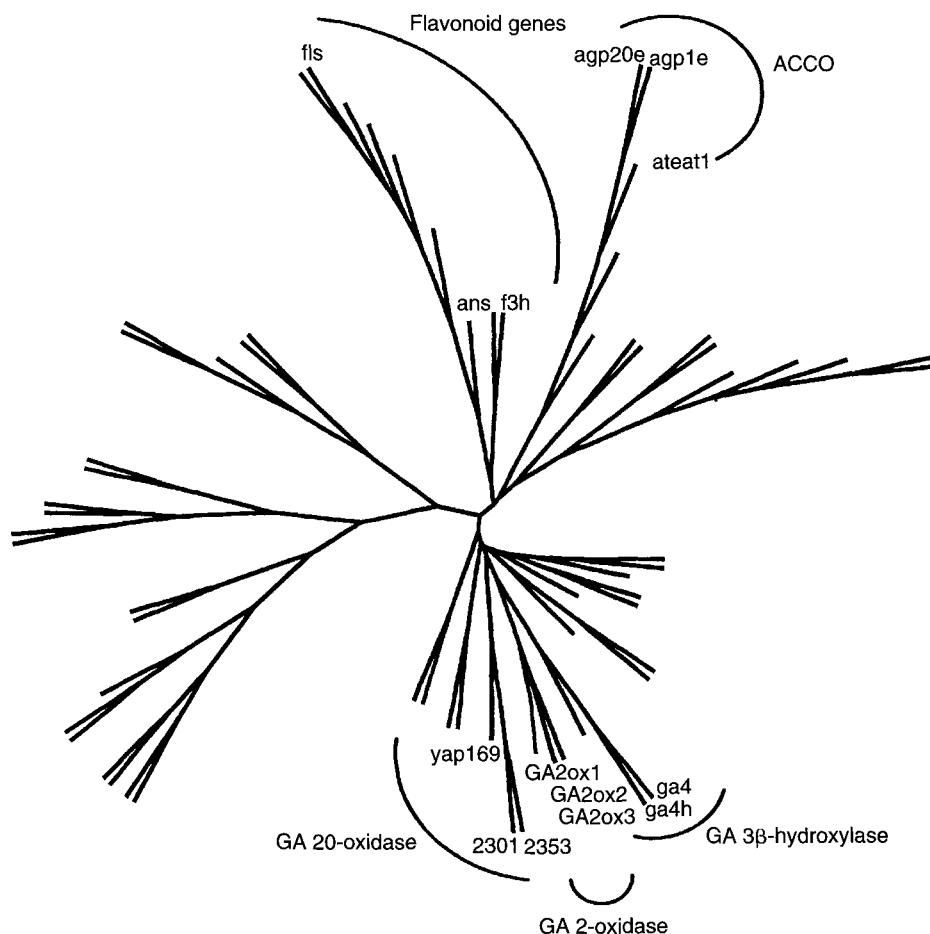


Figure 7.5 Phylogenetic tree of 2-oxoacid-dependent dioxygenases from the *Arabidopsis* genome. Each branch represents a single gene, the branches containing genes for ACC oxidases and for the enzymes involved in the syntheses of gibberellins and flavonoids are highlighted (Prescott, 2000).

ACC oxidase (Summers *et al.*, 1996). This secondary loss of ACC oxidase is likely to be related to the lack of a signalling function of ethylene in the aquatic environment of *P. pectinatus*.

In addition to ACC oxidase, the 2-ODD enzymes involved in anthocyanin biosynthesis also have their origin in the gymnosperms (Koes *et al.*, 1994). So it may have been that the Devonian period was one of divergence in the 2-ODD family of enzymes as a whole, with ACC oxidase activity representing one of many new catalytic activities among the 2-ODD enzyme family that arose in that period.

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8

Structural biomacromolecules in plants: what can be learnt from the fossil record?

Pim F van Bergen, Peter Blokker, Margaret E Collinson,
Jaap S Sinninghe Damsté and Jan W de Leeuw

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Introduction

When plants entered land they needed specific physiological adaptations to survive this new, hostile environment. Some of the main problems plants had to overcome included water loss or desiccation and exposure to increased UV radiation (Edwards, 2001). An increase in phenolic constituents in various plant tissues is believed to be one of the main adaptations to counteract the problem with UV radiation (Rozema *et al.*, 2001). With respect to desiccation, two important adaptations are suggested, i.e. (1) increased protection of the outer covering of the plant to prevent direct water loss and (2) enhanced water transport within the plant. The first aspect is directly related to the development of the cuticle on leaves and stems and resistant walls on the plant's reproductive tissues (i.e. spores), while the second relates to the formation of water-conducting elements such as tracheids and vessels.

Evidence for the evolution of these distinct structures is based primarily on morphological observations of plant remains dating as far back as the Ordovician for cuticles and distinct spore material (i.e. tetrads, dyads and monads) and the Middle Silurian for tracheids (Edwards, 2001 and references cited therein). In terms of evolution, it has been

suggested that spore walls may have evolved from algal cell walls (Hemsley, 1994), whereas the direct origin of the cuticle and tracheids is still unknown (Edwards, 2001).

Apart from evidence for the morphological evolution of these entities, many studies link microscopic observations directly to biochemical evolution in these structures enabling plants to colonize land. Commonly, the presence of fossil cuticles is suggested to be directly related to the occurrence of the biopolyester cutin (e.g. Edwards, 2001; Cooper-Driver, 2001), whereas the findings of fossil water-conducting tissues such as tracheids in xylem are often indirectly related to the biomacromolecule lignin (e.g. Edwards *et al.*, 1997; Edwards, 2001; Cooper-Driver, 2001). In addition, acid resistant fossil spore wall material is suggested mostly to be composed of or impregnated by the, yet chemically poorly defined, biomacromolecule sporopollenin (Cooper-Driver, 2001; Edwards, 2001). Additional resistant macromolecular compounds mentioned in connection with fossil plant remains are cutan in cuticles and algaenan in algal cell walls.

Despite the circumstantial evidence that the chemical compositions of fossil plant remains were very similar to those occurring in modern plant cuticles, spore and pollen walls or xylem elements, to date no *unequivocal* molecular chemical evidence exists proving that this was actually the case. The search for such evidence is complicated by a number of aspects including: (1) the chemically resistant nature of these, mainly macromolecular, entities in modern plants; (2) the intimate association of these molecules with other, often more abundant, compounds preventing purification of the specific biomacromolecules and thus hampering the elucidation of the detailed molecular composition of the extant counterparts (i.e. sporopollenin); (3) the biological and chemical transformation reactions occurring upon burial and subsequent fossilization (i.e. diagenesis); and (4) the inconsistent use of nomenclature for the various resistant macromolecules. Note that the term 'resistant macromolecules' refers to those compounds that are resistant against normal chemical treatments but are destroyed by chemical oxidation. This chemical resistance is believed to be the main underlying reason for their high preservation potential under most, but not all, depositional settings.

The following terms will be used to avoid confusion: *algaenans* for the chemically resistant macromolecules present in algal cell walls (including the walls of algal zygospores); *sporopollenin* is the chemically resistant macromolecule in spore and pollen walls of non-algae excluding fungal spores; *cutan* is the highly aliphatic macromolecule recognized in modern and fossil plant cuticles; and *lignin* will only be used for the ether-linked macromolecule based on monolignols, i.e. *p*-coumaryl alcohol, coniferyl alcohol and sinapyl alcohol. Terms such as 'lignin-like' and 'sporopollenin-like' should be avoided.

In this chapter we will first describe the main chemical methods that can provide detailed molecular insight into the chemical composition of both extant and fossil resistant (bio)macromolecules. Subsequently, the macromolecular composition of modern and fossil outer coverings, including algal cell walls, spore and pollen walls and cuticular tissues will be discussed both in terms of their chemical similarities and differences and in relation to the physiological adaptations and evolution of land plants. Finally, the chemical composition of water-conduction tissues of modern and fossil land plants will be evaluated also in terms of their physiology but with specific emphasis on the biomacromolecule lignin.

Characterizing resistant biomacromolecules

In order to study chemically resistant fossil (bio)macromolecules one has to have detailed molecular insight into their modern counterparts. In most cases, the modern counterparts

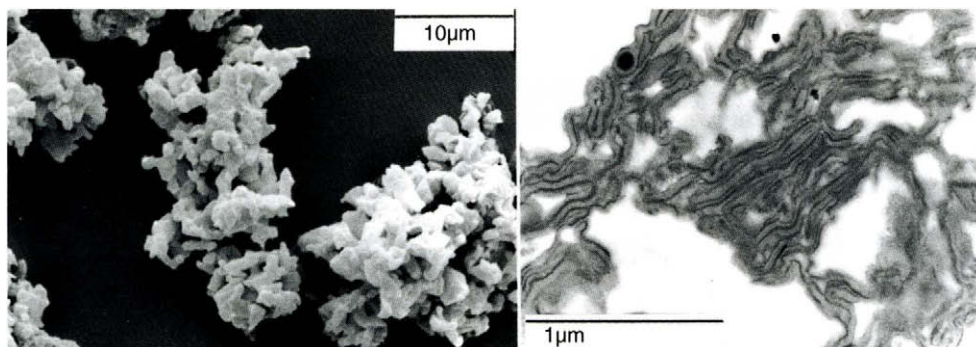


Figure 8.1 Example of purified algal cell walls of modern *Tetraedron minimum* obtained by Potter homogenization. For chemical analyses of this sample the reader is referred to Blokker *et al.* (1998).

can only be isolated by purification steps involving both physical and chemical separation methods. Mechanical stripping of cuticular membranes from leaf material (Mösle *et al.*, 1997) and ultrasonic destruction to obtain purified algal cell wall material (Potter homogenization; Blokker *et al.*, 1998; Figure 8.1) are effective physical pretreatments.

Numerous chemical extraction methods exist to obtain the most resistant organic molecules in plant remains. Although it should be emphasized that the organic remains recognized in the fossil record are not necessarily identical to the resistant material that one obtains after purification and extraction steps under experimental conditions. Normally a full extraction method involves the following steps:

1. solvent extraction to eliminate low molecular weight lipids
2. hydrolysis, either with base and/or acid, to remove ester-bound/amide-bound moieties; this treatment often consists of two sequential steps, a base hydrolysis and an acid hydrolysis
3. acid treatment using H_2SO_4 to remove polysaccharides
4. an ultimate base treatment to remove moieties that have become accessible after the previous steps (*cf.* the results of cutan isolation from *Agave* L. in Tegelaar *et al.* (1989: Figure 17) and the same sample after this final base treatment in de Leeuw *et al.* (1991: Figure 6) revealing unequivocally that the polysaccharides are *not* part of the resistant molecule in cuticles!).

However, it should be noted that some chemical pretreatments that are used to mimic fossilization processes (i.e. acetolysis) might drastically affect the chemical composition of extant material under examination. A prime example in this respect relates to the study of sporopollenin whereby the most commonly used treatment involves acetolysis (Graham and Gray, 2001; Hemsley *et al.*, 1996). Although the remains obtained after this treatment are morphologically identical to those found in the fossil record, detailed molecular information based on ^{13}C NMR revealed that the chemical composition of sporopollenin from pine pollen was severely altered (Hemsley *et al.*, 1996). Furthermore, Allard *et al.* (1997) showed that newly formed condensation products (i.e. melanoidin-like polymers derived from cell wall polysaccharides), that are not related to the resistant entities originally present, can easily be produced. Appropriate modified isolation procedures can avoid such problems (Allard *et al.*, 1998).

The various chemical methods that are used to study the resistant biomacromolecules can be subdivided into non-destructive and destructive techniques. The former includes solid state ^{13}C NMR and FTIR, both of which reveal information pertaining to the various carbon environments present (e.g. Hatcher *et al.*, 1999) and have revealed very informative data on lignin, cutan, algaenan and sporopollenin. Pyrolysis and chemolysis are both destructive methods that will reveal detailed molecular information about the specific building blocks (e.g. van Bergen, 1999). Pyrolysis is often used as an effective initial screening method as it reveals qualitative insights into the overall chemical composition of the macromolecule. Pyrolysis has been used on most resistant biomacromolecules (i.e. lignin, cutan, algaenan, sporopollenin). For further information about the various pyrolysis methods, the reader is referred to Stankiewicz *et al.* (1998) and van Bergen (1999). The advantage of chemolysis is that one can target specific molecular building blocks using specific chemical reagents. Chemolysis is often used in combination with pyrolysis as the latter reveals characteristic moieties that subsequently can be examined using appropriate reagents (van Bergen *et al.*, 1994a; Gelin *et al.*, 1997; Blokker *et al.*, 1998). The main specific reagents or methods used nowadays include treatment with CuO , RuO_4 , HI, KMnO_4 or thioacidolysis. For lignin the most commonly used methods include treatment with CuO and, more recently, thioacidolysis, for cutan RuO_4 , for algaenan RuO_4 and HI and for sporopollenin KMnO_4 , NO_x .

As both the destructive and non-destructive methods can yield biased results, a combination of these techniques should be used when possible, as this will provide greater insight into the composition of the biomacromolecule under investigation. Furthermore, it is always very instructive to undertake microscopic examinations prior to and after chemical treatments (Boon *et al.*, 1989; van Bergen *et al.*, 1995; Blokker *et al.*, 2001; Collinson *et al.*, 1998) as this can reveal where the actual resistant organic material resides in the plant organ. Also histochemical staining can reveal specific insight into the tissues that are preserved containing resistant macromolecules (van Bergen, 1994). However, a colourimetric method in itself does not *prove* the presence of one characteristic molecule as was shown in a study on modern and fossil chitin in arthropod cuticles (Bierstedt *et al.*, 1998). In particular with respect to fossil material specific histochemical colours cannot be used as chemical evidence for the presence of a certain macromolecule.

Resistant biomacromolecules in outer coverings

Algal cell walls

The abundant contribution of algal cell walls in many palynological microspore assemblages and coals (i.e. *Botryococcus* Kützing in boghead) has always been the main reason to suggest that these structures in the modern counterparts should contain a specific chemically resistant material. Subsequent chemical investigations, focusing initially on modern *Botryococcus* (Largeau *et al.*, 1986) and modern *Tetraedron* Kützing (Goth *et al.*, 1988), did indeed reveal the presence of distinct types of biomacromolecules nowadays called algaenans.

Recent studies on a number of modern freshwater and marine microalgae have shown that some, but not all, contain a resistant molecule in their cell walls (Gelin *et al.*, 1999; Blokker, 2000). To date, all algaenans obtained from modern algae are based on, so-called, aliphatic building blocks. However, large differences do exist. The algaenan obtained from *Botryococcus* is very different from those of all other algae studied to date (i.e. based on

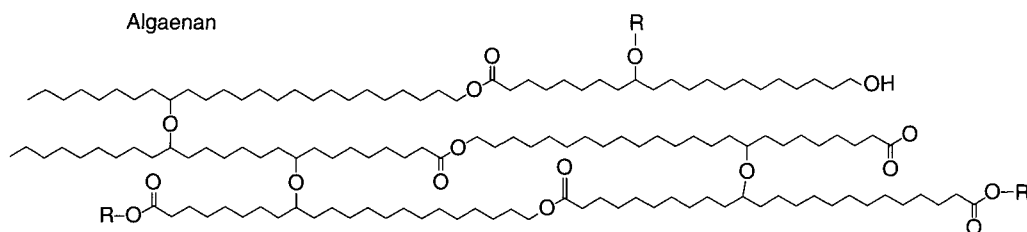


Figure 8.2 Simplified structure of an algaenan based on hydroxy and dihydroxy fatty acid units (modified after Blokker *et al.*, 1999).

ether-linked C_{32} dialdehyde monomers; Berthéas *et al.*, 1999). Detailed molecular data have shown that many biomacromolecule algaenans are based on long-chain (C_{36} and longer) ester- and ether-linked hydroxy fatty acids (Figure 8.2; *cf.* Blokker *et al.*, 1999). The building blocks of these biomacromolecules are connected by both ester and ether cross-links explaining why simple hydrolysis does not release the individual monomeric building blocks. The highly aliphatic nature of these compounds would provide an excellent hydrophobic outer covering preventing simple water transport across such a plant structure. With the exception of one case (see below), no molecular evidence exists of a contribution of aromatic or phenolic moieties to algaenans of modern algae.

Exceptions to the above are the finding of aromatic moieties in the algaenan of *Chlorella marina* (Derenne *et al.*, 1996) and in resistant material obtained from walls of the resting cysts of the dinoflagellate *Lingulodinium* (Stein) Dodge (Kokinos *et al.*, 1998). However, in contrast to the numerous studies of other algae, the resistant cell walls of dinoflagellates have rarely been studied using the detailed molecular approach outlined above and thus the extent to which one can generalize the dinoflagellate results is yet unknown. Current unpublished results have as yet not been able to verify the contribution of aromatic moieties in either the same species or in any others (van Mourik *et al.*, unpublished data).

One intriguing observation relates to the detection of 10,16-dihydroxy- C_{16} fatty acid as part of the resistant algaenan in the cell walls of the extant Zygnematalean alga, *Spirogyra* Link (Blokker, 2000). This compound is commonly found as an abundant building block in the biopolyester cutin present in higher land plant cuticles (Holloway, 1982; see below). In this respect it is interesting that the Zygnematalean algae have been suggested to have evolved from organisms that first colonized the land, after which they returned to grow under aquatic conditions (e.g. Stebbins and Hill, 1980; van Geel and Grenfell, 1996). If this were proved to be true then the presence of 10,16-dihydroxy- C_{16} fatty acid as a structural building block in outer coverings may be an adaptation to life on land.

Chemical evidence of algaenans in fossil algal cell walls is based mainly on data from microalgae preserved as ultralaminae (Derenne *et al.*, 1991, 1992a) and well-preserved cell wall material of *Tetraedron* (Goth *et al.*, 1988) and *Botryococcus* (Largeau *et al.*, 1986; Gatellier *et al.*, 1993). In these cases the fossil macromolecules are still very similar to those recognized in their modern counterparts. However, unequivocal taxon-specific data only relates to fossil material found in the Tertiary sediments. Various older samples have been studied, in particular *Pelta* (= *Botryococcus*), *Tasmanites* Newton etc., but apart from revealing macromolecules that are highly aliphatic in nature no specific chemical characteristics are known that prove that these represent original or only slightly modified algaenans. Thus, chemical evidence of Palaeozoic fossil algae has to be interpreted with caution in

relation to biochemical and physiological evolution. Even Tertiary material that is composed of rich monotypic algal cell wall material can be altered or interpreted incorrectly due to additional non-algal organic material that overprints the chemical signal. For example, chemical data from Late Miocene lacustrine oil shales containing up to 80% *Pediastrum* Meyen were used to suggest that the algaenan of this alga was mainly aromatic in nature (Sinninghe Damsté *et al.*, 1993). Subsequent analyses of modern *Pediastrum* cell walls revealed that the algaenan is based on C₃₀ and C₃₂ ω -hydroxy fatty acids clearly revealing its aliphatic nature similar to that of other Chlorococcales (Blokker *et al.*, 1998).

Pyrolysis data from a number of Tertiary, Palaeocene and Eocene, dinoflagellate cyst walls have revealed that these contain mainly aliphatic building blocks, although samples that appear to be more mature, based on colour, reveal an increase of aromatic pyrolysis products (Warnaar *et al.*, unpublished results). However, as no chemical data from modern counterparts are available yet, these data and, in particular, the significance of an aromatic contribution to algaenans, cannot be interpreted in terms of their original chemical composition or physiological differences.

Another distinct microfossil, without a direct modern counterpart, is *Gloeocapsomorpha prisca* Zalesky. The fossil remains of this organism constitute the bulk of the organic matter in Ordovician Kukersites (Foster *et al.*, 1989, 1990). Transmission electron microscopy revealed that the walls are composed of multiple sheets surrounding a void (Derenne *et al.*, 1992b; Blokker *et al.*, 2001). This distinct morphology, in combination with chemical data, means that the precise biological affinity of this organism is still unknown, although an algal or a bacterial affinity appears most likely (see Blokker *et al.*, 2001 and references cited therein). Based on the extremely low maturity of the Kukersite rocks it is believed that the organic matter preserved has not been altered significantly, which is an essential prerequisite for further meaningful interpretations. The building blocks of the resistant macromolecule present in the fossil remains of this organism are based on 5-*n*-alkyl-resorcinols with the alkyl side chain being mainly C₁₅, C₁₇ and C₁₉ (Derenne *et al.*, 1992b; Blokker *et al.*, 2001). If this organism were a true alga then this would be the first algaenan type containing 1,3-benzenediol (=resorcinol) building blocks (Metzger and Largeau, 1994). But if its closest biological affinity lies with the bacteria then the benzenediol contribution would be less surprising as these are known to produce resorcinols upon stress (Kozubek and Tyman, 1999). From a physiological point of view, however, the biological affinity is of minor importance as geological information has suggested that these organisms lived in a high UV light lagoonal or very near-shore environment (Foster *et al.*, 1990). This would probably induce the production of phenolic, i.e. resorcinol, moieties to counteract possible UV-induced damage.

Overall, the current molecular data on modern algaenans show primarily biomacromolecules based on long-chain (\geq C₂₆) hydrophobic moieties that most probably provide water repellent properties to the cell walls of these algae. Phenolic entities are not detected, but then one would not expect them in these aquatic organisms if their primary function was related to counteract high UV radiation levels.

Pollen and spore walls

In stark contrast to the algal material, spore and pollen walls are not protected constantly by an aqueous environment and thus will have been adapted to minimize water loss and to reduce UV damage. Studies of the chemical composition of the resistant material present in spore and pollen walls have a long history because acid resistant pollen and spores are

present in coals and other sedimentary rocks, dating as far back as the Ordovician (Edwards, 2001). This fossil record is believed to be due to the presence of a highly resistant macromolecule, namely sporopollenin. The term 'sporopollenin' was first used by Zetzsche and Kälin (1931) to represent the structural constituent of the outer walls (exines) of pollen and spores from vascular plants that is resistant to non-oxidative chemical treatments. As stated above, here we will use this definition of sporopollenin, excluding other remains such as algal and fungal spores, as well as avoiding the term 'sporopollenin-like'. However, we do include spore wall material from *Lycopodium clavatum* L. which has previously been mistaken for a marine alga (Bestougeff *et al.*, 1985).

Although earlier chemical studies had been undertaken (see Brooks and Shaw, 1978 for a review), it was Zetzsche and co-workers who started to analyse sporopollenin systematically (e.g. Zetzsche and Kälin, 1931; Zetzsche *et al.*, 1937). Using mainly oxidative techniques and elemental analyses, they concluded that all sporopollenins could be represented by a general empirical formula which they chose to formulate arbitrarily on a C₉₀ basis. Although no precise molecular structure was postulated, Zetzsche *et al.* (1937) assigned a polyterpenoid structure to sporopollenin.

Following this early work it was not until the 1960s that, in particular, Shaw and Brooks started to re-examine sporopollenin using, among a variety of techniques, FTIR and ozonolysis as their main methods (e.g. Shaw and Yeadon, 1964; Brooks and Shaw, 1968, 1978). Based on their results, they postulated that the structural building units of sporopollenin were carotenoids and/or carotenoid esters. In particular, the similarities observed in FTIR spectra of a variety of samples, ranging from fossil and extant spores and pollen to algal cysts (*Tasmanites*) and meteorites, were thought to reflect the omnipresence of carotenoid-composed sporopollenin. However, subsequent pyrolysis data from the algal cyst *Tasmanites* showed that the structural building units of these cysts are, primarily, linear long-chain aliphatics in addition to some terpenoids (Philip *et al.*, 1982). No evidence of carotenoids was found, thus questioning the inference of the presence of carotenoids from the FTIR spectra. Moreover, it is interesting to note that similar FTIR spectra have been interpreted as indicative of aliphatic (Hayatsu *et al.*, 1988) or emphasizing aromatic compounds (Schulze Osthoff and Wiermann, 1987) depending on the authors.

One of the possible reasons for Brooks and Shaw to postulate carotenoids as major constituents of sporopollenin might have been the fact that they expected the sporopollenin precursor to be a lipid abundantly present during the development of the pollen or spore wall (*cf.* Shaw, 1970, 1971). The finding of large amounts of carotenoids with, at that time, no apparent biological function (Shaw, 1970, 1971), may have led to the suggestion that these compounds were the precursors of sporopollenin. Nowadays, however, carotenoids are known to be involved in light harvesting and/or protection against photo-oxidation (e.g. Wiermann and Gubutz, 1992 and references cited therein). Moreover, subsequent studies of sporopollenin indicate that carotenoids and/or carotenoid esters do not contribute at all to sporopollenins and that their recognition in these earlier studies must be ascribed to an incomplete purification of sporopollenin isolates (e.g. incomplete removal of non-resistant material; for reviews see Wiermann and Gubutz, 1992 and de Leeuw and Largeau, 1993).

Studies using solid state ¹³C NMR, FTIR, oxidation and mild degradation techniques, tracer experiments and pyrolysis implied the presence of two chemically different types of sporopollenin. In one type the main building blocks are oxygenated aromatics (e.g. Schenck *et al.*, 1981; Schulze Osthoff and Wiermann, 1987; Herminghaus *et al.*, 1988; Wehling *et al.*, 1989; Mulder *et al.*, 1992), while in the other, the main building units are thought to

be predominately aliphatic in nature (e.g. Guilford *et al.*, 1988; Hayatsu *et al.*, 1988). It is interesting to note that in most of these studies evidence for both aliphatic and aromatic constituents can be found. Combining all these results from modern spore and pollen wall material reveals that cinnamic acids, such as *p*-coumaric acid and ferulic acid, are major aromatic units present in modern sporopollenin (see Figure 8.3a). These phenolic units are both ester- and ether-linked (Mulder *et al.*, 1992) within the macromolecular structure explaining why simple base hydrolysis will not remove all these cinnamic acids.

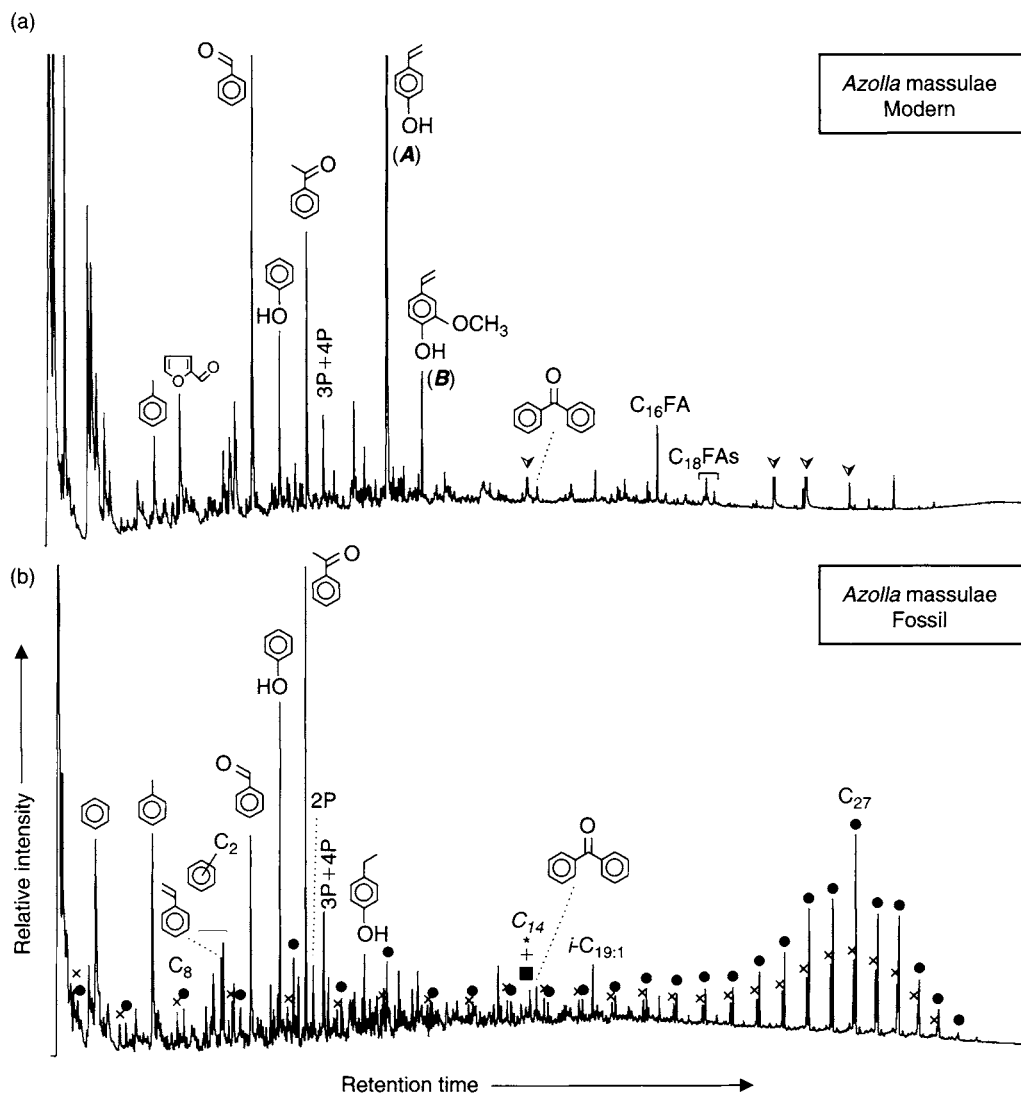


Figure 8.3 Gas chromatograms of pyrolysates (Curie-temperature 610°C) of sporopollenin from water ferns: (a) microspore massulae of modern *Azolla* Lam. (*cf.* van Bergen *et al.*, 1993) and (b) microspore massulae of fossil, K/T boundary, *Azolla*. The abundance of cinnamic acids in modern sporopollenin is evident from 4-vinylphenol (A) which is the main pyrolysis product derived from *p*-coumaric acid, while 4-vinyl-2-methoxyphenol (B) is the major pyrolysis product released from ferulic acid. Key: ● = *n*-alkanes, × = *n*-alk-1-enes, * = C₁₄-5-alkanone, ■ = C₁₄-6-alkanone, ▼ = contaminant. C₂₇ = heptacos-1-ene and heptacosane, 2P = 2-methylphenol, 3P + 4P = co-eluting 3- and 4-methylphenol, *i*-C_{19:1} = prist-1-ene, FA = fatty acid. For additional information regarding the samples the reader is referred to van Bergen *et al.* (1993).

With respect to fossil material, relatively few papers have been published on well-characterized spore and/or pollen wall material (e.g. Shaw, 1970; Brooks, 1971; Hemsley *et al.*, 1992, 1993, 1996; van Bergen *et al.*, 1993; Collinson *et al.*, 1994). Similar to the data from the modern samples, a combination of both aliphatic and aromatic, mainly phenolic, moieties were found. However, pyrolysis results of both modern and fossil well-preserved spore material of water ferns showed that aliphatic moieties became enriched upon fossilization but evidence of cinnamic acids as part of the resistant spore wall material remained (Figure 8.3; van Bergen *et al.*, 1993). Based on these early results in combination with pyrolysis data from fossil pollen clusters, van Bergen *et al.* (1995) suggested that sporopollenin is composed of both long-chain aliphatic and oxygenated aromatic (mainly phenolic) moieties. Subsequent solid state ^{13}C NMR (Figure 8.4) and RuO_4 data, in combination with the pyrolysis results of both spore and pollen wall material, have led to a tentative structure for sporopollenin in which long-chain ($\text{C}_{24}\text{--C}_{28}$) highly aliphatic units form the backbone of the sporopollenin and the cinnamic acids are the cross-linking units (Figure 8.5). In fossil sporopollenin the amount of cinnamic acids will have decreased substantially leading to a more aliphatic macromolecule. This may also explain some of the chemical data related to sporinities, the coal maceral which consists of fossil spore and pollen walls, showing a dominance of aliphatic moieties (Hayatsu *et al.*, 1988; Davis *et al.*, 1988; Kruge *et al.*, 1991). However, some pyrolysis data of sporinities showed phenols to be dominant, while aliphatic pyrolysis products were relatively less abundant

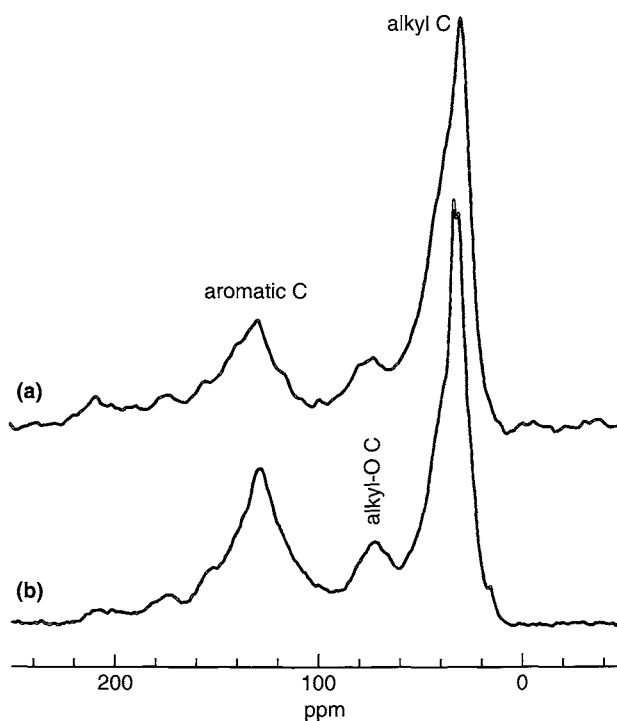


Figure 8.4 Solid state ^{13}C NMR spectra of fossil (K/T boundary) sporopollenin of (a) pollen clusters of the extinct angiosperm *Kurtzipites* Anderson and (b) microspore massulae of *Azolla* both obtained from the same bed. Both spectra are similar and reveal strong aliphatic carbons at 32 ppm, in addition to significant amounts of alkyl-O carbons at 71 ppm and aromatic carbons at 127 and 151 ppm. The pyrolysate of the *Azolla* sample is shown in Figure 8.3b. For additional information regarding the samples the reader is referred to van Bergen (1994).

(Nip *et al.*, 1988, 1992). Moreover, pyrolysates of handpicked Carboniferous microspores and megaspores also showed the abundant presence of simple aromatics and phenols (Collinson *et al.*, 1994). However, these phenols showed no direct structural link with cinnamic acids, making inferences in terms of the chemical composition of fossil sporopollenin somewhat hypothetical.

From a physiological point of view, the combination of a macromolecule based on a hydrophobic long-chain aliphatic backbone, that could provide water repellent properties, and cinnamic acid units, which could provide UV protection, would be ideal. Comparing the sporopollenin macromolecule with algaenan shows that both contain an aliphatic backbone which can be interpreted as evidence in favour of the evolutionary trend whereby sporopollenin is a modified algaenan. However, there are two significant differences, apart from the incorporation of cinnamic acid moieties into the structure of sporopollenin. First, the carbon chain length of the algaenans is longer than that of the sporopollenin and secondly the aliphatic chains in algaenans are generally based on hydroxy fatty acids, whereas those in sporopollenin appear to be based solely on ether-linked alkyl units. The difference in chain length is rather difficult to explain, although as yet it is not known whether the alkyl chains suggested for the water fern sporopollenin can be generalized. Maybe shorter chain alkyl units are an adaptation of life on land since the abundant biopolyester in cuticles, cutin, is based on much shorter alkyl groups (mainly C₁₆-units, see later). Shorter alkyl units may possibly have provided enough hydrophobicity in these structures to withstand desiccation. Reducing the amount of carbon needed to build these structural units that, under normal circumstances will not be used again by the plant, would be a major advantage in terms of energy.

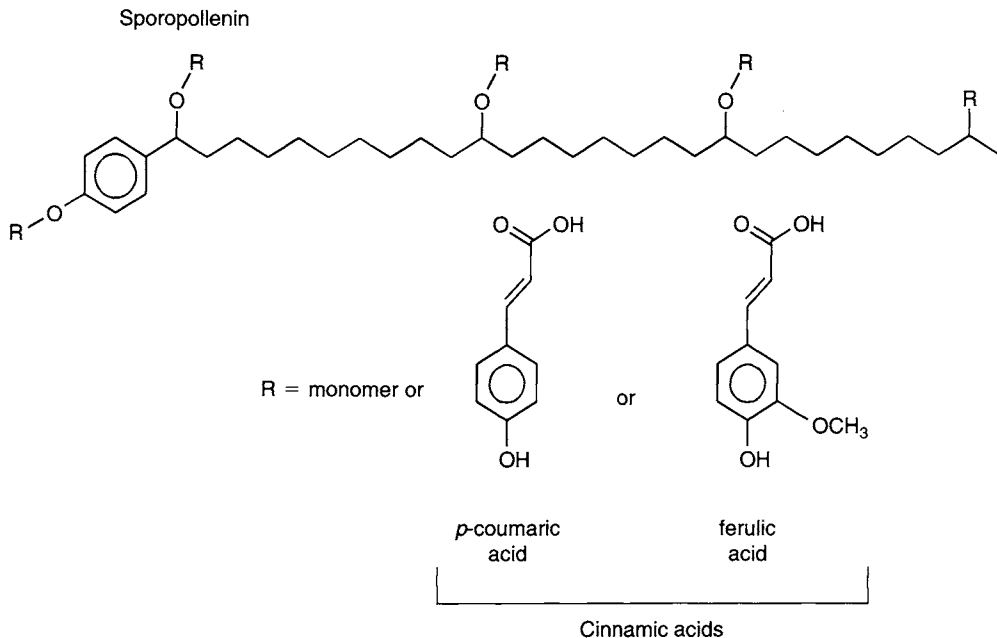


Figure 8.5 Tentative and simplified structure of a sporopollenin monomeric building block. The repeating units (R) can either be linked by ether bonds (C–O–C) or in case of the two acids also through ester bonds (C–O–CO–C).

Higher land plant leaf and stem cuticles

The bulk of the cuticle in modern plants is composed of a solvent-insoluble matrix (Tegelaar *et al.*, 1991). This matrix makes up the framework of the cuticle and is composed either of the hydrolysable biopolyester cutin, an insoluble non-hydrolysable macromolecule, named cutan, or (most commonly) a mixture of both (Tegelaar *et al.*, 1991).

The chemical structure of cutin in modern cuticles from both angiosperms and gymnosperms (Holloway, 1982; Collinson *et al.*, 1998) is well understood and is based primarily on ester-linked functionalized, mainly hydroxy- and dihydroxy-, C_{16} and C_{18} alkanolic acids (Figure 8.6; Holloway, 1982; Tegelaar *et al.*, 1991). Major building blocks include $10,\omega$ -dihydroxy C_{16} or C_{18} alkanolic acids and $9,10,\omega$ -trihydroxy C_{18} alkanolic acid, although the cutin is normally based either mainly on C_{16} or C_{18} units. In addition to the linear aliphatic moieties, the cutin fraction of some taxa also reveals the presence of cinnamic acids, in particular *p*-coumaric and ferulic acid (Kolattukudy, 1981; Holloway, 1982). Holloway (1982) however, stated that phenolic acids esterified to carbohydrates, rather than to cutin alkanolic acids, first have to be eliminated as a possible source for cinnamic acids in cutin. In particular, incorporation of cell wall material to the cuticle might lead to the apparent recognition of cinnamic acids as an integral part of cutin. Cinnamic acids have been suggested to be part of the cutin of only a small number of plants including *Lycopersicon esculentum* Mill. and *Ginkgo biloba* Linn. (Collinson *et al.*, 1998).

Unequivocal molecular evidence of cutin in fossil cuticles is sparse (Tegelaar *et al.*, 1991) as it would need to show the presence of the distinct C_{16} and C_{18} alkanolic acids; upon pyrolysis they occur as unsaturated alkanolic acids. In the few fossil cases where cutin was suggested, cinnamic acids were not recognized (Tegelaar *et al.*, 1991). In stark contrast, most fossil cuticles consist of macromolecular material composed of long-chain aliphatic compounds, termed cutan. The discovery of this material in fossil cuticles triggered a search to identify it in modern cuticles.

To date, the search for cutan in modern taxa has shown that only some modern cuticles reveal unequivocal evidence of this macromolecule (e.g. *Agave*, *Clivia* Lindl.). Moreover, the precise structure of cutan is still unclear. Pyrolysis data (de Leeuw *et al.*, 1991) in combination with chemolysis results (RuO_4 oxidation; Schouten *et al.*, 1998) indicate that it is based mainly on linear long-chain (C_{30} – C_{34}) aliphatic compounds, probably hydroxy-alkanoic acids. These could be bound covalently via non-hydrolysable ether bonds as well as via relatively labile ester bonds (Figure 8.7). Evidence of cinnamic acids in the cutan is

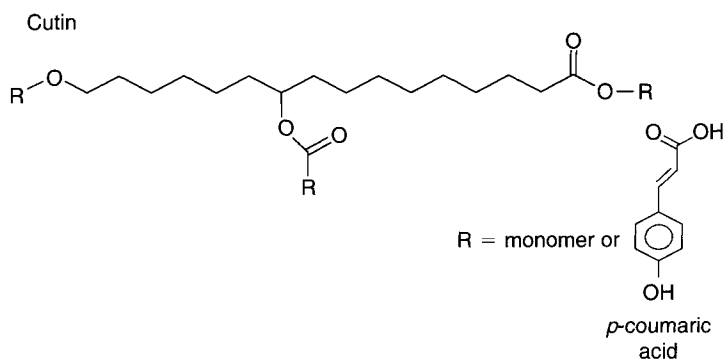


Figure 8.6 Structure of cutin building blocks based mainly on a dihydroxy fatty acid. See Figure 8.5 for additional information.

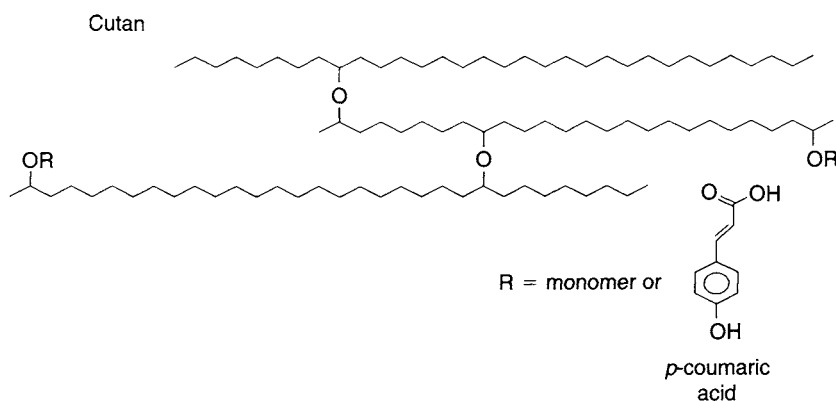


Figure 8.7 Tentative and simplified structure of a cutan based on long-chain alkyl units linked by ether moieties. See Figure 8.5 for additional information.

absent but the involvement of some small amounts of aromatic units in the cutan has been postulated based on ^{13}C NMR and thermochemolysis data (McKinney *et al.*, 1996).

Most fossil cuticles reveal the presence of a macromolecule based on long-chain aliphatic moieties often with distinct distribution patterns indicating mainly building blocks of even carbon chain length (C_{22} – C_{34} ; Tegelaar *et al.*, 1991; van Bergen, 1994; Collinson *et al.*, 1994, 1998; Zodrow and Mastalerz, 2001). Interestingly, however, these fossils include taxa (e.g. *Ginkgo*; Collinson *et al.*, 1998) the modern counterparts of which are devoid of cutan. This therefore implies that the long-chain aliphatic macromolecule in fossil cuticles may in some cases be selectively preserved cutan or alternatively, a newly formed highly aliphatic geomacromolecule. As these fossil remains still appear as cuticles based on morphology and ultrastructure, the compounds involved will have to be derived from very near the cuticle itself. Cutin has been suggested as a source but it is unlikely that stabilization of this biopolyester plays a major part as this should reveal evidence of distinct linear C_{16} and/or C_{18} moieties. More recently, these observations have led to the hypothesis of within-cuticle diagenetic stabilization of normally labile aliphatic constituents, i.e. cuticular waxes, for the formation of highly aliphatic macromolecules in fossil cuticles (Collinson *et al.*, 1998). Interestingly, the chemical composition of fossil cuticular layers of related taxa can reveal chemosystematic characteristics indicating that if the highly aliphatic macromolecule in most fossil cuticles were formed upon within-cuticle stabilization this process cannot be entirely random as this would mask the chemosystematic signal (*cf.* *Potamogeton* L. versus *Limnocarpaceae* Reid and Chandler, van Bergen, 1999; van Bergen *et al.*, 1999; *Typha* L. versus *Sparganium* L.; Collinson and van Bergen, 2003; differences between various Carboniferous plant cuticles, Mösle *et al.*, 2002). Maybe cutan in modern cuticles is also based on within-cuticle cross-linking of cuticular lipids including long-chain *n*-alkanols, aldehydes and alkanolic acids. Enzymatic reactions or photo-oxidation in the living cuticle may in that case aid the formation of cutan. Recent pyrolysis data of cuticular lipids in combination with a mineral matrix showed that a cutan-type signal was obtained, suggesting that such an aliphatic macromolecular signal found in fossil cuticles may be formed during early diagenesis from waxes, some of which are present as metal salts (Finch and Freeman, 2001). In addition, one has to be aware of pyrolytically-induced alterations that can affect the chemical composition of fossil cuticles leading to spurious interpretations with respect to their original chemical composition. In particular, older cuticles often reveal chemical data implying relatively short-chain aliphatic building blocks (C_{10} – C_{18} ; Collinson *et al.*,

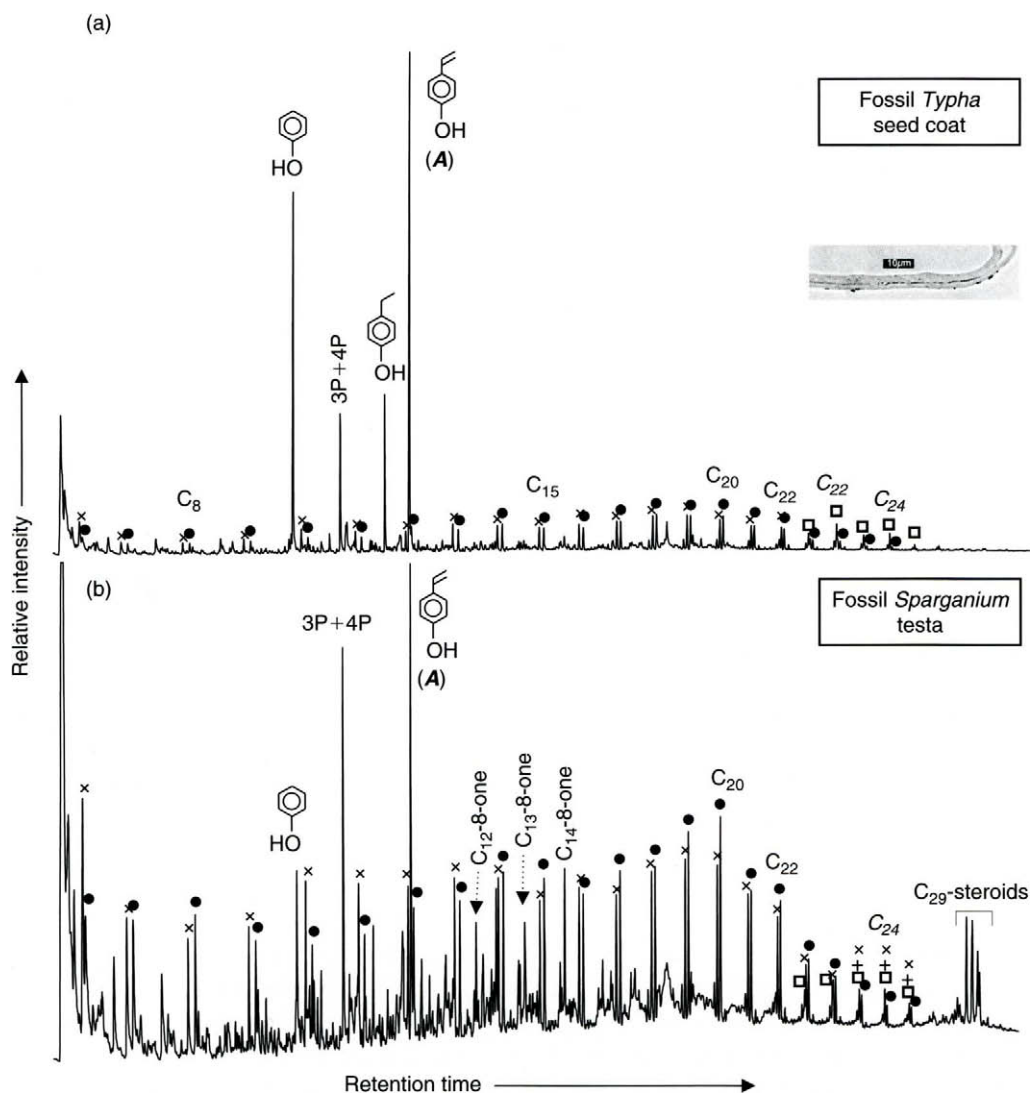


Figure 8.8 Gas chromatograms of pyrolysates (Curie-temperature 610°C) of (a) cuticular seed coat of fossil, Eocene (*ca.* 34 Ma), *Typha* and (b) fossil cuticular testa of *Sparganium*, obtained from the same bed, showing the presence of phenolic pyrolysis products (annotated P) derived from *p*-coumaric acid. Key: ● = *n*-alkanes, × = *n*-alk-1-enes, □ = 2-alkanones, C₉ = non-1-ene and nonane. C_{xx} indicate alkanones with *xx* representing the number of carbon atoms. P is phenol, 3P+4P = co-eluting 3- and 4-methylphenol. The LM photograph shows the two-layered cuticular composition of the fossil seed coat wall. For additional information regarding the samples, the reader is referred to van Bergen (1994a).

1994; Ewbank *et al.*, 1996; Zodrow and Mastalerz, 2001). These patterns are most probably solely due to increasing maturity of the organic material leading to an increase in cross-linking (van Bergen, 1999) and as such prevent unambiguous interpretations in terms of evolution and physiology.

In a few fossil cuticles evidence was found for the presence of phenolic compounds directly related to *p*-coumaric acid (Figure 8.8; seed cuticles of *Typha* and *Sparganium*;

leaf cuticles of *Ginkgo*; Collinson *et al.*, 1998). These phenolic compounds are linked both by ester- and ether-bonds. This could indicate that cinnamic acids are also incorporated in cutan of the cuticle, possibly with the same physiological function as in other plant organs (i.e. UV protection). However, the cuticular membrane of *Sparganium* is within a sclerotic endocarp making the need for UV protection less likely. Alternatively, these cinnamic acids have become incorporated upon within-cuticle stabilization during early diagenesis.

From a physiological point of view, the structural macromolecules in cuticles are, to some extent, similar and in other aspects rather different from those in algal, spore and pollen walls. First, the macromolecules in most cuticles do not contain abundant cinnamic acids, thus being distinct from sporopollenin. This despite the fact that most cuticles are directly exposed to direct sunlight. Most probably, other tissues underlying the cuticle (i.e. cell walls of the epidermis) may be involved in UV shielding, whereas no such protection exists for the cytoplasm in spores and pollen. A second difference is the presence of a structural polyester, i.e. cutin, that is based on relatively short-chain hydroxyacids. As long as these shorter alkyl units provide enough hydrophobicity in these structures, the production of such macromolecules may be advantageous. Reducing the amount of carbon needed to build these structural units, that under normal circumstances will not be used again by the plant, would be a major advantage. The similarity with algaenan and sporopollenin relates to cutan. This latter macromolecule is also based on long-chain aliphatic, ether- and possibly ester-linked (hydroxy) units. However, the paucity of in-depth molecular information of modern cutan prevents further physiological inferences.

Inner structural entities

Water-conducting and strengthening tissues

Lignin is most commonly associated with the secondary xylem in wood. This methoxyphenol-based biomacromolecule is often suggested to have been pivotal in an evolutionary sense for both water conductance and strengthening of the water-conducting elements (e.g. Edwards, 2001; Cooper-Driver, 2001). The findings of tracheids and other xylem-like elements in fossil settings as far back at the Silurian have been suggested to relate to the presence of lignin in these fossil remains (or at least lignification is implicated; Cook and Friedman, 1998; Cooper-Driver, 2001; Edwards, 2001).

The molecular building blocks of modern lignin are relatively well known despite the uncertainties of the overall molecular composition of the macromolecule itself. Lignin is based on monolignols including *p*-coumaryl alcohol, coniferyl alcohol and/or sinapyl alcohol (Figure 8.9) depending on the plant taxon (Sarkanen and Ludwig, 1971; Saiz-Jimenez and de Leeuw, 1986). In modern specimens the chemical composition of lignin is rather diverse with that of gymnosperms based mainly on coniferyl alcohol, whereas the angiosperms always contain coniferyl and sinapyl units. Monocotyledonous angiosperms and legumes may also contain abundant *p*-coumaryl units in addition to the other two. With respect to lower plants, such as bryophytes, controversy still exists as to whether the phenolic moieties present in these plants are indicative of real lignin (Lewis and Yamamoto, 1990). Proving the presence of lignin units is often dependent on detailed molecular analyses that can show intact and rather specific methoxyphenols. In particular the presence of intact side chains attached to the aromatic ring (Figure 8.10) is crucial

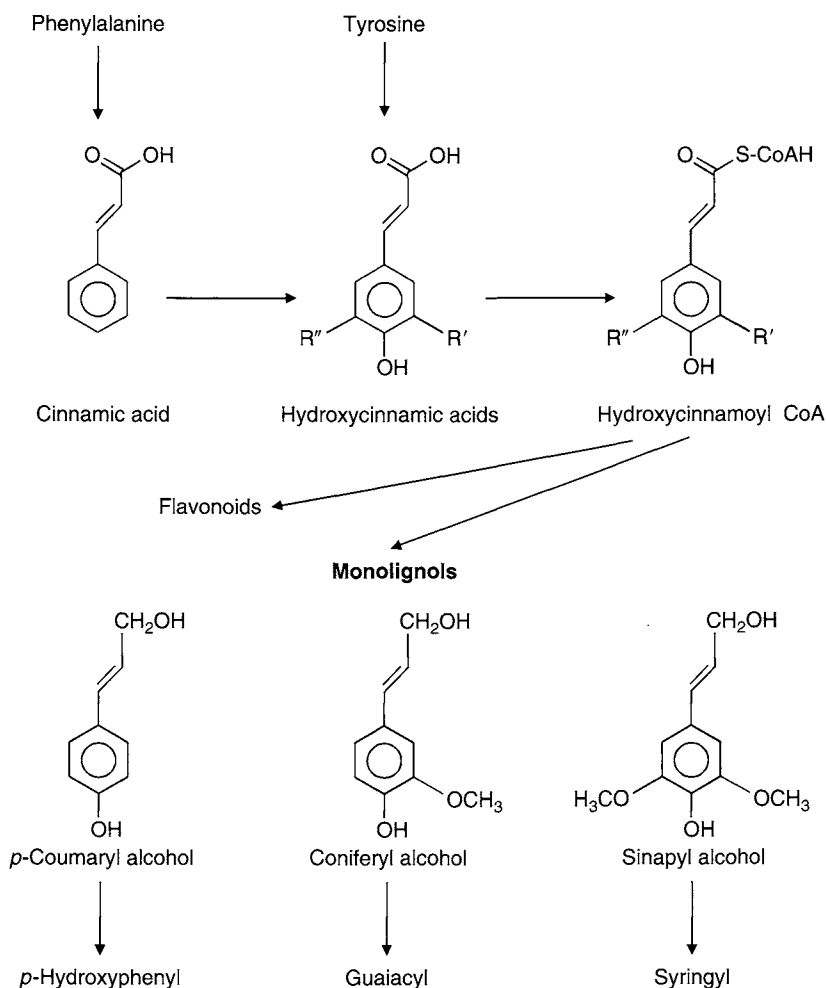


Figure 8.9 Biosynthesis of monolignols.

as various other biomolecules (e.g. tannins) can also release simple methoxyphenols (see below; Figure 8.11).

Evidence for lignin in fossil plant material is complicated by the fact that the characteristic (di)methoxyphenols undergo diagenetic transformation reactions leading to (3-methoxy)-1,2-benzenediols and ultimately phenols (Hatcher and Clifford, 1997; van Bergen *et al.*, 2000). These latter two compound classes are much less specific and can be derived from various sources other than lignin (i.e. from tannins, sporopollenin, proteins etc.). The distribution with time of original lignin units and their degradation products is shown in Figure 8.12. Unequivocal molecular evidence for lignin exists for wood samples as old Early Mesozoic, but most of these samples are characterized by lignin degradation products such as 1,2-benzenediols and phenols. In stark contrast, very few studies have shown the presence of methoxyphenols in Palaeozoic plant remains (e.g. wood material from the Moscow Basin; Hatcher and Lerch, 1991) and we know of no research revealing characteristic side chain evidence. For example Palaeozoic fossil wood from Carboniferous *Cordaites* only revealed phenols and to a smaller extent 1,2-benzenediols (Ewbank *et al.*,

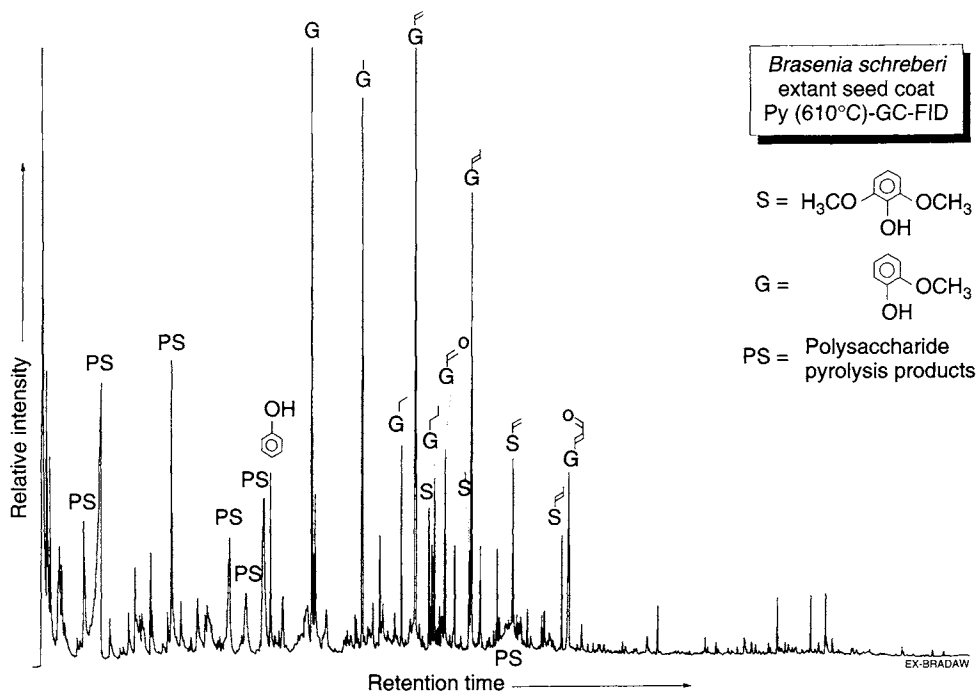


Figure 8.10 Gas chromatogram of the pyrolysate (Curie-temperature 610°C) of the ligno-cellulose complex present in seed coat of extant *Brasenia schreberi* J.F. Gmel. Note the abundance of products indicating 2-methoxyphenols (G) and 2,6-dimethoxyphenols (S) with side chains.

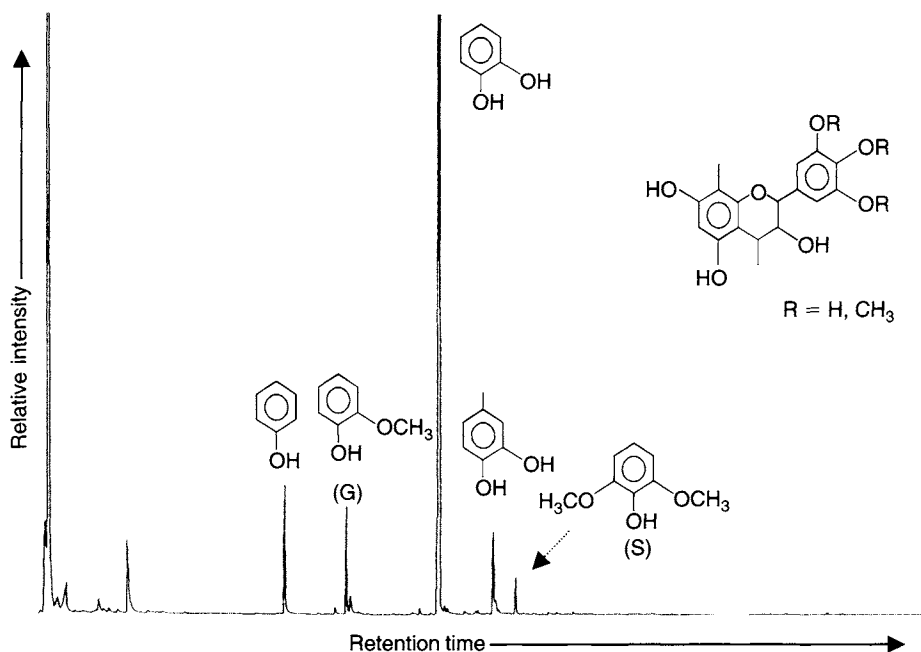


Figure 8.11 Gas chromatogram of the pyrolysate (Curie-temperature 610°C) of non-hydrolysable red wine tannin. Note the *absence* of 2-methoxyphenols and 2,6-dimethoxyphenols with side chains when compared with Figure 8.10.

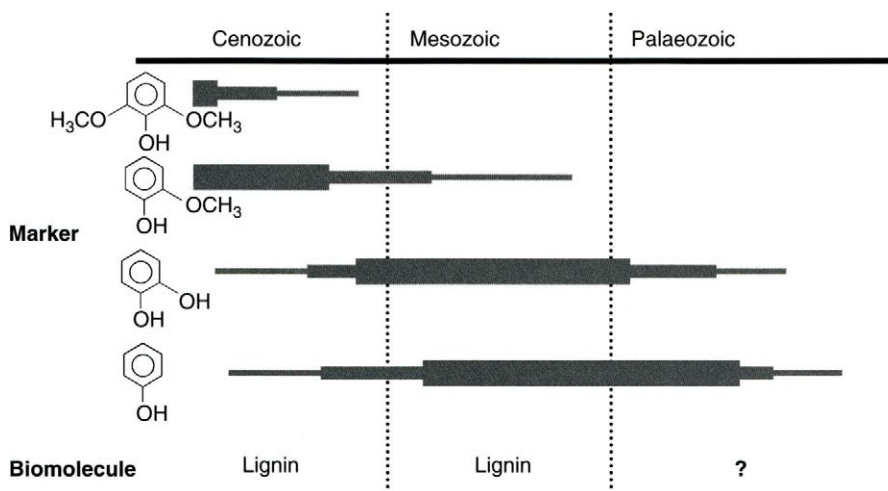


Figure 8.12 Abundance of (methoxy)phenols with time revealing the absence of characteristic methoxyphenols in the Palaeozoic.

1996), whereas older samples still, i.e. Devonian *Psilophyton*, *Prototaxites* Dawson and *Pachytheca* Hooker only showed evidence of phenols and simple aromatics (e.g. alkylbenzenes and naphthalenes; Edwards *et al.*, 1997; Abbott *et al.*, 1998). These data imply that the presence of lignin can only be proven in Mesozoic plant fossils and younger, minimizing the inferences that can be made on the physiological significance of the phenolic compounds present in early land plants in relation to strengthening and water conductivity.

However, xylem elements in modern plants also contain an abundant polysaccharide fraction *viz.* hemicellulose and cellulose which, together with lignin, form the ligno-cellulose complex. This complex in the main S-layers of the secondary wall is most probably composed of orientated cellulose fibres surrounded by a matrix of lignin and hemicelluloses (Terashima *et al.*, 1993; Salmén, 2000). Recent studies on the mechanical properties of wood imply that lignin in itself is not the sole factor determining cell wall strength (Hoffman *et al.*, 2000; Salmén, 2000). Cellulose fibres are now believed to be much more important for mechanical rigidity whereas lignin provides mainly a water proofing for the cellulose (Hoffman *et al.*, 2000). In addition, phenolic-based biomolecules such as condensed tannins (proanthocyanidins) have been suggested to play a structural role in some plant tissues (*cf.* Figure 3 in Shen *et al.*, 1986). Interestingly, pyrolysis data of condensed tannins revealed a number of distinct products including 1,2-benzenediols, 2-methoxyphenol and 2,6-dimethoxyphenol (see Figure 8.11). The presence of these latter two compounds is often used as evidence of lignin. However, the absence of additional (di)methoxyphenols containing characteristic side chains clearly shows that the sole occurrence of 2-methoxy- and/or 2,6-dimethoxyphenol cannot be used as unequivocal chemical evidence of lignin. An example that strengthening tissues of plants do not need lignin is provided by the molecular data of the fruit wall of *Nelumbo nucifera* Gaertn. (van Bergen *et al.*, 1997). These propagules are well-known for their extreme longevity (Shen-Miller *et al.*, 1995). Molecular analyses using solid state ^{13}C NMR in combination with pyrolysis revealed that the physically extremely hard sclerotic outer layer is based on a polysaccharide-tannin complex providing both physical strength as well as durability (van Bergen *et al.*, 1997). These molecular data from modern examples imply that lignin is not necessarily a physiological prerequisite for the evolution

of structural plant tissues and that other phenolic macromolecular biomolecules may have played a key role in the early evolution of land plants (*cf.* Abbott *et al.*, 1998).

Conclusions

The invasion of the land by plants may have forced the evolution of specific physiological adaptation to survive this hostile new environment. Two of the main problems plants had to overcome included an increase in the levels of UV radiation and water loss or desiccation. Studying the resistant macromolecular composition of outer coverings and strengthening tissues from both modern and fossil examples can reveal information on the molecular evolution of these structures. The resistant molecules in cuticles (i.e. cutin and cutan) and spore and pollen walls (sporopollenin) are all based on even carbon numbered long-chain aliphatic chemical building blocks providing sufficient hydrophobicity to reduce water loss. These aliphatic moieties are largely similar to those present in the resistant walls of algae (algaenan) from which the land plants may have evolved. Apart from the aliphatic material, sporopollenin and, to some degree, cutin and cutan from both modern and fossil examples also reveal the presence of cinnamic acids, which probably are responses to the enhanced levels of UV radiation on land. With respect to the strengthening tissues, lignin may have been an important biomolecule in early land plants but, to date, no unequivocal molecular evidence exists that it actually occurred in the oldest land plants. Moreover, molecular data from modern strengthening tissues indicate that lignin is not necessarily a physiological prerequisite for the evolution of plant tissues that provide physical strength and that other phenolic macromolecular biomolecules may have played an additional (key) role in the evolution of the early land plants.

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9

Early land plant adaptations to terrestrial stress: a focus on phenolics

Linda E Graham, Robin B Kodner, Madeline M Fisher,
James M Graham, Lee W Wilcox, John M Hackney,
John Obst, Peter C Bilkey, David T Hanson and
Martha E Cook

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Introduction

The first land plants no doubt found the terrestrial environment to be a bountiful source of carbon dioxide and light – resources that may have limited growth and reproduction of their aquatic algal ancestors (Graham, 1993). But there was probably a stressful ‘trade-off’ in the form of increased heat, desiccation, oxidative damage, harmful radiation and low soil

nutrient content. Interactions with terrestrial microbes may have been an additional new challenge. That early land plants successfully coped, is evidenced by more than a quarter million species of extant descendants and the many millions of terrestrial animal and microbial species that depend upon plants. Comparing the traits of modern representatives of ancient plant lineages offers insight into how plants developed ways of responding to terrestrial stressors. Molecular phylogenetic evidence indicates that modern land plants (embryophytes) are a monophyletic group descended from charophycean green algae (primarily aquatic) and that modern bryophyte lineages (primarily terrestrial) diverged prior to any group of modern vascular plants (data summarized by Kenrick and Crane, 1997).

The prevascular terrestrial biota probably included early plants that shared features with modern bryophytes. This assumption is supported not only by molecular sequence information, but also by an increasingly rich microfossil record of spores, tubes and cellular sheets that have been attributed to early, bryophyte-like plants (Edwards *et al.*, 1995; Graham and Gray, 2001 and references cited therein). Additional evidence is provided by morphometric similarity of ancient microfossils to remains of modern bryophytes that were subjected to procedures designed to mimic degradative changes that occur during fossilization (Kroken *et al.*, 1996; Kodner and Graham, 2001). Therefore, some traits of extant bryophytes may model early plant adaptations to terrestrial stress (though other features could be of more recent origin and thus not mirror the traits of ancient relatives).

One goal of our work was to map stress-related physiological traits onto a robust phylogeny for modern charophycean algae and bryophytes, because previous mapping of reproductive and structural developmental traits has provided useful insights into early plant evolution (Graham and Wilcox, 2000a; Graham *et al.*, 2000). Additional goals were to compare aspects of phenolic chemistry among charophyceans, bryophytes and pteridophytes and to estimate the extent to which non-vascular plants could have contributed to carbon sequestration prior to the origin of vascular plants. To these ends we: (1) used thioacidolysis as an assay for lignin-specific β -O-4 phenolic linkages in representative green algae and early-divergent land plants; (2) surveyed selected green algae and bryophytes for presence of resistant biomass; (3) quantitatively determined the percentages of resistant cell wall biomass for those that produced measurable amounts; and (4) used these and other data to estimate the amount of resistant organic carbon that might have been generated by early non-vascular land plants.

Materials and techniques

Trait mapping

We used phylogenies published by Karol *et al.* (2001) for charophyceans, Lewis *et al.* (1997) for liverworts, Qiu and Palmer (1999) and Nickrent *et al.* (2000) for bryophytes and Newton *et al.* (2000) for mosses, in order to compile a phylogram representing an emerging view of the early streptophyte (charophyceans plus embryophytes) radiation. Putative stress adaptation traits (whose occurrence was derived from the literature) were mapped onto the phylogram. Fourteen of these stress-related traits are described more fully in Appendix 9.1.

Thioacidolysis

Algae treated by thioacidolysis included unialgal, soil-free cultures representing major green algal lineages (Graham and Wilcox, 2000b): *Cladophora glomerata* (L.) Kütz.

(Ulvophyceae), *Chlorella vulgaris* Beij. (Trebouxiophyceae), *Oedogonium* Link sp. (Chlorophyceae), *Klebsormidium barlowii* (Silva *et al.*), *Spirogyra* Link sp., *Mougeotia* C.A. Agardh sp., *Staurastrum* Meyen sp., *Chara* L. sp. and *Coleochaete orbicularis* Pringsheim (Charophyceae). Soil-free laboratory cultures of the following bryophytes were examined: *Riccia fluitans* L. (liverwort), and the mosses *Sphagnum compactum* DC, *S. centrale* C. Jens, and *Polytrichum commune* Hedw. Moss cultures were generously provided by M. Sargent, University of Illinois. Cultures of the fern *Polypodium aureum* L. gametophytes and young sporophytes were grown from surface-sterilized spores in closed petri dishes containing mineral nutrient agar. The lycophytes *Lycopodium obscurum* L. and *Selaginella kraussiana* (Kunze) A. Braun and the pteridophytes *Pilotum nudum* L., *Tmesipteris vieillardii* Dangeard and *Adiantum capillus-veneris* L. were obtained from the University of Wisconsin Botany Department greenhouses and washed free of soil with sterile water prior to testing. Thioacidolysis was performed as described by LaPierre (1993). Freeze-dried plant or algal samples were boiled for 4 h under nitrogen in 0.2 M boron trifluoride etherate in 9:1 dioxane ethanethiol. After cooling, 10 μ l distilled water was added and the pH was adjusted to 3–4 with 0.4 M NaHCO₃. Five ml of a tetracosane solution (either 0.1 or 0.01 of tetracosane per ml of CH₂Cl₂ depending upon the anticipated concentration of derivatized product to be measured) was added as an internal standard. The sample was then extracted three times with 10 ml CH₂Cl₂. The combined extracts were dried over a small amount of MgSO₄ and paper filtered into an evaporation flask and the sample was rotationally evaporated to near dryness. The resultant residue was subsequently evaporated to dryness with a directed stream of N₂ gas and the products were derivatized by adding 1 drop of pyridine and 4–5 drops of BSTFA (bis trimethylsilyl trifluoroacetamide). Finally, the sample was reacted by warming at the lowest setting on a hot plate for 0.5 h prior to its analysis by GC/MS (Finnigan Model MAT 4500; 30 m DB-1 column).

Qualitative and quantitative assessment of acid hydrolysis-resistant biomass

Green algae studied included the desmids (Chlorophyta, Charophyceae) *Staurastrum* sp. Meyen, *Euastrum insigne* (Smith) Bréb. ex Ralfs, *Euastrum pinnatum* Ralfs, *Micrasterias truncata* (Corda) Bréb., *Closterium ehrenbergi* Meneghini, *Hyalotheca dissiliens* (Sm.) Bréb., *Desmidium majus* Lagerheim, and *D. grevillii* De Bary, and the co-occurring, non-desmid *Eremosphaera viridis* De Bary (Chlorophyta, Trebouxiophyceae). These taxa were isolated from Blueberry Lake (Vilas County, WI), Jyme Lake (Oneida Co., WI), or Birch Lake (Marquette Co., WI) and grown in unialgal culture in defined, soil-free medium DYIII (Lehman, 1976). Cultured algal cells of the same age were harvested by micropore filtration and lyophilized. Early-divergent mosses studied included two species of *Sphagnum*, *Andreaea rupestris* Hedw. and *Polytrichum ohioense* Ren and Card. Axenic cultures of *Sphagnum nemoreum* and *S. compactum* Lam. et Cand. (obtained from M. Sargent, University of Illinois) were grown in one-third strength Gamborg's B5 Basal Salts medium (Sigma) with 1% sucrose added, for an equivalent period of time, in both liquid or agar form. Small amounts of *Andreaea rupestris* were obtained from granitic rocks at 11 000 ft (3350 m) elevation from the Indian Peaks Wilderness Area, Boulder Co., CO (with permission of the US National Forest Service). Gametophytes of *Polytrichum ohioense* were collected from UW Madison Kemp Biological Station, Oneida Co., WI.

Qualitative assessment of resistance was performed by acetolysis treatment in Eppendorf tubes with concentrated acetic acid for 5–10 min at room temperature, then boiling in 9:1 acetic anhydride/concentrated sulphuric acid for 20 minutes, followed by washes in

concentrated acetic acid and then water (Kroken *et al.*, 1996). Each step was followed by gentle centrifugation in order to concentrate biomass so that supernatant could be removed by pipette. If plant or algal remains were present, they were visible as dark sediment in the tips of tubes and were removed for microscopic examination with a Pasteur pipette.

Quantitative determinations of resistant biomass were performed for the resistant-walled charophyceans *Staurostrum* spp., *Euastrum insigne*, *Desmidium grevillii* and *Hyalotheca dissiliens* and all of the mosses listed above. We first experimentally determined that the high temperature acid hydrolysis procedure described above did not significantly alter the weight of empty Eppendorf tubes. Lyophilized algae and mosses were thoroughly dried before being added to preweighed tubes, then tubes were re-weighed prior to acetolysis treatment. After treatment the tubes were dried for several days in an oven at 45°C before final weighing. Before and after treatment, plant or algal weight was calculated by subtracting the weight of the tube from the weight of the tube plus algae or plant material. The proportions of resistant to initial dry weights of specimens were computed and Student's *t*-test was used to identify significant differences among taxa.

Fluorescence, scanning and transmission electron microscopy

Acetolysed *Micrasterias* and *Andreaea* were examined with a Zeiss Axioplan fluorescence microscope in violet and UV excitation as described by Kroken *et al.* (1996). *Andreaea* was prepared for SEM by gold coating remains that had been affixed to aluminum stubs by double-sticky tape. Living specimens were prepared for transmission electron microscopy by soaking in 1% Triton-X for 1 h, followed by three quick rinses with phosphate buffer; fixation in 2% glutaraldehyde and 4% acrolein for 5 h, six buffer washes for 10 min each; fixation in 2% osmium tetroxide for 2 h, followed by six 10 min buffer rinses; dehydration in acetone, followed by infiltration with Spurr's resin over a period of 3 days; and polymerization at 70°C overnight. Images of sectioned, stained cells were obtained with a Zeiss 10 transmission electron microscope.

Global estimates of early Palaeozoic resistant and sequestered carbon

Estimates of sequesterable carbon produced by prevascular plants were developed by combining our quantitative determinations of acetolysis-resistant biomass in three early-divergent mosses (*Sphagnum*, *Polytrichum* and *Andreaea*) with literature-derived estimates for productivity and global area coverage. Minimal and maximal productivity estimates were 11–1656 g m⁻² year⁻¹ (Overbeck and Happach, 1957; Klinger *et al.*, 1994) for *Sphagnum* spp., 43–647 g m⁻² year⁻¹ (Davis, 1981; Cole and Monger, 1994) for *Polytrichum* spp. and 104 g m⁻² year⁻¹ (Longton, 1988) for *Andreaea alpina*. *Sphagnum* wetland habitat was estimated at 100% substrate cover (Vitt, 2000) of 3.5 · 10⁶ km² terrestrial surface, a minimal estimate for area occupied by this moss today (Gorham, 1991). *Polytrichum* substrate cover was estimated at 18%, based on measurements made on Signy Island, Antarctica (Longton, 1988). Today *Polytrichum* occupies a wide range of mesic to xeric environments, so its potential habitat prior to the origin and spread of vascular plants was conservatively estimated as 11 · 10⁶ km², 10% of the extent of modern humid tropical, temperate and dryland regions (Bailey, 1989). *Andreaea* substrate cover was estimated as 50% (Signy Island, Antarctica; Longton, 1988). Modern high latitude and altitude habitats occupy 38 · 10⁶ km² (Bailey, 1989). Since similar Ordovician-early Silurian habitat areas are difficult to estimate, we used 10% of this modern value (3.8 · 10⁶ km²) to estimate conservatively habitat area for ancient *Andreaea*-like mosses.

Molecular sequence divergence evidence suggests that earliest land plants may have appeared 700 million years ago (Ma) (Heckman *et al.*, 2001). However, our calculations were more conservatively made for the 40 Ma period ranging from the mid-Ordovician (Caradoc) (460 Ma) – when bryophyte-like microfossils first become apparent (Graham and Gray, 2001) – to the mid-Silurian (Wenlock-Ludlow boundary) (about 420 Ma) – when more complex plants began to dominate, as indicated by changes in the structure of dispersed spores and the appearance of higher plant-type cuticles in the fossil record (Wellman and Gray, 2000). Non-vascular decay-resistant plant carbon that could potentially be buried during this period was calculated as (biomass unit area⁻¹ year⁻¹ · % resistant biomass) · (% substrate cover · global habitat area) · 4 · 10⁷. From the amount of such resistant carbon that might have survived weathering and other loss processes, sequestered carbon was conservatively estimated at 1% of this value.

Results

Figure 9.1 represents a consolidation of recent phylogenetic information for charophyceans, bryophytes and early-divergent vascular plants, onto which we mapped traits associated with stress adaptation that have been reported in the literature (see Appendix 9.1 for additional details regarding 14 of these traits). No evidence of hydroxyphenyl, guaiacyl, or syringyl-propyl thioacidolysis products from β -O-4 ethers was detected by GC/MS for any of the green algae or bryophytes tested, nor for gametophytes of the leptosporangiate fern *Polypodium aureum*. However, guaiacyl thioacidolysis products were detected at the 1.7% (dry biomass) level in cultured *Polypodium* sporophytes. *p*-Hydroxyphenyl thioacidolysis products were detected in adventitious roots (0.12%) and stems (0.01%) of the lycophyte *Lycopodium obscurum*, and in adventitious roots (0.02%) and stems (0.01%) of the lycophyte *Selaginella kraussiana*. *p*-Hydroxyphenyl thioacidolysis products were not detected in any of the other pteridophytes examined, though guaiacyl residues were found in these pteridophytes.

Cell walls of both *Euastrum* species examined, *Staurastrum* spp., *Closterium ehrenbergii*, *Desmidium grevillii*, *Hyalotheca dissiliens* and *Micrasterias truncata* (Figure 9.2) survived acetolysis and were autofluorescent in violet and ultraviolet irradiation, suggesting the presence of phenolic constituents. Cell walls of the desmid *Desmidium majus* and the trebouxiophycean *Eremosphaera*, which occur in the same habitat as resistant desmids, did not survive acetolysis. Among the desmids whose walls were resistant and for which quantitative measurements were performed, the proportion of resistant biomass ranged from 5 to 17% and differences among genera were statistically significant (Table 9.1).

There was no significant difference in acetolysis resistance between the two *Sphagnum* species examined and there was no significant difference in the resistant carbon content of *Sphagnum compactum* grown in liquid versus that grown on agar media. Resistant biomass of the mosses examined ranged from 25 to 82% and differences among genera were statistically significant (Table 9.2). The extent of *Andreaea* resistance was surprising; entire stem and branch systems survived (Figure 9.3). Chlorophyll autofluorescence persisted in some cells, suggesting that acid did not penetrate to their cytoplasm. Cell walls of acetolysed *Andreaea* remains were strongly autofluorescent in V and UV excitation (Figure 9.4). Though some *Andreaea* cell walls were eroded by the acid treatment, many retained their typical morphology (Figures 9.3 and 9.4) and leaf fragments retained sufficient structure to be recognizable as moss leaves. Ultrastructural examination of non-acetolysed

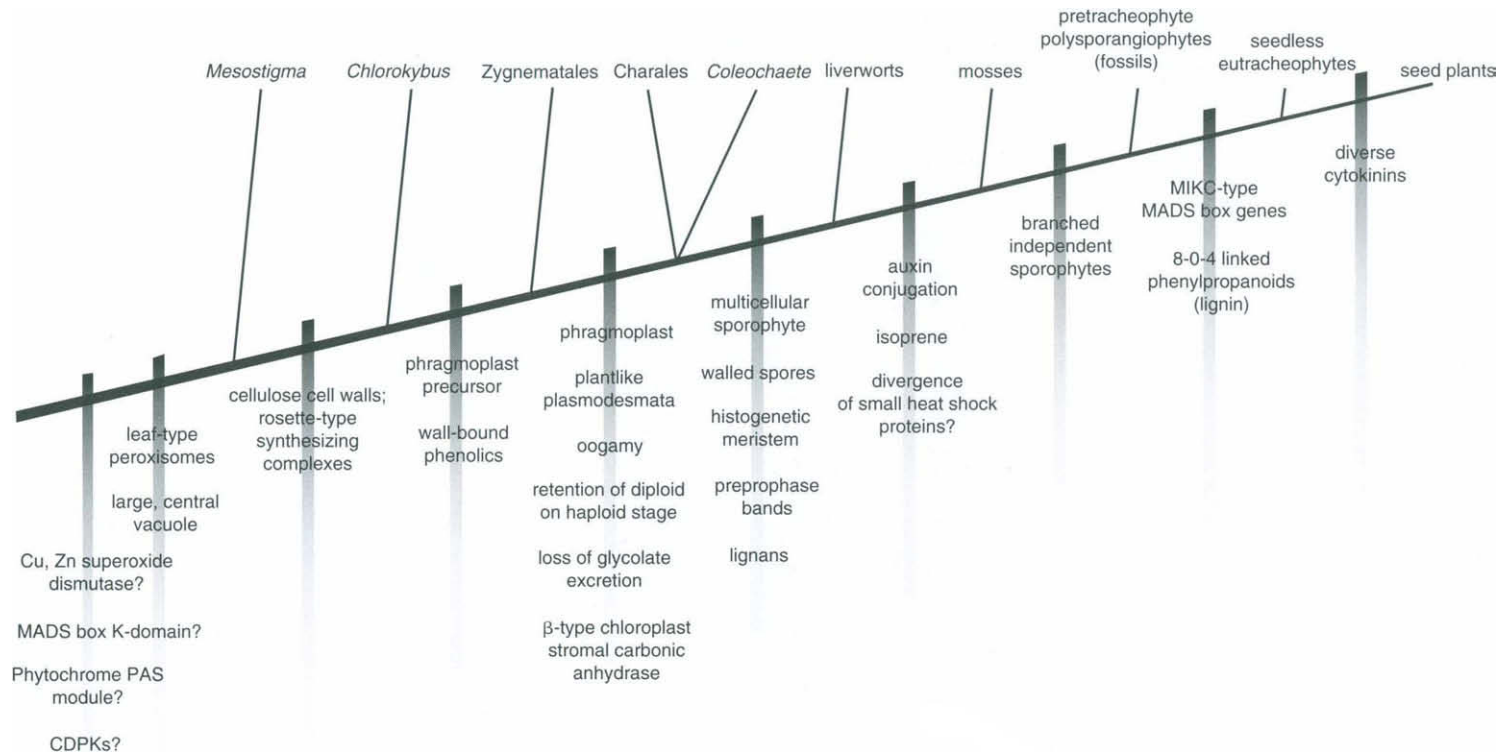
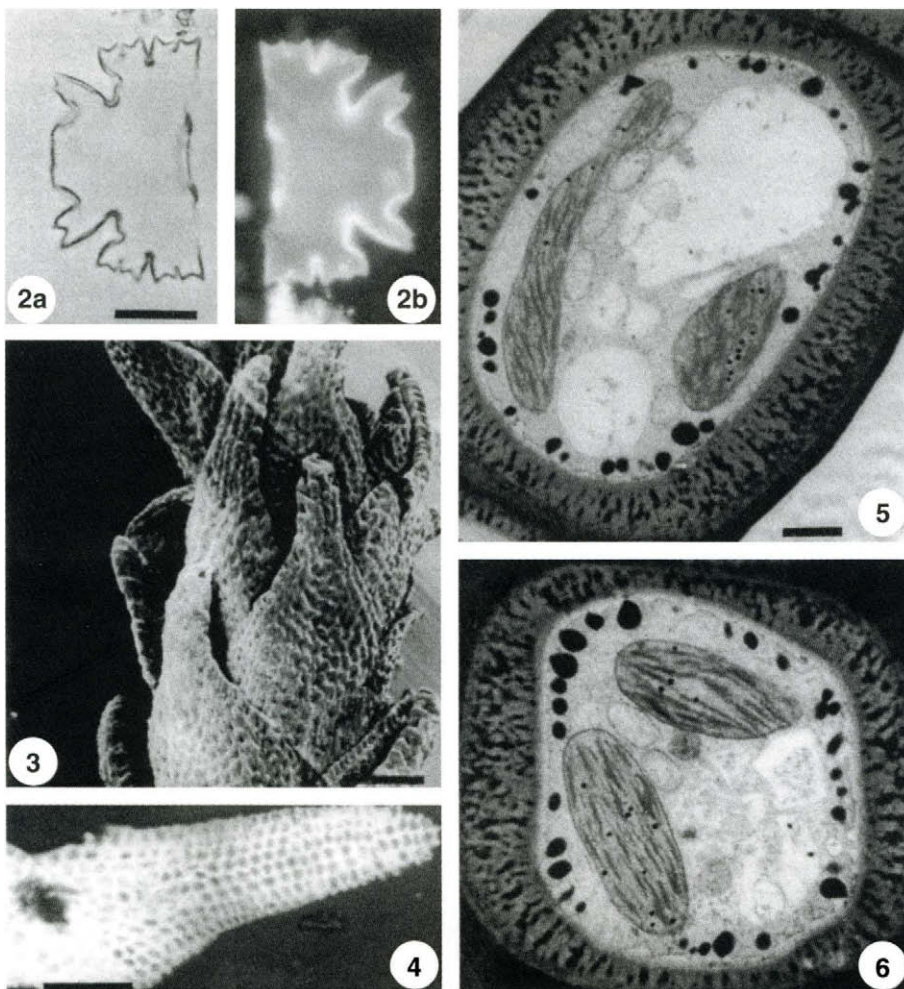


Figure 9.1 Traits 1–3 (see Appendix 9.1) appeared during the charophycean radiation, whereas traits 5 and 6 (as with the origin of the sporophyte) are associated with the dawn of embryophytes. Traits 7–12 likely appeared somewhat later, as their presence varies among bryophytes and trait 14 defines vascular plants. Though Karol *et al.* (2001) have linked Charales more closely with ancestry of land plants than *Coleochaete*, these taxa are figured together because they do not differ in the traits mapped here.



Figures 9.2–9.6 Resistant desmid and moss cells/tissues. Figure 9.2a Acetolysed *Micrasterias truncata*, LM view of semi-cell showing resistant wall having unaltered morphology. Scale bar = 50 μm . Figure 9.2b Same *M. truncata* specimen viewed with violet epifluorescence excitation. Figure 9.3 SEM of acetolysed *Andreaea* thallus showing intact branch and leaves. Scale bar = 23 μm . Figure 9.4 Autofluorescence of acetolysed *Andreaea* leaf in violet excitation suggests the presence of phenolics. Scale bar = 23 μm . Figures 9.5–9.6 TEMs of non-acetolysed *Andreaea* leaf cells showing extremely thick, densely-stained walls. Scale bar = 1.3 μm .

Table 9.1 Percent acetolysis-resistant biomass in selected charophycean green algae

| | Number of analyses | X | s ² |
|------------------------------|--------------------|------|----------------|
| <i>Staurastrum</i> sp. | 8 | 0.17 | 0.006 |
| <i>Desmidium grevillii</i> | 4 | 0.08 | 0.0001* |
| <i>Hyalotheca dissiliens</i> | 4 | 0.08 | 0.001* |
| <i>Euastrum insigne</i> | 4 | 0.05 | 0.0001** |

*Significantly different from *Staurastrum* at $P = 0.05$ level; **significantly different from *Staurastrum* at $P = 0.01$ level.

Table 9.2 Percent acetolysis-resistant biomass of early-divergent mosses

| | N | X | s ² |
|--|---|------|----------------|
| <i>Sphagnum nemoreum</i> (A) | 3 | 0.12 | 0.003 |
| <i>Sphagnum nemoreum</i> (L) | 3 | 0.24 | 0.002 |
| <i>Sphagnum compactum</i> (A) | 3 | 0.23 | 0.001 |
| <i>Sphagnum compactum</i> (L) | 3 | 0.26 | 0.006 |
| Pooled <i>Sphagnum</i> data ⁺ | 9 | 0.24 | 0.002* |
| <i>Polytrichum ohioense</i> | 5 | 0.45 | 0.04 |
| <i>Andreaea</i> sp. | 3 | 0.81 | 0.018** |

A, agar-grown cultures; L, liquid medium.

⁺All *S. compactum* plus liquid-grown *S. nemoreum* data pooled; *significantly greater than *Staurastrum* (and all desmids tested) at the $P = 0.05$ level;

**significantly greater than *Sphagnum* (and all desmids tested) at the $P = 0.025$ level.

Table 9.3 Estimates of early plant resistant carbon potentially produced during hypothesized 40 Ma period between appearance of moss-like bryophytes and rise of more complex plants

| | Minimum production rate | Maximum production rate |
|------------------------|----------------------------|----------------------------|
| I. <i>Sphagnum</i> | $485 \cdot 10^{18}$ g | $57\,960 \cdot 10^{18}$ g |
| II. <i>Polytrichum</i> | $1\,567 \cdot 10^{18}$ g | $23\,572 \cdot 10^{18}$ g |
| III. <i>Andreaea</i> | $6\,482 \cdot 10^{18}$ g | |
| Total | $8\,534 \cdot 10^{18}$ g | $88\,014 \cdot 10^{18}$ g |

Andreaea cell walls revealed that they were thick and stained densely with heavy metals (Figures 9.5 and 9.6).

Table 9.3 shows calculations of potential resistant organic carbon production by early non-vascular land plants. These are based on cover estimates and productivity data taken from the literature (cited in material and techniques section) and our measurements of percent resistant biomass for three early-divergent mosses: *Sphagnum* (characteristic of modern wetlands), *Andreaea* (an inhabitant of high altitude and high-latitude areas) and *Polytrichum* (which occurs in mesic to xeric regions).

Discussion

Mapping the traits of charophyceans and early-divergent land plants: (1) aids in discovery of new traits and provides an organizational basis for surveys of trait incidence in poorly-studied groups; (2) suggests the point of origin of traits, providing insight into adaptive or preadaptive function; (3) indicates how complex traits have evolved from simpler precursors, revealing systems that may be easier to investigate than those of higher plants; and (4) can be used to infer trait loss. One of the clearest signals emerging from the mapping exercise is the staged appearance of traits related to phenolic chemistry that contributed importantly to stress management.

Defensive phenolics associated with vegetative walls and sporopollenin-like materials (whose autofluorescence properties suggest presence of phenolic constituents) appear first in charophycean algae, prior to divergence of Zygnematales. Such defensive phenolics

(Appendix 9.1, Traits 2 and 4) were likely inherited by embryophytes (Graham, 1996), where they serve multiple roles (Cooper-Driver, 2001). Intracellular flavonoids and lignans, whose synthesis depends upon occurrence of the phenylpropanoid (C_6C_3) pathway, appear during divergence of bryophyte groups, but the order of first occurrence is unclear because flavonoids are not reported for hornworts and lignans have not been reported for mosses (Appendix 9.1, Traits 8 and 9). Presence of flavonoids and lignans implies the existence of biosynthetic pathways leading to production of cinnamic acid by deamination of phenylalanine, hydroxylation of cinnamate to *p*-coumaric acid and synthesis of ferulic acid and other compounds that are known intermediates and/or constituents of vascular plant lignin, cutin, suberin and sporopollenin (Cooper-Driver, 2001). O-methyl transferases, which confer greater reactivity to the number 4 carbon of phenylpropanoid monomers and thus facilitate formation of 8-O-4 (β -O-4) linkages (widely used to define lignin), appeared before the divergence of vascular plants (Appendix 9.1, Trait 14). The pattern of trait appearance suggests that biosynthetic and regulatory processes preadaptive to the evolution of lignin likely occurred in non-vascular plants, including ancestors of modern bryophyte and charophycean lineages. Trait mapping also suggests that early phenolics could have been preadaptive to the development of stable plant–microbe relationships (Appendix 9.1, Traits 3 and 11). As in modern plants, phenolic compounds may have controlled microbial behaviour, allowing microbes to live in close proximity to algae and early land plants without becoming pathogenic.

Our thioacidolysis data indicate that phenolic β -O-4 linkages are absent from non-vascular plants and charophyceans. Although lacking β -O-4 linkages, *Sphagnum* generates phenylpropanoids from phenylalanine via activity of phenylalanine ammonia lyase (PAL) and 4-cinnamic acid hydroxylase (4-CL) (Rasmussen *et al.*, 1995), as do vascular plants. Higher plant cell walls contain covalently bound non-lignin phenolics, which have been proposed to inhibit microbial degradation, serve as lignin-initiation sites, or limit wall extensibility (Wallace and Fry, 1994). We speculate that these higher plant wall phenolics may be related to those found in charophyceans and bryophytes. Our thioacidolysis results, that *p*-hydroxyphenyl monomers were more abundant in lycophytes (earliest-divergent extant vascular plants) than in the other vascular plants we examined, suggests that early lignins may similarly have been richer in non-methoxylated monomers than those characteristic of modern gymnosperms and angiosperms, where dominant monomers are methoxylated and β -O-4 bonds comprise 50–70% of linkages (Lewis, 1999). Quantitative assessment of resistant carbon content of selected desmids indicates that there is considerable variability among genera, with some desmids apparently lacking resistant walls, most tested species possessing resistant walls and *Staurastrum* having the largest proportion of resistant dry weight among desmid taxa tested. The latter result is consistent with: (1) the finding that *Staurastrum* is among the algae most resistant to microbial degradation (Gunnison and Alexander, 1975a); (2) the presence in *Staurastrum*'s walls of a 'lignin-like' phenolic polymer in amounts sufficient for chemical extraction, pyrolysis GC, thin-layer chromatography and other chemical analysis procedures (Gunnison and Alexander, 1975b); and (3) strong autofluorescence in V and UV excitation of *Staurastrum* cell walls both before and after high-temperature acid hydrolysis (Kroken *et al.*, 1996). Findings reported in the present work suggest that a variety of desmids (though not all) likely possess resistant cell walls that may protect them from microbial degradation and possibly also desiccation in some cases. Our data may explain: (1) the survival of desmid cells of several species after 3 months of drying and at depths to 6 cm in mud (Brook and Williamson, 1988); and (2) the occurrence of the

earliest known putative fossil desmid (*Paleoclosterium*) in Middle Devonian deposits (Baschnagel, 1966).

Desmidiium grevillii and *Hyalotheca dissiliens*, found in this study to possess similar levels (8% dry weight) of resistant biomass, are both known to harbour epibiotic bacterial communities. These communities include saprophytes having the potential to degrade carbohydrate components of cell walls, as well as members of the alpha proteobacteria (Fisher *et al.*, 1998a). Since many of the alpha proteobacteria are nitrogen fixers, it is possible (though not as yet demonstrated) that some desmid bacterial epibionts may provide nutritional advantage in nutrient-poor bog habitats. The presence of hydrolysis-resistant walls may reduce the chances that desmids might suffer deleterious effects, such as wall degradation, as a result of the activities of their epibiotic bacterial community. We propose that resistant wall phenolics originated in aquatic charophyceans as antimicrobial defences (that may also have provided some desiccation resistance), then acquired additional functions, including structural support and UV-screening in early land plants.

Our acetolysis analysis, demonstrating high proportions of resistant dry weight biomass in *Sphagnum* and other bryophytes, are consistent with conclusions that *Sphagnum* moss walls contain a 'lignin-like' 'polyphenolic network composed of *p*-hydroxyphenyl groups' 'linked with simple ether and ester bonds, interspersed with cinnamic acid' (Williams *et al.*, 1998). Our results help to explain why moss litter decomposes at 1–10% of the rate of vascular plants (Oechel and Van Cleve, 1986 and works cited therein).

Adaptive utility for high levels of wall phenolics in mosses might include: (1) resistance to attack by pathogenic bacteria, protists and fungi; (2) increased stability of cell walls, contributing to the ability to achieve increased height; (3) UV-damage resistance; and (4) desiccation resistance. *Sphagnum* occupies moist to waterlogged habitats in which bacterial abundance and activities are surprisingly high (Fisher *et al.*, 1998b). Hydrolysis-resistant cell walls may be helpful in resisting microbial attack and may also contribute to dimensional stability of the large, empty, porose water-holding hyaline cells of *Sphagnum*, which might otherwise tend to collapse. The relatively large size of *Polytrichum* erect shoots and the ability to form tall moss turf may be due in part to compression resistance aided by the presence of wall phenolics. Prevention of attack by microorganisms (and perhaps also invertebrate herbivores) provided by phenolic wall compounds may also contribute to biomass accumulation by *Polytrichum*, an inhabitant of mesic environments worldwide. The primary benefit obtained by *Andreaea* from its very high dry weight content of hydrolysis-resistant wall compounds may be UV protection afforded by phenolics, whose presence is implied by specific autofluorescence properties. This high altitude/high latitude moss occurs in some of the most UV-impacted habitats on earth. Resistant wall compounds may contribute to plant strength and formation of short, but high-density turfs and cushions, as may the extreme thickness of cell walls, an advantage in its harsh, wind-swept habitat. Wall phenolics may help maintain dimensional stability during cycles of hydration and dehydration, as well as contributing to desiccation and decay resistance. The bryalean moss, *Racomarpus purpurascens*, which likewise grows on rocks at high altitude, is reported to have cell walls consisting mainly of 'lignin', hemicellulose and cellulose in a ratio of about 9:8:5, that are architecturally adapted for absorption of water from fog, dew or rain (Edelmann *et al.*, 1998). The extreme habitats of *Andreaea* and *Racomarpus* model the exposed, bare rock surfaces that may have been occupied by some early land plants.

The rise of vascular land plants is regarded as an exceedingly important biogeochemical event in Earth's history because it is proposed to have played a dominant role in the dramatic Palaeozoic reduction in atmospheric CO₂ level (Berner, 1997) (but see Boucot

and Gray, 2001 for a critique). The twin mechanisms of this CO₂ reduction are suggested (by Berner, 1997) to have been: (1) an increase in weathering of silicate minerals by root systems; and (2) burial of decay-resistant organic carbon in the form of lignin. However, unequivocal fossil evidence for roots (assembled by Kenrick and Crane, 1997) does not occur until the Early Devonian, whereas Berner's (1997) models based on geochemical evidence suggest that atmospheric CO₂ drawdown began prior to that time. Analyses of carbon isotopic ratios in Fe(CO₃)OH of an upper Ordovician (440 myr) goethite dominating an oolitic ironstone (Neda Formation, Neda, Wisconsin, USA) suggest that the productivity of the prevascular biota was similar to that on modern soils (Yapp and Poths, 1994). Boucot and Gray (2001) point out that prevascular land plants, terrestrial and freshwater cyanobacteria, and marine algae may have generated considerable buried carbon during the Precambrian and Early Palaeozoic. Our results support Boucot and Gray's (2001) view that early bryophyte-like land plants may have played an important role in the carbon cycle prior to the rise of vascular plants.

Hanson *et al.* (1999) demonstrated that many mosses, including *Polytrichum* and *Sphagnum*, have the capacity to generate isoprene, which may protect the photosynthetic apparatus under high irradiance and temperature conditions. *Polytrichum* and *Andreaea* maintain the ability to photosynthesize at high levels after repeated cycles of dehydration and rehydration (Davey, 1997) associated with intermittently desiccating environments. Such environmental conditions probably prevailed in Ordovician-Silurian times, when a shady canopy of larger plants was absent. Soil-enriching, nutrient-binding humus is generated by modern mosses, which are the primary source of fresh organic matter in regions such as Antarctica (Beyer *et al.*, 1997).

Conclusions

The original data presented here, together with published molecular systematic and fossil evidence, suggest that productive bryophyte-like early land plants probably enriched soils with organic carbon and contributed to CO₂ sequestration for a long period prior to the rise of rooted plants. Even if as little as 1% of the 8540–82 000 × 10¹⁸ g resistant C that we estimated to have been produced by early bryophyte-like plants survived decomposition and weathering processes and was sequestered, early non-vascular plants could have removed 0.6–6% of atmospheric CO₂. If Berner's (1997) CO₂ drawdown models are correct in inferring that atmospheric CO₂ levels dropped from 20× present levels to 10× present levels during the 40 million year time period just prior to the rise of vascular plants, then our estimate for non-vascular plant resistant C sequestration (1% of resistant C produced) could account for 1–11% of this CO₂ drawdown.

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Appendix 9.1 Physiological traits related to early stress adaptation in land plants

Trait 1, Cu, Zn superoxide dismutase, is absent from most green algae, but occurs in all streptophytes that have been examined (DeJesus *et al.*, 1989). This enzyme prevents damage to cells by eliminating highly reactive oxygen radicals and may have contributed to early plant survival in the terrestrial atmosphere where O₂ diffuses into cells at a rate 10 000 times higher than in water.

Trait 2, decay- and acetolysis-resistant, autofluorescent phenolics in vegetative cell walls in some Zygnematales (Gunnison and Alexander, 1975a) as well as later-divergent charophyceans and bryophytes (Kroken *et al.*, 1996) is hypothesized to confer resistance to microbial attack (Gunnison and Alexander, 1975b), UV and possibly also desiccation and herbivory.

Trait 3, bacterial associations having putative beneficial properties with charophyceans (Fisher *et al.*, 1998a) and *Sphagnum* moss (Lau, 2000) suggest that such associations may be of ancient origin, possibly helping to alleviate low-nutrient stress.

Trait 4, sporopollenin-like layers in charophycean zygote walls (Delwiche *et al.*, 1989 and references cited therein), likely provide protection from microbial attack and possibly also desiccation during resting periods and dispersal of later-divergent charophyceans. Biochemical homology to embryophytic sporopollenin requires investigation.

Trait 5, sporopollenin-walled spores allowed meiospores of early plants (known as microfossils) to disperse via air. Sporopollenin provided dimensional stability, pathogen resistance (Graham and Gray, 2001) and possibly also desiccation resistance. Charophycean meiospores are devoid of sporopollenin wall layers (Graham, 1990).

Trait 6, tissues produced by a histogenetic meristem that cuts off derivatives in more than two directions is hypothesized as an adaptation that helped to prevent water loss from early plants by reducing surface area to volume ratio (Graham, 1996).

Trait 7, cuticle present on some moss tissues and hornwort sporophytes (Kroken *et al.*, 1996) (for which autofluorescence properties suggest inclusion of phenolics) likely reduces desiccation of underlying tissues, retards microbial attack and acts as a UV screen.

Trait 8, lignans – small, soluble polymers of phenylpropanoid monomers similar to those in lignin – characterize hornworts (Takeda *et al.*, 1990), liverworts (e.g. Cullman and Becker, 1999) and vascular plants (Lewis and Davin, 1994), but have not been reported from charophyceans or mosses. Because lignans function in lignin synthesis and perform as antibacterial, antifungal, or antiviral agents, or serve as antioxidants, they may have helped early land plants cope with the oxidative effects of higher terrestrial O₂ diffusion rates and pathogen attack or served as intracellular UV screens.

Trait 9, simple flavonoids, derived from the phenylpropanoid pathway and involving chalcone synthase, characterize at least some mosses and liverworts (but not hornworts, so far as is known) (Cooper-Driver and Bhattacharya, 1998). UV screening, antioxidant activity, mediation of plant-microbe symbioses and pathogen defences are among the roles played by flavonoids in modern plants that might be extrapolated to early plants.

Trait 10, vegetative desiccation tolerance, is present in at least some members of liverwort, hornwort and moss lineages, as well as some lycophytes, pteridophytes and angiosperms, and is regarded as a crucial step in the ecological transition from water to land (Oliver *et al.*, 2000; Proctor, 2000).

Trait 11 reflects associations of modern liverworts, hornworts and most groups of vascular plants (though not mosses or *Equisetum*, so far as is known) with ‘mycorrhizal’ fungi (Read *et al.*, 2000 and works cited therein). Presence of 455–460 My fossil glomalean fungi, which now occur only in symbiotic associations, together with molecular systematic analysis of the fungi (Redeker *et al.*, 2000 and works cited therein) and later fossil associations (Remy *et al.*, 1994) suggest that plants acquired fungal symbionts very early in their history. However, the time at which plants accrued positive benefits, namely increased access to nutrients and water, is not yet clear; the earliest plant–fungal associations may have been parasitic or saprophytic.

Trait 12, isoprene production in response to thermal and desiccation stress, has been demonstrated to occur in a hornwort, several mosses and various vascular plants, possibly functioning in protection of the photosynthetic apparatus. Absence of isoprene production from all liverworts examined (Hanson *et al.*, 1999) suggests that the enzyme isoprene synthase arose after the divergence of liverworts.

Trait 13, small heat shock proteins are produced by mosses and vascular plants in greater amounts as a response to high-temperature stress; they prevent thermal aggregation of cell proteins by maintaining them in a folded state. At least two of the five vascular plant families of small heat shock proteins occur in mosses (Waters and Vierling, 1999).

Trait 14, lignin, if defined as a polymer of phenylpropanoid monomers – at least some of which are linked by β -O-4 (8-O-4) bonds – occurs among extant plants only in vascular plants (Lewis, 1999) and coincides with the occurrence of O-methyl transferases.

Plant cuticles: multifunctional interfaces between plant and environment

Hendrik Bargel, Wilhelm Barthlott, Kerstin Koch,
Lukas Schreiber and Christoph Neinhuis

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Introduction

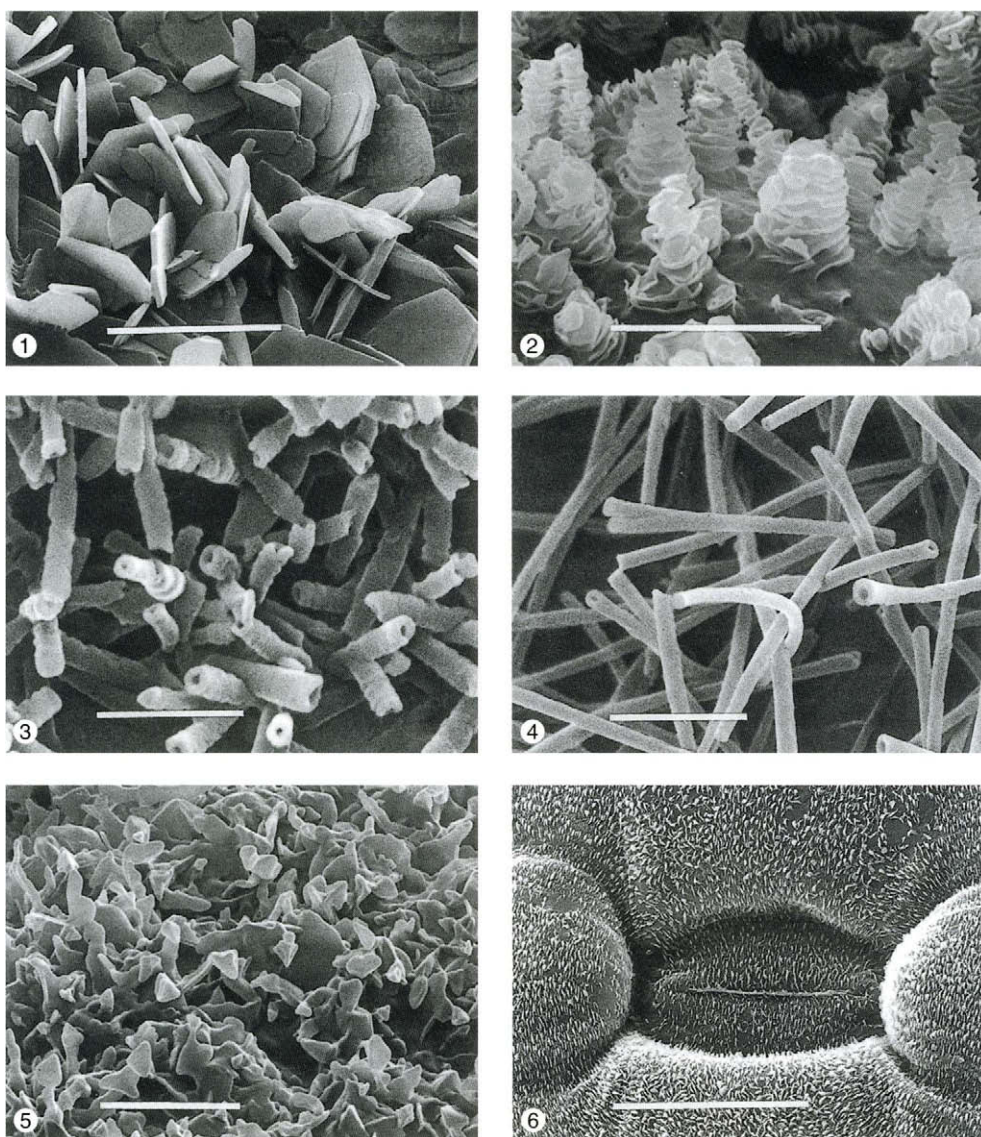
The plant cuticle covers all primary parts of vascular plants (except roots) and many bryophytes as a thin extracellular membrane. Deposited by the epidermis, the cuticle serves as the crucial protective layer between the organism and its environment, representing one of the largest interfaces between biosphere and atmosphere covering more than $1.2 \times 10^9 \text{ km}^2$ in total (Riederer and Schreiber, 1995). The cuticle has been of particular interest since it was first described by Brogniart (1834). In early studies, macroscopically visible features and fine structures were described and analysed. The reader is referred to Martin and Juniper (1970) or Holloway (1982b) for a summary. More recently, the ontogenetic development and chemical composition of both cuticle and waxes, as well as biosynthesis and transport phenomena have been subjects of major interest (Schönherr, 1982; Jeffree, 1986; Hamilton, 1995; Kolattukudy, 2001).

The cuticle represents a natural composite that consists mainly of two hydrophobic components, the insoluble biopolyester cutin and soluble lipids (Martin and Juniper, 1970). This type of lipid-derived polyester membrane is restricted to plants, whereas animals use carbohydrates or proteins as outer coverings (Kolattukudy, 2001). Minor amounts of cutinized polysaccharide fibre including cellulose and hemicellulose and a frequently abundant

pectin layer link the plant cuticle to the cell wall (Holloway, 1994). The basic framework is made of cutin, which is composed of saturated C₁₆ hydroxy- and unsaturated C₁₈ hydroxy-epoxy fatty acids (Kolattukudy, 1980a). The most common monomers are 16-hydroxy C₁₆ acid, 9- or 10,16-dihydroxy C₁₆ acid, 18-hydroxy-9,10-epoxy C₁₈ acid and 9,10,18-trihydroxy C₁₈ acid (Kolattukudy, 2001). The ratio of the C₁₆/C₁₈ monomers is organ as well as species specific and appears to vary. In general, fast growing plant organs, e.g. young leaves or fruits, seem to have a higher content of C₁₆ monomers (Espelie *et al.*, 1979; Baker *et al.*, 1982). The cutin monomers form a linear polyester by esterified primary hydroxyl groups, and cross-linking of esterified secondary hydroxyl groups yields a three-dimensional network (Kolattukudy, 1996; Ray *et al.*, 1998). In addition to cutin, a highly resistant residue, originally found in fossilized cuticles but also present in a few recent species, is named cutan (Nip *et al.*, 1986; Collinson *et al.*, 1998). Cutan has a high fossilization potential (van Bergen *et al.*, 1995; Möslle *et al.*, 1997) because of ether-linked long chain alkyl moieties mainly of even carbon chain length (C₂₂–C₃₄; Collinson *et al.*, 1998; Villena *et al.*, 1999). However, the origin and precise structure of this long chain aliphatic macromolecule still remains unclear (Jeffree, 1996; see also van Bergen *et al.*, Chapter 8).

The soluble lipids, collectively termed waxes, are embedded into the cutin matrix (intracuticular) as well as deposited onto the surface (epicuticular). Whereas the intracuticular waxes may be either amorphous or crystalline (Riederer and Schreiber, 1995), the epicuticular waxes often form complex three-dimensional crystalline structures in addition to thin amorphous films (Bianchi, 1995), a feature useful in plant systematics (Barthlott, 1981; Gülz, 1994; Barthlott *et al.*, 1998; Figures 10.1–10.6). Plant cuticular waxes are a complex mixture of aliphatic or cyclic components, including hydrocarbons, long chain fatty acids, aldehydes, primary and secondary alcohols, ketones, β -diketones and pentacyclic triterpenoids (Kolattukudy, 1980b; Walton, 1990), with overall dominating chain length for the aliphatic compounds of C₂₀ to C₃₅ (von Wettstein-Knowles, 1995). It has been shown in recrystallization studies that crystalline wax projections originate from self-assembly based on their specific chemistry (Jeffree *et al.*, 1975; Jetter and Riederer, 1995; Meusel *et al.*, 2000). Moreover, one predominating component often is responsible for a characteristic type of epicuticular wax crystals (von Wettstein-Knowles, 1972; Barthlott *et al.*, 1996; Meusel *et al.*, 1999). Nevertheless, wax composition is subject to great variation among plant species as well as during organ ontogeny, indicating a very customizable system (Markstädter, 1994; Riederer and Markstädter, 1996). It is worth noting that former studies on cuticular wax composition are more or less based on a mixture of intra- and epicuticular waxes caused by the extraction methods (Riederer and Markstädter, 1996), whereas it is now possible to separate both fractions distinctively with the help of a recently developed freeze-embedding method utilizing glycerol (Ensikat *et al.*, 2000; Jetter and Riederer, 2000).

Based on the results of chemical composition for different species, the plant cuticle appears to be without uniform composition and alterable during organ growth. Additional evidence for inhomogeneity was obtained from ultrastructural studies, e.g. the distinction between a thin outer 'cuticle proper' (<200 nm) with a variable number of layers supposed to contain either wax or cutin (lamellate region) and an inner 'cuticle layer' of variable thickness (up to 17 μ m) (Holloway, 1982b). The latter represents a mixture of waxes, cutin and polysaccharide fibres originating in the cell wall, in addition to pectins (reticulate region), resulting from the impregnation of the epidermal cell wall (Jeffree, 1996). There have been several attempts to summarize the structural variation, most prominently by Holloway (1982b, 1994). He identified six different types of lamellated cuticles and an extended review by Jeffree (1996) displays data of 119 species from 94 genera. From a systematic



Figures 10.1–10.6 Wax crystal forms. Figure 10.1 *Lecythis chartacea* Berg (Lecythidaceae): plates, widespread within plants. Figure 10.2 *Williamodendron quadriocellatum* (van der Werff) Kubitzki & H.G. Richt. (Lauraceae): transversely ridged rodlets, containing palmiton. Scale bars = 5 μm . Figure 10.3 *Lonicera korolkovii* Stapf (Caprifoliaceae): Nonacosan-10-tubules. Scale bar = 1 μm . Figure 10.4 *Columellia oblonga* Ruiz & Pav. ssp. *sericea* (Kunth) Brizicky (Columelliaceae): β -diketone-tubules. Figure 10.5 *Ledum glandulosum* Nutt. (Ericaceae): triangular rodlets, unknown chemistry. Scale bars = 2 μm . Figure 10.6 *Convallaria majalis* L. (Convallariaceae): locally restricted orientation pattern. Scale bar = 20 μm .

point of view, no particular cuticle type can be allocated to a particular taxon. Moreover, evidence is strong that the major differences between cuticle structural types may be assigned to the developmental stage of the cuticular membrane rather than to differences in mechanisms of formation, biochemistry or methodical approach (Jeffree, 1996).

Both cutin and the wax compounds, except the cyclic ones, are aliphatic homologues derived by *de-novo* synthesis of C₁₆ and C₁₈ fatty acids in plastids of epidermal cells (Post-Beitenmiller, 1996). Two enzyme complexes, namely fatty acid synthetase and palmitoyl (C₁₆)-elongase, have been found to be responsible for the basic fatty acid synthesis by a stepwise condensation-elongation mechanism (von Wettstein-Knowles, 1995). The resulting C₁₆/C₁₈ fatty acids are then processed or elongated extraplastidally by either several microsomal enzymes to the different wax derivatives, or by a family of mixed-function oxidases to cutin monomers (von Wettstein-Knowles, 1987; Post-Beitenmiller, 1996). However, little is known about either the molecular and genetic control, or the precise location where the biosynthetic processes take place.

The evolution of the cuticle is strongly interconnected to that of plant life on land. Pioneering land plants had to face the physical and physiological problems resulting from life out of water, such as gravity, desiccation, UV radiation or pollutant deposition on their surface (Edwards *et al.*, 1982). Earliest fossils, interpreted as cuticles, date back to the Ordovician and are more frequent in the Silurian (Edwards *et al.*, 1996; see also Raven and Edwards, Chapter 2). Due to their great functional importance, which will be discussed below, it is highly probable that early land plants were already covered, at least partially, by epicuticular wax crystals. These waxes, however, have no fossil record because of their delicate nature. Therefore, we only can speculate from extant plants what might have been the epicuticular coverings in extinct plants. The most widespread type of wax crystals to be found among all land plants is represented by small platelets, often composed of primary alcohols. Their most important function, especially in mosses and ferns, seems to be the protection against the formation of water films above stomata (Neinhuis and Barthlott, 1997) or other air-filled spaces, such as assimilation lamellae in Polytrichales (Neinhuis and Jetter, 1995), presumably to maintain high gas exchange rates. Another common type is represented by small tubules composed of, or dominated by, the secondary alcohol nonacosan-10-ol. It has been shown to occur in sporophytes of the moss family Polytrichaceae (Neinhuis and Jetter, 1995), in the fern genus *Pteris* L. (unpublished data), and characterizes most gymnosperms (Wilhelmi and Barthlott, 1997), as well as certain groups of angiosperms (Hennig *et al.*, 1994). This may indicate a mutual biosynthetic pathway once developed by the common ancestor of land plants. This question, however, remains open since the particular pathway leading to nonacosan-10-ol is unknown to date.

Generally, the diversity of wax crystals is lowest in lycophytes and mosses. In ferns a large number of substances have been analysed, but only for a limited number of species. Gymnosperms are characterized mainly by nonacosan-10-ol tubules and appear to be extremely uniform with respect to epicuticular waxes, except for cycads which show some degree of variability. On the other hand, angiosperms exhibit by far the greatest diversity, which even allows specific groups to be circumscribed on the ordinal level or above. A detailed description and discussion of the systematic significance of wax crystals among land plants is given by Barthlott *et al.* (2002).

Despite the variability of both cuticle and epicuticular wax, this chapter embraces the complex processes at the interface between plant and environment and thus stresses the relevance of this multifunctional hydrophobic coverage of land plants.

The multifunctional interface

The multiple constraints on the evolution of such an outer envelope, found on fossil and recent plant material, can be summarized by transport phenomena across the cuticle,

interaction with biotic and abiotic factors and biomechanical requirements. Most of the functions described in the following are related to the waxes within and upon the cuticle, a fact that stresses their importance in the interfacial interactions.

Transport properties of plant cuticles

One of the main functions of the cuticle, if not the most important, is the limitation of uncontrolled water loss via evaporation (Schönherr, 1982). Thus, selective pressure acted strongly on plants to develop an outer hydrophobic envelope functioning as a compromise between the contrary demands of desiccation avoidance and free gas exchange (Edwards *et al.*, 1996; Bateman *et al.*, 1998). Furthermore, leaching of organic and inorganic substances from the leaf interior has to be minimized (Tukey, 1970), as well as providing an effective barrier in terms of foliar uptake (Schönherr and Riederer, 1989) – a crucial role of the plant cuticle. Regarding the transport of organic and inorganic substances across plant cuticles, four different groups of compounds can be treated separately: (1) organic non-electrolytes (e.g. pesticides and xenobiotics); (2) water; (3) organic and inorganic ions; and, finally, (4) lipophilic molecules which form the cuticular waxes. The waxes are synthesized in epidermal cells and must cross the outer epidermal wall before they can be either sorbed into the cutin matrix or reach the surface of cuticles where they are deposited as epicuticular waxes. These different groups of compounds are considered in more detail below.

Organic non-electrolytes

In the past many investigations have been carried out analysing the transport of neutral organic compounds across plant cuticles (Schönherr and Riederer, 1989) and cuticular waxes (Riederer and Schreiber, 1995), and the effect of plasticizers on mobility in cuticles (Schönherr and Baur, 1994). Since plant cuticles themselves are essentially hydrophobic biopolymers, lipophilic substances generally have fairly high rates of diffusion across them. These investigations were either carried out in ecotoxicological research programmes, thus quantifying the uptake of lipophilic environmental xenobiotics entering the leaves by diffusion across the cuticle (Schönherr and Riederer, 1989), or it was intended to try to optimize the uptake of lipophilic herbicides into leaves (Hess and Foy, 2000). Transport across cuticles was studied using experimental systems at three different levels of complexity: (1) intact leaves (Schreiber and Schönherr, 1992a,b); (2) isolated cuticles (Riederer and Schönherr, 1984, 1985); and (3) isolated and subsequently recrystallized cuticular waxes (Schreiber and Schönherr, 1993). These studies showed that transport properties of recrystallized cuticular waxes, which represents a fairly artificial experimental system, were closely related to the transport properties of isolated cuticles and intact leaves (Kirsch *et al.*, 1997). From these results it may be concluded that the transport barrier of cuticles provided by waxes is a system which is self-organized, as waxes spontaneously recrystallize leading to similar barrier properties as found in intact leaves and isolated cuticles (Schreiber *et al.*, 1996a).

In steady state, transport of neutral molecules across cuticles is conveniently quantified using equation (1):

$$F = P \cdot A \cdot \Delta c \quad (1)$$

Flow, F , in mass per time (e.g. $\text{g} \cdot \text{s}^{-1}$), is directly proportional the area A (m^2) and the driving force Δc , in units mass per volume (e.g. $\text{g} \cdot \text{m}^{-3}$). The driving force Δc is defined as the concentration of substance in the donor minus that in the receiver ($c_{\text{donor}} - c_{\text{receiver}}$). The permeance P , in units of velocity ($\text{m} \cdot \text{s}^{-1}$), is the proportionality factor between F and

$A \cdot \Delta c$. The larger the value of P , the faster the transport across the cuticle and the poorer the barrier properties of cuticles. When P is known, barrier properties of cuticles from different species can be directly compared since they do not depend on experimental boundary conditions such as driving force, exposed cuticle area, or time (Kerler *et al.*, 1984). Cuticular permeances of different compounds across intact leaves and isolated cuticles varied between 10^{-11} to $10^{-7} \text{ m} \cdot \text{s}^{-1}$ and the extraction of cuticular waxes with organic solvents increased cuticular permeability for neutral organic compounds by factors varying between 10 and 10 000 (Schönherr and Riederer, 1989). This provides good evidence for cuticular waxes forming the transport-limiting barriers of plant cuticles for the diffusion of organic substances.

The permeance P ($\text{m} \cdot \text{s}^{-1}$) itself is a composite quantity (Equation (2)):

$$P = \frac{D \cdot K}{\Delta l} \quad (2)$$

The diffusion coefficient D ($\text{m}^2 \cdot \text{s}^{-1}$) describes the mobility of the compound in the cutin and amorphous wax phases. The partition coefficient K (dimensionless) is the ratio of the solubility of the compound in the lipophilic cuticle phase and in water (or other solvents), and Δl (m) gives the path length of diffusion across the cuticle (Kerler *et al.*, 1984). In simple homogeneous membranes, Δl would simply represent the thickness of the membrane, but since plant cuticles are heterogeneous membranes with the cuticular waxes basically establishing the transport-limiting barrier (Holloway, 1982a), the exact path length of diffusion is not known. There are, however, indications that the path length can be more than 2 orders of magnitude longer than the thickness of the cuticles due to the tortuosity of diffusion around impermeable, crystalline wax phases (Baur *et al.*, 1999; Buchholz and Schönherr, 2000). The partition coefficient K , a measure of the solubility of the compounds in the lipophilic cutin and wax phases, rises with increasing lipophilicity of the compounds and can vary by several orders of magnitude (Schönherr and Riederer, 1989). The diffusion coefficient D describes the mobility of the molecules in the cuticular membrane and decreases greatly when the molar volume of the solutes is increased (Baur *et al.*, 1997).

Water

Water permeability of plant cuticles has been studied repeatedly over the last decades (Schönherr, 1982; Kerstiens, 1996; Riederer and Schreiber, 2001). These ecophysiological investigations aimed at understanding the amount of water lost by cuticular transpiration under severe environmental conditions, such as water stress and high temperatures (Schreiber and Riederer, 1996), evaluating the limits of survival for plants. Most of these studies were carried out using isolated, stomatous cuticles. As water molecules are non-ionized, water transport across cuticles is normally analysed and quantified in the same way as with lipophilic organic compounds through equations (1) and (2).

Cuticular permeances for water, based on the concentration of liquid water as the driving force at 25°C, were measured on 24 species from different habitats (Schreiber and Riederer, 1996). Permeances varied between 10^{-11} and $10^{-9} \text{ m} \cdot \text{s}^{-1}$ and there was a clear relationship between cuticular transpiration and the natural habitat of the investigated species. Water permeability was lowest in tropical lianas reaching above the canopy with high radiation, increased in xeromorphic species naturally growing in the Mediterranean vegetation zone and were highest in mesomorphic species (Riederer and Schreiber, 2001). Most of the test specimens were not collected from their natural habitats but taken from

greenhouse grown plants. This may indicate that barrier properties are genetically determined and growth conditions have little, if any, effect. Extraction of cuticular waxes increased cuticular water permeability by factors varying between 10 and 1000 (Schönherr and Riederer, 1989), indicating that cuticular wax also forms the major transport barrier for small non-ionized molecules, such as water.

Other experiments have shown that the degree of water permeability of cuticles is significantly influenced by relative humidity (Schönherr and Merida, 1981). Increasing the relative humidity from 0 to 100% increased cuticular water permeability by factors varying between 2 and 3 (Schreiber *et al.*, 2001). This suggests that water molecules have to be adsorbed at the polar, non-esterified free carboxylic groups in the cutin polymer, leading to swelling and increased water permeability (Schönherr and Bukovac, 1973). Methylation of isolated cuticles, derivatizing free carboxyl groups to non-polar methyl ethers, in fact reduced the effect of relative humidity on cuticular water permeability by more than 50% (Schreiber *et al.*, 2001). How can the effect of air humidity best be explained on a molecular level? With increasing relative humidity an increasing amount of water is adsorbed. As a consequence the cutin polymer swells and the number and/or the diameter of water-filled polar pores traversing the cuticles increase, leading to higher rates of water diffusion across the cuticles. The occurrence of polar pores in plant cuticles has been demonstrated by Schönherr (1976). These observations suggest a cuticular transport model composed of two parallel paths by which water can cross cuticles (Figure 10.7): a polar path composed of polar aqueous pores and a lipophilic path composed of the hydrophobic cutin and amorphous wax domains (Schreiber *et al.*, 2001). Lipophilic compounds, having a high solubility in cutin and wax domains, will preferentially diffuse using the lipophilic path, while the polar path will be irrelevant for these compounds as their water solubility is limited. However, water has access to both paths of diffusion, namely through the lipophilic waxy domains and water-filled polar pores. This kind of interpretation is supported by the

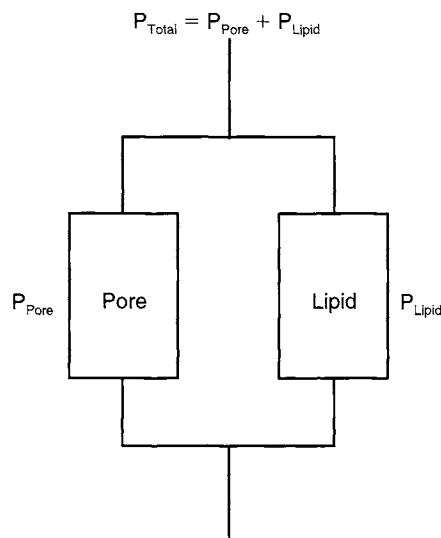


Figure 10.7 Schematic picture of the two parallel pathways of diffusion through plant cuticles: a polar path of diffusion across water-filled aqueous pores (Pore) and a lipophilic path of diffusion across cutin and waxes domains (Lipid). P_{Total} is the sum of the permeance across polar pores (P_{Pore}) and the permeance across lipid phases in the cuticle (P_{Lipid}).

experimental results for the transport of water across cuticles since water transport is strongly influenced by cuticular waxes. This indicates a transport across the lipophilic path, which is also affected by humidity, and suggests a parallel transport across polar pores. This transport model is further supported by studies of the transport of charged organic molecules and inorganic ions across cuticles, as discussed below.

Ions

Transport of ions (Schönherr, 2000, 2001) and charged organic molecules (Schönherr, personal communication) across cuticles isolated from several different plant species has been recently studied. Most interestingly, rates of cuticular penetration of these charged compounds were not significantly affected by temperature, since activation energies of permeation estimated from Arrhenius plots (P vs. $1/T$) were close to 0 kJ mol^{-1} (Schönherr, 2000). In contrast to water transport properties, wax extraction for ion transport had only a small effect on rates of penetration, as they increased by factors between 2 and 3. These findings are fundamentally different from those previously published on transport of water and lipophilic molecules across plant cuticles. Activation energies for these two compounds vary between 20 and 200 kJ mol^{-1} (Schönherr *et al.*, 1979; Baur *et al.*, 1997; Schreiber, 2001) and permeability always increased by factors from 10 to 10 000 upon wax extraction (Schönherr and Riederer, 1989). Furthermore, typical lipophilic plasticizers significantly increased the mobility of organic compounds and water in cuticles and waxes (Riederer and Schönherr, 1990; Schönherr *et al.*, 2001), whereas wax domains of the cuticle had no effect on the penetration rates of charged molecules (Schreiber *et al.*, 1996b). As observed with water permeability, the rates of penetration of inorganic ions increased with increasing humidity by factors of 2 to 3. Since hydrated charged molecules are neither soluble in oils nor in the lipophilic cutin and wax domains, nor can lose their hydration shells, these experimental results focusing on the transport of hydrated ions can only be explained when the existence of water-filled polar pores traversing the cuticles is postulated.

Wax movement: a simple solution?

In addition to organic non-electrolytes, water as well as ions, the transport of wax molecules through and onto the cuticles has gained special attention. The presence of epicuticular wax crystals has been continuously associated with the question of how these move through the cuticle. From a physicochemical point of view, the transport of wax compounds should follow the same rules as described for organic non-electrolytes, since cuticular waxes are either composed of linear, long-chain, aliphatic molecules or of pentacyclic, lipophilic aliphatic triterpenoids (Walton, 1990). After more than a century of intensive work, only recent studies have revealed some essential answers to the potential molecular mechanisms for wax transport through the plant cuticle (Riederer and Schreiber, 1995; Neinhuis *et al.*, 2001).

'Pores' or 'microchannels'

One old, yet still cited, model suggests wax transport across the cuticle via 'pores' or 'microchannels'. Deduced from the observation that single wax crystals lie a certain distance from the neighbouring crystals on the cuticle surface, numerous authors have postulated that channels or pore-like structures in the cuticle are involved in the process of wax movement (de Bary, 1871; Hall and Donaldson, 1962; Baker, 1982; Anton *et al.*, 1994). However, the concept of pores is inconsistent with the findings of transmission electron microscopy studies, where structures similar to such 'pores' show a greater electron-density

than the cutin matrix (Jeffree, 1996). Wax 'microchannels' were reported in freeze-etched cuticles of *Trifolium repens* L. and using carbon replica of *Brassica oleracea* var. *botrytis* L. cuticles, in addition to other species, by Hall (1967). Miller (1986) reported the existence of 'transcuticular pores' in leaves of more than 50 species but, unfortunately, his observations were made with the light microscope having a resolution which is inadequate to confirm the identity of such features (Jeffree, 1996).

Lipid transfer proteins

Lipid transfer proteins (LTPs) are small, abundant proteins, first identified in animals as possible intercellular carriers of lipids between membranes and organelles (see also van Dongen and Borstlap, Chapter 6). Later, they were found in microorganisms (Yamada, 1992) and plants (Sterk *et al.*, 1991), where they have been isolated from epidermal tissues and leaf surface waxes, e.g. of *Brassica oleracea* L. ssp. *oleracea* convar. *botrytis* (L.) Alef. var. *italica* Plenck (sprouting broccoli). This led to speculation that they may be involved in transporting cutin monomers and wax compounds across the aqueous cell wall phase (Pyee *et al.*, 1994; Kader, 1997). As the transport direction of LTPs is from the inside of the cuticle to the outside and the LTPs do not return for additional transfer, the number of lipid molecules to be transferred is limited (Post-Beitenmiller, 1996). As LTPs cannot explain all transport phenomena of the wax components across the cuticle, '... another, as yet unidentified mechanism must also be involved'. (Jeffree, 1996: 58).

Wax transport via diffusion

Model experiments analysing the transport properties of wax compounds in isolated plant cuticles (Baur *et al.*, 1996, 1997, 1999; Buchholz *et al.*, 1998; Buchholz and Schönherr, 2000) or in recrystallized cuticular waxes (Schreiber and Schönherr, 1993; Schreiber *et al.*, 1996a) from several plants species did show that waxes behave like lipophilic aliphatic and aromatic substances, supporting the conclusion that waxes behave in a similar way to organic non-electrolytes. Linear, long-chain alkanes, alcohols, fatty acids and cholesterol, serving as model compounds for triterpenoids, had mobilities in both cuticular waxes and isolated cuticular membranes in a similar range to that observed for typical organic compounds such as herbicides (Baur *et al.*, 1996; Schreiber *et al.*, 1996a). Typical plasticizers, which increase mobilities of organic non-electrolytes (Riederer and Schönherr, 1990) and water (Schönherr and Baur, 1994) diffusion across plant cuticles, had the same effect on characteristic wax compounds, e.g. tetracosanoic acid, where mobility was increased in recrystallized cuticular wax of *Hordeum vulgare* L. (barley) by factors ranging from 22 to 315 (Schreiber, 1995; Schreiber *et al.*, 1996b). From these experimental observations, it can be concluded that wax molecules themselves can passively diffuse within the lipophilic cutin matrix and wax domains along the concentration gradient, from the inner side to the sink, in the cutin matrix (intracuticular waxes) and onto the surface (epicuticular waxes).

Yet another possible mechanism for wax transport was proposed by Jeffree *et al.* (1976) and Baker (1982). These authors discussed diffusion of the wax precursors with an unknown carrier solvent through the cuticle with the wax components crystallizing after evaporation of the solvent. More recently, Neinhuis *et al.* (2001) argued that the solvent permeating through the cutin polymer could be water vapour. Thereby the cuticular transpiration is the major driving force for the movement of wax through the plant cuticle. However, this model requires the assumption that the lipid wax components interact with water. This could possibly result from a process similar to steam distillation in chemistry

where substances with high boiling-points (e.g. hydrocarbons) can be separated by water vapour at ambient pressure so long as they are insoluble, or virtually insoluble, in water (Onken and Weiland, 1984; Falbe and Regitz, 1999). This holds true for the wax components. During this distillation process, the partial pressure of the water vapour is added to the partial pressure of the wax components, therefore their mobility should increase, allowing a facilitated movement through the lipophilic cuticle. As advantageous as the model of co-transport of the cuticular waxes with water sounds, up until now not all phenomena are properly understood and data concerning the general kinetics are lacking.

Interactions with the biotic and abiotic environment

Due to their sessile nature, plants have evolved defence and resistance strategies that are adapted to the selective pressures of offending organisms such as pathogenic fungi, bacteria or herbivorous animals in different environments (Edwards, 1992). Generally, colonization of microorganisms or feeding by herbivores on plant surfaces involve constitutive factors like contact and adhesion, recognition markers based on surface chemistry, cuticle thickness and toughness as well as tissue toughness, or is set by environmental factors such as the pH of surface water, water availability over time, wind exposure and speed as well as temperature (Southwood, 1986; Bernays, 1991; Juniper, 1991; Butler, 1996). With regard to the problem of adhesion for example, the offending pathogens and insects have to face self-cleaning and slippery plant surfaces caused by epicuticular waxes (Knoll, 1914; Barthlott and Neinhuis, 1997; Markstädter *et al.*, 2000). Additional hindering of successful growth or feeding is evoked by a range of inhibitory substances metabolized by plants, toxic for both pathogens and herbivores (Juniper, 1991; Zangerl and Berenbaum, 1993). Among these antimicrobial substances, mainly phenols and terpenoids, and to a lesser extent organic acids, flavones, polyacetylenes and alkaloids can be found. All appear to have three sources of origin: (1) deposited on the plant surface via diffusion; (2) associated with waxes; or (3) stored in glandular trichomes (Weinhold and Hancock, 1980; Blakeman and Atkinson, 1981). Again, the epicuticular waxes play a particularly important role in plant-animal interactions where the lipid components can determine growth and feeding (Woodhead and Chapman, 1986; Schwab *et al.*, 1995; Espelie, 1996). Instead of constitutive plant defence mechanisms, induced ones are based on cellular cascade mechanisms, e.g. hypersensitive response (Strange, 1992).

However, microorganisms themselves have developed strategies for establishment on their host plants, such as attaching structures (appressorium) (Wheeler, 1981), contact mediating proteins such as hydrophobins (Wessels, 1994), slime production often including glycoproteins to form adhesion pads (Mendgen, 1996) and an enzymatic repertoire of cutinases and esterases (Kolattukudy, 1985). Herbivores, particularly insects, exhibit tarsal pulvilli or mucilaginous secretion (Eigenbrode, 1996), morphological adaptations of the mandibles (Bernays, 1991), or pathways to metabolize toxic plant deterrents (Zangerl and Berenbaum, 1993). The interactions between defending plants and insects have been categorized to be co-evolutionary based on selective pressure on plant secondary metabolites. However, this is not necessarily correlated since plants are subject to multiple selective pressures from vertebrate to invertebrate herbivores, pathogens, other plants and from the abiotic environment (Edwards, 1992).

Interactions between insolation and UV-B radiation and the plant cuticle must also be noted. Evidence is strong that in visible (400–700 nm) and in infrared light (700–3000 nm), leaf epicuticular waxes increase reflectance of solar radiation leading to

reduced photoinhibition of photosynthesis, reduced rates of transpiration and thereby increased leaf water-use efficiency (Eller, 1979; Robinson *et al.*, 1993; Barnes and Cadoso-Vilhena, 1996). Ultraviolet radiation (280–400 nm), especially UV-B, appears to be rather attenuated by flavonoids incorporated in the cuticle matrix or the surface waxes (Day *et al.*, 1993; Krauss *et al.*, 1997) and may have stimulating effects on epicuticular wax production and cuticle thickness (Steinmüller and Tevini, 1985; Givnish, 1988; Barnes *et al.*, 1994). The same holds true for visible radiation (Hallam, 1970; Reed and Tuckey, 1982).

Water repellency and self-cleaning property

Water repellency

In addition to the hydrophobic nature of the cuticle, the presence of epicuticular wax crystals often leads to water repellency instead of wetting. This is due to their hydrophobic nature and microroughness of about 1–5 μm (Baker, 1982; Jeffree, 1986). This is frequently observed in the garden cabbage patch where the leaves of *Brassica oleracea* L. are simply not wetted during rainfall. The physical principles of surface wetting were solved at the beginning of the last century and summarized by Wenzel (1936) and Cassie and Baxter (1944). Basically, roughening a hydrophilic solid increases wetting, in contrast, roughening of a hydrophobic solid causes water-repellency. Since then, the wetting properties of surfaces have been studied intensively and later reviewed in the fields of biology and physics (Holloway, 1970; Adamson, 1990; Myers, 1991; Herminghaus, 2000). Whenever a liquid is applied on a solid, the process of wetting involves three different interfacial boundaries, i.e. solid–liquid, solid–air and liquid–air. The epicuticular waxes minimize the contact area between water (liquid) and the plant surface (solid) by the combination of hydrophobic chemistry and microroughness and form an enlarged water/air interface, thus constituting a composite surface with air enclosures between the epicuticular wax crystals (Dettre and Johnson, 1964). On such ‘low energy’ surfaces water forms spherical droplets due to the surface tension and rest on the outermost tips of the wax crystals, a phenomenon called water repellency. Water repellency is generally expressed as the contact angle θ [°] between the water droplet and the surface (Holloway, 1970). A contact angle of 0° implies complete wetting, while an angle of 180° describes complete non-wetting, but neither extreme is apparent in plants. Barthlott and Neinhuis (1997) categorized the contact angle of wettable leaves <110°, whereas water-repellent species often display contact angles >150°.

Influence of biotic and non-biotic factors on water repellency

As mentioned above, plant surfaces are living habitats for microorganisms such as pathogenic fungi and bacteria, all of which have more or less pronounced effects on wetting (Martin, 1964; Dickinson, 1976; Juniper, 1991). Knoll and Schreiber (2000) have shown that epiphytic microorganisms can mask the native wetting properties of leaf surfaces and discuss a pH-dependent decrease of contact angles due to carboxylic groups of the microorganisms (Bunster *et al.*, 1989).

As well as organisms, plant surfaces are subject to large quantities of airborne pollutants such as organic and inorganic dust. These contaminants are the product of natural erosion or have anthropogenic origin (Pye, 1987) and may cause considerable damage to leaf surface morphology as well as influencing physiological processes, depending upon their size and chemical nature (Crossley and Fowler, 1986; Eveling, 1986; Farmer, 1993).

It was shown by Eller (1977, 1985) that road dust deposits lead to increased leaf temperature under insolation and result in reduced photosynthesis and higher transpiration rates, as well as influencing stomatal diffusive resistance (Flückiger *et al.*, 1979). Direct impact on plant surfaces and their waxes derives from mechanical abrasion due to dust particles or even snow crystals (van Gardingen *et al.*, 1991; Grace and van Gardingen, 1996).

The influence of acidic rain or ozone on the micromorphology of plant surfaces has been extensively studied and reviewed in forest decline research (Mengel *et al.*, 1989; Turunen and Huttunen, 1990; Percy *et al.*, 1994; Huttunen, 1996) but no uniform picture could be drawn from the findings. Whereas several authors claim that air pollutants cause severe damage, e.g. destruction of waxes caused by chemical degradation processes (Mudd *et al.*, 1982), other researchers reached the conclusion that the observed effects cannot be separated from natural environmental influences (Grill *et al.*, 1987; Euteneuer-Macher, 1990; Neinhuis *et al.*, 1994; Neinhuis and Barthlott, 1998) or even could not be found (Riederer, 1989). Burkhardt *et al.* (2001) argue that fused wax patterns described in forest decline research are more likely caused by deliquescent hygroscopic aerosol particles. It can be concluded that air pollution effects can alter the structural and physiological appearance of epicuticular waxes at least in combination with other factors, but strongly depend on the species investigated (Kim and Lee, 1990).

Concerning pesticides, influences of tensides, which enable the uptake of active ingredients by decreasing the surface tension of water (Stevens and Bukovac, 1985; Knoche and Bukovac, 1993), can cause considerable damage to the wax ultrastructure and thus decreased water repellency (Noga *et al.*, 1987, 1991; Wolter *et al.*, 1988). Contrary to the aim of pesticide application, contaminating particles including spores and conidia are also found within the areas of altered waxes, a condition that enhances the probability of infection (Neinhuis *et al.*, 1992).

Self-cleaning property: the 'lotus-effect'

Despite numerous types of potential contamination, there are several plants that appear to be almost completely clean throughout the year. This can be most impressively demonstrated with the large peltate leaves of *Nelumbo nucifera* Adan. (sacred lotus, Figure 10.8), which are characterized by large epidermal papillae densely covered with wax tubules (Barthlott *et al.*, 1996). There has been a vague knowledge of the correlation between water repellency and reduced contamination for more than a century (Lundström, 1884), but enlightenment in this aspect can be assigned to Barthlott and co-workers (Barthlott and Ehler, 1977; Barthlott and Wollenweber, 1981; Barthlott, 1990; Barthlott and Neinhuis, 1997), who named this self-cleaning ability of plant surfaces the 'lotus-effect'. After screening the leaf surfaces of some 15 000 species by electron microscopy, it can be demonstrated that microrough water repellent plant surfaces display the ability to self-clean, whereas smooth wettable cuticles accumulate particulate contaminations. This holds true even for very small particles, independent of their chemical nature. Again, minimization of the contact area is the clue, here, between particle and plant surface or epicuticular waxes, respectively, leading to a quantitative higher contact area between water droplet and particle. The consequence is greater adhesion of particles to the water droplet instead of to the plant surface (Barthlott and Neinhuis, 1997). Thus, depositions are removed from the plant surface by rainwater droplets and even dew droplets are able to clean such leaves (see Figures 10.9 and 10.10). During recent studies on structural parameters for an optimized 'lotus-effect', evidence was found that a single surface microstructure built up by epicuticular waxes and a secondary, larger scaled papillose cellular



Figure 10.8 The Lotus plant (*Nelumbo nucifera* Adan.), a symbol of purity and model for technical surfaces with a self-cleaning property.

structure underneath the waxes are necessary, as characterized for *Nelumbo nucifera* Adan. (Wagner *et al.*, 2002).

Hence, double-structured plant surfaces represent an optimized compromise. Epicuticular waxes are fragile and thus limit the self-cleaning property of plants. As outlined above, the waxes can be altered, destructed or removed mechanically. To some extent plants can regenerate their coverage to compensate for the damage (Hallam, 1970;

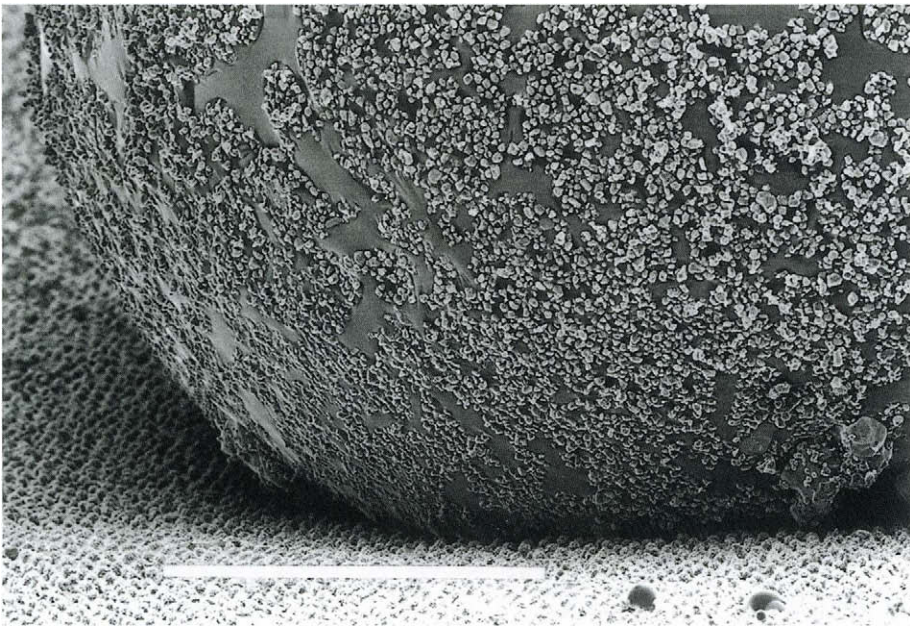


Figure 10.9 Droplet of saturated sugar solution on the papillose adaxial epidermal surface of *Alocasia macrorrhiza* (L.) G. Don f., demonstrating the 'lotus-effect'. Contaminating particles adhere more to the surface of the droplet than to the plant's surface. Scale bar = 500 μm .

Wolter *et al.*, 1988) but this is not ubiquitous. Considerable differences in the ability to regenerate wax were discussed with respect to seasonal changes of leaf contamination of *Fagus sylvatica* L., *Quercus robur* L. and *Ginkgo biloba* L. by Neinhuis and Barthlott (1998). Recently, Neinhuis *et al.* (2001) studied the regeneration ability of 24 plant species and classified four categories: (1) regeneration occurs at all stages of development; (2) regeneration occurs only during leaf expansion; (3) regeneration occurs only in fully developed leaves; and (4) plants were not able to regenerate wax at all. What is the strategy of these plants that do not regenerate their epicuticular wax crystals? Conifers and evergreen as well as deciduous angiosperm trees can be generally distinguished by three different surface syndromes via the evolution of two protection mechanisms, either the 'lotus-effect' or the development of a thick cuticle (Neinhuis and Barthlott, 1997). Moreover, several species replace the surface waxes by developing thick cuticles during ontogeny.

The biological implications of the self-cleaning property are obvious – contaminations of multiple origin and their impacts are effectively excluded. Moreover, the 'lotus-effect' plays an important role in the green arms race against pathogens. The availability of water is a crucial factor for adhesion, germination and growth of spores and conidia (Rogers, 1979; Campbell *et al.*, 1980; Juniper, 1991). Due to extremely low water-capacity, self-cleaning plant surfaces are virtually dry, a condition not very beneficial for the majority of plant invaders, except for a few pathogens causing powdery mildew (Wheeler, 1981). But the self-cleaning property is not only restricted to plants. It can also be found on large insect wings that cannot be cleaned by legs (Wagner *et al.*, 1996). In this case, the maintenance of flight capability seems to be the major evolutionary impetus. Thus, the 'lotus-effect' has an overall biological importance, which is based only on physicochemical properties of natural surfaces. It was therefore possible to initiate a joint project together with several

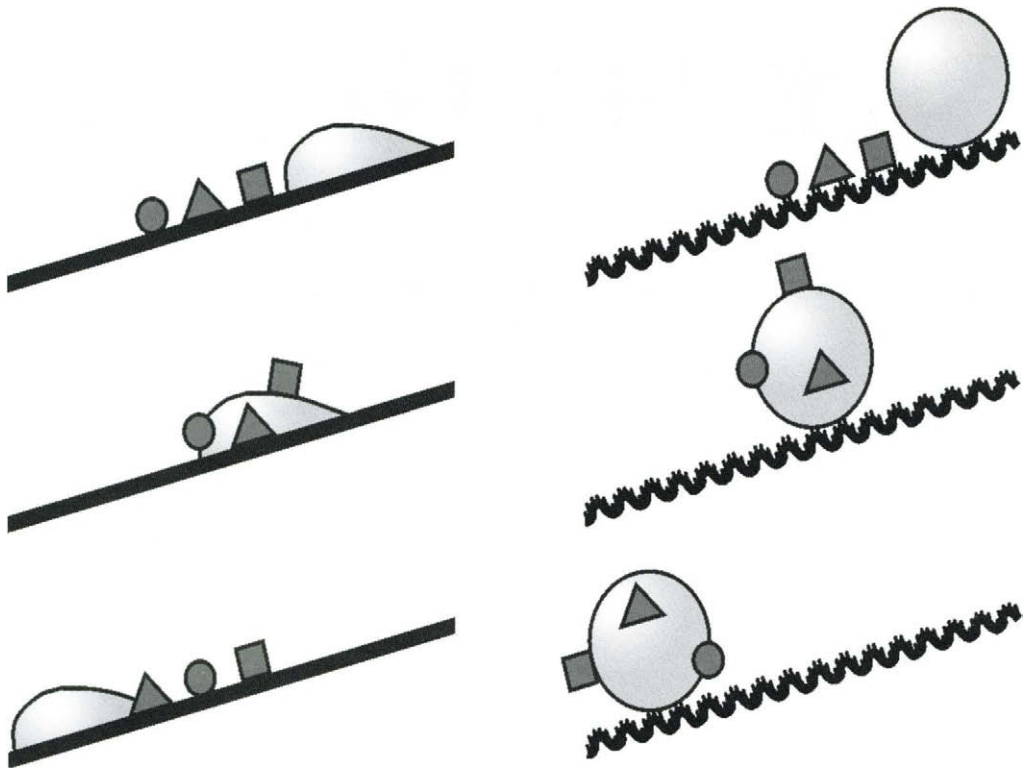


Figure 10.10 Schematic diagram summarizing the 'lotus-effect'. (Left) On smooth surfaces particles are mainly distributed by water, (right) while they adhere more strongly to the water droplet than to the rough hydrophobic surface on 'lotus-effect' leaves and are efficiently removed when the droplets roll off.

industrial companies to transfer the 'lotus-effect' seen in plants, to biomimetic self-cleaning technical products, such as wall paint or roof tiles.

Biomechanical properties

As mentioned above, the plant cuticle can also be seen as the first mechanical barrier against microorganisms and herbivores. The overall cuticle thickness can be very great, even more so than the epidermal cell wall, which in fact may be heavily encrusted (Fritz, 1935; Holloway, 1994). From a mechanical point of view, the location at the outer perimeter of plant organs and its rigid appearance, at least in succulents, indicates that the cuticle may function as external structural element that possibly adds mechanical support for tissue integrity and impacts on morphogenetic processes (Edelmann and Neinhuis, 1997), as is proposed for bark (Niklas, 1999). Hoffmann-Benning and Kende (1994) showed that the cuticle of rice coleoptiles is under tension during coleoptile elongation and thus could provide a mechanical constraint on organ growth. The mechanical properties of the cuticle, especially of fruits, are of commercial significance. To prove this, surface cracking studies on tomato or cherry (*Prunus avium* L.) fruits have been carried out but none of these distinguished between epidermis and cuticle (Voisey *et al.*, 1970; Sekse, 1995). Up to now, only very few reports have addressed the biomechanical properties

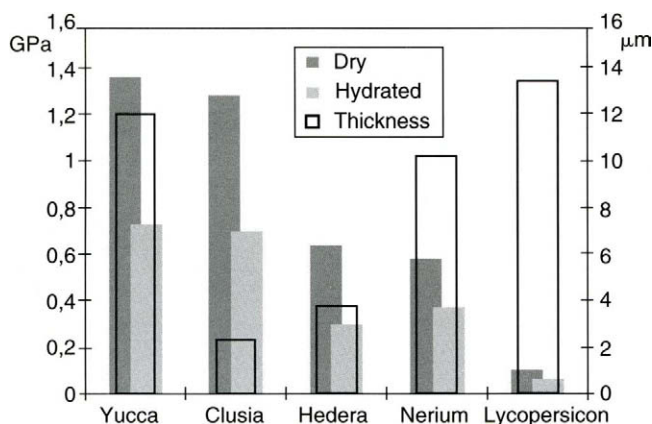


Figure 10.11 Young's moduli and thicknesses of isolated plant cuticles of leaves of *Yucca aloifolia* L., *Clusia fluminensis* Planch. & Triana, *Hedera helix* L., *Nerium oleander* L. and fruit of *Lycopersicon esculentum* Mill. Hydration causes a decrease in stiffness of about 35–50%.

solely of the plant cuticle in an experimental approach (Pescareta *et al.*, 1991; Hasenstein *et al.*, 1993; Wiedemann and Neinhuis, 1997), whereas the literature on general plant biomechanics does not refer to cuticles (Niklas, 1992). Hence, available data are rare, although vague indications of significance can be found in the literature. Wiedemann and Neinhuis (1997) studied the mechanical properties of isolated cuticles from five leaf and one fruit species by means of a one-dimensional tension test and reported Young's modulus – a measure of stiffness – ranging between 0.6 and 1.3 GPa for the leaves and 0.1 GPa for tomato fruit (Figure 10.11). Direct comparison with polyethylene (0.2 GPa) reveals remarkably high stiffness values for these cuticles (Czichos, 1989). Hydration of the cuticles caused a decrease in the Young's modulus of about 35–50%. As a consequence, water acts as a plasticizer, a fact supported by studies focusing on the decrease in the surface elastic modulus of isolated tomato fruit cutin in relation to water content (Round *et al.*, 2000). This is most likely to occur if water binds to polar groups or disrupts hydrogen-bonded cross-links in the cutin matrix (Dominguez and Heredia, 1999; Round *et al.*, 2000). Recently, Marga *et al.* (2001) undertook chemical analyses of cuticles from different organs, each having different biomechanical properties, and found that rigid cuticles can be classified by C₁₆ cutin monomers, while elastic cuticles correspond to mixed C₁₆/C₁₈ cutin monomers. During tomato fruit ripening, the elasticity of the cuticle seem to decrease as the stiffness increases (Bargel *et al.*, 2000; Bargel and Neinhuis, unpublished data), an aspect that may be determined by a decrease in the relative amount of trihydroxy C₁₈ fatty acids (Baker *et al.*, 1982). Thus, hydroxyl groups seem to enhance the hydrophilic character and the hydration state of the cutin matrix, which in turn results in higher plasticity.

The few results outlined here could be seen as the first steps towards an understanding of cuticle biomechanics. Even from an engineering approach, the outer plant coverage appears to be a highly diverse composite with biomechanical properties determined by its dynamic structure and chemical composition. From these, the cuticle may inherit a possible role in stabilization of plant organs and thus a newly described function that has been largely neglected so far to date.

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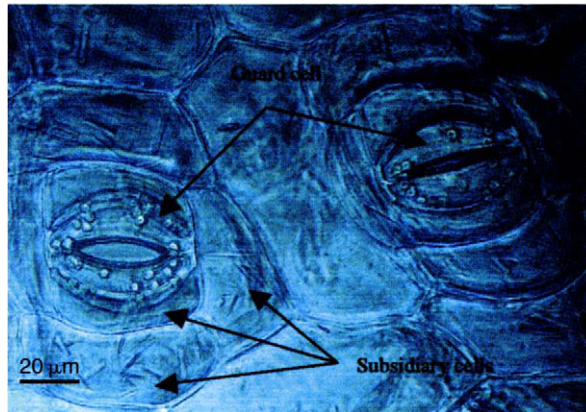


Plate 1 A reflected light image of a section of epidermal peel taken from *Commelina communis* showing two stomatal complexes within the epidermis.

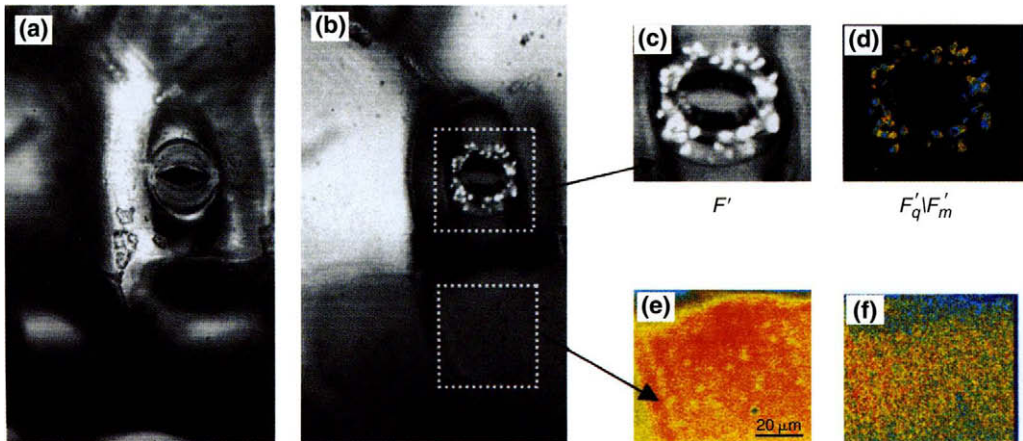


Plate 2 Example of the use of high resolution imaging of chlorophyll fluorescence in guard cells in *Tradescantia* (a) reflected light image, (b) whole fluorescence image, (c) and (e) isolated areas of mesophyll and guard cells showing steady state fluorescence images, (d) and (f) built up image of F_q'/F_m' .

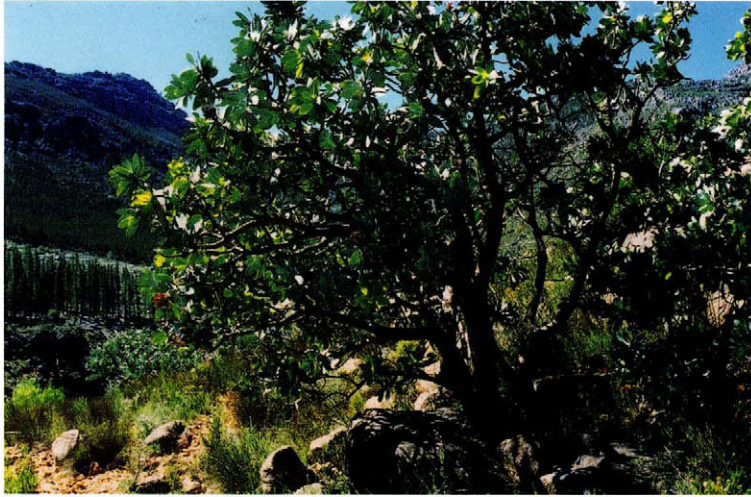


Plate 3 Three growth forms of the Fynbos Mediterranean ecosystem: the deeper rooted proteoid growth form, *Protea nitida* Mill. (top), the more shallow rooted ericoid form, *Phaeocomma prolifera* (L.) D.Don. (bottom left) and also the shallow rooted restioid form, *ELEGIA juncea* L. (bottom right).



Plate 4 The open vegetation of the Fynbos, characterized by its low density with representatives of the families Restionaceae, Compositae, Ericaceae and Proteaceae (left). Vertical distribution of the open vegetation is accentuated by *Erica chloroloma* Lindley (right) and other species.



Plate 5 Three *Pelargonium* species with entire, deeply lobed and parsley-like leaf forms. From left to right: *P. cucullatum* (L.) L.'Her., *P. scabrum* (L.) L.'Her. and *P. triste* (L.) L.'Her.

Falling atmospheric CO₂ – the key to megaphyll leaf origins

Colin P Osborne, William G Chaloner
and David J Beerling

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Introduction

The vast majority of contemporary leaves are megaphylls, derived from the ‘planation’ of determinate, overtopped side-branch systems and their ‘webbing’ by a thin lamina of photosynthetic mesophyll tissue, a sequence first documented as the telome theory (Figure 11.1; Zimmermann, 1930, 1952). Since the independent occurrence of this evolutionary innovation in four clades around 360 million years ago (Ma) (Boyce and Knoll, 2002), megaphyll leaves have become such a successful and ubiquitous feature of plants that it is difficult to imagine a world without them. Primary production by these organs is the energy source for almost all terrestrial life, with tetrapod and arthropod foliar herbivores key components in most modern ecosystems. Similarly, leaf carbon, energy and water exchanges are key driving steps in biogeochemical cycles and the regional climate system. Despite this pivotal role in ecosystem energy flows, laminate megaphyll leaves did not become widespread in plant assemblages until the Late Devonian, some 40–50 million years after the first appearance of their vascular ancestors on land (Figure 11.2) (Chaloner and Sheerin, 1979). Quite why this evolutionary change took so long to accomplish has hitherto been unclear (Robinson, 1991; van der Burgh, 1996; Niklas, 1997).

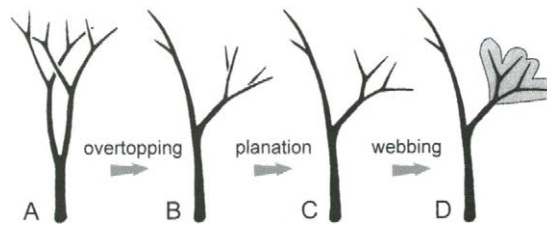


Figure 11.1 Stages in the evolution of the megaphyll leaf, as documented by the telome theory (Zimmermann, 1930, 1952). A. The ancestral form – a dichotomizing axis branching in three dimensions (e.g. rhyniophytes); B. Evolutionary ‘overtopping’ produces a main axis bearing reduced, lateral, determinate, photosynthetic stem systems, each branching in three dimensions (e.g. trimerophytes); C. ‘planation’ flattens these lateral systems of terete stem segments to a single plane (e.g. some cladoxylaleans); D. ‘webbing’ joins the segments of lateral branches with a lamina of photosynthetic mesophyll tissue (e.g. some progymnosperms).

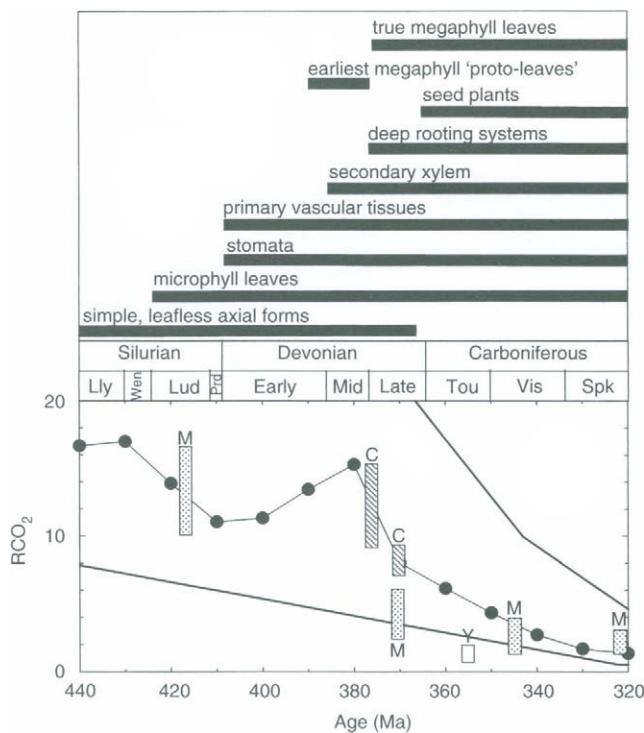


Figure 11.2 Evolutionary innovations of early terrestrial plants (upper panel) and concurrent changes in atmospheric CO_2 (lower panel). Intervals for the emergence and duration of characteristics in the fossil record were obtained for: axial forms, stomata, microphylls, primary vascular tissues (Edwards and Wellman, 2001); secondary xylem, megaphylls (Chaloner and Sheerin, 1979); ‘proto-leaves’ (Berry and Fairon-Demaret, 2001; Hao and Gensel, 2001); root systems (Raven and Edwards, 2001); and seed plants (Chaloner *et al.*, 1977). Periods and epochs of the geological timescale are shown for reference, with the abbreviations: Lly, Llandovery; Wen, Wenlock; Lud, Ludlow; Prd, Pridoli; Tou, Tournaisian; Vis, Viséan; and Spk, Serpukhovian (Harland *et al.*, 1990). Modelled atmospheric CO_2 concentration, relative to today’s value (RCO_2), is plotted with a range of error (bold lines) based on sensitivity analyses (Bernier and Kothavala, 2001). RCO_2 ranges estimated from the carbon isotope composition of palaeosols (C, Cox *et al.*, 2001; M, Mora *et al.*, 1996; Y, Yapp and Poths, 1996) are also shown.

The earliest vascular land plants consisted of dichotomizing, leafless, cylindrical (terete) axes, but megaphyll evolution soon gave rise to determinate, non-laminate but presumably photosynthetic, lateral branch systems in their successors (Edwards and Wellman, 2001; Figure 11.2). The deeply lobed laminate leaves of the enigmatic Early Devonian fossil *Eophyllophyton bellum* Hao and Beck (1993) demonstrate that such evolution had occurred by around 390 Ma, although most plants at this time were still leafless or possessed microphyll leaves (Gensel *et al.*, 2001; Figure 11.2). Laminate megaphyll leaves had therefore clearly evolved in at least one lineage by the Early Devonian, but were not widely adopted until the appearance of the first forests in the Late Devonian (Gensel and Andrews, 1984; see Figure 11.2). This delay is puzzling, because the Devonian period witnessed far more complex polyphyletic evolutionary innovations in other plant structures (see Figure 11.2): the development of the seed habit and heterospory from homosporous ancestors (Chaloner *et al.*, 1977) and the rise of arboreal forms from tiny herbaceous ancestors (Chaloner and Sheerin, 1979). The latter feature, in itself, necessitated the origin of extensive root systems (Algeo and Scheckler, 1998; Algeo *et al.*, 2001; Gensel *et al.*, 2001; Raven and Edwards, 2001). It therefore seems appropriate to enquire why the widespread adoption of laminate megaphyll leaves took so long to come about.

The close coupling between leaves and the atmosphere suggests a likely role for some change in the aerial environment in early megaphyll evolution. A dramatic drop in atmospheric CO₂ is a good candidate, because simulations of the long-term geochemical carbon cycle indicate a fifteen-fold reduction during the Devonian (Bernier and Kothavala, 2001). These model predictions are supported by independent estimates of atmospheric CO₂ (see Figure 11.2) using the stable carbon isotope composition of fossil soils (Mora *et al.*, 1996; Yapp and Poths, 1996; Cox *et al.*, 2001). We previously proposed a causal link between the delayed evolution of megaphyll leaves and this pronounced decline in atmospheric CO₂ (Beerling *et al.*, 2001a), mediated on a timescale of at least 40 Ma by the tight inverse relationship between the density of stomata on photosynthetic structures and atmospheric CO₂ concentrations (Beerling and Royer, 2002).

This chapter further details the mechanism of this linkage and the distinctive trends in early megaphyll leaf evolution expected to arise from it. The validity of these expected trends is explored by a preliminary quantitative analysis of megafossils, using published works on the Devonian and Early Carboniferous fossil record. In this analysis, we have focused on the origin, occurrence and form of megaphyll leaves to provide a preliminary test of the Beerling *et al.* (2001a) hypothesis.

A mechanism coupling Devonian megaphyll evolution with falling CO₂

In developing our hypothesized link between CO₂ and plant evolution we have used a model of leaf biophysics and physiology to investigate the functional consequences for Devonian megaphylls of the evolutionary relationship between stomatal density and atmospheric CO₂ (Beerling *et al.*, 2001a) (Figure 11.3). The model mathematically describes the key interactions between photosynthetic CO₂ fixation, stomatal conductance, transpiration and the leaf energy balance (Beerling and Woodward, 1997). It explicitly accounts for the influences of temperature, light, humidity and CO₂ on these processes, and their feedbacks operating via CO₂ concentration in the sub-stomatal cavities, leaf temperature and the leaf–air vapour pressure deficit.

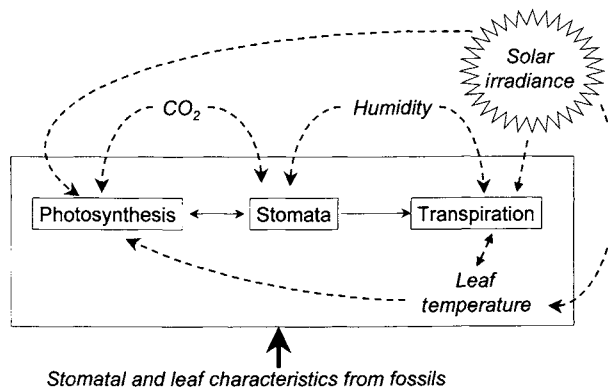


Figure 11.3 Model of leaf biophysics. A biochemical model of leaf photosynthesis is fully coupled with a model of stomatal functioning and leaf energy balance, incorporating the environmental feedbacks from atmospheric humidity, temperature and CO_2 (Beerling and Woodward, 1997). The stomatal model is initialized using measurements of density and pore dimensions from fossil cuticles, and simulations of the leaf laminar boundary layer (see Figure 11.4C) utilize estimates of fossil leaf dimensions (for details, see Beerling *et al.*, 2001a).

Leaf temperature in the model is determined by the absorption of solar energy, and the extent to which this is dissipated through: (1) the emission of longwave radiation; (2) the loss of latent heat by the evaporation of water in transpiration; and (3) the transfer of sensible heat to air flowing across the leaf surface. Stomatal density and pore size are key inputs into the model (see Figure 11.3) because they constrain the maximum stomatal conductance and therefore the rates of transpiration and latent heat loss. Leaf size is also a critical model input because it influences leaf boundary layer characteristics and hence the transfer of sensible heat to the atmosphere. A detailed model description is provided by Beerling and Woodward (1997).

Model simulations show that hypothetical large megaphyll leaves would have provided no selective advantage over branched axes for photosynthesis in the Early Devonian high CO_2 atmosphere (Beerling *et al.*, 2001a). This is because the exceptionally low stomatal densities observed for this period (Edwards, 1998) would have restricted transpiration rates in these leaves, and so dissipated very little absorbed solar energy as latent heat (Figure 11.4B), with a high associated risk of lethal overheating (Beerling *et al.*, 2001a). Efficient absorption of solar energy for photosynthesis by laminate leaves (Figure 11.4A) may therefore have carried a fatal cost.

By contrast, the erect forms of early axial land plants protected them from overheating by minimizing solar energy absorption around midday (Figure 11.4A), when the sun is high in the sky and air temperature approaches its maximum. This ecological strategy is used to good effect by some modern aridland plants, which have secondarily evolved reduced, erect leaves (Valladares and Pearcy, 1997) or photosynthetic stems (Haase *et al.*, 1999). Further protection from overheating in early plants could have been obtained by small, cylindrical, or highly dissected photosynthetic surfaces, which minimize the thickness of the laminar boundary layer and therefore aerodynamic resistance to sensible heat dissipation (Gurevitch and Schuepp, 1990; Figure 11.4C).

This mechanism leads us to predict that the earliest megaphyll leaves would have been erect, small and/or highly dissected (Beerling *et al.*, 2001a). In the absence of significant cooling through sensible and latent heat fluxes, we calculate that the effective solar energy

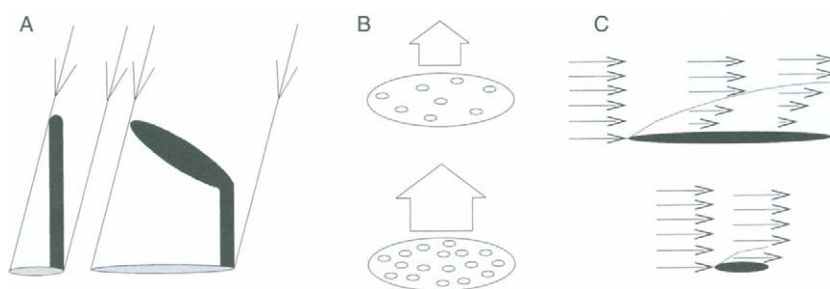


Figure 11.4 Mechanisms coupling leaf morphology and stomatal density with energy balance. If energy gained by the absorption of solar radiation exceeds that lost as latent heat and longwave radiation, leaf temperature will rise (Gates, 1979). A. Interception of solar energy is lower in an erect axis (left) and it therefore casts a smaller shadow (grey) than a more horizontal laminate leaf (right) when the sun is high in the sky. B. Rising stomatal density leads to greater stomatal conductance and therefore transpiration (top to bottom) indicated by relative size of the arrows. This higher transpiration rate increases the energy lost as latent heat. C. Air flowing over the leaf surface is slowed by friction (wind velocity indicated by length and direction of arrows), forming a laminar boundary layer that impedes energy losses as latent heat and convection. This resistance to energy loss increases with the thickness of the layer, which is greater for a large leaf (top) than a cylindrical stem, or a small (bottom) or dissected leaf (both examples are viewed in cross-section).

interception by large megaphylls would have caused lethal overheating. However, drastic evolutionary increases in stomatal density tracking the decline in Late Palaeozoic CO₂ concentration would have markedly increased latent heat loss, thus reducing the inherent biophysical constraints for developing larger and more entire megaphylls.

Our calculations show that, as stomatal density rose in response to falling CO₂ levels through the Devonian and Early Carboniferous, large laminate megaphyll leaves increasingly gained a selective advantage in terms of carbon gain by photosynthesis, without the penalty of high temperature damage. The advantage to photosynthesis was gained through a higher leaf conductance to CO₂ and more effective absorption of solar energy by the planated and more horizontal lamina (see Figure 11.4A). Higher stomatal densities would also have permitted greater transpiration rates, dissipating more absorbed solar energy as latent heat (Figure 11.4B), thereby lowering leaf temperatures and reducing the risk of lethal overheating (Beerling *et al.*, 2001a). It is important to note that cooling by transpiration is a fortuitous, but inevitable, consequence of stomatal guard cell opening for CO₂-fixation, rather than a prescribed primary function for stomata in our model (Beerling *et al.*, 2001b). The mechanism is widely accepted as a cooling device in leaves (Gates, 1968; Burke and Upchurch, 1989). Experiments using soybean clones differing in stomatal density provide strong support for our proposed linkage between cooling and stomatal characteristics. Rapid transpiration rates significantly reduced leaf temperatures in a high stomatal density soybean clone compared with those of its low stomatal density counterpart (Tan and Buttery, 1995).

Cooling by transpiration is an ecological strategy adopted by large-leaved plants in today's deserts, savannah, Mediterranean and riparian zones, and may be critical in avoiding lethal overheating during exposure to high solar radiation (Gates, 1979; Ehleringer, 1988; Matsumoto *et al.*, 2000). Latent heat losses from these rapidly transpiring leaves may lower their temperature by up to 15°C below that of the air (Lange, 1959). By contrast, leaves coated experimentally with a compound designed to prevent transpiration quickly

reach lethal temperatures under these conditions (Lange, 1959). Although the application of such 'anti-transpirants' may alter leaf surface properties such as absorptance, significant cooling by transpiration has also been demonstrated using independent techniques causing stomatal closure, such as the exogenous application of abscisic acid (Kitano *et al.*, 1995).

The cooling strategy requires an adequate water supply and rapid rates of uptake and vascular transport (Beerling *et al.*, 2001b). Studies of stem conductance in extant species with a similar primitive stelar anatomy to those of early land plants, show that these would have been insufficient to meet the high hydraulic demands of laminate leaves simulated by our model (Beerling *et al.*, 2001a). The requirement for effective water transport would therefore have provided a strong selection pressure for the coevolution of xylem tissues and leaves in Late Devonian plants. The fossil record supports this prediction, showing that the simulated increase in water demand was met by the evolution of secondary xylem tissues (see Figure 11.2), and an increase in the complexity of stelar anatomy (Niklas, 1997) and root systems (Algeo and Scheckler, 1998; Raven and Edwards, 2001) in several clades. However, the appearance of leaves is unlikely to have been the sole selection pressure acting on the evolution of Devonian vascular anatomy. Selection for the efficient operation of xylem tissues in mechanical support (Niklas, 1994) and the transport of mineral nutrients for growth (Niklas, 1997) would also have been driven by an increase in plant height and size. An evolutionary trend in height and size is observed in plant fossils throughout the Devonian (Chaloner and Sheerin, 1979), and would have increased fitness by promoting the dispersal of propagules and minimizing the shading of photosynthetic organs (Niklas, 1997).

Our simulations therefore provide a biologically plausible mechanism for the observed delay in megaphyll leaf evolution and suggest strong selection pressures for Devonian stem evolution. Critically, they also make important predictions about the morphology of the earliest leaves that are testable using the plant fossil record. These predictions are characterized by two key expectations:

1. Biophysical constraints would have restricted the earliest megaphylls to small or highly dissected shapes, which maximize the dissipation of absorbed solar energy (see Figure 11.4C).
2. Concurrent changes in atmospheric CO₂ and stomatal density during the Devonian would have permitted the evolution of larger and less dissected laminate megaphylls by increasing the energy lost in transpiration (see Figure 11.4B).

We next turn to evidence from the fossil record in an effort to test these expectations, using the first quantitative analysis of leaf size in Late Palaeozoic fossil floras.

Early evolution of the megaphyll leaf

The earliest tracheophyte land plants had neither lateral branches nor leaves, and consisted merely of naked, dichotomous-branching, three-dimensional stem systems (Edwards and Wellman, 2001). Their simple axial form is typified by *Cooksonia* Lang (Edwards *et al.*, 1983), initially appearing as macrofossils in the Mid-Silurian plant fossil record (Wenlock), but inferred from earlier spore assemblages (Edwards and Wellman, 2001). The discovery of stomata in Late Silurian (Pridoli) plants with a similar morphology is strongly suggestive of photosynthetic function in these axes, with gas exchange occurring through stomatal openings in an otherwise impermeable cuticle (Edwards *et al.*, 1996).

The evolution of primary xylem tissues appeared, significantly, at the same time (see Figure 11.2; Edwards *et al.*, 1992), facilitating the delivery of water to photosynthetic tissue, and permitting the evaporation that must inevitably accompany stomata-based CO₂-fixation in a land plant. Rhyniophyte species with the leafless axial form diversified and became common in the Late Silurian–Early Devonian (Edwards and Wellman, 2001). Although rare after the Early Devonian, they did not entirely disappear until the close of the Devonian with the widespread appearance of megaphyll leaves and rise of arborescence (Berry and Fairon-Demaret, 2001).

In contrast with the slow arrival of megaphylls, microphylls make an early appearance in the terrestrial plant fossil record – for many years *Baragwanathia longifolia* Lang and Cookson (1935), with its covering of narrow, linear microphylls, was the earliest known land plant (Ludlow-Lochkovian). Microphylls are presumed to have evolved from the enlargement of tiny spine-like ‘enations’ (Bower, 1935) and, despite their name (literally ‘small leaves’), are defined by their simple, usually unbranched venation and distinctive vascular anatomy (Stewart and Rothwell, 1993). These ultimately reached lengths of up to a metre in the arborescent lycophytes dominating Carboniferous forest communities, but always remained narrow (Kosanke, 1979; Chaloner and Meyer-Berthaud, 1983). The importance of microphylls subsequently declined with the rise of gymnosperms and they are retained only in relict groups within the extant flora (Wikström and Kenrick, 2001). Our analysis of leaf evolution is currently confined to megaphyll leaves, although the physical principles involved could equally be applied to microphylls.

Rapid evolutionary radiation in Early Devonian (Pragian–Emsian) plants apparently led to the widespread appearance of determinate lateral branch systems (Gensel *et al.*, 2001), the necessary precursors to megaphyll leaves (see Figure 11.1; Zimmermann, 1930, 1952). However, interpretation of these ‘proto-leaves’ as lateral appendages may be problematic because the natural orientation of major axes may be difficult to ascertain in some fossils – i.e. whether they were more or less upright, axial systems or an overtopped, determinate, lateral branch. This difficulty is illustrated well by the recent re-interpretation of the Mid-Devonian plant *Hyenia ‘complexa’* Leclercq (Fairon-Demaret and Berry, 2000). Similarly, fossilization by compression may artificially flatten shoot systems, making it difficult to determine whether planation in lateral branches is a taphonomic or a truly developmental phenomenon. Nevertheless, the ‘overtopping’ of dichotomous branch systems (see Figure 11.1) and appearance of determinate laterals does seem to have been an evolutionary innovation within the Early Devonian trimerophytes. The dichotomizing axial form remained common, but was confined to rhyniophytes such as *Horneophyton lignieri* (Kidston and Lang) Barghoorn and Darrah (Figure 11.5; Eggert, 1974).

Morphology is diverse in Early Devonian fossil branched axial systems (Figure 11.5), but all are characterized by repeated dichotomies and a lack of evidence in most cases of pronounced planation (Gensel *et al.*, 2001). Termination may be either in sterile tips or sporangia, but branching is nearly always three-dimensional (Figure 11.5), as in the trimerophytes *Pertica quadrifaria* Kasper and Andrews (1972) and *Psilophyton* Dawson (Gensel and Andrews, 1984). Vascular traces in permineralized proto-leaves of *Psilophyton coniculum* Trant and Gensel (1985) are markedly differentiated from those of the main axis, providing anatomical evidence that they were indeed developed laterally (Gensel, 1984).

Proto-leaves from the Chinese Yunnan (Pragian) fossil assemblages are markedly more differentiated than most Early Devonian examples, with those of *Eophyllophyton bellum*

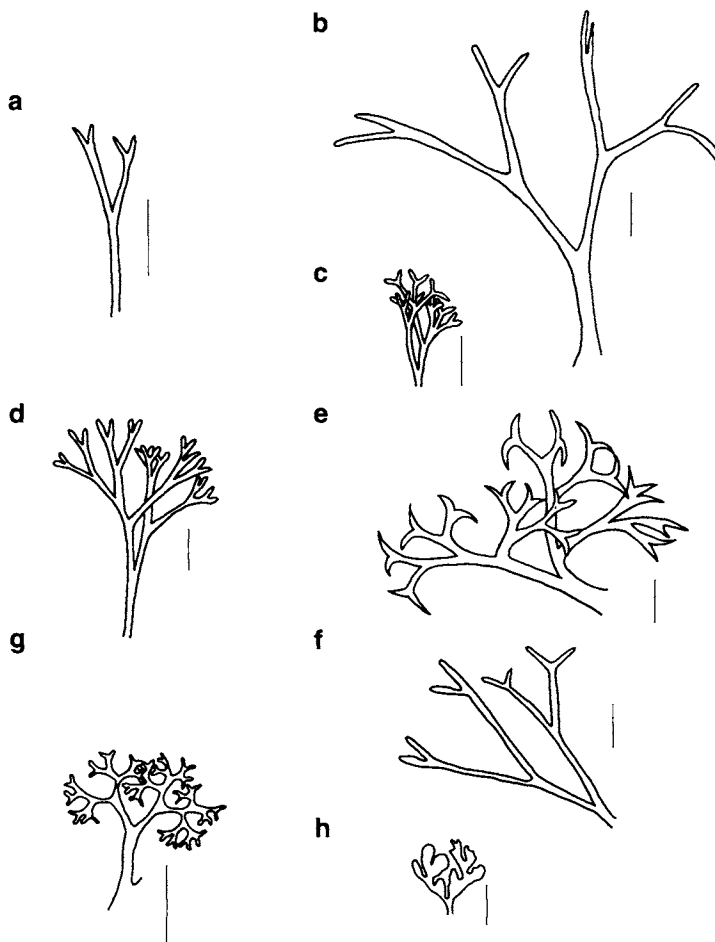


Figure 11.5 Photosynthetic structures of characteristic Early Devonian genera. Asterisks indicate that the genus is confined to this epoch, following Chaloner and Sheerin (1979). Each outline is redrawn from either a: †reconstruction; ‡drawing of a fossil; or ¶photograph of a fossil. Scale bars represent 10 mm for a–g and 1 mm for h. Estimated stages follow the time scale of Harland *et al.* (1990) and are abbreviated as: Pra, Pragian; Ems, Emsian; Eif, Eifelian; Giv, Givetian; Frs, Frasnian; and Fam, Famennian.

- a *Psilophyton dapsile* Andrews *et al.* (1977)[†] (Ems/Eif).
- b *Ps. forbesii* Gensel[†] Andrews *et al.* (1977).
- c *Ps. dawsonii* Banks *et al.* (1975)[†] (Ems/Eif).
- d *Horneophyton lignieri* (Kidston and Lang) Barghoorn and Darrah[†] (Pra/Ems) Eggert (1974).
- e *Pertica quadrifaria* Kaspar and Andrews[†] (Ems/Eif) Andrews *et al.* (1977).
- f *Psilophyton microspinosum* Andrews *et al.* (1977)[†] (Ems/Eif).
- g *Ps. charientos* Gensel (1979)[†] (Ems/Eif).
- h *Eophyllophyton bellum* Hao and Beck (1993)[‡] (Pra).

being both planated and webbed by a thin lamina (thickness = 40–200 μm ; Hao and Beck, 1993). Unlike similar laminar structures in *Adoketophyton subverticillatum* Li and Edwards and *Calatheca beckii* Hao and Gensel from the same flora, which are always confined to terminal fertile regions, those in *E. bellum* are lateral and may be fertile or

sterile (Hao and Gensel, 2001). Thus the proto-leaves of *E. bellum* seem to have no clear role in protecting the sporangia and are interpreted as true photosynthetic megaphyll leaves (Hao, 1988; Hao and Beck, 1993). Their morphology is distinctive, being deeply divided into lobes and reach only 2.0 to 4.5 mm in length and 1.4 to 4.0 mm in width (see Figure 11.5; Hao and Beck, 1993). While their interpretation as the earliest photosynthetic megaphyll leaves may be contentious, these structures undoubtedly demonstrate that there were no developmental barriers to megaphyll leaf evolution in at least one lineage by the Early Devonian.

A transition towards later forest floras began in the Mid-Devonian (Eifelian-Givetian), with large increases in plant stature of up to 3–5 m permitted by the evolution of secondary xylem or cortex tissues (Chaloner and Sheerin, 1979; Niklas, 1997; Berry and Fairon-Demaret, 2001). Laminate megaphyll leaves also appear at this time in several rare genera of uncertain affinity (Høeg, 1967), and may be present in the early progymnosperm *Svalbardia polymorpha* Høeg, although the finely dissected ultimate appendages of this species have been variously interpreted as branches (Beck, 1970) and laminae (Høeg, 1942). However, species with non-laminate, determinate, lateral branch systems (proto-leaves) remain the norm in fossil assemblages (Figure 11.6), and naked, dichotomizing rhyniophyte axes are still present, but restricted in distribution (Berry and Fairon-Demaret, 2001).

Non-laminate proto-leaves of Mid-Devonian age are more commonly planated than are their Early Devonian counterparts (see Figures 11.1 and 11.6), as illustrated by the cladoxylalean *Cladoxylon scoparium* Kräusel and Weyland (Leclercq, 1970) and the putative sphenophyte *Ibyka amphikoma* Skog and Banks (1973). Nevertheless, some species retain three-dimensional branching structures, for example the progymnosperms *Tetraxylopteris schmidtii* Beck (1957) and *Actinoxylon banksii* Matten (1968) and the cladoxylalean *Pseudosprochnus nodosus* Leclercq and Banks (Berry and Fairon-Demaret, 1997). While the terminal segments of these fossils may appear laminate, this is as likely to be taphonomic as developmental, the result of compression of an initially terete structure during preservation (Chaloner, 1999).

Large laminate megaphylls are atypical of Mid-Devonian fossil floras and, in most cases, are species or genera known only from a single locality or specimen. The morphogenus *Platyphyllum* (Dawson) White is represented by five Mid-Devonian species, all characterized by fan-shaped (flabelliform) laminae between 2 and 15 cm in length and 1 and 5 cm in width (Høeg, 1967). Apparent evidence of tracheids in at least three of these specimens confirms their status as tracheophytes (Høeg, 1967), but details are lacking on the plants that bore them. *Ginkgophytopsis gilkinetii* (Leclercq, 1928) Høeg is similar in morphology and similarly enigmatic. However, the largest leaves of late Mid-Devonian (or earliest Late Devonian) age belong to the flabelliform *Enigmophyton superbum* Høeg (1942) and measure at least 16 cm by 12 cm. They demonstrate that, although rare, large laminate megaphylls were clearly a viable adaptation in some environments at this time. It would be of great interest to know more of the ecology of these rare Mid-Devonian flabelliform leaves.

True laminate megaphyll leaves finally came of age in the forest floras of the Late Devonian (Frasnian-Famennian) dominated by the progymnosperm genus *Archaeopteris* Dawson (Beck, 1964). Fossil megaphylls of these trees vary considerably in size and morphology, ranging from the laminate leaves of *A. hibernica* (Forbes) Dawson to the finely dissected branching structures of *A. fissilis* Schmalhausen (Figure 11.7). Thus, although laminate leaves were widespread, plants bearing non-laminate proto-leaves remained key components of forest floras (Gensel and Andrews, 1984). The planated, finely branching,

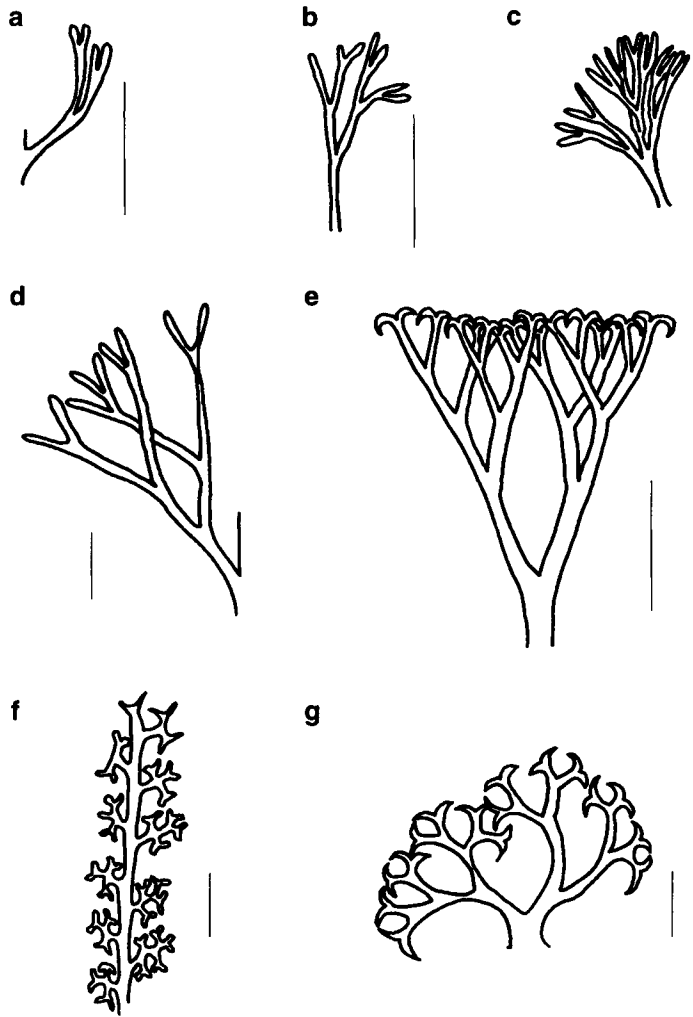


Figure 11.6 Photosynthetic structures of characteristic Middle Devonian genera. See Figure 11.5 for key. Scale bars represent 10 mm for all except a, where the scale is 5 mm.

- a *Calamophyton bicephalum* Leclercq and Andrews (1960)^{*†} (Giv).
- b *Hyenia elegans* Kräusel and Weyland^{*‡} (Eif) Hirmer (1927).
- c *Pseudosprochnus nodosus* Leclercq and Banks (1962)[†] (Giv).
- d *Actinoxylon banksii* Matten (1968)^{*†} (Giv).
- e *Ibyka amphikoma* Skog and Banks (1973)[†] (Giv).
- f *Arctophyton gracile* Schweitzer (1968)[†] (Eif).
- g *Protocephalopteris praecox* (Høeg) Ananiev^{*†} (Eif/Giv) Schweitzer (1968).

non-laminate 'fronds' of the putative primitive fern *Rhacophyton ceratangium* Andrews and Phillips are particularly noteworthy in this context (see Figure 11.7), many reaching lengths in excess of 30 cm (Andrews and Phillips, 1968). They demonstrate that laminate megaphylls, although obviously successful, were by no means a ubiquitous adaptation in Late Devonian plants.

Fossil floras of the Early Carboniferous (Tournaisian-Viséan) are dominated by arborescent lycophytes with long microphyll leaves, and megaphylls are confined to the

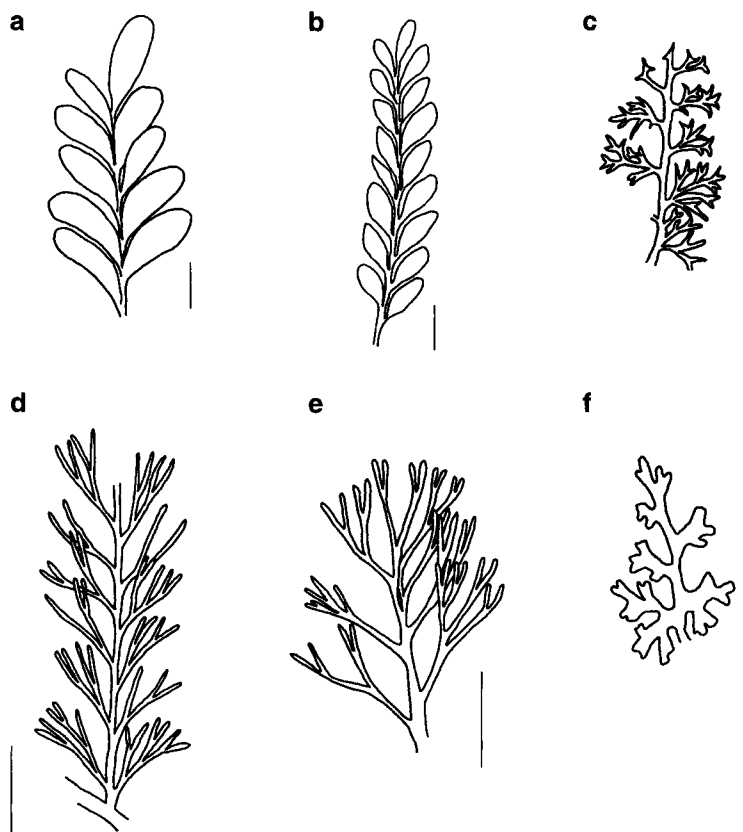


Figure 11.7 Photosynthetic structures of characteristic Late Devonian genera. See Figure 11.5 for key. Scale bars represent 10 mm.

- a *Archaeopteris hibernica* (Forbes) Dawson*[¶] (Fam) Original.
- b *A. halliana* (Goeppert) Lesquereux*[¶] (Fam) Hirmer (1927).
- c *Rhacophyton ceratangium* Andrews and Phillips[¶] (Fam) Cornet *et al.* (1976).
- d *Archaeopteris fissilis* Schmalhausen*[†] (Stage uncertain) Andrews *et al.* (1965).
- e *Svalbardia banksii* Matten (1981)[†] (Frs).
- f *Sphenopteridium rigidum* (Ludwig) Potonié[¶] (Fam) Cleal and Thomas (1995).

ferns, pteridosperms and sphenophytes. In early sphenophytes such as *Archaeocalamites radiatus* (Brongniart) Stur and *Sphenophyllum tenerrimum* Ettingshausen, these remain as lateral branching systems without laminae (Figure 11.8), although laminate forms appear in later genera of this group (Stewart and Rothwell, 1993). By contrast, megaphylls of the pteridosperms *Diplopteridium teilianum* (Kidston) Walton and *Rhacopteris circularis* Walton, and likely members of the same group *Charbeckia macrophylla* Knaus *et al.* (2000) and *Genselia compacta* Knaus and Gillespie (2001), are exclusively laminate, closely resembling those of modern ferns (Figure 11.8). However, Lower Carboniferous leaves display a wide diversity in form (Boyce and Knoll, 2002), with the broad laminae of pteridosperms such as *C. macrophylla* coexisting with the highly dissected terete forms of examples such as *Diplopteridium holdenii*. With the decline of arborescent lycophytes in the Late Carboniferous and rise of gymnosperms, the laminate megaphyll leaf became firmly established in subsequent floras, as it is today (Beck, 1970).

Quantifying the trends in early megaphyll leaf evolution

Qualitative trends in megaphyll evolution support the two key expectations arising from our model, with small, dissected, non-laminate lateral branches being followed by larger laminate megaphylls during the course of the Devonian. Here we quantify these trends using simple observations on a representative selection of Late Palaeozoic photosynthetic structures. For each, we have measured the maximum width of the ultimate segment of branches or laminae, the key determinant of laminar boundary layer thickness for both terete and laminate structures (Jones, 1992). The results presented here are the preliminary findings of a more comprehensive study currently underway.

Figures 11.5 to 11.8 illustrate examples of the ultimate vegetative appendages of Devonian and Early Carboniferous genera. They are chosen quite subjectively to represent what may fairly be regarded as the characteristic range of leaf form (or branched axial systems) shown by plants of each age. We have indicated in the list of genera (Figures 11.5–11.8) those regarded as characteristic of a particular epoch (Chaloner and Sheerin, 1979). We include only forms which are megaphylls with an evident branched vein system, or what appear to be determinate lateral branch systems of the kind which are widely regarded as the antecedents to such megaphylls. We deliberately exclude lycopsid microphylls since our interest is in the origin of the megaphyll leaf.

In interpreting the photosynthetic physiology of these fossils, one weakness is our uncertainty about the extent to which some of the individual small-diameter branch divisions are laminate, since the narrow segment of a laminate structure will develop

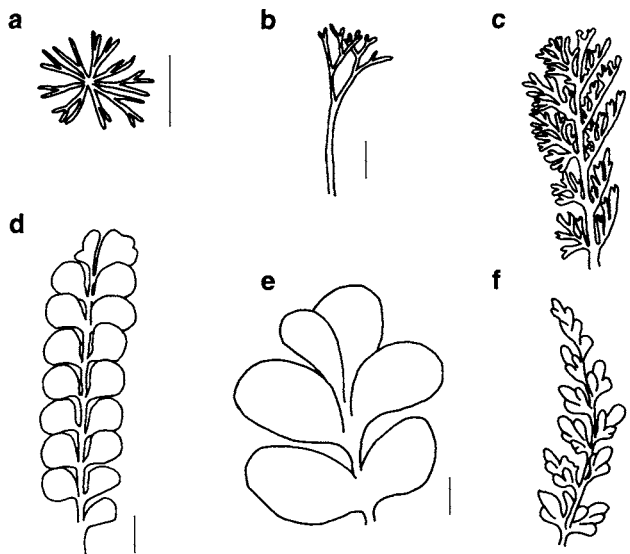


Figure 11.8 Photosynthetic structures of characteristic Early Carboniferous genera. See Figure 11.5 for key. Vis, Viséan; Tou, Tournaisian, other epoch abbreviations as Figure 11.2. Scale bars represent 10 mm.

- a *Sphenophyllum tenerrimum* Ettinghausen[‡] (Vis) Hirmer (1927).
- b *Archaeocalamites radiatus* (Brongniart) Stur[‡] (Tou through Vis) Hirmer (1927).
- c *Diplopteridium teilianum* (Kidston) Walton[¶] (Vis) Andrews *et al.* (1970).
- d *Rhacopteris circularis* Walton[¶] Andrews *et al.* (1970).
- e *Charbeckia macrophylla* Knaus *et al.* (2000)[‡] (Tou).
- f *Genselia compacta* Knaus and Gillespie (2001)[‡] (Tou).

different boundary layer conditions to a terete stem (Jones, 1992). As they are almost exclusively compression fossils (Chaloner, 1999), we may be either viewing a finely divided lamina, or an extensively branched terete lateral branch system. For most of the Early Devonian plants, the circumstantial evidence is that these are terete, branched axial systems; for the Mid-Devonian, some of the finely branched laterals may indeed be laminate (i.e. each final unit being broader than deep in cross-section). We accept that we probably cannot resolve this uncertainty for all the material figured.

The morphological nature of the units illustrated and analysed is also uncertain, but this does not affect the parameter (ultimate segment width) we have recorded. In some cases we are simply making measurements on part of a branched axial system, either the whole subaerial plant (Figure 11.5d) or simply part of it (Figure 11.5a–c) or what appears to be a determinate lateral branch system (Figure 11.5h). In that the pattern is generally repeated within each axial system as a whole, the parameter we are measuring will be the same, whether we base it on a representative sample or the entire system.

A further problem concerns the three-dimensional nature of the branching of some of the Early and Mid-Devonian plants; in many such cases part of a branch system may be hidden within the rock matrix, as a single fracture plane in a rock reveals only two dimensions. Palaeobotanists investigating such a fossil will normally excavate into the matrix to reveal the three-dimensional character of the plant ('degagement' – see Fairon-Demaret *et al.*, 1999). The whole three-dimensional form of the plant may then be presented as a reconstruction by the author, and where available, we have used such a reconstruction in making our figures of characteristic forms (Figures 11.5–11.8). Of course, we may have part of a branch system missing in such a reconstruction, but again, as the part seen is presumably a representative sample of the whole, this will not affect the value of the ultimate branch width.

Maximum widths of the ultimate segment of branches or laminae were measured to the nearest millimetre in the examples from Figures 11.5–11.8, and displayed in Figure 11.9 after a correction for scale. These observations provide preliminary quantitative support

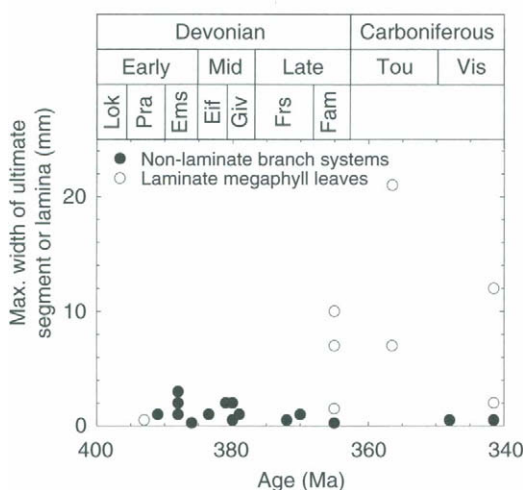


Figure 11.9 Maximum width of the terminal segment of branches or leaves in genera characterizing floras of the Devonian and Early Carboniferous. All are illustrated in Figures 11.5–11.8 and plotted against age, estimated as the Stage mid-point after Harland *et al.* (1990). The symbols distinguish non-laminate axial branching structures from laminate megaphyll leaves.

for the qualitative trends already described. Branched, non-laminate photosynthetic organs of the selected species were less than 3 mm in width throughout the Early and Mid-Devonian. *Eophyllophyton bellum* is the only example of a species with laminate megaphyll leaves during this interval and also conforms to the pattern, with leaf lobes less than 1 mm in width. Only late in the Late Devonian (Famennian) did photosynthetic organs increase in size, with the widespread appearance in progymnosperms of laminate megaphyll leaves up to 10 mm in width. This upward trend in maximum leaf size continued in the Carboniferous, as illustrated by the pteridosperms in Figure 11.8, exceeding 20 mm in *Charbeckia macrophylla*. Narrow, non-laminate 'proto-leaves' were still present in sphenophyte floras at this time, but were rare by comparison with their heyday in the late Early and Mid-Devonian.

Discussion

The quantitative analysis presented in Figure 11.9 must be regarded as a preliminary investigation of the trends occurring in megaphyll evolution during the Late Palaeozoic, rather than a rigorous objective test of our hypothesis. Nevertheless, it supports the two key predictions arising from our model simulations. First, that biophysical constraints restricted the earliest megaphylls to narrow or highly dissected shapes. Evidence from the fossil record suggests that these constraints were important limiters of megaphyll evolution, confining proto-leaves to (predominantly) non-laminate forms throughout the Early and Mid-Devonian, and delaying the widespread appearance of laminate forms until the Late Devonian (see Figure 11.9).

Secondly, evidence supports our expectation that the evolution of larger and less dissected laminate megaphylls tracked the Devonian and Early Carboniferous decline in atmospheric CO₂. Our examples show a clear trend in maximum lamina width and suggest that this evolutionary change was particularly rapid in the 20 Ma spanning the latest Devonian and earliest Carboniferous (see Figure 11.9). This period must therefore be regarded as an important focus for future studies. The precise starting date for these increases in lamina size is uncertain, because of the combination of examples selected for study and the small sample size. Future examination of Givetian-Frasnian fossils will be important to date this evolutionary event with greater accuracy (see Figure 11.9).

For the reasons set out earlier (Beerling *et al.*, 2001a) and amplified here, we therefore interpret the late appearance of laminate foliage as a response to changes in atmospheric composition, largely brought about by terrestrial plant life itself (Algeo and Scheckler, 1998; Algeo *et al.*, 2001). However, we accept that this simple interpretation must be qualified with some provisos.

The first of these is the presence from the Early Devonian onwards of very few records of flabelliform leaves, seemingly produced by vascular plants. Unfortunately we know next to nothing of the ecology of the plants producing them, or indeed the ecology of most of the plants that we use here to illustrate this theme. The interpretation of these 'anomalous' leaves is complicated by the fact that the vascular nature of the plants themselves is not in every case securely documented. The flabelliform phylloids attributed to *Prototaxites* Dawson (see e.g. Schweitzer, 1987: Figure 11.7), a structure now considered to be fungal (Hueber, 2001), are a reminder of the possible confusion between such structures and the leaves of terrestrial land plants.

A further proviso relates to the fact that, synchronously with the early pinnate megaphylls of the Late Devonian/Early Carboniferous progymnosperms, pteridosperms

and/or pteridophytes, there remains a range of deeply divided leaves or leaf-like lateral systems (e.g. Figure 11.8a–c). This is hardly surprising because modern plants with very different light-trapping and transpiration strategies evidently coexist in the same communities (e.g. Gates, 1979). The presence of rosette weeds such as the deeply dissected *Achillea millefolium* L. in juxtaposition in the same sward as the entire-leaved *Plantago major* L. reminds us that we need to be guarded in offering simplistic explanations attributing ‘adaptive’ merit of one strategy against another. Leaves of small herbaceous plants encounter a wide range of environmental variables through the course of a seasonal climate and equally must ward off a range of different herbivores. While one plant may use chemical means of achieving this, others may use structural modifications, so producing very different leaf forms for different reasons. It is evident that the appearance of megaphyll leaves did not ‘displace’ their more deeply dissected antecedents, but while expanding in diversity in different plant groups, came to coexist with them.

As we see in other aspects of evolutionary change in plants, an innovation may partially displace the version that had preceded it, but only rarely supplants it entirely. In a simple way, this might be argued of the development of heterospory from homospority and subsequently of the seed habit from heterospory. Those three types of life cycle are still well represented in the present-day flora, although clearly the representation of each has changed drastically since Devonian time.

Further expectations from our biophysical simulations relate to the ecology and spatial distribution of early megaphyll evolution. First, we might expect laminate leaves to have appeared first in species of shaded forest understorey habitats, where the adverse effects of high solar irradiance are reduced, but not eliminated (Woodward, 1980). Unfortunately, the Devonian plant fossil record does not permit a test of this expectation, since sun leaves cannot be distinguished from the shade leaves of early forests. Secondly, we might expect to see significant latitudinal trends in the early evolution of leaves. Since solar elevation declines with latitude, the interception of solar energy by a planated lamina must also decrease (see Figure 11.4A), reducing the risk of overheating and the necessity for effective heat dissipation. The laminate megaphyll leaf is therefore likely to have become widespread first at high latitudes, an expectation that may be tested using palaeolatitude data for fossil localities (Scotese and McKerrow, 1990). Furthermore, the temporal trend in leaf width seen in Figure 11.9 is likely to be paralleled by a spatial pattern, with megaphyll laminae becoming larger and less dissected with latitude. Such expectations offer another key test of our simulations, requiring the assembly of a larger set of observations.

Acknowledgements

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12

Stomatal function and physiology

Tracy Lawson and James I L Morison

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Introduction

Stomata are small adjustable pores found in large numbers on the surface of most aerial parts of higher plants (Spermatophyta), and also in the Pteridophyta. They have been recorded in the fossil record from as early as 411 million years (Ma) ago in the late Silurian-early Devonian period (Edwards *et al.*, 1992). During this period plants colonized terrestrial environments which led to the evolution of thin cuticles, stomata and vascular systems in order to avoid desiccation and to transport water while still allowing CO₂ exchange (Chaloner, 1970). Stomata are formed from two specialized cells in the epidermis, known as guard cells and, in many species, there are also other adjacent subsidiary cells, that are

morphologically distinguishable from the general epidermal cell (see Colour Plate 1). The central role of stomata is in regulating gas exchange between the inside of the leaf and the external environment (Cowan and Troughton, 1971; Jones, 1992) because the cuticle is almost impermeable to water vapour and CO₂. The plant needs sufficient CO₂ to enter the leaf to photosynthesize, while conserving water to avoid tissue dehydration and metabolic disruption. Even though the stomatal pores as a whole occupy only 0.5–5% of the leaf surface when fully open, almost all the water transpired as well as the CO₂ absorbed for photosynthesis passes through these pores, so stomatal function has huge importance in the global hydrological and carbon cycles.

Stomatal aperture is regulated by both internal physiological and external environmental factors. Pore opening through guard cell movements is stimulated by illumination with light in the photosynthetically effective waveband (particularly the blue waveband), low CO₂ concentrations and high humidity, while closure is promoted by darkness, low humidity, high temperature and high CO₂ concentrations (see reviews by Assmann, 1993 and Willmer and Fricker, 1996). Such guard cell movements are brought about through changes in guard cell turgor (Heath, 1938) and through changes in the difference between guard and epidermal cell turgor (Weyers and Meidner, 1990). These turgor changes require the loss or accumulation of K⁺ or other cations and the parallel exchange of anions including organic solutes such as malate and sucrose (e.g. Willmer and Fricker, 1996; Outlaw, 1996; Asai *et al.*, 2000).

The numbers of stomata per unit leaf area (referred to as stomatal density or frequency) vary with species and conditions and range from 0 to 2000 or more stomata mm⁻² (Willmer and Fricker, 1996). In herbaceous plants, stomata are found on both the upper (usually adaxial) and lower (usually abaxial) surfaces of leaves and are termed amphistomatous, although there are usually more stomata on the lower surface. However, many tree species have stomata only on the lower surface (hypostomatous) and aquatic plants with floating leaves, such as water lilies have stomata only on the upper surface (epi- or hyperstomatous).

Interest in stomata and their evolution, function, anatomy and physiology have been the subjects of intense studies over the last century. In this chapter, we will concentrate on their function and physiology, in order to clarify the role stomata play in determining carbon assimilation. First, we look at the role of stomata in controlling gas exchange and the limitation stomata can impose on leaf CO₂ uptake. We then describe the effects of three key environmental factors on stomatal movements and the consequences for photosynthesis. In the last section we will briefly discuss some new approaches for studying stomatal function and physiology. Our approach is illustrative, rather than an extensive review as there are many detailed texts and reviews available (e.g. Assmann, 1993; Willmer and Fricker, 1996; Zeiger, 2000).

Stomatal control of leaf gas exchange

The diffusion rate of gases into or out of the leaf, or any other plant part, depends on the concentration gradient and the diffusive resistance of the pathway. For water loss from the mesophyll cells inside the leaf, the major pathway is therefore from the mesophyll cell walls through the sub-stomatal cavity to the pore and then out through the layer of air immediately surrounding the leaf, to the mixed air stream (Figure 12.1). The pathway for the uptake of CO₂ by the mesophyll during photosynthesis is essentially the same, in the converse direction, but with an additional component of diffusion through the mesophyll cell into the chloroplast. The resistance of the stomatal pathway depends on the geometry

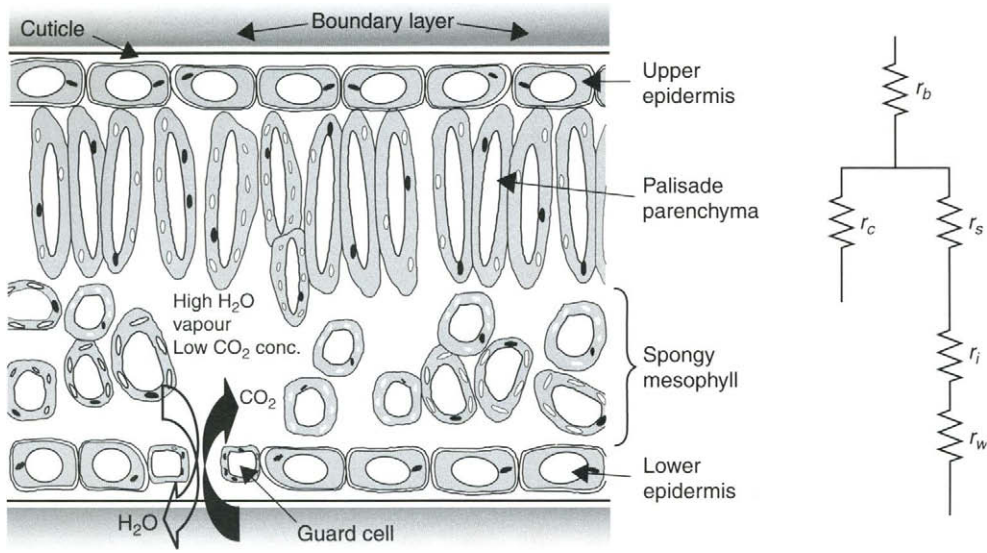


Figure 12.1 Diagrammatic cross-section of leaf showing pathway for diffusion for CO₂ and H₂O into and out of a leaf, respectively. The resistance network analogy is also shown for one surface combining the boundary layer (r_b), cuticular (r_c), stomatal (r_s) and intercellular space (r_i) resistances (see Nobel, 1991: 143, for details and typical values of the resistances).

of the pores as well as their frequency. Note that although the pore area when open may only be at maximum a few percent of the total leaf area, the rates of evaporation can be about half that of a wet surface of similar dimensions; this is due to the ‘edge effect’ of diffusion through multiple pores (see Willmer and Fricker, 1996: 116). The stomatal resistance, r_s , can be calculated from pore dimensions of elliptical pores as:

$$r_s = \frac{(d + 2c)}{D_w \cdot A_s \cdot SF} \tag{1}$$

where: D_w = water diffusivity in air (mm^2s^{-1}); A_s = mean pore area (mm^2), d = pore depth (mm), c = an ‘end correction’ for the edge effect (mm) and SF = stomatal frequency (mm^{-2}). Different end corrections are necessary for different shaped pores (for further details see Weyers and Meidner, 1990: 57). The units of r_s are therefore in time taken to diffuse unit distance (s mm^{-1}). However, it is common to use the reciprocal of resistance, termed a ‘conductance’, g_s , and to express fluxes as a molar density, giving conductance in $\text{mmol m}^{-2}\text{s}^{-1}$. To convert r_s in s mm^{-1} to g_s in $\text{mmol m}^{-2}\text{s}^{-1}$, the $1/r_s$ is multiplied by (P/RT) where P = atmospheric pressure (Pa), R = gas constant ($\text{J mol}^{-1}\text{K}^{-1}$), and T = mean of leaf and air temperature (K, see Jones, 1992: 56). Equation (1) can be used to examine the sensitivity of r_s or g_s to changes in the component parameters (Weyers and Lawson, 1997). Using a plausible range of values for the stomatal parameters for *Phaseolus vulgaris* L. (Figure 12.2), it is clear that the various stomatal characters influence the calculation of g_s to different extents. The main determinant of g_s is pore aperture (width), and stomatal frequency makes a smaller contribution, although more than pore depth or length.

Equation (1) only describes the stomatal part of the pathway for diffusion of water vapour from inside the leaf into the mixed air stream. First, the ‘boundary layer’ of air close

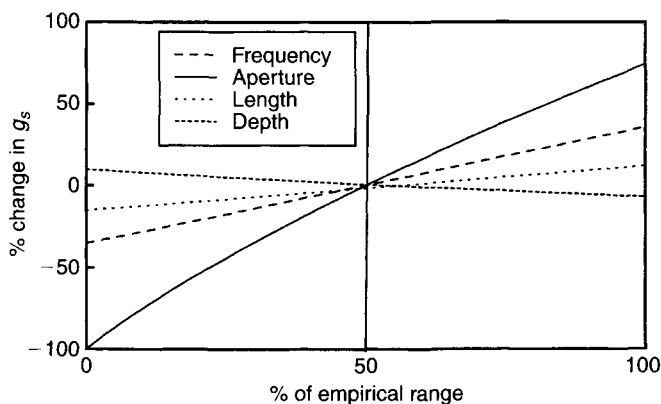


Figure 12.2 Predicted sensitivity of stomatal conductance, g_s , to changes in pore dimension and frequency within empirically derived ranges. Effects of adjusting each anatomical character within its estimated range on g_s were calculated using Equation (1) in the text, keeping the other values constant according to Weyers and Lawson (1997). The analysis follows typical ranges of values derived from observation of *Phaseolus vulgaris*: stomatal aperture, 0–15 μm ; stomatal frequency, 35–65 mm^{-2} ; pore length, 33.3–40 μm ; pore depth, 15–25 μm . Values within each range were used to calculate stomatal conductance, g_s , using Equation (1). The vertical line represents the g_s obtained using the median values for each variable, which was 431 $\text{mmol m}^{-2} \text{s}^{-1}$.

to the leaf (see Figure 12.1) also poses a resistance to diffusion (r_b) and this resistance is not uniform across the leaf, increasing with increasing downwind distance across the leaf (Grace and Wilson, 1976). The magnitude of r_b varies widely depending upon surface characteristics, such as presence of hairs, leaf size and shape and also wind speed and turbulence. For large leaves the boundary layer can be a few millimetres thick even under moderate wind speeds (e.g. Aphalo and Jarvis, 1993). Lobes and serrations reduce the effective downwind leaf dimension and reduce the average r_b compared with a leaf with equal surface area, but smooth margins (Gottschlich and Smith, 1982). Narrow grass leaves and needle-shaped leaves obviously have the lowest r_b values (see Jones, 1992: 65, for graphs showing the relationships between r_b , wind speed and size). Secondly, while the cuticle is relatively impermeable, some water is lost through it (varying with species and conditions, see review by Kerstiens, 1996) giving a ‘cuticular resistance’, r_c , in parallel to and usually much higher than r_s . There is also an internal resistance, r_i , for the pathway from cell wall to pore, but this is normally small compared to r_s and r_b . The rate of diffusion of water from a leaf (E , $\text{mmol m}^{-2} \text{s}^{-1}$) can therefore be calculated (Equation (2)) from the difference in water vapour pressure between the inside and outside of the leaf, ($\text{VPD} = w_i - w_a$, kPa) and the leaf resistance, r_l ($\text{m}^2 \cdot \text{s mol}^{-1}$), or conductance, g_l ($\text{mmol m}^{-2} \text{s}^{-1}$), which is given by the sum of the various resistances in series and/or parallel as appropriate as shown in Figure 12.3, and expressed in Equation (3):

$$E = \frac{w_i - w_a}{r_l \cdot P} \quad \text{or} \quad E = \frac{w_i - w_a}{P} \cdot g_l \quad (2)$$

$$r_l = \frac{r_c(r_s + r_i)}{r_c + r_s + r_i} + r_b \quad (3)$$

Figure 12.3a shows that if the cuticular resistance is very low, then r_l becomes curvilinearly related to r_s and Figure 12.3b shows that r_b only influences E when $r_s < r_b$. Note

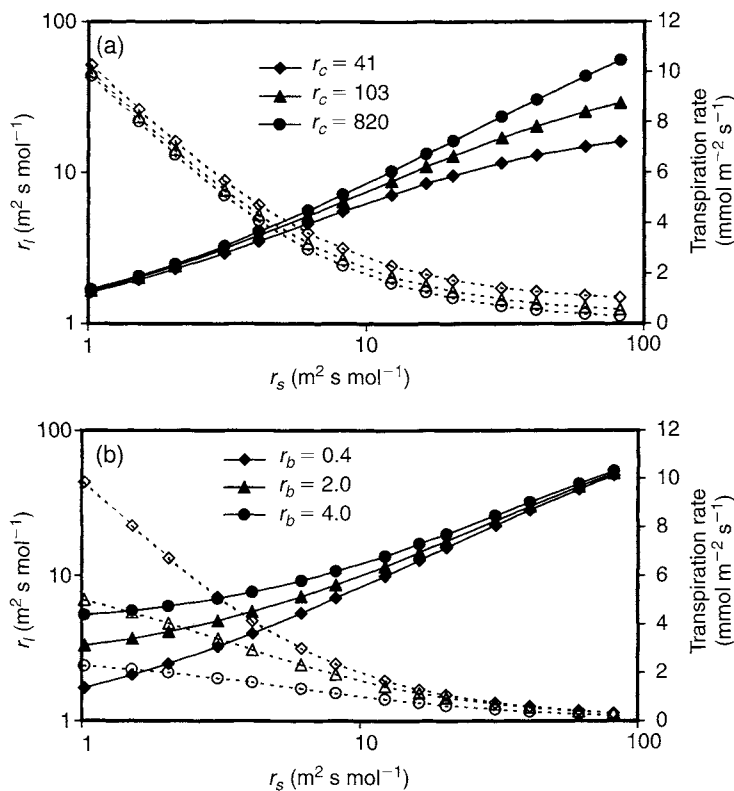


Figure 12.3 Effect of changes of (a) cuticular resistance and (b) boundary layer resistance on leaf resistance (closed symbols) and transpiration rate (open symbols) (calculated from Equations (2) and (3), see text). Calculations in both used a ratio of abaxial to adaxial r_s of 0.3, $r_l = 0.82 \text{ m}^2 \text{ s mol}^{-1}$ and a VPD of 1.0 kPa. In (a) fixed $r_b = 0.41$ and in (b) $r_c = 410 \text{ m}^2 \text{ s mol}^{-1}$.

that equations can be written for leaf net CO_2 assimilation rate (A) that are essentially very similar to Equations (2) and (3), but replace the partial pressure of water vapour by that for CO_2 and take into account that the resistance for CO_2 diffusion in air is slower than that for water vapour (see Jones, 1992: 185).

The above diffusion equations can be used for simple analyses, but in practice the leaf temperature is not independent of the transpiration rate, thereby affecting convective heat transfer, the long-wave radiation balance and the internal water vapour pressure, w_i . In particular, this latter determines the driving gradient for evaporation (Equation (2)). Because of these 'feedbacks' it is necessary to consider a more complete 'energy balance' equation, such as the Penman-Monteith equation (see, for example Jones, 1992: 112) in order to examine the relative control that stomata exert on transpiration, compared with other components. Analyses with these additional aspects show (Figure 12.4) that the important feature is the degree of 'coupling' of the leaf to the air stream (Monteith, 1981); if the leaf has a small r_b compared to r_s then the leaf is 'well-coupled' and leaf temperature will not increase substantially, and changes in r_s will be reflected in E (for other example calculations see Morison and Gifford, 1984). This is typically the case with small, needle-shaped leaves, at the top of the canopy with relatively high wind speeds. The opposite situation occurs with large, broad leaves within short, dense canopies, in still conditions when

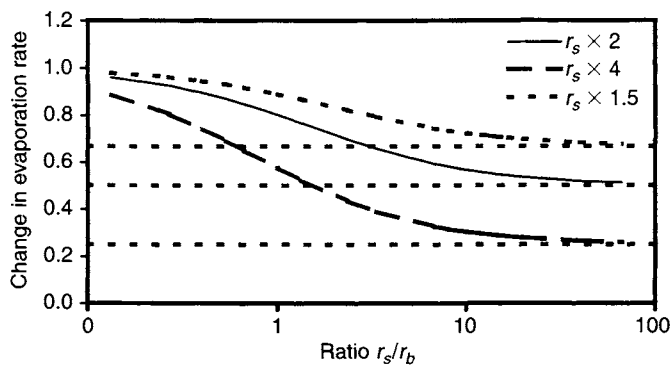


Figure 12.4 Effect of set increases in the stomatal resistance, r_s , on the relative evaporation rate, E , from a leaf. Calculated using energy-balance equations, with air temperature of 20°C, radiative resistance of 5.1 and a boundary layer resistance of 0.37 m² s mol⁻¹. Evaporation rate is shown as the changed rate, relative to original. Dotted lines indicate E change equals r_s change.

E will not closely reflect changes in r_s as r_s/r_b is small (Figure 12.4). Indeed, it has recently been suggested that the evolution of larger planate leaves from the earlier leafless branched shapes became possible during the late Devonian period because increased stomatal frequencies, in response to declining atmospheric CO₂ concentrations, produced greater evaporative cooling and kept leaves below their lethal temperature limit (Beerling *et al.*, 2001; see Chapter 11).

Role of stomata in leaf gas exchange

Stomatal behaviour is obviously important because it directly modifies the CO₂ assimilation rate and transpiration rate and consequently affects plant water and carbon status. In addition, there are also other less obvious indirect effects such as on nutrient status and leaf temperature, caused by stomatal control of transpiration. Here we will concentrate on the direct effects. Clearly, the restriction that stomata (and the other parts of the diffusion pathway) place on CO₂ assimilation rate, A , (i.e. the diffusion limitation) depends upon environmental conditions and plant photosynthetic characteristics. Assimilation rate, A , is not linearly related to leaf conductance, g_l (Figure 12.5), because under all but very restricted CO₂ supply rates, assimilation rate is colimited by other factors (primarily light). When CO₂ supply through the stomata is high, further increase in g_l has little or no effect (in this example, when $g_l > 500$ mmol m⁻² s⁻¹). The limitation that stomata can pose to leaf CO₂ assimilation rate can best be examined through the now traditional 'A/C_i' analysis, where A is measured at a range of external CO₂ concentrations, usually at high light and constant temperature and the intercellular space CO₂ concentration (C_i) is calculated from the diffusion equations (see Farquhar and Sharkey, 1982). Typical A/C_i response curves constructed for *Zea mays* L. (C₄) and *Phaseolus vulgaris* (C₃) show (Figure 12.6) that A is near saturation at low C_i concentrations in maize, but in bean A increases up to C_i of ca. 500 μmol mol⁻¹. The intersections of the dashed vertical line with the A curves indicate the assimilation rates if there was no diffusion limitation by stomata ($g_l = \infty$ so $C_i = C_a$) at present atmospheric CO₂ concentration. The dashed diagonal lines therefore represent 'supply functions' corresponding to lower g_l values and, as g_l is reduced, supply of CO₂ is reduced, so C_i declines, possibly affecting A depending on the

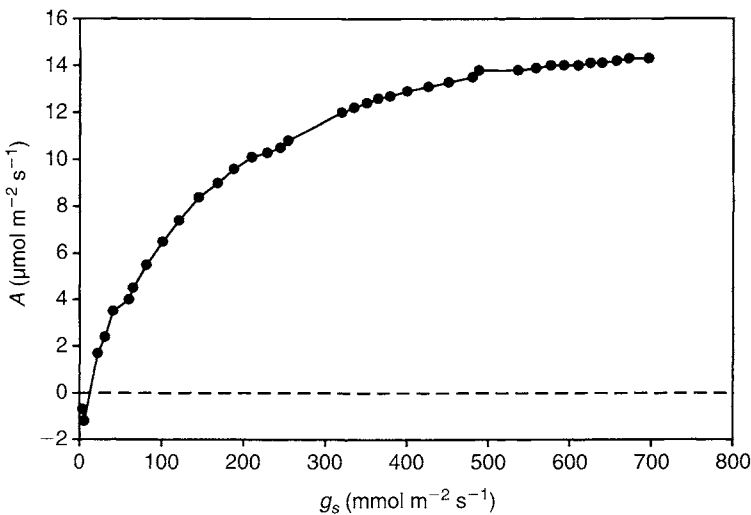


Figure 12.5 The relationship between net CO₂ assimilation rate, *A*, and stomatal conductance, *g_s* for a single *Phaseolus vulgaris* (bean) leaf. The change in *g_s* was caused by illuminating a pre-darkened plant (to ensure stomatal closure), at a saturating PPFD of 1200 μmol m⁻² s⁻¹. Cuvette CO₂ concentration was maintained at 356 μmol mol⁻¹ and VPD of 1.4 kPa and temperature 24°C.

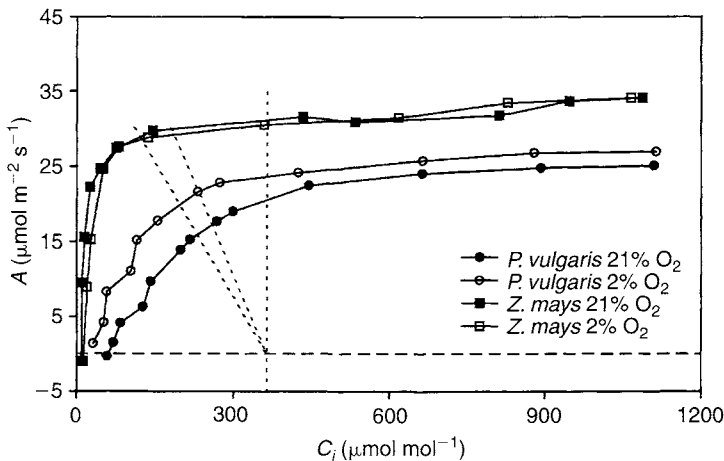


Figure 12.6 Relationship between net CO₂ assimilation rate (*A*) and internal CO₂ concentration (*C_i*) in *Phaseolus vulgaris* (*C₃*) and *Zea mays* (*C₄*) measured in 2 and 21% oxygen concentration. Solid curves represent the CO₂ ‘demand function’ and dashed lines indicated the ‘supply function’ depending upon *g_l* as described by Farquhar and Sharkey (1982).

degree of photosynthetic CO₂ saturation. In the example curves shown (in normal 21% O₂), all values of *g_l* limit *A* significantly in the bean leaf, whereas the maize leaf would still photosynthesize at close to the maximum rate with *g_l* of 150 mmol m⁻² s⁻¹. Another contrast between these *C₃* and *C₄* photosynthetic types is shown by the effect of low O₂ concentration (2.1%), which is also of palaeontological interest. Obviously there is no effect of reduced O₂ on *A* in the *C₄* species but, in the *C₃* species, it causes a 6–25% increase in *A* and reduces the sensitivity to stomatal diffusion limitation markedly. While in Figure 12.6 the major contrast between the two species in photosynthetic sensitivity to

stomatal limitation is due to different photosynthetic pathways, similar but less dramatic differences can be seen between leaves with nitrogen content differences (Wong, 1979), or sun and shade morphologies and biochemistry (Patterson, 1980) and with other stresses, e.g. water (von Caemmerer and Farquhar, 1984). It should be clear from these examples that to understand the control that stomata exert on photosynthesis requires a quantitative description of photosynthesis in the particular conditions being examined. This in turn will help in understanding stomatal responses to changes of environmental conditions in the geological past.

Heterogeneity in stomatal characters

It is probable that when they measure g_l or r_l on a whole leaf in a cuvette, many researchers imagine that it is a simple bulk version of that shown in the stylized diffusion pathway diagram of Figure 12.1. However, we should always remember that it is a bulk measurement across a very large and heterogeneous population of stomata and mesophyll. As mentioned in the introduction, there are likely to be differences between the leaf surfaces, as well as other important sources of variation. Heterogeneity in both space and in time of stomatal anatomical characteristics (or related variables such as g_s) is found at many scales, from the size, frequency and behaviour of small groups of guard cells to the gas exchange of whole plants or stands of plants, even in apparently homogeneous environments (Tichá, 1982; Solárová and Pospíšilová, 1983; Pospíšilová and Šantrucek, 1994; Weyers and Lawson, 1997). For example, Colour Plate 1 shows two guard cell complexes adjacent to one other, yet one pore is open and the other is closed, illustrating variation in stomatal behaviour even at this small scale. Heterogeneity in stomatal characters is not confined to anatomical features but has also been observed in function. As an example, Figure 12.7 shows substantial variation in g_s , A and C_i over the adaxial surface of a *Phaseolus* leaf interpolated from approximately 30 spot readings on 1.25 cm² areas (see Lawson and Weyers, 1999 for other examples). It is also obvious that g_l and A are not consistently well correlated in different areas, although a general pattern exists. Comparison with the map of C_i emphasizes that areas of high g_s may have high C_i and low A rates, indicating that g_s and C_i are not limiting the rate of A . Other common examples of stomatal variation can be found between leaves on plants, due probably to age and ontogenetic effects. Figure 12.8 shows that in grape vine shoots g_s values increased with leaf number from the base (and therefore also with decreasing age) up to leaf 10, but newer leaves further along the shoot had progressively lower g_s . This was not simply related to illumination as all apart from the basal few leaves were well lit, nor was it related to leaf size, but was probably largely due to leaf age effects, with a peak g_s in mature but not old leaves (see also Solárová and Pospíšilová, 1983).

It is important to recognize the extent of spatial and temporal variability in stomatal and photosynthetic parameters because experimental treatment effects are often examined with 'snapshot' measurements of gas exchange with small chambers that sample only small proportions of the total area of a leaf, at least on broadleaved species. The existence of spatial and temporal heterogeneity in stomatal behaviour necessitates that we ensure that: (1) measurements are taken from a similar area in each sample leaf; (2) conditions around the leaf under examination are kept constant; and (3) a large sampling area is employed. Furthermore, we should remember that leaves on a plant are not independent particularly because of hydraulic connections and therefore ideally conditions around the whole plant

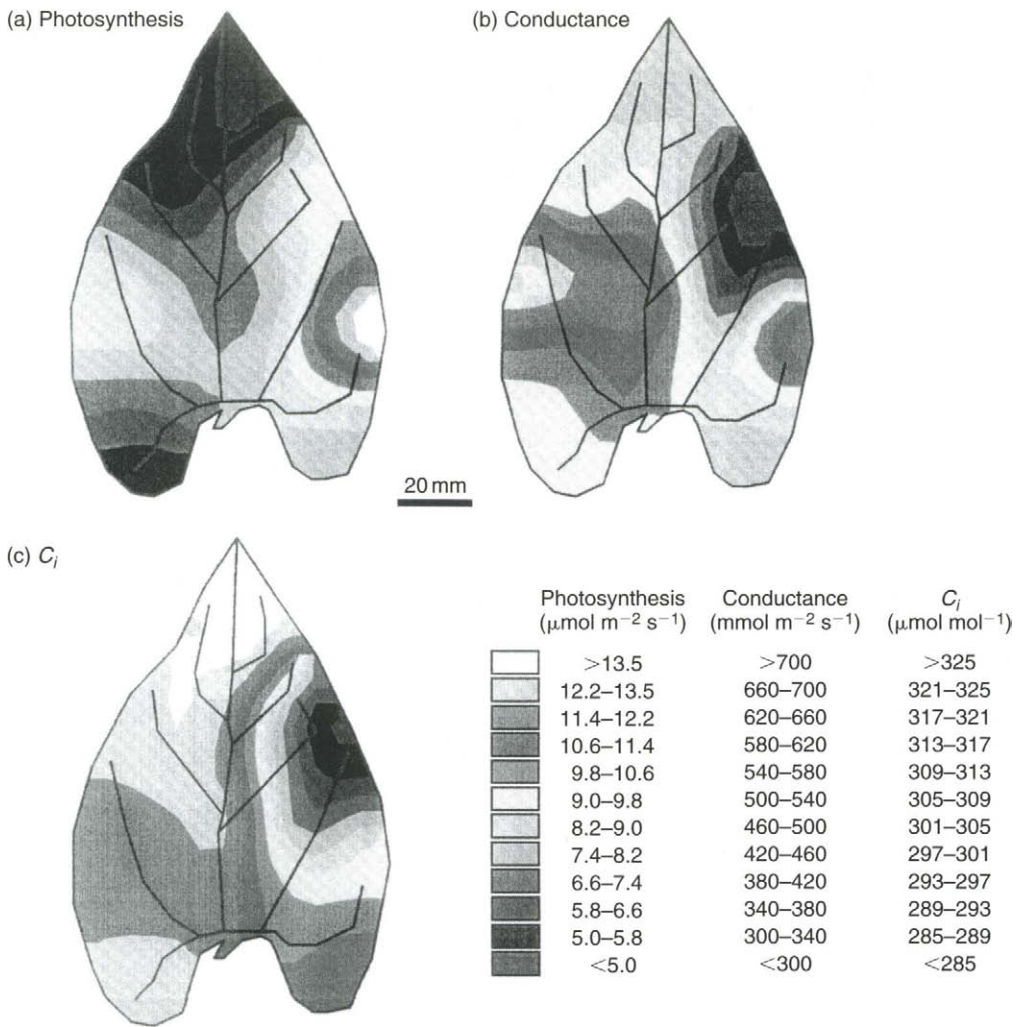


Figure 12.7 Contour maps showing spatial variation in gas exchange in *Phaseolus vulgaris*: (a) net CO₂ assimilation rate, (b) leaf conductance and (c) estimated internal CO₂. Data were collected from 24 sites using a 125 mm² cuvette. Cuvette conditions were set at an external CO₂ concentration of 347 ± 3 μmol mol⁻¹ and relative humidity of 47.2 ± 4%.

should be controlled. We need also to bear in mind that treatments or stresses may change the variability, which may be interesting in itself (e.g. Weyers *et al.*, 1997).

Effect of environmental variables on stomata and photosynthesis

Part of the fascination with stomata is that apertures respond to many different environmental stimuli because the guard cells sense several stimuli, most notably CO₂, humidity, light and temperature. While many studies have tried to isolate individual factors for study, it is important to realize that naturally it is often the combination of environmental stimuli that affect stomatal behaviour. This is as important to studies on palaeoecophysiology as to those on modern plants. For example, a change in light intensity may

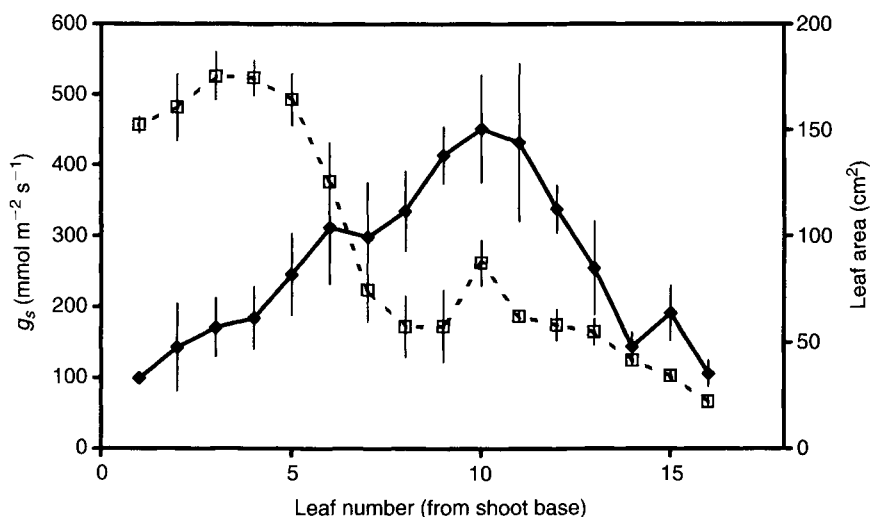


Figure 12.8 Stomatal conductance (solid symbols) and area (open symbols) of leaves along well-illuminated shoots of grape vine. Measurements taken on 27th June 1991, in a vineyard in central Spain, in clear sky conditions, around midday (PPFD > $1700 \mu\text{mol m}^{-2} \text{s}^{-1}$, air temperature = 35°C , VPD = 3.2 kPa) measured with a diffusion porometer. Average of four similar shoots, bars are standard error of the mean.

simultaneously change photosynthetic rate and therefore C_i , and change leaf temperature, which will modify transpiration and the leaf water status. In addition, the majority of work has examined steady-state responses and has not looked at dynamic changes, which are probably critical in natural environments. Below we show the effect of CO_2 , humidity and light and discuss both the direct and indirect effects and also illustrate the effect of interactions between a couple of environmental parameters on stomatal and photosynthetic responses.

Stomatal response to CO_2

The stomatal response to CO_2 has gained great attention over the last couple of decades due to concerns about the effect of rising atmospheric CO_2 concentrations ($[\text{CO}_2]$) caused by industrialization and land use change (Mansfield *et al.*, 1990). However, an explanation of how stomata respond to CO_2 has been a central question in stomatal physiology since the earliest observations of Linsbauer (1916) and Freudenberg (1940). It is now clear that there are at least two different parts to the question of how stomata respond to increased ambient $[\text{CO}_2]$. The first part is a medium- to long-term morphological and developmental response where the numbers and frequency of stomata may change after growth of the plant in high or low ambient CO_2 partial pressure (see review by Woodward and Kelly, 1995). Stomatal characters have now been used to infer past climatic conditions (McElwain, 1998) which are then used in the evaluation of global palaeoclimatic models (e.g. Beerling *et al.*, 1998). Therefore, it is important to understand more about the mechanisms determining stomatal patterns and the variation that exists between taxa and conditions. This stomatal patterning response to $[\text{CO}_2]$ has two particularly interesting features. First, in both experimental work and analysis of herbarium specimens the largest increase in stomatal frequency is usually at concentrations *lower*

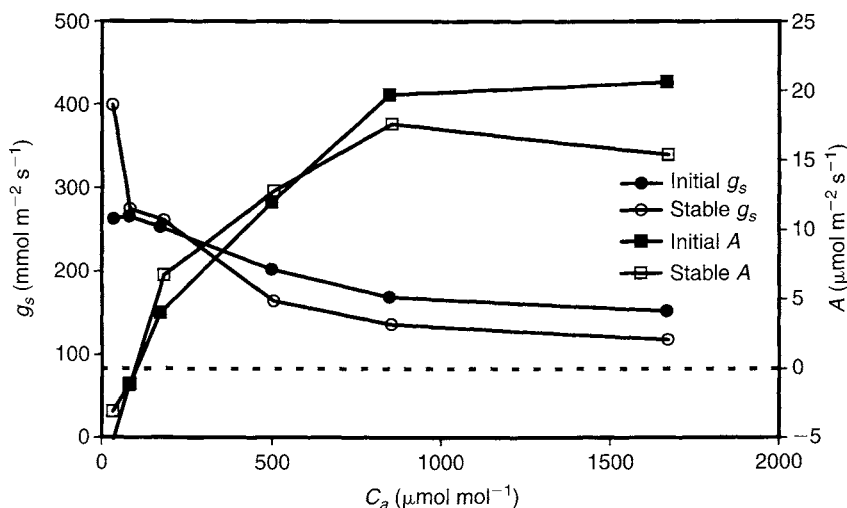


Figure 12.9 The response of stomatal conductance and net CO_2 assimilation rate to external CO_2 concentration, C_a , in *Phaseolus vulgaris*. Cuvette conditions were maintained at 1.24 kPa VPD at a temperature of 25°C and a PPFD of 1200 $\mu\text{mol m}^{-2} \text{s}^{-1}$.

than present atmospheric (Woodward, 1988). Secondly, it has been shown in one experimental system with *Arabidopsis*, that the signal for the altered patterning is not generated by the $[\text{CO}_2]$ around the developing leaf, but is a transmissible signal generated in older leaves exposed to different concentrations (Lake *et al.*, 2001). The second part of the stomatal response to $[\text{CO}_2]$ increase is a short-term, reversible, physiological response where aperture reduces with an increase in ambient $[\text{CO}_2]$ in most situations and, in most species examined, with a typical reduction of g_s of about 40% with a twofold increase in ambient $[\text{CO}_2]$ (see Morison, 1987, 1998). However, there is some notable variation in the response, from high to zero sensitivity, dependent on environmental conditions, pre-conditioning and, apparently, with species, although it has to be noted that there are very few detailed side-by-side comparisons of species sensitivity to CO_2 . In addition, it should be appreciated that the response of g_s is commonly not linear, showing usually a reduced sensitivity at higher than present atmospheric concentrations (Morison, 1987, 2001). Whether this is simply because that with reduced apertures, there is less scope for further movements, or whether it is an important consequence of the underlying physiological mechanism is not yet clear. Another point of concern is the general assumption that stomata respond in a similar manner today as they did many millions of years ago (e.g. using 'nearest living equivalent' material, e.g. McElwain, 1998). However, there is some evidence that long-term growth in high $[\text{CO}_2]$ can cause an acclimation (a 'physiological change in response', Drake *et al.*, 1997) in the stomatal response to $[\text{CO}_2]$ with changed stomatal sensitivity (Morison, 1998; Assmann, 1999; Lodge *et al.*, 2001).

An example of the effect of ambient $[\text{CO}_2]$ on both g_s and A in a leaf of the C_3 species *Commelina communis* L. is shown in Figure 12.9. Initial measurements were taken only a few minutes after changing $[\text{CO}_2]$, and A showed the typical increase at higher C_a which levelled off at ca. 1000 $\mu\text{mol mol}^{-1} C_a$, whereas g_s showed a decline with increasing C_a , particularly between 50 and 500 $\mu\text{mol mol}^{-1} C_a$. Measurements taken after the leaf had stabilized (20–30 minutes) are similar, but g_s increased at lower C_a and decreased at

higher C_a . This slower change in g_s than in A resulted in small but significant modifications in A , emphasizing the effect of stomatal behaviour on CO_2 diffusion into the leaf.

There is no clear explanation for how changes in $[\text{CO}_2]$ cause the change in aperture or conductance (Assmann, 1999), although it can be demonstrated that several physiological processes in guard cells, whether intact, in peels or in protoplast suspensions (e.g. carbon fixation, ion transport, chlorophyll fluorescence) can be affected by large changes in ambient $[\text{CO}_2]$. However, this is not the same as elucidating the signalling pathway for how a 50 or 100 $\mu\text{mol mol}^{-1}$ change in C_i results in a marked change of aperture in the intact leaf (e.g. Figure 12.9). It has been demonstrated convincingly that it is C_i not the external concentration that affects guard cells (Mott, 1988). From an entirely teleological point of view, a control mechanism based on C_i could be a way to link demand for CO_2 by the mesophyll cells with supply by the guard cells (Raschke, 1976). However, the many observations showing that g_s is largely independent of the rate of mesophyll carbon fixation over a short period of time (e.g. Figure 12.5) show that this is not the only controlling factor. Clearly, C_i interacts with other environmental signals and C_i is not *constant*, although it is *conservative*, because it is both the result and an effector of stomatal aperture (Jarvis and Morison, 1981).

Stomatal response to humidity

As described above in Equation (2), diffusion of gases through stomata depends on the difference in concentration between the inside and outside of the pore. If g_s is unchanged, the transpiration rate increases linearly with leaf-to-air vapour pressure difference (VPD) caused either by changes in air vapour pressure (w_a) or by leaf temperature affecting the vapour pressure inside the leaf, w_i . However, it is important to realize that VPD has a direct effect on stomata independently of the effect on whole leaf transpiration and it is the VPD, rather than relative humidity to which the stomata respond (Aphalo and Jarvis, 1991). Stomatal responses to VPD were first described by Schulze *et al.* (1972) who studied species living in desert habitats. In general, stomatal aperture declines as the VPD increases and, under certain conditions, the reduction in aperture may be so large as to reduce the transpiration rate (Farquhar, 1978), thereby preventing extreme water loss from the plant. In an elegant experiment, Mott and Parkhurst (1991) used differences in the diffusion rate of water vapour in different gases to conclude that stomata respond to water loss rates and do not directly sense, and respond to, the water vapour pressure near the leaf. Therefore, the most widely accepted mechanism for stomatal responses to humidity is that evaporation directly from epidermal and guard cells near the stomatal pore alters the guard cell water potential independently from that of mesophyll cells.

Figure 12.10a shows the effect of increasing w_a from 0.8 kPa to 1.88 kPa on leaf gas exchange in *Phaseolus vulgaris*. This caused a drop in the VPD and resulted in an initial drop in g_s for about 5 min, a 'hydropassive' stomatal behaviour (Stålfelt, 1955), caused by a change in water balance between epidermal and guard cells. This was followed by a steady increase for the subsequent 35 min, after which time g_s remained stable with an overall increase in g_s of 57%. The initial drop in g_s was matched by a fall in C_i , which then rose with increased g_s and remained stable after about 20 min; but further increase in g_s did not result in a further increase in C_i . A remained stable for the first 20 min, after which A increased with increasing g_s , which accounts for the lack of increasing C_i with increasing g_s . These results show that decreased VPD eventually allowed a larger g_s and consequently a higher C_i and higher A . An opposite increase in VPD (Figure 12.10b) led to an

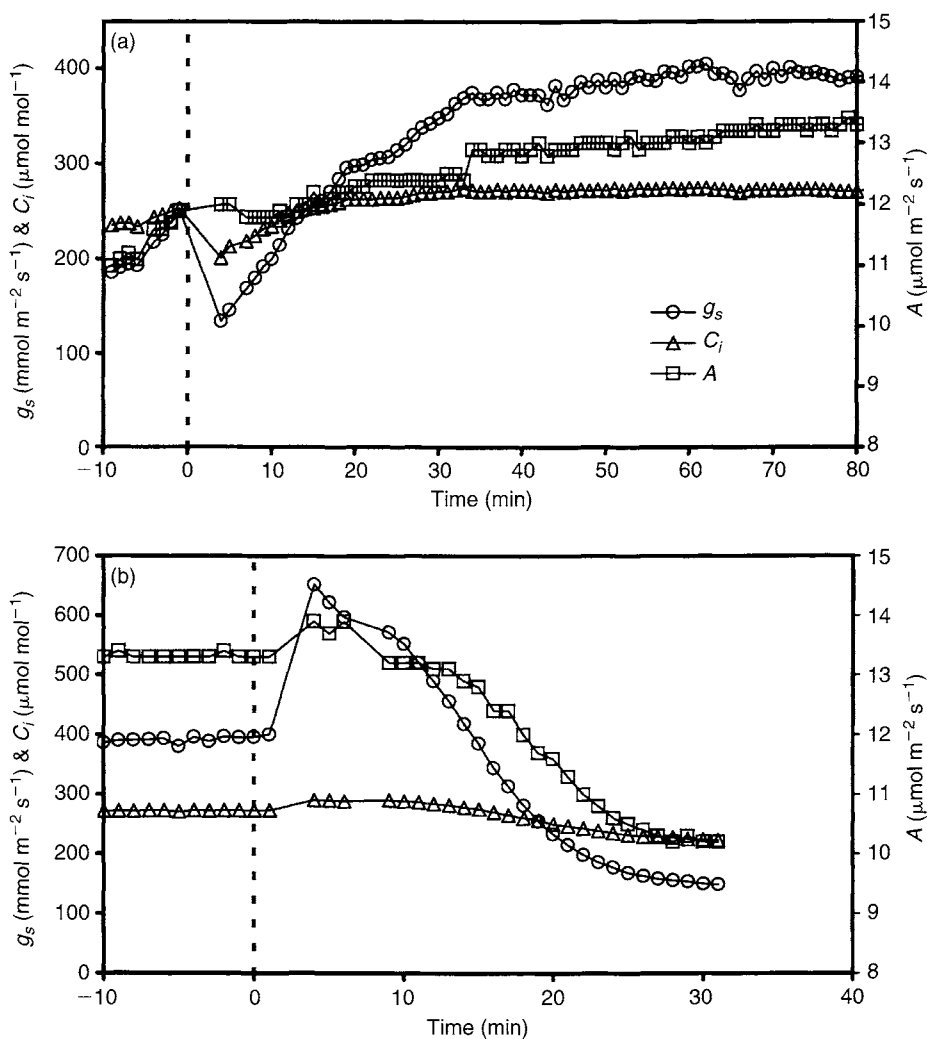


Figure 12.10 Effect of changes in humidity on stomatal conductance and CO₂ assimilation rate in *Phaseolus vulgaris*. (a) VPD was decreased from 1.74 to 0.77 kPa. (b) VPD was increased from 0.77 to 1.83 kPa. Assimilation rate is represented by squares, conductance by circles and C_i by triangles. Cuvette conditions were temperature of 21.6°C, with a PPFD of 245 μ mol m⁻² s⁻¹, and CO₂ concentration at 352 μ mol mol⁻¹.

increase in g_s for the first 5 min, followed by a slow decline. C_i initially increased slightly, then after 10 min it declined and remained stable around 270 μ mol mol⁻¹. A remained approximately constant during the initial 5 min, but dropped with decreasing g_s . Comparison of the figures (noting different axis scales) indicates that the rate of stomatal response was slightly faster with increasing, rather than decreasing, VPD. These examples show once again that changing environmental conditions can directly affect stomatal behaviour, which in turn affects assimilation rate, with consequent changes in C_i also affecting stomata. Disentangling these effects is difficult and has led to various 'systems analysis' approaches over the years, such as the feedback analysis of Farquhar and colleagues (Dubbe *et al.*, 1978).

Stomatal response to light

The effect of light on stomata was first recognized by Francis Darwin (1898), who noticed that a leaf facing a bright window had open stomata, while a leaf in the dark had closed stomata. Guard cells have chlorophyll (which is unusual, as other epidermal cells in many species do not) and stomata usually open in response to light in the photosynthetically effective wavelengths (blue through to red, 400–700 nm). It is incontrovertible that photosynthesis in the guard cells results in ATP and NADPH⁺ production, which can then be used in ion transport and possibly in carbon assimilation, although this last point is hotly debated (see Willmer and Fricker, 1996; Zeiger, 2002). In addition, there is high sensitivity to blue light (e.g. Zeiger *et al.*, 1981), as well as to various UV wavelengths (Eisinger *et al.*, 2000). The blue light response could be acting through various flavonoid and carotenoid pigments found in guard cells (Lu *et al.*, 1993). It has faster dynamics than the red light response, (particularly in grass (Poaceae) and sedge (Cyperaceae) species) and it may be involved in the rapid opening of stomata at dawn (Zeiger *et al.*, 1981) and during sun flecks (Kirschbaum *et al.*, 1988).

The overall light response is made up of several components. There is a direct photosynthetic and blue light response which is clearly evidenced by guard cell responses in epidermal strips and in guard cell protoplasts (e.g. Zeiger and Zhu, 1998). An indirect effect of light is also caused by the response of guard cells to CO₂ depletion in the intercellular air spaces due to mesophyll photosynthesis and in intact leaves it is hard to disentangle this from the direct response. A third effect of light may be through a signal transmitted from the mesophyll cells to the guard cells such that mesophyll photosynthesis controls the degree of stomatal opening (Heath and Russell, 1954; Wong *et al.*, 1979; Lee and Bowling, 1993). Such a messenger would explain the usually close positive correlation between photosynthetic rate and conductance, but the nature of any messengers is not yet clear. Sucrose movement is a possibility, as recently Outlaw and colleagues have shown that sucrose transport in the transpiration stream of *Vicia faba* can be a major source of organic carbon for the guard cells and can also exert an osmotic effect by accumulation in the cell apoplast (Lu *et al.*, 1997; Outlaw and De Vlieghere-He, 2001). However, the extent of this process in other species and its role in stomatal regulation in the field needs clarification.

In the whole leaf, the overall light response varies between species and growing conditions. Typically, whole leaf conductance saturates at photosynthetic photon flux density (PPFD) between 300 $\mu\text{mol m}^{-2}\text{s}^{-1}$ and 800 $\mu\text{mol m}^{-2}\text{s}^{-1}$ (Figure 12.11). In general g_l increases in parallel with A with increasing PPFD, whereas C_i remains relatively constant, except at low PPFD levels, when A is limited by light more than by CO₂ diffusion. However, there are marked differences between stomata on the upper and lower surfaces in response to light, as the latter open at much lower light intensities and have wider apertures (see review by Pemadasa, 1981), presumably reflecting their lower light environment. Sun and shade leaves also differ in responses (e.g. Turner, 1979), echoing the differences between upper and lower surfaces. In addition, it should be noted that the opening response to light is not universal, as plants with CAM photosynthetic metabolism show stomatal closure during daylight and opening at night. This is thought to be because the light response is overridden by the control of aperture by C_i (Willmer and Fricker, 1996).

As with other environmental factors affecting stomata, much of the work on light responses has been directed at examining 'steady-state' responses, where the stomatal aperture or conductance value has been measured when it has reached a quasi-constant value after a long period (tens of minutes) at a constant light intensity. However, under natural

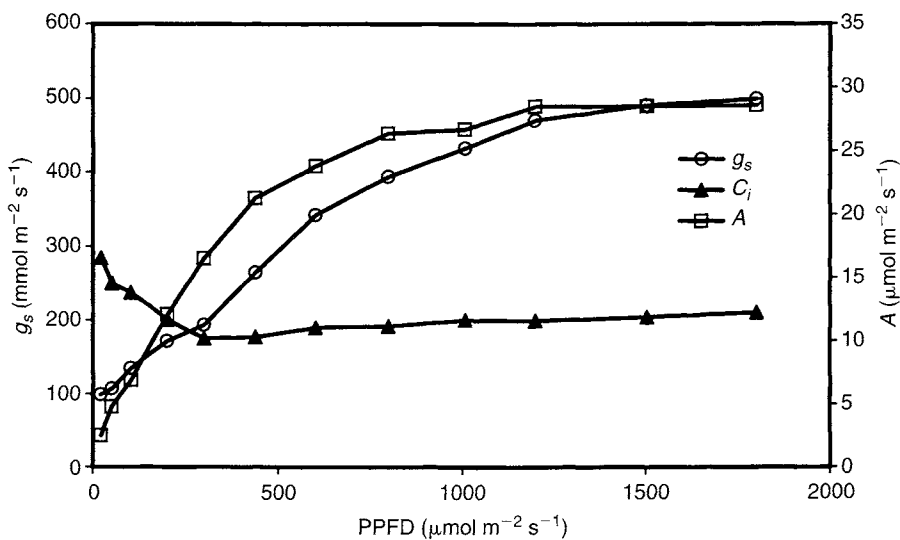


Figure 12.11 The response of stomatal conductance (g_s), assimilation rate (A) and internal CO_2 concentration (C_i) to PPFD in *Phaseolus vulgaris*. Cuvette conditions were 1.27 kPa VPD, temperature of 25°C and a CO_2 concentration of 372 $\mu\text{mol mol}^{-1}$.

condition leaves are exposed to a highly fluctuating light environment, with sun and shade flecks ranging between seconds and minutes (Barradas and Jones, 1996) caused by canopy movement and leaf flutter in the wind (Tang *et al.*, 1988) and cloud movements (Knapp and Smith, 1987). After a period of low light, an increase in irradiance does not result in an immediate increase in A , but shows a delay before maximum A is achieved. This lag period is due to both mesophyll photosynthetic induction (which involves the light regulation of key enzymes and changes in metabolite pool sizes) and changes in stomatal aperture (Percy, 1990). The changes in the metabolite pool sizes are relatively rapid, altering within seconds, compared to light regulation of enzymes which is in the order of minutes (Percy, 1990). Although the increase of g_s in response to a light increase during sun flecks is faster than the decreasing response to a drop in light, stomatal movements which can take up to tens of minutes, and can 'overshoot' – continuing to open after the sun fleck has passed (Kirschbaum *et al.*, 1988; Tinoco-Ojanguren and Percy, 1993). Therefore, stomata could limit assimilation rate during sun flecks. However, most work has indicated that the main control of assimilation during the first 10 min of induction is within the biochemistry of photosynthesis and that stomata do not cause a major limitation (Percy, 1990; Barradas and Jones, 1996).

To illustrate the effect of these different time lags on A and g_s , the effects of 5 and 15 minute artificial 'sun flecks' of 615 $\mu\text{mol m}^{-2} \text{s}^{-1}$ PPFD on a *Phaseolus vulgaris* leaf adapted to 215 $\mu\text{mol m}^{-2} \text{s}^{-1}$ are shown in Figure 12.12. During the 5 min 'sun fleck' (Figure 12.12a), A increased rapidly within the first minute and showed a reduced rate of increase up to a maximum at the end of the fleck after which A immediately dropped back to the original value. The largest change in g_s occurred after the sun fleck finished, with g_s continuing to rise steadily. During a 15 min 'sun fleck' (Figure 12.12), after starting with similar A and g_s values to those during the 5 min sun fleck, A showed a similar initial increase after 5 min, but continued to increase with increasing g_s to a value some 11% higher after 15 min. The stomatal conductance increased throughout the whole 15 minutes, although it was

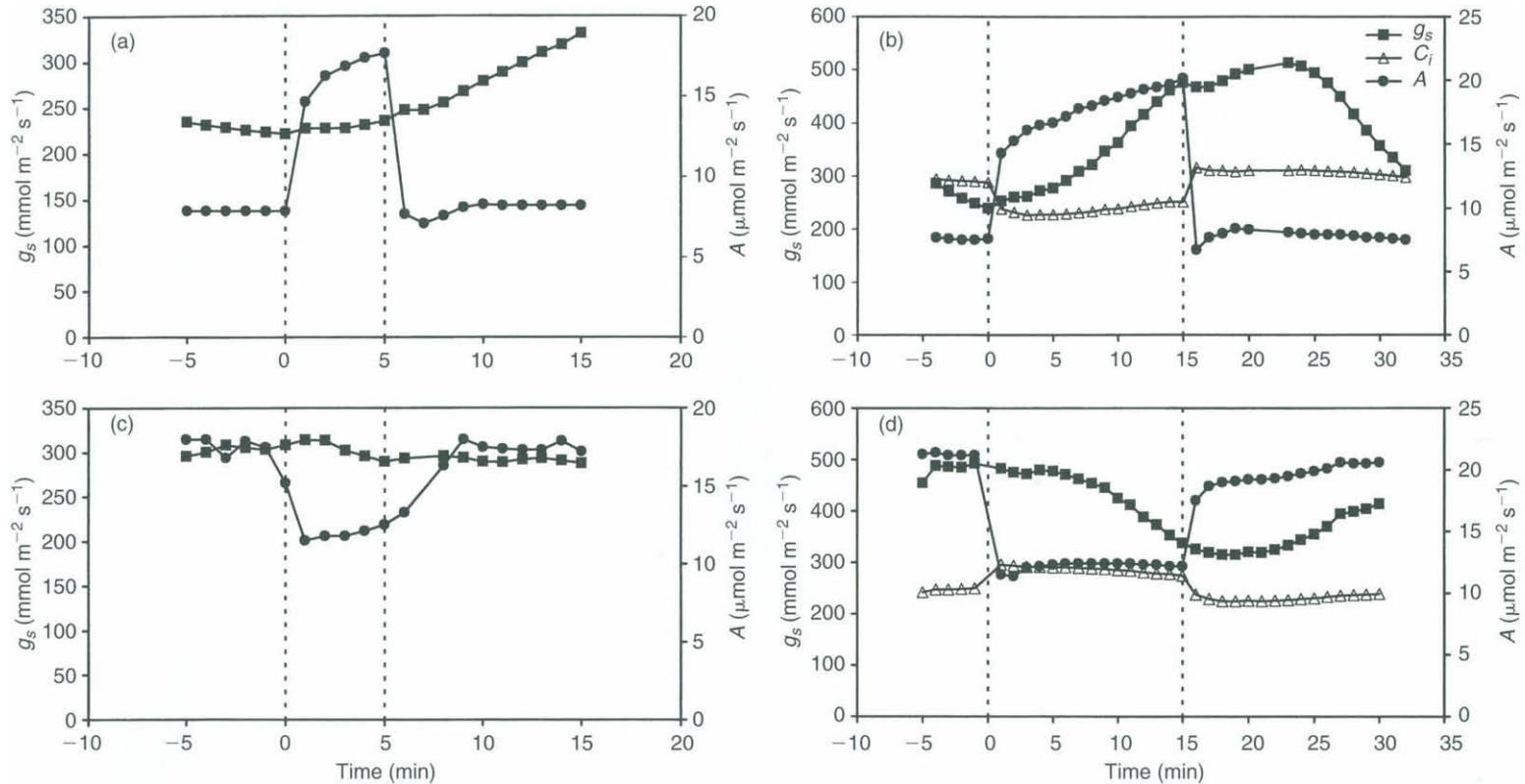


Figure 12.12 The effect of sun and shade flecks on CO₂ assimilation rate and stomatal conductance. (a) and (b) *Sun flecks*: PPFD was increased from 230 to 615 μmol m⁻² s⁻¹ at time zero for (a) 5 min, (b) 15 min. Measurements of assimilation rate (solid circles) and stomatal conductance (solid squares) made every 5 s. (c) and (d) *Shade fleck*: PPFD was decreased from 615 to 230 μmol m⁻² s⁻¹ at time zero. Photosynthesis is represented by solid circles and leaf conductance by solid squares. C_i is represented by open triangles. Readings were taken at 5 s intervals with cuvette CO₂ maintained at 357 μmol mol⁻¹. Cuvette conditions were maintained at 1.34 kPa VPD at a temperature of 25°C.

appearing to slow down by the end. After the sun fleck, the increase was small and continued for only 6–8 min, before closure started. Note that most of the C_i change occurred when illumination changed and, although there were subsequent small changes, changes in g_s were not sufficient to keep C_i constant. Of course, the effect of ‘shade flecks’ on A and g_s are potentially as important as sun flecks (Figure 12.12c and d). During both 5 and 15 min duration ‘shade flecks’ there was an immediate decrease in A which recovered back to its original value when the PPFD was restored in the 5 min fleck. However, A took about 15 min to return back to the original value after a 15 min shade fleck, because of the decrease in g_s that had occurred with the lower light. The start of a stomatal opening response was delayed for about 8 min after the end of the shade period and the slow recovery back towards its original value took over 15 min, consequently limiting A through CO_2 diffusion by some 10%. Such sun and shade fleck data emphasize the importance of the different dynamic behaviour of stomata and photosynthesis which can result in A and g_s being uncorrelated with each other, in the natural, changeable environment. Note also that these examples are with a species with rapidly responsive stomata, but there are other species with much more sluggish stomata (e.g. some conifers, Ng and Jarvis, 1980), where the correlation must be much less complete or frequent (Jarvis and Morison, 1981).

Environmental interactions and stomatal responses

Clearly, stomata are sensitive to a large number of environmental factors, but these rarely vary singly in nature, so the interaction between factors must be borne in mind. Two examples are shown here: the interaction between $[\text{CO}_2]$ and light (PPFD) and between $[\text{CO}_2]$ and VPD. Figure 12.13 shows the effect of increasing the ambient CO_2 concentration from 360 to 700 $\mu\text{mol mol}^{-1}$ on g_s and A (Figure 12.13a and b) at different PPFD. At each PPFD the leaf was left to stabilize for a minimum of 30 min before C_a was increased. Clearly, the initial rates of g_s and A depended upon the initial PPFD as we expect from the steady-state responses discussed above. In addition, the effect of increased C_a was much larger at higher (1000 and 500 $\mu\text{mol m}^{-2}\text{s}^{-1}$) than at lower PPFD. Note again that g_s took considerably longer than A to reach a new approximately steady value following the step change. As expected, A increased in response to high C_a as the extra CO_2 removed the supply limitation at high light. In contrast, the larger g_s at high PPFD was largely suppressed by the high C_a , showing clearly the interactive effect of CO_2 and PPFD on stomatal behaviour.

While light and $[\text{CO}_2]$ both influence assimilation rates directly, VPD does not. However, VPD affects stomata, so can indirectly affect assimilation rate through influencing CO_2 supply, as exemplified in Figure 12.13c and d. At normal ambient $[\text{CO}_2]$, g_s differed by about 70% between a VPD of 0.95 and 2.2 kPa, so that at the higher VPD assimilation rate was reduced by about 25% (as suggested by the example A/C_i curve in Figure 12.6). However, when the C_a was increased, A increased and the stomatal limitation was reduced. In all three VPD conditions, g_s followed a similar time course when CO_2 was doubled, taking over 20 min to come to a new steady value, twice that for A . Note also that the marked effect of VPD on g_s was reduced in high $[\text{CO}_2]$ which is in part due to the reduction in aperture and the consequent reduced ‘scope’ for stomatal aperture changes (see Morison and Gifford, 1984 for other examples).

Several other interactions have been examined, but there is obviously a potentially bewildering range of possible combinations and few studies have examined any more than two or three, and usually on only one or two species in the same conditions. For this

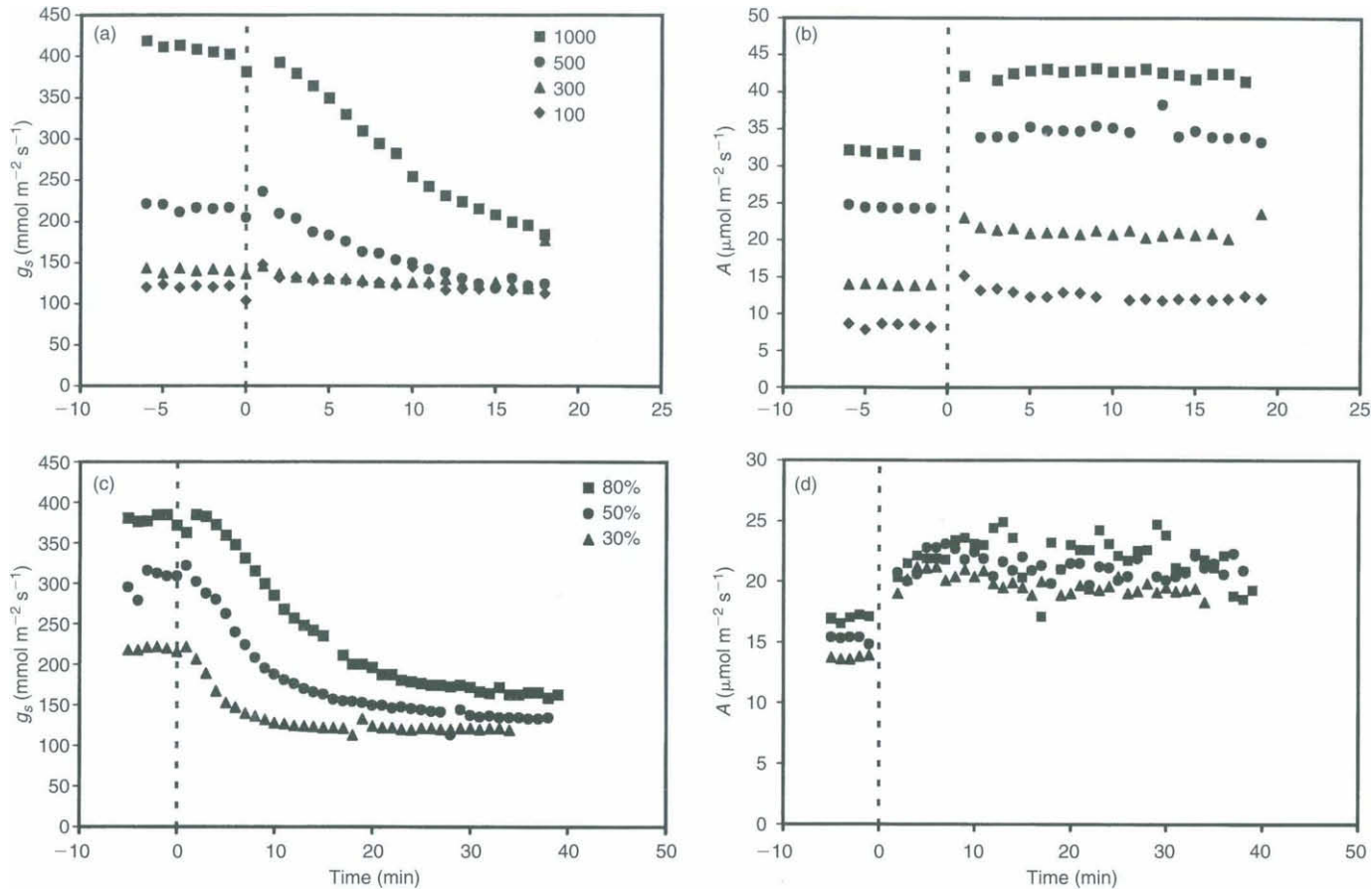


Figure 12.13 The effect of increasing CO₂ from 360 to 720 μ mol mol⁻¹ on stomatal conductance (a and c) and CO₂ assimilation rate (b and d) in *Phaseolus vulgaris* at different (a and b) PPFD or (c and d) humidity. Apart from the variable under investigation other cuvette conditions remained constant, at a PPFD of 1200 μ mol mol⁻¹; temperature 25°C; VPD 1.27 kPa, C_a 365 μ mol mol⁻¹. Initially the leaf was left to stabilize (for 20–30 min) at the conditions described above before a step increase in CO₂.

reason, much reliance has been placed on models for prediction of leaf gas exchange under different environmental conditions. There are two common types: the empirical model pioneered by Jarvis (1976) and the assimilation rate linked model suggested originally by Ball *et al.* (1987). The Jarvis-type model is based upon a multiplicative combined reduction of conductance from a maximum value depending on functions derived from measurements of stomatal responses to environmental factors. The Ball-Berry model and its subsequent refinements assume that stomata respond in order to regulate C_i . While both approaches have had considerable success for particular tasks, the problem is that they are not based upon the various mechanisms involved in stomatal behaviour and so will not capture the sorts of interactions demonstrated above, nor the dynamic changes illustrated. Some models that are more closely based on the physiological processes in guard cells have been devised (e.g. Farquhar and Wong, 1984; Jarvis and Davies, 1998), but while very promising, they are at the moment experimental and limited. It may be unrealistic to include all the many observed environmental sensitivities, but we urgently require physiological models that can integrate the key responses of stomata that are so readily demonstrated in the laboratory and the field.

Modern techniques for ecophysiological stomatal research

Although gas exchange techniques, such as exemplified here, have been important in monitoring stomatal behaviour in relation to mesophyll photosynthesis, they obviously do not probe what is going on at the single cell scale. Clearly, physiological and molecular biology work on guard cells in epidermal strips and other isolated systems is invaluable for probing the complexity of guard cell ion transport and metabolism. However, results from that research do need to be placed in the *in vivo*, *in natura* context of intact, photosynthesizing and transpiring leaves. Recently, there have been a number of cell scale investigations that open up new possibilities. First, microscope work with pressure probes has investigated guard cell mechanics and water relations (e.g. Franks and Farquhar, 2001). Secondly, imaging systems are now available to measure aperture changes in intact leaves, in the laboratory and even in the field as the environment changes and consequently relate these to gas exchange (e.g. Kaiser and Kappen, 2000). Thirdly, it is now possible to study guard cell photosynthesis *in vivo* using high resolution chlorophyll fluorescence imaging (Oxborough and Baker, 1997; Lawson *et al.*, 2002) which allows possible links between stomatal behaviour and underlying mesophyll photosynthesis to be studied. The basic principle in this latter technique is that fluorescence measurements are made with an imaging system through a microscope under the experimental conditions and under different light conditions when photosystem II (PSII) is in defined states (for details see Baker *et al.*, 2001). This results in measurements of the parameter F'_q/F'_m which estimates the yield of PSII photochemistry, often described as a measure of the efficiency of photosynthetic electron transport. Values from guard cell chloroplasts can be compared with those from the adjacent underlying mesophyll, and an example is shown in Colour Plate 2. This work has revealed that the photosynthetic Calvin cycle is functional in guard cells in a range of species and that the guard cell photosynthetic electron transport efficiency responds to light, water stress and $[CO_2]$ in a quantitatively similar way to that in mesophyll chloroplasts. The technique now makes it possible to compare a range of species which are different ecologically, taxonomically and evolutionarily, which will help build more mechanistic models of stomatal behaviour and the relationship with mesophyll assimilation.

Evolutionary context

Fossilized stomatal characters such as stomatal density (SD) have been used to determine past CO₂ concentrations (Kürschner, 1996; McElwain, 1998) and, in conjunction with other parameters, can be used to estimate g_s (Kürschner *et al.*, 1997). While in some species SD has been shown to correlate closely with g_s under certain conditions (e.g. Woodward and Bazzaz, 1988) the influence of SD on g_s is secondary to that of stomatal aperture (see Figure 12.2) which may not always be obtainable from fossil material. It may be possible to use a surrogate measure of aperture (Lawson *et al.*, 1998) or use pore length to give an indication of the maximum possible aperture (Beerling and Woodward, 1987). Maximum g_s can then be used to predict water-use efficiency and carbon balance (Beerling and Woodward, 1993), given assumptions about the link between conductance and assimilation rate (for example, see survey in Leuning *et al.*, 1995). While a general correlation between A and g_s certainly exists, there are many situations where stomatal aperture does not always correlate with assimilation rate (Jarvis and Morison, 1981) and as exemplified in the A/g_s curve shown in Figure 12.5. Furthermore, the evolution of stomatal behaviour may not have been driven by an optimization of instantaneous A with a minimum E , but may have been driven by other very different pressures such as the need to avoid water stress and runaway xylem embolism, or to avoid lethal high temperatures. Clearly, adaptation pressures may have differed according to habitats and conditions as they do in modern plants today. Therefore, we should be cautious about simple correlative models.

An example of how physiological understanding of plants can help to identify and explain plant fossils records has been shown by McElwain *et al.* (1999) who suggested that high levels of atmospheric CO₂ at the end of the Triassic may have resulted in the observed decrease in floral species in the fossil record. They suggest that high atmospheric CO₂ led to low stomatal density/index which resulted in reduced evapotranspiration rates and lethal increases in leaf temperature. Further evidence to support the theory that forced-CO₂ global warming can explain selective floral extinctions found in the fossil record has been published recently by Beerling (2002). Although this shows the important use of modern physiological understanding in determining past floral distribution, it has been pointed out that the use of anatomical parameters of fossil plants to reconstruct palaeoatmospheric CO₂ concentrations, which are then used to evaluate consequences for palaeoplant physiology, may compound errors already associated with assuming a direct correlation between stomatal properties and CO₂ (Beerling and Chaloner, 1993; Cowling, 2001).

It has been suggested that evolution of land plants should be considered in terms of the plants' ability to balance photosynthetic carbon gain versus water loss (Knoll, 1984). As both of these processes are under the direct influence of stomata, this implies that stomata could play a key role in plant evolution and distribution. Therefore, a full understanding of the physiology and function of stomata is important not only in modern plant physiology but could provide vital information about plant evolutionary processes. The study of modern plants compared with those that grew in the geological past could give an indication of the direction of such processes. For example, comparing ferns and cycads (which could be taken as representing evolutionary end points) with evolutionarily modern plants (such as angiosperms) could indicate evolutionary changes and adaptations. Such an experimental approach has suggested that ancient taxa such as ferns and cycads evolving under periods of high [O₂] and low [CO₂] may have photosynthetic systems with lower sensitivities to O₂ when compared with modern day angiosperms (Beerling *et al.*, 1998). However, when studying modern plant physiology it is important to remember that they have

developed over many hundreds of years, therefore biochemical and biophysical adjustments (acclimation) to their current environmental conditions could have taken place as well as genetic adaptation (Cowling, 2001) and therefore, behave differently today than in the past.

By studying modern C_3 species, C_4 and CAM plants we are looking at the results of millions of years of evolution and we can begin to understand how plants have adapted to changing environmental conditions. For example, the development of a CO_2 concentrating mechanism in C_4 and CAM photosynthetic pathways was probably the evolutionary result of changed atmospheric gas concentrations (Sage, 2001). Whether a particular species evolved a CAM or C_4 pathway depended upon the initial steps in the evolutionary sequence. These initial steps involved different selection pressures, in CAM this was associated with the scavenging of respiratory CO_2 during the dark, while in C_4 evolution it was associated with scavenging photorespiratory CO_2 in the light (Sage, 2002). Due to the CO_2 concentrating mechanisms displayed in CAM and C_4 plants, changes in atmospheric CO_2 concentrations have little effect on stomatal density, index or pore areas (Raven and Ramsden, 1988). Royer (2001) documented that only one out of nine C_4 plants studied revealed an inverse relationship with CO_2 concentration. Although stomatal responses to changing CO_2 concentration may be less significant in C_4 and CAM plants, it is possible that they still played a critical role in C_4 distribution and evolution. Sage (2001) suggested that many ecological factors, including CO_2 concentration, influenced where and when C_4 photosynthesis evolved. One such factor was the influence that low $[CO_2]$ contributed to aridification, through increased transpiration resulting from an increase in stomata aperture – such conditions would benefit the establishment of C_4 over C_3 plants.

The use of modern techniques may help to establish the timing of evolutionary processes, for example, the oldest C_4 fossils are believed to be 12 My old, however, phylogenetic evidence from molecular clock interpretations of genetic similarities in various grass lineages, indicates divergence of C_4 taxa was occurring 20–30 My ago (Kellogg, 1999). Further, comparison of responses of angiosperms with gymnosperms, CAM with C_3 plants and also plants from different environmental locations could reveal novel and interesting information about the evolutionary development of plant physiology. By combining palaeoecophysiology with modern plant physiology we will enhance our predictions of both future and past interpretations of global climate environments and associated plant responses. For example, recent work by Sage (2002) suggests that low CO_2 probably acted as a significant evolutionary agent, selecting plants adapted to CO_2 deficiency and that adaptations to low CO_2 might still exist in plants today, which might constrain responses to the current rising CO_2 concentration. Such acclimation would have important implications for current agriculture, water-use efficiency and natural selection.

Conclusion

A detailed understanding of stomatal behaviour and function is essential to understand both past and present impacts of environmental change on global carbon and hydrological cycles. In this chapter we have looked at the role of stomata in leaf gas exchange and the way they respond to three key environmental variables of light, CO_2 and humidity. Other factors such as leaf water status are obviously as important as these to plant growth and survival, but are less directly amenable to short-term experiments. We have tried to

emphasize the role of stomata in determining mesophyll CO₂ assimilation rates. Examples of physiological variation at a number of different levels stress the importance of appropriate sampling strategies for both modern plant physiologists and those attempting to use stomatal characters as indicators of past environments. The effects of a number of environmental factors, both singly and when interacting, on both stomatal and photosynthetic behaviour have exemplified the complexity of the relationship between mesophyll assimilation and stomatal function. In the final sections, examples of modern techniques for studying stomatal physiology have been illustrated. Use of such techniques should improve the elucidation of stomatal behaviour and mechanisms in a range of species (from ancient to modern), and hence reveal evolutionary changes.

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13

The photosynthesis–transpiration compromise and other aspects of leaf morphology and leaf functioning within an evolutionary and ecological context of changes in CO₂ and H₂O availability

Pieter J C Kuiper and Cécile M H Lapré

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Introduction

This chapter begins with a description of the 'trade-off' between photosynthesis and water loss by transpiration in desert plants. Is saving water a key strategy for desert plants? The stomatal response to atmospheric CO₂, recently and in the past, is another factor affecting the water-use efficiency of plants. The 'trade-off' between photosynthesis and transpiration is only one of several compromises and a strategy with emphasis on a single environmental factor should be replaced by strategies in which the whole environment, including the ecosystem (vegetation) structure and seasonality, is taken into account.

Modern Mediterranean ecosystems are chosen as an example of seasonality, in which the strategies concerning maintenance of photosynthesis and water relations should operate in the dry and hot summer as well as in the wet and cool winter.

During the wet season rainfall and fog may inhibit photosynthesis by blocking stomatal opening with raindrops. Also, a dripping point of the leaf and positioning of the leaf within the canopy are important in rainfall interception and possible inhibition of photosynthesis. Other factors, such as wettability of leaves and the presence of wax plugs in the stomatal pore and a hair layer over the leaf, also may alleviate negative effects of rainfall on photosynthesis. Next to inhibition of photosynthesis by rain, wet leaves are susceptible to fungal leaf pathogens; also, during rainfall leaves may leach nutritional ions.

Photosynthesis of wet leaves may be restored by drying of the leaves by evaporation of the intercepted rain. The rate of drying of wet surfaces of different sizes is formulated under conditions of an ample energy supply (diffusion law of Fick, see Martin (1943), e.g. in an open vegetation) and under energy limitation (Penman-Monteith equation, see Monteith and Unsworth (1990), e.g. in a dense forest). The role of the (wet) leaf size in the rate of drying is included in these formulations. In addition vegetation structure is included as a factor influencing the rate of drying of wet vegetation: drying of wet leaves in a dense forest with a closed canopy will be limited by available energy for evaporation sooner than an open and heterogeneous Mediterranean ecosystem with small leaves and a low leaf area index. A simulation example will be presented.

The review concludes with a discussion regarding how differences in leaf morphology could be functionally related to 'trade-off' between photosynthesis and water relations in a Mediterranean ecosystem.

'Trade-off' between photosynthesis and transpiration

Desert plants

In the desert environment saving water via a 'trade-off' between photosynthesis and water loss by transpiration is a popular view (Gibson, 1998). Indeed CAM succulents such as cacti, euphorbs and stapeliads exhibit a high degree of water saving by having closed

stomata during the daytime and open stomata during the night (Gibson and Nobel, 1986). They tolerate extremely high temperatures in the stem at midday because of a lack of evaporative cooling.

Non-succulent desert plants that have both leaves and photosynthetic stems are very well suited for maximizing photosynthesis: under an adequate water supply the leaves are best suited for absorption of light and CO₂ together with evaporative cooling. At the same time stems are better at saving water, as evident from the less negative carbon isotope ratio of the stem tissue (Ehleringer *et al.*, 1987). The relationship between water-use efficiency and carbon isotope ratio can be explained as follows. During photosynthesis, CO₂ diffuses into the leaf via the stomata, while simultaneously H₂O diffuses outwards to the atmosphere. Because of its lower molecular weight, ¹²CO₂ diffuses at a faster rate into the leaf than the heavier ¹³CO₂ causing an enrichment of ¹²C in the plant tissue, compared with atmospheric CO₂. This effect of ¹²C enrichment in plants can be counterbalanced by an increase in affinity for CO₂ in the plant tissue, resulting in a change in the carbon isotope ratio, as mentioned above for stems of desert plants.

Non-succulent desert plants exhibit a wide range of responses to the desert climate. In a careful review, Gibson (1998) doubts that saving water is the key strategy for these plants, but that maximizing carbon gain by photosynthesis is at least as important. Annual desert plants show the highest P_{\max} values, followed by shrubs with drought-deciduous leaves and evergreen shrubs. The longer time span of photosynthesis of an evergreen, such as the creosote bush (*Larrea divaricata* Cav.), compensates for a lower rate of photosynthesis with its deep root system (Table 1 in Gibson, 1998). In conclusion, there is no single and generally valid 'trade-off' between photosynthesis and transpiration in desert plants.

Stomatal response to atmospheric CO₂

In 1987 Woodward published his observation that stomatal numbers were sensitive to an increase of atmospheric CO₂; since pre-industrial times a gradual decrease in stomatal density with time has been observed. More recently Wagner *et al.* (1996) demonstrated that global climate change, in the form of an effect of increasing atmospheric CO₂ on plant performance, is clearly present: analysis of the buried leaves of an isolated birch tree (*Betula pendula* Roth) in a Dutch bog revealed that stomatal density of the leaves from this tree had decreased from 270 per mm² in 1955 to 180 per mm² in 1995, a response in line with the increase in atmospheric CO₂ during these 40 years. In the same birch leaves a similar reduction in the stomatal index, the proportion of stomata as a percentage of the total number of cells (epidermal cells + stomatal cells), was noted.

Other tree species, e.g. durmast oak (*Quercus petraea* Liebl.), showed a similar reduction in stomatal density and index when tree seedlings were exposed to elevated CO₂ in a climate room experiment (Kürschner *et al.*, 1998). Transpiration was reduced at elevated CO₂ by partial closure of the stomata, as has been observed in other tree species (Ceulemans and Mousseau, 1994; Curtis, 1996). There is an evolutionary effect of the response of stomata to elevated CO₂; some coniferous species show no reaction to elevated CO₂ (Eames and Jarvis, 1989). In modern species stomata close as a response to high CO₂. Stomatal opening is stimulated at low CO₂, an adaptive response to the limiting CO₂ concentrations during the ice age intervals (Kuiper, 1998).

It is important to test the validity of the field observations on the above mentioned birch tree (Wagner *et al.*, 1996) on other tree species (e.g. *Eucalyptus*). By their high transpiration rate eucalypt forests maintain a low level of saline groundwater and thus

prevent dry land salinization. Removal of forest for agriculture may result in rising groundwater and consequent salinization. A similar stomatal response to increasing atmospheric CO₂ in *Eucalyptus*, as observed in birch, would contribute to dry land salinization and provide part of the explanation for the present-day dramatic dry land salinization in south and western Australia.

So far, discussion has focused on the effect of atmospheric CO₂ on stomata. The tree fern, *Cibotium schiedeii* (Dicksoniaceae), has a leaf epidermis almost exclusively of stomata with an occasional parenchyma cell (Eschrich, 1995). In this ancient group of plants epidermal parenchyma cells may have evolved during times of adequate atmospheric CO₂, eventually leading to the modern pattern of a CO₂-dependent stomata/parenchyma ratio of the leaf epidermal cells.

C₄-photosynthesis is the evolutionary answer to low atmospheric CO₂ concentrations in more recent geological times (see an extensive review by Sage, 2001).

Analysis of the 'trade-off' response within a broader context

As mentioned in the introduction, the 'trade-off' between photosynthesis and transpiration is only one of several compromises and a strategy with emphasis on a single environmental factor such as water and CO₂ availability should be replaced by strategies in which the whole environment, including ecosystem (vegetation) structure and seasonality of the environment, is taken into account. Such an approach is also in line with an evolutionary approach: evolutionary changes in plant populations and species occur within an ecosystem context. Sage (2001) also stresses the point that the evolution of C₄-photosynthesis should be analysed within the context of a combination of environmental conditions, such as high light and high temperature, salinity and aridity.

Modern Mediterranean ecosystems were chosen as an example of seasonality, in which the adaptive strategies concerning maintenance of photosynthesis and water relations should operate during the dry and hot summer as well as in the wet and cool winter. Besides, in line with continental drift, many land surfaces have moved from a situation with a relatively constant environment to one with seasonality, as demonstrated by the splitting up of the ancient Gondwana palaeo-continent.

Modern Mediterranean ecosystems

Dry season

During the dry season, water availability will determine photosynthesis and transpiration of the plant species. In the South African Fynbos, a Mediterranean ecosystem with a relatively low drought stress, three growth forms are recognized: shallow-rooted restios and ericoid shrubs (see Colour Plate 3) exhibit a greater response to drought in the form of a strong reduction in water potential and photosynthesis in summer; the deeper rooted proteoid growth form maintains photosynthesis and water potential in summer (Stock *et al.*, 1992). Wind will cool the foliage of the restioid and ericoid growth forms by heat exchange between leaf and air and wind will cool the leaves of the proteoid growth form by heat exchange as well as by increased transpiration, under ample water availability. The so-called sclerophylly in the above growth forms has not been entirely attributed as an adaptive trait to drought stress: in particular in South Africa and Australia a sclerophyllous

leaf structure has been related to nutrient use efficiency under the relatively nutrient-poor habitats of these regions, phosphate in particular (Beadle, 1966; Specht and Rundel, 1990; Stock *et al.*, 1992; Keeley, 1992). Sclerophylly and low nutrient levels in the leaf are linked together since sclerophyllous leaves are rich in structural carbon compounds such as cellulose and lignins, which lead to a dilution of nutrient concentration per unit leaf weight. Moreover, low nutrient levels in combination with a high content of cellulose and lignin will reduce edibility and act as a defence against herbivory (see also Johnson, 1992).

Wet season

During the wet season the above mentioned growth forms of the Fynbos and similar growth forms of other Mediterranean ecosystems are physiologically active due to an ample water supply and moderate temperatures. No 'trade-off' between photosynthesis and transpiration is evident, on the contrary, CO₂ exchange between the atmosphere and leaf and, consequently, photosynthesis may be hindered by rain and fog: leaves may become wet and covered with a layer of water. The rate of diffusion of CO₂ in liquid water is around 10 000 times slower than in air (Nobel, 1991), which explains an inhibition of CO₂ exchange in rain-wetted leaves. To give an example, after a short rain shower the leaves of an ash tree (*Fraxinus excelsior* L.), located at the Biological Centre, Haren, The Netherlands, were covered with a layer of water of 0.03 mm, measured by radar (de Jong, 2001). Diffusion of CO₂ through this liquid film layer is equal to diffusion through a gaseous leaf boundary air layer of 30 cm; in reality the latter is around 0.5 to 1 cm under still air conditions in a room (Kuiper, 1961; Gates, 1980; Monteith and Unsworth, 1990). In this case the wet ash leaf has no appreciable rate of CO₂ exchange and a strong wind is essential to dry the leaf quickly for restoration of photosynthesis.

The problem of the low rate of diffusion of CO₂ in liquid water has long been recognized in studies on CO₂ exchange in submerged aquatic higher plants and algae. Various mechanisms to counteract this negative effect on photosynthesis have been described: (1) a more effective CO₂ concentrating mechanism; (2) conversion of bicarbonate into extra CO₂ by acidification of the boundary layer of water surrounding the leaf of higher plants; and (3) direct uptake of bicarbonate in algae (Elzenga and Prins, 1989). Surprisingly, the possible negative effects of a layer of water or water droplets on photosynthesis in wet leaves due to rain have never attracted much attention and, with the exception of Horton (1919), have only recently been adequately studied (Smith and McClean, 1989; Ishibashi and Terashima, 1995; Brewer and Smith, 1997; Feild *et al.*, 1998; Hill and Brodribb, 2001). The next section deals with the various strategies plants have developed to reduce prolonged wetting of leaves and facilitation of drying. In addition to inhibition of photosynthesis in wet leaves, rain may also leach nutritional ions from the wet leaves and stimulate fungal leaf pathogens.

Prevention of wetting of leaves by rain and facilitation of drying

Dripping tip of leaves and position of the leaf

In tropical rainforests, leaves of several species possess a *dripping tip*, which may facilitate water runoff (Berrie *et al.*, 1987). In addition, such leaves are often hanging – the perpendicular position may facilitate runoff of water by gravity and result in a thinner water film on the leaf. In Mediterranean ecosystems numerous *Eucalyptus* species exhibit vertical leaves, with a possible similar function. The sleeping position of leaflets of many Leguminosae at night may also facilitate rainwater runoff during the night.

Wettability of leaves

Wettability of the leaf is involved in the prevention of the formation of a water film on a leaf; a low adhesion between leaf surface and raindrops is created by a waxy cuticle which, as in *Ficus elastica* Roxb.ex Hornem. and *Asplenium scolopendrium* L., may prevent the formation of a thin film of water on the leaf. On the other hand, leaf surface roughness may facilitate the aggregation of small rain droplets into larger ones which eventually may roll off the leaf surface. In several plant species leaf veins are slightly sunken in the leaf surface and thus may function as 'gutters'.

Horton (1919) was the first to describe raindrop interception on leaves of various species. Smith and McClean (1989) observed an adaptive relationship between leaf water repellency and stomatal distribution over both sides of the leaves. In 50 out of 57 species the leaf side with the highest stomatal frequency was also the side with the highest water repellency. Application of a fine mist to bean plants caused closure of the stomata within 2 min and partial recovery of stomatal opening within 60 min (Ishibashi and Terashima, 1995). Photosynthesis was reduced similarly: an abrupt reduction at the onset of rain, followed by partial recovery. In addition, a long-term negative effect of 24 h of rain was measured under dry conditions, indicating that leaf wetness caused not only immediate reduction of photosynthesis, but also damage to the chloroplast. Brewer and Smith (1997) introduced an elegant method to determine leaf water repellency: a droplet of 5 mm³ water was placed on a horizontal leaf and its contact angle was determined by measuring the line tangent to droplet through the point of contact between droplet and leaf surface. In the montane and alpine area of the Rocky Mountains (USA) plant species of open meadows were characterized by more frequent leaf wetting by rain and dew than species of the forest understorey. As an adaptive response, species of open meadows were less wettable and had more stomata than species of the understorey. Also, in species of the open meadows the presence of leaf trichomes reduced the area of leaf surface covered with water.

Avoidance of direct contact between water film and stomata: effect of a hair layer and stomatal wax plugs

CO₂ exchange in a leaf takes place via the stomatal pores in the epidermis and wetting of the leaves may block the stomatal pore with liquid water. Bosveld (1999) observed a difference between measured and calculated transpiration in a partially wet Douglas-fir (*Pseudotsugo menziesii* Franco) forest: this difference could be explained by assuming that one-third of the stomata were blocked by rain under wet conditions.

Blockage of the pores by rainwater may possibly be prevented by several adaptive strategies, e.g. an interrupted water film on the leaf by low wettability of the cuticular surrounding of the stomatal guard cell in combination with a sunken location of the stomata. Feild *et al.* (1998) observed that stomatal cutin plugs of *Drimys winteri* Forster and Forster, a tropical cloudforest species, protect leaves from mist: removal of the stomatal plugs resulted in a marked increase of wettability together with reduced photosynthesis. Leaves without stomatal plugs showed closure of the stomata with increasing evaporative demand, a usual reaction for higher plants. Intact leaves with stomatal plugs remained open, as expected from a cloudforest tree.

Another possibility is the presence of a hair layer on the leaf, which prevents formation of a continuous water film so that direct contact of the rain drops with the stomatal pores is no longer possible (Brewer and Smith, 1997). From the Australian macrofossil record of Proteaceae and Casuarinaceae, Hill and Brodribb (2001) conclude that a dense covering of the leaves with trichomes primarily functions as mechanism of keeping the

water off the leaf surface. *Hypericum elodes* L., an Atlantic species of shallow lakesides, is characterized by tomentose leaves and it is very unlikely that the hair layer has a function in drought tolerance. Alpine vegetation in Papua New Guinea with a continuous wet climate also contain many species with tomentose leaves (Hope, 1986; Lapré, personal observation). In alpine plants pubescence of the leaves is a major factor for retention of rain and fog, thus contributing to the negative effects of acid rain to natural vegetation (Monson *et al.*, 1992).

Grasses like *Ammophila arenaria* (L.) Link create a microatmosphere, which is inaccessible for raindrops by lengthwise rolling of the leaf blades (Massart, 1907).

Another way to avoid direct blockage of stomatal pores by raindrops is the exclusive location of stomata on the underside (hypostomatous) of the leaves as evident in numerous deciduous trees: raindrops will hit and wet the upper surface, while the lower surface with stomata will only be wetted after a more prolonged period of rain.

Further ecological consequences: effect of fungal leaf pathogens and ion leaching

In the above sections the beneficial effects of prevention of wetting and fast drying of leaves on photosynthesis have been discussed. Prevention of wetting and fast drying of leaves may also be of ecological significance in other cases, e.g. as in prevention of infection of leaves by microbes and fungal pathogens: the longer the leaves stay wet, the more likely a possible fungal infection of the leaves, especially at the end of the wet season when temperature rises. Presence of water on the leaf surface is important for germination of spores and for hyphal growth of fungal pathogens through stomatal pores (Butler, 1996). Wind speed is important in leaf wetness, since the amount of intercepted water on the leaf will be limited by increasing wind speed. In addition, the rate of drying increases with wind velocity as noted herein.

Another negative effect of continued wetting of the leaf canopy is loss of nutritional ions by leaching (Tukey, 1962, 1970; Ovington, 1968; De Luca d'Oro and Trippi, 1987). Leaching of young plants with distilled water for 24 h resulted, in some species, in considerable losses of Ca (up to 31%), Mg (up to 27%), K (up to 20%) and P (up to 16%; Tukey, 1962). Measurements of K, Ca and Mg in precipitation at ground level in hardwood and coniferous forests demonstrated high concentrations of these ions under the forest canopy, compared with an open field (Ovington, 1968). More recently Gordon *et al.* (2000) showed species-specific differences in red spruce (*Picea rubens* Sarg.), black spruce (*Picea mariana* (Mill.) Britton) and white spruce (*Picea glauca* (Moench.) Voss) plantations in the nutrient content of rainwater that had reached ground level after passage through the canopy: beneath white spruce precipitation was enriched in total N, organic N, NH_4^+ and K, compared with the other two species; the canopy of all three plantations acted as a sink for P during rain. Species differences in nutrient cycling were ascribed to differences in morphology and crown structure of the spruce species.

Drying of wet leaves by evaporation

Formulation of the rate of drying of a wet surface

In this section, first a description of the effect of leaf size and shape on evaporation from wet leaves is given, following Fick's diffusion law. Afterwards the effect of energy limitation of evaporation is discussed, using the Penman–Monteith equation (Monteith and Unsworth, 1990).

Under conditions when there is a direct contact between the water film and leaf surface and the stomatal pores are blocked by liquid water, the resulting question is how quickly a leaf becomes dry again and CO₂ exchange can be restored. Climatic factors, which determine the rate of evaporation from a wet surface are wind velocity and the difference between the saturation vapour pressure of the evaporating wet surface (at leaf surface temperature) and the actual vapour pressure of the atmosphere.

Leaf size and shape are important for the rate of drying. Aerodynamic properties of the evaporating leaf will determine the thickness of the boundary air layer adjacent to the evaporating surface: under identical external conditions small leaves will dry more quickly than leaves with a larger surface and, as will be shown below, long, narrow leaves will dry quicker than short and wide leaves of the same surface area (under condition that the wind direction is parallel to the length of the leaf). In addition, the leaf edge is important: a serrate or dentate leaf edge will facilitate air turbulence thus reducing the boundary air layer with its laminar flow of water vapour. No data on evaporation of wet leaves are available, but as a first approximation evaporation of rectangular pieces of wet blotting paper has been measured and formulated (Martin, 1943; Penman, 1948; Raschke, 1956, 1960; Kuiper, 1961; Gates, 1980) in wind tunnel experiments under the condition of ample energy supply for evaporation. More recently, Brenner and Jarvis (1995) confirmed these earlier results.

Formulation of the combined results yields:

$$E = 0.73 \cdot k \cdot (e_s - e_d) \cdot W^{-0.2} \cdot L^{-0.3} \cdot V^{+\alpha} \quad (1)$$

in which E = evaporation of a rectangular piece of blotting paper ($\text{g} \cdot \text{h}^{-1} \cdot 100 \text{ cm}^{-2}$), k = coefficient of diffusion of water vapour ($\text{cm}^2 \cdot \text{sec}^{-1}$), $e_s - e_d$ = vapour pressure difference in mmHg at atmospheric pressure, W = width (cm) of the evaporating area at right angles to the wind direction, L = length (cm) of the evaporating area parallel to the wind direction, V = wind velocity ($\text{cm} \cdot \text{sec}^{-1}$), α = a proportionality factor, 0.78 (Raschke, 1956) or 0.76 (Kuiper, 1961) for turbulent air flow and 0.5 (Kuiper, 1961) for laminar air flow (see also Brenner and Jarvis, 1995). Reformulation of (1) gives:

$$E/0.73 \cdot k \cdot (e_s - e_d) \cdot V^{+\alpha} = W^{-0.2} \cdot L^{-0.3} \quad (2)$$

A proportionality factor, K , is introduced for the first part of equation (3):

$$K = W^{-0.2} \cdot L^{-0.3} \quad (3)$$

in which $K = 1$ for W and L both being 1 cm, a quadrant of 1 cm^2 .

Figure 13.1 summarizes values for K for evaporating surfaces with W and L varying between 0.1 and 100 cm. High values of K implicate a relatively high rate of evaporation and fast drying of a wet leaf surface and low values of K the reverse. $K = 3.16$ for W and L both being 0.1 cm, a quadrant of 0.01 cm^2 , indicative for a very small leaf (upper left corner of Figure 13.1). K = intermediate for long and narrow quadrants, e.g. $K = 0.47$ for $W = 0.5 \text{ cm}$ and $L = 20 \text{ cm}$, representative for grass- and rush-like leaves and very low for larger quadrants (lower right corner of Figure 13.1). In this figure the curves for varying width merge at increasing length, indicating that differences between the rate of drying of wet surfaces are most visible at low values of W and L .

It is important to realize that the effect of large differences between high and low values of K on drying may be considerably reduced, because heat exchange is also dependent

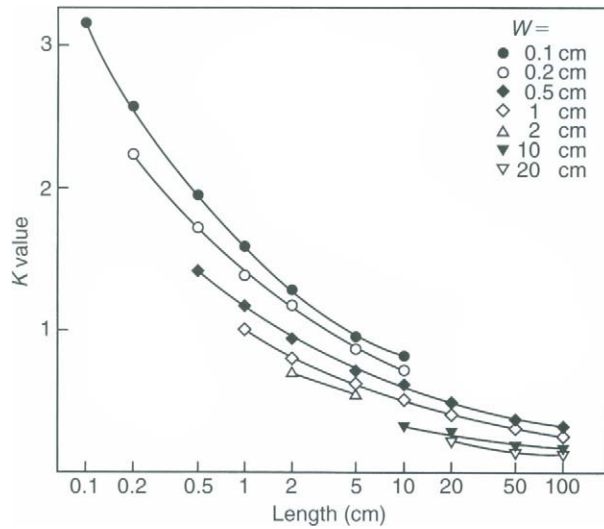


Figure 13.1 Effect of length (L) and width (W) of a rectangular wet surface, as a model for plant leaves, on K , the evaporation rate per unit surface and per unit vapour pressure difference between wet surface and air, including an effect of wind velocity. On the ordinate K is expressed as:

$$K = E/0.73 \cdot k \cdot (e_s - e_d) \cdot V^{+\alpha} = W^{-0.2} \cdot L^{-0.3}$$

in which E = evaporation per unit time and surface, k = diffusion coefficient of water vapour, $e_s - e_d$ = vapour pressure difference between wet surface and air, V = wind velocity and α = a proportionality factor. High and low values of K represent respectively high and low rates of evaporation, due to the length and width of the evaporating surface.

on leaf morphology, causing a counterbalancing effect of the surface water vapour saturation pressure, e_s , which is highly temperature dependent. Even more important, the energy demand for evaporation of water may easily limit evaporation and transpiration of partially wet leaves and leaf temperature and leaf vapour pressure will drop, resulting in reduced effects of leaf size on the rate of drying. Also, competition for energy between transpiration and evaporation of the partially wet needle surface of Douglas fir was evident (Bosveld, 1999). Water storage by leaves during rain and the simultaneous evaporation of components of rainfall interception are analysed by Lankreijer *et al.* (1993) and Klaassen *et al.* (1998).

The Penman–Monteith equation removes the problem of the strong temperature dependence of the vapour pressure of water: two components, the energy requirement for evaporation and the water vapour pressure difference between leaf and air, are combined in a single formulation (see Monteith and Unsworth, 1990). Heat transport by convection (sensible heat):

$$C = \rho c_p (T - T_0) / r_H \tag{4}$$

and heat transport by evaporation (latent heat):

$$\lambda E = \rho c_p (e_s - e_d) / \gamma^* r_H \tag{5}$$

With incoming radiation, R_n , the heat balance is:

$$\lambda E + C = R_n \quad (6)$$

Combination of (4), (5) and (6) yields:

$$\lambda E = \frac{\Delta R_n + \rho c_p (e_s - e_d) r_H^{-1}}{\Delta + \gamma^*} \quad (7)$$

in which: C = flux of heat per unit area, ρ = density of air, c_p = specific heat of air, T = temperature, T_0 = temperature of surface, r_H = resistance for heat transfer for convection, λ = latent heat of vaporization, E = evaporation rate, γ^* = apparent value of psychrometer constant, R_n = net radiation flux density and Δ = rate of change of saturation vapour pressure with temperature.

Equation (7) contains a radiation component and a vapour pressure component. The contribution of these components to drying of wet vegetation now will strongly depend upon the following factors:

1. The amount of incoming radiation for energy supply
2. The conductance of the vegetation to heat and evaporation fluxes, which in turn will depend upon leaf area and leaf distribution within the vegetation.

Clearly vegetation structure will be a decisive factor for the rate of drying: a high vegetation density as in crops and forests with a closed canopy will easily lead to a limitation of drying, due to lack of energy, especially at low net radiation (see Lankreijer *et al.*, 1993; Klaassen *et al.*, 1998; Bosveld, 1999; Klaassen, 2001).

The situation in Mediterranean vegetations (see Colour Plate 4) is quite different from the above situation with:

1. A low vegetation density
2. A low leaf area index
3. An effective vertical distribution of the leaves
4. Strong wind during the rainy winter.

The radiation component will not so easily limit drying of wet Mediterranean vegetation as is the case with high yield crops and forests with a closed canopy.

A detailed study of drying of wet forest in relation to canopy wetness, canopy cover and net radiation is given by Klaassen (2001). A micrometeorological model with a detailed representation of a forest canopy, including a vertical distribution of the leaf area index (LAI) over 20 m, was evaluated against experimental results. The vertical distribution of sensible and latent heat (equations (4) and (5) respectively) in a wet and homogeneous forest was simulated. Evaporation of the wet canopy increased with canopy cover and, because of increasing energy limitation and increasing canopy roughness, the increase in canopy evaporation with canopy cover gradually slowed down. As expected, a patchy distribution of canopy leaves facilitates drying of the leaves and the drying rate in a patchy canopy cover was stimulated more at low canopy cover: 28% (cover = 0.2), decreasing to 4% (cover = 0.8).

A similar procedure was followed for canopy evaporation versus canopy wetness: after the start of rain, canopy evaporation strongly increased with canopy wetness and reached

a rather constant level when canopy wetness approached 0.5. When the rain stopped, a fully wet canopy exhibited the fastest drying rate, which strongly decreased until a wetness value of 0.6 was reached.

In conclusion, the above described methodology may be very helpful to compare forests and shrub vegetations with different heights and canopy structures – a closed canopy versus an open and often patchy structure – and to relate these canopy structures to possible adaptive leaf morphology.

Ecological consequences of leaf morphology on the rate of drying of wet leaves

Wetted small leaves are characterized by high values of K and thus will dry relatively quickly (see Figure 13.1). In the Mediterranean Fynbos of South Africa wetted leaves of the restioid and ericoid growth form will dry quicker than leaves of the proteoid growth form, enabling the first two growth forms to perform photosynthesis even during cool winter days with frequent showers, provided wind velocity sufficiently compensates for the low vapour pressure deficit during these cool and rainy days.

The same conclusion also applies for other Mediterranean ecosystems. As a consequence the so-called sclerophyllous leaf structure also functions adequately in the fast drying process as a possible adaptation to frequent rainy and windy conditions. The above ericoid leaf growth form exhibits small leaves, which dry quickly. In addition, many species are characterized by hairs on the surface, which will reduce wetting of the leaf during rain, as in many *Erica* species (Schumann *et al.*, 1992), including the European Atlantic *Erica tetralix* L.

A comparison of entire and compound leaves with respect to drying seems appropriate. Wet compound leaves will dry quicker than entire leaves of a similar leaf area as the combined leaflets, because of the spatial distribution of the small and separate leaflets, have a reduced leaf boundary layer thickness. We will give an example, based on the data of K in Figure 13.1. Assume an entire large leaf, corresponding to a rectangular evaporating surface of $W = 10$ cm and $L = 20$ cm and $K = 0.26$. In a narrower leaf of $W = 5$ cm and L remains 20 cm the K value is 0.295 and the drying rate will be enhanced by 14% because of the narrower leaf shape. If the leaf is divided into leaflets of a size comparable with $W = 2$ cm and $L = 5$ cm, the K value is 0.26 and the drying rate is more than double that of the original entire leaf, all under non-limiting energy supply.

There is some experimental evidence that exchange processes, like evaporation, proceed faster in compound leaves. Taylor (1975) reported that the first pinnation in the leaf of *Pteridium aquilinum* (L.) Kuhn, the bracken fern, was the fundamental unit (boundary layer) for heat exchange. The third and second pinnation had a common boundary air layer and heat exchange was only enhanced up to the second pinnation. Unfortunately, no experimental data are given. The large leaves of Musaceae (*Heliconia latispatha* Berth. and *Strelitzia nicotia* Regel and Koern.) commonly show leaf tearing by wind and the boundary layer may be reduced by tearing as evident from leaf temperature measurements: in exposed sites entire leaves often exhibited nearly lethal leaf temperatures, with even a reduced photosynthesis, while plants with damaged leaves were at an advantage in this respect (Taylor and Sexton, 1972).

With respect to drying, a similar comparison can be made between leaves with an entire leaf margin and leaves with a serrate or dentate leaf margin. The aerodynamic properties of the latter category would facilitate the drying process but, at the same time, transpiration of

the leaf would be enhanced under dry weather conditions. In this respect it is of interest that the veins of serrate or dentate leaves in general extend to the leaf margin, while the major veins of leaves with an entire margin in general loop within the margins (Parrish, 1998: 140). As mentioned earlier, veins of leaves of many species may act as gutters for runoff for water during rain and thus facilitate drying afterwards, as in many species of Melastomataceae.

Adaptive changes in leaf morphology in relation to the 'trade-off' between photosynthesis and transpiration

As mentioned earlier, the wet season in Mediterranean ecosystems is characterized by a relatively cool and rainy period, where drying of leaves is determined by a compensatory effect of strong wind to the relatively low water vapour pressure deficit. Palaeoclimate has regularly fluctuated from warmer to cooler conditions and vice versa, including the last 2 million years (Deacon *et al.*, 1992). Changes in climatological conditions in the past, e.g. from a geological period with a high transpiration demand to a cool and rainy period with a low transpiration demand could exert selective pressure on leaf morphology via the drying process: genotypes with entire and simple leaves might possibly be replaced by genotypes with an increased leaf rim as for example in dentate and serrate leaves or lobed leaves, palmately or pinnately. Under further selective pressure, genotypes with compound leaves divided into separate leaflets may have evolved. As mentioned earlier, a reduction of leaf surface is much less effective in drying after rain when the entire leaf shape is maintained than when the reduction in leaf area is realized by development of leaflets of compound leaves. It is also evident, that the selection pressure will only be profound in open vegetations, as occurs after a catastrophic event in which all vegetation has been destroyed.

The effectiveness of selective pressure of climate changes will depend on the number of genes involved in the adaptive change. The genetics of leaf shape seems to be controlled by relatively few genes e.g., *Sambucus nigra* L. with its one time pinnate leaves also has a variety *laciniata* with a delicate parsley-like leaf structure. Variability of leaf shape within a single plant may be large, e.g. in the case of *Acacia heterophylla* Vassal (Ursem, personal communication). Some plant species have two different leaf forms, during the vegetative and generative phase, as in ivy (*Hedera helix* L.), indicating developmental timing in leaf development. A similar case has been described by Stebbins (1974) for the effect of developmental timing on flower petal shape: a large fused and sympetalous flower corolla is formed if the intercalary cells develop along with the petal primordia. If they develop after the petal primordia have grown, the result is well separated petal lobes of the flower. Clearly, modification of leaf and petal shape is, genetically speaking, not complicated.

Closely related species are important since they may exhibit strikingly different leaf forms. An example within a very limited area as the Fynbos, South Africa, is the genus *Pelargonium* and the following species may be mentioned: *P. cuculatum* (L.) L.'Her. with entire leaves, *P. scabrum* (Burm.f) L.'Her. with deeply lobed leaves with coarse hairs, *P. triste* (L.) L.'Her. with a delicate parsley-like leaf form (see Colour Plate 5). A study of the microclimate of sites of these species may reveal whether these different leaf forms exhibit any consistent adaptive value to differences in microclimate.

To take another example, wet, cool and shady habitats with practically no wind exhibit a low transpiration demand. Such a habitat may contain ferns, mosses and liverworts.

Ferns like *Athyrium filix-femina* (L.) Roth and *Woodwardia radicans* (L.) Sw. exhibit compound leaves, sometimes divided two or three times, with only a relatively small reduction in leaf area. This morphology allows a large leaf area for photosynthesis in the shade, together with the advantage of numerous small leaflets, which are adapted to relatively quick drying under these humid conditions. A fern species with entire leaves from a humid and wet habitat such as *Asplenium scolopendrium* L. has shiny leaves, with probably a low wettability. Mosses and foliose liverworts with their minute foliage will also be able to dry quickly under these otherwise unfavourable evaporative conditions.

Finally, evolutionary aspects should be taken into account. As an example for Mediterranean ecosystems, the Fynbos is situated along the rim, where two oceans meet: the Atlantic Ocean with cold water and the Indian Ocean with warm water. This situation causes an extreme average westerly wind speed and together with the large number of climate fluctuations in the past may have led to a species-rich flora (Cowling, 1992), in which the rate of speciation exceeded the rate of extinction. The combination of a continuous strong westerly wind since the breakup of Gondwana, together with regular climatic fluctuations, may also have caused a relatively low degree of variation of leaf shape: the larger the number of repeated climatic fluctuations the lower the number of adaptive traits which will facilitate in survival of the species.

Final remarks

In the above, leaf morphology is related to drying after rain in order: (1) to restore photosynthesis; (2) to minimize nutritional ion leaching; and (3) to reduce infection by fungal leaf pathogens. Physical formulations of evaporation from wet surfaces under energy limiting conditions, as in closed forest canopies, and under less energy limiting conditions, as in open Mediterranean ecosystems, are used in order to evaluate the effect of vegetation structure on leaf morphology. In addition, suggestions are made for prevention of wetting by various adaptive leaf morphology traits.

The presented experimental evidence and observations in the field are, in general, limited to recent publications and the approach is to collect more experimental data in the field to test the applicability of the hypothesis. Within this perspective a number of questions may be asked and the answer to these questions may help to understand leaf morphology within the photosynthesis–transpiration compromise.

The following questions among others may be asked. Is photosynthesis reduced in rain-wetted leaves of varying morphology? How thick is the water layer of wet leaves of different sizes? Which species have a low/high wettability of leaves? Does photosynthesis continue during rain in velvet and tomentose leaves? Can the simulation of rainfall interception and evaporation of Klaassen (2001) be used for Fynbos vegetation and other Mediterranean ecosystems?

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Xylem hydraulics and angiosperm success: a test using broad-leafed species

Norman W Pammenter, Guy F Midgley and William J Bond

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Introduction

Angiosperms are both more numerous in species and more diverse in form than other seed plants. They also dominate low latitude and low altitude biomes of the world in terms of biomass. Angiosperms began to diversify only in the Cretaceous when they also began to infiltrate, and eventually displace, vegetation dominated by *Ginkgo* L., conifers, cycads, Gnetales and extinct gymnosperm groups (Crane *et al.*, 1995; Wing and Boucher, 1998). Over the last ten years, there has been considerable progress in identifying attributes of early angiosperms from remarkable fossil discoveries and in identifying sister taxa from molecular and morphological phylogenetic analyses (Crane *et al.*, 1995; Crepet, 2000; Friedman and Floyd, 2001). There has also been a growing number of studies on angiosperm traits that might account for their ecological 'success'. Diversification of angiosperms has been linked to co-evolution with insect pollinators in many angiosperm groups (Grimaldi, 1999; Crepet, 2000) and to the evolution of growth forms with short generation times (Eriksson and Bremer, 1992; Tiffney and Mazer, 1995; Dodd *et al.*, 1999). However, diversity does not necessarily translate to ecological dominance, as measured by the area occupied or the proportion of biomass present (Bond, 1989). It is this aspect of angiosperm success that we consider here.

Extant gymnosperms are all woody and most have long-lived leaves with low surface area to weight ratios. Many angiosperms resemble gymnosperms, with long-lived evergreen

leaves similar to extant conifers (although few have needle-like leaves) and often grow together with them. However, angiosperms also include trees, shrubs and herbs with leaves of a short life span and high surface area to weight ratios (Reich 1998), unlike any extant (or extinct) gymnosperms. Plants with these leaf properties dominate ecosystems where gymnosperms are rare or absent, including temperate deciduous forests, lowland tropical forests (which, although evergreen, are ever-growing), savannas, and herb-dominated grasslands and ruderal communities. Bond (1989) suggested that the presence of vessels and of distinctive, anastomosing leaf venation in angiosperms, coupled with a reproductive system that allowed them rapidly to produce seeds, permitted flowering plants to develop novel growth forms that could out-grow gymnosperms. He suggested that these features might be most important in the recruitment stage, before gymnosperms could build up a large leaf mass by accumulating successive cohorts of evergreen leaves. Rapid growth and rapid reproduction would be most favoured in productive environments and it is these that are now dominated by flowering plants. Angiosperms would have little advantage over gymnosperms in environments with chronic stress such as cold, or nutrient poor areas and here gymnosperms would still be able to persist.

There are a number of recent studies of hydraulic properties and leaf physiology of angiosperms and gymnosperms (Becker *et al.*, 1999; Brodribb and Feild, 2000) and of angiosperms which lack vessels (Feild *et al.*, 1998, 2000; Feild and Holbrook, 2000). Becker (2000) has reviewed such studies in the context of Bond's (1989) hypothesis, that differences in the hydraulic system partly account for angiosperm 'success'. The general conclusion is that angiosperms can attain higher conductance and photosynthetic capacity than gymnosperms but that values measured for many angiosperms overlap those obtained for gymnosperms. Becker (2000) argues that the results do not support the contention that angiosperm innovations in the hydraulic system influence leaf physiology and competitive performance. However, most of the studies compared angiosperm and gymnosperm trees growing together in low productivity environments including cloud forests, nutrient poor soils and other sites with chronic stress, where the hydraulic characteristics of angiosperms would have little advantage.

A more appropriate test of the importance of hydraulic design in limiting gymnosperm success, would be to compare relative performance of species with similar growth forms in productive environments, from which gymnosperms have been excluded. The question is whether gymnosperm-like vascular systems can support fast-growing angiosperm-like canopies. Could a gymnosperm-type hydraulic system, made up of tracheids, supply water fast enough to sustain uniquely angiosperm-type leaves with short life spans, a high surface area to mass ratio and the capacity to rapidly grow and fill space? Could these uniquely angiosperm leaf properties function with gymnosperm-like leaf vasculature (Roth-Nebelsick *et al.*, 2001)? Because we do not have the technology to design plants to specification, with mixed angiosperm and gymnosperm elements, we explored the first question, of hydraulic limitations on leaf function, by searching for gymnosperms with leaves most similar to those of angiosperms growing in productive environments. There are very few of them: most extant conifers have needle-like or scale leaves. *Ginkgo* is the only extant gymnosperm which grows a broad-leaf and discards it in a single season. We compared this species with broad-leaved deciduous angiosperm trees of genera that occur in temperate deciduous forests. Although the earliest angiosperms were probably evergreen, we chose deciduous rather than evergreen angiosperms for three reasons: (1) the hypothesis concerns the dominance, not the origin of angiosperms, and deciduous broad-leaved ginkgo-phytes were displaced by deciduous angiosperms from the Cretaceous (Upchurch *et al.*,

1998); (2) we wished to ensure that the study species had short-lived leaves (leaf life span of evergreen plants is extremely variable among species and growth conditions); and (3) the angiosperm and 'model' gymnosperm species to be compared would have the same growth form. To extend the database for the gymnosperms, we also included data from two further species, *Agathis* Salisb. and *Podocarpus* L' Hér. ex Pers. which, although evergreen, are relatively broad-leaved. All the trees were growing in resource-rich productive sites. We wished to determine whether: (1) the hydraulic properties of these broad-leaved gymnosperms differ from those of angiosperms with a similar tree growth form; (2) whether any such differences are correlated with leaf physiology; and (3) whether differences in leaf physiology could potentially influence competitive ability.

Materials and methods

Choice of species

To assess the influence of anatomy or morphology on physiological characteristics, the plants chosen should ideally be growing together, so as to avoid potentially confounding effects of different growth conditions. This requirement for species of similar habit occurring in the same habitat imposed constraints on the species available for study. However, in a municipal park in Cape Town, South Africa, some suitable species have been planted. In addition to *Ginkgo biloba* L., the following broad-leaved deciduous angiosperm species were studied: *Quercus robur* L. (English oak), *Quercus palustris* Münchh. (southern pin oak), *Ulmus procera* Salisb. (English elm), and the London plane (*Platanus* × *acerifolia* (Aiton) Willd., a hybrid between *Platanus orientalis* L. and *P. occidentalis* L., sometimes also referred to as *Platanus* × *hispanica* (Pakenham, 1996)). The park where the plants were growing was irrigated during the dry summer months, when the measurements were taken. To broaden the base of the gymnosperm data set, studies were undertaken on *Agathis robusta* (Moore) Bailey: although not deciduous this species has relatively broad leaves in comparison with most gymnosperms. Additionally, some data collected for a different study on the gymnosperm *Podocarpus latifolius* (Thunb.) R. Br. ex Mirb. were included in the data set.

The constraints imposed on the choice of species create statistical difficulties in that truly independent samples were not available for all species. Studies were conducted on four individual young trees of *G. biloba*, three mature *Q. robur* trees, two young specimens of *Q. palustris* and a single young but mature tree of each of *U. procera* and *P. × acerifolia*. A single specimen of *A. robusta* growing as a street tree near the park in Cape Town was used for the water relations studies and a young specimen in the Durban Botanic Gardens was used to investigate photosynthesis characteristics. The specimen of *P. latifolius* was growing in an irrigated domestic garden in Durban. In all cases, measurements were made on several (up to five) individual leaves or twigs from the trees available. Where appropriate, analyses of variance were conducted and data that were not normally distributed were log-transformed prior to analysis.

Physiological studies

Water relations

Hydraulic conductivities of excised young branches were measured using the procedure of Sperry *et al.* (1987). Briefly, conductivity was calculated by measuring the mass flow of water passing through a branch segment connected to a constant pressure reservoir (2 kPa).

It was ensured that all branch segments were longer than the longest xylem vessel. Hydraulic conductivity was measured after flushing the branch segment at a pressure of approximately 200 kPa until no further increase in flow rate occurred and values reported here are maximum values. A 0.01 M HCl solution (with filtered (0.22 μm membrane filter), degassed, distilled water) was used throughout. After measurement, safranin dye (0.05%) was passed into the stem to stain the functional xylem, the area of which was measured and the data were expressed as specific conductivity (k_s). Leaf area distal to the measured stem was measured and data also expressed as leaf specific conductivity (k_l).

Stomatal conductances and transpiration rates of leaves in their natural orientation were measured using a LiCor 1600 porometer (LiCor, Lincoln, Nebraska) and light intensity incident on the leaf surface noted. The water potential of each leaf on which transpiration rate had been measured was determined immediately afterwards by means of a pressure chamber (Soil Moisture Corporation, Santa Barbara, California). Measurements were taken from pre-dawn to shortly after midday to establish the relationship between transpiration rate and leaf water potential (measurements were not taken in the afternoon to avoid hysteresis effects).

Photosynthesis studies

Rates of net carbon dioxide assimilation (A) were measured using a LiCor 6200 portable photosynthesis system with a 1 l chamber volume. Leaves were maintained in their natural orientation during measurements and light intensity incident on the surface was measured. Data were collected from pre-dawn to shortly after midday and measured rates were plotted against incident photosynthetic photon flux density (PPFD), so that photosynthesis rate at saturating light intensity (A_{max}) could be assessed. We used a number of leaves to construct a single 'light curve' and replicates consisted of measurements on different days.

The response of rates of assimilation to intercellular CO_2 concentration (C_i) were measured using the same equipment (except for *A. robusta* and *P. latifolius*). A fully illuminated leaf was enclosed in the chamber at normal atmospheric CO_2 concentration and photosynthetic rates measured as the CO_2 concentration in the chamber was drawn down by the photosynthesis of the leaf. Once the compensation point had been approached the CO_2 concentration in the chamber was increased by the injection of a small quantity of pure CO_2 , and photosynthetic rates measured as CO_2 concentration was drawn down to ambient levels. Data sets in which there were differences between the two measures of assimilation rate at ambient CO_2 concentration (at the start and end of the experiment) were rejected. Relative humidity in the chamber was maintained close to ambient levels by controlling the proportion of the airflow passing through a desiccant. Leaf temperatures increased during the course of measurements, but did not exceed air temperature by more than 5°C. $A:C_i$ curves and light response curves of *A. robusta* and *P. latifolius* were measured using a LiCor 6400 portable photosynthesis measuring system. The initial slope of the response of A to C_i was taken as a measure of the carboxylation coefficient, which has been related to the activity of ribulosebiphosphate (RuBP) carboxylase (Farquhar and Sharkey, 1982). The saturation value (J_{max}) indicates the maximum rate of RuBP regeneration and thus electron flow through the photosynthetic electron transport pathway (Farquhar and Sharkey, 1982). To have sufficient data points for calculation and analysis of initial slopes, the data from individual $A:C_i$ curves were combined.

Stomatal limitation to photosynthesis was assessed according to Farquhar and Sharkey (1982). Instead of calculating a stomatal limitation for each individual $A:C_i$ curve for each species, a single value was estimated by combining all the curves for a species and using

the mean value of stomatal conductances at PPFD values greater than $200 \mu\text{mol m}^{-2} \text{s}^{-1}$ (the light intensity by which stomatal conductance had saturated). Although this means that replicate values were not available, it was felt to be more representative of the leaves in their natural state, as stomatal conductances of leaves in these conditions, rather than in a chamber, were used.

Results

The maximum hydraulic conductivity of minor branches expressed per unit area of functional xylem is shown in Figure 14.1. The values separated into two statistically distinct groups, with the angiosperms having specific conductivities an order of magnitude higher than those of the gymnosperms. In terms of water supply to leaves, inefficient xylem could be compensated for by decreases in the leaf area supplied per unit area of xylem. In the species studied, although the gymnosperms did have lower leaf area/xylem area ratios, the differences were not large (data not shown). Consequently, when hydraulic conductivity was expressed per unit leaf area supplied, the differences apparent in specific conductivity were maintained, with the angiosperms having leaf specific conductivities five to ten times higher than those of the gymnosperms (Figure 14.2).

The tension developed in the transpiration stream is directly related to the transpiration rate and inversely proportional to the leaf specific conductivity (Tyree and Ewers, 1991). Thus a low leaf specific conductivity would imply either, or both, low transpiration rates and low leaf water potential. The relationship between transpiration rate and the leaf water potential developed support this (Figure 14.3; the lines for individual species are extended to the maximum transpiration rates measured). The slopes of this relationship for the species studied fall into two statistically distinct groups, with the gymnosperms showing the steeper slopes. Consequently, the gymnosperm species had midday transpiration rates

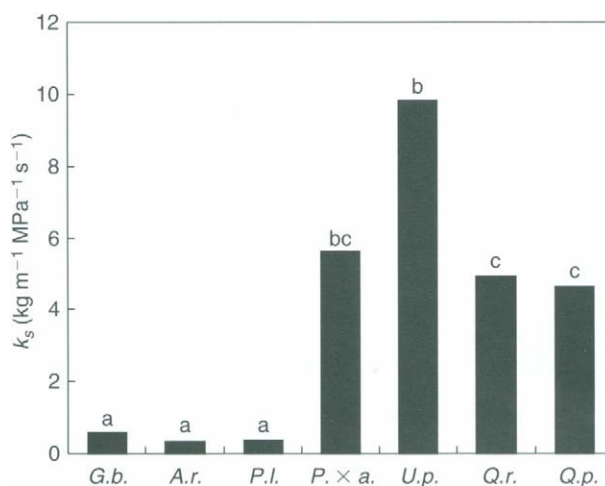


Figure 14.1 Maximum hydraulic conductivity per unit xylem area (k_s) of three gymnosperm and four angiosperm species. Abbreviations: *G.b.*, *Ginkgo biloba*; *A.r.*, *Agathis robusta*; *P.l.*, *Podocarpus latifolius*; *P. × a.*, *Platanus × acerifolia*; *U.p.*, *Ulmus procera*; *Q.r.*, *Quercus robur*; *Q.p.*, *Quercus palustris*. The same abbreviations are used in all the figures. Values with the same letter are not statistically different (analysis of variance followed by Tukey's multiple range test, $P < 0.05$).

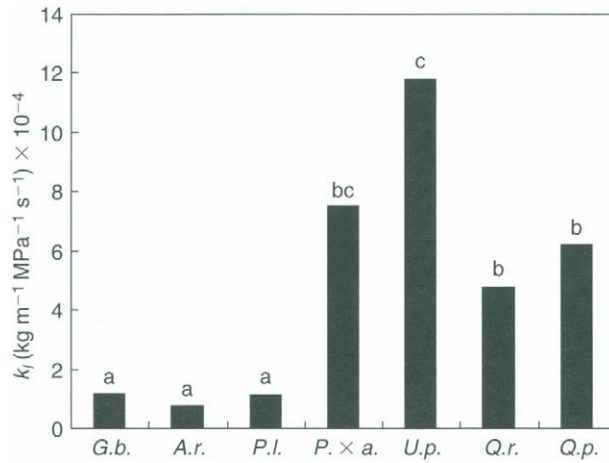


Figure 14.2 Maximum hydraulic conductivity per unit leaf area supplied (k_l). Abbreviations as in Figure 14.1. Values with the same letter are not statistically different (analysis of variance followed by Tukey's multiple range test, $P < 0.05$).

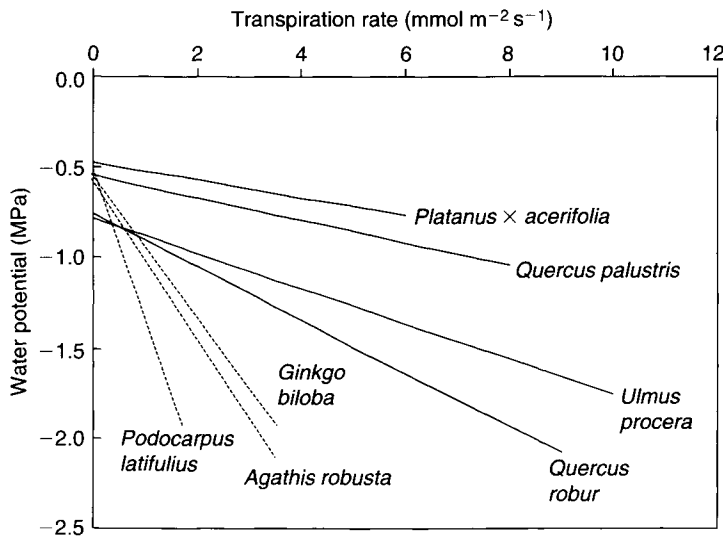


Figure 14.3 Relationship between transpiration rate and leaf water potential developed. For clarity only the fitted linear regressions, rather than all individual data points, are shown. The lines have been extended to the maximum transpiration rates measured. The solid lines and the dotted lines constitute two statistically distinct groups of regressions (T' -method for unplanned comparisons among a set of regression coefficients at 95% confidence level; Sokal and Rohlf, 1981).

considerably less than those of the angiosperms. The leaf water potentials developed by the gymnosperms tended to be lower than those experienced by the angiosperms, although there was some overlap.

The fact that the gymnosperms exhibited lower transpiration rates than the angiosperms, under the same soil water supply and atmospheric vapour pressure deficit conditions,

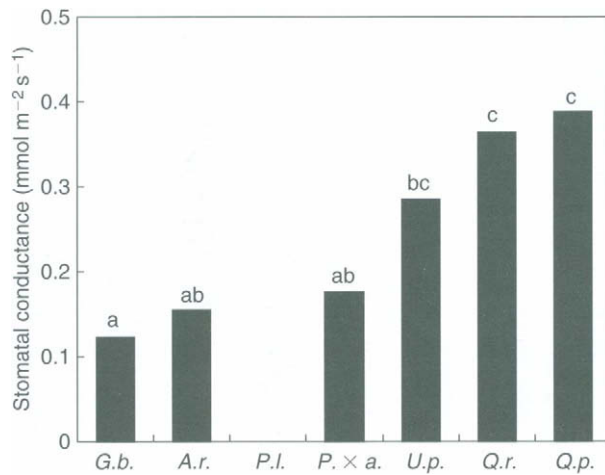


Figure 14.4 Average stomatal conductances measured in the field at incident photosynthetically active radiation intensities greater than $200 \mu\text{mol photons m}^{-2} \text{s}^{-1}$. Abbreviations as in Figure 14.1. Values with the same letter are not statistically different (analysis of variance followed by Tukey's multiple range test, $P < 0.05$).

suggests that the stomatal conductances to vapour diffusion were lower in the gymnosperms. Stomatal conductance is dependent upon incident light intensity, but in the species studied the effect was generally saturated by a PPFD of $200 \mu\text{mol m}^{-2} \text{s}^{-1}$. The mean stomatal conductance measured for each species at incident radiation higher than this is shown in Figure 14.4 (as the data for *P. latifolius* were collected for a different study, stomatal conductance values are not available). Although the differences were not as marked as reported for the hydraulic conductivities, there is a tendency for the gymnosperms to have lower stomatal conductances than the angiosperms, with *P. × acerifolia* showing intermediate values closer to the gymnosperms than the other angiosperms.

Lower stomatal conductances could possibly limit CO_2 diffusion into the leaf and hence reduce maximum photosynthetic rates. Light-saturated rates of photosynthesis (A_{max} , PPFD $> 1000 \mu\text{mol m}^{-2} \text{s}^{-1}$) fell into distinct groups (Figure 14.5), although *P. × acerifolia* again grouped with the gymnosperms, not the angiosperms. To assess whether the lower maximum photosynthetic rates were a consequence of stomata-imposed limitations on rates of CO_2 diffusion, stomatal limitations were calculated (Figure 14.6). Although the method of calculation did not permit statistical analysis of the data, it is clear that there are no major differences in stomatal limitation between the two groups of plants. To assess whether photosynthetic capacity, rather than stomata, limited light-saturated photosynthetic rates, the carboxylation coefficients (Figure 14.7) and maximum rates of RuBP regeneration (Figure 14.8) were measured from $A:C_i$ curves. Differences in carboxylation coefficients were not clear, although there was a tendency for the species with high rates of light-saturated photosynthesis to have high carboxylation coefficients (Figure 14.7). In terms of maximum rates of RuBP regeneration, although *P. × acerifolia* had the lowest value among the angiosperms, the gymnosperms constituted a single group with values lower than any of the angiosperms (Figure 14.8). Thus the low maximum photosynthetic rates of the gymnosperms is a consequence of low photosynthetic capacity, rather than directly due to stomatal limitation to CO_2 diffusion. In terms of photosynthetic characteristics, *P. × acerifolia* is more similar to the gymnosperms, than to the angiosperms.

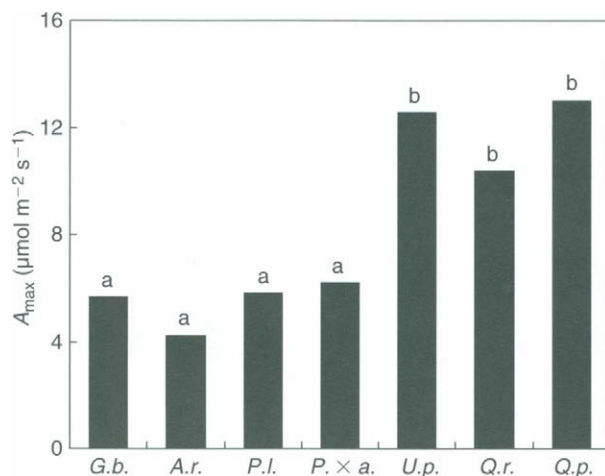


Figure 14.5 Average rates of photosynthesis measured in the field at incident photosynthetically active radiation intensities greater than $1000 \mu\text{mol photons m}^{-2} \text{s}^{-1}$ (A_{\max}). Abbreviations as in Figure 14.1. Values with the same letter are not statistically different (analysis of variance followed by Tukey's multiple range test, $P < 0.05$).

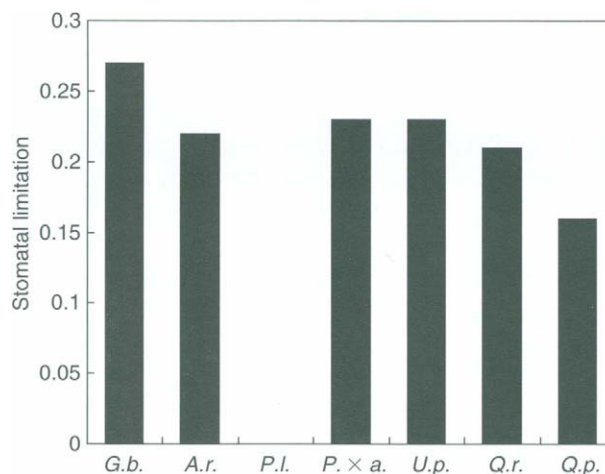


Figure 14.6 Stomatal limitation to photosynthesis, assessed from bulked replicate $A:C_i$ curves and average stomatal conductance measured in the field at incident photosynthetically active radiation intensities greater than $200 \mu\text{mol photons m}^{-2} \text{s}^{-1}$. Abbreviations as in Figure 14.1.

Discussion

The essence of the argument presented here concerning the current dominance of the angiosperms is that the more efficient hydraulic characteristics conferred by the vessels of the angiosperms could provide this group with a competitive advantage *under conditions of high potential productivity*. Thus, when subjecting this hypothesis to test, the choice of species is critical. Additionally, although there is considerable overlap in the hydraulic properties of gymnosperms and angiosperms (Tyree and Ewers, 1996; Becker, 2000), hydraulic

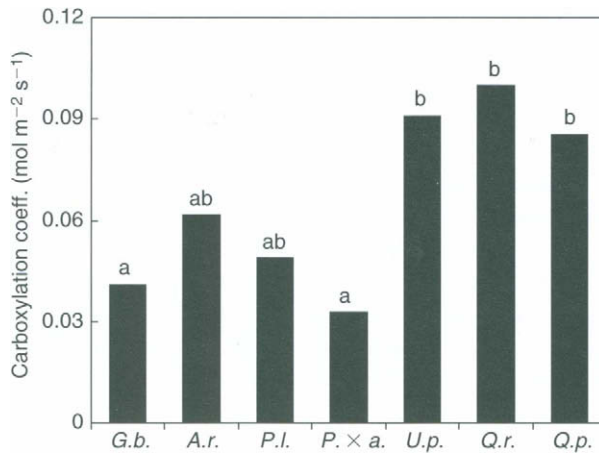


Figure 14.7 Carboxylation coefficient measured as the slope of bulked $A:C_i$ curves. Abbreviations as in Figure 14.1. Values with the same letter are not statistically different (T'-method for unplanned comparisons among a set of regression coefficients at 95% confidence level).

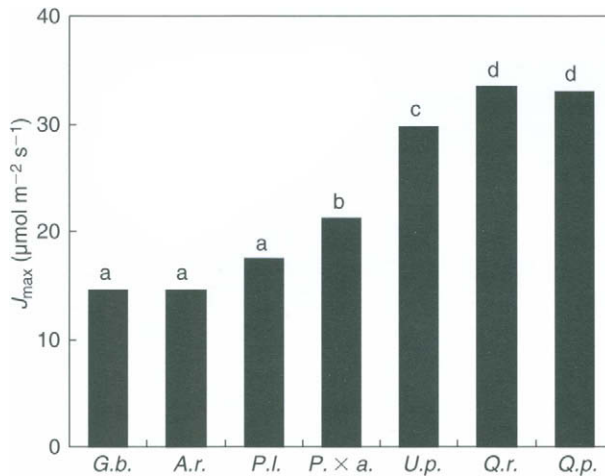


Figure 14.8 Maximum rates of RuBP regeneration (J_{max}) measured from $A:C_i$ curves. Abbreviations as in Figure 14.1. Values with the same letter are not statistically different (analysis of variance followed by Tukey's multiple range test, $P < 0.05$).

architecture and conductivities may respond to growth conditions (Shumway *et al.*, 1993; Mencuccini and Grace, 1995; Heath *et al.*, 1997; Vander Willigen and Pammenter, 1998; Brodribb and Feild, 2000). This reinforces the need to select, not only the test species, but also the individual plants, with care. Becker *et al.* (1999) showed similar whole plant hydraulic conductances of sapling-sized angiosperms and gymnosperms, but these were growing on nitrogen- and phosphorus-poor white sands in Borneo. Similarly, Feild and Holbrook (2000) showed no difference in leaf specific conductivities between the vessel-less *Drimys granadensis* L. (Winteraceae) and co-occurring vessel-bearing angiosperms in a cloud forest of Costa Rica. However, the conductivities measured were at the low end of the range for angiosperms and were, in fact, similar to the values we measured for the

gymnosperms in our study. Thus we chose angiosperms with broad, short-lived leaves from productive habitats, the species being selected specifically because they were growing together with a gymnosperm of similar growth habit. Two additional relatively broad-leaved gymno-sperm species were also included; the specimens of both species were growing in productive environments.

There were clear differences between the angiosperms and gymnosperms in our study in terms of hydraulic conductivities of twigs, expressed on either xylem area (see Figure 14.1) or leaf area (see Figure 14.2). Although the roots may constitute the main resistance to water flow, particularly in small plants (Tyree and Ewers, 1991), these differences in minor branches in our study translated into differences at the leaf level of intact plants, in terms of rates of transpiration and leaf water potentials developed (see Figure 14.3). Becker *et al.* (1999) have pointed out the importance of measuring whole plant conductance when assessing the significance of hydraulic characteristics. We were unable to do this directly because it is difficult to measure root conductance using a high pressure flow meter (Tyree *et al.*, 1995) on plants with basal diameters in excess of a few centimetres (over-and-above problems of getting permission to fell trees in a park!). Meinzer *et al.* (1999) have estimated whole plant conductance from stem sap flow measurements and leaf water potentials: we were unable to do this because of the range in stem size we were dealing with. However, the slope of the relationship between leaf transpiration rate and leaf water potential frequently has been taken as a measure of whole plant conductance (see Table 4 in Becker *et al.* (1999) for references). This measure indicated significantly lower whole plant conductances of the gymnosperms when compared with the angiosperms in our study (see Figure 14.3).

The lower hydraulic conductances and lower transpiration rates of the gymnosperms were associated with lower stomatal conductances to diffusion (see Figure 14.4). It has been suggested that a function of stomata is to maintain transpiration rates at levels preventing excessive tensions in the xylem water columns, so preventing runaway embolism cycles (Tyree and Sperry, 1989) and hydraulic constraints on stomatal conductance and transpiration have been reported (Whitehead, 1998; Kolb and Sperry, 1999; Meinzer *et al.*, 1999; Clearwater and Meinzer, 2001). The clear exception here is the low stomatal conductance of *P. × acerifolia*, which is obviously not a consequence of low plant hydraulic conductance. The reason for this is unknown: it should not be related to the diffuse-porous nature of the wood of this species (the other angiosperm species are ring-porous), because hydraulic conductivities were measured directly; additionally, the existence of scalariform perforation plates would suggest that the species has low vulnerability to cavitation and so transpiration rates would not have to be kept low to maintain low xylem tensions, as would be the case if vulnerability was high.

Low stomatal conductances could possibly limit photosynthesis by restricting CO₂ diffusion into the leaf intercellular air spaces (Farquhar and Sharkey, 1982) and light-saturated rates of photosynthesis of the gymnosperms were indeed lower than those of the angiosperms (except *P. × acerifolia*; see Figure 14.5) and there was a significant linear relationship between stomatal conductance and A_{\max} ($R^2 = 0.81$, $P < 0.02$). However, these lower photosynthetic rates in the gymnosperms do not appear to be a direct consequence of stomatal limitations as there were no clear differences between the gymnosperms and angiosperms in stomatal limitations to carbon assimilation rate (as assessed from A:C_i curves and average light-saturated value of stomatal conductances (see Figure 14.6)). Rather, the differences in photosynthetic rates between the groups seem a consequence simply of higher photosynthetic capacity in the angiosperms (i.e. higher carboxylation coefficients and RuBP regeneration rates, see Figures 14.7 and 14.8).

It is now well known that the relationship at leaf level between maximum stomatal conductance and maximum assimilation rate is strongly conserved across many plant species and functional types (Schulze *et al.*, 1994), as is the relationship between carboxylation capacity and RuBP regeneration rate (Wullschlegel, 1993). Therefore, a functional limitation on stomatal conductance imposed by hydraulic constraints should translate to lower leaf level photosynthetic capacity. Our data indeed show a significant positive relationship between A_{\max} and k_l ($R^2 = 0.59$, $P < 0.05$). This result echoes that of Brodribb and Feild (2000), who showed a significant positive relationship between k_l and photosynthetic capacity (assessed from electron flux through PSII) in a wide range of angiosperm and gymnosperm trees. In that study, although there were some overlaps, the gymnosperms and vessel-less angiosperms had significantly lower hydraulic conductivities and photosynthetic capacities (from which was inferred lower stomatal conductances) than the vessel-bearing angiosperms.

Thus, in our study the hydraulic conductivities of broad-leaved deciduous angiosperms were higher than those of broad-leaved gymnosperms growing in the same or similar environments and these differences in hydraulic characteristics were associated with differences in leaf physiology, with the angiosperms having higher photosynthetic capacities and light-saturated photosynthetic rates (with the exception of *P. × acerifolia*). Growth rates, of course, depend upon more than photosynthesis rates per unit leaf area, with characteristics such as partitioning of photosynthate (Körner, 1991) and respiration rates and relative sink strengths (Farrar, 1999) being important. It is thus unlikely that differences in leaf physiology are the sole factor underlying the dominance of woody angiosperms over gymnosperms in these environments. However, among species of similar growth habit (woody, broad-leaved, short leaf life span) and habitat (productive environments), a higher photosynthetic capacity could translate into potentially higher growth rates and hence competitive ability.

We speculate that hydraulic limitations are likely to emerge as constraints on plant growth and there is evidence of a general trend of high whole plant hydraulic conductivities being associated with fast growth rates of pioneer species (Tyree and Ewers, 1996). In addition to any affect at the level of leaf physiology, hydraulic limitations are likely to constrain canopy development rates as well. Because of the lower stem hydraulic conductivity per unit xylem area of the gymnosperms, to maintain high capacity for water delivery to leaves with high transpiration rates (and thus photosynthetic rates), more wood has to be allocated to support unit leaf area. (This ratio, like other aspects of hydraulic architecture, is influenced by locality and growth conditions, and so comparisons should be made among similar growth forms and habitat type.) Allocation of more carbon to non-photosynthetic conducting tissue is likely to reduce total photosynthate assimilation and so potential growth rate. To develop both the physiological and allometric arguments fully, information on maximum individual growth rate, maximum canopy leaf area index and canopy development rate are needed, not only at the seedling level (the 'slow seedling' hypothesis of Bond (1989)), but also at the sapling stage. It is well known, for example, that pine (gymnosperm) plantation forests are highly productive, but the key to this apparent paradox is that they require time to develop the leaf area indexes and canopies to achieve high production rates once established.

An additional possible factor in the competitive potential of angiosperms is that our casual (unrecorded) observations suggest that the growth form of angiosperm trees is more plastic than that of gymnosperms. This plasticity would be an advantage in exploiting stochastically available resources (such as gaps in a forest canopy), leading to a competitive edge. However, high growth rates are a prerequisite for plants to be able to take advantage of this plasticity.

In summary, our data set pertaining to angiosperms and gymnosperms species of similar growth habit in similar habitats are consistent with the concept of a hydraulic limitation to the growth rate of gymnosperms and hence their relative lack of success in productive habitats in the face of competition from 'better-plumbed' angiosperms. The ultimate test of manipulating the hydraulic conductivities of the test species and observing the effects on leaf physiology and growth rates awaits to be done.

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15

Evolution of xylem physiology

Pieter Baas, Frank W Ewers, Stephen D Davis and
Elisabeth A Wheeler

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Introduction

This chapter summarizes how, according to our present understanding, certain wood anatomical traits evolved in woody angiosperms and how during the geological past clear ecological preferences can be recognized for a number of arbitrarily defined wood functional types. That summary will be related to experimental evidence of how key wood anatomical traits affect hydraulic (and to a lesser extent mechanical) functioning of woody trees, shrubs and vines. In conclusion, the ecological patterns in xylem anatomy will be discussed in terms of their functional significance.

‘Trade-off’ triangle

Xylem evolution can be viewed in the context of a ‘trade-off’ triangle (Figure 15.1), with different adaptive solutions to the structure/function problems depending on environmental demands as well as phylogenetic constraints (taxonomic history). Evolution of xylem physiology is complicated by the fact that, ever since the early evolution of land plants, xylem has simultaneously performed multiple functions. These include not only water transport but also mechanical support of the plant and storage of water, minerals and carbohydrates. The optimal structures for each of the xylem functions most likely differ and selection to optimize for one function could lead to suboptimal performance or even complete failure of another function. An extreme example would be wood maximized for efficient water conduction with many wide thin-walled conduits, but which is so mechanically weak that stem failure occurs prior to seed production.

Herein, conductive efficiency of xylem is expressed in terms of specific conductivity (k_s in $\text{m}^2\text{MPa}^{-1}\text{s}^{-1}$). Specific conductivity is the hydraulic conductivity divided by the xylem transverse area, with hydraulic conductivity equal to the flow rate per pressure gradient (Wagner *et al.*, 1998). Resistance to embolism (air blockage) is defined as the water potential (in MPa) at which there is 50% loss in conductivity due to embolism (Tyree *et al.*, 1994). We define mechanical strength both in terms of stem strength and vessel wall strength. For stem strength we use modulus of rupture (MOR in GN m^{-2}) of fresh stems, as measured with a cantilever test. This test is relevant since breakage of stems in nature would impair reproductive output by a plant (Wagner *et al.*, 1998). For vessel strength we use $(t/b)^2$, which is the square of vessel wall thickness (t) divided by the diameter of the conduit (b). This parameter is relevant to determining the resistance to vessel implosion (Hacke *et al.*, 2001), which is a danger to plants since the xylem sap is transported under negative pressure (= tension).

Early beginnings

The first woody plants had vessel-less wood and the first forests were comprised of trees with vessel-less woods (Meyer-Berthaud *et al.*, 1999). Vessels arose independently in

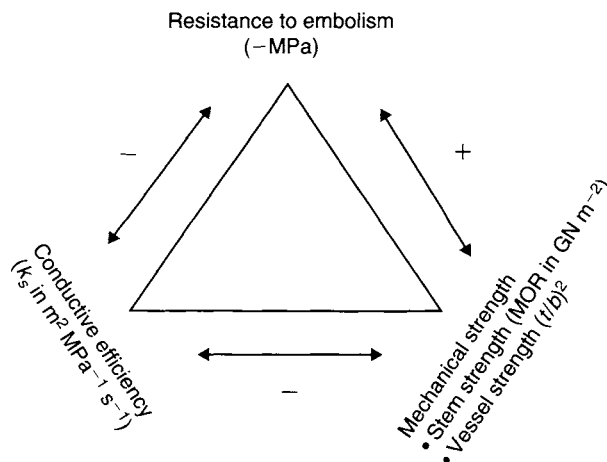


Figure 15.1 Xylem evolution can be viewed in the context of a ‘trade-off’ triangle, where conductive efficiency is inversely proportional to both mechanical strength and resistance to dysfunction via embolism.

several clades (e.g. ferns, horse tails, *Selaginella* Pal., Gnetales, angiosperms). Recent molecular phylogenies indicate angiosperms were primitively vessel-less (Soltis *et al.*, 2000). The vessel-less *Amborella* Baillon, a monotypic genus of evergreen shrubs from New Caledonia, is basal to all angiosperms. Thus, presumably, vessel-less angiosperms gave rise to vessel-bearing ones. Bailey and Tupper (1918) proposed a transformation series for this evolutionary event. It is debated whether vessel-lessness in Winteraceae and *Tetracentron* Oliver, which are nested in more advanced angiosperm clades, represents a reversion or whether these woods are primitively vessel-less with several independent, unparsimonious origins of vessels in the lower clades (Baas and Wheeler, 1996; Herendeen *et al.*, 1999; Doyle and Endress, 2000). Comparison of the most successful of modern vessel-less woody plants, the conifers or softwoods, with the most successful modern vessel-bearing plants, the dicotyledons or hardwoods, shows two alternative strategies.

Conifers have longitudinal tracheids interconnected by large bordered intertracheary pits whose pit membranes (modified primary walls) usually are differentiated into a porous margo and a central thickened non-porous torus. The porous margo allows for effective sap transport. The non-porous torus provides safety against embolisms spreading from one tracheid to another. If an embolism occurs, the pit membrane, if flexible, deflects so that the torus presses against the pit border and seals the opening (aperture) between adjacent cells. This process effectively limits the spread of the embolism, as long as pressure differences across the pit aperture do not exceed a threshold that disrupts the seal (Sperry and Tyree, 1990). Individual longitudinal tracheids are approximately 100 times longer (2–4 mm) than wide (0.02–0.04 mm). Vessel-less angiosperms also rely on long tracheids, but their pit membranes are much less porous than conifers and the membranes generally lack tori.

Angiosperms have vessels for xylem sap transport and fibres for support. Vessels are comprised of a series of vessel elements with perforated common end walls. Vessels, while considerably longer than longitudinal tracheids, are finite in length, varying from a few millimetres to several metres long (Zimmermann, 1983). Vessel diameter ranges from 0.025–0.3 mm, most commonly more than 0.05 mm. Intact sapwood intervessel pits have relatively compact, seemingly non-porous pit membranes which would not readily allow movement of embolisms from vessel to vessel.

Distribution of wood anatomical features

Vessel element perforations

According to the transformation series of Bailey and Tupper (1918), in angiosperms, vessel elements with scalariform perforation plates and scalariform pitting were derived from imperforate elements with scalariform pitting. Subsequently, vessel elements with simple perforation plates were derived from vessel elements with scalariform perforations. Support from the fossil record for this transformation series includes: (1) the angiosperms as a whole, a higher incidence of scalariform perforations in the Cretaceous than in Tertiary (Figure 15.2) (Wheeler and Baas, 1991, 1993); (2) the Platanaceae, Cretaceous and Early Tertiary species have exclusively scalariform perforation plates, while Late Tertiary and modern species have a mixture of simple and scalariform perforations (Wheeler, 1991). However, the fossil record also shows ambiguities with respect to the Bailey transformation series: *Paraphyllanthoxylon* is one of the two oldest types of angiosperm and it has vessel elements with simple perforations (about 100 million years

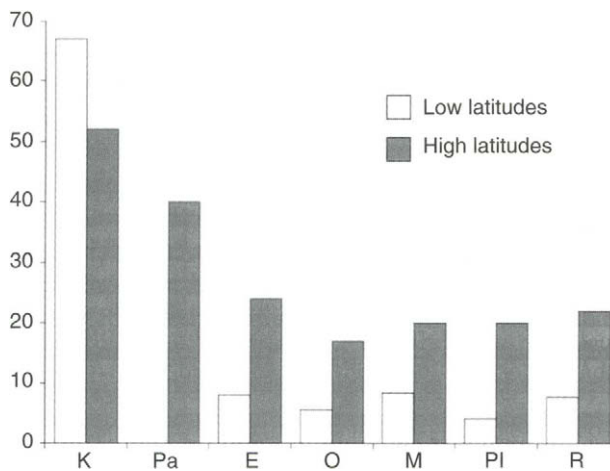


Figure 15.2 Incidence of scalariform perforations (%) through time, percentage of number of reports per time unit for Gondwana (primarily low latitudes) and Laurasia (primarily high latitudes). K = Cretaceous (from Aptian/Albian onward, ca. 100–65 Ma); Tertiary records reported for epoch: Pa = Palaeocene, E = Eocene; O = Oligocene; M = Miocene; Pl = Pliocene, R = Recent. Data from Wheeler *et al.* (1986) and Wheeler and Baas (1991) plus updates.

before present, Ma) (TX, USA, Serlin, 1982; Utah, Thayne *et al.*, 1983). *Paraphyllanthoxylon*-type woods occur throughout the Cretaceous, are abundant at some localities and some were more than 1 m in diameter (e.g. Mogollon Rim, AZ, USA, E.A. Wheeler and J.A. Wolfe, personal observation; AL, USA, Cahoon, 1972). Vessel-less angiosperm woods are not known until approximately 20 million years later (Antarctica, Poole and Francis, 2000). There is a need to examine additional collections of Aptian-Albian woods as well as later stages of Cretaceous, especially from low latitudes, to elucidate the wood anatomical characteristics of early angiosperms and changes in wood structure throughout the Cretaceous. Examination of charcoalfied woods with well-preserved detail and potentially representing plants of small stature should be particularly useful (e.g. Herendeen, 1991a,b).

Modern ecological trends show scalariform perforations have been preferentially retained in cool mesic conditions and to be rare in warmer more xeric climates and among lowland tropical trees (Baas, 1982, 1986). That trend can be traced in the fossil record as the differences in incidences of scalariform perforation plates (see Figure 15.2) between temperate-zone Europe, North America and Asia and tropical Africa, Malesia and Central and South America are similar from the Eocene onwards. After the globally warm early Eocene interval, the middle Eocene marks the beginning of modernization of climates in the northern hemisphere (Graham, 1999).

A simple functional explanation for the decline in the incidence of scalariform perforation plates is that the bars offer some resistance to flow and, in hot and seasonally dry climates, simple perforations would be advantageous as they would have low resistance to flow. Recent experimental work has shown that the additional resistance to water flow offered by perforation plates is least for simple perforation plates (from 1.7 to 5.1% of the total resistance; Ellerby and Ennos, 1998), greater for scalariform plates with about 4 bars (about 8% in *Liriodendron tulipifera* L.; Schulte and Castle, 1993) and still greater in a species with about 19 bars (about 22% in *Liquidambar styraciflua* L.; Schulte, 1999).

Models have not been tested for species with a large number (more than 50) of closely-spaced bars and with steeply inclined end walls, but it is possible that this 'primitive' type of perforation plate would offer still greater resistance to flow. Zimmermann offered an alternative explanation (1983), that the retention of scalariform perforations was adaptive in regions with frost as the bars in the perforations would trap embolisms during the thawing of frozen water. That explanation may apply to Arctic and cool temperate regions, but would not explain the high incidence of scalariform plates in tropical montane regions that, although cool, usually are frost free at 1500–2000 m. More experimental work is needed by physiologists and biophysicists to understand the role of scalariform perforations in the hydraulic architecture of woody plants. Monocots also show a trend from vessel-less to scalariform perforations to the elimination of scalariform perforations, especially within roots and basal parts of stems (e.g. Cheadle, 1944; Thorsch, 2000).

Diameter, density and length of vessels

Vessel diameter and density are less controversial wood functional parameters than vessel element length and perforation plate type. Numerous studies show that in erect trees and shrubs vessel density is roughly inversely proportional to vessel diameter (at least the logarithmically transformed data, Baas, 1973; van der Oever *et al.*, 1981; Noshiro and Baas, 1998, 2000; Klaassen, 1999). Only lianas often have both wide vessels and high vessel densities (Carlquist, 1991). Another correlation that has been established several times is that wide vessels are associated with much greater total vessel length than narrow vessels (Zimmermann and Jeje, 1981). Thus, vessel diameter might be considered a proxy for vessel length for purposes of discussion of hydraulic efficiency.

We have delimited a number of somewhat arbitrary hydraulic types and analysed their occurrence in modern floras and, when data permit, in the geological past:

1. Diffuse-porous woods with very wide vessels ($>200\ \mu\text{m}$) (Figure 15.3a, see Figure 15.6)
2. Diffuse-porous woods with narrow vessels ($<100\ \mu\text{m}$) (Figure 15.3b, see Figure 15.7)
3. Ring-porous woods with very wide earlywood vessels and narrow latewood vessels. (Figure 15.3c, see Figure 15.8)
4. Wood with mixed wide and narrow vessels throughout the growth ring (Figure 15.3d, see Figure 15.5)
5. Lianas: vessels both wide and frequent; often with clusters or multiples of narrow vessels (Figure 15.3e, see Figure 15.5)

Ecological preferences in modern woods

Type 1

Wide-vesselled diffuse-porous species are typical of lowland evergreen rainforest trees (Baas and Wheeler, 2000). The syndrome of wide, infrequent vessels also occurs in some drought tolerant deep-rooting desert trees (e.g. *Acacia gerrardii* Benth. subsp. *negevensis*, *Balanites aegyptiaca* (L.) Del., Fahn *et al.*, 1986). This wood type is not known to occur in shrubs and trees of high latitudes.

Type 2

Narrow-vesselled diffuse-porous species are typical of both evergreen tropical montane and temperate trees and many deciduous temperate species of trees and shrubs. In widespread

taxa the narrowest vessels occur in Arctic and alpine shrubs and in highly xeric desert shrubs. This has been demonstrated for florulas within California (Carlquist and Hoekman, 1985), for the Middle East (Baas *et al.*, 1983), North America, and for temperate versus tropical world woods (Wheeler and Baas, 1991, 1993). Within the genera *Cornus*, *Ilex*, *Symplocos*, or the family Theaceae, the widest vessels occur in tropical lowlands, the narrowest at high latitudes and intermediate at high tropical altitudes.

Type 3

Ring-porosity is typical of seasonal climates in the Northern Hemisphere (Gilbert, 1940; Woodcock, 1994; Woodcock and Ignas, 1994) and subtropical/mediterranean climates. In the tropics, only some deciduous species of seasonally dry monsoon forests are ring-porous; Teak, *Tectona grandis* L. is the most well known example. Ring-porosity tends to be restricted to certain clades (families, genera). Some typically ring-porous genera include diffuse-porous species at the tropical periphery of their primarily temperate distribution (e.g. *Celtis*, *Fraxinus*, *Sapindus*, *Ulmus*) (Baas *et al.*, 1988; Zhong *et al.*, 1992; Klaassen,

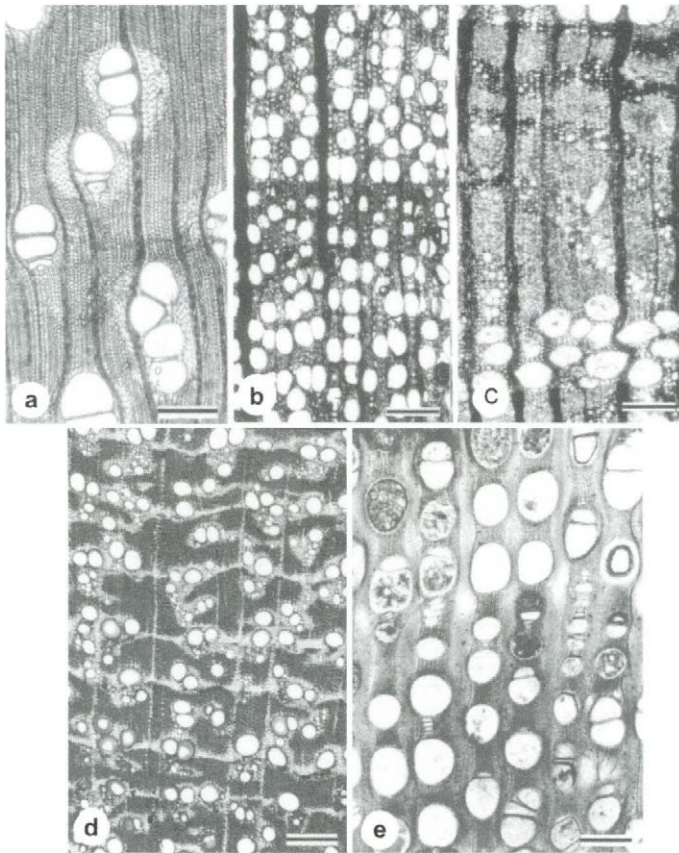


Figure 15.3 Wood types 1, 2, 3, 4 and 5. (a) Diffuse-porous wood with few wide vessels (Clarno Nut Beds, Oregon, Middle Eocene). (b) Diffuse-porous wood with many narrow vessels (Clarno Nut Beds, Oregon, Middle Eocene). (c) Ring-porous wood (Florissant Fossil Beds, Late Eocene). (d) Diffuse-porous wood with different vessel size classes intermingled, *Nitraria retusa* (Forssk.) Aschers. (Recent). (e) Liana-type wood (Clarno Nut Beds, Oregon, Middle Eocene).

1999). Some typically diffuse-porous clades (e.g. Bignoniaceae, *Morus*) include extratropical deciduous genera or species that are ring-porous (e.g. *Catalpa*, *Morus rubra* L.). However, there are also many families that are entirely diffuse-porous, even though they include deciduous species that occur in north temperate or seasonally dry monsoonal forests (e.g. Cornaceae, Hamamelidaceae, Magnoliaceae).

As far as we know ring-porosity only occurs in deciduous plants. A subsequent section on xylem physiology discusses the strategy of ring-porosity and its correlation with phenology.

Type 4

The wood type with different vessel size classes intermingled throughout a growth ring is rather poorly documented, but analyses of ecological trends within the woody floras in the Middle-East and Europe have revealed the common occurrence of this syndrome (Baas *et al.*, 1983; Baas and Schweingruber, 1987). In some ecosystems of temperate and dry Mediterranean to desert regions, 80–97% of all species have a dual strategy for safe and efficient transport with conduits for both (relatively) efficient and (relatively) safe transport.

Type 5

Lianas also show two vessel size classes with large and small vessels intermixed throughout the wood (Carlquist, 1991; Gasson and Dobbins, 1991). This contradicts the notion that efficiency of xylem sap transport is all that matters in lianas with long, slender climbing stems and massive foliage many tens of metres from the root system. There is apparently a need for a safety network through narrow (often clustered) and assumedly short vessels as well.

In Figures 15.4 and 15.5, the incidence of trees and shrubs with two vessel size classes in the European, Middle Eastern and tropical Javanese floras is illustrated. The category of

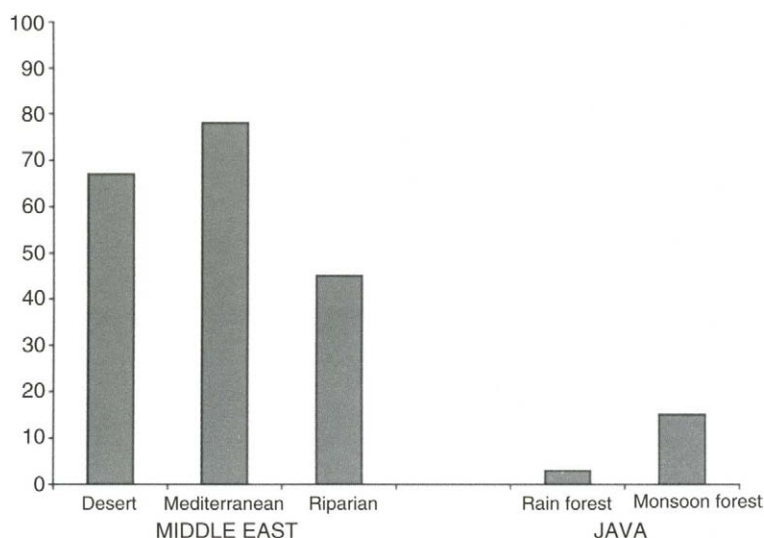


Figure 15.4 Incidence of species (%) with two or more vessel size classes (caused by ring- or semi-ring porosity or by different size classes throughout the growth rings) in different ecological categories in the Middle East and in tropical Java (data from Table 2 in Baas *et al.*, 1983).

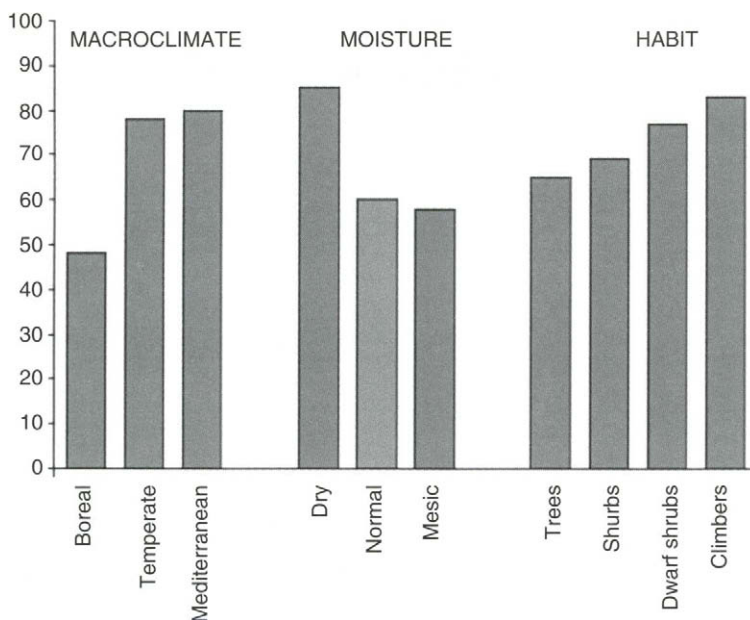


Figure 15.5 Incidence of species (%) with two or more vessel size classes (caused by ring- or semi-ring porosity or by different size categories throughout the growth rings) in different latitudinal (macroclimatic), moisture availability and habit categories in the European flora (redrawn from Baas and Schweingruber, 1987).

two vessel size classes combines wood types 3 and 4 and a subset of type 5. It is striking how much more common the syndrome of different vessel size classes is in seasonal (European and Middle Eastern) than in aseasonal tropical floras. Within these broad geographical categories there are distinct ecological trends showing that the dual strategy for efficiency (relatively wide vessels) and safety (relatively narrow vessels) is most developed in ecological categories that ‘need’ it most (Mediterranean shrubs in Europe and the Middle East and monsoon species in Java) (Baas *et al.*, 1983; Baas and Schweingruber, 1987).

Geological record

We have updated a database for fossil angiosperm woods that was used in earlier analyses of the incidence of selected wood anatomical features through time (Wheeler and Baas, 1991, 1993). Fossil roots and twigs are excluded from consideration because vessel diameter varies from pith to bark and between stem and root. For the diffuse-porous woods within this group, we included only those records with data for vessel diameter. This leaves about 1100 records (Table 15.1).

Very wide vessels, as now commonly occur in tropical lowland floras, were less common in the Cretaceous than in the Tertiary (Figure 15.6). Narrow vessels were more common (Figure 15.7), especially in the Cretaceous, even at tropical palaeolatitudes. In the Tertiary, the statistics fluctuate but, overall, they are not very different from the modern flora. Does this mean that in the Cretaceous there was a greater need to opt for safety rather than efficiency of xylem sap transport? We do not think so. There are at least

Table 15.1 Reports of dicotyledonous woods that are not roots or twigs and have information about anatomy and age (derived from database described in Wheeler and Baas 1991, including updates)

| <i>Age</i> | <i>Number of records</i> |
|------------|--------------------------|
| Pliocene | 192 |
| Miocene | 449 |
| Oligocene | 129 |
| Eocene | 219 |
| Palaeocene | 39 |
| Cretaceous | 80 |
| Total | 1108 |

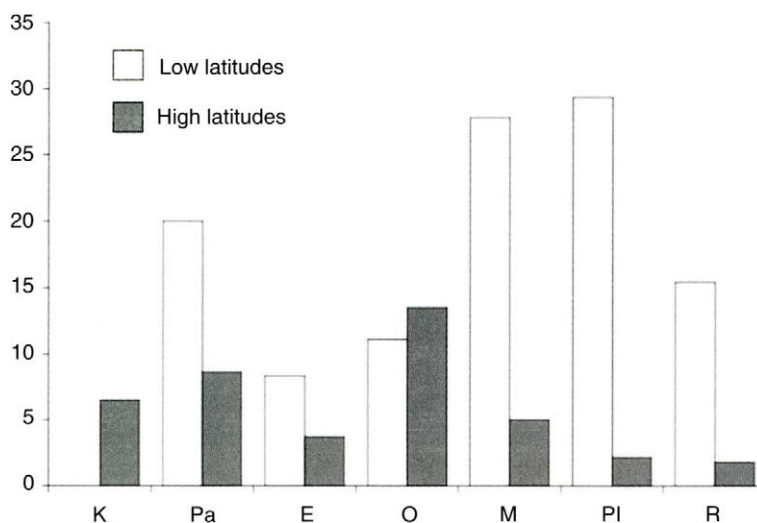


Figure 15.6 Incidence of vessels (%) with average tangential diameter more than 200 μm through time, per time unit for Gondwana (primarily low latitudes) and Laurasia (primarily high latitudes). K = Cretaceous (from Aptian/Albian onward, *ca.* 100–65 Ma); Tertiary records reported for epoch: Pa = Palaeocene, E = Eocene; O = Oligocene; M = Miocene; Pl = Pliocene, R = Recent. Data from Wheeler *et al.* (1986) and Wheeler and Baas (1991) plus updates.

two possible explanations. One is that there was a higher incidence of smaller trees in the Cretaceous than at present; very wide vessels are extremely rare in small trees and shrubs in the recent flora (Wheeler, 1991). Multistratal tropical rain forests with large emergents are believed not to have appeared until the Tertiary (Graham, 1999). A second explanation is related to the higher incidence of scalariform perforation plates in the Cretaceous than in the Tertiary and present-day. A link between scalariform perforations and narrow maximum vessel diameter, as occurs in the present-day, may have constrained the development of wide diameter vessels. However, there are notable exceptions in the fossil record. The Albian *Paraphyllanthoxylon* Bailey from Texas has wide vessels (Serlin, 1982; Wheeler, 1991). Also, one combination of features that is more common in Cretaceous than at present is the co-occurrence of a few wide vessels with scalariform perforation plates, albeit the bars are few and widely spaced. This combination is extraordinarily rare in the extant flora (Wheeler *et al.*, 1987; Wheeler and Lehman, 2000). Why this

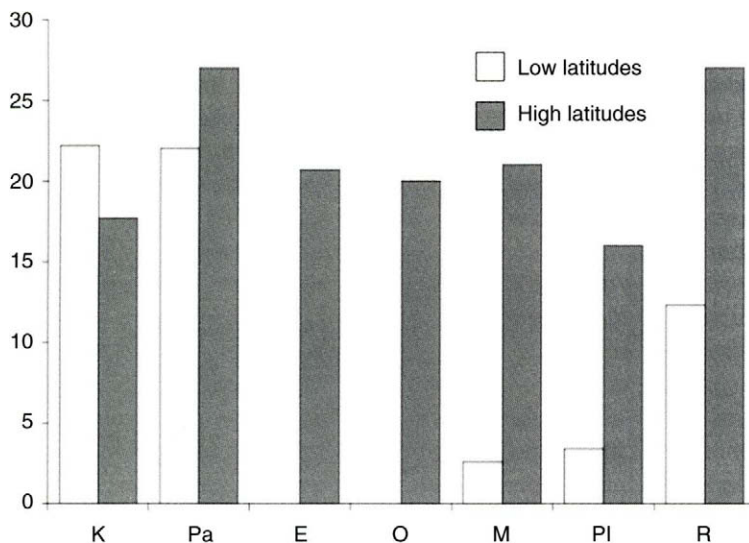


Figure 15.7 Incidence of vessels (%) with average tangential diameter less than $50\ \mu\text{m}$ through time, per time unit for Gondwana (primarily low latitudes) and Laurasia (primarily high latitudes). K = Cretaceous (from Aptian/Albian onward, *ca.* 100–65 Ma); Tertiary records reported for epoch: Pa = Palaeocene, E = Eocene; O = Oligocene; M = Miocene; Pl = Pliocene, R = Recent. Data from Wheeler *et al.* (1986) and Wheeler and Baas (1991) plus updates.

combination should occur in trees of the Cretaceous is unclear. The Cretaceous world was not similar to the present-day, e.g. CO_2 levels were higher and large herbivores were more common. Whether and how these different conditions might affect leaf development and characteristics of Cretaceous plants and, consequently, their xylem differentiation and physiology is a fascinating, but poorly understood topic.

From the Palaeocene onwards the percentage of woods with very wide vessels is at, above, or slightly below modern values, with a clear predominance of widest vesselled species at tropical (palaeo) latitudes (Figure 15.6). Complementarily, a relatively high incidence of tree species with very narrow vessels is typical for high latitudes from the Eocene onwards (Figure 15.7). In the Cretaceous, the Northern (Laurasia) and Southern (Gondwana) floras do not differ in their incidence of narrow vessels. The Cretaceous records for diffuse-porous woods with wide vessels are from Laurasia, in contrast with the Tertiary and Recent. However, during the Albian, Texas and Utah, where these wide-vesselled woods occur, were at lower latitudes than today.

The oldest known ring-porous wood is from the Late Cretaceous of Antarctica (Poole *et al.*, 2000). This report of *Sassafras*-like wood is remarkable and supports the suggestion that high latitude trees of the Cretaceous were deciduous as an adaptation to a seasonal climate, with a long, dark and relatively cool season. The first occurrence of ring-porous woods being in the southern hemisphere is also remarkable because ring-porous woods are now rare in temperate zones of the Southern Hemisphere (data from the OPCN database, Wheeler *et al.*, 1986). In both the Cretaceous and Palaeogene, the incidence of ring-porosity was low and reached more or less modern levels in the Miocene of the northern hemisphere (Figure 15.8), coincident with development of temperate seasonal climates (Graham, 1999).

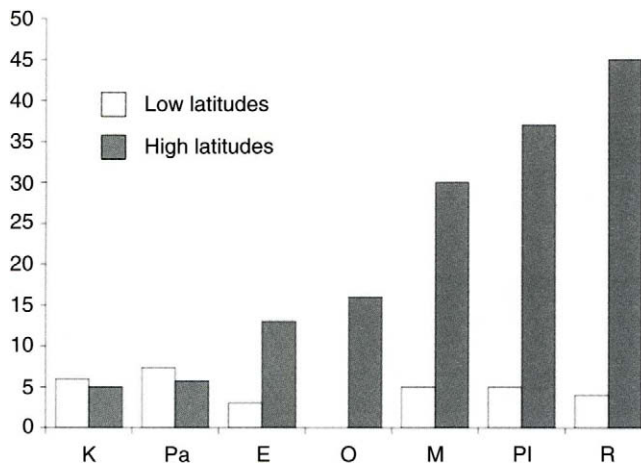


Figure 15.8 Incidence of ring-porous wood (%) through time, per time unit for Gondwana (primarily low latitudes) and Laurasia (primarily high latitudes). K = Cretaceous (from Aptian/Albian onward, *ca.* 100–65 Ma); Tertiary records reported for epoch: Pa = Palaeocene, E = Eocene; O = Oligocene; M = Miocene; Pl = Pliocene, R = Recent. Data from Wheeler *et al.* (1986) and Wheeler and Baas (1991) plus updates.

Unfortunately, data for the fossil wood database were not recorded in a way that allows systematic review for the occurrence through time of the syndrome of two vessel size classes, mixed throughout the rings.

The syndrome of numerous wide vessels mixed together with multiples of very narrow vessels, characteristic of vines, is known from the Late Cretaceous onwards (e.g. Page, 1970; Wheeler and LaPasha, 1994; Melchior, 1998; Poole, 2000; Wheeler and Lehman, 2000). The number of well-documented fossil liana samples is, however, too low for quantitative analysis of trends through time.

This brief summary of hydraulic structure of fossil woods shows that by the end of the Cretaceous the basic types listed above were present. Also, during the Tertiary the general pattern of incidences of vessel diameter categories and ring-porosity corresponds with general changes in climate. More detailed studies of sequences from specific geographic areas, as is being done for Antarctica (e.g. Poole, 2000), are needed to better understand the nature and timing of changes in wood structure through time.

Experimental work

Conductive efficiency versus vulnerability to embolism

Herein, regression analysis of various physiological and anatomical characters is used to look for trends and possible ‘trade-offs’ between various extant species. The temporal and spatial distribution of the wood types discussed above are largely explained by the ‘trade-offs’ between xylem conductive efficiency and vulnerability to embolism. The two major causes of embolism in plants appear to be freezing and water stress, but the mechanism for embolism formation differs in these two cases (Figure 15.9). For drought-induced embolism, the size of pores in the pit membranes appears to be the critical factor to resisting embolism (Sperry and Tyree 1988; Jarbeau *et al.*, 1995; Sperry *et al.*, 1996). In Figure 15.9, B has few relatively wide pores and therefore the capillary forces preventing

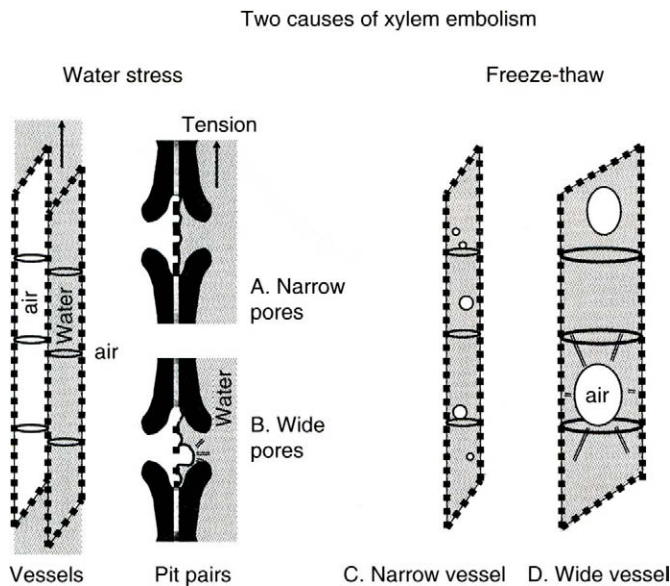


Figure 15.9 Water stress and freezing stress are the two major causes of embolism in plants. Vulnerability to water-stress induced embolism is proportional to the diameter of pores in the pit membrane; case B is more vulnerable than A. Vulnerability to freeze-induced embolism is proportional to the diameter of the vessel, and so case D is more vulnerable than C. Modified from Langan *et al.* (1997) and Sperry *et al.* (1996).

the entry of air from an adjacent air-filled conduit are not as great as in A, which has relatively narrow pores.

The positive correlation between conductive efficiency and vulnerability to freezing-induced embolism is clear (Figure 15.10); wider conduits, which are the most efficient in transport, are also the most prone to embolism caused by freezing (Ewers, 1985; Cochard and Tyree, 1990; Sperry and Sullivan, 1992; Davis *et al.*, 1999b). Evidently, vessel diameter determines the size of gas bubbles forced out of solution during a freezing event and wider vessels produced larger bubbles which are more resistant to dissolution at the time of thaw. Such bubbles are more likely to result in embolism. Why would plants evolve such risky vessels? There must be a selective advantage for wide vessels, as they occur early in the history of angiosperm wood. According to Poiseuille's law for ideal capillaries, conductive efficiency should be proportional to the sum of vessel diameters to the fourth power (Zimmermann, 1983). This means that doubling the number of vessels per cross-sectional area should double k_s , whereas doubling vessel diameter should increase k_s by 16-fold.

The amount of embolism with freeze-thaw depends not just on vessel diameter but also the water stress that the plants experience during the freeze-thaw event (Sperry and Sullivan, 1992; Sperry *et al.*, 1994), the rate of thaw (Langan *et al.*, 1997) and the position of the stem within the plant (Lemoine *et al.*, 1999). Davis *et al.* (1999b) found that for isolated 8–12 mm diameter stem segments under a water stress of -0.5 MPa, vessels wider than $44 \mu\text{m}$ will always embolize upon freeze/thaw (Figure 15.10). There may be a similar vessel diameter threshold for 8–12 mm stems with regard to phenology of wood production and leaf maturation.

It is well established that temperate ring-porous trees, which have very wide earlywood vessels, have delayed bud break and delayed leaf maturation in the spring (Lechowicz,

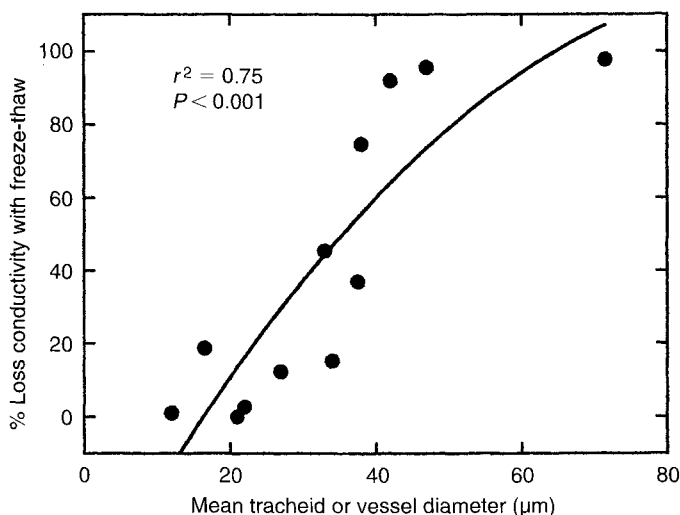


Figure 15.10 Percent loss conductivity with freeze–thaw as a function of mean tracheid diameter (in the case of two conifer species) or mean vessel diameter (in the case of 10 angiosperm species) in 8–12 mm diameter stems. Each point is an average for one species, the curve fit was a second order polynomial. Modified from Davis *et al.* (1999a).

1984; Wang *et al.*, 1992). Why is vessel diameter related to leaf phenology in temperate woody plants? In north temperate areas, where freezing now occurs every year, the wide earlywood vessels of ring-porous trees remain conductive for only one growing season (Zimmermann, 1983; Ellmore and Ewers, 1986; Cochard and Tyree, 1990; Sperry *et al.*, 1994; Hacke and Sauter, 1996; Jaquish and Ewers, 2001). The narrow latewood vessels often remain conductive for several years, but they contribute little to the total hydraulic conductance of the stem. For instance, in *Ulmus americana* L. the latewood vessels of wood more than a year old contributed only about 6% of the total axial conductance (Ellmore and Ewers, 1986). The total conductive efficiency of stems of ring-porous trees tends to be quite high, but there is a cost because their current year's wood does not mature until danger of freezing is past in the late spring and thus they have a shorter growing season. Apparently there has been natural selection for ring-porous trees to delay maturation of their leaves and their new xylem, thus allowing the mature leaves to have a reliable water supply (Zimmermann, 1983; Lechowicz, 1984; Wang *et al.*, 1992). While ring-porous wood can be extremely efficient in conduction, the wide vessels tend to be prone to dysfunction during periods of freezing or severe drought. For species with wider vessels, bud break and leaf maturation is delayed in the spring and the leaves also tend to senesce earlier in the autumn, with the result that there is an inverse correlation between vessel diameter and the length of the growing season. In Figure 15.11, growing season length was defined as the number of weeks between bud break in the spring and leaf colour change or leaf abscission in autumn. Note that in Figures 15.10 and 15.11, narrow stems were used; there is research currently underway to examine the same relationships in the trunk xylem of trees, which have much wider vessels than those of the narrow stems.

At the other extreme from ring-porous trees and shrubs in North America, evergreen plants have the longest potential growing season and they have consistently narrow conduits. At the North American sites for the study shown in Figure 15.11, there were also seven evergreen species present (not shown in Figure 15.11), five conifers and two

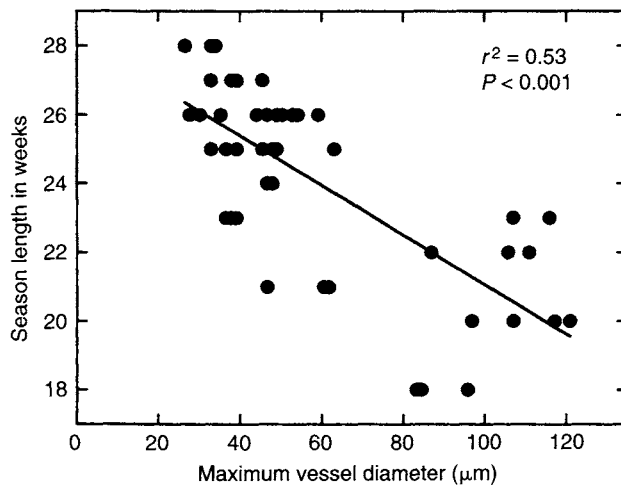


Figure 15.11 Growing season length in weeks as a function of maximum vessel diameter in 7–11 mm diameter stems of woody deciduous plants in Michigan, USA. Seventeen species were sampled in 1991, with three individuals per species. Season length was defined as the number of weeks between bud break in the spring and leaf colour change or leaf abscission in autumn. Wide vessel species tend to have late bud break and early leaf senescence, resulting in a short growing season. Data from a study by Billingsley *et al.* (1994).

dicotyledons, whose stems had a maximum conduit diameter of $23.1 \mu\text{m}$ (SE = 4.3, $n = 7$), well below the freeze/thaw embolism threshold for stems of that size. Their freezing resistant transport system allows evergreen plants to have the relatively long growing seasons, albeit with low conductive efficiency that can limit their maximum photosynthetic rates (Pallardy *et al.*, 1995).

Resistance to drought-induced embolism, like freezing embolism, appears to be inversely related to conductive efficiency. For instance, Tyree *et al.* (1994), in a literature review on vulnerability curves comprising over 60 plant species from tropical, Mediterranean, and temperate habitats, found a weak, but significant ($r^2 = 0.18$) linear regression between resistance to embolism and conductive efficiency. The correlation is statistically significant due to the high number of species sampled, but the weakness of the correlation is not satisfying to those who wish to understand the mechanisms. However, when only plants from a particular habitat are considered, for instance, chaparral plants of southern California, the inverse correlation is stronger (Figure 15.12; $r^2 = 0.31$) but still not as strong as for freeze-thaw embolism. However, it appears that drought-induced embolism is indirectly related to vessel diameter and directly related to the size of pit membrane pores (Sperry and Tyree, 1988; Pockman *et al.*, 1995; Jarbeau *et al.*, 1995; Sperry *et al.*, 1996). In tracheids, the pit membrane resistance to flow is often as great or greater than lumen resistance (Gibson *et al.*, 1985; Calkin *et al.*, 1986) and the same may be true for vessels. In angiosperms, as in conifers and ferns, the measured conductance is typically about 20–80% of the theoretical axial conductance defined by Poiseuille's law (Tyree and Zimmermann, 1971; Chiu and Ewers, 1992; Hargrave *et al.*, 1994; Wagner *et al.*, 1998). The reduction in measured conductance below the theoretical axial conductance may be due largely to pit membrane resistance. It follows that if pit membrane resistance is one of the major limiting factors to water flow, k_s should be proportional to the diameter of pores in pit membranes and inversely proportional to resistance to drought-induced embolism.

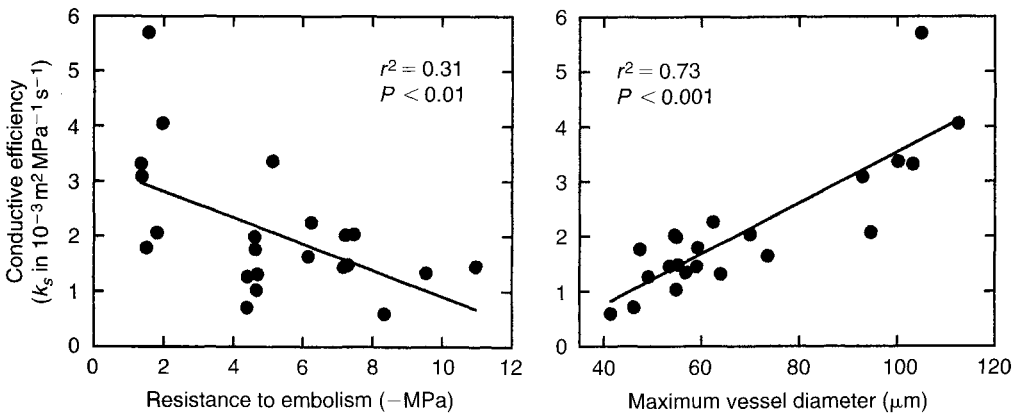


Figure 15.12 Conductive efficiency (k_s in $10^{-3} \text{ m}^2 \text{ MPa}^{-1} \text{ s}^{-1}$) as a function of resistance to embolism and maximum vessel diameter in species of chaparral shrubs in southern California. Each point is an average for one species. Resistance to embolism is equal to the water potential at 50% embolism, based upon dehydration vulnerability curves. Unpublished data from studies by Davis *et al.*

As with freezing embolism, the narrow vessels and tracheids in a stem are generally more resistant to drought-induced embolism than are wide vessels in the same stem (Salleo and Lo Gullo, 1986, 1989; Sperry and Tyree, 1988; Lo Gullo and Salleo, 1991; Hargrave *et al.*, 1994). Thus there seems to be validity to Carlquist's hypothesis (1988) that narrow vessels and tracheids serve as high resistance auxiliary pathways for use when wide vessels become embolized. This would explain the common occurrence of narrow conduits closely associated with wide ones within the woods from arid regions.

The common occurrence of wood with strictly narrow vessels or tracheids in Arctic, alpine and dryland habitats is best explained in terms of resistance to freezing or drought-induced embolism. Diffuse-porous wood, a characteristic of the earliest known dicot woods, appears to be adapted to a wide range of habitats. Ring-porous wood appears to be more narrowly adapted to strongly seasonal environments where growing conditions are suitable for only a part of the year. This is consistent with the first appearance of ring-porous woods in the Late Cretaceous of Antarctica and the increased incidence of this wood type in the Late Tertiary of the Northern Hemisphere when seasonality becomes more pronounced (see Figure 15.8).

Conductive efficiency versus mechanical strength

Increasing vessel diameter or vessel frequency, both of which would increase conductive efficiency, should tend to make less dense, weaker wood, since vessels represent weak areas of the xylem. For modulus of rupture, it is the 'weakest link' that determines the point of failure and for wood the weakest mechanical link could be the vessels (Wagner *et al.*, 1998). An inverse relationship between mechanical strength and conductive efficiency has been found when comparisons are made between closely related taxa. For instance, when co-occurring species pairs of *Ceanothus* L. and *Adenostoma* Hook. and Arn. were examined, *C. spinosus* Nutt. and *A. sparsifolium* Torr. had greater k_s values but lower stem mechanical strength than the co-occurring *C. megacarpus* Nutt. and *A. fasciculatum* Hook. and Arn. (Wagner *et al.*, 1998). Similarly, within *Toxicodendron diversilobum* (Torr. and Gray) Greene, a species that produces heteromorphic stems, viney (supported)

and shrubby (unsupported) stems showed a similar trend, viney stems having greater conductive efficiency but being mechanically weaker than shrubby stems (Gartner, 1991a,b). However, in angiosperms there can be mechanical compensation for wide vessel lumens with the evolution of thicker-walled fibres, tracheids and vessels. Although the entire range from very thin-walled to very thick-walled fibres occurs in emergent tropical trees, thin-walled fibres are more typical for rapidly growing early successional species and thick-walled fibres more common in climax species (Swaine and Whitmore, 1988).

When comparisons are made between lianas and free-standing trees and shrubs of the same genus or within the same family, the liana taxa usually have wider conduits (Figure 15.13, route B), or greater vessel frequency per cross-sectional area (Figure 15.13, route A), depending on the taxa (ter Welle, 1985; Fisher and Ewers, 1995; Ewers *et al.*, 1997). The wide vessels and high vessel frequency have resulted in lianas having greater conductive efficiency but weaker stems than free-standing growth forms (Ewers *et al.*, 1991; Chiu and Ewers, 1992). The weaker stems for lianas are not a liability since they depend on host plants or other external objects for mechanical support.

As mentioned above, for free-standing growth forms, vessel frequency per transverse area is inversely proportional to vessel diameter. This may be because wood with relatively few wide vessels may be as mechanically strong and have greater conductive efficiency, than wood with many narrow vessels (Figure 15.13, route C). Conversely, if vessel diameter is increased without decreasing vessel frequency (Figure 15.13, route B), then mechanically weak, liana-like wood is produced. However, in angiosperms, the evolution of thicker-walled fibres can help to compensate for the mechanical weakness brought on by wider vessels. Therefore vessel evolution is likely to affect fibre evolution and vice versa.

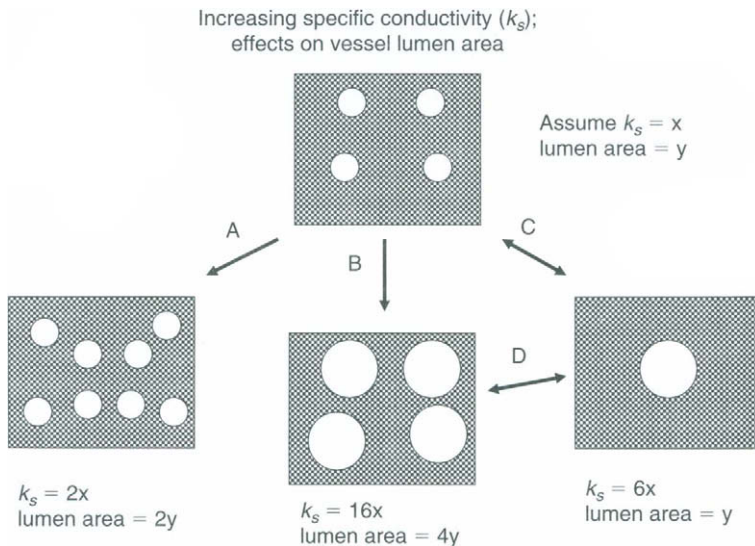


Figure 15.13 Effect of relative vessel diameter and frequency per transverse area on theoretical hydraulic efficiency (k_s) and lumen area. Large lumen area per transverse area can weaken the wood. According to Poiseuille's law for ideal capillaries, doubling the vessel frequency doubles the lumen area and merely doubles the k_s (route A), whereas doubling the diameter increases lumen area 4-fold and increases k_s by 16-fold (route B). Route C shows a method of increasing k_s without increasing lumen area, perhaps explaining the typical inverse relationship between vessel frequency and vessel diameter in most taxa.

Cohesion–tension theory and sap ascent

The cohesion–tension theory has been the leading explanation for xylem transport in plants for more than a century and, despite recent challenges (Balling and Zimmermann, 1990; Canny, 1995, 1997) support for the theory is quite robust (Tyree, 1997; Steudle, 2001). The theory is critical to our understanding of xylem evolution, since it means that the water is normally under considerable negative pressure (tension) during transport. The vessels thus must be constructed to avoid the entry of air, which can cause cavitation and embolism (air blockage) of the water columns. They also have to withstand the negative pressure (tension) of the water column to avoid implosion. The cohesion–tension theory also implies that parenchyma cells, which are present in most woods, do not have a direct role in transport but function in storage of photosynthates, water and minerals and for resisting the invasion of pathogens.

Mechanical strength, implosion resistance and resistance to embolism

In a recent study including 36 species of angiosperms and 12 species of conifers from North America, it was found that there was a positive correlation between wood density and resistance to drought-induced embolism (Hacke *et al.*, 2001). The authors offered the explanation that resistance to drought-induced embolism required, among other things, that vessels or tracheids must be able to resist implosion due to the water tension (negative pressure) that the conduits experience. They argued that the critical factor was $(t/b)^2$, that is, the square of the thickness of the vessel or tracheid wall (t) divided by the diameter of the conduit (b), as used in Figures 15.1 and 15.14. Therefore, to resist collapse of the conduits, plants that experience very low water potentials must have thick walls and narrow lumens, with high wood density as a corollary. However, it may not be avoidance of vessel implosion itself that causes the observed correlation, but characteristics of the pit membranes. Stretching of pit membranes under extreme negative pressures may enlarge the pit membrane pores so that they are more prone to air seeding. Is deflection of pit membranes in high density woods less likely to enlarge pit membrane pores?

Another possible explanation is that selection for resistance to embolism would tend to result in narrower conduits, as argued above. Wood with narrower conduits will tend to have greater branch strength. Among chaparral shrubs we have found that resistance to

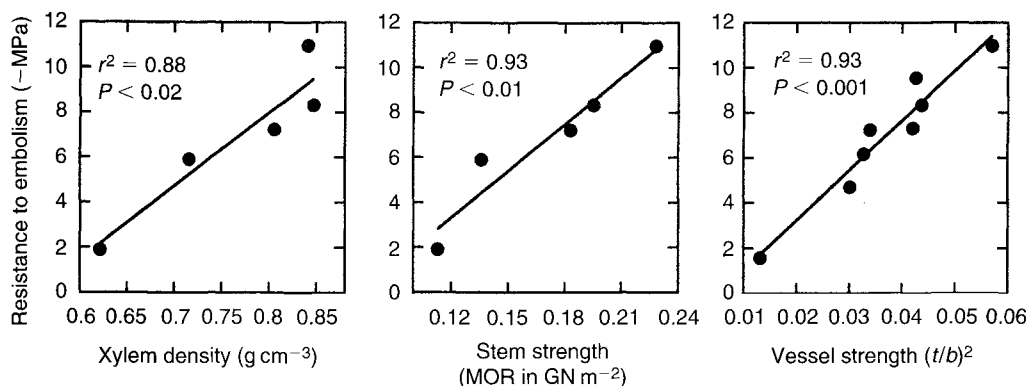


Figure 15.14 Resistance to embolism as functions of xylem dry density, stem strength (modulus of rupture, MOR in GN m^{-2}), and vessel strength $(t/b)^2$ in chaparral shrubs. Each point represents the mean for one species. Based upon data from Wagner *et al.* (1998) and Hacke *et al.* (2001).

embolism correlates well with wood density, stem strength and vessel strength (Figure 15.14). It is well known that wood density is correlated with wood strength, but in some cases high wood density may have been selected for as a mechanism to resist vessel embolism or as a mechanism to resist vessel implosion. It remains to be determined which of these factors are most critical in wood evolution.

Hacke *et al.* (2001) note that fibre evolution would impact implosion resistance of vessels and they suggest that, as a result, $(t/b)^2$ values are fundamentally different for angiosperms versus gymnosperms. In angiosperms, fibres offer some of the support against implosion, whereas in conifers only tracheid walls are involved with implosion resistance.

Roots versus stems

The aforementioned studies have all involved the xylem of stems. In considering the 'trade-off' triangle, it should be noted that, compared to stems, roots would normally not experience such low water potentials, nor would the mechanical support demands be as great as for free-standing stems; the soil surrounding roots provides some of the mechanical support for plants. In addition, roots tend to play a greater role in carbohydrate storage than do stems. Given the above considerations, it is not surprising that woody roots in many plants have wider vessels than woody stems, more parenchyma and less fibre (Fegél, 1941; Zimmermann and Potter, 1982; Gasson, 1985; Gartner, 1995; Pate *et al.*, 1995). Based upon vulnerability curves, roots tend to be more vulnerable to embolism than are stems, as might be predicted by their wider vessel diameters (Sperry and Saliendra, 1994; Alder *et al.*, 1996). However, some species' roots have fewer, narrower vessels than do their stems and so their roots may prove to have similar or lower vulnerability to embolism than the stems (see illustrations in Cutler *et al.*, 1987). Another exception may be for lianas, where the mechanical demands on the stems are much reduced. In lianas, unlike free-standing growth forms, vessel diameters of stems tend to be similar to those found in roots (Ewers *et al.*, 1997).

Parenchyma

In this chapter little attention was paid to the role of parenchyma (ray and axial). The abundance and distribution of parenchyma varies considerably within the angiosperms. Some schemes suggesting functional significance in variation of parenchyma distribution and abundance have been devised (e.g. Braun, 1970).

Woods with abundant (diffuse) parenchyma and high ray volumes are common in some Cretaceous floras (Wheeler *et al.*, 1987; Wheeler and Lehman, 2000): were these woods indicative of xeric conditions and the need for water storage? In modern floras such types are exemplified by certain Bombacaceae (*Adansonia* L., abundant axial parenchyma) and Cucurbitales (very broad rays). Alternative strategies for water storage are found in woods that lack axial parenchyma but have septate fibres which take over the water storage and other living functions of parenchyma, e.g. in many Sapindales (Burseraceae, Sapindaceae, Meliaceae). This wood type is among the earliest known and is abundant throughout the Cretaceous (Wheeler and Baas, 1991), but today tends to be most common in mesic tropical rain forests (Baas, 1982).

Towards a synthesis: the evolution of hydraulic structure and function

What sense do the global ecological trends in xylem anatomy make in terms of the experimentally demonstrated 'trade-offs' between vessel diameter, vessel density and dual vessel

diameter strategies and hydraulic efficiency (conductivity) and safety (control and repair of tension and freezing embolisms)?

In frost-prone areas it has been abundantly shown that the freezing/thawing embolisms must somehow be controlled and prevented from spreading. That possibility exists in short- and narrow-vessel woods (often provided with scalariform perforations), which is the most common type in these regions. Woods with a dual safety-efficiency strategy (ring-porosity; two vessel size classes in diffuse-porous condition) are extremely common in ecosystems characterized by seasonality where temporary high demands for conductive efficiency alternate with demands for safety and localization and/or repair of drought-stress embolism. Finally, wide-vessel diffuse-porous woods, opting for high conductive efficiency, without apparent safety provisions characterize the rapidly dwindling tropical rainforest flora. This should perhaps surprise us rather than confirm our intuitive ideas. After all tall rainforest emergents have their transpiring crowns above the canopy, with daily sunny periods when the humidity on the forest floor stays high, but becomes quite low above the canopy. According to the cohesion theory these tall trees have to develop considerable xylem sap tensions, although the tensions are less than those that shorter, arid habitat plants experience during droughts. The experimental work summarized in this chapter provides a basis for understanding the variations in the incidences of vessel diameters as related to freezing and drought. However, additional experimental work is needed to elucidate the physiological significance of variations in vessel wall thickness, vessel perforation type, wood density and parenchyma distribution.

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16

Hydraulics and mechanics of plants: novelty, innovation and evolution*

Nick Rowe and Thomas Speck

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Introduction

Physiological traits are exceptionally difficult to ascertain in fossils of long extinct plant groups. In this chapter we discuss the significance of hydraulic and mechanical novelties, both of which are physiologically interrelated and show high levels of interdependence

* Dedicated to Hanns-Christof Spatz on the occasion of his 65th birthday.

and constraint. Mechanical tissue for generating stiffness is most effective if comprised of thick-walled fibre elements, but these are physiologically expensive; water-conducting elements are most effective if composed of wide diameter elements but such tissues are less dense and thus contribute less effectively to mechanical stiffness. These two functions could, therefore be viewed to be in conflict. Another example of conflicting functional roles is seen in stem systems without specialized mechanical tissues, where mechanical structure is upheld by the hydraulic system and maintenance of turgor. Turgor requires living parenchyma cells and when these are fully hydrated the stem/leaf system is mechanically sound and does not wilt. If water supply becomes limited the stem/leaf system will wilt if there are no other mechanical strengthening elements within the stem. Advanced wilting and water stress can have a range of deleterious effects including embolism and mechanical failure. There is a conflict in this system because turgescence water-filled cells are heavy when fully hydrated and thus mechanically load the plant. These are some simple examples of potential compromises between mechanics and hydraulics that have probably played important roles during radiations of terrestrial plants. These kinds of processes are potentially relevant to a wide range of plant organs as well as entire plants of different complexity. The 'trade-offs' associated with turgor, such as weight load, were probably important for many early land plants and are still so for almost any organ or young developmental stage of any plant. The compromise between hydraulic and mechanical functions in terms of water-conducting elements and fibre tissues has many examples among more complex architectures such as the appearance of hypodermal steromes and lateral meristems. Hydraulic and mechanical constraints can be seen as coupled functions and, to a large extent, physiological compromises which have to be met for adaptive modification of the plant body. Many authors have commented on this, but in our view co-organization of these constraints is especially crucial for morphological radiation of growth forms. By growth form we mean the size, shape and, importantly, the mechanical posture or orientation of the plant. Self-supporting plants can show quite different mechanical and hydraulic specializations from non-self-supporting plants. Trees, shrubs, herbs, climbers, epiphytes and hemi-epiphytes show an incredible variety of hydraulically and mechanically coupled innovations. Given the recent advances in plant biomechanics and hydraulics as well as more decisive and testable methods for establishing phylogenetic and historical contexts, the combination of functional biology and phylogenetic studies offers the opportunity for more accurately understanding how diverse plant forms and their underlying functioning have evolved.

Terminology and evolution

An accepted terminology for unambiguously referring to functional aspects of an organism and their evolutionary significance is a rich area of controversy. The main point of discussion is historically and in our opinion misleadingly centred around adaptation. In the latter half of the 20th century, such discussions focused particularly on whether it is possible to identify traits which are of adaptive significance (Gould and Lewontin, 1979; Gould and Vrba, 1982; Rose and Lauder, 1996). This has more recently led to an extension of this theme questioning the adaptive significance of morphological radiations in plants as well as the gradual or punctual processes characterizing them (Bateman and DiMichele, 1994; Bateman, 1999a,b). While these arguments are of great interest in resolving high level evolutionary processes we argue that significant avenues of doubt still remains over

the actual functioning of many observed traits, let alone whether they are adaptive or not. This is particularly relevant to systems of long extinct organisms which might show little in the way of modern potential analogues (Novacek, 1996).

Another term directly relevant to hydraulic and mechanical function is that of 'evolutionary innovation' or 'key innovation', terms that are generally deployed to describe a trait or syndrome which launched a significant morphological radiation. Perceived interpretation of a key innovation without a (1) functional, (2) phylogenetic and historical framework is not proof. Recent observers point to more combined and testable approaches which usually include (1) a functional argument to assess the 'performance' of the structure and (2) a phylogenetic framework to observe and test patterns of traits within the context of a chosen evolutionary hypothesis (Jablonski and Bottjer, 1990; Sanderson and Donoghue 1996; Bateman, 1999b). Both analyses of function and evolutionary pattern are complicated by the fact that many plants are complex structures and that a single trait or structure might have several functions and confer several benefits to the life history of the plant which might interplay in any number of ecological, evolutionary or adaptive scenarios. This aspect of functional biology is one of several bugbears at the heart of controversies concerning function and evolution where functional studies attempting to assess the significance of one of a range of functions stand accused of implicitly 'atomizing' an organism's biology (Gould and Lewontin, 1979).

Inferred physiological constraints and innovations have formed the basis of many evolutionary discussions concerning plant radiations, especially the appearance and morphological radiations of land plants. A convincing interpretation of function is a basic requirement for any higher-level conceptual interpretation of evolutionary process. In this chapter we discuss the impact of biophysical studies for interpreting specific functions of plants and how knowledge of basic biomechanical results from experimental work on living plants can be usefully applied, with caution, to long extinct but nevertheless, pivotal representatives of major radiations. The second aim of this chapter is to demonstrate the complex nature of interpreting functional process – let alone innovatory or adaptive significance – and to investigate examples from different levels and periods of plant evolution. While specific functional and evolutionary approaches have been traditionally the realm of zoologists, e.g. (Lauder, 1990, 1991; Larson and Losos, 1996), we hope that this contribution might help to consolidate and open the field more into studies based on plants.

Hydraulic and mechanical functioning are two of several areas of functions commonly believed to be essential for invading and establishing major plant lineages on the land. Hydraulic innovations have been popularly stated to include: an outer envelope of cuticle to control evaporation from the plant body; the appearance of primary xylem or xylem-like water-conducting tissue to increase conductance over less efficient parenchyma; the appearance of stomata to control and optimize gaseous exchange; and the appearance of roots or rhizoids for water uptake. Mechanical innovations have been stated to include an outer envelope to act as a mechanical wall of tissue surrounding soft tissues – in mechanical parlance, a pneumatic structure; the appearance of fibre tissue comprising a hypodermal sterome towards the outside of the stem for added mechanical stiffness; the appearance of xylem tissue to impart additional mechanical stiffness and toughness; and the appearance of lateral meristems for augmenting and mechanically sustaining increased body size and complex branched architectures. From both the relatively simple viewpoint of perceived function up to the more conceptually challenging levels of perceived ecological performance and evolutionary fitness many, if not all, of the major structures discussed have therefore been attributed both hydraulic and mechanical significance.

There exists a range of interpretational levels when discussing trait appearance, function, innovatory significance and what is best described as interactedness between other functional parts. Within the context of the hydraulic and mechanical aspects discussed here we provisionally define five terms for the sake of argument, for this chapter.

Function How a structure or tissue functions; a single tissue or combination of tissues may of course have more than one function. Biophysical approaches based on living plants can measure hydraulic conductance and mechanical properties and can be integrated within biophysical models of fossil plants.

Novelty A *de novo* structure or a derived modification of an existing one which significantly alters its function. Its identification requires empirical functional analyses as well as a historical perspective ideally from phylogenetic and/or temporal references.

Innovation A hypothesis in which a novelty arose that 'triggered' a major diversification. Its identification requires empirical functional analysis and a historical perspective as well as a test of competing hypotheses as would be performed by integration of functional traits in an experimental and comprehensive phylogenetic analysis. The word 'triggered' is important here, there could be two possible inferences to this: (1) the trigger was immediate in a temporal sense and the diversification followed the appearance of the novelty; (2) the key innovation represented only part of a 'template' for potential evolutionary diversification but still required either or both (a) intrinsic element(s): additional novelties or (b) extrinsic element(s): a favourable ecological regime.

Integration Implicit or necessary functional coordination of novelties within the organism; the introduction of one functional novelty, such as secondary growth of xylem, may 'require' further developmental characteristics, e.g. compensatory tissue arrangement and additional lateral cambia such as a periderm for accompanying expansion of the stem.

Modularity Dissociation of novelties within a functional whole; the introduction of one functional novelty; e.g. *de novo* production of sclerenchyma fibres within a parenchymatous cortex might not require coupled developmental processes to accommodate or function with other novelties or existing structures.

We have selected four divergent themes relevant to discussions on the evolution of hydraulic and mechanical functioning. These range from: (1) recent biomechanical analyses of early land plants, tracheophytes and lignophytes; (2) a consideration of the mechanical significance of lignin in plant cell walls; (3) an example from the Permian of the relatively 'late' appearance of reaction wood representing physiologically mediated mechanical modulation of a wood cylinder. Finally we present a brief discussion on the hydraulic and mechanical 'trade-offs' involved in specialized climbing growth habits and the fact that recent phylogenetic analyses (Bowe *et al.*, 2000; Chaw *et al.*, 2000; Donoghue and Doyle, 2000) now point to potentially separate origins of sophisticated lianoid growth habits in Gnetales and angiosperms.

Turgor

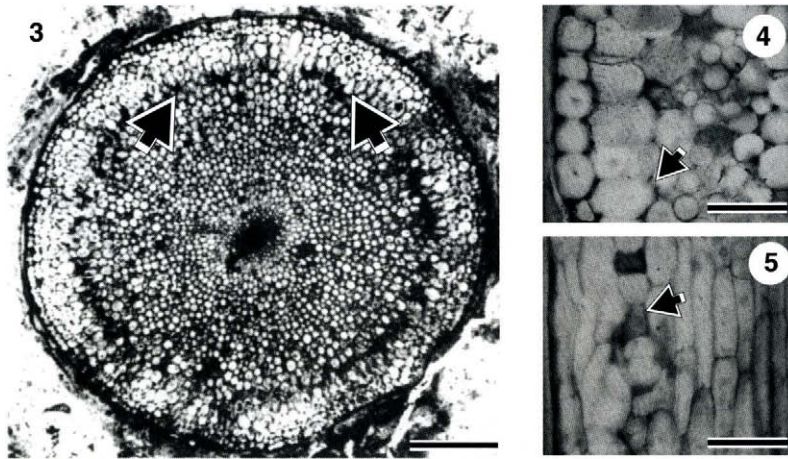
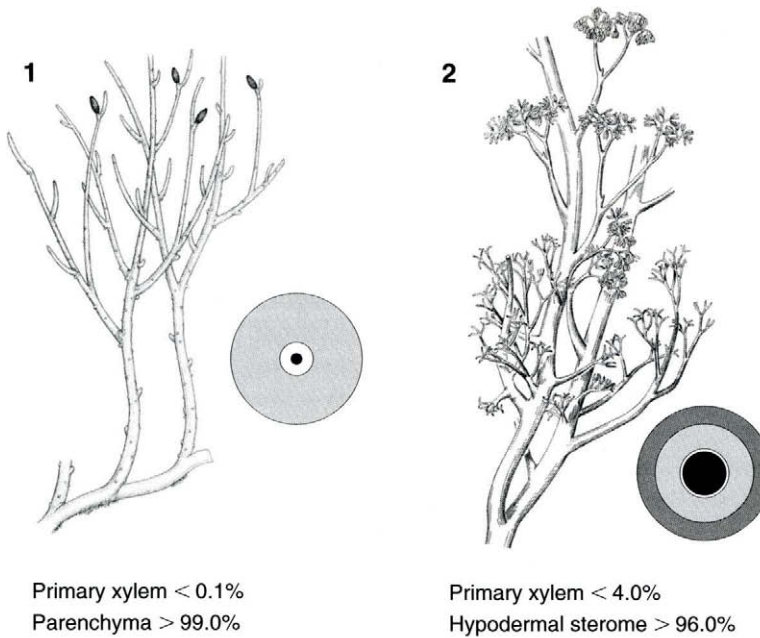
Recent discussions focusing on the probable ecology of early land plants suggest that the initial phases of land colonization were undoubtedly dominated by hydraulic novelties (e.g. Raven, 1984, 1994; Edwards, 1996, 1999). Homoiohydry (physiological maintenance of a hydrated plant body) resulted from the appearance of an outer cutinized envelope surrounding hydrated, living plant tissues. The initial function of the outer cuticle of putative Palaeozoic bryophytes and 'protracheophytes' was almost certainly hydraulic in

combating desiccation. Hydrated turgor systems in plant stems were mechanically important for upright columnar growth and spreading semi-recumbent clonal axes in plants such as *Aglaophyton major* Kidston and Lang, *Horneophyton lignieri* Barghorn and Darrah and *Rhynia gwynne-vaughanii* Kidston and Lang. In such stems, hydrated parenchyma was the principal contributing element to stem stiffness (Speck and Vogelhehner, 1988, 1994; Niklas 1989, 1992; Speck *et al.*, 1990). Thickened epidermal cell walls combined with a layer of cuticle could also have functioned as a peripheral tension bracing system and mechanical pneumatic structure (Speck and Vogelhehner, 1988; Speck *et al.*, 1990). Quantitative biomechanical models of this grade of organization indicate that central conducting strands were small and too centrally placed within the stem to contribute significantly to stem flexural stiffness (Speck *et al.*, 1990; Speck and Vogelhehner, 1994) (Figure 16.1); stem stiffness permitting upright and probably mutually supported stands of upright stems was almost entirely a result of turgor systems. The central conducting strands of such early land plants represent primarily hydraulic functional novelties (Roth and Mosbrugger, 1996; Bateman *et al.*, 1998; Roth *et al.*, 1994, 1998; Roth-Nebelsick *et al.*, 2000; Konrad *et al.*, 2000) compared with aquatic and dorsoventral body plans of antecedent forms. Hydraulic optimization via tracheid-like central strands and the increased and more rapid supply (Raven, 1994) to more extensive and more distant parts of the plant body probably conferred improved hydraulic supply for more extensive clonal systems of horizontal axes.

Central conducting strands did not contribute considerably in a direct manner to stem stiffness and were primarily hydraulic in function. On the face of it, this appears to have permitted greater hydraulic supply to a larger plant body and ensuring greater degrees of intrinsic homoiohydricity. The appearance of a small central strand of tracheids is effectively a 'modular novelty' in that its appearance did not necessitate further simultaneous developmental novelties for the new structure to function; this holds true if one assumes that a cuticle and epidermis with stomata for regulating transpiration and rhizoids for water uptake had already been put in place. Within limits, the appearance of a conducting strand could be interpreted as a 'key innovation' in as much as it set a template for optimized conductance in a number of putative lineages (Kenrick and Crane, 1997a,b). In terms of an immediate morphological diversification, the appearance of the conducting strand probably represented an innovation for extensive clonal growth and low-lying vegetation commonly depicted for the early land plant radiation, but not one for significantly enlarging and expanding plant architectures into the aerial realm.

The hypodermal sterome

By the Middle Devonian many lineages of land plants possessed members with thick-walled fibre tissues at or towards the outside of the stem cross-section. Biomechanical investigations indicate that these types of tissue configurations represented significant contributions to the stiffness of the stem (Speck *et al.*, 1990; Speck and Vogelhehner, 1994) potentially allowing greater height and more diverse branched architectures than were possible from turgor systems. Devonian plants such as *Psilophyton dawsonii* Banks, Leclercq and Hueber (Figure 16.2), *Goslingia breconensis* Heard and species of *Zosterophyllum* Penhallow show that the hypodermal sterome generally contributes to over 95% of the flexural stiffness of the stem. In these plants at least, the hypodermal sterome undoubtedly had a mechanical function.



Figures 16.1–16.5 Biomechanics of the central conducting strand and hypodermal sterome in two representative early land plants. Figures 16.1 and 16.2 Reconstructions of *Rhynia gwynne-vaughanii* and *Psilophyton dawsonii*, cross-sections: black = central xylem tissue; white = phloem; light grey = parenchymatous cortex; dark grey = hypodermal sterome. Figure 16.1 *Rhynia gwynne-vaughanii*; the central xylem elements contribute less than 0.1% to the flexural stiffness of the stem, the parenchymatous cortex dominates in contributing to the stem stiffness which is dependent on turgor. Figure 16.2 *Psilophyton dawsonii*, an enlarged xylem cylinder contributes minimally to flexural stiffness whereas the hypodermal sterome dominates the mechanical contribution to flexural stiffness. Figure 16.3 *Aglaophyton major*, entire stem (TS), arrows, indicate slightly thicker-walled cells of the parenchymatous cortex. Scale bar = 1 mm. Figures 16.4 and 16.5 *Rhynia gwynne-vaughanii*. Figure 16.4 Outer stem and cortex of stem with thicker walled elements in mid-cortex (TS). Scale bar = 0.25 mm. Figure 16.5 Outer stem and cortex, showing epidermis, enlarged cortical elements with thicker cell walls than surrounding parenchyma (LS). Scale bar = 0.25 mm.

A developmental change from parenchymatous cells to fibre elements is a relatively simple novelty requiring little coordinated developmental features. Biomechanical models have focused on early land plants with turgor systems and well-developed hypodermal steromes and it is clear that some plants, such as *Aglaophyton major* and *Rhynia gwynne-vaughanii*, show cortical differentiation with moderately thicker-walled parenchyma towards the outer part of the inner cortex (Figures 16.3–16.5). Kidston and Lang (1917, 1920) refer to a ‘mechanical hypoderma’ comprising the outer cortex which, in both species, consists of larger cortical cells with relatively thin walls. In our observations, thicker-walled and smaller cells are found just to the inside of this zone forming a boundary with smaller cells of the inner cortex and appear more well developed in stems attributed to *A. major* (Figures 16.3–16.5). The origin(s) of the hypodermal sterome among different basal clades is probably complex and involved a range of integrated features to combine both mechanical and photosynthetic functioning of the plant. Photosynthetic tissue would either have had to be placed outside the hypodermal sterome – but somehow communicating with the xylary and translocatory apparatus – or placed on more distal appendages of the branch system where a hypodermal sterome was less completely developed. Interestingly, Kidston and Lang (1917) observe gaps in the outer cortex of *R. gwynne-vaughanii* where the inner cortex extends to areas subtending stomata, suggesting some degree of functional differentiation between possible mechanical, assimilatory and transpirational functioning.

The origins and exact functioning of early hypodermal steromes and their surrounding tissues deserve much more attention from a functional and evolutionary perspective. Are certain kinds of sterome ‘hydraulic’ in function, whereby a layer of thicker-walled parenchyma cells prevented evapotranspiration? Were mechanical steromes modifications of this via wall thickening and elongation into fibre elements?

Whether the hypodermal sterome was initially hydraulic or mechanical in function, derived hypodermal systems are predominantly mechanical and permitted a wider range of branched architectures and almost certainly higher plant growth forms. Branch attachments and wider branch angles are more effectively articulated compared with those of parenchymatous turgor systems. From this perspective the appearance of hypodermal steromes among basal plant groups permitted an increase in architectural diversity in the Lower to Middle Devonian. It is notable that hypodermal systems could have increased the height of not only independent self-supporting plant species, but also contributed to the robustness and height of clonal, mutually supporting upright stems.

Functional studies of the early land plant radiation can be combined with phylogenetic analyses and provide invaluable references for assessing single or multiple origins of functional novelties such as conducting strand tissue, hypodermal steromes and secondary growth. Analysis of Kenrick and Crane’s (1997a) representative cladogram (Kenrick and Crane 1997a; Figure 4.32) and consensus tree of polysporangiophytes (Figure 16.6) indicates an early accumulation of hydraulic and mechanical novelties. Basal groups such as Horneophytosida, *Aglaophyton* and Rhyniopsida show a range of modifications of a possibly homologous central conducting strand, which vary mostly in terms of the types of wall thickening (Kenrick and Crane, 1991, 1997a). These are mostly turgor-maintained mechanical axial systems. A hypodermal sterome, as defined by these authors – present as a single centripetal ring of thick-walled cells several cells thick – is characteristic of eutracheophytes (Euphyllophytes, Lycopsidea, Zosterophylloids and a grade basal to zosterophylls including cooksonioid species) with a loss of the sterome in *Asteroxylon* Kidston and Lang, *Huperzia Bernhardtii* and *Nothia* Lyon. Among Euphyllophytes, basal members such as *Psilophyton* possess an entire sterome, whereas more derived types of sterome

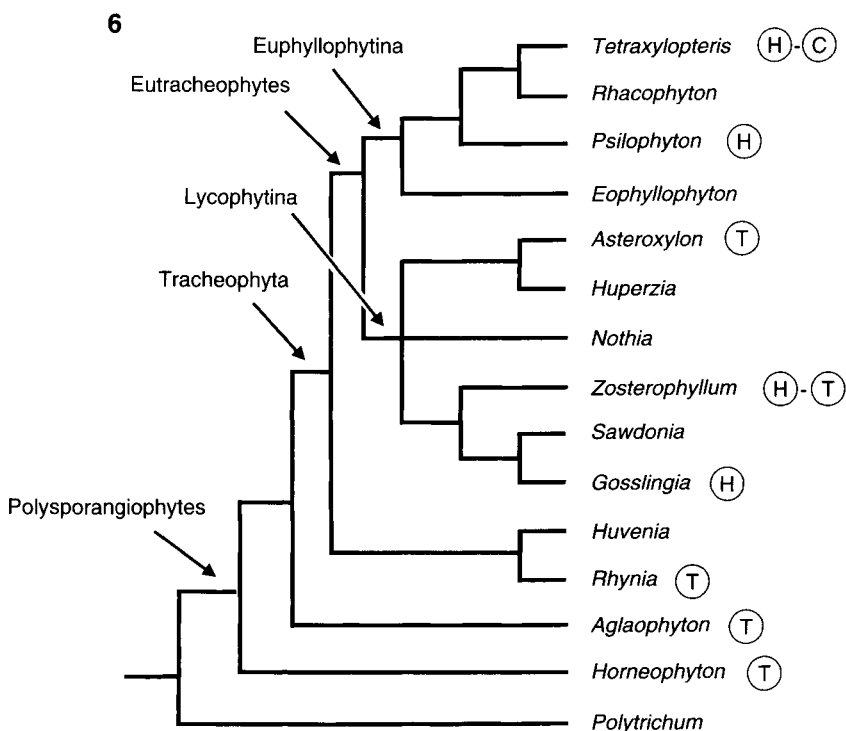


Figure 16.6 Phylogenetic relationships of early polysporangiophytes (based on Kenrick and Crane, 1997a; Figure 4.33). Biomechanical structures are indicated for tested taxa: T = turgor system, H = hypodermal sterome, C = vascular cambium.

characterize both lignophyte taxa and the putative basal sphenopsid *Ibkyia* Skogg and Banks. Finally, a hypodermal sterome and a bifacial vascular cambium characterizes *Tetraxylopteris* Beck and other lignophytes (Figure 16.6).

Secondary growth

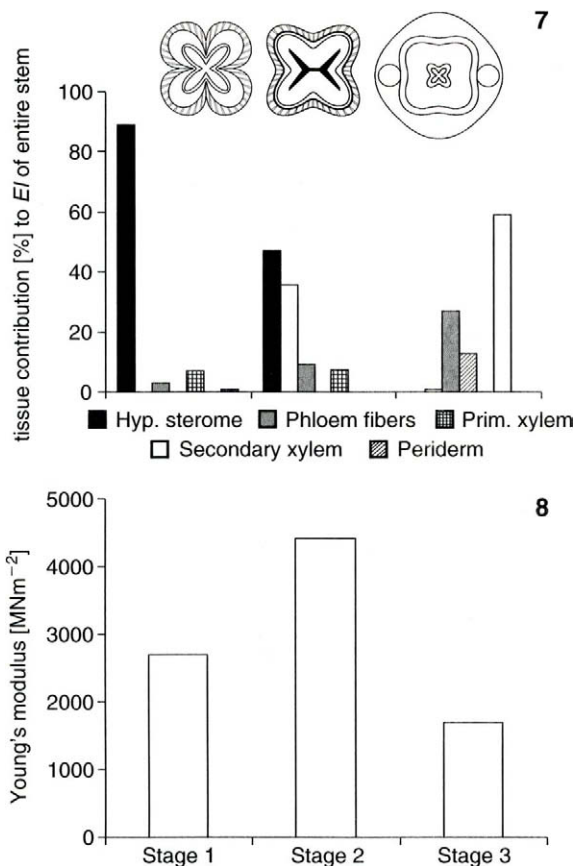
Secondary xylem evolved separately in possibly five different lineages and four of these, lycopsids, lignophytes, *Rhacophyton* Crépin and cladoxylls, had appeared by the Middle Devonian. The vascular cambia generally differ considerably in organization and products formed (Cichan, 1986a; Cichan and Taylor, 1990) as well as in actual function. Biomechanical analyses indicate that wood in arboreous lycophytes is nearly entirely hydraulic in function (Speck and Vogellehner, 1992; Speck, 1994b). Wood in some stem-group lignophytes, such as *Archaeopteris* Dawson, was clearly both hydraulic and mechanical (Galtier *et al.*, 1999). The functional and ecological significance of these separate origins is of extreme interest for understanding the post-homoiohydric phase of diversification in land plants and has been discussed briefly in a broadly biomechanical perspective elsewhere (Rowe, 2000). In this section and below we focus on the bifacial vascular cambium in lignophytes and consider the functional significance of the appearance of secondary xylem in basal members of the group.

Preliminary biomechanical findings based on *Tetraxylopteris schmidtii* Beck indicate that young stages of growth possess a mechanical hypoderm which contributes dominantly to the flexural stiffness of the stem (Galtier *et al.*, 1999; Rowe *et al.*, 2000) (Figure 16.7). A second stage of development where secondary xylem and phloem mostly fill the previously mid-cortical region up to the outer hypodermal tissue (Figure 16.7) shows a higher contribution of secondary xylem to the flexural stiffness of the stem and, by the final stage of development tested, in which an extensive layer of peridermal tissue has formed, the contribution of the wood to flexural stiffness approaches 60% in the model selected (Figure 16.7). Despite the increase in contribution of flexural stiffness of the wood, the pattern of Young's modulus for these three stages is not typical for a self-supporting plant (Figure 16.8) (Speck, 1994a; Speck and Rowe, 1999a,b). Self-supporting plants show basal or older stages of development having higher Young's elastic moduli than more distal segments, which therefore represent relatively stiffer bases for supporting the distal load. In *T. schmidtii*, periderm forms a wide, parenchymatous and disorganized tissue continuing outwards from the secondary phloem (Figure 16.12) (Scheckler and Banks, 1971; Scheckler, 1976). Preliminary results based on a model of ontogeny of *Triloboxylon arnoldii* Matten indicate a similar general pattern in the trend of calculated Young's modulus, although material under investigation of this plant (Stein and Beck, 1983) shows a different pattern of secondary development (Figures 16.9–16.10) where fibre and sclereid reorganization of the hypoderm as well as periderm formation differ from that in *Tetraxylopteris*. The results for *Tetraxylopteris* and *Triloboxylon* will be presented in detail elsewhere.

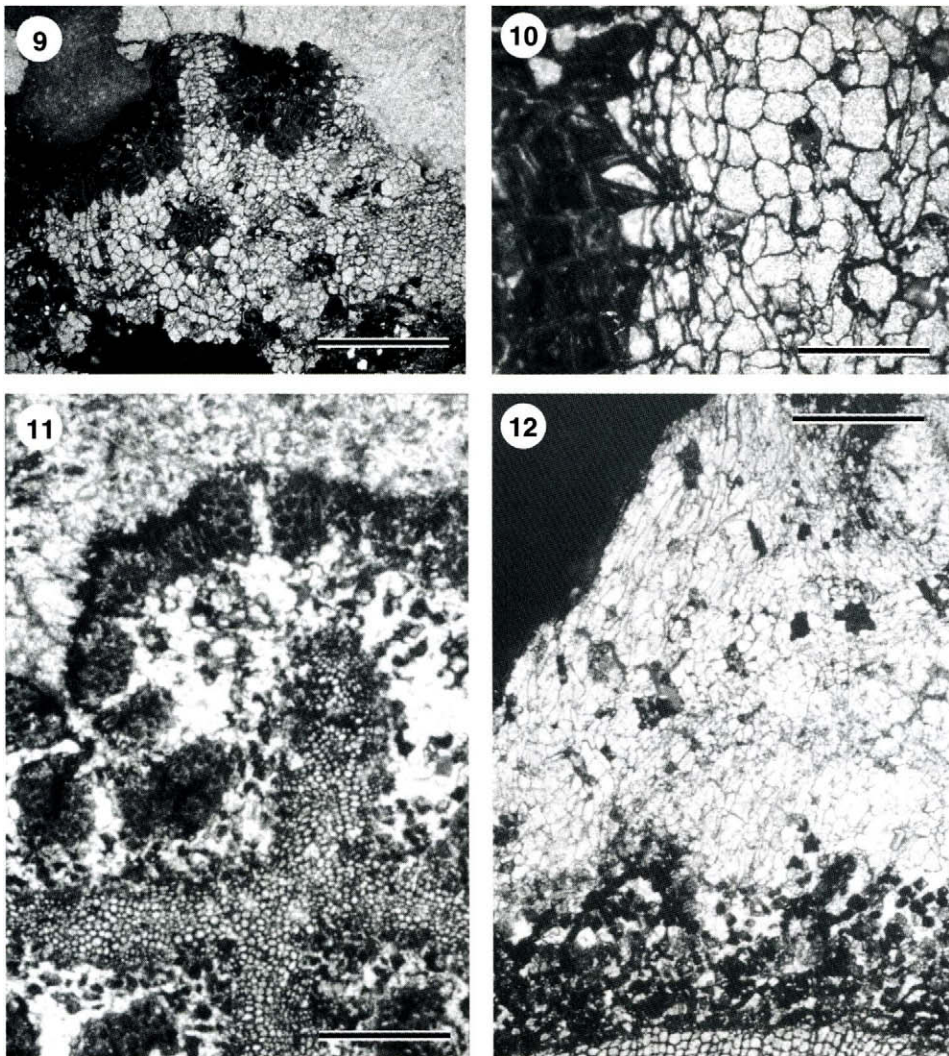
Both *T. schmidtii* and *T. arnoldii* show a hypodermal sterome organization (Figures 16.9 and 16.11) which dominates the mechanical architecture in youngest growth stages. This organization is comparable with some basal land plants relying on a hypodermal sterome for mechanical support. The hypodermal sterome in these aneurophytes consists of longitudinal ribs around the perimeter of the stem in the form of a 'sparganum' cortex differing from the entire hypodermal steromes seen in plants such as *Psilophyton* and *Leclercqia* Banks, Bonamo and Grierson.

Current phylogenetic analyses (Rothwell and Serbet, 1994; Kenrick and Crane, 1997a) most commonly place aneurophytealean progymnosperms at the base of the lignophyte clade. These plants are usually depicted as small statured, possibly understory shrubs, with relatively determinate growth (Scheckler, 2001). Significant development of the bifacial vascular cambium enlarges and modifies the geometry of the primary body and is developmentally complex. Secondary development within the primary plant body clearly requires integral developmental pathways for expansion, containment, prevention of rupturing and sealing-off of the outside of the primary body (Figures 16.9 and 16.12). Neither of the aneurophytes examined shows a secondary growth process, which produces trends in Young's modulus typical of a self-supporting plant. In other words, the production of secondary xylem and secondary phloem fibres in the latest stage of growth does not occupy a sufficiently large proportion of the stem to 'optimize' the stiffness of the basal stem part to a value which would be normal for a self-supporting plant (Speck and Rowe, 1999a). In *T. schmidtii*, there is a wide zone of outer secondary cortical tissue (Figure 16.12) and in *T. arnoldii* there are one or more zones of proliferated primary and secondary cortex (Figure 16.9).

At the risk of generalization, many woody self-supporting gymnosperms and angiosperms have a woody cylinder surrounded by a zone of bark tissue which is relatively thin compared with the diameter of the wood. A relatively stiff woody component



Figures 16.7 and 16.8 Preliminary biomechanical analysis of the basal lignophyte *Tetraxylopteris schmidtii*. Figure 16.7 Contribution of main tissues to flexural stiffness [%] (graph shading: black = hypodermal sterome, white = secondary xylem, grey = phloem, cross hatched = primary xylem, diagonal hatched = periderm). Three tested stages of *Tetraxylopteris schmidtii*, from young to old stages (left to right) are depicted at top. Stage 1, tissues from outside to inside: banded hypodermal sterome, primary parenchymatous cortex, band of primary phloem, primary xylem; Stage 2, tissues from outside to inside: banded hypodermal sterome, thin band of secondary phloem and compressed cortical parenchyma, wide band of secondary xylem, inner star-shaped band of primary xylem; Stage 3, tissues from outside to inside: periderm (mostly parenchymatous), pair of branch traces consisting mostly of xylem and phloem (the model presented here does not compute the branch traces as these are not continuous along stem, depart at wide angles from the axis and so do not contribute continuous longitudinal resistance to the entire stem), thin band of secondary phloem, broad band of outer secondary xylem, band of inner secondary xylem, primary xylem. Calculations are based on centrisymmetric models and data gathering protocols explained in Rowe *et al.*, 1993; Speck, 1994b; Speck and Vogellehner, 1994. Stage 1 is dominated by contribution by the outer sparganum cortex; stage 2 combines relatively high values of both hypodermal sparganum cortex and secondary xylem. Further development of the axis involves loss of the hypodermal sterome and development of a broad and mostly parenchymatous periderm around the wood cylinder; a contribution of wood to flexural stiffness of around 60% is not typical of self-supporting plants in which mechanical support is dominantly provided by the wood cylinder. Figure 16.8 Young's modulus for stages 1–3. The trend observed with a lower value of Young's modulus for the oldest ontogenetic stage compared with younger ontogenetic stages is not typical of self-supporting plants. This corresponds to the loss of the hypodermal sterome, the relatively large proportion of parenchymatous periderm towards the outside of the stem and the consequently relatively central position of the wood cylinder.



Figures 16.9–16.12 Secondary development in aneurophyte progymnosperms (basal lignophytes). Figures 16.9 and 16.10 *Triloboxylon arnoldii*. Figure 16.9 Hypodermal sterome with packets of thick-walled fibre cells and sclereids alternating with parenchyma; secondary development has already developed and accompanied by fissuring of the outermost cortex and at least two types of secondarily produced cortex (Stein and Beck, 1983), scale bar = 0.6 mm. Figure 16.10 Secondary cortex, produced around the periphery of longitudinal fissures and within meristematic zones between the outer secondary phloem and the hypodermal sterome, scale bar = 0.2 mm. Figures 16.11 and 16.12. *Tetraxylopteris schmidtii*. Figure 16.11 Hypodermal sterome comprising alternating packets of thick-walled fibres and parenchymatous tissue, scale bar = 0.5 mm; Figure 16.12 peridermal development in outer part of stage 3 (see Figure 16.7), heterogeneous parenchymatous tissue with scattered thick-walled cells extends from the limit of secondary phloem fibres. Scale bar = 0.8 mm.

occupies a relatively large proportion of older stems and such axes have a correspondingly higher stiffness than younger axes with less wood and more cortex. We cannot overstate the fact that these are initial findings based on the few adequately preserved specimens of this rare plant group, but ones which are, nevertheless, of great interest

for determining the constraints and possible functioning of the appearance of the vascular cambium in lignophytes. We summarize the functional significance of these findings below.

1. Young stages of aneurophytes are mainly supported by a hypodermal sterome of the sparganum type of longitudinal ribs of sclerenchymatous fibres that are fused or nearly fused in young ontogenetic stages. This organization can be viewed as a structural and developmental novelty compared with entire hypodermal steromes of antecedent tracheophytes. The modified function of such an arrangement could be linked to integral processes such as (i) coordination of mechanics with externally placed photosynthetic tissues (as seen in many extant herbaceous plants), (ii) reduced expenditure of physiologically costly thick-walled fibres, or (iii) integral optimization permitting expansion of the outer parts of the primary body as a result of expansion of secondary growth from inside. Whatever the functional innovation operating here, there are several possibilities which would confer developmental innovation over archetypal forms with entire hypodermal steromes.
2. The xylem produced by the secondary vascular cambium, at least in *Tetraxylopteris*, exceeds the original area of primary xylem within the primary body. This is clearly a developmental novelty over other body plans with only primary xylem development. Some authors have interpreted xylem in *Psilophyton* to show possible secondary development as seen from aligned primary xylem elements (Banks *et al.*, 1975). The actual function of the wood cylinder seen in *Tetraxylopteris* and *Triloboxylon* is difficult to demonstrate unequivocally from a physiological, hydraulic or mechanical point of view. The result from the biomechanical models of these plants indicate that the increase in xylem volume *does not* contribute 'mechanically' to a self-supporting growth form. All tested self-supporting plants show an increase in Young's modulus during ontogeny or towards the base of the plant so that older development stages are constructed from a stiffer material (Speck and Rowe, 1999a; Figure 1). Despite the comparatively large volume of wood in the old stage of *Tetraxylopteris*, the Young's modulus for this stage is less than that calculated for younger stages. The larger size and the presence of wood in older stages of *Tetraxylopteris* do produce a stiffer stem in terms of flexural stiffness and this is true for many lianas in which older stages are simply larger (Speck and Rowe, 1999a; Figure 2) but, as in lianas and other non-self-supporting stems, the material properties of the older basal parts of the stem are not stiffer than those younger or above in terms of Young's modulus and this is not an optimal design for a self-supporting plant. Xylem volume increase is possibly some kind of hydraulic novelty but it is difficult to examine further in what precise way. It is unknown, for example, whether the entire wood cylinder remained water conducting throughout development or only in the most recently formed wood. It is possible that an increased xylem cylinder either optimized conductivity over a longer time or over more extended aerial portions of the plant or both. Uncertainty of how much of a wood cylinder remained conductive and, indeed, how much of the distal part of the fossil plant was supplied, remain major obstacles for models of hydraulic functioning in fossil plants (Cichan, 1986a; Rowe and Speck, 1998).
3. Cortical proliferation and peridermal activity also represent developmental novelties compared with most earlier tracheophyte body plans. Evidence exists of localized cellular proliferation in plants lacking other secondary growth and this is sometimes referred to as 'wound tissue' (Banks, 1981) such as in *Psilophyton*. This is quite different

- (though possibly developmentally allied) with fully circumferential periderm growth coupled as integral developmental traits with bifacial vascular cambium activity. Periderm and cortical development in *T. schmidtii* and *T. arnoldii* appear to ‘surround’ and ‘seal off’ respectively the geometric change brought about by secondary growth. They could justifiably be cited as examples of integral functional novelties formed *de novo* and coupled with expansion of the wood and secondary phloem. Interestingly, the peridermal systems observed in *Tetraxylopteris* and *Triloboxylon* are different (see Figures 16.9–16.12) suggesting that integral pathways linked with secondary development of the bifacial vascular cambium were derived independently. Once again the functional origin of periderm is an interesting one of possible co-option where an original wound repair mechanism possibly related to phytophagy was later modified and deployed as an integral developmental system linked to growth of the bifacial vascular cambium. Certainly this is hinted at by the cortical activity in *T. arnoldii* where at least two broadly defined meristematic areas occur just inside the outer hypodermal sterome and flanking longitudinal fissures caused by internal expansion of the axis (see Figure 16.9).
4. Secondary phloem proliferation and significant volumes of tissues in both species suggest prolonged translocatory function over a more extensive plant body. Fibre cells within the secondary phloem might have had a mechanical role, but rows of fibres are not interconnected and contain rounded sclereids, which are not as mechanically efficient as fibres.

Does the appearance of the bifacial vascular cambium represent a key innovation?

More derived progymnosperms, Archaeopteridales, co-occurred with aneurophytes, became globally distributed and developed additional vegetative novelties including leaves, complex branch organization and large-bodied architectures (Scheckler, 2001). Lignophytes from the Late Devonian onwards possess a bifacial vascular cambium and show increasing architectural diversity. From this viewpoint, the combination of novelties built around the bifacial vascular cambium represents further consolidation of the lignophyte body plan and accompany a morphological and architectural diversification. In terms of the long-term success of lignophytes, the implication that the bifacial vascular cambium was a key innovation triggering a morphological radiation of lignophytes must be considered with caution. The appearance of the seed habit – perhaps a clear example of a modular innovation in relation to vegetative novelties associated with secondary growth – is another of the ‘key innovations’ widely believed to have contributed to the radiation of the group.

The bifacial vascular cambium emerged prior to the seed. We have shown that, among at least two representatives, cambial activity might have optimized hydraulic conductivity and photosynthate translocation. Both are physiological novelties compared to basal groups with a more static primary growth trajectory.

Phylogenetic resolution of basal lignophytes and seed plants remains unsubstantiated in any great detail. However, biomechanical analyses of some early seed plant representatives indicate that some and probably many show body plans with a vascular cambium in which secondary growth is mainly confined within the primary plant body and that mechanical stiffness was conferred by a hypodermal sterome of the sparganium or dictyoxylon type (Speck and Vogellehner, 1992, 1994; Rowe *et al.*, 1993). Other early seed plants following the Late Devonian were undoubtedly capable of woody self-supporting

growth habits (Speck and Rowe, 1994a). Interestingly, the earliest known seed plants are small-bodied plants with limited secondary growth but extensive leaves (Rothwell and Scheckler, 1988; Serbet and Rothwell, 1992), a later novelty which could be interpreted as an innovation added to the cambial body plan organization observed in aneurophytes. Furthermore, the possession of planated leaf surfaces could be considered as integrated novelties linked to the enhanced hydraulic potential of cambial activity. The appearance of seed structures represents a modular innovation in relation to the bifacial cambium body plan.

In summary, the appearance of the bifacial vascular cambium appears to have been an innovation for lignophyte architecture with a primary functional novelty concerned with optimizing hydraulics rather than immediately conferring mechanical advantages as witnessed with the limited stature of tested aneurophytes and the biomechanical results suggesting poor optimization for self-supporting habits. Self-supporting architectures are seen among Archaeopteridales and other putative progymnosperms, such as *Protospitys*, and it is a fascinating aspect of the early evolution of lignophytes that this lignophyte stem group became largely extinct by the early Carboniferous – apart from one possible Lazarus taxon *Cecropsis lunulata* Stubblefield and Rothwell (Stubblefield and Rothwell, 1989). Only members of the clade having the seed habit, an unrelated additional modular innovation, diversified after this point.

Lignification and biomechanics of the plant cell wall

Chemical and physical complexity of plant phenolics and especially lignin have been of particular interest to evolutionary studies, where their primary roles as either light screens, phytophagy deterrents, water-repellents or mechanical stiffening have been discussed recently (Cooper-Driver, 2001). In addition to the difficulties of understanding what lignin actually does in living plant cell walls, is the notorious problem of unequivocally identifying lignin derivatives in fossil plants, especially in the early land plant radiation. There are several elements to the potential functional roles of lignin in plant tissue mechanics. First is its potential role as a 'water proofing agent'; as a hydrophobic molecule it 'chemically dries' the embedded cellulose microfibrils which can maintain more intramolecular H-bonds and so retain higher stiffness. Second, as a matrix structure or filler (Wainwright *et al.*, 1976; Niklas, 1992) in composite cell walls, it can only act mechanically in combination with longitudinally orientated molecules, such as cellulose. These two functions of lignin discussed in the recent experimental literature (Hoffmann *et al.*, 2000a,b; Spatz and Speck, 2000) are significant for interpreting the origins of such roles in early land plants. Retaining stiffness in water-conducting cell walls is important to prevent collapse resulting from negative pressures building up inside the lumen; it is possible that lignin might have played a crucial role at some point in the evolution of water-conducting strands where lignin maintained stiffness in cell walls constantly in contact with water. Somewhat different constraints might have characterized lignin functioning in hypodermal steromes. As these tissues do not conduct water after cell death they might have been less reliant on lignified cell walls for protecting cellulose wall-stiffening elements from contact with water. The role of lignin in strengthening water-conducting cells is possibly linked with other geometric features of the cell wall including spiral, annular and scalariform thickenings, which can also locally reinforce and strengthen water-conducting cells against negative pressure.

Recent biomechanical research on a wide systematic range of plants from algae to seed plants is one way of observing how different clades deploy varying mechanical strategies at the architectural, morphological, anatomical, ultrastructural and biochemical levels (Spatz and Speck, 2000). In this case study we briefly report some recent results from studies on *Equisetum* L. where mechanical support is provided from non-lignified fibre cells.

Equisetum hyemale L., is a herbaceous, clonal plant with a hollow stem, nodal structure and a mechanical architecture which provides a lightweight, ‘cheap’ organization that resists mechanical failure by local buckling (Speck *et al.*, 1998). In *Equisetum hyemale* ovalization of the stem and local buckling are resisted by relatively high Young’s elastic moduli of the hollow stem wall. The wall includes a non-lignified hypodermal sterome in combination with a double layer of relatively thick-walled endodermis cells with lignified casparian rings (Figure 16.13). Detailed staining techniques with phloroglucinol (Gerlach, 1984) and other reagents indicate that cells of the hypodermal sterome are living and unlig-nified (Figure 16.13). This tissue has Young’s moduli varying from 4.5 GPa in basal inter-nodes to lower values of 1.5 GPa towards the apex (Speck *et al.*, 1998). All values in these data are lower than the usual range for sclerenchyma which range in values from 5 to 30 GPa (Wainwright *et al.*, 1976; Niklas, 1992) and higher than values measured for collenchyma ranging from 1 to 2 GPa (Nachtigall *et al.*, 1988; Spatz and Speck, 1995).

Unlignified tissues can provide significant mechanical stiffness approaching values to those of lignified tissues. One of the uncertainties of any kind of biomechanical modelling of fossil plants is that biochemical composition of cell walls is unknown or unknow-able and that measurements on living plants might not be confidently applied to fossils (Edwards *et al.*, 1997). Of course, cell wall chemistry, ultrastructure and degrees of hydra-tion as well as numerous other factors influence cell wall mechanical properties and values of Young’s moduli assigned to fossil tissues can only be hoped to represent a similar mag-nitude of value. The point we wish to make here is that lignification is not a prerequisite for conferring ‘stiffness’ to a plant tissue and that a thick-walled tissue comprising fibre-shaped elements would not necessarily require lignin in its walls to have a high value of Young’s modulus. This has direct implications for interpreting the functioning of hypodermal

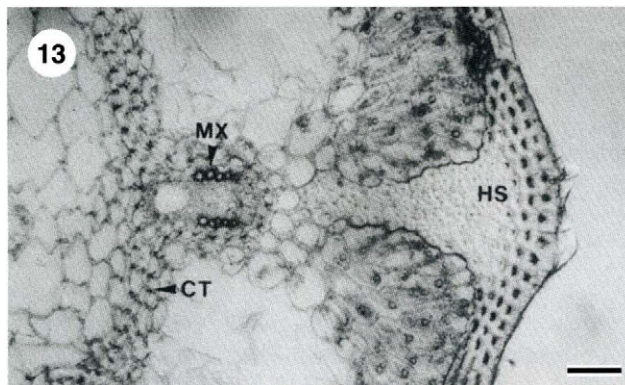


Figure 16.13 Transverse section of *Equisetum hyemale*, stained for lignin, based on Speck *et al.* (1998). A segment of dense thick-walled tissue representing a hypodermal sterome (HS) represents an important contribution to the mechanical stability of the stem but is unlignified; other lignified tissues in the same stem cross-section include the metaxylem (MX) and the casparian thickenings (CT). Scale bar = 0,1 mm.

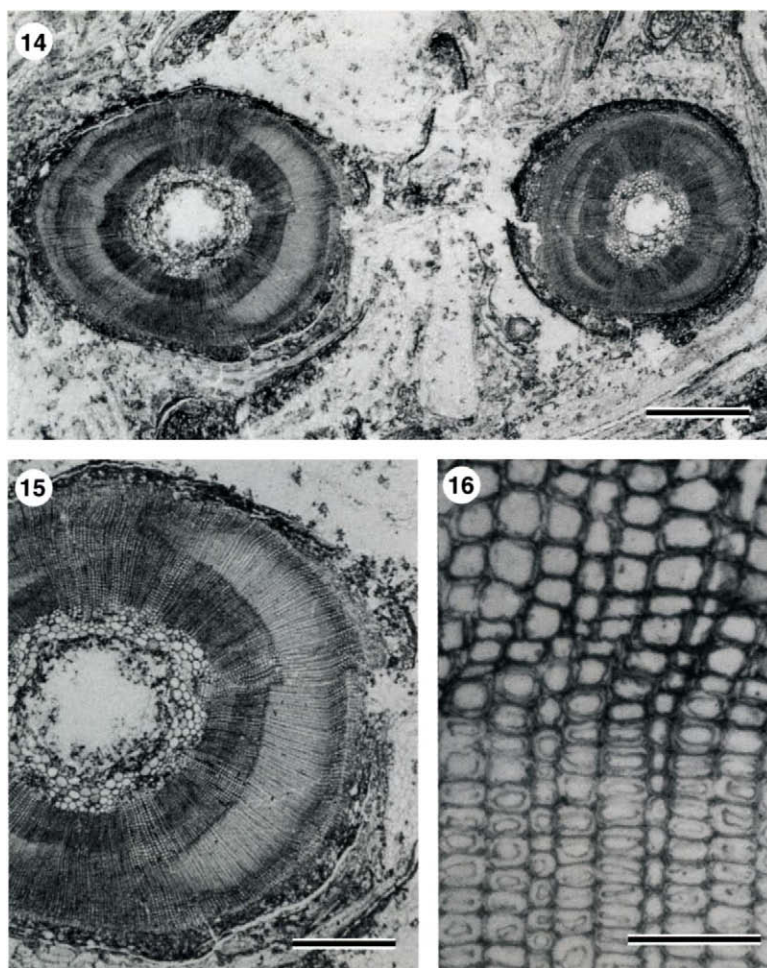
steromes in early land plant diversifications where lignification of cell walls would not be prerequisite for a mechanically significant structure. The physiological and mechanical roles assigned to phenolics in general, from light screens, pathogen/phytophage deterrence, water repellent and mechanical properties of cell walls underline the difficulty of confidently assigning function to biochemical processes and such areas undoubtedly require more research based on carefully deployed studies of living plants.

Reaction wood

Reaction wood is a developmental modification of normal wood concerned with righting, adjusting and maintaining the orientation of woody leaders and branches (Archer and Wilson, 1970; Wilson and Archer, 1977). Two types of reaction wood exist, including compression wood in extant conifers and tension wood in angiosperms. Both show cellular characteristics differing from normal wood and often occur in crescent-shaped areas within the wood cylinder. Generally, compression wood is often, but not always, found on the underside of conifer lateral branches and tension wood is often found in the upper side of angiosperm branches.

The fact that reaction wood is so widespread and common in extant trees and shrubs, prompted some workers to consider when it first appeared in the evolution of woody plants (Timell, 1983, 1986). Considering the large body size and architectural complexity of woody lignophytes since the Devonian and the dependence of extant woody architectures on reaction wood, this is an interesting question. Was compression wood present since the appearance of Middle Devonian plants such as *Archaeopteris* or was it a more derived feature that was 'added' to an increasingly complex and derived system comprising the bifacial vascular cambium? Timell (1983) located only low to moderate wood heterogeneity in *Callixylon* wood and nothing that he could confidently ascribe to reaction wood. His survey of wood illustrations from the Permian fossil woods of eastern Europe (Greguss, 1967) also revealed difficulties in definitely establishing the presence of reaction wood.

Re-investigation of cordaite shoots of *Pennsylvanioxylon tianii* from the Early Permian of China (Baolin and Wang, 1988; Li, 1995; Baolin *et al.*, 1996) and *Pennsylvanioxylon* from the Permian of France, indicate crescent-shaped zones of modified wood within several small branches (Figures 16.14–16.16). One branch pair includes reaction wood, on opposite sides of each branch. Areas of reaction wood are sharply differentiated with transitions to normal wood (Figure 16.15). Cell wall outlines of individual tracheids within crescent-shaped areas have more elliptical outlines, thicker walls and a different type of preservation in which the wall content between the outline of the lumen and the outer part of the secondary wall is usually empty (Figure 16.16). Opposite and temporally alternating areas of reaction wood are a well-known phenomenon in extant compression wood formation during 'over-correction' of upright or lateral branches (Timell, 1986). In cases of over-correction, physiological and mechanical reactions in one direction exceed the 'programmed' equilibrium position for the orientation of the stem and result in the formation of reaction wood in the opposite direction to readjust the position of the branch. This finding suggests a relatively late appearance of reaction wood among lignophytes. It remains to be seen whether reaction wood can be detected in other seed plant groups or among progymnosperms. Analyses of *Archaeopteris*, *Tetraxylopteris* and *Triloboxylon* as well as wood of putative seed plants *Pitus* Witham, *Endoxylon* Kidston and *Eristophyton* Zallesky have not yet yielded the levels of differentiation found among these cordaites.



Figures 16.14–16.16 *Pennsylvanioxylon tianni*, a late Permian cordaitalean plant from China. Figure 16.14 A pair of twigs showing at least three crescent-shaped developments of reaction wood represented by the lighter areas of wood. Reaction wood is organized in the same (left to right) orientation for both axes and is very similar to opposite reaction wood formed in extant conifers, scale bar = 1.2 mm. Figure 16.15 Crescent-shaped area of reaction wood alternating with segments of normal wood, scale bar = 0.6 mm. Figure 16.16 Transverse section of transition between reaction wood, below and normal wood above. As in many instances of compression wood in extant plants, the compression wood consists of tracheids with thicker cell walls and elliptical outlines. Scale bar = 80 μm.

Whether reaction wood will be identified in basal groups of lignophytes or not, it represents a physiological and mechanical novelty superposed on the vascular cambium body plan. This highlights a crucial aspect of the vascular cambium as a developmental plan that can accommodate added functional novelties. Reaction wood is clearly mechanical in function in extant plants and almost certainly so for cordaites. Interestingly, recent studies on extant conifers indicate that conductance might be reduced by compression wood (Spicer and Gartner, 1998) suggesting that the appearance of reaction wood might have been of mechanical advantage for the development of complex architectures but with some integrative burden and ‘trade-off’ with stem hydraulics.

Hydraulics, mechanics and evolution of the climbing habit

Types of climbing strategy

Climbing plants can show quite different mechanical and hydraulic constraints from self-supporting plants (Gartner, 1991; Putz and Holbrook, 1991; Ewers *et al.*, 1991; Speck, 1991, 1994b; Speck and Rowe, 1999a). Furthermore, different types of climbing strategy also show quite different trends in mechanical and hydraulic properties during development. Recent and ongoing biomechanical and hydraulic analyses have shown that climbing strategies involve a range of interrelated constraints based on hydraulics and mechanics. A plant that has reduced the physiological cost of mechanical support by producing slender stems needs to augment its hydraulic conductance if it is to maintain a comparable photosynthetic surface. This is often accomplished via large-diameter conducting elements; water conductance, generally speaking, scales to the fourth power of the radius of the conducting element. However, slender stems with large-diameter conducting elements supplying a large leaf area can be a risky business. Climbing stems can be easily damaged by host tree movement, canopy sway, tree falls, host branch failure from epiphyte loading and so on. A relatively small amount of damage to the stem in terms of surface area because of damage by excessive bending or torsion could potentially cut off a large portion of hydraulic supply; this situation is made worse by the fact that large diameter elements are more prone to dysfunction on account of cavitation (embolism) and that relatively limited damage such as localized cracks passing in the vicinity of vessels might be enough to cause embolism and hydraulic dysfunction (Putz and Holbrook, 1991).

At the risk of over generalization, there are basically three different strategies to overcome these conflicts involved in minimizing stem diameter and maximizing stem conductance. First, grow very close to or actually fixed along the surface of large-bodied supports and reduce free movement between the host support and the climber (root climbers, e.g. Araceae, *Hedera* L.). Second, protect a non-self-supporting stem with stiff mechanical properties, these plants (sometimes referred to as semi-self-supporters (Speck, 1994a,b; Speck and Rowe, 1999a) can exceed their stable critical buckling height and maintain a relatively 'loose' more-or-less vertical scrambling position in the surrounding vegetation. Third, attach firmly along host plants by stem twining, winding and tendrils etc. and produce specialized compliant stems, which dissipate mechanical energy under bending or torsional stresses. This latter strategy can increase survival of the attached plant during swaying and movement of the host as well as raising the probability of survival during dramatic rearrangements of the surrounding forest structure during tree falls. Arguably the most complex strategy in terms of hydraulics and mechanics is the third kind of strategy which is found among many families of lianoid angiosperms and the gnetalean genera *Gnetum* L. and *Ephedra* L.

Appearance of the lianoid habit

The possibility of elucidating specific growth habits of long extinct organisms and the implications for interpreting community structure and evolutionary patterns is appealing. Progress has been made in recent years with the discovery of cuticular plant compressions with attachment organs from the late Palaeozoic indicating probable climbing strategies of the first type described above (Kerp and Krings, 1997; Krings and Kerp, 1997, 1999) as well as biomechanical models indicating climbing strategies of the second type (Speck and Vogellehner, 1992; Rowe *et al.*, 1993; Speck, 1994b; Speck and Rowe 1999b). In addition to these are the quantitative hydraulic studies, particularly of Michael Cichan,

investigating the potential hydraulic traits of a range of fossil plants discussed in reference to climbing habits. These include sphenophylls (Cichan, 1985) and sphenopsids, (Cichan and Taylor, 1983; Cichan, 1986b), and medullosans and cordaites (Cichan, 1986a). Other recent discoveries include remarkable vessel-like structures among gigantopterid seed plants (Li *et al.*, 1996), which have also been suggested as representing climbing plants.

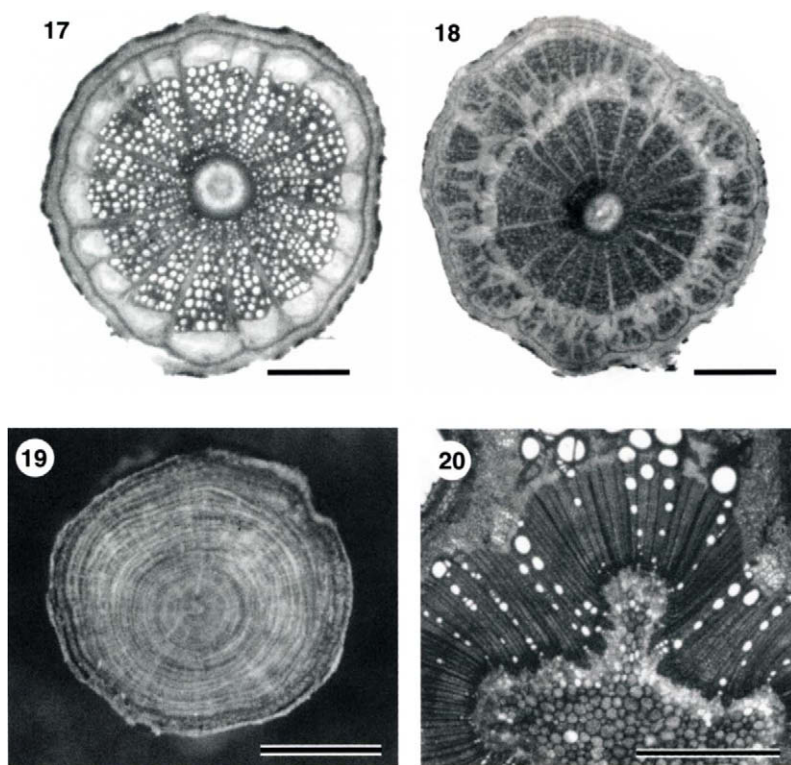
In many forms of lianoid development there is an ontogenetic shift from young, relatively stiff stems with dense wood and low specific conductance, to larger diameter, relatively flexible stems with high specific conductance (Figures 16.17–16.20). Old stages of many lianas also have high degrees of built in toughness (not to be confused with stiffness), where stems can be bent or twisted to a remarkable extent before becoming mechanically and hydraulically dysfunctional (Putz and Holbrook, 1991). This is often accomplished by predisposed segmentation via alternation of hydraulically active tissues with wedges of deformable and fracture-confining tissues (Figure 16.18). During early development, mechanical stiffness is a prerequisite for searching leaders or self-supporting young individuals (Speck and Rowe, 1999a). Following secure attachment to the host, increased hydraulic optimization, seen in large surface areas of large diameter vessels, is combined with mechanical compliance and segmentation of the wood cylinder. Our examples here serve to illustrate the complex developmental patterns in a species of the angiosperm *Bauhinia* L. and the gymnosperm *Gnetum*. Mechanical data based on Young's modulus measured in bending for the two species illustrates that both species show a significant drop in Young's modulus during ontogeny.

Vessels and the climbing habit

Enhanced hydraulic conductance via vessels can be viewed as a functional novelty compared with the type of organization seen among most basal lignophytes and non-lignophytes, most of which show little differentiation of tracheid elements in terms of wide size variation. Like the appearance of reaction wood discussed above, vessel appearance could be seen as another supplementary novelty to the existing bifacial cambium body plan. The appearance of vessels has often been cited as a specialization, whereby hydraulic and mechanical functioning within the wood cylinder could be performed by more specialized cell types within the same tissue (Carlquist, 1975).

It is not clear whether, climbers with lianoid (type 3 climbing strategy) existed in early or later Palaeozoic ecosystems. Certain forms such as *Sphenophyllum* Koenig, *Medullosa* Cotta and cordaites have been observed to combine large diameter xylem elements with slender stems, but biomechanical studies are yet to provide evidence whether such forms comprised biomechanically sophisticated flexibility and toughness or adopted the arguably more conservative mechanical strategies such as root climbers or relatively rigid 'semi-self-supporting' types mentioned above. Xylem element diameter taken on its own is not a clear indicator of a lianoid habit.

The appearance of wide conducting elements and vessels in the lignophyte body plan could have potentially elevated leaf surface to stem diameter ratios. The three types of climbing habit outlined above would require different integral and modular novelties. For example, semi-self-supporting plants produce and retain stiff stem mechanics, which would require either stiffening elements in the wood cylinder or in the cortex. This situation contrasts with the necessary integral novelties associated with lianoid climbers, which require compliancy and toughness. Lianoid features arose independently many times among angiosperms and with many variations in terms of ways in which the older stages of development



Figures 16.17–16.20 Anatomy and organization in extant lianas. Figures 16.17 and 16.18 Anatomy in young and older stems of the gymnospermous liana *Gnetum africanum*. Figure 16.17 The young stage has relatively dense wood with increasing numbers of vessels towards the outside of the wood cylinder, scale bar = 2.5 mm. Figure 16.18 The older stage shows segmentation of the wood cylinder via variant cambium development, which has formed segments of secondary phloem and cortical tissue around and between the second ring of wood formed. This type of organization is similar to many angiosperm lianas and is typical of older stem development showing high compliance and protection of the hydraulic system via fracture-limiting areas between xylem segments, scale bar = 5.0 mm. Figures 16.19 and 16.20 Transverse sections of the angiosperm liana *Baubinia guianense* (see also Speck and Rowe, 1999a). Figure 16.19 Young developmental stage of a self-supporting individual prior to the climbing phase. The young growth has produced dense wood with many fibres and very few vessels, scale bar = 3.0 mm. Figure 16.20 Anatomical transition marking shift from young self-supporting phase to lianoid phase and a shift from dense stiff wood to a second wood type with many large vessels. In this species, the vascular cambium is limited to growth in two areas of the perimeter forming a ribbon-shaped lianoid stem, in this figure the continued growth of lianoid wood is at the top. Scale bar = 0.85 mm.

produce flexibility and partition their hydraulic system for protection against hydraulic failure following mechanical perturbation (Gentry, 1991; Ewers *et al.*, 1991; Speck, 1991, 1994a; Speck *et al.*, 1996a; Speck and Rowe, 1999a). Close inspection of any modern habitat, temperate or tropical, indicates enormous differences in position, volume, length, branching, shading and many other characteristics between root climbers, self-supporters and lianas. Dense swathes of semi-self-supporting ferns of *Gleichenia* Smith in the tropics, for example, up to eight or so metres high occupy an entirely different niche from lianoid forest climbers which produce the bulk of their leaves in the canopy via stems that

are entirely dependent on self-supporting hosts. These kinds of differences are detectable from consideration of stem mechanics and hydraulics of fossil plants. A putative fossil liana with wide xylem elements, but with a thick fibrous hypodermal sterome around the periphery of old stages of growth can not have been a flexible lianoid climber but more probably a semi-self-supporter.

Since the earliest land plant radiations, the different kinds of potential climbing strategy we have outlined above would have required similar mechanical functioning as plants today involved in attachment and stem stiffness and compliancy. If the entire suite of characters contributing to a climbing syndrome is viewed as a complex character, clearly some climbing strategies are more complex than others.

Growth forms requiring minimal associated novelties probably include root climbing and semi-self-supporting growth forms where increased hydraulic capacity could be accompanied by either retainment of existing mechanical tissues, e.g. the hypoderm for stem stiffness (semi-self supporters) and repositioning of rooting meristems or other attachment organs on cauline parts of the stem (root climbers). Both of these types of climbing strategy could be attained by simple body plans without secondary growth, with relatively little organizational rearrangement and few additional novelties. Heterotopic expression of rooting meristems on stem axes could provide root climbers with the necessary anchorage mechanism. Given adequate hydraulic supply, semi-self-supporters would probably require even fewer additional novelties. If growth was confined to the primary body and primary tissues of the hypodermal sterome, simple continuation of apical meristems would rapidly exceed the plant's critical buckling length, the plant would no longer be 'self-supporting' and would then require support from a dense stand of similar neighbours perhaps of the same clonal stand or larger bodied self-supporters. Simply leaning or interlocking branches to maintain a relatively stiff plant stem upright would not require sophisticated attachment devices and, indeed, many modern tested semi-self-supporters simply interlock branches or petioles with the surrounding vegetation to stay upright (Gallenmüller *et al.*, 2001).

At least these two 'types' of climbing strategies require relatively little modification from bauplans based on turgor, hypodermal or lignophyte body plans. It is possible that such forms were appearing relatively early in the land plant radiations and that such climbing growth forms were just a short physiological step away from the underlying thigmomorphogenetic economizing (Jaffe, 1973) that a great many green plants appear inherently to possess. A shift to a root-climbing or semi-self-supporting growth form would not require a large input of secondary novelties. A number of early land plants do possess both horizontal axes with rhizoids as well as upright aerial axes and this organization might be relatively predisposed towards at least potentially facultative root-climbing, if not on host plants then on local ground topography. Root climbing is possibly more amenable to turgor systems as once firmly attached to a solid host support there is little requirement for stem stiffness or toughness, indeed many root-climber stems are actually brittle structures. Semi-self-supporting strategies among turgor systems were probably more limited as local bending moments on a cylindrical turgor system from greatly extended upright stems may be prone to mechanical and hydraulic failure. It is possible that 'climbing' strategies among such forms were more limited to optimizing and opportunistically modulating safety factors (variable stem lengths *below* the critical buckling length) than greatly developing long semi-self-supporting axes (Speck and Vogellegner, 1994).

The appearance of the hypodermal sterome is often talked about as being an innovation for stem stiffening among early land plants and for optimizing self-supporting growth habits (Niklas, 1992; Speck and Vogellegner, 1994; Rowe and Speck, 1997).

In fact, the appearance of the hypodermal sterome could have also facilitated climbing semi-self-supporters by imparting sufficient integral stiffness in stems exceeding their critical buckling length but laterally supported by other supports. Evolutionary transitions between self-supporting and semi-self supporting among plants with simple body plans therefore probably require little added novelties and little extra integrated developmental characters.

What is special about lianas?

The mechanical and hydraulic properties present in lianoid climbing habits require numerous integral and modular novelties superimposed on a basal early land plant or early ligno-phyte body plan. Like other climbing growth forms an optimized hydraulic supply and vessel-sized elements is probably essential to maintain hydraulic supply to a large effective leaf surface via a narrow stem. Essential mechanical features include stiff young stages as searchers and increasing compliancy during development associated with stem toughness. This mechanical shift in properties is an essential feature and one that distinguishes the lianoid habit from root climbing and semi-self-supporting strategies. A basal lianoid body plan would require relatively high levels of integral novelties including optimized hydraulic conductance, compliance and toughness.

Conclusions

Following the establishment of basic embryophyte organization and physiology (Graham *et al.*, 2000), the early phase of terrestrialization concerned mostly hydraulic novelties and the establishment of physiologically homoiohydric body plans. Columnar growth forms depending on physiologically maintained turgor were hydraulically optimized by outer cutinized sheaths and central conducting strands. Both represented primary hydraulic novelties. Biomechanical studies demonstrate that many early conducting strands did not contribute directly to the bending mechanics of the stem and thus clarify that mechanical support of columns relied on physiologically mediated water maintenance. The central position of the xylem tissue and its relatively small contribution to the second moment of area of the stem indicate a small direct contribution to flexural stiffness of the stem (see Figure 16.1). It is possible that xylem strands and cutinized sheaths might, however, have been of secondary mechanical significance in ensuring turgor and hydraulically stable axes for larger bodies or during intermittent water availability (e.g. Bateman *et al.*, 1998).

Representatives of basal clades show of a range of hypodermal steromes. Biomechanical analyses indicate that the hypodermal sterome became important for contributing directly to the stiffness of the stem and represented a functional novelty for larger and more diverse architectures. Mechanical hypodermal steromes were important in optimizing self-supporting growth forms as well as semi-self-supporting forms. There is some indication that cell wall thickening in the mid to outer cortex in some early representatives such as *Rhynia gwynne-vaughanii* and *Aglaophyton major* may have initially been concerned with limiting evapotranspiration and had little or no effect on directly contributing to stem stiffness. It is possible that such steromes could have secondarily optimized mechanical stability by better controlling turgor pressure over larger and/or taller body plans. Further functional analyses coupled with more inclusive phylogenetic histories might indicate that 'hypodermal steromes' were primarily hydraulic novelties co-opted for mechanical support.

Basal lignophytes possess an undoubtedly mechanical hypoderm in young stages of growth, which are fused or nearly fused in young ontogenetic stages. Development of the bifacial vascular cambium conferred several functional breakthroughs including increased hydraulic, mechanical and translocatory functions. Biomechanical studies suggest that large surface areas of cortex at the 'mechanically influential' outside of the stem meant that the relatively centrally positioned wood cylinder contributed little to flexural stiffness of the stem and the products of the secondary vascular cambium were not optimized for a mechanically self-supporting stem system.

Whereas many authors agree that the Siluro-Devonian radiations were periods of 'innovation' characterized by appearances of many new physiological and morphological 'novelties', precisely how these new novelties actually functioned is more easy to speculate on than to test empirically. This is undoubtedly true in terms of 'hydraulic' or 'mechanical' functioning. A major part of the problem lies in the potentially rapid integral functioning of hydraulic and mechanical features among these early body plans. Conducting strands may have represented *secondary* mechanical novelties by increasing turgor potential over a larger or taller plant body; thick-walled hypodermal sterome fibres might represent modified cortical cells initially concerned with restricting evapotranspiration; secondary xylem acting initially as hydraulic optimization became mechanically important for self-supporting architectures after the addition of further integrated novelties such as entire periderm formation. Further additional novelties to the cambial body plan then permitted yet greater diversification of growth forms via novelties, such as reaction wood and vessels.

The appearance of the bifacial vascular cambium as seen in basal aneurophytalan lignophytes provided a developmental template for a wide range of further physiological and structural novelties including indeterminate, self-supporting as well as non-self-supporting woody architectures. Reaction wood could be said to represent one of the 'bells and whistles' or additional novelties added to the basic cambial body plan, which can be perhaps more directly interpreted as a purely mechanical novelty. The evidence from observed aneurophytes and Archaeopteridales – up to present – indicates that reaction wood was not present in these basal lignophytes. Reaction wood, is an example where a functional novelty is more easily ascribed a single function, compared with more primordial novelties more basic to the emerging polysporangiophyte and tracheophyte body plan where emerging developmental templates share a range of functions.

Phylogenetic investigations of recent radiations, which include functional observations based on living plants, potentially carry higher expectancies for resolving evolutionary processes. A combination of molecular and morphological character traits as well as empirical experimentation on the physiology, hydraulics and mechanics can offer more robust frameworks for interpreting correctly the significance of novelties, innovations and adaptations. Functional studies combined with phylogenetic frameworks can more easily address patterns in hydraulic and mechanical developmental strategies such as transitions between self-supporting plants and climbers. Derived seed plants such as Gnetales and angiosperms have complex and integrated hydraulic and mechanical novelties adapted for lianoid climbing strategies.

Such studies are demonstrating whether transitions in growth form during evolution from self-supporters to climbers occurs via hydraulically and mechanically 'intermediate' forms (e.g. semi-self-supporters) and whether a clade that has once established a lianoid body plan can revert back to a self-supporting architecture (Speck *et al.*, 1997; Civeyrel and Rowe, 2001). In the latter case ongoing work is indicating that the hydraulic and mechanical novelties accumulated during the development of climbing body plans carries

high degrees of developmental and ecological burden which is rarely reversed and that subsequent 'self-supporting' plants show differences in trends of mechanical or geometrical characters.

Developmental burden and the ability to 'turn back' having accumulated a specialized mechanical and hydraulic body plan is a fascinating aspect of plant evolution and one which might be relevant to many patterns of body plan complexity and intrinsic constraint from the Devonian radiations onward. The rise and decline of the lycopsid, sphenopsid and filicopsid clades are monumental examples of rises in complexity, long-term ecological success, potential intrinsic constraint and canalization and, finally, disastrous extinction. We argue that an increased application of biophysical modelling on such topics based on parallel studies of living plants that can be appropriately applied to fossils can potentially offer much in the way of explaining or at least reducing the number of speculative scenarios.

Key innovations are widely believed to initiate major radiations and observers are surely correct in emphasizing the necessary phylogenetic framework for testing potentially conflicting or alternative hypotheses based on phylogenetic pattern rather than simply mapping *a priori* assessed functional related traits (Bateman, 1999b). The above examples illustrate the wide range of data-related and conceptual difficulties centred on interpreting function, let alone interpretation of higher conceptual levels such as innovatory significance and adaptation. We propose that function-level interpretations lie at the core of any study involving assessments of innovation and adaptation.

A recurring theme in identifying key innovations in plants is the significance of vegetative and architectural novelties as opposed to reproductive novelties. Was the bifacial vascular cambium a key innovation for the radiation of lignophytes? Or can the ultimate success of lignophytes be attributed to a step-wise series of innovations including acquisition of heterospory and the seed habit. What was the key innovation underlining the radiation of angiosperms? Was it based on sophisticated hydraulic and mechanical architectures generated by deployment of vessels? Or was it due more to coevolution with insect pollinators or rapid generation turnover? Given an appropriate phylogenetic framework as well as rigorous functional and biophysical studies these kinds of questions might be answerable. Many functional traits or processes are 'taken for granted' in the literature as well as in the relatively few systematic studies that comment on functional and ecological traits in any mechanistic detail. Examples include the supposed mechanical significance of the appearance of wood, the functioning of the seed plant ovule and seed and the appearance and developmental significance of periderm, to name just a few. If one of the aims of evolutionary studies is to infer functional process-related dynamics, such as the radiation of early land plants and the radiation of angiosperms and Gnetales, such studies are potentially just as lost without functional analysis as they are without a phylogenetic and historical background.

Functional studies and biophysical modelling allow investigations of structural evolution over long time periods and across gradients of complexity. The origin(s) of complex structures have been one of the most discussed aspects of evolutionary biology. Some of the findings discussed above show potentially interesting patterns of function and complexity. Simple *de novo* appearance of the conducting strand, the hypoderm and the bifacial vascular cambium show relatively clear primary functional characteristics, which are possibly either coupled initially with secondary functional implications or closely followed by that secondary function. In all three novelties this has been argued as a shift from primarily hydraulic to hydraulic *and* mechanical functioning.

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Becoming fruitful and diversifying: DNA sequence phylogenetics and reproductive physiology of land plants

Martin Ingrouille and Mark W Chase

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Introduction

The use of physiological data in systematics and phylogeny reconstruction has been limited because the physiology of relatively few species has been examined in any detail. However, morphology, anatomy and biochemistry have provided many useful surrogate physiological characters, if physiology is regarded in the broadest sense as covering all aspects of the plant in action. For example, photosynthetic pathways in grasses have been usually inferred from the anatomy of the leaf (Clayton and Renvoize, 1986). The same broad approach to physiology has been used in this chapter.

In general, though, systematists have not valued physiological characters highly. Physiological characters have been regarded as being somewhat different from other characters, more influenced by the environment and not normally a source of phylogenetic information. This was not always the case. Early systematists emphasized the importance of

functional characters that had the greatest influence on the organism. This essentialist approach, pioneered perhaps by Cesalpino (1583), achieved great success since it directed early taxonomists to the importance of reproductive characters and led to the establishment of many 'natural' groups still recognized today. One example of the success of this *a priori* approach was the recognition by Ray of the taxonomic significance of the number of cotyledons, although it has become clear that only the monocot type is useful phylogenetically (Chase *et al.*, 1993).

However, the realization that it was difficult to identify the most essential, or most important functional, characters led to the rejection of this *a priori* approach. De Candolle, who at first propounded *a priori* principles (de Candolle, 1813) later, in his *Prodromus systematis naturalis regni vegetabilis* (de Candolle, 1824-onward), adopted a more pragmatic Adansonian approach, utilizing the distribution of characters to identify the most essential, i.e. those with the most constancy within groups were obviously the most essential.

Most post-Darwinian taxonomists adopted this pragmatic approach (Cain, 1959), even those who explicitly indicated that their classification was phylogenetic. Now they had a theoretical framework to explain why physiological/functional characters might not indicate the phylogenetic history of large groups: they were regarded as having been particularly strongly selected and especially exposed to parallel and convergent evolution that might obscure a phylogenetic signal. For example, a fundamental physiological difference like the presence or absence of crassulacean acid metabolism (CAM) was found to be seemingly randomly distributed in ferns and a range of angiosperm families. Many of the physiological characters have a scattered distribution across all angiosperms and even within families. For example, in a well-defined family such as Poaceae, distinct kinds of PS ('starch' or parenchyma sheath) type and MS (inner or mestome sheath) type Kranz anatomy associated with variation in C₄ photosynthesis, have been shown to have evolved independently at least five times (Clayton and Renvoize, 1986). Nevertheless, surrogate physiological characters, especially from reproductive physiology have remained important.

This rejection of the use of functional characters did not occur at species rank at which adaptation of a species to its environment came to be viewed as an important validation for an evolutionary species concept. Biosystematists such as Clausen and his colleagues (Clausen, 1951) emphasized that the process of speciation involved adaptation to different environments so that at first ecotypes and then sister species differed in their functional characters. The use of that important tool of the biosystematist, the botanic garden, can be regarded as a 20th century attempt to identify what are the 'essential characters' of different groups by cultivating plants in order to minimize environmentally induced variation. In addition, functional differences, relating to pollination and fertilization, which might confer reproductive isolation, were regarded as important for the recognition of what came to be called the 'biological species' (Mayr, 1942). In general, though, taxonomic practice changed little from the morphological-geographical approach long used by most systematists.

However, the availability of stable phylogenetic trees derived from genetic (DNA sequence) data (Chase *et al.*, 1993; Soltis *et al.*, 1997, 1999, 2000; Qiu *et al.*, 2000; Savolainen *et al.*, 2000; Pryer *et al.*, 2001), has provided a new opportunity to study physiological data in a phylogenetic context and thereby evaluate the importance of physiological diversification in the adaptive radiation of plants (Weller *et al.*, 1995).

At first glance, physiological characters, such as the distribution of CAM (crassulacean acid metabolism) or C₄ photosynthesis, widespread in many lineages, do not appear to be constant within clades in the new phylogenetic trees. This scattered distribution in the new

classification (APG, 1998) is also true of some other kinds of characters and is not just a feature of physiological characters. A striking example of sister taxa that is counterintuitive is the placing of such morphologically distinct families as the Nelumbonaceae (lotus), Proteaceae (*Banksia* L.f., *Grevillea* R.Br. ex J.Knight, *Leucadendron* R.Br etc.) and Platanaceae (planes) as close lineages in Proteales. The new phylogeny does point out some remarkable associations such as all the nitrogen fixing taxa in the same major clade of the rosids.

In a similar way, an understanding of the evolution of self-incompatibility (SI) in angiosperms has been hindered by two things: first, by a lack of information outside a few well studied crop plants and, secondly, by the use of broad and imprecise categories. As a consequence, little correlation between phylogeny and SI has been detected. A traditional dichotomy has been made between sporophytic and gametophytic SI on the basis of whether the haploid gametophytic tissue in the pollen grain or the diploid sporophytic tissue in the tapetum mediates the SI specificity, but as we will explain below this is misleading. Gametophytic SI is so widespread that it does not appear to carry any phylogenetic signal (Figure 17.1). Likewise it has been perplexing that the more narrowly distributed sporophytic SI is found in such diverse and unrelated groups as Asteraceae, Betulaceae and Brassicaceae. The perceived low taxonomic value of SI data is being challenged by recent research that has studied the physiology of SI in greater depth and in a greater range of organisms. Below we discuss the distribution of different kinds of SI in the angiosperms in the light of the latest phylogenetic trees derived from DNA sequence data.

First we explore some examples from the reproductive physiology of plants in the broadest sense, utilizing surrogate characters from gynoecial and androecial morphology, anatomy, cytology and development. In fact, although there are some remarkable differences between the phylogenies derived from molecular data and pre-existing phylogenetic classifications, there is, overall, broad agreement. Ironically, this is largely because taxonomists, who have rejected the *a priori* choice of physiologically important characters, have discovered the self-same ‘biologically important’, ‘organizational’ or ‘constitutive’ characters, *a posteriori*, because of their constancy within groups (Davis and Heywood, 1973). The importance of a number of surrogate physiological reproductive characters has been confirmed or re-established and that of a number of new ones indicated. Several of these reproductive characters are also highly significant in the evolution of self-incompatibility in the angiosperms.

Reproductive ‘Physiology’

In the following, we review some of the most remarkable examples of the distribution of surrogate reproductive physiological characters in relation to the phylogenies derived from molecular sequence data (some data from Watson and Dallwitz, 1992 onwards). The examples may be placed into one of two contrasting categories. One category includes lineages/groups that have long been established on the basis of reproductive characters and which find support in the new phylogenetic trees, though frequently the circumscription of the lineage/group has undergone minor changes. The other category includes lineages/groups that have been previously unsuspected but which are supported by some characters from the reproductive physiology of plants previously undervalued or ignored.

One of the most remarkable results of molecular phylogenetics has been the re-establishment of extant gymnosperms as a monophyletic group (Bowe *et al.*, 2000; Pryer *et al.*, 2001). In recent decades gymnosperms have been regarded as a paraphyletic group

| | | | |
|-------|----------------------------|-------------|------------------------------------|
| S | Fagales | | Rosids |
| G | Cucurbitales | | |
| G | Rosales | | |
| G | Fabales | | |
| | Oxalidales | Eurosid 1 | |
| | Malpighiales | | |
| | Celastrales | | |
| | Zygophyllales | | |
| | Sapindales | Eurosid 2 | |
| S & G | Malvales | | |
| S | Brassicales | | |
| S | Myrtales | | |
| | Crossosomatales | | |
| | Geraniales | | |
| | Vitaceae | | |
| G | Saxifragales | | |
| | Berberidopsidales | | |
| G | Dipsacales | | Asterids |
| S | Apiales | Euasterid 1 | |
| S | Asterales | | |
| S | Aquifoliales | | |
| G | Lamiales | | |
| S | Solanales | | |
| S & G | Gentianales | Euasterid 2 | |
| | Garryales | | |
| S & G | Cornales | | |
| S & G | Ericales | | |
| | Caryophyllales | | |
| | Dilleniaceae | | |
| | Santalales | | |
| | Myrothamnaceae/Gunneraceae | | |
| | Trochodendraceae | | |
| | Buxaceae/Didymeleaceae | | |
| | Sabiaceae | | |
| | Proteales | | |
| G | Ranunculales | | |
| S | Ceratophyllales | | |
| G | Winterales | | Magnoliids (including monocots) |
| | Magnoliales | | |
| | Laurales | | |
| | Piperales | | |
| G | Monocots | | |
| | Chloranthaceae | | |
| S | Illiciaceae | | ANITA grade |
| | Schisandraceae | | |
| | Austrobaileyaceae | | |
| | Nymphaeaceae | | |
| | Amborellaceae | | |

S = Sporophytic
G = Gametophytic

Figure 17.1 A simplified taxonomic arrangement of the angiosperms, based on APG (1998), with the distribution of sporophytic (S) and gametophytic (G) self-incompatibility indicated.

with at least two distinct clades, the cycads plus *Ginkgo* L. and the conifers plus gnetophytes differing in seed and other characters. Similarly, the angiosperms are maintained as a monophyletic group for which there has been little doubt, although there are few synapomorphic characters. The production of a specialized pollen reception tissue, the stigma, along with the closure or semi-closure of the carpel, is confirmed as an apomorphic or 'essential' character for the angiosperms, although in some basal angiosperms the carpel is not fused but sealed by mucilage (Endress, 2001). The presence of a stigma greatly enhanced the possibilities of the evolution of self-incompatibility and the evolution of a distinct, elongated style in eudicots and advanced monocots greatly expanded this potential.

Alternatively the gnetophytes rather than being a distinct gymnosperm lineage, ‘anthophyte’ sister to the angiosperms, are now placed as a highly derived lineage of pines (Pinopsida) in the new phylogenetic trees (Bowe *et al.*, 2000). A bisexual fertile apex is now relegated from being an ‘essential’ character.

The new phylogeny does not support the dichotomy of monocots versus dicots, although it does support the essential nature of the possession of a single cotyledon because, although the monocots are demonstrated to be a monophyletic lineage, they are now subsumed as a clade within a broader magnolioid clade close to the base of the angiosperm tree. Closer to the root of the angiosperm tree are a set of minor relatively primitive minor clades, sometimes called the ANITA group from the constituent taxa (Amborellaceae, Nymphaeales, Illiciaceales, Trimeniaceae, Austrobaileyaceae). Therefore, the possession of more than one, usually two cotyledons is shown to be unreliable and dicots are shown to be a paraphyletic group (Savolainen *et al.*, 2000). The exclusion of the magnoliid dicots and various other relatively primitive dicot taxa leaves a monophyletic eudicot group defined by several features. This major lineage of flowering plants has triaperturate pollen with furrows running in parallel to the polar axis, compared to the monoporate/monosulcate pollen of the non-eudicots. They also have a secretory anther tapetum, simultaneous microsporogenesis, well-differentiated stamens with a filament much longer than the anther and two leaf traces. Within the eudicot clade there are two major clades, the rosids and asterids, each divided into two large subclades (euasterid 1 and 2, eurosid 1 and 2) and several other clades such as the ranunculids and caryophyllids (see Figure 17.1).

Non-eudicots frequently have pollen produced by successive microsporogenesis and rather flattened or petaloid stamens. The relationship of some of these characters to the evolution of SI may prove a fruitful avenue for research. An obvious relationship is suggested between SI and a secretory tapetum producing and depositing molecules carrying self-incompatibility specificity into the complex exine of eudicot pollen. One wonders if there is any relationship between type of SI and the distinction that can be made in Asparagales between those (including the Orchidaceae, Iridaceae, Asphodelaceae) with simultaneous microsporogenesis and those with successive microsporogenesis (Alliaceae, Amaryllidaceae, Agavaceae, Hyacinthaceae, Asparagaceae and Rusaceae).

Other reproductive characters do not relate to SI. For example, the core of the caryophyllid clade has been long recognized on the basis of their gynoeceum and developing embryo as the Centrospermae as well as by their betalain pigments and particular phloem sieve-tube plastids. The new phylogenetic trees have revolutionized the circumscription of the main mass of eudicots with the recognition of two main lineages the asterids and rosids, the former including nearly all tenuinucellate and unitegmic eudicots and the latter being mostly crassinucellate and bitegmic. These last two features are also present in non-eudicots but are homoplasious. There are several kinds of crassinucellate conditions, each separately derived. This distinction between the apomorphic tenuinucellate unitegmic state and a plesiomorphic crassinucellate bitegmic state is paralleled by the distinction between the monophyletic leptosporangiate ferns and the paraphyletic eusporangiate ferns and fern allies (Pryer *et al.*, 2001). Remarkably the horsetails are now included within this diverse set of eusporangiate taxa, amply demonstrating that the ‘eusporangiate’ condition includes many different types of sporangia, in the same way as the crassinucellate condition includes many different kinds of nucellus.

Other aspects of reproductive physiology are valuable phylogenetically at lower taxonomic ranks. For example, tetrasporic development is found in a number of groups of families indicated by molecular data. It links the families now subsumed within the

Adoxaceae (Adoxaceae, Sambucaceae and Viburnaceae). Similar groups of sister lineages can be identified that are bisporic such as the Alismataceae plus Potamogetonaceae, Zannichelliaceae and Limnocharitaceae. Another useful character is the nature of the endothecium. For example, the endothecium has fibrous thickenings in a number of groups including Ericaceae (including Empetraceae, Epacridaceae and Monotropaceae) and Myrsinaceae. Girdling thickenings are widespread in monocots but spiral thickenings are found in some sets of families such as the Anarthiaceae–Eriocaulaceae–Flagellariaceae–Poaceae–Joinvilleaceae.

Sexual incompatibility systems

Self-incompatibility (SI) as a defining angiosperm characteristic

Among plants, self-incompatibility (SI) is almost uniquely an angiosperm phenomenon. The presence of mating types in isogamous algae is closer to the determination of two different sexes than the self-recognition characteristic of self-incompatibility. Self-incompatibility has not been reported from bryophytes and pteridophytes. Outcrossing rates that have been measured are relatively high, but this is largely due to alternative mechanisms such as herkogamy (dicliny or dioecy) or dichogamy (protogyny or protandry), or in other cases to inbreeding depression. The single documented case of SI in the fern *Pteridium* Gled. Ex Scop. (Hiscock and Kuess, 2000) is geographically variable within the species and has been doubted by several other workers. Among the gymnosperms self-incompatibility would not be expected in the dioecious cycads, *Ginkgo* and gnetophytes. Owens *et al.* (1998) report SI in *Picea*, active at the stage of pollen tube penetration of the nucellus. This is the only attested example in the gymnosperms, but a similar acting interspecific incompatibility is widespread in Pinaceae (Hiscock and Kuess, 2000).

The relationship between self-incompatibility and the remarkable Cretaceous radiation of the angiosperms has been fertile ground for speculation. *Clavatipollenites* Couper, Forster and Forster grains provide the earliest direct evidence for the presence of angiosperms because they have a complex tectate structure (Friis *et al.*, 1987). *Clavatipollenites* is like the pollen of the living *Ascarina* (Chloranthaceae). Its columellate tectate pollen is associated in living angiosperms with sporophytic incompatibility; the chemical signal mediating one part of the incompatibility is carried within the tectum. In a similar way, the presence of an earlier analogous complex, *Classopollis* Pfl., of the extinct conifers Cheirolepidaceae has been related to their high Jurassic and Cretaceous diversity as if they had had SI (Alvin, 1982). However, SI, especially sporophytic SI, is unlikely to have been ancestral in angiosperms. The distribution of SI among 'primitive' living angiosperms is haphazard as far as has been determined. For example, *Amborella* Baillon., sister to all other angiosperms (Qiu *et al.*, 1999), is dioecious. Many of the ANITA group and relatively primitive eumagnoliids are self-compatible, but some do have SI: *Saururus* L. (Saururaceae) has SI acting at the dry stigma (Pontieri and Sage, 1999), and some members of Winteraceae have late-acting SI.

In recent years information has accumulated from more diverse groups, and molecular techniques have enabled a more precise definition of SI types. In harness with the new phylogeny (Chase *et al.*, 1993), these advances are beginning to demonstrate that the evolution of SI has been far from haphazard. Indeed it reinforces the importance of the evolution of different kinds of SI in the evolutionary diversification of angiosperms, enabled as it was by the key event of the closure of the carpel. The closure of the carpel and evolution of the stigma and style provided a greatly magnified stage on which self-incompatibility

could act. Pollination became a complex sequence of physiological events, from early to late stages, that provided many potential opportunities for the evolution of SI specificity. SI can become manifest at any stage (Wilhemi and Preuss, 1999):

- adhesion of the pollen to the stigma
- hydration of the pollen grain
- germination of the pollen grain
- penetration of the stigma
- growth of the pollen tube in the style
- guidance of the pollen tube to the ovule and embryo sac.

There are also post-fertilization SI mechanisms (Lipow and Wyatt, 2000; Vervaeke *et al.*, 2001).

Each of these stages requires several coordinated physiological events involving cell signalling, and there is a complex cascade of activity with a large potential for specificity at any stage. Different groups have evolved SI by introducing specificity at one or more stages. Although, it seems in some families, or groups of families, a particular form of SI was adopted early in their diversification and has become widespread within all derived lineages, it is clear that some groups have multiple types of SI present. Even in the single species of Asteraceae, *Rutidosia leptorrhynchoidea* F. Muell., the rejection of self-pollen takes place at multiple stages, by the cumulative reduction in adherence of pollen to stigma, pollen germination and pollen-tube penetration of stigma and fertilization (Young *et al.*, 2000). In heterostylous Turneraceae SI is mediated by a different mechanism in short-styled plants from long-styled plants (Tamari *et al.*, 2001). Polemoniaceae have two distinct kinds of SI in different genera (Goodwillie, 1997) as shown by their distinct pattern of inheritance: in *Phlox* L. there is gametophytic SI and *Linanthus* Benth. sporophytic SI (Goodwillie, 1999).

SI acting at the stigma

Brassicaceae exhibit a self-incompatibility type termed sporophytic because the pollen carries the male determinant of both SI alleles present in the diploid anther. However, as has been pointed out (Doughty *et al.*, 1998, 1999) this may simply be because the developing haploid produces pollen coat proteins (PCPs) gametophytically but which are freed and mix within the anther with PCPs from alternative alleles so that all pollen grains carry both signals. In this case SI is only pseudo-sporophytic. Alternatively the diploid tapetum cells might express both alleles in a heterozygote, which would be true sporophytic SI.

The complex cascade of physiological activity involved in SI has been studied in most detail in Brassicaceae. The complex S-locus in Brassicaceae contains several genes expressed in the stigma, including an S glycoprotein (SLG) and a related S receptor protein kinase (SRK) located in the plasma membrane. SLG and SRK function in the stigma and not in the pollen (Conner *et al.*, 1997). SRK is a transmembrane protein, one of a class of receptor-like protein kinases (RLKs) (McCubbin and Kao, 2000). The SLG acts to transfer pollen-coat proteins to the SRK and the SRK mediates the SI (Luu *et al.*, 1997, 1999). SLG also has an adhesive function. Pollen coat proteins, called SCRs or SP11, which are the male determinants of SI, are also part of the complex S-locus (Watanabe *et al.*, 2000). The SCRs act by releasing the inhibition of SRK autophosphorylation normally caused by a stigma thioredoxin (THL1) (Cabrillac *et al.*, 2001). Hypervariable regions that have evolved by point substitutions and intragenic recombination may provide specificity. Phylogenetic trees suggest possible co-evolution of genes coding for male and female determinants (Watanabe *et al.*, 2000).

Pre-existing signal and receptor molecules have been taken over to mediate the SI response. The SLG (S locus glycoprotein) is similar to the extracellular part of SRK. SLG diversification predates speciation of *B. oleracea* L. and *B. campestris* L. (Kusaba *et al.*, 1997), and also SLGs diverged before the *Brassica/Raphanus* split. SLR is an SLG-like receptor targeted to the stigma cell wall. SLR1 is expressed and highly conserved among *Brassica* and functions in pollen–stigma adhesion. SRK and SLG are part of an S-gene family that includes several receptor-like kinase genes (RLKs). The other classes of RLKs are either leucine-rich (LRR) or epidermal growth factor-like (EGF) RLKs. These three have kinase domains that share 40% identity at the amino acid sequence but have different patterns of expression. Other S-gene family kinases are not related to the SI response. In *Brassica* there are other SLRs that are not factors in cell adhesion such as SLR2 and SLR3: the latter is transcribed in leaves, cotyledons and developing anthers and is not linked to the S-locus (Luu *et al.*, 2001). *Orychophragmus violaceus* Bunge and more distantly related Brassicaceae had similar SLR1 sequences to *Brassica/Raphanus* but lacked closely-related SLG sequences (Sakamoto *et al.*, 1998). One, a kinase called SFR2, is implicated in plant defence and accumulates rapidly in response to bacterial infection (Pastuglia *et al.*, 1997), emphasizing the availability of molecules with the potential to mediate SI.

Other genes associated with SI activity in Brassicaceae are an aquaporin gene related to water channel for water transfer to pollen (Ikeda *et al.*, 1997), a kinase-associated protein phosphatase (KAPP) gene (McCubbin and Kao, 2000) and a gene coding for a molecule with thioredoxin activity. Each has a role as an effector molecule in the SI signal cascade as targets of the SRK, but again they have a more general biological function in plant tissues.

In *Ipomoea trifida* (Kunth) G. Don (Convolvulaceae), there is a single S-locus homologous to that in *Brassica* and with molecules called IPG and IRK that have 40–46% similarity to *Brassica* SLGs and SRKs, respectively. IPG expression is developmentally regulated in stigma and anther tissue (Kowyama *et al.*, 1995; Kakeda and Kowyama, 1996). However IRK1, a putative receptor kinase, appears not to be primarily involved in the self-incompatibility system and shows no linkage to the S-locus (Kowyama *et al.*, 1996, 2000). Similar S-gene RLKs have been detected in maize (*Zea* L.).

Asteraceae also have sporophytic SI, but this is different in detail from that in Brassicaceae and is likely to be of independent origin. For at least some species it acts in a different way to Brassicaceae SI: by cumulative reductions in the adherence of pollen, pollen germination, pollen tube penetration of the stigma and fertilization (Young *et al.*, 2000). Betulaceae also have a sporophytically determined SI but, at least in *Corylus avellana* L. it does not show any relationship to that in Brassicaceae (Hampson *et al.*, 1996).

More similar to the *Brassica* model of SI is that exhibited by Papaveraceae, yet this is genetically determined gametophytically. The lack of any extensive style precludes a stylar recognition mechanism reported for some other gametophytic systems. Here SI occurs in two stages. The first stage is as if the process of pollen hydration and germination have been taken over by the SI mechanism as it has in Brassicaceae, at least in part because it involves Ca^{2+} and calmodulin dependent increased phosphorylation of a 26-kD glycoprotein (Rudd *et al.*, 1997; Hearn *et al.*, 1996). However, it also involves DNA degradation (Jordan *et al.*, 2000b).

SI acting in the style

One of the most common and phylogenetically widespread SI mechanisms has arisen by the taking over of specific RNases acting in the style to degrade RNA produced by the

growing pollen tube, especially in eudicots and more derived monocots because they have an elongated style. This may prove to be the most common form of SI because elongated style S-RNase-mediated incompatibility is associated with wet stigmas and pollen binucleate at anthesis, conditions that are widespread in eudicots and monocots. RNases have been widely detected in plants and fungi as regulators of development but here have gained specificity to mediate the SI response (Bower *et al.*, 1996).

The assessment of homologies is complex, and it may be that there has been substantial independent evolution of stylar-acting SI mediated by RNases. Lack of homology may be indicated by variations in the precise mechanism of S-RNase-mediated SI in diverse families such as Solanaceae, Scrophulariaceae, Campanulaceae and Rosaceae. This is gametophytically determined because the RNA targets are being transcribed in the haploid pollen tube. One could predict that S-RNase-mediated SI is likely to be discovered in other groups with gametophytic SI with stylar inhibition such as Fabaceae. However, Fabaceae may differ because they seem to be capable of having an order of magnitude greater number of SI alleles than Rosaceae (Kowyama *et al.*, 1996) and, at least in *Acacia Senegal* (L.) Willd., self-incompatibility acts in the embryo sac (Tandon *et al.*, 2001). S-RNase-mediated SI is also likely to be present in monocot families with stylar gametophytic SI such as Bromeliaceae (Vervaeke *et al.*, 2001). The question of whether S-RNase activity is a plesiomorphic or apomorphic feature for these groups has been addressed by a detailed study of the phylogeny of the S-RNase sequences in three families, Solanaceae, Scrophulariaceae and Rosaceae and fungi (Richman *et al.*, 1997). RNases not linked to the S-locus and with functions other than SI were included in the study. RNases that function in defence against pathogens in *Petunia* Juss. are related to S-proteins in Solanaceae (Lee *et al.*, 1992). Angiosperm non-S-RNases form one monophyletic group, and the S-RNases of each of the families form three more monophyletic groups.

Independent origin of stylar RNase-mediated SI does seem likely in the rosids and asterids. The S-RNases of Rosaceae are sufficiently distinct from those in Solanaceae/Scrophulariaceae to indicate an independent origin (Sassa *et al.*, 1996). S-RNases have been studied in most detail in Solanaceae and Rosaceae and, in both, diverged at an early stage of, or even before, the diversification of these families. S-RNases have only 25% sequence identity to non-S-RNases in Rosaceae (Norioka *et al.*, 1996). In each family there has been extensive trans-specific and trans-generic evolution of alleles (Richman *et al.*, 1995, 1996).

SI in euasterid I

In Solanaceae, sets of alleles among *Physalis* L., *Nicotiana* L., *Petunia* and a species of tomato, *Solanum peruvianum* L., are more similar to each other than alleles are intraspecifically. The alleles of a different species of tomato, *Solanum esculentum*, have an independent origin (Richman and Kohn, 1996) as do those in the distant genus *Momordica* L. (Cucurbitaceae). Similarly, in Rosaceae, pairs of alleles between *Pyrus* L. and *Malus* L. are more similar to each other than those within either genus (Ishimizu *et al.*, 1998). S-RNases have also been detected in *Prunus* L. (Burgos *et al.*, 1998).

Even though the S-RNases of Solanaceae and Scrophulariaceae appear to be related and these families are both in the major angiosperm clade euasterid I, the origin of S-RNase-mediated SI may be independent in these two families (Xue *et al.*, 1996).

S-RNase-mediated SI is not universal in the euasterid I clade. Although S-RNase-determined SI is widespread, alternative mechanisms exist even within those families in

which S-RNase has been detected. *Lycium cestroides* Schtdl. (Solanaceae) has ovarian SI (Aguilar and Bernadello, 2001). *Asclepias exaltata* L. in another family, Apocynaceae, has a different kind of SI involving post-zygotic rejection of self-fertilized ovules due to a single late-acting SI locus (Lipow and Wyatt, 2000). Apocynaceae are different in another way in having a complex pollen presentation mechanism and are also members of Gentianales in which heterostyly is widely distributed; four out of five Gentianales are heterostylous (Gelsemiaceae, Gentianaceae, Rubiaceae, Loganiaceae). In addition, another kind of SI is found in the unplaced euasterid family Boraginaceae, detected because it has polygenic inheritance.

In another euasterid I family, Lamiaceae, SI has not been demonstrated convincingly (Owens and Ubera-Jiménez, 1992), although differences between the two major lineages in pollination strongly indicate that, if it is present, there are at least two distinct kinds. It is no accident that detected SI is rare in this family in which there are highly developed mechanisms of dichogamy and herkogamy, preventing self-pollination. Another example of a family where SI is rare is Orchidaceae in which a complex floral morphology mechanically limits the opportunities for selfing (Johnson and Edwards, 2000). Perhaps the lack of SI in Lamiaceae is more a consequence of it not having been looked for. Certainly the clear separation of two major lineages of Lamiaceae, Lamioideae and Nepetoideae, is suggestive of a differential distribution of types of SI: Lamioideae have binucleate pollen at anthesis and wet stigmas, conditions normally associated with gametophytic SI and Nepetoideae have tri-nucleate pollen and dry stigmas, which are normally associated with sporophytic SI (though exceptionally *Lavandula* L. has wet stigmas).

Alternative kinds of SI

Various other kinds of SI have been detected. In *Lilium* L., SI is associated with the activity of 1-aminocyclopropane-1-carboxylate (ACC) oxidase. Self-incompatibility provokes a kind of stress response with reduced levels of ethylene and superoxide dismutase and enhanced catalase, ascorbate peroxidase, dehydrogenase reductase and glutathione reductase (Suzuki *et al.*, 2001). In *Narcissus* L. there is another kind of SI resulting from embryo sac degeneration following self-pollination (Sage *et al.*, 1999).

Grasses exhibit yet another kind of gametophytically-determined SI, two-locus SI. The two loci, S and Z, are unlinked showing remarkable synteny and conservation of gene order among grasses, in Triticeae, Poaceae and Avenae, possibly all Poaceae (Baumann *et al.*, 1999). The S-gene is present throughout grasses in all subfamilies regardless of self-compatibility (Li *et al.*, 1997). The S-gene in grasses has thioredoxin activity and an allele-specific portion as well as catalytic domain. SI in grasses is associated with trinucleate pollen, high respiration, short viability, difficult growth *in vitro*, dry stigma papillae with entire cuticle, inhibition at stigma surface, and callose deposited in exine.

In *Theobroma cacao* L. (Malvaceae), the incompatibility response is modulated by auxin (Hasenstein and Zavada, 2001). Multilocus SI is found in Ranunculaceae (*Ranunculus* L.) and Amaranthaceae (*Beta* L.). Concealed genes for self-incompatibility have also been detected in Caryophyllaceae (Lundqvist, 1995).

Heteromorphic SI

It is clear that SI has diverse origins but, at lower taxonomic ranks, shows homology. This fits in well with what is understood about the evolution of heteromorphic SI, which is present in 24 families and has clearly multiple origins (Ganders, 1979). Here too confusion

has been created because of the sloppy use of terms to describe non-homologous adaptations. Differential pollen tube growth in styles of different lengths (heterostyly) mediates SI in some heterostylous groups, as in many *Primula* L. (Primulaceae) species (Richards, 1986). However, in *P. obconica* Hance and *P. vulgaris* Hudson most of the SI is mediated at the stigma. In heterostylous *Linum* L. (Linaceae) differences in osmotic pressure ratio between the morphs of pollen and style leads either to non-germination of the pollen or bursting of the pollen tube in alternative incompatible pollinations (Lewis, 1943).

‘Heterostyly’ is recorded from about 13 families, including Plumbaginaceae but, in this family at least, it is weak and rare and has little to do with SI, which is instead associated with dimorphism of the stigma and pollen and the adhesion of pollen to the stigma. Similarly in *Narcissus triandrus* L., which is tristylous, there are no significant differences in pollen tube growth in the style, and SI is mediated by differential ovule development in self- and cross-pollinated plants (Sage *et al.*, 1999).

Heteromorphic SI provides more examples of the complexity of SI mechanisms in the angiosperms, and the lack of homology among groups at higher taxonomic ranks but more evidence for the homology of individual mechanisms at lower taxonomic ranks.

Self-compatibility

Self-incompatibility of different sorts has arisen many times, obscuring any simple phylogenetic pattern. An additional confusion is the widespread distribution of self-compatibility (SC). This has evolved even more times than SI. It may be ancestral (plesiomorphous) in the angiosperms, but it is clear that most SC has arisen from SI independently in each group, even within sets of closely related species, for example in *Linanthus* Benth. (Goodwillie, 1999). Phylogenetic analysis has demonstrated that the presence of SI is the ancestral state in one section of *Linanthus* within which there have been three to four transitions to self-compatibility. The distinct self-compatible lineages exhibit convergent evolution with respect to morphological and behavioural traits associated with self-compatibility (Goodwillie, 1997).

SC has sometimes arisen by deletion of SI loci as in *Arabidopsis* Heynh. (Conner *et al.*, 1998) but, gametophytic mutants have also been detected that impose SI in various stages from pollen tube growth to endosperm development and even fruit development (Wilhelmi and Preuss, 1999). Of great evolutionary significance is what Hiscock (2000), from studies of *Senecio*, has called pseudo-self-compatibility. In *Senecio squalidus* L. SI is weakened by a cryptic unlinked gametophytic modifying element (G gene). S-locus function is retained but modified under selection, thereby furnishing the possibility of a flexible evolutionary response to selection for greater or lesser rates of out-crossing. It also clouds the perception of the distribution of SI within groups because of the perception of the phylogenetic intermingling of SI and SC species.

Another seemingly paradoxical pattern is the frequent presence of dioecy and SI in sister lineages. A clue has been provided by the failure of SI due to polyploidy, which has, for example, been demonstrated in *Rosa* L. recently (Ueda and Akimoto, 2001). Charlesworth (2001) has suggested that dioecy provides a mechanism by which a lineage can maintain high rates of outbreeding after polyploidization and SI breakdown.

Conclusion

In the light of the new DNA sequence derived phylogenetic trees, physiological traits have been shown to have patterns of distribution substantially the same as other types of

characters. They are neither more nor less essential or more or less functional than other characters. They potentially have as much value in classification as other characters.

The distribution of different kinds of SI across groups at different ranks in the taxonomic hierarchy appears complicated because of a hitherto unsuspected flexibility in SI response. The assessment of homology is complicated because a shared common origin of an SI mechanism (true homology) may be obscured by multiple loss of SI and the presence of other kinds of SI in the families. Gene duplication and transpacific evolution, so that lineages of different SI genes in an individual taxon pre-date the divergence of the taxon itself, further complicate evolutionary patterns. Parallel evolution of identical SI mechanisms has also occurred by the subversion of the same cell-signalling process at multiple stages of pollination. In the latter cases detailed study of the arrangement of genes, or of the gene sequences, has indicated a lack of true homology. For example detailed analysis of the S-RNases between Solanaceae, Scrophulariaceae and Rosaceae has indicated homology between the first two and a remarkable convergence between them and Rosaceae. As with other kinds of characters the absence of a feature, of a certain kind of physiology, provides a rather weak kind of taxonomic information. In fact the detailed study of the physiology of self-incompatibility is revealing hitherto unsuspected variation, potentially the source of many new characters.

The amazing physiological diversity of self-incompatibility between and within groups emphasizes the importance of self-incompatibility in the diversification of the angiosperms. It also reflects on the fundamental importance of the closure of the carpel and evolution of the style and stigma within angiosperms because it was these events that provided the locus from which many kinds of self-incompatibility could arise.

There are examples of similar kinds of SI being present across widely divergent taxa, such as the distribution of S-RNases between Solanaceae/Scrophulariaceae and Rosaceae, but detailed analysis often indicates a remarkable convergence rather than a shared ancestral state. As with other characters, absence of a feature, of a certain kind of physiology, provides a rather weak kind of taxonomic information.

The mistrust with which physiological characters, such as the distribution of types of SI, have been regarded by systematists is unjustified. In relation to self-incompatibility it rests only on ignorance of the precise nature of the physiology and of the distribution of such characteristics across a broad range of taxa. More generally the 'essential' nature of physiological characters depends upon the group in which they are being examined.

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18

Evolution of angiosperm fruit and seed dispersal biology and ecophysiology: morphological, anatomical and chemical evidence from fossils

Margaret E Collinson and Pim F van Bergen

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Introduction

Angiosperms, or flowering plants, become obvious in the fossil record from about 125 million years ago (Ma) and dominant from about 90 Ma (Crane *et al.*, 1995; Wing and Boucher, 1998). A second radiation (Collinson, 1990) takes place in the latest Cretaceous and early Tertiary (Palaeogene). This chapter documents those fruit and seed palaeoecophysiological strategies (especially dispersal biology, embryo and endosperm development, germination and establishment, dormancy and resistance) which had evolved by, or during, the Palaeogene (65 to 34 Ma) after this second radiation. Evidence for these strategies comes largely from the gross morphology and internal anatomy of fossils as well as the chemical composition of resistant layers (e.g. fruit wall, seed coat) which survive in fossils. Where possible the evolutionary status in the Palaeogene will be compared with that of the earlier stages of angiosperm evolution in the Cretaceous.

Dispersal biology

Abiotic – plumes

Details of Palaeogene plumed disseminules are given in Appendix 18.1. These details reveal that there is an extremely low diversity of plumed disseminules in contrast to the high diversity of winged forms (Figure 18.1, Appendix 18.2) in the Palaeogene. Only a single type of plumed seed is clearly represented (the *Apocynospermum*-like plume; see Appendix 18.1), in which the plume probably functioned to control orientation on landing. This seed morphology is very uncommon although it is widespread in Palaeogene floras across Europe and North America. The only other clearly documented example is hairs on achenes of middle Eocene *Platanus*, hairs being lacking on earlier platanaceous fossils. The plumed strategy of influencing speed or orientation of fall seems to have been scarcely exploited in the Palaeogene and has not been reported in the preceding Cretaceous.

Palaeogene floras seem to have exhibited an extremely limited array of plumed disseminules by comparison with the modern variety (e.g. Ridley, 1930; Burrows, 1986). Taphonomic bias (including shedding and loss of pappus, lack of recognition of small plumed seeds, inadequate study of bedding surfaces of sufficiently fine grained sediments) could all have affected the fossil record of plumes. However, small seeds are well known in the Cretaceous and Palaeogene record where SEM studies should have revealed plume scars and the numerous winged disseminules (see Figure 18.1, Appendix 18.2) testify to extensive study (at least for the Palaeogene) of fine grained bedding surfaces. Plumes may be more recent evolutionary innovations as is seen in the fossil record of Platanaceae (Appendix 18.1) and also by the fact that many modern plumed taxa belong in relatively advanced clades (e.g. Asteraceae) with almost no Palaeogene fossil record (Collinson *et al.*, 1993). Higher humidities may not have favoured the early evolution of this strategy.

Abiotic – wings

In striking contrast to plumes, Figure 18.1 shows that a huge variety of wings are known in the Palaeogene, encompassing all major modern categories and having the potential to

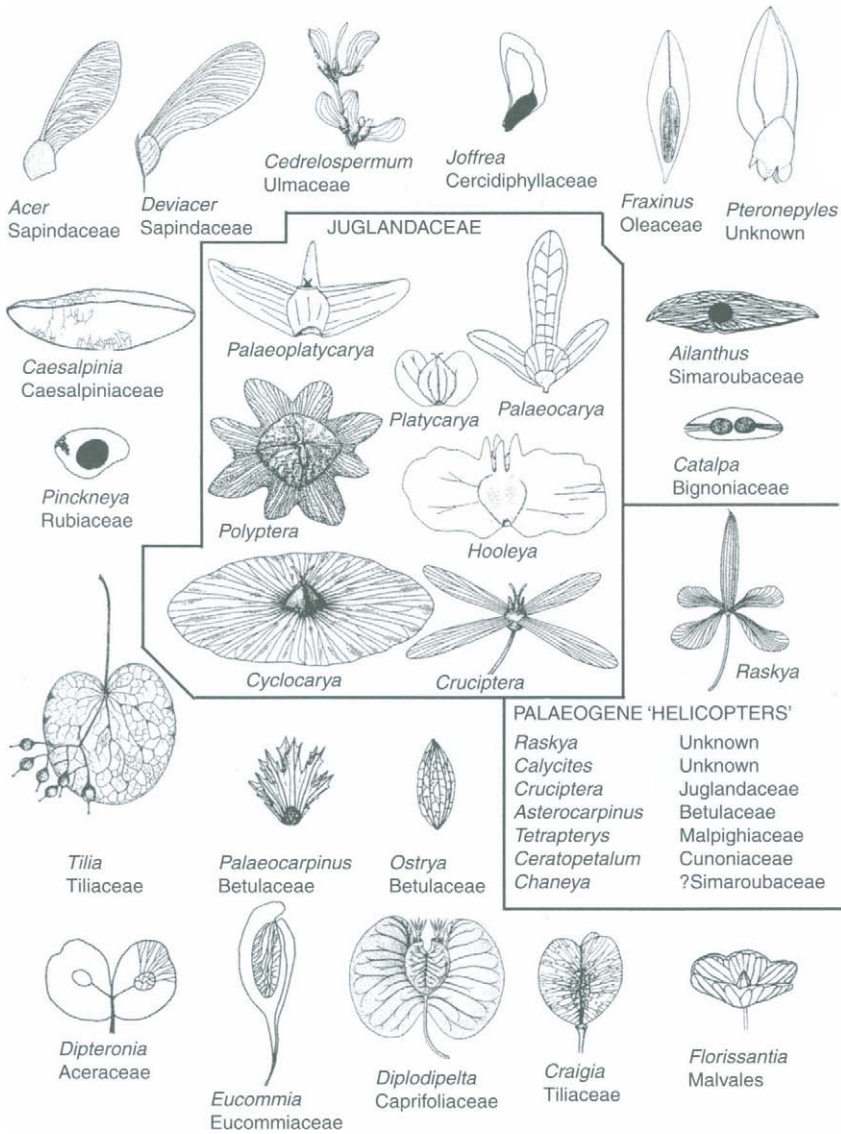


Figure 18.1 The variety of wings in Palaeogene disseminules. All illustrations are based on actual fossils. References to the sources may be found in the text of Appendix 18.2. The figure demonstrates the following points, which are also discussed in the main text:

1. That there is a wide variety of winged disseminules in the Palaeogene. None of this variety is present in the Cretaceous
2. That a wide variety of wings may be seen within a single family, e.g. Juglandaceae
3. That a given wing morphology (e.g. 'helicopters') may be found across a wide variety of taxonomic affinities
4. That wings are derived from a wide variety of different parts of the inflorescence, flower, fruit wall and seed coat
5. That the variety of wing morphology which was established by the Palaeogene encompasses the vast majority of modern strategies.

exploit a wide variety of flight paths, trajectories and attitudes in flight (Burrows, 1986). Details of Palaeogene winged disseminules are summarized in Appendix 18.2. Although the summary is extensive it may not be totally comprehensive. However, it does serve to demonstrate the following key points: (1) that individual flowering plant families in the Palaeogene may show considerable variety of wing form (e.g. Juglandaceae, Figure 18.1, Appendix 18.2); (2) that comparable wing morphologies in the Palaeogene occur across systematic boundaries (e.g. helicopters, Figure 18.1, Appendix 18.2); (3) that comparable wing morphologies in the Palaeogene were developed from different anatomical entities (many examples, Figure 18.1, Appendix 18.2); and (4) that a very wide variety of wing morphology, on seeds and fruits, and derived from a variety of organs, had already evolved in the Palaeogene (many examples, Figure 18.1, Appendix 18.2).

In sharp contrast to the situation in the Palaeogene, Cretaceous relatives (where well-understood and based on actual fossils) tend to show unaided dispersal (Crane *et al.*, 1995). Taphonomic bias may partly account for this observation as many studies of Cretaceous angiosperm fruits and seeds are based on sieved residues where wings would be less obvious than they are when preserved on bedding surfaces. However, scars of former wing attachment should be obvious from SEM studies of specimens from sieved residues. Many Cretaceous angiosperms, especially those of the earlier phases of angiosperm evolution, were probably plants of small stature, which produced small fruits and seeds. Neither of these attributes would have favoured the earlier development of winged disseminules. Eriksson *et al.* (2000a) recorded up to 15% wind dispersal in late Cretaceous fruit and seed floras with lower levels (0–7%) in Palaeogene floras rising again in the Neogene. However, they inferred dispersal based on extrapolation from nearest living relatives, whereas in this chapter we document actual presence of wings in the fossils themselves. In addition, it is not clear if they included small dry fruits/seeds lacking wings in their wind-dispersed category. There are few examples of seeds with wings preserved in the literature on Cretaceous seeds, those that are documented (e.g. Crepet and Nixon, 1994) are minute wings on small seeds. The exceptionally preserved early Cretaceous angiosperms from the Crato Formation (bedding plane assemblages in fine grained limestone where wings would typically be preserved) do not include any winged disseminules (Mohr and Friis, 2000). Even if Cretaceous floras contained significant amounts of wind dispersed disseminules, any wings were small and certainly did not show the elaborate variety documented in Appendix 18.2 for the Palaeogene.

Abiotic – dust and microseeds

Cretaceous seeds, almost without exception, are small (volume less than 1 mm³) (Crane *et al.*, 1995; Friis *et al.*, 1999, 2000; Eriksson *et al.*, 2000a,b). Only in the latest Cretaceous (Maastrichtian) is a small increase in size evident (Knobloch and Mai, 1986, 1991). Small seeds persist in the Palaeogene where they are joined by a wide variety of larger forms.

Cretaceous seed assemblages are only just beginning to be studied in detail so systematic affinities are mostly not yet clear enough to assess more detailed evolutionary patterns. Certainly, some families, with larger seeds today, are represented in the late Cretaceous (e.g. Campanian and Santonian) by extinct forms with small seeds. These include: members of the Fagales *sensu* APG, 1998 and Fagaceae (Herendeen *et al.*, 1995; Sims *et al.*, 1998; Schönenberger *et al.*, 2001); the Juglandaceae (Friis, 1983; Knobloch and Mai, 1986, 1991; Herendeen *et al.*, 1999) and the Mastixioid Cornaceae in the Maastrichtian (Knobloch and Mai, 1986, 1991). Less clear is the Cretaceous representation of taxa with small seeds

today but it certainly includes members of the Ericales (Friis, 1985b; Knobloch and Mai, 1986, 1991; Schönenberger and Friis, 2001). Small seed size in the Cretaceous may in part be related to small stature (e.g. Wing and Boucher, 1998) of early angiosperms, a suggestion also made by Eriksson *et al.* (2000a), rather than reflecting the adoption of a particular dispersal strategy. Nevertheless, small seeds would be appropriate for the colonizing strategy implied by the occurrence of early angiosperm fossils in depositional environments representing disturbed habitats (e.g. Wing and Boucher, 1998).

Palaeogene floras also include small seeds and many of these belong to families which are typified by small seeds today. Examples include:

1. *Rhododendron* (Ericaceae) with elongate seeds, with small terminal prolongations of the seed coat which are recorded from the Palaeocene/Eocene transition in England (Collinson and Crane, 1978; Collinson and Cleal, 2001c) and in the late Eocene of California (Wang and Tiffney, 2001)
2. *Hydrangea* (Hydrangeaceae) and *Emmenopterys* (Rubiaceae) with tiny winged seeds from the middle Eocene of the USA (Manchester, 1994b)
3. Hydrangeaceae, Ericaceae and Onagraceae from the middle Oligocene and Miocene of Europe (Mai, 1985b, 1998)
4. Restionaceae (Dettman and Clifford, 2000) from the Eocene/Oligocene of Australia
5. Juncaceae from the Eocene/Oligocene transition of England (Collinson, 1983a)
6. Lythraceae and Flacourtiaceae, with seeds less than 1 mm in length, from the early Eocene London Clay flora of England and middle Eocene Clarno flora of the USA (Reid and Chandler, 1933; Manchester, 1994b; Collinson and Cleal, 2001b).

Based on their size the smallest of these Cretaceous and Palaeogene seeds can only have had minimal stored reserves in the endosperm or embryo (no preservation known). Therefore, they may have relied on alternative nutritional associations like some of their living relatives.

Abiotic – flotation

The fibrous fruit wall of fossil *Nypa* fruits is indicative of the potential for dispersal by flotation similar to modern fruits of the genus. This is consistent with the fossil record of the fruits, including their sedimentary facies and strand-line-like accumulations (e.g. Collinson, 1993, 1996) and the overall comparability of the reproductive biology between modern and fossil examples. Some fossil seeds exhibit thin-walled tissues in the seed coat often referred to as ‘spongy’, which may represent light tissue aiding flotation. Examples include *Decodon* (Cevallos-Ferriz and Stockey, 1988a and references therein) and to a lesser extent *Keratosperma* (Cevallos-Ferriz and Stockey, 1988b). Water tight seed-coat layers, such as columnar palisades, may aid dispersal in water (see discussion on dormancy).

Biotic – dry fruits and seeds

Potential dispersal by animals is likely (based on inference via living relatives) for large dry nuts with an extensive food reserve, e.g. Juglandaceae, Betulaceae, Fagaceae which became abundant in the late Eocene, coincident with a radiation of rodents (Collinson, 1999; Collinson and Hooker, 2000; Hooker and Collinson, 2001). In the late Cretaceous, Juglandaceae and Fagaceae are represented only by small dry nuts, while Betulaceae have not yet been reported (see Appendix 18.2 and Chen *et al.*, 1999).

Previous work has shown that the earliest examples of larder hoarding are Miocene (Collinson, 1999; Hooker and Collinson, 2001; Gee *et al.*, in press). Except in cases of hoarding, direct fossil evidence of animal interaction with nuts and dry seeds represents seed predation. These include rodent-gnawed seeds dating from the latest Eocene and insect borings in Eocene Rutaceae seeds (Collinson, 1999; Collinson and Hooker, 2000; Hooker and Collinson, 2001).

Biotic – spines

Spines or hook-like protrusions are known on Eocene *Fagus* cupules (Meyer and Manchester, 1997), on Eocene *Castanea* cupules (Crepet and Dahlgian, 1980), on Oligocene *Sloanea* cupules (Kvacek *et al.*, 2001) and on Palaeocene *Aesculus* cupules (Manchester, 2000b, 2001). They are also recorded on small (2 mm maximum dimension) early Cretaceous fruits named *Appomattoxia* (Friis *et al.*, 1995). Larger fruits of *Trapa*, *Ceratostratiotes* Gregor and *Ceratophyllum* (from the Palaeocene and Eocene) also bear short to long spines, sometimes hooked (Mai, 1985a, 1995; Herendeen *et al.*, 1990; Wehr, 1995; Manchester, 2000a; Meller and van Bergen, 2003), some of which are known attached to complete plants confirming their hydrophytic biology (Herendeen *et al.*, 1990). Spines of aquatic disseminules can function as anchors and this has also been argued (Collinson, 1999) as their probable role in *Appomattoxia*. In other cases, especially when unhooked (e.g. *Sloanea*), spines probably had a protective function. A dispersal function seems less likely in cases where fruits shed seeds on splitting.

The only known proof of fossil animal epizoochory of an angiosperm of which we are aware is that where mammalian hairs are entwined in the hooks of a fruit identified as the bamboo *Pharus* from late Eocene Dominican amber (Poinar and Columbus, 1992). Modifications favouring epizoochory seem to be exceptionally rare in Cretaceous and Palaeogene fruits and seeds in spite of the logical origin of mammalian fur in the late Triassic (Collinson, 1999; Hooker and Collinson, 2001).

Biotic – fleshy tissues

Fossil evidence for endozoochory has recently been documented elsewhere (Collinson, 1999; Hooker and Collinson, 2001). This evidence is derived from exceptionally preserved fossils including mammalian and bird gut contents and evidence for the former presence of soft tissues on fruits (e.g. from the middle Eocene of Messel, Germany, Richter, 1987; Schaal and Ziegler, 1992) as well as from dental morphology and microwear of mammalian fossils. This evidence shows that, by the early Palaeogene, large fleshy fruit existed (e.g. Lauraceae, Vitaceae, Menispermaceae). Fleshy fruits were consumed by mammals and, at least in one case, the seeds survived uncrushed in the stomach and so could have been voided in faeces in a viable condition. Seed predation is also evident in the form of crushed seeds in the gut content of one mammal. However, these examples are rare and, although they prove seed predation and potential endozoochorous seed dispersal in the Palaeogene, most interpretation of animal dispersal is still based on inference from the presence (or inferred presence) of fleshy tissues. Mack (2000) argued that fleshy fruit pulp might have evolved as a defence against seed predation and secondarily become structures to promote seed dispersal.

By contrast to the Palaeogene, Cretaceous fruits and seeds were small and none were enclosed in large fleshy fruits. Some fruits possessed thin layers of fleshy tissue which may originally have been rather leathery (Knobloch and Mai, 1986, 1991; Crane *et al.*, 1995; Friis *et al.*, 1999, 2000; Eriksson *et al.*, 2000a). Eriksson *et al.* (2000a) documented that

proportions of animal dispersal increased steadily from about 80 million years ago and reached a peak in the Palaeogene. However, their study relied mainly on living relatives to assign a dispersal system for the fossils (see also Abiotic – wings above). In one very well-studied early Cretaceous fruit and seed flora (Eriksson *et al.*, 2000b) it was determined that about one quarter of the taxa could be interpreted as drupes or berries with specialised tissues for animal dispersal. This interpretation was based on the presence of thin outer fruit walls. The average fruit volume was 2.22 mm³ (range 0.31–7.58 mm³). Thus, the Cretaceous fleshy fruit resource is of an utterly different category, in terms of size and tissue volume, compared with that of the Palaeogene. The significance of Cretaceous birds or reptiles for fruit and seed dispersal is uncertain but small Mesozoic mammals, such as multituberculates, haramyids and docodonts all had dentitions indicative of potential fruit or seed eating (Eriksson *et al.*, 2000a,b; Hooker and Collinson, 2001). However, direct evidence of a dispersal (as opposed to predation) role for Cretaceous fruit and seed feeders is lacking, though it could be argued that such small seeds would avoid tooth crushing and survive to be defecated much as tomato seeds do in humans today.

Germination and establishment

Embryo and endosperm

Radicle emergence

Emergence of a radicle (hence proof not only of presence of an embryo but also of mode of germination) is recorded in fossil seeds but very rarely. We are aware of only three examples. One is a late Cretaceous charred seed (Friis, 1985a), the second the seeds named *Microphallus* Manchester (affinity unknown) from the middle Eocene Clarno flora (Manchester, 1994b) and the third three specimens of *Joffrea* (Cercidiphyllaceae) from a compression flora with numerous seedlings (see seedlings below). *Microphallus* is quite common (25 specimens in a flora where a number of taxa are represented by fewer than ten specimens but others by 200+). The number of specimens showing radicle emergence probably indicates more or less simultaneous germination. (See also under seedlings below.)

Dispersal strategies (above) suggest utilization of appropriate orientations, positionings, or placements, at least in the Palaeogene. However, there is little or no evidence of the elaboration of this such as is associated with many dry land herbs today (e.g. no twisting awns etc).

Embryo and endosperm development, dormancy and establishment

Sources of evidence. Charred fossils from the Cretaceous do preserve anatomical details of delicate tissues such as endosperm. However, the emphasis in the study of the Cretaceous charred angiosperms has so far been largely on the flowers which yield ovule rather than seed detail (e.g. Drinnan *et al.*, 1990). Friis (personal communication 2001) considers that the Cretaceous dispersed charred seeds (e.g. Eriksson *et al.*, 2000b) do hold potential for investigation of embryo and endosperm characteristics.

The degree of development of the embryo and endosperm (see following section) can be assessed from several permineralized floras from the Cretaceous/Tertiary transition and Palaeogene. These are summarized below. Where embryo and endosperm preservation is lacking this can sometimes be shown to be due to the activities of fungi prior to permineralization (Kalgutkar *et al.*, 1993; Stockey *et al.*, 1998).

The floras from the Deccan Intertrappean series have been variously considered as Tertiary or Cretaceous in the past, but recent dating suggests that the Intertrappean Series

spans the Cretaceous/Tertiary (Maastrichtian/Palaeocene) transition including the boundary itself (Courtilot *et al.*, 1988; Jaeger *et al.*, 1989; Courtilot, 1999; Guleria and Srivastava, 2001). Reviews of the flora and floral lists may be found in Prakash (1960, 1978), Bande *et al.* (1988) and Band and Chandra (1990). The fossils are preserved as silica permineralizations or petrifications and hence preserve some anatomical details not found in compression floras. There are relatively few fruits or seeds described from this flora. A large number of these have suggested affinity with the palms (e.g. lists in Prakash, 1960, 1978). Some of these do not preserve internal seed details (e.g. Mehrotra, 1987) while others do (e.g. Bonde, 1990), the latter having a small, apical undifferentiated embryo within extensive non-ruminate endosperm. Seeds of *Viracarpou* (?Araceae, Prakash, 1978) are unfortunately not sufficiently well-preserved to reveal embryo or endosperm detail (Chitale, 1958). Fruits of *Tricocites* Rode are known in organic attachment to plants with *Cyclanthodendron* Sahnii and Surange stems (Bonde, 1985; Biradar and Bonde, 1990) with affinity with Musaceae/Strelitziaceae. Their anatomical preservation is discussed below. In addition, fruits of *Enigmocarpou* (?Myrtales) are also discussed below.

The middle Eocene flora from the Princeton Chert is reviewed by Cevallos-Ferriz *et al.* (1991) and Pigg and Stockey (1996) and is currently under active study (e.g. Stockey *et al.*, 1998). This flora is permineralized by silica and provides excellent anatomical detail (e.g. Cevallos-Ferriz *et al.*, 1991; LePage *et al.*, 1997; Stockey *et al.*, 1998). The early Eocene London Clay flora was reviewed by Collinson (1983b) and Collinson and Cleal (2001b). The fossils are preserved by pyrite permineralization and many yield good anatomical detail. The middle Eocene flora of the Clarno Chert was monographed by Manchester (1994b). The specimens are mostly preserved as silica petrifications with some permineralization yielding anatomical details of less resistant tissues. The late Palaeocene Almont flora was reviewed by Crane *et al.* (1990) and also contains some anatomical details as the result of siliceous preservation.

Fossil evidence of Palaeogene endosperm and embryos. The following list summarizes the direct fossil evidence for Palaeogene endosperm and embryos taking all the above floras into account. Additional references are cited where necessary.

Small undifferentiated embryos with large amounts of endosperm occur in Musaceae from the middle Eocene of Clarno (Manchester and Kress, 1993; Manchester, 1994b) and some seeds assigned to palms from the Deccan flora (Bonde, 1990).

Direct evidence of the presence of an extensive ruminate endosperm filling the seeds, and hence indirect evidence of a small embryo, is known from numerous seeds. In particular these are numerous forms of Anonaceae seeds (*Anonasperrum*) in the Eocene Clarno and London Clay floras and also from the Palaeocene of Pakistan with earliest records in the Maastrichtian of Nigeria (review in Tiffney and McClamer, 1988; see also As-Saruri *et al.*, 1999). Vitaceae from the London Clay and Clarno are another example.

Perisperm (a product of the nucellus) accompanied by a small embryo has also been inferred on the basis of cellular preservation with a suitably placed embryo cavity in seeds of *Allenbya* (Nymphaeaceae) from the middle Eocene Princeton Chert (Cevallos-Ferriz and Stockey, 1989) and in seeds of the dicotyledon *Princetonia* Stockey from the same flora (Stockey and Pigg, 1991). In the absence of anatomical preservation it is not possible to establish degree of differentiation of embryo in these examples.

Small differentiated embryos are represented in *Tricocites* from the Deccan flora (see Zingiberales below).

Larger differentiated embryos with little endosperm include *Decodon* and *Enigmocarpou* (Lythraceae and extinct relative) (see Myrtales below) and *Keratosperma* (Araceae) from

the middle Eocene Princeton Chert (Cevallos-Ferriz and Stockey, 1988a,b; Cevallos-Ferriz *et al.*, 1991) and *Trema* (Ulmaceae) from the middle Eocene of Clarno which has a large coiled embryo with a prominent curved radicle. A weakly differentiated (possibly immature) embryo not associated with endosperm is preserved in *Palaeorosa* Basinger (Rosaceae) from the Princeton Chert (Cevallos-Ferriz *et al.*, 1993).

Embryos filling the seed and fully differentiated are found in Sapindaceae and *Juglans* from the Eocene of the London Clay and Clarno. The Sapindaceae are diverse and represented by several clearly distinct forms (distinct at generic level and with several morphologies within each form genus). Each shows a fully differentiated radicle and paired folded cotyledons comparable with modern members of the family. In the case of *Juglans* the nature of the embryo with two large relatively smooth cotyledons is inferred from chalcidony casts of the embryo cavity (Manchester, 1994b) which conform to those of living relatives.

Although relatively limited, these data show examples of a range of strategy in the early Palaeogene. At one extreme this variety includes potentially dormant embryos and embryos with slow germination as the food reserve is mobilized from a large endosperm store to the embryo. However, the examples are ruminant, thus increasing the surface area for this mobilization. In contrast, there are larger differentiated embryos indicating faster germination and establishment. The bent or folded embryos, where the curved structures stretch on germination, increasing the chances of reaching the soil surface and placing cotyledons above adjacent vegetation or litter etc., are indicative of strategies enhancing the chance of rapid establishment (Bewley and Black, 1994).

Embryo and endosperm in the order Myrtales

In the case of the *Decodon* seeds, the relationships to the Lythraceae and to *Decodon* are confirmed by the reconstruction of two partial to complete whole plants from the middle Miocene (Kvacek and Sakala, 1999) and middle Eocene, (Little and Stockey, 2000). These also confirm the habitat of the fossils based on facies and anatomical evidence as being from aquatic to swampy conditions similar to the extant relative *D. verticillatus* (the former) and in submerged conditions with fluctuating water levels at the edge of a shallow water system (the latter). Seeds of *Decodon* and extinct relatives have an extensive Palaeogene fossil record (Tiffney, 1981; Cevallos-Ferriz and Stockey, 1988a; Manchester, 1994b).

The Cretaceous/Tertiary transitional Deccan fruit *Enigmocarpon* and its allied flowers are a very abundant element in the Deccan intertrappean flora. These have been compared to modern members of the Myrtales especially Sonneratiaceae and Lythraceae (Sahni, 1943; Shukla, 1944; Chitale, 1977; see also discussion in Friis and Crepet, 1987:161). In combination these two examples offer an opportunity to consider the evolution of embryo and endosperm in the order Myrtales.

Both *Enigmocarpon* (Sahni, 1943) and *Decodon allenbeyensis* (Cevallos-Ferriz and Stockey, 1988a) have relatively large developed, differentiated embryos with little endosperm or no endosperm. This is comparable to seeds of extant *Decodon* (Baskin and Baskin, 1998) and typical of Myrtales (Cronquist, 1981) showing that the modern strategy had developed by the Palaeogene. Unfortunately, other permineralized Myrtales do not exhibit embryo or endosperm preservation (e.g. berries of *Paleomyrtinaea* from the Eocene Princeton Chert and Palaeocene Almont floras, Pigg *et al.*, 1993). Further details on the fossil record of the order, including another partially reconstructed plant from a compression flora, are given by Manchester *et al.* (1998).

Embryo, endosperm and seed internal organization in the order Zingiberales

Rodriguez-de la Rosa and Cevallos-Ferriz (1994) described two types of late Cretaceous (Campanian) permineralized zingiberalean fruits from Mexico. The Formation in which these fossils were found contains dinosaurs and is biostratigraphically dated as latest Campanian (Kirkland *et al.*, 2000). One fruit, *Striatornata*, Rodriguez-de la Rosa and Cevallos-Ferriz, was judged to be closely related to the extinct *Spirematospermum* (represented by compression fossils of seeds and fruits from the late Cretaceous to Neogene, see summary in Manchester and Kress, 1993). Both of these genera are assigned to the Musaceae based on the presence of a chalazal chamber and hilar cavity. In *Striatornata* the embryo sac itself is represented by an internal mould but cells are preserved in the chalazal chamber. Middle Eocene permineralized seeds of *Ensete oregonense* Manchester and Kress (1993), (see also Manchester, 1994b, 1995) of the Musaceae, also contain a chalazal chamber and hilar cavity. In addition, a few specimens reveal preservation of an endosperm chamber containing a small, straight bulbous embryo suspended from the hilar end. A small cylindrical to bulb-shaped embryo in this position is characteristic for the Zingiberales (Manchester and Kress, 1993) while the enlarged chalazal chamber appeared to be diagnostic for the family Musaceae (Manchester and Kress, 1993) where it may function to assist germination. In modern Musaceae, the embryo is usually shorter than in other families of the order, as it is in the fossil *Ensete* (Manchester and Kress, 1993).

The other Mexican Cretaceous fruit, *Tricostatocarpon* Rodriguez-de la Rosa and Cevallos-Ferriz, was of uncertain affinity within the order, lacking the musaceous chalazal chamber. Other fossil seeds of uncertain affinity within the order, are permineralized specimens formerly assigned to *Musa* but which lack the chalazal chamber, originally described by Jain from the Cretaceous/Tertiary transitional strata of the Deccan Intertrappean Beds of India (Manchester and Kress, 1993). Fruits of *Callistemonites* Bande *et al.*, also from the Deccan, were assigned to Musaceae and are certainly similar to that illustrated by Manchester (1995) but, although seed morphology is well-preserved, no internal anatomical detail could be discerned (Bande *et al.*, 1993). The genus *Cyclanthodendron*, a reconstructed whole plant said to combine features of the Musaceae and the Strelitziaceae, bore the fruits described as *Tricoccites* (Bonde, 1985; Biradar and Bonde, 1990) and comes from the Cretaceous/Tertiary transition of the Deccan Intertrappean Beds. Seeds contained within these fruits reveal a large cellular endosperm containing a small elongate, cylindrical (ribbon-like in section), cellular embryo in which some possible differentiation was observed (Bonde, 1985). However, the fruit structure is considered as controversial and poorly known by Biradar and Bonde (1990) suggesting a need for re-investigation of the details.

Overall these fossils suggest that the internal organization characteristic of modern seeds of Zingiberales was in place by the late Cretaceous. Specific internal structures (e.g. large chalazal chamber), combined with embryo and endosperm preservation, proves organization comparable with modern Musaceae at least by the middle Eocene.

Seedlings

Vivipary in Rhizophoraceae

Direct evidence of establishment strategy comes from fossil seedlings. Viviparous embryo hypocotyls of *Ceriops* and *Palaeobruquiara* (Rhizophoraceae), first recognized by Chandler, were fully described by Wilkinson (1981, 1983). That work shows beyond doubt that vivipary was exhibited by early Eocene Rhizophoraceae and specimens are preserved at two localities (Herne Bay and Sheppey) of slightly different ages (Collinson, 1983b). The London

Clay flora is perhaps the only fossil flora likely to demonstrate this as rapid permineralization with anatomical preservation is required in a flora from a brackish to marine facies. Another possibility is the Deccan Traps where the mangrove palm *Nypa* is found (see below) but *Nypa* is very rare. In sites such as Bracklesham (Collinson, 1996), rich in *Nypa*, but with preservation in the form of carbonaceous compression, mangrove hypocotyls probably would not have been preserved, or if preserved might not be recognizable as such.

Tomlinson and Cox (2000) showed, by experimentation, that modern viviparous seedlings were able rapidly to erect after rooting if dispersed by floating on the seawater surface and then, as likely, stranded horizontally. This strategy raises the plumule above tidal influence. It leaves anatomical evidence in the form of reaction (tension) wood fibres and morphological evidence in a resultant hook-like base. Wilkinson (1981) emphasized that the fossils were fragments of the hypocotyl, although she indicated that 'the fragments most frequently found suggest that they come from the distal tip'. The fossil *Ceriops* from Sheppey is sufficiently abundant (one per visit, personal observation) to examine for evidence of self-erection, but studies would have to be accompanied by detailed anatomical work to demonstrate that the true base of the seedling was preserved. Collinson has not encountered a hooked *Ceriops* during years of collecting in the London Clay of Sheppey. Although specimens figured by Chandler (1961, 1978) and Collinson (1983a) are curved, this is likely to be merely normal curvature of the hypocotyl axis. Only one is strongly curved (Chandler 1978: plate 7 Figure 1) and Chandler (1978) stated specifically that this is broken and incomplete at the radicle end. Therefore, the erection strategy to raise the plumule, is not demonstrated in the Palaeogene embryos.

Vivipary in Nypa

The mangrove palm *Nypa* also exhibits vivipary today (Tomlinson, 1986; Farnsworth, 2000). Fossil *Nypa* fruits have been shown to be exactly comparable to modern fruits in many ways (Collinson, 1993). The only obvious difference being that the fossil fruits reached larger maximum sizes than the modern fruits. However, in all *Nypa* fossils, even permineralizations, the seed-containing cavity has a mineral (amorphous silica, micro or macrocrystalline pyrite) or, rarely, sediment infill. This is the case for the London Clay flora based on many different workers who have studied hundreds of specimens (Bowerbank, 1840; Reid and Chandler, 1933; Chandler, 1961; Collinson, 1993, personal observation); and for the Belgian Brussels Sands specimens where again, many have been studied (Stockmans, 1936; Collinson, 1993). It is also the case for the Deccan Intertrappean flora, although only four specimens have been studied anatomically and *Nypa* is apparently very rare in this flora (Sahni and Rode, 1937; Chitale, 1960a,b; Nambudiri, 1966; Chitale and Nambudiri, 1969). Three-dimensionally preserved specimens with carbonaceous preservation, e.g. from the Eocene of Bracklesham, West Sussex, England (Collinson, 1996) have sediment infill in the seed cavity.

These *Nypa* fossils have a basal aperture (Reid and Chandler, 1933; Sahni and Rode, 1937) which is comparable with the germinal aperture in modern fruits. Mature modern fruits have a small undifferentiated embryo which then differentiates while still attached to the fruiting head. The detachment of the fruits from the fruiting head is assisted by the growth of the plumule, which breaks through the fruit at the base (Tomlinson, 1986). The fact that many fossils are found dispersed is not in itself proof of shedding through viviparous seedling growth because many aborted fruits, lacking seeds or former seed-containing cavities, are also found fossil (Collinson, 1993, 1996; Pole and MacPhail, 1996).

There are several explanations why embryo preservation has not been documented in any permineralized *Nypa* fruit specimens. First, once the plumule has penetrated the fibrous

mesocarp at the base of the fruit, the germinal aperture is formed. At this stage, failing establishment, the soft tissues would be readily vulnerable to decomposition. Secondly, once established and the food resource used up, the fruit could be released from the seedling yielding an empty husk with high preservation potential. Thirdly, it should be noted that relatively few of the mineralized specimens have actually been cut and polished or thin-sectioned so that future work might reveal embryos in *Nypa* fossils. Such work on some silicified specimens from Curacao is in progress (Collinson). In conclusion, although vivipary cannot be proven in the case of *Nypa* fossils, the lack of seed and embryo preservation in many fossils is consistent with a viviparous habit comparable with the modern *Nypa*.

Other seedlings

Numerous (over 8000) fossil seedlings have been found in a thin layer, traced laterally for 50–60 m in the Palaeocene of Joffre Bridge in Canada; 99% of these seedlings are of *Joffrea*, a Cercidiphyllaceae (Stockey and Crane, 1983), reconstructed as a partial whole plant (Crane and Stockey, 1985). Other seedlings (about 100 specimens) are platanaceous (Stockey and Crane, 1983; Crane and Stockey, 1985; Pigg and Stockey, 1991). Almost all seedlings are preserved upright in growth position. They are preserved at many different stages of development but none have more than three pairs of leaves suggesting that they all germinated at approximately the same time. Germination was epigeal producing an emerging radicle, thin hypocotyl and two cotyledons 2–4 mm long (*Joffrea*) and epigeal yielding two cotyledons with other parts obscure (Platanaceae). These two plants have small seeds, produced in large numbers (especially *Joffrea* and relatives with small winged seeds) and their remains are frequently found in association in fluvial sediments. Together with the seedling biology these features suggest that the plants represent early successional communities, with rapid seedling establishment that occurred in disturbed habitats along stream margins (Stockey and Crane, 1983; Pigg and Stockey, 1991). Hoffman and Stockey (1999) place the seedling bed at Joffre Bridge in detailed sedimentological context which supports this interpretation. *Joffrea* seedlings have also been recorded, along with seedlings of a taxodiaceous *Metasequoia*-like plant, in the Palaeocene of two other Palaeogene localities in Canada in comparable situations (Falder *et al.*, 1999).

Dormancy versus germination

Categories of dormancy can be generally summarized as follows (Bewley and Black, 1994: Chapter 5; Baskin and Baskin, 1998: 28)

Endogeneous (sensu Baskin and Baskin); broadly equal to embryo controlled

Physiological – physiological inhibition

Morphological – underdeveloped embryo; released by suitable conditions for growth

Morphophysiological – the above combined.

Exogenous = coat imposed/enhanced

Physical – coat impermeable to water; released by specialized opening mechanism

Chemical – germination inhibitors; released by leaching

Mechanical – restriction (physical constraint) on growth by woody tissues; released by stratification.

Germination may be defined as beginning with water uptake and ending with axis (usually radicle) emergence from the seed (Bewley and Black, 1994). Direct fossil evidence of radicle emergence during germination does exist but is very limited (see above). Fossil

evidence of vivipary and of other seedlings (see above) also prove germination mechanisms in fossils. Preserved embryo evidence (see above) clearly shows differentiated embryos which are unlikely to have had morphological dormancy as well as probable dormant seeds with poorly developed embryos.

Exogenous, coat-imposed/enhanced, dormancy can act by interference with water uptake, mechanical restraint, influencing gas exchange, preventing exit of inhibitors (from/to) embryo and supplying inhibitors. In exogenous dormancy the fruit or seed coat is extremely important. To quote Boesewinkel and Bouman (1995:9): 'The seed coat is not just a protective covering but a multifunctional organ that transports, transforms and secretes metabolites and oxygen to the embryo sac'. The fossil record provides a huge amount of evidence on the fruit and seed coat.

Hard seed coats have long been considered to be a widespread cause of dormancy. Many fossils have hard seed (or endocarp) coats so it seems highly likely that many (or at least some) were probably dormant. Direct evidence of the specialist opening mechanisms that are associated with physical dormancy, as defined above, is readily available and widespread in the Palaeogene fossil record. These include plugs or valves in endocarps of Anacardiaceae, Mastixioid and other Cornaceae, Nyssaceae, Potamogetonaceae and in seeds of Nymphaeaceae (Collinson, 1983b; Mai, 1993, 1998; Manchester, 1994b; van Bergen *et al.*, 1996; Tiffney and Haggard, 1996; Stockey *et al.*, 1998; Manchester *et al.*, 1999). Maastrichtian (latest Cretaceous) mastixioid Cornaceae are also recorded (Knobloch and Mai, 1986, 1991).

Bewley and Black (1994) cite experiments that indicate the importance of the osteosclereid layer beneath the palisade layer where, in dormant seeds, puncture of the osteosclerieds is necessary for germination to occur. However, most of this experimental evidence is derived from agriculturally important plants, few of which have a Palaeogene record. Therefore, this experimental evidence cannot easily be extrapolated to the past.

Columnar palisades and thick tough seed/endocarp coats with potential for imposing or enhancing coat dormancy, are very common in Cretaceous and Palaeogene seeds. However, these are very variable and the relative potential for dormancy conferred by each of the following, for example, is uncertain:

- (i) Extremely close-packed columnar palisades (e.g. modern legumes) are represented in endocarp walls such as Neogene *Cinnamomum* (Pingen *et al.*, 1994) and many Lauraceae endocarps are known in the Palaeogene. There is an extensive record of fossil legumes (Herendeen and Dilcher, 1992). However, data on seed coat anatomy seem to be lacking, seeds are generally poorly preserved and much of the work has relied on fruits. In their study of *Nelumbo* disseminule preservation, van Bergen *et al.* (1997) showed that a closely packed columnar structure need not imply high chemical resistance and argued that a distinctive tannin polysaccharide composition might account for the fossil record lacking *Nelumbo* disseminules while *Nelumbo*-like leaves occur from the Cretaceous onwards. It is possible that legume seed-coat chemistry may have affected the preservation potential of legume seeds.
- (ii) Closely packed columnar sclerotesta, e.g. Nymphaeaceae, widespread in the Palaeogene which, if water tight, would aid aquatic dispersal and could also confer dormancy (Collinson, 1980; Cevallos-Ferriz and Stockey, 1989; Cevallos-Ferriz *et al.*, 1991; van Bergen *et al.*, 1996).
- (iii) Rather porous or open columnar sclerotesta, where palisade cells have larger lumina than in i, or ii, as, for example, in some Nymphaeaceae (Collinson, 1980; van Bergen

et al., 1996) and Moraceae (Collinson, 1989) and also commonly present in Cretaceous seeds (Friis *et al.*, 1999, 2000).

- (iv) More equiaxial as opposed to columnar sclereids (e.g. *Toricellia* endocarp, Collinson 1988 (as ?Lythraceae); Meller and Collinson unpublished data).
- (v) Varying thickness of these layers, which are obviously relatively thinner, (though not necessarily proportionately thinner in terms of overall seed dimension) in smaller Cretaceous seeds.

Overall, the common occurrence of specialized plugs and valves as well as distinctive dehiscence patterns, combined with thick, tough, compact seed (or endocarp) coat organization, combined with the presence of fossils of both germinated and non-germinated but of mature size seeds (Collinson, 1999), all argue strongly for the presence of coat-induced dormancy of a wide diversity in Palaeogene seeds. Until more detailed studies are made on large assemblages of Cretaceous seeds, with this question in mind, the evidence prior to the Palaeogene is less clear but suggestive of the existence of fewer specializations at that time.

Of course we cannot easily address seed longevity in these fossil assemblages as we do not know if the germinated fossils germinated immediately on shedding or how long the last to germinate existed prior to germination. The more recent fossil/archaeological record does have the potential to address longevity in the natural environment (see review in Bewley and Black, 1994). There are claims for Arctic lupin to 10 000 years and *Nelumbo* to 1700 years (but both have no direct dating). Reliable evidence would depend on accurate dating of associated materials or of a molecule from the seed coat that has not been metabolically active thus avoiding recycled carbon. Shen-Miller *et al.* (1995) reported a *Nelumbo* seed which was germinated and then destroyed for dating giving an age of 1288 ± 271 years old. Another specimen, the plant of which was still growing in 1994, was dated from a fragment of pericarp as 332 ± 135 years old. As van Bergen *et al.* (1997) have shown the *Nelumbo* disseminule wall is not only physically very tough but also has a distinctive tannin/polysaccharide composition which may play a role in seed longevity. Legume seeds from herbarium sheets of known age, were shown to germinate after some 150 years when affected by fire extinguishing water at the British Museum of Natural History. The best-authenticated large scale example seems to be the Beal experiment where, of 21 species buried, only one retained viability after 100 years (1979), although several extended to 35–50 years. (Bewley and Black, 1994). However, this experiment was undertaken mainly on small-seeded, economically important arable plants or weeds and so gives little indication as to the link between seed coat anatomy and longevity in the vast majority of angiosperms.

A broadly-based study of the relationship between the anatomy of the seed (or endocarp) coat, the possession of specialist physical opening structures and other physical attributes of the seed coat and the longevity of seeds or their dormancy mechanisms would be of considerable value. In this regard the fossil record is able to provide direct evidence of the location, morphology and structure of the physically and chemically resistant layers (which survive as fossils). These include cuticles and thick-walled sclerotic tissues (Figures 18.2 and 18.3) both of which are common constituents of fossil seeds and endocarps (Collinson, 1980, 1983b; Collinson and Gregor, 1988; van Bergen *et al.*, 1996, 1999; and see the following section).

Chemical composition of resistant layers

Fossil fruits and seeds highlight those layers which are sufficiently resistant (e.g. to decomposition) to have survived in the fossil record. These layers can, therefore, be argued to have had a function in life related to their survival ability, such as in dormancy (physical

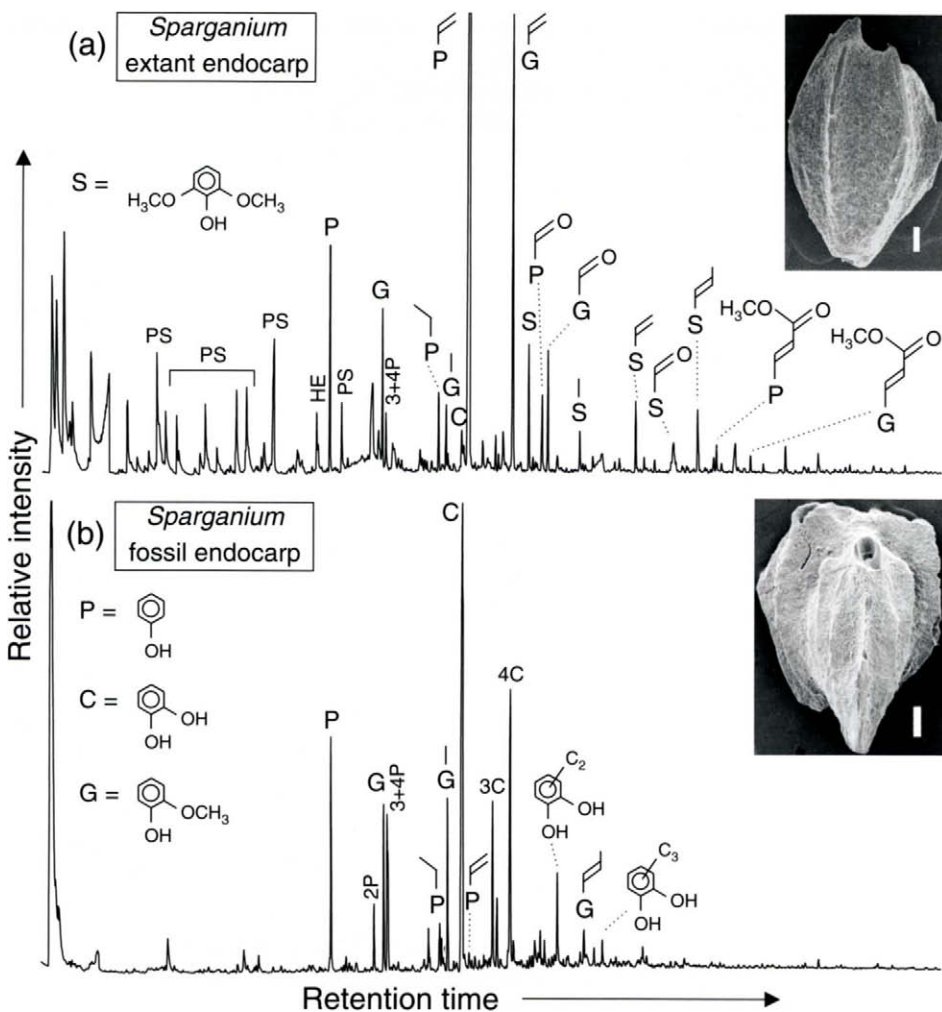


Figure 18.2 Total ion chromatograms of the pyrolysates of sclerotic endocarps of *Sparganium* (a) modern and (b) fossil showing a typical modern lignin–cellulose–hemicellulose complex and its modified condition in Eocene material with loss of PS (polysaccharides) and HE (hemicelluloses) but with the retention of most lignin markers, i.e. G (guaiacyl) and P (*p*-hydroxyphenol units). S = Syringyl and C = Catechol units. 2P = 2-methyl-phenol; 3 + 4P = co-eluting 3-methyl- and 4-methyl-phenol. Scale bars on SEM illustrations of endocarps represent 500 μm .

constraint, impermeability to gas or water, inhibition etc. see above) or for protection (from physical or biological hazards). Fossil fruits and seeds often include both cuticular layers and thick-walled sclerotic tissues. These fossils therefore provide perfect packages in which the chemical composition of both can be studied. Examples include *Sparganium* endocarps figured herein (Figure 18.3c); *Potamogeton* and *Limnocarpus* (van Bergen *et al.*, 1999) and Nymphaeaceae (van Bergen *et al.*, 1996). Resin composition has also been documented for fossil endocarps of mastixioid Cornaceae (van Aarssen *et al.*, 1994).

The different coats of fossil fruits and seeds retain distinctive chemical signatures, which can be recognized in modern relatives both in cuticles and in thick-walled, sclerotic tissues. This has been shown for Nymphaeaceae (van Bergen *et al.*, 1996) and for *Stratiotes*

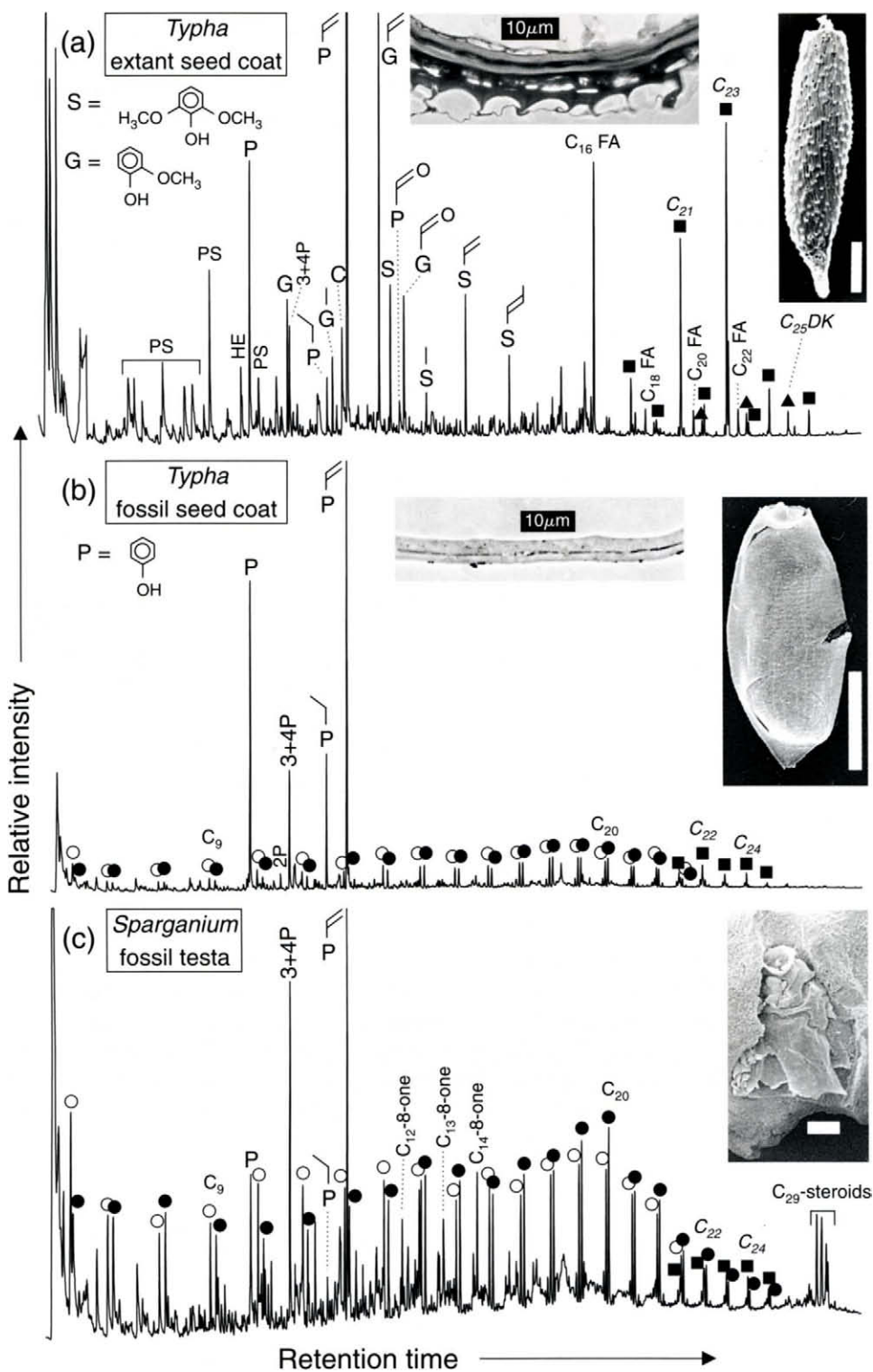


Figure 18.3 Total ion chromatograms of the pyrolysates of the extant *Typha* seed coat (a) dominated by monocotyledonous lignin markers (peaks labelled P and G) and the fossil seed coats of

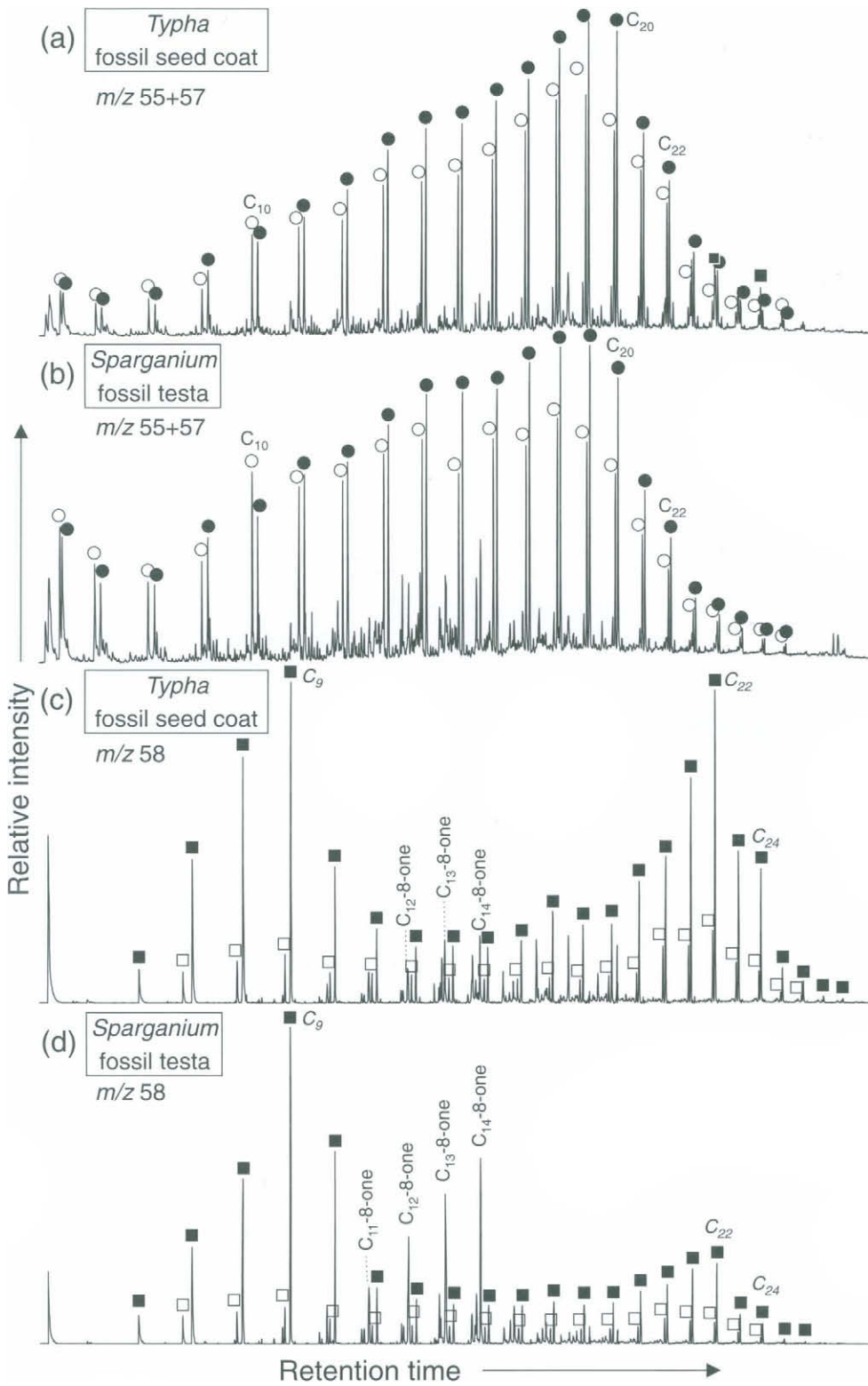
(van Bergen *et al.*, 1999). These results demonstrate that the particular biosynthetic pathways producing resistant fruit and seed coats, which exist today, also existed in the past (at least as far back as mid-Palaeogene). The results also add a new dimension of chemosystematics to the study of fossil seeds. For example, *Potamogeton* and the extinct *Limnocarpus* show clear chemosystematic relationship, which supports that previously based only on morphological comparisons (van Bergen *et al.*, 1999).

Here we report new results from *Sparganium* and *Typha* (families Sparganiaceae and Typhaceae, together order Typhales *sensu* Dahlgren *et al.*, 1985; and family Sparganiaceae, order Poales *sensu* APG, 1998). The fossil samples are derived from the Bembridge Limestone Formation, Gurnard Ledge, Thorness Bay, Isle of Wight, from the top 20 cm of Daley and Edwards (1990) bed 6C (see Collinson and Cleal, 2001a for details of the flora and references) and thus are from the Eocene/Oligocene transitional interval, probably latest Eocene in age.

The chemical composition of the modern and fossil resistant disseminule walls was examined by pyrolysis, which is a destructive method using thermal decomposition to yield characteristic building blocks (van Bergen, 1999). Details of methods are as reported in van Bergen *et al.* (1996, 1999). The fossil *Sparganium* sclerotic endocarp shows, as expected, modification of the lignin-hemicellulose complex (compared with the modern endocarp) with the loss of PS (polysaccharides) and HE (hemicellulose) but with the retention of most lignin markers, i.e. G (Guaiacyl) and P (*p*-hydroxy-phenol) units (See Figure 18.2). The chromatogram of the pyrolysate (Figure 18.2b) is quite distinct from that of the sclerotesta of fossil *Stratiotes* (order Alismatales Dumort. *sensu* APG, 1998) reported by van Bergen *et al.*, (1999) from the same fossil assemblage.

The fossil *Typha* and *Sparganium* seed coats are thin cuticles and their total ion chromatograms of the pyrolysates are dominated by alkene-alkane doublets (see Figure 18.3), which are long straight chain aliphatic hydrocarbon units that are typical of fossil cuticles (see Chapter 8). In addition, there are phenolic pyrolysis products, most probably derived from *p*-coumaric acid moieties. The extant *Typha* seed coat, in contrast, is multilayered, including cuticular layers and cellular layers. The chromatogram is dominated by monocotyledonous lignin markers (e.g. peaks labelled P and G on Figure 18.3a) which swamp out a small signal from the cuticles which can be detected in appropriate mass chromatograms. Detailed mass chromatograms (Figure 18.4) are essential to provide chemical evidence for similarities. These show that the fossil *Typha* and *Sparganium* seed coats are strikingly similar in distribution patterns of specific constituents such as the alkanes (sharp drop off after C₂₀ and C₂₂) and the 2-alkanones (sharp drop off after C₂₂ and C₂₄). These latter compounds are also characteristic long-chain aliphatic units in cuticular membranes and in combination with alkanes/alkenes provide crucial evidence of chemical similarities. In addition, there is a distinct group of 8-alkanones which are dominated by the C₁₄-8-alkanone

Typha (b) and *Sparganium* (c), dominated by alkene-alkane doublets with phenolic pyrolysis products most probably derived from *p*-coumaric acid moieties. The difference between fossil and modern is largely explained by the fact that the modern *Typha* has a multilayered seed-coat as shown in the transverse section, while the resistant material which survives in both the fossils is translucent cuticles only (as shown in the fossil *Typha* transverse section). Entire seed cuticles are illustrated by scanning electron micrographs; that of *Sparganium* is in place within a fractured endocarp. Transverse sections are transmitted light micrographs. Open circles: *n*-alk-1-enes, closed circles: *n*-alkanes, closed squares: 2-alkanones (= methyl ketones), Closed triangles: diketones, other abbreviations as in Figure 18.2. Scale bars on SEM illustrations of specimens represent 250 μm.



in both taxa. Furthermore, these mass chromatograms are very different from those documented in cuticular seed coats of *Potamogeton* and *Limnocarpus* (order Alismatales Dumort. *sensu* APG, 1998) from the same fossil assemblage where alkanes show drop off at C₂₆ and 2-alkanones (=methyl ketones) at C₂₈ (van Bergen *et al.*, 1999).

Fossils readily reveal resistant fruit wall and seed coat layers, the morphology and anatomy of which can easily be studied. Our results, from a variety of flowering plants (Nymphaeaceae, Potamogetonaceae, *Stratiotes*, *Typha* and *Sparganium* – see above), demonstrate that markers of chemosystematic and biosynthetic significance are surviving in Palaeogene fossils. These results indicate the potential to investigate any links (in fossils and their living relatives) between chemical composition, physical nature and the anatomy of fruit and seed coats and their functional role. This new combined information can be applied, in future, not only to understand the evolution of exogenous dormancy via the fossil record but also further to understand controls on dormancy and longevity at the present day.

Conclusions

The aim of this chapter was to document the grade of evolution of dispersal biology and ecophysiological strategies that had been attained by the Palaeogene and to compare this, where possible, with that in the preceding Cretaceous. The following are the main conclusions from this study.

Plumes, which may enable ‘floatation’ in air or may assist in orientation on landing, were lacking in Cretaceous floras and were of extremely low diversity in the Palaeogene. Only one form, *Apocynospermum*-like with a long tufted plume, is reasonably widespread, although rare, in Palaeogene floras. Taphonomic bias is unlikely to explain this low diversity which may be related to higher humidities which would not favour plumed disseminules or to phylogenetic constraints (many plumed disseminules occur in clades with first fossil occurrences in the Neogene).

In contrast to plumes, wings, which may confer aerodynamic properties favouring varied dispersal trajectories and behaviour, were very widespread, very abundant and highly diverse in Palaeogene floras (See Figure 18.1). A wide variety of wings may be seen within a single family, e.g. Juglandaceae and a given wing morphology (e.g. ‘helicopters’) may be found across a wide variety of taxonomic affinities. Wings were derived from a wide variety of different parts of the inflorescence, flower, fruit wall and seed coat. The variety of wing morphology, which was established by the Palaeogene, encompasses many, if not most, of the modern strategies. By comparison wings were insignificant in preceding Cretaceous floras.

Small seeds, including dust and microseeds, were dominant throughout the Cretaceous. This may be as much related to parent plant stature as to a specific dispersal strategy. There are clear examples where clades with larger seeds in the Palaeogene and Recent had small seeds in the Cretaceous, thus excluding a simple phylogenetic constraint. Dust seeds are

Figure 18.4 Mass chromatograms of the pyrolysates of the fossil *Typha* (a, c) and *Sparganium* (b, d) seed coats which demonstrate a striking similarity in distribution patterns of specific constituents such as the alkanes (sharp drop off after C₂₀ and C₂₂) and the 2-alkanones (sharp drop off after C₂₂ and C₂₄). In addition there is a distinct group of 8-alkanones, which are dominated by the C₁₄-8-alkanone in both taxa. Open circles: *n*-alk-1-enes, closed circles: *n*-alkanes, closed squares: 2-alkanones (= methyl ketones), open squares: monounsaturated 2-alkanones, other abbreviations as in Figure 18.2.

also found in Palaeogene floras, representing families and often genera that have similar seeds today. Potential dispersal by floatation in water is indicated by the fibrous fruit wall of Palaeogene *Nypa* fruits.

Like plumes, spines on disseminules are very rare in Palaeogene (and in Cretaceous) floras. Only one example can be convincingly argued to be adapted for animal dispersal (a small fruit entwined in mammal hair within late Eocene amber) in spite of the logical origin for mammalian fur in the late Triassic. Spines may also function in anchoring to other substrates such as wet mud.

Large dry nuts and large fleshy fruits exhibit an Eocene radiation, the former coincident with a radiation of rodents. Seed predation (gnawing by rodents; crushing by mammal teeth and boring by insects) and survival of seeds from fleshy fruit in mammalian gut contents both indicate interactions, including potential dispersal, by mammals. However, these examples are still rare in the Palaeogene. As Cretaceous fruits and seeds were all so small, large dry nuts were lacking. Furthermore, fleshy tissue attractants or food resources and indeed food resources from seed contents were at an utterly different, very small scale compared with those of the Palaeogene.

Rare Cretaceous and Palaeogene fossils prove radicle emergence and seedling strategy including rapid simultaneous germination from small seeds in Palaeogene examples. Preserved (permineralized) embryo and endosperm in Palaeogene fossils indicates a wide range of strategies, comparable to those of the present day. These range from underdeveloped potentially dormant embryos including those with extensive ruminant endosperm tissue, to large embryos, including those which are fully differentiated, filling the seed and having folded cotyledons adapted for rapid establishment. In the very few examples where several members of one order can be assessed (e.g. Myrtales, Zingiberales), the indications are that modern strategies already existed in Palaeogene or even in Late Cretaceous examples. Vivipary is also documented in the Palaeogene.

Fossil fruit walls and seed coats provide a wealth of information pertinent to the further understanding of the evolution of dormancy. Physical dormancy, released by specialized opening mechanisms, is widely indicated in Palaeogene floras by the presence of distinct plugs and valves and germinated and non-germinated but mature seeds. This strategy was present in at least a few late Cretaceous examples but is much more common in the Palaeogene. Resistant layers, which are evident by their preservation as organic compression fossils, have the potential to confer not only physical but also mechanical and chemical dormancy. These layers include both sclerotic and cuticular materials in fruit walls and seed coats. Palaeogene sclerotic tissues vary considerably in their compactness and sclereid form (columnar to equiaxial) as well as in thickness, whereas earlier Cretaceous examples tend to be thin, single-layered and with sclereids with relatively open lumina. This may indicate differences in dormancy potential. Surprisingly, in spite of the excellent Palaeogene record of legume fruits, containing evidence of the former presence of seeds, no well-preserved legume seed coat (the classic columnar palisade of text-books) is known to us. This might indicate a distinctive underlying chemistry as we found in columnar sclereids of modern *Nelumbo* disseminules.

Resistant cuticular layers, which may function in dormancy as physical constraints or impermeable barriers, are particularly easily revealed in fossils. An assemblage of fossils with examples comprising both sclerotic and cuticular layers provides ideal packages in which to study the chemical composition of fossils because variations due to conditions of fossilization are eliminated. Our studies of Palaeogene fossils, including new data on co-occurring *Sparganium* and *Typha* reported here, show that different taxa and different tissue types carry distinctive chemical signatures indicative of underlying distinctive

biosynthetic pathways. These distinctive chemistries may all have played a role in mechanical and chemical (and possibly endogenous) dormancy in the Palaeogene. In combination, morphology, anatomy and chemistry of the resistant layers of disseminules have future potential as a powerful tool to investigate aspects of fruit and seed physiology in the past and in the present. This combined approach would not only improve our understanding of the evolution of fruit and seed ecophysiology but also help to understand controls on dormancy and on seed longevity at the present day.

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Appendix 18.1 Palaeogene Plumed Disseminules

Apocynospermum-like plumes

Fossil seeds named *Apocynospermum* Reid and Chandler, *Cypselites* Heer and *Echitonium* Unger represent seeds similar to those in modern Apocynaceae and Asclepiadaceae (Reid and Chandler, 1926; Hably *et al.*, 2000: 65). These occurred in the Eocene Green River flora of North America (e.g. Brown, 1934: pl. 10 fig 3; MacGinitie, 1969: pl. 18 fig 4); the Eocene of Clarno and Florissant, North America (Manchester, 1999); the Eocene/Oligocene transition flora of England (Reid and Chandler, 1926; Collinson and Cleal, 2001a); the middle Eocene of Eckfeld (Wilde and Frankenhäuser, 1998: pl. 18 fig 12); the middle Eocene of Messel, Germany (Manchester, 1999: 476) and the late Eocene to early Oligocene floras of Hungary and the Czech Republic and the middle Miocene of Switzerland and Czech Republic (Kvacek and Walther, 1998; Manchester, 1999; Hably *et al.*, 2000; Sakala, 2000). Rather similar seeds were assigned to *Phyllanthera* (Asclepiadaceae) by Reid and Chandler (1926). Together these demonstrate one plumed seed morphology (possibly including several species, genera or even more than one family) in the Palaeogene, very uncommon but widespread, sometimes represented by just a single or few specimens though Reid and Chandler (1926) had 15 *Apocynospermum* specimens (13 with coma of hairs preserved) and also 6 (2 with coma) which they assigned to *Phyllanthera*.

In the terminology of Hoerner, applied by Burrows (1986), this terminal coma of long simple uniseriate hairs (Reid and Chandler, 1926) may act as a guide parachute rather than a drag parachute. A major functional role may have been in orientation on landing rather than influencing the aerodynamic properties of the seed.

Other plumes

Typha fossils are widespread and very common seeds (e.g. Collinson, 1983a; Mai, 1997 and references cited therein; Dettman and Clifford, 2000). Intact fruiting heads are known in the Eocene Green River flora (Grande, 1984: 265) and early Miocene infructescence fragments are figured by Mai (1999: Taf. 29 figs 5 and 6). These figures do not indicate any pappus hairs. Whereas isolated seeds are to be expected (Collinson, 1999), in view of the dehiscence of fruits, the fruiting heads should show pappus hairs on the fruits had they existed. However, I know of no detailed study of the Green River *Typha* fossils and Mai (1999, Taf. 29) refers to the infructescences in the plate explanation as 'Teile von fruchtständen. Achse mit ansitzenden Samen'.

Hairs are not preserved on the awns of the fruits named *Clematis* by Reid and Chandler (1926). The middle Eocene Clarno plane and platanaceous plants from the Canadian Palaeocene at Joffre Bridge (*Macginicarpa* Manchester) lack pappus hairs on the achenes as do earlier Cretaceous relatives (Pigg and Stockey, 1991; Manchester, 1994b). However, middle Eocene achenes from the Clarno flora and nearby sites possess pappus hairs and have been identified as the earliest true *Platanus* (Manchester, 1986, 1994b). We are not aware of any umbrella-like drag parachutes in Palaeogene floras.

Appendix 18.2 Palaeogene Winged Disseminules

Selected examples are illustrated in Figure 18.1 to demonstrate the variety of Palaeogene winged disseminules.

Betulaceae (see review in Chen *et al.*, 1999 and also Takhtajan, 1982)

Ostrya nutlets, surrounded by bladder-like involucre, occur in the early Oligocene of the USA and Europe and Miocene of USA, China and Japan (Meyer and Manchester, 1997; Manchester, 1999; Hably *et al.*, 2000). The extinct *Cranea* also produced a nutlet surrounded by an involucre and is known from the Palaeocene of the USA (Manchester and Chen, 1998).

The extinct *Asterocarpinus* (Manchester and Crane, 1987) produced four to seven-winged helicopters with *Carpinus*-like nutlets and occurred in the late Eocene and Oligocene of the USA (Manchester and Crane, 1987; Meyer and Manchester, 1997). The extinct *Palaeocarpinus* Crane (1981) produced small *Carpinus*-like nutlets with paired deeply dissected to entire margined bracts similar to those of modern *Corylus*. *Palaeocarpinus* was circumboreal in the Palaeocene, persisting into the Eocene of western North America and eastern Asia (Manchester, 1999).

The asymmetrical trilobed bracts of *Carpinus*, with basal nutlets, occurred in the middle Eocene and later in Europe (Wilde and Frankenhäuser, 1998; Manchester, 1999) and doubtfully in the USA (Manchester, 1999). *Betula*, with bilaterally winged fruits, is rare in the Oligocene of U.S.A, Europe and Asia (Hably *et al.*, 2000), though the earliest records of trilobed bracts (characteristic of *Betula*) are from the middle Eocene of the USA and Palaeocene of Asia (Manchester, 1999).

Juglandaceae (see Manchester, 1987, 1989a; Budantsev, 1994)

The extinct *Cruciptera* Manchester (1991) produced four-winged propeller-like samaras with central nutlets and occurred in the middle Eocene of Messel, Germany (Manchester *et al.*, 1994) and middle Eocene to Oligocene of western USA (Manchester *et al.*, 1994; Meyer and Manchester, 1997; Manchester, 1999, 2000a). The extinct *Palaeocarya* Saporta produced trilobed bracts with triveined lobes with a basal nutlet and occurred in the middle Eocene of Europe and North America and the Oligocene to Miocene of Europe and the USA (Hably *et al.*, 2000; Manchester 1999, 2000a). *Casholdia*, from Palaeocene/Eocene transitional strata in England and France, has an unlobed wing but is otherwise similar to engalhardioid fruits (Manchester, 1989a). *Pterocarya* has bilaterally winged nutlets and occurred in the early Oligocene and Miocene of the USA and Oligocene to Pliocene of Europe and Asia (Meyer and Manchester, 1997; Manchester, 1999). The extinct *Hooleyia* produced bilaterally winged nutlets with large wings and occurred in the middle Eocene of North America and at the Eocene and Oligocene transition of Germany, England and Hungary (Manchester, 1987; Wilde and Frankenhäuser, 1998). Bilaterally winged nutlets of *Platycarya* occurred in the early Eocene of England and North America (Manchester, 1999) while those of *Palaeoplatycarya* Manchester occurred in the early Eocene of the USA (Manchester, 1987).

Cyclocarya nutlets with large, entire disk-like encircling wings, occurred in the Palaeocene and Eocene of North America and in the Oligocene to Pliocene of Eurasia (Manchester, 1999; Hably *et al.*, 2000). The extinct *Polyptera* Manchester and Dilcher produced nutlets with large, multilobed, disc-like encircling wings and occurred in the Palaeocene of western North America (Manchester and Dilcher, 1997). *Polyptera* is the oldest unequivocal Juglandaceae and is known from a reconstructed partial plant. In spite of this wealth of wing variety in late Palaeocene and Eocene Juglandaceae, Cretaceous relatives of this family show no similar modifications for dispersal and have only small nuts (Friis, 1983; Crane and Herendeen, 1996; Manchester and Dilcher, 1997: 649)

Oleaceae

Fruits of *Fraxinus*, with flattened single-seeded samaras with lateral and apically extended wings, occurred in the middle Eocene and Oligocene of North America and the Oligocene of Europe and Kazakhstan. They became much more common in the Miocene of Europe and North America and Miocene to Pliocene of Asia (Call and Dilcher, 1992; Meyer and Manchester, 1997; Manchester, 1999; Hably *et al.*, 2000).

Tiliaceae and Tilioideae

Tilia bracts, subtending fruiting peduncles, are documented by Manchester (1994a) with the earliest record in the late Eocene of North America (Manchester, 1994a, 1999). Distinctive extinct round-bracted forms, with peduncle attached at the base (type A), occur in the early Oligocene of Oregon. Round bracted forms and oval bracted forms occur together in the Bridge Creek flora, early Oligocene USA (Meyer and Manchester, 1997). Type B bracts, ovate with peduncle fused at the base, occur in the late Eocene to Miocene of North America and Miocene to Pliocene of Europe. Type C bracts, ovate with peduncle fused midway like most extant species, occur in the Asian Oligocene onwards. Broad bracted forms also occur in the European Oligocene (Hably *et al.*, 2000).

Craigia (Malvaceae : Tilioideae) produced capsular fruit with five two-winged valves, often represented by single valves, formerly named *Koelreuteria*, *Ptelea* and *Pteleaearcypum* Weyland (Buzek *et al.*, 1989; Kvacek *et al.*, 1991, 2002; Pinggen *et al.*, 2001). The fruits are recorded in the Eocene and Oligocene of the USA, the Eocene to Miocene of Asia and the Oligocene to Pliocene of Europe (Manchester 1994a, 1999, 2000a; Meyer and Manchester, 1997; Hably *et al.*, 2000; and references above). Kvacek *et al.*, (1991) described fossil *Craigia* as three-winged and this led Vassiliev *et al.*, (1995) to question the inclusion of some Asian Miocene fossils in *Craigia*, arguing for an affinity with *Tripterygium* (Celastraceae) on the basis of the presence of three wings, while modern *Craigia* has five wings. However, continuing investigations and studies of new material have enabled Pinggen *et al.*, (2001) and Kvacek *et al.*, (2002) to demonstrate conclusively the five-winged nature of fossil *Craigia* fruits.

Some authors assign *Florissantia* Manchester to the Tilioideae or Malvaceae (see section on other winged disseminules).

Ulmaceae

Cedrelospermum (an extinct Ulmaceae known from a partial reconstruction of the plant) produced a small samara. It occurred in the Eocene and Oligocene of western North America, the Neogene of South Mexico and Eocene to Miocene of Europe (Manchester, 1989b,c, 1999; Magallón-Puebla and Cevallos-Ferriz, 1994b; Meyer and Manchester, 1997; Hably *et al.*, 2000; Hably and Thiébaud, 2002). Winged fruits of *Ulmus* are known from the late Eocene to Oligocene of the USA and the Oligocene to Miocene of Europe (Manchester, 1989b,c, 2000b; Meyer and Manchester, 1997; Hably *et al.*, 2000).

Aceraceae/Sapindaceae

Acer, large schizocarpic samaras with one-sided wings, are represented by at least six morphotypes in the early Oligocene Bridge Creek flora, USA (Meyer and Manchester, 1997). The earliest record is from the late Palaeocene of the USA (Crane *et al.*, 1990). There is an extensive Palaeogene record in North America and Eurasia but links between fruits and leaves to establish species diversity more accurately are still needed (Wolfe and Tanai, 1987;

Manchester, 1999; Hably *et al.*, 2000). *Deviacer* Manchester (1994b) produced schizocarpic samaras with backward orientation of fruits compared to *Acer* (Manchester, 1994b: text-fig 17) and occurred in the middle Eocene of the USA (Manchester, 1994b, 1999).

Dipteronia (sometimes previously named *Bohlenia*) produced schizocarpic samaras with encircling wing, paired or grouped in threes. Though the genus is now restricted to Asia in the Palaeocene it occurred only in the USA in the late Palaeocene, Eocene and Oligocene (Wolfe and Tanai, 1987; Meyer and Manchester, 1997; Manchester, 1999, 2000a; McClain and Manchester, 2001). Isolated valves of the inflated winged capsules of *Koelreuteria* occurred in the Eocene of the USA, Oligocene to Miocene of Europe and Miocene of China (Manchester, 1999).

Caprifoliaceae

Dipelta, large fruits with large bracts forming lateral wings and a smaller median wing, occurred in the Eocene/Oligocene transition of England (Reid and Chandler, 1926; Manchester and Donoghue, 1995; Collinson and Cleal, 2001a). The extinct *Diplodipelta* Manchester and Donoghue produced a samara-like, two-fruited infructescence with two larger bract wings and smaller median wings. *Diplodipelta* occurred in the late Eocene to Miocene of western USA (Manchester and Donoghue, 1995; Meyer and Manchester, 1997; Manchester, 1999).

Other winged disseminules

One-sided lateral wings

Seeds of *Saportaspermum* Meyer and Manchester (1997), sometimes named '*Embothrium*', (affinity unknown) are recorded from the Oligocene of the USA and late Eocene and Oligocene of Europe (Hably *et al.*, 2000).

Large winged seeds of *Gordonia* (Theaceae) are recorded from the middle Eocene and Oligocene of the USA and late Eocene of Europe (Grote and Dilcher, 1992). The related extinct genus *Gordonopsis* Grote and Dilcher lacked wings.

Small seeds of *Joffrea/Nyssidium* (Cercidiphyllaceae) are up to 5 mm long with an elliptic seed body and a curved terminal wing. These were widespread in the late Cretaceous to Eocene of North America; late Cretaceous to Palaeocene of Asia and Palaeocene to Eocene of Europe, Greenland and Spitsbergen (Crane, 1984; Crane and Stockey, 1985; Manchester, 1999; Feng *et al.*, 2000). *Cercidiphyllum* itself, with similar winged seeds, is known from the Oligocene onwards (Kvacek and Konzalova, 1996; Meyer and Manchester, 1997).

Winged *Liquidambar*-like seeds (Hamamelidaceae) are recorded from the Eocene of Washington State, USA (Wehr, 1995).

Fruits of *Pteronepyles* Manchester (1994b) (affinity unknown) are samaras with a laterally flattened endocarp and an elongate almost elliptical wing. They are recorded from the middle Eocene of Clarno and Republic, USA (Manchester, 1994b; Wehr, 1995).

Fruits of *Liriodendron* (Magnoliaceae) are two-seeded winged samaras. They are recorded from the Eocene/Oligocene transition onwards in Europe (Buzek *et al.*, 1976; Kvacek and Walther, 1995; Bellon *et al.*, 1998).

Encircling wings

Potanispira Meyer and Manchester (1997) (affinity unknown) were small, 7 mm diameter, winged seeds. They are recorded from the early Oligocene of western North America.

Small (up to 10 mm in maximum dimension) seeds of *Bekerosperma* Meyer and Manchester (1997) and Manchester (2000a) (affinity unknown) are recorded from the late Eocene and Oligocene of western North America.

The small seeds (6 mm) of *Pinckneya* (Rubiaceae) were surrounded by a reticulate membranous wing. They are recorded from the early Oligocene of the USA (Meyer and Manchester, 1997).

The small winged seeds of *Landeenia* (Sapindales) are recorded from the middle Eocene of Wyoming, USA (Manchester and Hermsen, 2000).

The small winged seeds of *Hydrangea* are recorded from the middle Eocene of Oregon, USA (Manchester, 1994b, 1999).

Fruits of *Eucommia* (Eucommiaceae) are large elongate samaras, with an asymmetrical wing surrounding a single-seeded fruit body. They are recorded from the Eocene of the USA and Asia, Neogene of South Mexico and Oligocene of Asia (Takhtajan 1974; Magallón-Puebla and Cevallos-Ferriz, 1994a; Call and Dilcher, 1997; Manchester, 1999, 2000a; Wang *et al.*, 2003).

Fruits of *Florissantia* Manchester (1992) (assigned to extinct Sterculiaceae by Meyer and Manchester, 1997; to Tilioideae by Hably *et al.*, 2000 and to the broadly circumscribed Malvaceae by Manchester, 2000b) have a round encircling wing formed from the persistent calyx. They occurred in the middle Eocene to early Oligocene of the USA (Meyer and Manchester, 1997; Manchester, 2000b) with one record in the Miocene of eastern Asia (Manchester, 1999).

Fruits of *Micropodium* are winged legume pods that occurred in the early Oligocene of the USA (Meyer and Manchester, 1997). Rather bladder-like fruits of the legume *Caesalpinia* subgenus *Mezoneuron* are recorded from the Eocene and Miocene of the USA and Eocene of England (Herendeen and Dilcher, 1991, 1992).

Fruits of *Paliurus* (Rhamnaceae) are large nuts with a round encircling wing, convergent with *Cyclocarya* (see Manchester, 1999, for distinguishing characteristics). *Paliurus* occurred in the early Eocene to Miocene of the USA, middle Eocene to late Miocene of Asia and Upper Oligocene and Miocene of Europe (Manchester, 1999).

Fruits of *Buzekia* Manchester (1999) of unknown affinity (formerly named *Pterocarpus*), had a large encircling wing. They are recorded from the Oligocene to early Miocene of Europe and Miocene of North America.

Fruits of *Ptelea* (Rutaceae) are two- (rarely three-) loculed samaras with an ovoid fruit body and two (rarely three) broad surrounding papery wings. They are recorded from the Mid-Miocene of Oregon-Idaho, USA (Call and Dilcher, 1995).

Bilateral wings

The small (6 mm long) seeds of *Catalpa* (Bignoniaceae) have transversely elongate striate wings (?composed of stiff coarse hairs), the striae radiating straight out from the sides of the seed body. They are recorded from the late Eocene and Oligocene of the USA (Meyer and Manchester, 1997; Manchester, 2000b); Eocene/Oligocene transition of England (Reid and Chandler, 1926) and Oligocene to Miocene of continental Europe (Hably *et al.*, 2000). Seeds assigned to *Radermachera* (Bignoniaceae) with long striate wings (15 mm broad) are recorded from the Eocene/Oligocene transition in England (Reid and Chandler, 1926; Collinson Cleal, 2001a). Bilaterally winged seeds assigned to Bignoniaceae are also recorded from the Eocene of Washington State, USA (Wehr, 1995).

Fruits of *Terminalia* (Combretaceae) have two wide longitudinal wings, lacking marginal veins and radiating from a fusiform body. They are recorded from the Eocene and early Oligocene of western USA (Meyer and Manchester, 1997).

Large fruits of *Ailanthus* have large twisted lateral wings. They are recorded from the middle Eocene to middle Miocene of Europe, the Oligocene to Miocene of Asia and Eocene and Miocene of the USA (Mosbrugger, 1996; Manchester 1999, 2000a; Hably, 2001).

Multiple wings ('helicopters/propellers')

Tetrapterys (Malpighiaceae) has a four winged fruit with a central nutlet. It is recorded from the Oligocene of Slovenia and Hungary (Hably *et al.*, 2000).

The extinct *Calycites* Crane (affinity unknown) was a fruit with six wings and an elongate fruit body. It is recorded from the Palaeocene of Scotland and Wyoming and middle Eocene of western North America (Crane, 1988; Manchester; 1999).

Raskya Manchester and Hably was a four-winged, hypogynous, propeller-like fruit, with a fusiform body. It is recorded from the latest Eocene to early Oligocene of England, Czech Republic, France and Hungary (Manchester and Hably, 1997; Thiebaut, 1999).

'*Abelia*'-like fruits (affinity unknown) are three to five winged propeller-like fruits originally described as *Abelia* but now all rejected as not belonging to this genus (Collinson and Cleal, 2001a: table 9.3). They are from the Eocene/Oligocene transition, England.

Chaneya Wang and Manchester (2000) (affinity uncertain, similar to some Simaroubaceae) was a five-winged accrescent calyx with one or more globose fruit bodies. It is recorded from the middle to late Eocene of the USA, the Eocene of China and the Miocene of eastern Asia. It is possibly congeneric with forms named *Porana* and *Monotes* in Europe (Wang and Manchester, 2000).

Ceratopetalum (Cunoniaceae), is a fruit with four to six woody sepals radiating from a central disc. It is recorded from the middle Eocene of South Australia (Barnes and Hill, 1999).

Multiple wings (longitudinally arranged)

Halesia (Styracaceae) has fruits with two to four longitudinal wings (with intramarginal veins) radiating from a spindle-shaped body. It is recorded from the late Tertiary of Europe (Manchester, 1999).

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The rise and fall of the Podocarpaceae in Australia – a physiological explanation

Tim Brodribb and Robert S Hill

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Introduction

Two large families dominate conifer diversity today. Remarkably, the distributions of these two families are almost perfectly separated by the equator, with the southern hemisphere dominated by Podocarpaceae and the northern hemisphere by Pinaceae. These families are ecologically and physiologically distinct and have apparently remained separate for many tens of millions of years. Reasons for this extraordinary pattern of hemispheric preference are likely to be associated with climatic factors, particularly the current predominance of highly seasonal continental type climate in the northern hemisphere compared with the oceanic climate in the southern landmasses where seasonal extremes are greatly attenuated.

In this chapter we will examine the history of the Podocarpaceae in Australia and discuss how the distribution and diversity of this family has changed since the rise to dominance of the angiosperms. Using evidence from the physiology of extant members of the Podocarpaceae, we will postulate how this family was able to compete successfully with angiosperms until later in the Cenozoic, when drying of the Australian continent led to large scale range contractions and extinction.

A physiological approach to the interpretation of the fossil record

The fundamental assumption in arguments we present here is that the physiology of living Podocarpaceae species can be taken as representative of closely related extinct fossil taxa. Clearly this assumption would be entirely unjustified when dealing with many angiosperm families, which show aggressive speciation and span large morphological ranges. An example of such a group is the Proteaceae, an extremely common family in Australia today and one that is typically associated with Podocarpaceae fossils. Unlike Podocarpaceae, clades within the Proteaceae today occupy a great range of environments, and exhibit enormous variation in their tolerance to environmental stresses (Beadle, 1981). Because of this, it is often difficult to use the physiology of living Proteaceae species to interpret conditions (including climate) that were prevailing when related fossil species were growing tens of millions of years ago (e.g. Carpenter *et al.*, 1994).

In contrast to families like the Proteaceae, Podocarpaceae species provide an ideal opportunity for nearest living relative research. The reasons for this are that the morphology of many extant genera have remained highly conserved over evolutionary time, so much so in some cases that fossil species from as far back as the Eocene or Early Oligocene are almost identical in morphology to extant species (e.g. the fossil *Microstrobos microfolius* P. Wells and R.S. Hill *cf.* the extant *M. niphophilus* J. Garden and L.A.S. Johnson (Wells and Hill, 1989) and the fossil *Phyllocladus aberensis* R.S. Hill *cf.* the extant *P. aspleniifolius* (Labill.) Hook. f. (Hill, 1989)). Secondly, clades within the Podocarpaceae are generally small and well resolved (Conran *et al.*, 2000), meaning that macrofossils that can be identified to genus are likely to exhibit a high degree of similarity to extant members of that genus. Finally, the community associations of podocarps have remained largely conserved over time and these associations typically involve other morphologically conservative taxa such as the conifer families Araucariaceae and Cupressaceae and the angiosperm families Nothofagaceae, Winteraceae and Cunoniaceae. Even at the extremes of their distributions, in regions such as Central America, podocarps are found associated with their palaeo-counterparts *Drimys* Forster and Forster f. (Winteraceae) and *Weinmannia* L. (Cunoniaceae). This further strengthens confidence that the physiology and even ecology of the extant species of Podocarpaceae are very similar to their fossil relatives.

There are currently 19 extant genera in the Podocarpaceae, although not everyone accepts this (Hill and Brodrigg, 1999). Most extant species belong to *Podocarpus* L'Hér. ex Pers., but there is evidence to suggest that *Podocarpus* may not be monophyletic (Kelch, 1997). The morphological diversity among extant Podocarpaceae is large, although its ecological and environmental range is relatively small, with almost all species restricted to rainforest or wet montane environments. Within these forests regeneration can take the form of either a continuous recruitment of shade tolerant seedlings (Midgley *et al.*, 1990; Lusk and Smith, 1998) or in less shade tolerant species it is reliant on small scale disturbances or topographic features such as rivers or ridgelines to open the canopy. A few Podocarpaceae possess leaves that are imbricate and microphyllous and they grow as low shrubs in sub-alpine vegetation across the full range of southern latitudes available for woody vegetation.

The fossil record of the Podocarpaceae

The Podocarpaceae are one of the oldest of the extant conifer families, with several Early Triassic species reported (e.g. Townrow, 1967; Axsmith *et al.*, 1998). More research is

needed to clarify the relationships of the earliest probable podocarp fossils, but the inference is that the Podocarpaceae or near relatives may have been widespread and important early in the Mesozoic. Podocarps were still prominent in the southern hemisphere in the Cretaceous and they are usually assigned to extinct genera. There are other Mesozoic and Cenozoic macrofossil records of the Podocarpaceae from the northern hemisphere that require further research (e.g. Dilcher, 1969; Krassilov, 1974; Reymanówna, 1987; Kimura *et al.*, 1988), but the fossil record of the family in the Cenozoic is overwhelmingly southern hemispheric.

The Cretaceous and Cenozoic Podocarpaceae macrofossil record is mainly based on vegetative material, but many diagnostic vegetative characters of the extant genera fossilize well, thus enabling reliable identification of most fossils. Among the extant genera, only *Afrocarpus* (J. Buchholz and N.E. Gray) C.N. Page, *Halocarpus* Quinn, *Nageia* Gaertn., *Parasitaxus* de Laub. and *Saxegothaea* Lindl. have no macrofossil presence in the Cenozoic.

Podocarpaceae diversity through time

The macrofossil record for south-eastern Australia is now well enough understood to be able to attempt a preliminary assessment of the diversity of Podocarpaceae through time. This can only be done for the Cenozoic, since relevant Cretaceous fossils are sporadic. Every Cenozoic site that has been reasonably well documented has been included (Table 19.1), but the data from each locality are not always strictly comparable for a variety of reasons, including:

1. Research is incomplete on most sites, so some species numbers underestimate reality
2. Some localities represent limited catchments, whereas others (notably the Latrobe Valley coal) represent very large catchment areas and several different vegetation types
3. Some sites probably represent relatively small time intervals, whereas others represent long periods of continual fossil accumulation and therefore a greater potential for mixing of different community types.

The overall picture is of a high diversity of Podocarpaceae at cooler locations throughout the region from the Late Paleocene through to the present day, although there has been an obvious decline in diversity since the peaks of the Late Eocene–Early Oligocene. However, with the possible exception of Anglesea, the warmer Middle Eocene sites on mainland south-eastern Australia stand as an anomaly, with very low Podocarpaceae diversity. Podocarpaceae today enjoy only limited success in such lowland broad-leaf forest and this may be due to mechanical limitations on the maximum size of leaves and shoots. Although it has not been tested, it is probable that the cost of increasing leaf or shoot size in Podocarpaceae is higher than in angiosperms given the common limitations of having only a single vein per leaf and relatively low specific leaf area. This would result in podocarps being unable to reach the maximum growth potential of angiosperms and thus being excluded from highly productive forest. *Nageia* (the only podocarp with multiveined leaves) come closest to competing with angiosperms in these environments. The single tropical species of *Phyllocladus* Rich. ex Mirb., with its multiveined phylloclades, also competes with angiosperms. Other species in the Podocarpaceae have alternative strategies to allow them to compete and these are outlined later. The overall difference between diversity and range of Podocarpaceae in the Palaeogene and today is clear, with species numbers uniformly much lower today (see Table 19.1), and this is matched by range decreases.

Podocarpaceae shoots are rare in the Australian Cretaceous, but those that have been recovered usually have narrow, imbricate foliage. The first appearance of podocarps with

Table 19.1 Podocarpaceae diversity in selected Cenozoic macrofossil localities in south-eastern Australia (modified from Hill and Brodrigg, 1999).

| <i>Site</i> | <i>Age</i> | <i>Fossil Podocarpaceae</i> | <i>Podocarpaceae living in catchment</i> |
|------------------------------|--|-----------------------------|--|
| SE mainland Australia | | | |
| Lake Bungarby | Late Paleocene | 6 (4) | 0 |
| Anglesea | Middle Eocene | 5 (4) | 0 |
| Golden Grove | Middle Eocene | 2 (2) | 0 |
| Nelly Creek | Middle Eocene | 2 (1) | 0 |
| Nerriga | Middle Eocene | * 1 (1) | 0 |
| Berwick Quarry | Late Oligocene-earliest Early Miocene | 1 (1) | 0 |
| Latrobe Valley | Oligocene-Miocene | 5 (4) | 0 |
| Tasmania | | | |
| Buckland | Early Eocene | 7 (4) | 0 |
| Regatta Point | Early Eocene | 5 (4) | 1 |
| Hasties | mid-Late Eocene | 10 (7) | 1 |
| Loch Aber | mid-Late Eocene | 4 (4) | 1 |
| Cethana | Early Oligocene | 10 (6) | 2(2) |
| Lea River | Early Oligocene | 6 (4) | 2(2) |
| Little Rapid River | Early Oligocene | 19 (9) | 1 |
| Monpeelyata | Late Oligocene-earliest Miocene | 2 (2) | 1 |
| Pioneer | Late Oligocene-earliest Miocene | 7 (5) | 0 |
| Regatta Point | Early Pleistocene | 5 (5) | 1 |

For each location the number of fossil species is given followed by the number of genera in brackets. Also shown is the number of podocarp species growing within these catchments today. Data from Cookson and Pike (1953a,b, 1954), Townrow (1965), Blackburn (1981, 1985), Hill (1982, 1989, 1995), Hill and Macphail (1983, 1985), Christophel *et al.* (1987), Greenwood (1987), Wells and Hill (1989), Carpenter (1991), Hill and Carpenter (1991), Hill and Pole (1992), Pole (1992), Macphail *et al.* (1993), Pole *et al.* (1993), Carpenter *et al.* (1994a), Jordan (1995), Hill and Scriven (1997, 1999), Hill and Christophel (2001).

*Identification as a conifer not confirmed.

significantly bilaterally flattened shoots was in southern Australia during the Palaeogene and it is probable that the podocarp genera with bilateral shoots first evolved under the low solar angle conditions experienced during the Cretaceous/Palaeogene.

Although the data are not precise, there do seem to be patterns of extinction that vary among the podocarp genera present in the Australian macrofossil record. At the extremes, we know that several genera have persisted into the living Australian vegetation, although only a few of these have broad photosynthetic surfaces (e.g. *Phyllocladus*, *Podocarpus*, *Prumnopitys* Phil., *Sundacarpus* (J. Buchholz and N.E. Gray) C.N. Page), while there are broad-leaved genera in the fossil record that are now globally extinct (e.g. *Smithtonia* R.S. Hill and M.S. Pole, *Willungia* R.S. Hill and M.S. Pole) that have no known fossil record past the Early Oligocene. Although the physiology of these extinct genera is unknown, predictions about some aspects may be possible as our understanding of the physiological responses of living podocarps improves and we develop good correlations between morphology and physiological response. Other podocarp taxa with a broad photosynthetic area seem to have become extinct in Australia at different times. For example, *Acropyle* Pilg. is quite common in the macrofossil record up until the Early Oligocene,

but has not yet been recorded after that time, *Phyllocladus* with compound phylloclades (now extinct in Australia) persists in the macrofossil record until at least the Early Miocene and *Dacrycarpus* (Endl.) de Laub. persists until the Early Pleistocene. It is important not to assume that genera that are present today have a long history in the region. For example, the monospecific genus *Lagarostrobos* Quinn only occurs in Tasmania today and has a good Quaternary macrofossil record, but is sparse or possibly absent as a macrofossil before this time. The only formally described fossil in this genus, *L. marginatus* P. Wells and R.S. Hill (Wells and Hill, 1989), was described before *Manoao colensoi* (Hook.) Molloy was separated off from *Lagarostrobos* and this fossil is not particularly close to the single extant species, *L. franklinii* (Hook. f.) Quinn. This may reflect either a minor presence in the vegetation earlier, a different ecological niche, or a relatively late evolution of the genus.

Large contractions and extinctions of Podocarpaceae species have been a feature of the Holocene, although there is no useful macrofossil record of this. Several glacial cycles have no doubt played a significant role in this process. However, there is strong evidence to show that the effects of humans on coniferous species have been particularly severe. The most important influence of humans has been to greatly increase the frequency of forest fire (Figure 19.1), and this has had serious consequences for conifers. Few conifer taxa survive fire and of these none are capable of the type of rapid maturation seen in some angiosperms. Hence, in regions such as south-eastern Australia and Papua New Guinea, there has been a large scale decrease in conifer abundance and even local extinction (e.g. *Phyllocladus* on King Island in the 19th century, probably following European settlement, Jennings (1959)).



Figure 19.1 *Eucalyptus* L'Hérit. dominated forest in south-western Australia with *Podocarpus drouynianus* in the foreground. This is one of the few podocarp species that can regenerate vegetatively following fire.

Adaptation to low light

The Australian fossil record suggests that, although conifers have been a major component of the southern hemisphere flora since the early Cretaceous, it was not until the Palaeogene (Palaeocene to Oligocene), more than 80 million years later, that conifer diversity reached its peak in southern Australia (Hill, 1995; Wells and Hill, 1989; see Table 19.1). Conditions at this time are believed to have been highly equable, with a prevailing wet, temperate climate which apparently enabled the coexistence of a high diversity of conifer and angiosperm taxa (Quilty, 1994). Angiosperms also thrived under these conditions and the shift from conifer dominated forest to largely evergreen conifer/angiosperm mixed forest around this time is likely to have had a profound effect on light availability. This effect would have been most profound in the understorey of such forests where canopy closure by angiosperms most likely decreased light availability (Stenberg, 1996). The bulk of the coniferous taxa recorded from this period are araucarians and podocarpaceous genera bearing broad shoots, such as *Acmopyle*, *Dacrycarpus*, *Phyllocladus*, *Podocarpus*, *Prumnopitys*, *Retrophyllum* C.N. Page and extinct genera *Smithtonia* and *Willungia* (Hill and Pole, 1992).

In general, conifer leaves possess only a single vein and this has been suggested as a major factor limiting conifer success in competition with broad leaved angiosperms in the regeneration niche (Bond, 1989). An impressive number of podocarps appear to have overcome this limitation, having evolved an array of bilaterally flattened photosynthetic structures derived from short shoots, branch tips and leaves. Some 135 species of extant podocarps from 11 genera exhibit flattened or composite photosynthetic structures. We hypothesize that the predominance of conifers with flattened shoots during the Palaeogene demonstrates that these species were growing in forests with relatively dark understoreys and that the flattening of podocarp shoots represents a convergence in conifer foliage towards the efficient light harvesting broad leaf of angiosperms.

Strong support for this hypothesis was provided in a study that illustrated a functional correlation between shoot morphology and light requirements in conifer species (Brodrribb and Hill, 1997a). A study of the living relatives of these broad-shoot bearing conifers, as well as other conifer species in the southern hemisphere, revealed that the flattening of shoots was associated with the light saturation characteristics of leaves (Brodrribb and Hill, 1997a). Photosynthesis in species with highly flattened leaves was found to become light saturated at low light intensities (Figure 19.2) relative to conifers that did not exhibit shoot flattening, indicating their suitability to low light environments. Given these findings, it is surprising that the podocarps that have multiple-veined photosynthetic organs (leaves in *Nageia* and phylloclades in *Phyllocladus*) have not become more successful and widespread in comparison to the single-veined genera. For *Nageia* this may be because it has evolved relatively recently (it has no macrofossil record and a relatively restricted diversity and distribution), but *Phyllocladus* has been extant for at least 50 million years.

Other evidence points towards a change in the availability of light in forests of the Palaeogene, such as the distribution of stomata in leaves of fossil *Acmopyle*, another podocarp bearing broad shoots (Hill and Carpenter, 1991; Hill, 1994). The oldest fossil species (Late Paleocene) has fully formed stomata evenly distributed over both leaf surfaces and no partially formed (hypoplastic) stomata. However, Eocene and Early Oligocene species show varying levels of loss of fully formed stomata from one leaf surface and varying densities and distributions of hypoplastic stomata. *Acmopyle* has not been recorded as a fossil in Australia after the Early Oligocene.

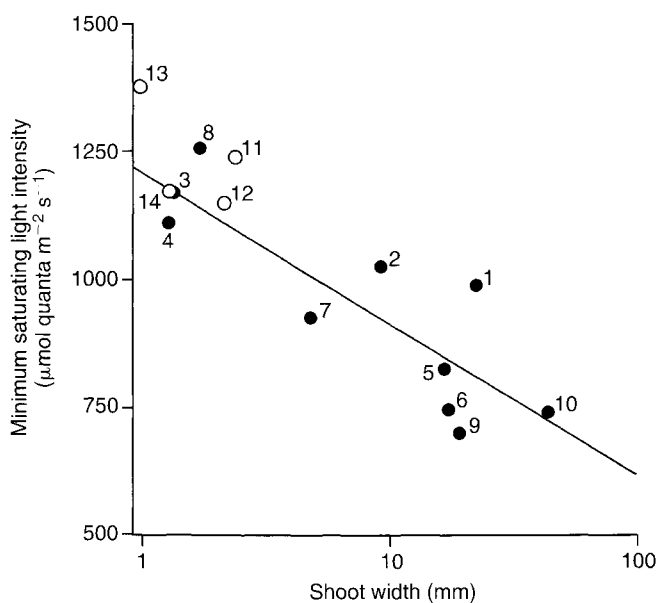


Figure 19.2 Mean light saturation requirements for shoots from 10 species of Podocarpaceae; 1 *Acmopyle pancheri* (Brongn. and Gris) Pilg., 2 *Dacrycarpus dacrydioides* (A. Rich.) de Laub., 3 *Lagarostrobos franklinii*, 4 *Microstrobos niphophilus*, 5 *Phyllocladus aspleniifolius*, 6 *Podocarpus dispermus* C.T. White, 7 *Podocarpus drouynianus* F. Muell., 8 *Podocarpus lawrencei* Hook. f., 9 *Prumnopitys ferruginea* (G. Benn. ex D. Don) de Laub., and 10 *Retrophyllum comptonii* (J. Buchholz) C.N. Page, and four species of Cupressaceae; 11 *Athrotaxis cupressioides* D. Don, 12 *Athrotaxis selaginoides* D. Don, 13 *Callitris rhomboidea* R.Br. ex Rich. and A. Rich., 14 *Diselma archeri* Hook. f. All plants were grown under identical glasshouse conditions.

Hypoplastic stomata may be evidence for stages in either the loss or acquisition of stomata from one leaf surface, but given the relative ages of the species, the former is more probable. The living *Acmopyle* species both have fully formed stomata in two rows along the entire length of the underside of the leaf, but on the upper surface stomata are usually restricted to small areas near the leaf base and apex with hypoplastic stomata in between. However, glasshouse grown specimens are very plastic in their stomatal distribution. In low light, shoots are almost hypostomatic, whereas in high light shoots tend towards amphistomatic (Brodrribb, unpublished data), demonstrating that stomatal distribution on the leaf undersurface is not genetically fixed and it probably was not during the Cenozoic. Therefore, the trends seen in stomatal distribution among the fossils probably represent a plastic response to the environment in which they occurred.

It has been demonstrated that leaf diffusive conductance is well correlated with maximum assimilation (Wong *et al.*, 1985; Körner *et al.*, 1991) and that a linear relationship exists between stomatal density and maximum conductance in extant podocarps (Brodrribb and Hill, 1997b). Thus the decrease in stomatal density in *Acmopyle* is almost certainly a response to decreased maximum assimilation rate in leaves and this most likely came about either through a decrease in temperature, or a decrease in light availability. Evidence supporting a decrease in light intensity comes from the fact that stomatal numbers were found to become differentially reduced on the upper surface of the shoot when compared to the lower surface, thus causing a shift from amphistomy to hypostomy in shoots. Hypostomy in extant *Acmopyle* species is most common in shade plants, while amphistomy

is generally associated with leaves from open habitats (Peat and Fitter, 1994). Thus a shift from amphistomy to hypostomy in *Acmopyle* fossils tends to suggest a reduction of the available light in the *Acmopyle* habitat. A closure of the canopy by invading angiosperms as Australia moved into lower latitudes may have been the cause for such a change in the distribution and availability of light from forest crown to floor. There is also evidence of a decrease in temperature by the time of the last occurrences of *Acmopyle* in Australia (Early Oligocene). Although this temperature change apparently did not affect the shoot size of *Acmopyle*, it is possible that it may have contributed to stomatal loss in the later history of the genus in Australia.

Another fossil Podocarpaceae genus that exhibits a trend in foliar morphology during the Cenozoic is *Dacrycarpus*. Two groups of fossil *Dacrycarpus* species cover a substantial time range and probably represent separate phylogenetic lines. The first is *D. mucronatus* P. Wells and R.S. Hill (Early Eocene to the Early Oligocene in Tasmania), where the oldest specimens have both bilaterally and bifacially flattened foliage and stomata are distributed equally over all leaf surfaces. Bifacially flattened foliage is the normal form of leaf flattening, with a distinct abaxial and adaxial surface, whereas bilaterally flattened foliage in the Podocarpaceae leads to each functional leaf surface containing part of the original adaxial and abaxial surface, with the original leaf margins being approximately in the middle of each flattened surface. By the Early Oligocene *D. mucronatus* is known only as bifacially flattened foliage. The phylogenetically more remote *D. latrobensis* R.S. Hill and R. Carpenter, from Victoria, occurs substantially later in time and still has both foliage types (Cookson and Pike, 1953a). However, stomata are more common on one leaf surface of the bilaterally flattened foliage than the other (Hill and Carpenter, 1991). Therefore, in these species there is evidence for loss of stomata on bilaterally flattened foliage and, in Tasmania at least, loss of the bilaterally flattened foliage type. Recently discovered fossils from Miocene sediments in northern New South Wales further demonstrate that *Dacrycarpus* from lower latitudes did not lose the bilaterally flattened foliage as quickly as those species in Tasmania (Hill and Whang, 2000).

The second phylogenetic line is *Dacrycarpus linifolius* P. Wells and R.S. Hill which occurs in Early Eocene and Early Oligocene sediments in Tasmania and has only been recovered as bifacially flattened foliage, which in gross leaf morphology is identical across the time range. However, while the Early Eocene leaves have stomata along the entire length of both leaf surfaces, the Early Oligocene leaves have stomata restricted to less than the basal third of the abaxial surface, providing further evidence for reduction in stomatal distribution, but this time on bifacially flattened leaves.

While the phylogenetic relationships of the remaining *Dacrycarpus* fossils are not clear, they do exhibit general leaf morphology trends. Six *Dacrycarpus* species have been described from lowland Oligocene–Early Miocene Tasmanian sites, all with imbricate, bifacially flattened foliage. Five are amphistomatic, but the sixth, *D. linearis* P. Wells and R.S. Hill, probably has stomata restricted to the adaxial surface. All bifacially flattened foliage of extant *Dacrycarpus* species is amphistomatic, although in *D. compactus* (Wasscher) de Laub., which occurs at high altitudes in New Guinea and is the only extant species to lack bilaterally flattened foliage, there are extremely few stomata on the abaxial surface. Significantly, the extant podocarpaceous alpine Tasmanian species *Microstrobos niphophilus* and *Microcachrys tetragona* (Hook.) Hook. f. are closely imbricate and epistomatic. A similar *Microstrobos* J. Garden and L.A.S. Johnson species co-occurs with *Dacrycarpus* in two Tasmanian fossil localities.

The last Australian *Dacrycarpus* macrofossil record is from Early Pleistocene Tasmanian sediments (Jordan, 1995). Only bifacially flattened foliage of this fossil has been recorded,

with stomata restricted to the adaxial leaf surface. *Dacrycarpus* must have become extinct in Australia soon after this. The change in leaf morphology in *Dacrycarpus* is more complex than in *Acmopyle*, but some aspects of it are similar. The ability to change foliage form completely (from bilaterally to bifacially flattened leaves) gave *Dacrycarpus* more flexibility and may explain its longer known time range in Australia.

Cretaceous forest structure at high southern latitudes

There has been some speculation as to the structure and function of the high latitude conifer/angiosperm dominated forests that apparently extended across much of Australia and Antarctica during the mid-late Cretaceous (e.g. Specht *et al.*, 1992) and this has considerable bearing on the evolution of today's conifer flora. During the mid-Cretaceous, Australia and Antarctica were joined and most of Australia was south of 60°S. Occurrence of these forests at such high latitudes has led to the hypothesis that they were dominated by stands of pyramidal crown conifers, as this crown shape most efficiently harvests light at low solar angles (Terborgh, 1985; Specht *et al.*, 1992), forming a forest structure similar to that seen at high latitudes in the northern hemisphere boreal zone today. However, apart from the Araucariaceae, few of the southern hemisphere conifers today have the classical conical shape of many northern hemisphere conifers and it is unlikely that this conical shape was common in the southern hemisphere in the past. Many southern conifers have foliage cascading down the sides of the tree in a more random fashion and this is a more likely model for high southern latitude forests in the past. Evidence suggests that many of the angiosperms (and gymnosperms) present in these Cretaceous forests were deciduous (Pole, 1992; McLoughlin *et al.*, 1995; Hill and Scriven, 1995), also conforming to the observed pattern in the boreal forests, where a mixture of deciduous angiosperms and evergreen conifers predominates. Obviously there are difficulties in ascertaining the relative importance of the deciduous element in these fossil floras, as the taxonomic affinities of the species present are often poorly understood and the identification of deciduousness is sometimes based on fairly meagre evidence. However, it is widely accepted that a mixture of deciduous and evergreen angiosperms was present. Extant evergreen broad-leaved taxa with affinities to fossil species have been shown to survive long dark periods such as those that would have prevailed during winter at high latitudes (Read and Francis, 1992). The presence of a significant high latitude evergreen angiosperm flora contrasts with the situation today at high latitudes in the northern hemisphere. Persistence of the evergreen element was probably allowed by the fact that mild winter temperatures (Dorman, 1966) did not cause significant damage to overwintering broad-leaves while not allowing excessive dark respiration.

Constraints on extant Podocarpaceae distributions

It appears that podocarps, by virtue of their relatively large foliar morphological range, are well adapted to survive under most light conditions. Changing light conditions therefore do not appear to account for the restriction and extinction of conifers during the Cenozoic in the southern hemisphere. To address better the question of why podocarps, and indeed all conifer families, suffered during the later Cenozoic it is instructive to examine what constrains the distribution of extant southern conifer species.

The fundamental requirement for genetic survival in conifers, and indeed all plants, is the production of enough photosynthate to allow growth and reproduction. Thus, limitations

imposed by the physical environment on the distribution of plant species will usually manifest themselves through their effects on leaf photosynthesis. The environmental parameters most commonly used to explain presence or absence of species are temperature and rainfall regimes and their inferred effects are almost invariably on the photosynthetic capacity of the species in question. Because of their characteristic longevity (Loehle, 1989), sporadic reproduction, slow rates of dispersal and short-term viability of soil-borne seed, (Gibson *et al.*, 1994; Veblen *et al.*, 1995), conifers are particularly closely linked to long-term environmental characteristics. Unlike the fast growing, highly fecund angiosperms, most conifers do not have the capacity to move in and out of marginal environments in response to short-term physical fluctuations and are thus confined at the physiological extremes of their distributions by the long-term survival of adult plants.

Despite the anecdotal and quantitative evidence of environmental constraints on species distributions, few studies have demonstrated the actual mechanisms by which physical limitations affect the survival and reproduction of plant species. Over the last 50 years, research into the characteristics and control of leaf gas exchange has been prolific and the use of leaf photosynthetic characters to explain simple successional changes in the vegetation has become common in the literature, typically using water-use efficiency (e.g. Meinzer *et al.*, 1984; Dias-Filho and Dawson, 1995), drought response (e.g. Kubiske and Abrams, 1993) and light response (e.g. Field, 1988; Knapp and Smith, 1991; Ashton and Berlyn, 1994) to account for small-scale competitive outcomes. This type of ecophysiological research generally takes the form of comparisons between small numbers of species and, as such, does not attempt to make a quantitative link between photosynthetic function and environmental parameters. Indeed, with a few exceptions (e.g. Teeri and Stowe, 1976), large-scale relationships between physiology, environment and species distributions have been overlooked by ecophysicologists.

In order to understand how the distribution of species is likely to have been affected by historical climate change and might be affected by future climatic perturbations, it is important to be able to describe current, known distributions in terms of the most likely physiological constraint. This was undertaken for a group of 13 conifer species, including eight podocarps, in a study by Brodribb and Hill (1998). We took species from a diverse geographical range spanning Australia (Figures 19.3 and 19.4), New Zealand, New Guinea, South Africa and New Caledonia (Figure 19.5) and a broad spectrum of habitats and compared the water-use efficiency and drought tolerance of photosynthesis with environmental water availability. Water availability was considered the most likely constraint on distribution given that podocarp forest in the southern hemisphere is almost exclusively restricted to areas of high rainfall, with the most extensive and diverse conifer forests occurring on the west coasts of New Zealand and Tasmania (Figure 19.4), in New Caledonia (Figure 19.5) and highland New Guinea, where rainfall is commonly over 5000 mm per annum (van Royen, 1979).

We measured two physiological parameters related to the drought tolerance of leaves and the xylem vascular system. The first of these was maximum water-use efficiency during drought (Brodribb and Hill, 1996) and the second was the vulnerability of xylem to cavitation by water stress (Brodribb and Hill, 1999). These parameters were compared with minimum rainfall data over the ranges of the selected species. A good correlation between maximum water-use efficiency (see Brodribb, 1996 for details) and rainfall parameters (Figure 19.6) suggests that the climatic ranges of these conifers are closely controlled by their photosynthetic and gas-exchange characteristics during drought. This is an important result as it provides a direct link between instantaneous gas-exchange characteristics



Figure 19.3 Cool temperate rainforest in western Tasmania containing mature trees of *Phyllocladus aspleniifolius*. These trees have been recently exposed on a road cutting.

and species distribution. The cost to the plant of maximizing water-use efficiency, as measured by these two parameters, may come as an increase in the saturation light requirement (discussed below) and respiration rate of leaves. Increases in leaf nitrogen associated with higher carboxylation efficiency and photosynthetic capacity in species have commonly been linked to increases in the dark respiration rate and this has been explained by suggestions of higher tissue maintenance costs, or costs associated with phloem loading and carbohydrate export from the leaf (Connor *et al.*, 1993; Bouma *et al.*, 1995; Baxter *et al.*, 1995). Thus, there must be a significant selective pressure on these species to maintain their physiological drought tolerance to a level dictated by the long-term water availability in the environment and an insufficient drought response must become limiting by causing local extinction of a species, either by death during periods of extreme conditions, or by competitive exclusion by more efficient species. Either way, the importance of rainfall is emphasized in constraining the distribution of these species.

Strong correlation also exists between xylem vulnerability to cavitation and average rainfall during the dry season in the Podocarpaceae and southern hemisphere Cupressaceae

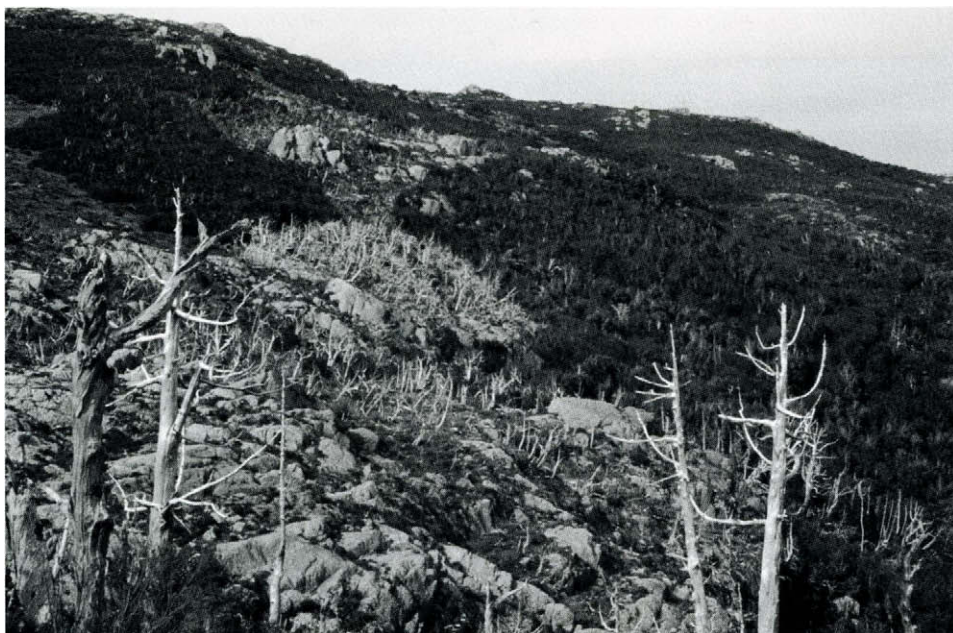


Figure 19.4 High altitude vegetation on Mt Read in western Tasmania. Six conifer species coexist in this area, with four belonging to the Podocarpaceae. The very damaging impact of fire can be seen in the foreground.



Figure 19.5 Conifer dominated forest in New Caledonia, with *Dacrycarpus* and Araucariaceae among the prominent trees.

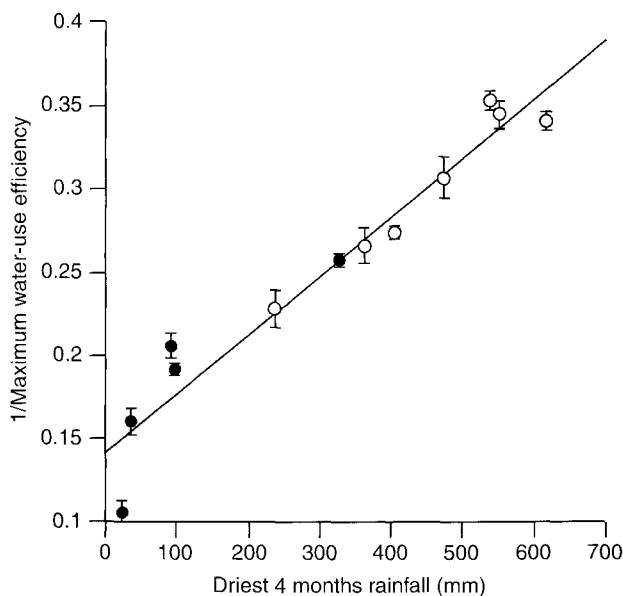


Figure 19.6 Shows highly significant correlation ($r = 0.97$) between mean rainfall during the driest four months within the distributions of 8 Podocarpaceae (○) and 5 Cupressaceae (●), and leaf drought tolerance. The drought tolerance parameter is the lowest ratio of mesophyll to ambient CO_2 concentration measured in leaves during imposed drought. This ratio is inversely proportional to the maximum water-use efficiency attainable by leaves.

(Brodribb and Hill, 1999). Species from wet environments were highly vulnerable to cavitation while species from the semi-arid zone produced stem xylem that was extremely resistant to pressure-induced cavitation. Clearly this indicates an important role for xylem vulnerability in determining the distributional limits of these plants in terms of minimum water availability. This also provides evidence for linkage between leaf drought tolerance and stem cavitation characteristics, as shown by a highly significant regression relating maximum water-use efficiency with the water potential (Ψ_{50}) that caused 50% cavitation in stem xylem (Figure 19.7). One possible inference from this is that a loss of hydraulic conductance in the xylem during water shortage is the causal factor dictating a loss of leaf function during drought in these species. Such a hypothesis is supported by data suggesting that plants may operate close to the point of 'runaway cavitation', where a positive feed-back following xylem embolism has the potential to cause the vascular system rapidly to lose hydraulic conductivity unless transpiration is reduced (Sperry *et al.*, 1993; Alder *et al.*, 1996). However, several pieces of evidence point away from xylem dysfunction as the primary cause of leaf failure during drought, particularly for species from drier habitats. This evidence comes from research illustrating that complete stomatal closure and a loss of optimal quantum yield (indicating damage to photosystem II) in these species both occurred at leaf water potentials above the value corresponding to Ψ_{50} (Brodribb and Hill, 1999). Given this large 'safety margin' between stem water potential during active photosynthesis and that which would induce significant (or possibly runaway) cavitation, it seems unlikely that the xylem water potential would approach Ψ_{50} unless plants were subject to severe water shortage. Considering that none of the species investigated are likely to suffer significant embolism by freeze-thaw cycles (Sperry *et al.*, 1994), large-scale stem xylem cavitation probably only occurs when plants experience soil moisture conditions approaching Ψ_{50} .

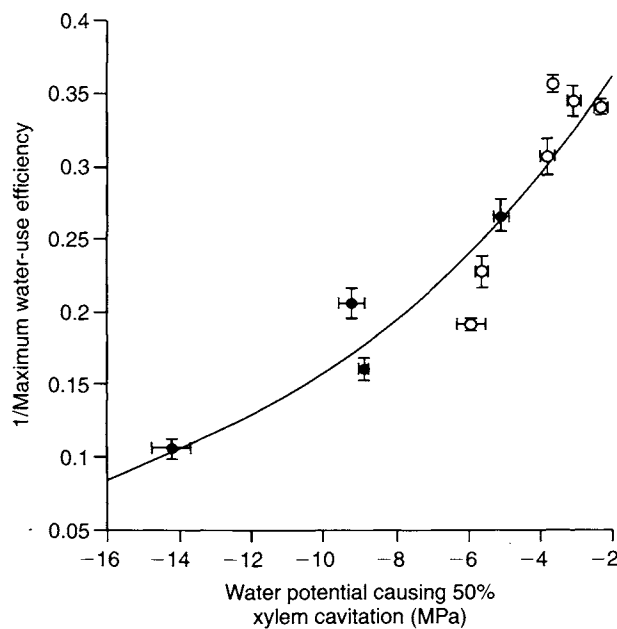


Figure 19.7 The relationship between average ($n = 5$) water potential which reduced stem hydraulic conductance to 50% of the measured maximum (Ψ_{50}), and the average ($n = 3$) leaf drought tolerance index (Figure 19.6) (Brodrribb, 1996) in each of 6 species of Podocarpaceae (O) and 4 Cupressaceae (●). A highly significant exponential regression ($y = 0.71x^{-0.66}$; $r^2 = 0.89$, $P < 0.001$) is shown.

The conclusion here, that conifer distributions are commonly water limited, has significant implications for interpretation of the conifer history in the southern hemisphere. Large-scale changes in the rainfall and other climatic characteristics are proposed to have occurred throughout the Cenozoic and these changes must have significantly influenced conifer survival. This is reflected in the species numbers shown in Table 19.1 and in the staged extinction of Podocarpaceae genera with broad photosynthetic surfaces discussed earlier. The Australian environment changed from one that could support widespread rainforest in the Palaeogene to the current arid-dominated landscape over a long time period, but it was not a smooth transition. As the fossil record improves we will better be able to test the correlation of extinctions in the Podocarpaceae against periods of intense aridification.

Synthesis

The most common conifer shoots from Australian Cretaceous deposits are narrow, imbricate foliage from genera within the Araucariaceae, Cupressaceae (including Taxodiaceae) and Podocarpaceae. The first appearance of podocarps with significantly bilaterally flattened shoots is in southern Australia during the early Cenozoic (probably *Acropyle* in the Palaeocene; Hill and Carpenter, 1991) and it is probable that the podocarp genera with bilateral shoots first evolved under the low solar angle conditions experienced during the Cretaceous/Palaeogene. Shoot arrangement in many conifers from high latitudes in the northern hemisphere today shows a progression from cylindrical in the canopy, to bilaterally flattened in the understorey (Sorrenson-Cothorn *et al.*, 1993) and it is conceivable that a similar pressure to allow penetration of light deep into the canopy (Stenberg, 1996) led

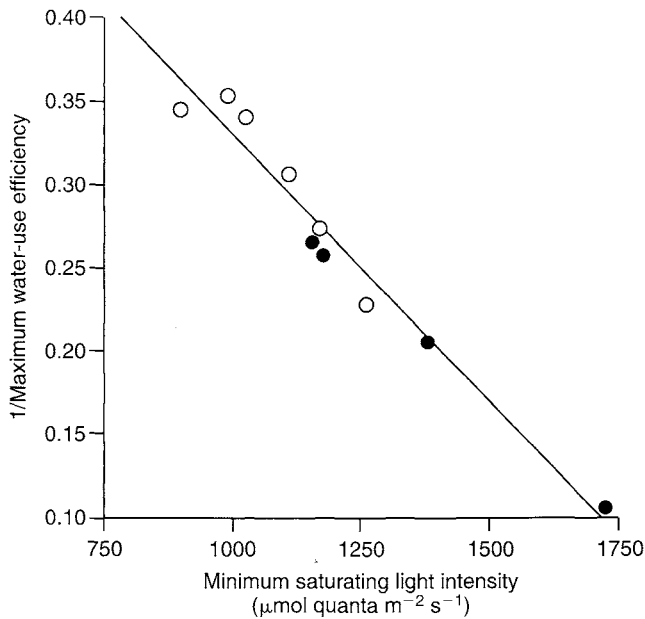


Figure 19.8 A strong negative correlation exists between saturation light intensity and $(\text{water-use efficiency})^{-1}$ in 6 species of Podocarpaceae (○) and 4 southern hemisphere Cupressaceae (●) indicating a probable 'trade-off' between these two physiological properties of leaves.

to the evolution of morphological plasticity in podocarp shoots. Of course, as Australia drifted north and the solar angle increased, allowing the winter period to become relatively light, the advantages of producing a deep crown and deciduousness were reduced. Higher incident radiation year round would have allowed broad-leaved, evergreen angiosperms to flourish in the canopy. Podocarps would have been 'exapted' (Gould and Vrba, 1981) to this change and the expansion of genera with broad shoots at this time would have taken advantage of the change in light conditions. The morphological plasticity of these broad shoots allowed efficient light harvesting in the understorey as well as the ability to adopt a three-dimensional helical leaf arrangement in the canopy that is suited to harvesting high light intensities.

The foliar convergence between broad-shoot podocarps and angiosperm broad-leaves appears to have enabled the persistence, and possibly the radiation, of conifer taxa bearing this shoot morphology as the Cenozoic forest structure in southern Australia apparently shifted from high latitude conifer/deciduous to evergreen broad-leaf forest. A significant liability does, however, seem to be linked to this adaptation, that being a low tolerance of water stress. Virtually all extant podocarps exhibit quite low drought tolerance and the broad-shoot genera, such as *Dacrycarpus* and *Acmopyle*, are particularly intolerant of drought, a feature evident in the physiology of both leaf and xylem tissues (see Figures 19.6 and 19.7) of these species. Interestingly, there appears to be an inverse relationship between shade tolerance and drought tolerance of leaves of southern hemisphere conifers when data from 10 species are compared (Figure 19.8). A 'trade-off' between drought and shade tolerance has been frequently hypothesized but rarely shown (Smith and Huston, 1989; Holmgren *et al.*, 1997) and the significant inverse correlation between drought and shade (Figure 19.8) provides strong support for a drought/shade tolerance 'trade-off' in southern hemisphere conifers. The outcome of this is that while flattened broad-shoots

may have allowed podocarps to retain codominance in high rainfall, mixed Cenozoic forests, these taxa were left extremely vulnerable to the progressive decrease in water availability that followed.

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20

The adaptive physiology of *Metasequoia* to Eocene high-latitude environments

Richard Jagels and Michael E Day

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Introduction

Both conceptual (Wolfe, 1979, 1985) and predictive (Iverson and Prasad, 1998) models that relate the distribution of plant species to palaeoclimates have focused primarily on temperature and hydrological parameters, with the assumption that these are the principal forces driving latitudinal changes in distribution. While the predominance of these two factors in determining the ranges of species under current global climate regimes is

well established (Woodward, 1987), in the climatic regime of the palaeo-Arctic, the light environment may have exerted a major control over plant distribution. Such conditions, which have no contemporary analogue, would occur during global temperature maxima, when plant growth at high latitudes would not be restricted by low temperatures. Plant populations migrating toward polar regions, would encounter light environments characterized not only by reduced maximum irradiance but unique temporal patterns on both diel and seasonal scales. Optimizing growth in warm, high-latitude irradiance regimes would require specialized approaches to carbon balance physiology that would be both qualitatively and quantitatively distinct from those optimal at low and middle latitudes.

Because assimilation of CO₂ necessitates water loss through stomata, success in a particular environment requires an appropriate balance between carbon gain and hydraulic stresses. Many modern plants that are adapted to both high-temperature and high-light environments employ crassulacean acid metabolism (CAM) or C₄ photosynthetic pathways to maximize water-use efficiency. However, CAM requires a diurnal dark period to operate effectively and, to our knowledge, the C₄ pathway has not been demonstrated in any conifer, therefore these adaptations would have not been available to conifers in high-latitude Eocene environments. The greater distances between roots and photosynthetic organs in trees, increases the potential for hydraulic limitations to photosynthesis in that lifeform. In addition, success in the continuous light of the high-Arctic summer requires effective mechanisms for protection from detrimental effects of excess light energy. When solar energy is captured by photosynthetic pigment complexes it must be dissipated by photosynthetic or non-photochemical systems to prevent damage to the photosynthetic apparatus. In a continuous light environment, these systems must function without restriction (i.e. inhibition by accumulation of a product at any stage) and without a non-photosynthetic period for maintenance of structural and enzymatic components offered by diurnal day-night cycles. Although various photoprotective mechanisms have been studied in herbaceous species, their function in tree canopies is poorly understood.

The high-Arctic, swamp palaeoforests of the Eocene were often dominated by the gymnosperm *Metasequoia* Miki, while the extant natural range of this genus, consisting of a single species *M. glyptostroboides* Hu et Cheng (Figure 20.1), is restricted to deep valleys in the Hupeh province of China. This contrast poses some interesting and important questions. What physiological and morphological attributes did *Metasequoia* possess that enabled it to dominate the unique environment offered by the Eocene high-latitude sites, in the face of competitors such as *Larix* P. Mill, a genus that now dominates many northern boreal forest ecosystems? Can the morphology and physiology of extant *M. glyptostroboides* provide clues to ecophysiological attributes that confer competitive advantages to temperate, continuous-light habitats? In this chapter we explore the physiological basis for *Metasequoia*'s success in the Eocene high-Arctic and, in so doing, hope to shed light on ecophysiological attributes which imparted adaptive value to trees inhabiting that unique environment. In our attempt at answering these questions, we have adopted a comparative ecophysiological approach contrasting attributes of *Metasequoia* with *Larix* and other conifers that now occupy boreal forests.

Background

The Eocene lowland forests of Axel Heiberg Island

At least 28 layers of buried fossil forests from the middle Eocene, about 45 million years ago (McIntyre, 1991; Ricketts and McIntyre, 1986), have been found on the Canadian



Figure 20.1 *Metasequoia glyptostroboides* growing at approximately 44.8°N latitude in Hancock County, Maine, USA.

Arctic Island of Axel Heiberg (79°55'N, 88°58'W), where wood, cones, fruits and leaves have been mummified and possess extraordinary preservation (Basinger, 1986; LePage and Basinger, 1991; Young, 1991) (Figure 20.2). Current evidence indicates that these were wet-site forests, dominated by the gymnosperm *Metasequoia* (Estes and Hutchison, 1980; Basinger, 1991; Francis, 1991; Irving and Wynne, 1991; Greenwood and Basinger, 1994). Other fossil *Metasequoia* sites have been found throughout the high Arctic (Christie and McMillan, 1991; Monohara, 1994).

At an estimated palaeolatitude of $78 \pm 5^\circ\text{N}$ (Irving and Wynne, 1991), the Axel Heiberg palaeoforests grew under light/temperature combinations not currently found on the planet (Figure 20.3). During the months of May, June, July and parts of April and August, the forests received continuous, albeit low to moderate intensity, low-angle sunlight combined



Figure 20.2 *Metasequoia* trunk excavated at the Axel Heiberg palaeosite.

with warm temperatures. However, during the months of November, December, January and February the forests were exposed to either no light or light levels below the compensation point of photosynthesis (Berner, 1991; Pielou, 1994; Greenwood and Wing, 1995). Despite the restricted growing season, productivity of the polar *Metasequoia* forests approximated that of contemporary lower latitude *Metasequoia glyptostroboides* stands (see Chapter 21).

The palaeoforests of Axel Heiberg have been characterized as swamp-type wetlands between river systems (Francis, 1991; Tarnocai and Smith, 1991). In forested horizons, autochthonous leaf-litter mats represent the ancient forest floors of poorly-drained floodplains and associated swamps (Ricketts, 1986, 1991; Basinger, 1991). Megafloral remains in forest-floor mats include the fertile and vegetative remains of the dominant *Metasequoia* and, to a lesser extent, *Glyptostrobus* (Endlicher), with minor occurrences of *Larix*, *Picea* A. Dietr., *Pseudolarix* Gordon, *Pinus* L., *Betula* L., *Alnus* P. Mill., *Juglans* L., *Chamaecyparis* Spach., *Tsuga* Carr., *Osmunda* L. and several unidentified angiosperm taxa (Ricketts and McIntyre, 1986; Basinger, 1991; LePage and Basinger, 1991; McIntyre, 1991).

Among the tree stems and stumps which we have identified to genus (approximately 30), most are *Metasequoia*, a few are *Larix* (Jagels *et al.*, 2001) and one is a dicotyledon (possibly a member of the Lauraceae; unpublished data, R. Jagels). This suggests that *Larix* grew as a minor component in a *Metasequoia*-dominated lowland wet-site forest. As no large woody fossils of the other conifers have been found at the Axel Heiberg site, it is possible that their cones and small branches were transported from higher elevations and deposited at the site. Preliminary analysis of the tree trunks and stumps (see Chapter 21) provide estimates of forest structure and biomass. The diameters of the preserved tree trunks indicate that the *Metasequoia*-dominated swamp and floodplain forests were 33–39 m tall (see Figure 20.2).

The only fossil *Larix* species found at Tertiary high-latitude sites are those with short-bracted cones (LePage and Basinger, 1991). Hu (1980) lists *Larix potaninii* Batalin as a member of the *Metasequoia* flora in China. However, according to Wang and Zhong (1995) *L. potaninii* is not found at altitudes below 2300 m and is only common between

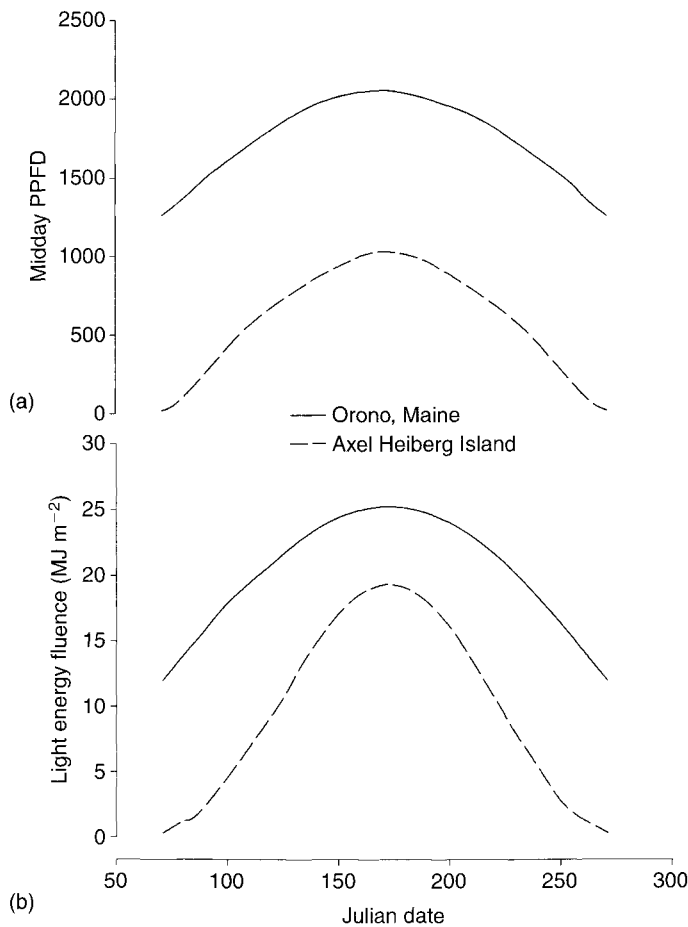


Figure 20.3 Modelled seasonal patterns of (a) midday photosynthetic photon flux density (PPFD, $\mu\text{mol m}^{-2} \text{s}^{-1}$) and (b) light energy fluence for 44.8°N latitude (Orono, Maine) and 78°N latitude (Axel Heiberg Is.). Model is based on equations in Gates (1980), with the assumptions of 70% atmospheric transmission, 50% of incident light energy in PAR (400–700 nm) wavelengths and an average of $4.6 \mu\text{mol photons J}^{-1} \text{PAR}$ (Jones, 1992).

2600 and 4000 m elevation, where it is sympatric with *Picea* and *Abies* P. Mill. species. In contrast, natural populations of *Metasequoia* occur between 1000 and 1100 m, elevations that are mostly devoid of *Picea* and *Abies* (Hu, 1980). In any case, *L. potaninii* is a long-bracted species, a characteristic typical of montane *Larix* species. For our comparative studies, we therefore have chosen *Larix laricina* (DuRoi) K. Koch as representative of the circumboreal short-bracted *Larix* species, to compare with *M. glyptostroboides*. However, it should be noted that *Larix* is a widely distributed genus with 10 species and numerous varieties, ecotypes and hybrids (Schmidt and McDonald, 1995). Therefore, the likelihood that any modern *Larix* species is representative of the fossil *Larix* is presumptuous. Nevertheless, since the modern species are all deciduous and have numerous morphological and physiological attributes in common, a comparison with *Metasequoia* can provide some insight into their relative adaptability and competitiveness in the Axel Heiberg palaeoenvironment.

Characteristics of the temperature regime at the 45 million year old palaeosite are very speculative at present. Rates of primary production in the palaeoforest suggest a temperate climate (see Chapter 21). Bassinger (1991) suggested that winter temperatures seldom dropped below freezing, based on identification of vegetative remains and the lack of significant latewood production in *Metasequoia*. However, *Larix* from the same site shows significant latewood production (Jagels *et al.*, 2001). McIntyre (1991) concluded that the climate was warm-temperate, based on the pollen assemblage.

Metasequoia: fossil and living

Extant natural stands of *Metasequoia* (*M. glyptostroboides*) are restricted to a small, remote area of Hupeh province near the border of Szechuan, China at an approximate latitude of 30°10'N (Chu and Cooper, 1950). Although in the fossil high-latitude forests *Metasequoia* was the upper-canopy dominant species and exhibited growth rates more typical of shade-intolerant conifers, characteristics of both the natural forests in China (Chu and Cooper, 1950; Florin, 1952; Li, 1957) and planted stands of extant *Metasequoia* (personal observations) suggest that the species maximizes growth in partially shaded rather than full sun conditions. This paradox will be explored further in later sections of this chapter.

Several authors have proposed that the deciduous habit of *Metasequoia* provided a competitive advantage for the Arctic winter (Francis, 1991; Basinger, 1991; McIntyre, 1991; see Chapter 21). Chaney (1948) postulated that the deciduous forest type originated in the north during the Cretaceous and migrated south during the later Tertiary and Vann *et al.* (see Chapter 21) suggest the possibility that *Metasequoia* evolved at high-latitudes. Hu (1980) disputed Chaney's hypothesis on the basis that a deciduous habit can also be adaptive to wet/dry periodicity in relatively frost-free areas, such as the Hupeh Forest. Fossil evidence demonstrates that *Metasequoia* was well established at lower latitudes during the Cretaceous (Figure 20.4), predating the Eocene Arctic swamp forests. This is consistent with Hu's argument for the evolution of the deciduous habit under a wet-dry climate regime, particularly since this is a rare attribute in conifers. Therefore, in high-latitude Eocene forests deciduousness may have been adaptive not only by reducing metabolic carbon demand during winter dark periods, but by reducing water stress during the dry phase of soil moisture cycles. In a contemporary analogy, *Taxodium distichum* (L.)

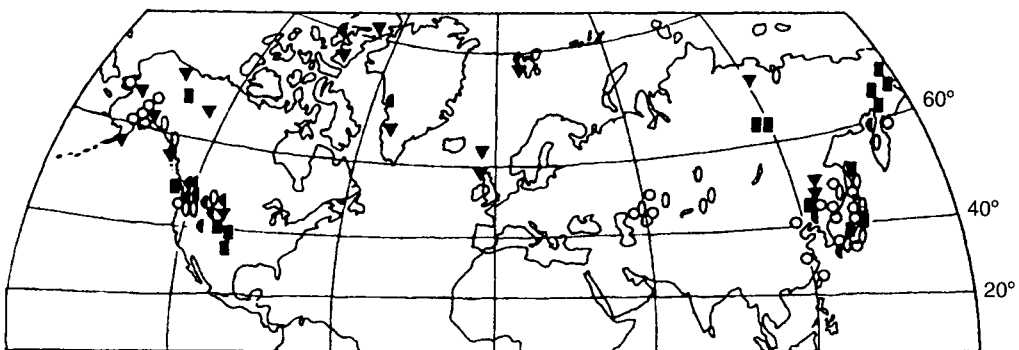


Figure 20.4 Distribution of *Metasequoia* fossil palaeosites. Key to sites: ■ = Late Cretaceous, ▼ = Palaeocene, ▲ = Eocene, ○ = Oligocene, ○ = Miocene. Map adapted from Monohara (1994).

Richard is deciduous and is restricted to floodplains in the southern USA where wet-dry periods dominate the soil moisture regime.

Metasequoia was well distributed above the Arctic circle from the Palaeocene through the Eocene. As the earth cooled and became drier during the Oligocene and Miocene (Wolfe, 1985) its distribution became more limited to temperate zones of North America and Asia (see Figure 20.4) and eventually became restricted to moist, narrow river valleys in China (Chu and Cooper, 1950; Li, 1957; Momohara, 1994). In contrast, short-bracted *Larix*, an equally poor competitor, but one having a greater low-temperature tolerance (either a preadaptation or one evolving over time) was able to successfully occupy north temperate and boreal hydric sites where competition is minimal.

Vann *et al.* (see Chapter 21) develop a compelling case that *Metasequoia glyptostroboides* retains a substantial array of morphological and, by implication, physiological traits present in the Eocene *Metasequoia*, and is, therefore, a useful nearest living relative (NLR) for comparative studies. As discussed in subsequent sections, extant *M. glyptostroboides* appears to have 'relict' traits; and unusual combinations of morphological and physiological characteristics would have particular adaptive value in the unique environment suggested for the Eocene high-Arctic forests. In modern temperate forests these same traits would have little adaptive value and may even be maladaptive. This collection of morphological and physiological traits is consistent with the suggestion of Konoe (1960) that establishment of this species may depend on successful responses to multiple environmental factors. While also consistent with a high latitude evolution for *Metasequoia*, it does not rule out the possibility that these adaptations evolved at lower latitudes and provided *Metasequoia* with a preadaptive advantage in Eocene Arctic swamp-forests.

Physiological challenges of a high-latitude, continuous-light environment

Our research has focused on physiological and morphological traits which would be adaptive to a tree species growing in a temperate climate with a season of continuous light (CL) alternating with a season of continuous darkness (CD). Light intensities during the summer were assumed to be substantially less than at lower latitudes (see Figure 20.3). Categories of attributes considered adaptive to the Eocene palaeoarctic forest environment were: (1) the ability to photosynthesize efficiently under the light intensities and temporal patterns of high latitudes; (2) possession of an integrated carbon balance strategy (photosynthesis, respiration and allocation) that would be adaptive to growth and reproduction in that unique environment; (3) an expeditious system for translocation of photosynthetic products to prevent direct or indirect feedback inhibition of photosynthesis; (4) processes for dissipation of excess light energy through non-photochemical pathways or photorespiration in order to protect photosynthetic systems; and (5) strategies for minimizing limitations to photosynthesis from water stress imposed by continuous transpiration in CL.

Adaptations to Arctic light regimes

During the CD period of winter, the tree must be able to effectively minimize metabolic activity, which would provide strong selective pressure for a deciduous habit. Because the polar early spring provides only minimal light intensity, the tree should be able to produce

new foliage and initiate chlorophyll synthesis at very low light levels. Foliar characteristics along the sun-foliage to shade-foliage continuum are determined by the light environment under which the leaves develop and are adaptive in that carbon and nutrient resources are allocated in proportion to potential carbon gain (Givnish, 1988; Küppers, 1994). However, at high latitudes spring light intensity is far lower than the summer intensity (see Figure 20.3), decreasing the efficiency of this mechanism. This great variation in light intensities over the growing season would impart adaptive advantage to photosynthetic physiology that provided for efficient carbon fixation (relative to the allocation costs) over a wide range of light intensities. While photosynthetic traits associated with shade-adapted foliage would be optimal during the Arctic spring and autumn, shade-adaptation has been shown to be negatively correlated with attributes necessary to address the physiological stresses of higher irradiance environments (Givnish, 1988; Johnson *et al.*, 1993; Tyree *et al.*, 1998).

Under selective pressure to optimize carbon gain in the Arctic light regime, this incompatibility between shade- and sun-adaptation could be addressed through one or more of the following mechanisms: (1) by modifying the photosynthetic physiology of existing foliage; (2) by production of new foliage as the light environment changes; or (3) by producing foliage that exhibits relatively efficient photosynthesis over a wide range of light intensities (i.e. an 'optimal compromise' of sun- and shade-adaptations). Adaptive adjustment of photosynthetic attributes of existing foliage to increasing light intensities has been demonstrated in conifers (Brooks *et al.*, 1996), however, this process has not been examined in *Metasequoia* or closely related species. Trees with an indeterminate growth habit, such as *Metasequoia*, continuously produce foliage through the growing season (Table 20.1). This provides a mechanism for adjusting the light-adaptation status of foliage to changing light environments (Pothier and Margolis, 1991). Furthermore, as discussed in the next section, the photosynthetic light-response curve of *Metasequoia* suggests that both shade-adapted and sun-adapted foliage of that species is photosynthetically efficient over a wide range of low to moderate light intensities, roughly coincident with those found in the high Arctic.

Carbon balance physiology

Efficient carbon balance physiology requires synchronization of photosynthetic, carbohydrate sink and storage activity with diel and seasonal cycles of resource availability (Luxmoore, 1991). A deciduous habit is obviously adaptive to extended periods when photosynthesis is resource (light or moisture) limited. In annual cycles at middle latitudes aspects of carbon balance are controlled by phenological processes (Cannell, 1990), which are induced by environmental cues such as day length and temperature. Adaptation to a temperate high-latitude environment would require modification of these processes in a manner that has no extant analogue.

During the midsummer period of maximum light intensity, down-regulation of photosynthesis by end-product buildup may become a challenge to plants growing in a CL environment. End-product down-regulation may decrease carboxylation rates through several potential pathways that may act directly on biochemical mechanisms or indirectly by genetic induction (Stitt, 1991; Krapp and Stitt, 1995) and have a two-fold deleterious influence by concurrently limiting potential carbon gain and restricting the carboxylation pathway for dissipation of intercepted light energy. Pathways leading to end-product inhibition of photosynthesis may be specific to metabolites accumulated (Schaffer *et al.*, 1986; Stitt, 1991; Huber and Huber, 1992; Krapp and Stitt, 1995; Jang *et al.*, 1997). Therefore, physiological strategies to limit feedback inhibition of photosynthesis under CL conditions must

Table 20.1 Comparison of attributes of crown architecture and foliar morphology among the gymnosperms *Metasequoia glyptostroboides*, *Larix laricina* and *Picea rubens*. SLA data for *P. rubens* from Day *et al.* (2001) and for *Larix laricina* from Hutchison *et al.* (1990).

| Attribute | <i>Metasequoia glyptostroboides</i> | <i>Larix laricina</i> | <i>Picea rubens</i> |
|--|---|---|---|
| Shade-tolerance | Mixed (see text) | Intolerant | Tolerant |
| Regeneration | Seedling rapidly captures growing space | Seedling rapidly captures growing space | Advance regeneration captures canopy gap |
| Juvenile growth | Rapid | Rapid | Slow |
| Leaf area index (m ² m ⁻²) | 5.0 | 1.4–1.8 | 4.0–5.0 |
| Mutual shading by foliage on shoot | Low | Low | High |
| Leaf orientation | Horizontal | Multidirectional | Multidirectional |
| Shoot development | Indeterminate throughout growing season, recurrently flushing | Limited indeterminate | Determinate (second flushes rare in natural environments) |
| Foliar longevity | 1 year | 1 year | 7–15 years |
| Specific leaf area of sun-foliage (cm ² g ⁻¹) | 110 | 100 | 40 |
| Specific leaf area of shade-foliage (cm ² g ⁻¹) | 315 | – | 46 |
| Plasticity of foliage morphology with respect to light environment | Very high | Low | Intermediate |
| Surface-to-volume ratio | High | High | Low |
| Epidermal walls | Adaxial moderately thick, abaxial thin | Uniformly thick | Uniformly thick |
| Hypodermis | Absent | Present | Present |
| Chloroplasts in epidermal cells | Present on abaxial surface | Absent | Absent |
| Stomata: Arrangement | Hypostomatic | Amphistomatic | Amphistomatic |
| Position | Surface | Sunken | Sunken |
| Density | Relatively low | Relatively high | Relatively high |

address not only rate of export of photosynthetic products, but also form of end products (sugars, starch).

Site dominance

Dominant tree species in forested ecosystems are characterized by a suite of attributes that enable capture and retention of available growing space, while limiting the ability of potential competitors to infringe on that growing space (Oliver and Larson, 1990). Such attributes include rapid shoot growth, production of high leaf area index (LAI) and longevity (Küppers, 1994). When growing on nutrient-poor soils, such as the spodosols suggested for the Axel Heiberg palaeoforest (Tarnocai and Smith, 1991), a competitive advantage would be gained by species that could develop and maintain a high LAI with minimal investment of carbon and nutrient resources for supporting woody structures and species with a high nutrient-use efficiency for production and maintenance of photosynthetic organs.

Photoprotection

Photoprotective mechanisms prevent damage to components of the photosynthetic system from excessive energy input and are presumed to have significant adaptive value to species currently inhabiting high-latitude or alpine environments where incident solar radiation is substantially in excess of photosynthetic capacities (Lloyd and Woolhouse, 1979; Mawson *et al.*, 1986; Streb *et al.*, 1998; Manuel *et al.*, 1999). However, a direct comparison is complicated by the lower temperature regimes in those regions under current-climate conditions.

Challenges associated with protection from photoinhibitory damage include dissipation of excess incident light energy through non-photochemical (thermal) quenching pathways or photorespiration to prevent damage to photosynthetic pigment-protein complexes (Johnson *et al.*, 1993; Demmig-Adams and Adams, 1996; Park *et al.*, 1996; Streb *et al.*, 1998); maintaining protective transthylakoid proton gradients to control cross-membrane pH gradients in the grana of chloroplasts (Briantais *et al.*, 1979; Manuel *et al.*, 1999); preventing the buildup of excess ATP and reducing capacity (Kozaki and Takeba, 1996); and accumulation of antioxidants such as glutathione, which is a byproduct of photorespiratory activity (Noctor *et al.*, 1999).

Water balance

The challenges associated with water relations for trees in a temperate CL habitat are poorly understood, but may be qualitatively different from those faced by trees under normal day-night cycles. In a CL regime, optimum growth requires continuous gas exchange without excessive risk of catastrophic xylem embolism (see Sperry *et al.* (1993) for a review of water relations challenges associated with high energy environments).

Extant conifers in temperate and boreal biomes generally use stem capacitance as an integral part of water relations strategies (Pallardy *et al.*, 1995). High transpirational demand during the day is supported, in part, by water stored in stem sapwood, with losses from storage sapwood replenished during the night when transpirational demand is low. This may permit the maintenance of a larger leaf area with a root system that would not be capable of directly meeting transpirational demands. However, in the continuous light of a high-latitude warm summer the value of a water-relations strategy employing capacitance and overnight recharging is lost, placing a greater adaptive value on higher water-use efficiency (WUE; unit CO₂ fixed per unit H₂O transpired).

The Eocene *Metasequoia*-dominated forests have been described as swamp-forest types, which would imply minimal restriction on productivity by soil moisture limitations. However, in extant temperate swamp-forest analogues, productivity is subject to both seasonal (predictable) and stochastic (unpredictable) limitations to water availability and/or uptake. These processes can be restricted by drought and/or limitations to root uptake due to anoxic conditions in the rooting zone during flooding (Kozłowski *et al.*, 1991; Oren *et al.*, 2001).

Comparative ecophysiology of *Metasequoia glyptostroboides*

The following presentation of morphological and physiological traits of *Metasequoia* that may have imparted an adaptive advantage in a temperate, high-latitude forest is based on our experiments and observations, as well as information adapted from the literature. Potential relationships between traits and adaptive advantages for success in the Eocene temperate, high-latitude forest are explored. Where appropriate, attributes of *Metasequoia*

are contrasted with those of *Larix*, which was a minor component of Eocene high-latitude palaeoforests (but commonly dominates extant forests in sub-Arctic biomes) and with other conifers that are currently important components of boreal forests.

Foliar morphology and crown architecture

Metasequoia has both long (persistent) and short (deciduous) shoots. Foliage is decussately arranged (Figure 20.5). Leaves are uninerved with bifacially flattened, linear lamina and decurrent bases. The hypostomatic leaves have two bands of stomata on the abaxil surface and guard cells are surrounded by four to eight subsidiary cells. Non-stomatal epidermal cells have ‘undulating’ walls (Sterling, 1949; Florin, 1952; Srinivasan and Friis, 1989). Examination with light microscopy of freehand sections of living leaves (Figure 20.6) and



Figure 20.5 Shoot and foliage structure of *Metasequoia glyptostroboides*. Arrow points to a late season shoot arising from a leaf base and the scale bar is 2 cm long.

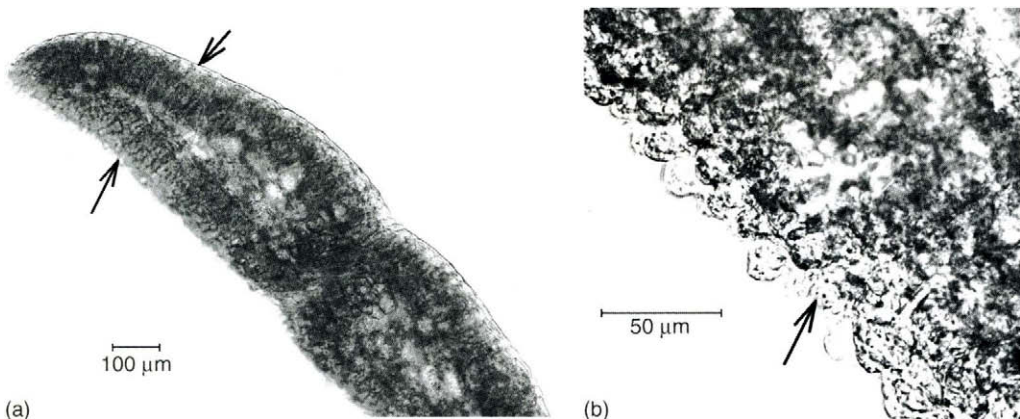


Figure 20.6 Microphotographs of cross-sections from *Metasequoia glyptostroboides* leaves. (a) Presents a comparison of abaxial and adaxial epidermal cell morphology. Note the thin exterior walls on the abaxial cells and thicker adaxial walls (arrows). (b) The protruding nature of adaxial epidermal cells and the presence of chloroplasts are apparent. The arrow points to a protruding abaxial epidermal cell containing chloroplasts.

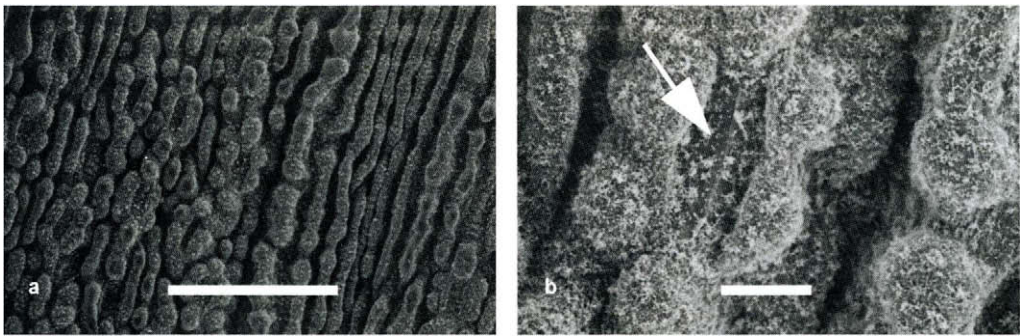


Figure 20.7 Scanning electron micrographs of the adaxial surface of a *Metasequoia glyptostroboides* leaf. (a) The convoluted surface arising from protruding epidermal cells and (b) stomatal guard cells (arrow) surrounded by raised subsidiary cells. Scale bars: (a) = 100 μm ; (b) = 10 μm .

scanning electron micrographs of leaf surfaces (Figure 20.7), revealed that *Metasequoia* leaves have thickened outer walls on adaxial epidermal cells, while those on the abaxial surface have very thin walls. Abaxial epidermal cells contain chloroplasts, which seem to be lacking in their adaxial counterparts. In addition, the abaxial epidermis is characterized by protruding cells interspersed with deep grooves (Figures 20.6 and 20.7a). Stomata are hypostomatic and not sunken, while subsidiary cells are raised (Figure 20.7b).

In Table 20.1 we compare crown morphological features and foliar attributes for *Metasequoia glyptostroboides*, *Larix laricina* and *Picea rubens* Sarg. An interspecific comparison of photosynthetic response to light is shown in Figure 20.8. For these comparisons, *P. rubens* was included as a representative cold-temperate, shade-tolerant species and *Pinus banksiana* Lamb. as a cold-temperate, shade-intolerant species adapted to water-limited habitats.

Metasequoia has some characteristics which are typical of extant shade-intolerant species, such as crown architecture, regeneration strategy, indeterminate shoot growth, rapid shoot elongation, high photosynthetic efficiency (initial slope of light response curve) and low diversity in photosynthetic light-response and foliar display between sun- and shade-adapted foliage (Givnish, 1988; Walters and Reich, 1999). In contrast, traits such as large differences in specific leaf area between sun- and shade-adapted foliage, horizontal leaf orientation, planar foliage display and high foliar density and LAI are generally characteristic of shade-tolerant species (Givnish, 1988). Other attributes, such as unusual presence of chloroplasts in the epidermal cells of abaxil leaf surfaces and their thin outer walls, are more characteristic of shade-loving ferns and other pteridophytes.

Several anatomical characteristics suggest that *Metasequoia* is best adapted to an environment with high atmospheric humidity. These include a high surface-to-volume ratio, enhanced by a convoluted lower epidermis (see Figure 20.6 and 20.7a), non-sunken stomata with raised subsidiary cells (see Figure 20.7b), thin epidermal walls on the abaxial surface and lack of a hypodermis. This contrasts with the more xerophytic leaf anatomy of *Larix*, *Picea* and many other conifers. Whether the humid palaeoenvironment included fog is not known, but we know that *Sequoia sempervirens* (D. Don) Endl. satisfies a large portion of its water needs through fog capture (Dawson, 1998). Dew capture might be a valuable asset during periods of reduced soil moisture (Munne-Bosch and Alegre, 1999). In *Metasequoia*, condensation of dew from humid air would be enhanced by the spreading

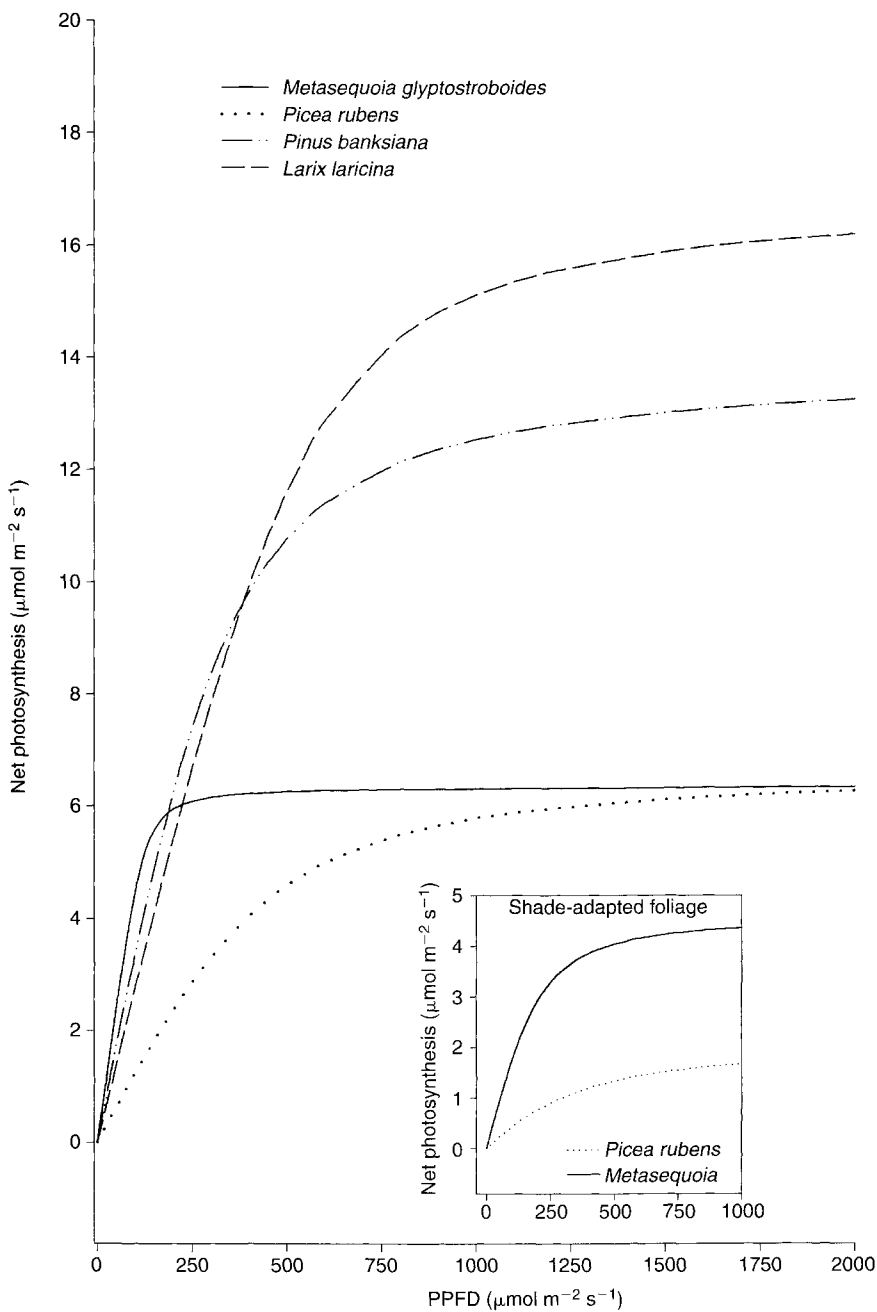


Figure 20.8 Photosynthetic light-response curves for sun-adapted foliage of *Metasequoia glyptostroboides*, red spruce (*Picea rubens*), jack pine (*Pinus banksiana*) and eastern larch (*Larix laricina*) at photosynthetic photon flux densities (PPFD) $< 1000 \mu\text{mol m}^{-2} \text{s}^{-1}$. *M. glyptostroboides* shows photosynthetic efficiencies similar to shade-intolerant species (*P. banksiana* and *L. laricina*) at low PPFD levels and a rapid approach to the light-saturated state, giving it much greater carbon gain per unit leaf area than the shade-tolerant *P. rubens*. Insert compares shade-adapted foliage of *M. glyptostroboides* and *P. rubens*, showing the similarities between response curves of sun- and shade-adapted *M. glyptostroboides* and their great divergence in *P. rubens*. Curves were modelled from light-response data for 6 saplings of each species, with data collected under standardized conditions.

foliar display which would increase the rate of radiational heat loss and more effectively lower foliar temperatures. In addition, the increased surface-to-volume ratio resulting from the convoluted abaxial surface may increase convective heat transfer and enhance the efficiency of dew collection (Muselli *et al.*, 2001).

Most monopodial conifers have determinate growth (Kozłowski *et al.*, 1991). However, *Metasequoia* shows both a monopodial growth habit and indeterminate shoot growth, a pattern more common in angiosperms. With adequate moisture, new leaves are produced throughout the growing season. This growth habit provides a continuous sink (new shoot growth) for carbohydrates produced in neighbouring leaves (Luxmore, 1991), which would reduce sink limitations to photosynthesis and associated challenges. Although sink-limitation of photosynthesis has been relatively unexplored in conifers, indirect evidence has been reported for *Pseudotsuga menziesii* (Mirb.) Franco (Leverenz, 1981) and *Picea rubens* (Day, 2000) and experimentally demonstrated for *Pinus taeda* L. (Myers *et al.*, 1999).

In *Metasequoia*, new foliage can arise on epicormic shoots on small stems or branches, as side shoots on short shoots, as new flushes on long shoots and occasionally from the bases of leaflets on short shoots. These subsequent shoots may often be oriented in a vertical direction to maximize light capture in the crown (see Figures 20.1 and 20.5). In general, carbon allocation to foliage is positively associated with high relative growth rates and competitive advantage in woody species (Poorter, 1989, 1999; Cornelissen *et al.*, 1996). The indeterminate growth habit of *Metasequoia* may be a critical component of its ability to dominate the high-Arctic forests by: (1) enhancing growth potential and capture of growing space in the long growing season; (2) permitting new foliage to be produced and displayed to adjust effectively to changing angular light distribution; (3) producing new foliage with morphological and physiological traits optimized to the particular light intensity at the time of development; and (4) providing a constant sink for photosynthate, thereby reducing problems associated with end-product down-regulation of photosynthesis. This growth habit confers the ability to develop foliage that is morphologically and physiologically optimized for the light environment in which it develops. As we presented in our list of challenges faced by species in the unique high-Arctic light regime, this is theoretically an important adaptation to low-intensity, low-angle incident light during the Arctic spring and fall and moderate intensities during the summer. In addition, this habit permits exploitation of changing distributions of light resources due to the effects of competition and small-scale disturbances (Canham and Marks, 1985; Canham, 1989).

Based on our greenhouse and field experiments and observations of planted trees in various environments, *Metasequoia* trees rapidly develop extensive crowns with high foliar density (FD) under favourable conditions of moisture and light, which contrasts with the much lower FD and LAI of *Larix* (see Table 20.1). In three 20+ year-old planted *Metasequoia* stands in Japan, Vann *et al.* (see Chapter 21) reported only 1–3% of incident PPFd penetrated to the forest floor. This degree of light interception is typical of climax forests (Landsburg and Gower, 1997). In *Metasequoia* plantings receiving full sunlight for only part of the day, we have observed the rapid development of a closed canopy. However, if trees are growing under full sunlight for more than 6 hours per day, shoot elongation is greatly reduced and the leaves are smaller and appear chlorotic. Under these conditions both crown size and foliar density are reduced. Under dense shade, shoot elongation is favoured over leaf production, a characteristic typical of shade-intolerant species (Kozłowski *et al.*, 1991). Typical shade-tolerant conifers such as *P. rubens* can survive for decades under dense shade as suppressed saplings (Davis, 1991). This advance regeneration is then available to respond to small-scale disturbances to attain upper canopy status.

In contrast, *Metasquoia* appears to perform poorly under dense shade, which is, in part, a consequence of producing foliage with a one-year lifespan. In temperate gymnosperms, leaf longevity is correlated with shade-tolerance (*P. rubens* can retain needles for up to 16 years). Among the genera of deciduous gymnosperms native to North America, *Larix* is classified as very shade-intolerant and *Taxodium* is described as requiring full sunlight for maximum growth (Burns and Honkala, 1990). Ida (1981) found that *Metasequoia* increased its leaf weight much more effectively at low light intensities than did *T. distichum*.

Among conifers, specific gravity (sp. gr.) of wood varies considerably. Those adapted to windy sites generally produce denser wood. *Metasequoia* has an unusually low sp. gr. among conifers. Depending on source, sp. gr. ranges from 0.29 to 0.31, while that of *Larix* species ranges from 0.57 to 0.59 (Alden, 1997; Polman *et al.*, 1999). Thus, considerably less photosynthetic resources are needed to create the same height of supporting trunk for a *Metasequoia* tree compared to *Larix*. *Metasequoia* partially compensates for low wood density by producing a fluted trunk near its base to provide additional lateral stability. *Metasequoia* may also be competitively favoured over *Larix* in crown attributes such as FD and LAI (see Table 20.1). These traits in *Metasequoia* more closely resemble those of the more shade-tolerant conifers (e.g. *Picea* and *Abies* spp.) that often dominate better quality sites in boreal and cool-temperate forests to the exclusion of *Larix*. Extant *Larix* is often relegated to bogs or steep mountain slopes due to competition on better sites from conifers producing substantial shade (Schmidt and McDonald, 1995).

Photosynthetic light-response

As with its morphology and growth habits, *M. glyptostroboides* shows characteristics in its photosynthetic light-response curve (see Figure 20.8) that are typical of both sun- and shade-adapted species. Photosynthetic quantum efficiency (QE: mol CO₂ fixed per mol photons absorbed, i.e. the initial linear slope of the response curve) is similar to the shade-intolerant *L. laricina* and *P. banksiana* at low light levels, but its maximum photosynthetic rates are more comparable with the shade-tolerant *P. rubens*. However, in contrast to *P. rubens*, maximum photosynthetic rates are reached at a far lower light intensity, resulting in a response curve with much higher convexity (lower loss in apparent quantum efficiency between its initial linear portion and light-saturation). Additionally, the photosynthetic light-responses of sun- and shade-adapted foliage show remarkably less divergence in characteristics than those of typical shade-tolerant conifers (compare with *P. rubens*, insert in Figure 20.8). These physiological traits contrast with the significant sun-shade divergence with respect to specific leaf area (see Table 20.1). This inconsistency between high morphological and low physiological variation along the sun-shade adaptation continuum can be interpreted as adaptations to two opposing challenges in the model of the high-latitude light environment. The stochastic probability that foliage produced under low light intensity might become 'obsolete' as light resources are limited by shading within the canopy warrants minimal investment of resources. Opposing this is the predictable seasonal cycle of foliar development under low light intensities followed by exposure to much greater intensities as the growing season progresses (see Figure 20.3). In this interpretation, *Metasequoia* evolved a compromise response by minimizing carbon investment in foliage developed under low illumination, while investing greater resources in the photosynthetic apparatus than prevailing light conditions would warrant. Carbon investment in foliage is permanent and is lost if a leaf senesces due to low light availability. However, most nutrients (e.g. N, P, K, Mg) are mobile and are commonly retranslocated from senescing foliage before abscission. Therefore, investment of nutrients in

the photosynthetic system has a lower risk of permanent loss to the tree and provides the potential for delivering higher photosynthetic output if, or when, higher light resources become available. Interestingly, this strategy, while adaptive to the unique light regime of the high-Arctic, would have much reduced value under forest light regimes in current temperate zones, where the light environment under which foliage develops would be highly correlated with the light environment throughout the growing season. In species of similar leaf longevity, investment in nutrient resources closely parallels investment in carbon and nutrient resources (Evans, 1989; Hikosaka and Terishima, 1995; Hollinger, 1996) and may be an important physiological trait for shade-tolerance (Niinemets *et al.*, 1998). The unusual composite of sun-shade characteristics in *M. glyptostrobooides* may be a conserved relict trait from cloudier, wetter paleoclimates in the lower latitudes.

The saturating light intensity for *M. glyptostrobooides* (see Figure 20.8) is close to the predicted maximum intensity for 78°N latitude (see Figure 20.3). This suggests that investment of resources in maximum photosynthetic capacity may be optimal for the high-latitude light environment. The high convexity of the light-response curve for individual leaves compared with those of the other conifers (see Figure 20.8) shows a minimal loss of quantum efficiency from self-shading at the leaf level. A Blackman-response curve, where response increases linearly with input of resource (in this case light) until a saturation point is reached, describes an optimal utilization of resource. Any departure from that response, indicated by a bending of the response curve (non-linearity in the presaturation portion), suggests less than optimal resource utilization. In the case of photosynthetic light-response, this departure from optimality is associated with mutual shading of light harvesting elements (Hikosaka and Terashima, 1995). At the leaf-level, mutual shading is due to the architecture of mesophyll cells and the chloroplasts they contain. Under high intensity illumination, mutual shading may permit capture of 'excess' light by photosynthetic elements lying deeper in the mesophyll as surface elements become light-saturated. However, in an environment characterized by low-angle direct solar radiation at low to moderate intensities, architecture that minimizes mutual shading (e.g. *Metasequoia*) would provide for more efficient resource utilization compared to conifers with more robust needle morphology (e.g. *Larix* or *Picea*). Although we have not developed whole-tree light-response curves, this structural minimization of mutual shading is carried to the shoot-level, where flat-bladed foliage would be more efficient in this respect than the multidirectional, clustered foliar architecture common to many extant conifers, including *Larix* and *Picea*. A flattened foliar display pattern is also associated with more efficient light harvesting in environments of low to moderate light intensity (Canham and Marks, 1985; Givnish, 1988). The presence of chloroplasts in the protruding abaxial epidermal cells of *Metasequoia* (see Figure 20.6b), could additionally enhance capture of low-intensity diffuse light.

Efficiency in CO₂ fixation might be enhanced by transport and/or sink physiology that reduces potential inhibition from the direct and/or indirect effects of photosynthetic end-product accumulation, such as a reduction of starch production in foliage. From an experiment with 72 one-year-old *M. glyptostrobooides* trees planted in a forest opening in Orono, Maine, in which half the trees received natural diurnal light (NL) and half were under CL (supplied by high intensity metal halide lights at night), we found that by mid-July leaves from the NL side had abundant starch, but leaves from trees on the CL side produced little or no starch (as determined microscopically with IKI staining). Omitting starch synthesis in leaves would improve energy efficiency and might reduce investments in catalytic systems. The avoidance of starch production and subsequent catabolism reduces energy costs to the plant but, under CL conditions, requires a continuous and rapid transport of

soluble carbohydrates to a sink, which is provided by indeterminate shoot growth. Furthermore, reducing starch storage in chloroplasts permits more efficient light capture and decreases the possibility of physical damage to grana from accumulating starch (Schaffer *et al.*, 1986).

Water-use efficiency: direct measurements and stable isotope data

High water-use efficiency (WUE; Figure 20.9) and hypostomatic leaves with non-sunken stomates suggest that *Metasequoia* efficiently regulates gas exchange through stomata. This ability would contribute to maximizing carbon fixation under continuous illumination and to adapting to possible wet/dry soil moisture cycles associated with seasonally flooded habitats. By contrast *Taxodium* and *Sequoia* have amphistomatic leaves (Srinivansan and Friis, 1989).

Table 20.2 shows data for stable isotopes of carbon which we have determined for cellulose extracted from fossil *Metasequoia* and *Larix* wood (method of Loader *et al.*, 1997) collected from Axel Heiberg Island (80°N latitude). This is compared to extant

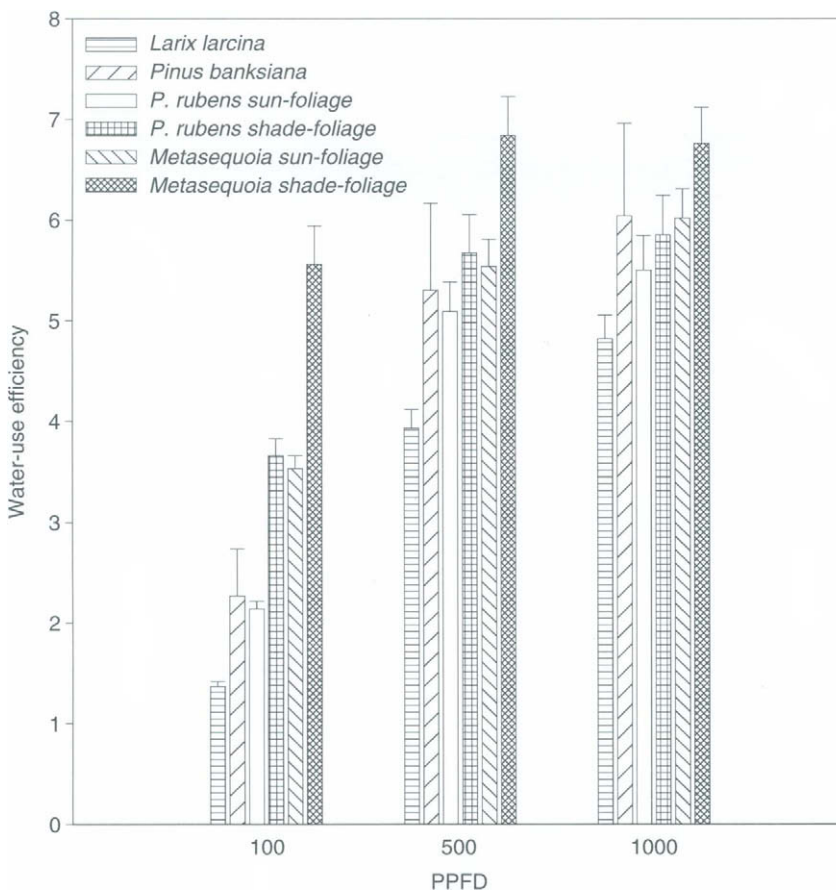


Figure 20.9 Photosynthetic water-use efficiency ($\mu\text{mol CO}_2$ fixed per $\text{mmol H}_2\text{O}$ transpired) for saplings of selected gymnosperms. Species and sampling were the same as those as described for the light-response curves in Figure 20.8.

Table 20.2 Mean $\delta^{13}\text{C}$ values (‰) extracted α -cellulose from both fossil and extant specimens. Means calculated from 6 samples. Standard deviations are given in parentheses.

| | <i>Fossil species</i> | | <i>Extant species</i> | |
|---|-----------------------|--------------|----------------------------|--------------------|
| | <i>Metasequoia</i> | <i>Larix</i> | <i>M. glyptostroboides</i> | <i>L. laricina</i> |
| α -cellulose $\delta^{13}\text{C}$ (‰) | -20.53 | -21.70 | -22.42 | -24.80 |
| Standard deviation | (0.066) | (0.137) | (0.187) | (0.133) |

Metasequoia glyptostroboides and *Larix laricina* wood. Of note is the fact that both fossil woods have more positive values than the modern counterparts and *Metasequoia* is more positive than *Larix* (both fossil and modern).

The $\delta^{13}\text{C}$ values for extant *Metasequoia* and *Larix* species fall mostly within the expected range for extant C_3 species, including conifers (-23 to -34%), although the *Metasequoia* value (-22.42) is slightly more positive. Values for fossil *Metasequoia* and *Larix* are more positive than those usually associated with C_3 plants (O'Leary, 1988). Several environmental factors can promote more positive $\delta^{13}\text{C}$ values by influencing stomatal conductance. These include high irradiance and factors elevating plant water stress such as moisture limitations (Guy and Reid, 1986; Farquhar *et al.*, 1989; Ehleringer, 1993; Gröcke, 1998). Yet the fossil trees are believed to have grown under low level irradiance and were wet site species. Growth rates of the fossil trees (our ring width data and Vann *et al.* (see Chapter 21)) are comparable to that observed in extant *Metasequoia* growing on good sites (Jagels *et al.*, 2003).

Gröcke *et al.* (1999) have recently proposed that $\delta^{13}\text{C}$ values of fossil plant remains are primarily controlled by concentration and isotopic composition of CO_2 in the ocean-atmosphere system (as registered in marine carbonates) and not by palaeoenvironmental and palaeoecological factors. For the 40 million years ago time period, several studies have suggested that atmospheric CO_2 concentrations were higher than at present and more negative values in marine carbonate sources have been reported (Berner, 1991; Cerling, 1992; Freeman and Hayes, 1992). If true, one would expect that $\delta^{13}\text{C}$ values from the fossil specimens would be more negative than that of their NLRs, the opposite of what we have observed. In contrast, the reconstruction of Eocene atmospheric CO_2 concentrations by Royer *et al.* (2001) indicates values comparable with present-day levels. In addition, Pearson *et al.* (2001) have recently proposed that surface water temperatures of tropical seas during the Eocene were 10°C higher than previous estimates. Therefore, if Eocene CO_2 concentrations were near those of the present and global temperatures were driven by factors other than CO_2 concentrations, $\delta^{13}\text{C}$ values of fossil cellulose are likely to be driven by the same ecophysiological factors influencing stable carbon isotope values at present. These would include, stomatal resistance to gas exchange, external and internal water relations, form and rate of photosynthetic end-product production and photorespiration rates. All of these factors would be potentially influenced by light regime periodicity (CL versus diurnal cycle).

WUE and $\delta^{13}\text{C}$ values have been shown to be highly correlated by numerous studies (e.g. Silim *et al.*, 2001). Therefore, $\delta^{13}\text{C}$ values have been widely used as indicators of integrated water-use efficiency. The more negative $\delta^{13}\text{C}$ values of *Larix* compared with *Metasequoia*, for both living and fossil specimens (see Table 20.2), suggest that *Metasequoia* had greater WUE than the second most abundant tree species in the Axel Heiberg Eocene lowland forests. This agrees with the extensive survey by Kloeppel *et al.* (1998) who found that WUE in most *Larix* species, determined from $\delta^{13}\text{C}$ values, were lower than for co-occurring conifers.

Two principal factors control WUE: (1) stomatal control that minimizes transpiration and limits to photosynthesis by decreasing internal CO₂ concentration (C_i); and (2) the photosynthetic efficiency and rates of carbon fixation at a given C_i . To evaluate the first factor, we tested photosynthetic WUE ($\mu\text{mol CO}_2$ fixed per mmol H₂O transpired) for sapling-sized *M. glyptostroboides*, *L. laricina*, *P. rubens* and *P. banksiana* (see Figure 20.9) over light intensities of 100 to 1000 $\mu\text{mol m}^{-2} \text{s}^{-1}$ PPFD. Since none of these trees were drought-stressed and measurements were made under ambient CO₂ levels, the differences are presumed to be characteristic of the genetic potential of each species (Donovan and Ehleringer, 1994; Osório and Pereira, 1994). Shade-adapted *M. glyptostroboides* demonstrated the highest WUE values throughout the range of light intensities. *Metasequoia* sun-foliage had WUE values as high as or higher than all other species by foliar type combinations. In contrast, *Larix* displayed the lowest WUE values. In addition, the relative consistency in WUE of *Metasequoia*, especially its shade-adapted foliage, across the range of light intensities indicates that stomatal conductance is more sensitive to photosynthetic rates. At low light intensity, stomatal conductance decreases to conform with reduced photosynthetic rates in *Metasequoia*, while stomatal conductance remains high in the other species, decreasing their WUE at low light levels. Thus, *M. glyptostroboides* appears to be better able to finely tune its gas exchange to minimize water loss at low to medium light intensities, intensities comparable with those through most of the high-latitude growing season. At high light intensity, *Metasequoia* shade-foliage still maintained the greatest WUE, followed by *Pinus banksiana*, a species adapted to dry, well drained sandy soils (Burns and Honkala, 1990).

While a conceptual model based on stomatal optimization of gas exchange rates is consistent with the interspecific variations in WUE observed at lower light intensities (i.e. *Metasequoia* showing greater optimization efficiency than the other species), its continued maintenance of high WUE at greater light intensities is somewhat surprising, given the much higher light-saturated photosynthetic rates of *L. laricina* and *P. banksiana* (see Figure 20.8). A possible contributor to the high WUE and $\delta^{13}\text{C}$ values in *Metasequoia* is an enhanced level of photorespiratory activity which has been shown to influence net carbon isotope discrimination (Gillon and Griffiths, 1997). This possibility suggests that photorespiration may be a critical process for optimization of carbon utilization, phosphorus cycling and photoprotective functions under CL conditions. We are currently conducting research that will provide a better understanding of these potential mechanisms.

Photosynthetic and accessory pigments

The low intensity light environment of the Arctic spring would make chlorophyll production at low light levels advantageous in temperate high-latitude forests. Wieckowski and Goodwin (1967) and Mitrakos (1963) have reported that chlorophyll synthesis can occur in complete darkness in gymnosperms, including *Metasequoia* (Nikolaeva *et al.*, 1979). Ida (1981) has shown that in the Taxodiaceae chlorophyll production actually increases as light intensity decreases, down to a minimum of about 7% of full daylight. *Metasequoia* produced more chlorophyll at all light intensities than three other members of the Taxodiaceae and, in particular, produced significantly more at low light intensities when compared with the other deciduous member, *Taxodium distichum*. Ida (1981) also found chlorophyll a:b ratios to be negatively correlated with light intensity. Decreases in the chlorophyll a:b ratio have been associated with improved photosynthetic efficiency at low light levels (Kirk and Tilney-Bassett, 1978; Fedtke, 1979).

We have noted differences in leaf colour between trees growing in NL and CL regimes in our experiments. However, we have not determined whether these differences are related to absolute or relative concentrations of chlorophylls and/or accessory pigments. In a preliminary trial where *Metasequoia* seedlings were grown under natural and continuous light regimes under the ambient light intensities of 45°N latitude, the continuous light cycle led to significant production of a red accessory pigment. This pigment in *Metasequoia* has been identified as rhodoxanthin (Czeczuga, 1987), the production of which has been shown to be a photoprotective response (Jagels, 1970; Czeczuga, 1987).

Summary: a conceptual model for the success of *Metasequoia* in Eocene high-Arctic forests

Over millions of years plants have evolved adaptations to environments with widely differing light, temperature and water regimes. These environments can be characterized by both levels (intensities) and temporal distribution of inputs. The environment that has been advanced for Eocene high-latitude forests provided a unique combination of light, temperature and water regimes not currently found on the planet and, therefore, offered an unparalleled set of adaptive challenges to plant species. The temperate continuous light (CL) regime offered both substantial physiological challenges and the potential benefit of high growth rates for species able to overcome those physiological hurdles. Our research to date has allowed us to develop a compelling, although admittedly incomplete, understanding of how the morphological and physiological traits of *Metasequoia* permitted this now relict species to attain such complete dominance of many Eocene high-latitude lowland swamp forests.

Among the unusual aspects of *Metasequoia* is a collection of characteristics that do not fit contemporary models of shade-adaptation or sun-adaptation. Typically, species successful in these opposing light-niches generally do not possess a complete set of shade-adapted or sun-adapted characteristics (Givnish, 1988; Hättenschwiler, 2001). However, *Metasequoia* possesses an aggregation of characteristics not generally associated with either adaptive strategy, but competitively adaptive at high latitudes. While its photosynthetic rates under light intensities typical of the growing season at middle latitudes are clearly much lower than those of sun-adapted conifers (see Figure 20.8), its resource allocation to photosynthetic systems (leaf-level) is close to optimum for the moderate light intensities of the Arctic lowland forests, or cloudier lower palaeolatitudes.

Our research indicates that fixed carbon is rapidly translocated as sugars under CL conditions, a process driven by strong growth sinks and an indeterminate growth habit, minimizing end-product inhibition of photosynthesis. These mechanisms, combined with flexibility in production of photoprotective pigments and perhaps enhanced photorespiratory systems, would be highly adaptive by minimizing damage from excess fluence of light in a CL environment. The indeterminate growth habit would also provide plasticity in leaf display and physiology to permit continual adjustment to changing light resources during the growing season light conditions in an environment characterized by high seasonal variability in light intensity. Finally, the deciduous habit would eliminate carbon losses by leaf respiration during the winter dark period.

In the Eocene lowland forests, *Metasequoia* could rapidly produce canopies with extensive leaf area, not only efficiently capturing incident light for photosynthesis, but minimizing the transmitted light that would be available for competitors. Production of

low density stem wood permitted *Metasequoia* rapidly to overtop potential competitors that were establishing concurrently (Jagels *et al.*, 2003).

It is possible that photorespiration is of adaptive value to growth in a CL environment through several pathways, including photoprotection, enhancing internal cycling and production of antioxidants that lessen the effects of free radicals. Enhanced levels of photorespiration would be consistent with the unusually positive $\delta^{13}\text{C}$ values for extracted α -cellulose. We are currently exploring the importance of photorespiration and alternate pathways for photoenergetics in *Metasequoia* and other species growing under CL regimes.

Acknowledgements

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21

Experimental evaluation of photosystem parameters and their role in the evolution of stand structure and deciduousness in response to palaeoclimate seasonality in *Metasequoia glyptostroboides* (Hu *et* Cheng)

David R Vann, Christopher J Williams and Ben A LePage

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Introduction

Reconstruction of palaeoenvironments is frequently accomplished using floristic analogy based on nearest-living-relatives (NLRs), comparing the species assemblages from the fossil record and inferring the palaeoclimate from that of the closest extant ecosystem.

This method has been applied widely to interpret Tertiary environments (e.g. MacGinitie, 1941; Hickey, 1977; Mosbrugger and Utescher, 1997). In this approach, it is assumed that the physiological requirements and climatic tolerances of the fossil representatives have not changed appreciably through geological time. Reliability of NLR use increases when: (1) there is a close morphological relationship between a fossil species and its NLRs; (2) there are a large number of NLRs represented in a fossil flora that have similar climatic affinities today; (3) the living representatives belong to widespread and diverse groups; and (4) the plant groups used possess anatomical and morphological features related to their climatic tolerances (Wing and Greenwood, 1993). Wolfe (1978) has observed that this approach can be flawed, as many modern NLRs have relictual distributions that fail to reflect the full potential range of the species. More recently, Jordan (1997) has returned to this issue, presenting comparisons of modern and reconstructed climates using species ranges, concluding that temperature estimates for many Tertiary sites are probably overestimated.

The presence of a well-preserved Eocene (45–50 million years ago (Ma)) flora within the ‘Fossil Forest’ at the Napartulik site in the Geodetic Hills site of Axel Heiberg Island, Nunavut, Canada (Francis and McMillan, 1987; Basinger *et al.*, 1994) has presented an unparalleled opportunity to explore some of the difficulties and test some of the assumptions inherent in floristic analogy and the NLR approach to palaeoenvironmental reconstruction. The dominant species at the Fossil Forest site is *Metasequoia occidentalis* (Newberry) Chaney (McIver and Basinger, 1999, and references therein), which is morphologically indistinguishable from the modern NLR, *Metasequoia glyptostroboides* Hu *et* Cheng. The site consists of coaly layers alternating with sandy layers; the coaly layers contain, *in situ*, rooted stumps and forest floor litter layer with palaeosols and *ex situ*, detached logs in sandy layers as well as in the coaly layers (see chapter 20; McIver and Basinger, 1999 and other references herein for additional details).

Application of the floristic analogy approach to the Axel Heiberg site has led some authors to conclude that the palaeoenvironment at this site was warm-temperate (Francis, 1988; Basinger *et al.*, 1994; McIver and Basinger, 1999); it has been described in the popular press as a ‘tropical’ or ‘Carolinian’ flora (Basinger, 1986; Struzik, 1999). However, the absence of certain important genera which define the modern ‘Carolinian’ flora of the southeastern USA (e.g. *Magnolia* L., *Sabal* Adans., *Serenoa* Hook. f.) underscores the limitations of floristic analogy, as without these clearly warm-temperate genera, the Arctic flora is no different from a ‘Pennsylvanian’ flora (Table 21.1).

An alternate approach to the use of floristic analogy is the use of the autecological characters of the NLR to infer the fossil species’ ecological environments (Wing and DiMichele, 1992). These authors outline three assumptions underlying this approach: (1) there must be a close morphological/anatomical relationship between the fossil and extant taxa; (2) the extant species can be found in a full range of suitable environmental conditions; and (3) there has been little evolution in the organism’s ecological tolerance. The Napartulik site provides an opportunity to apply these rules in as close a fashion as is likely possible with the fossil record. The living and fossil *Metasequoia* Miki used in our analyses are morphologically indistinguishable and based on this similarity, the Eocene *Metasequoia* could be assigned to the extant species. The specimens from this site are ‘mummified’ material, consisting of compressed and desiccated, but not decayed, original plant matter (Basinger *et al.*, 1994; McIver and Basinger, 1999). Consequently, we have high confidence that the two species are identical from a morphological standpoint, satisfying the first condition.

Table 21.1 List of genera found in three floras

| Genus | Flora | | | Genus | Flora | | |
|-----------------------|--------|-----|-----|----------------------|--------|-----|-----|
| | Arctic | PA | GA | | Arctic | PA | GA |
| <i>Acer</i> | + | + | + | <i>Metasequoia</i> | + | - | - |
| <i>Alnus</i> | + | + | - | <i>Myrica</i> | + | + | + |
| <i>Betula</i> | + | + | + | <i>Nyssa</i> | + | + | + |
| <i>Carya</i> | + | + | + | <i>Pachysandra</i> | + | - | + |
| <i>Castanea</i> | + | + | + | <i>Picea</i> | + | + | - |
| <i>Cercidiphyllum</i> | + | - | - | <i>Pinus</i> | + | + | + |
| <i>Corylus</i> | + | + | - | <i>Pterocarya</i> | + | - | - |
| <i>Diervilla</i> | + | + | - | <i>Quercus</i> (E/D) | -/+ | -/+ | +/+ |
| <i>Engelhardtia</i> | + | - | - | <i>Sabal</i> | - | - | + |
| <i>Fagus</i> | + | + | + | <i>Serenoa</i> | - | - | + |
| <i>Fraxinus</i> | + | + | + | <i>Salix</i> | + | + | + |
| <i>Ilex</i> | + | + | + | <i>Tilia</i> | + | + | + |
| <i>Juglans</i> | + | + | - | <i>Tsuga</i> | + | + | - |
| <i>Magnolia</i> (E/D) | -/- | -/+ | +/- | <i>Ulmus</i> | + | + | + |

Arctic flora is the fossil assemblage found at the Napartulik site, Axel Heiberg Island; PA is Pennsylvania, USA, a cool-temperate region (Mean annual temperature (MAT) 10–12°C, with frost); GA is coastal lowland Georgia, USA, a warm-temperate region ('Carolinian'; MAT 12–15°C, frost-free). + = present, - = absent, E = evergreen, D = deciduous. Data from Sargent, 1965; McIntyre, 1991).

Subsequent to the discovery of a large relict population of *M. glyptostroboides* in China in the late 1940s (Chaney, 1948), seeds were widely distributed to botanical gardens and academic institutions throughout the world. Horticultural interest has resulted in its propagation at many additional sites since that time. As a result, we have access to mature individuals growing under a very wide range of conditions that appear to exceed the range over which *Metasequoia* has been found in the fossil record, thus fulfilling the second condition.

The third condition, little evolutionary change in the ecological preference of the species, is more difficult to assess. There are no morphological/anatomical differences; consequently, any such features tied to environmental conditions are present in both the extant and fossil forms. From this aspect, their ecological constraints are the same. Changes in physiological responses to the environment could well arise from more subtle responses, such as changes in the cell biochemistry or enzyme efficiencies. Such biochemical differences are more difficult to assess, as organic compounds tend to deteriorate with time. At this point, we have extracted some residual biochemicals and expect to analyse this question in some detail. Preliminary analyses of amber resin suggest that, apart from the expected oxidation products, there is little difference between the chemical composition of fossil and the extant amber (Anderson and LePage, 1995). We do not expect to detect many differences in the biochemical constituents of the two species.

Metasequoia first appears in the fossil record near the Arctic circle (60–63°N) during the Cenomanian (about 95 Ma) and became widely distributed during the Palaeogene (Yang and Jin, 2000); the modern form was found growing around 30°N latitude in a small portion of central China. It is reasonable to ask whether any physiological changes occurred during the expansion and contraction of this species' distribution, as global climate cooled and the species went from a largely aseasonal to a strongly seasonal habitat with a markedly different light regime. To address this question, we have begun exploring the

physiological ecology of *M. glyptostrobooides*. If we find evidence of changes in the species' physiological response in the modern environment, this forms a model system from which to infer physiological evolution in response to changing climates. Conversely, there may be no evidence for physiological differences between the modern *M. glyptostrobooides* and the fossil genus. In this case, as *Metasequoia* is a major component of the northern hemisphere flora throughout the Tertiary (Tidwell, 1998), understanding the ecology of *M. glyptostrobooides* should provide important insights into the nature of many Tertiary palaeoenvironments.

In particular, we would like to consider three questions that this system seems well suited to address, as follows: (1) are the environmental tolerances of the modern species consistent with its distribution above the Arctic circle? (2) how do physiology and environment interact to influence stand structure? and (3) could deciduousness have evolved in response to light/dark seasonality?

The high latitude light regime is substantially different from that of the lower latitudes and may represent a selection pressure inducing physiological responses particular to this environment. We conducted a survey of gas exchange parameters to evaluate response to various environmental conditions. The responses were then used to delineate the optimal environment for the species and to identify conditions under which the tree would likely be unsuccessful. From this range of conditions it is possible to infer fossil palaeoenvironments. Conversely, if the fossils are found under conditions for which supporting evidence indicates a climate outside of the modern species' tolerances, one can infer some sort of physiological evolution. Second, we use the species' light-response data, along with observations on modern plantation-grown trees and a seed germination study, to examine how physiology affects stand structure and dynamics. Light levels, branching patterns and tree density provide insights into stand structure. Finally, we examine the possibility that the Arctic light and climatic regime determined leaf turnover rates in this species.

Wing and DiMichele (1992) observe that the floristic analogy approach is more reliable if there are physical characters that are associated with specific climatic tolerances. *M. glyptostrobooides* is an annually deciduous member of the Pinopsida. Conifers are almost exclusively evergreen (defined as holding a cohort of leaves for more than a year) and have been successful in exploiting habitats from the sub-Arctic to the tropics. Seasonal deciduousness has evolved in only four other genera in the Pinopsida; *Taxodium* Rich., *Glyptostrobus* Endl., *Larix* Mill. and *Pseudolarix* Gordon. The closest relatives of each of these taxa are evergreen. The first two taxa are closely related to *Metasequoia*; all three were once separated into a distinct family (Taxodiaceae); recently, this family has been consolidated into the Cupressaceae (Hart and Price, 1990). *Taxodium* is seasonally deciduous in a temperate climate; this genus appears to be very closely related to *Glyptostrobus* (Kusumi *et al.*, 2000), which in turn seems to be largely drought-deciduous, although the ecology of this species is poorly known at this time. Similarly, from known locations, *M. glyptostrobooides* is seasonally deciduous. The evolution of deciduousness in an otherwise evergreen and widely-dispersed clade implies the existence of particular selection pressures. It has been inferred that *Metasequoia* evolved near the Arctic circle (Yang and Jin, 2000). In the high latitudes, it has been suggested that deciduousness may have evolved in response to the seasonal pattern of light rather than temperature (Axelrod, 1984; Read and Francis, 1992). The rationale behind this is that, during a mild winter, the respiratory activity of the leaves would exceed their carbon reserves fixed during the short growing season. At some point, the cost of replacing a leaf would be less than respiratory loss of carbon if the leaf were retained. Using the gas exchange data and biomass estimates from

reconstructed and plantation stands, we provide a simple evaluation of the leaf carbon balance under various winter scenarios.

Materials and methods

Gas-exchange measurements

All gas-exchange measurements were conducted using a portable photosynthesis and transpiration measurement system (LCA-4; Analytical Development Company, Ltd., Hertfordshire, UK) outfitted with PLC4B cuvette. Leaf area was determined using a digital scanner (ScanJet IICx, Hewlett-Packard, Inc., Palo Alto, CA, USA) and image analysis software (SigmaScan Pro 5, SPSS, Inc., Chicago, IL, USA). The LCA-4 was calibrated using standard gases of known CO₂ concentration prior to each use. Light-response curves were obtained using the LCA-4's built-in light and neutral density filters or using 15-cm diameter Tiffen brand neutral density filters placed over the cuvette when in full sunlight in the field. Filter densities ranged from ND0.1 to ND1.0 and were stacked to obtain intervening values. Light intensities were determined using the PLC4B's internal light sensor. CO₂ and humidity were regulated using the LCA-4's built-in controls. Temperature response was determined using the built-in temperature controls for portions of the range. The LCA-4's internal system was unable to regulate the chamber temperature beyond about 10°C above or below the ambient temperature. The response to temperatures below about 15°C was assessed in the laboratory, with the cuvette's Peltier cooler embedded in ice or with the entire cuvette in a refrigerator. Laboratory measurements were performed on branches cut in the field and re-cut immediately in water. Such measurements were made on samples collected from trees at the University of Pennsylvania campus or from the Morris Arboretum, USA.

Field measurements of light and temperature (15–35°C) response were conducted at arboreta or educational institutions spanning a range of latitudes (Table 21.2) using mature trees from the 1948 plant distribution (Wyman, 1970) or their vegetatively propagated

Table 21.2 List of locations used in gas-exchange experiments and whole-tree metrics. All locations in the continental USA or on Honshu Island, Japan

| <i>Location</i> | <i>MAT</i> °C | <i>MAP</i> cm | <i>Elev.</i> m | <i>Lat.</i> °N | <i>N</i> |
|--|------------------|------------------|-------------------|-------------------|----------|
| Morris Arboretum, Philadelphia, PA | 12 | 1040 | 40 | 40 | 8 |
| University of Pennsylvania, Philadelphia, PA | 12 | 1040 | 18 | 40 | 2 |
| National Arboretum, Washington, DC | 13 | 990 | 12 | 39 | 3 |
| Sarah P. Duke Gardens, Duke University, Durham, NC | 15 | 1150 | 120 | 36 | 1 |
| J. C. Raulston Arboretum, Raleigh, NC | 15 | 1150 | 137 | 36 | 3 |
| North Carolina State University, Raleigh, NC | 15 | 1150 | 133 | 36 | 1 |
| Tulane University, New Orleans, LA | 20 | 1665 | 3 | 30 | 1 |
| Loyola University, New Orleans, LA | 20 | 1665 | 3 | 30 | 1 |
| Tanashi Experimental Station, Tanashi, Japan | 13.7 | 1400 | 60 | 35 | na |
| Aono Experimental Forest, Minamiizu, Japan | 15 | 2300 | 180 | 34 | na |
| Kamigamo Experimental Forest, Kyoto, Japan | 15.3 | 1700 | 150 | 35 | na |

MAT = approximate mean annual temperature. MAP = mean annual precipitation. Lat = approximate latitude. *N* = number of individual trees at each site (gas-exchange measurements only). In the USA, all trees are from the 1948 seed collection except at Raulston (3) and Morris (3) Arboreta, which were derived from cuttings. Slope is 0% for all sites except Izu (15%) and Kyoto (20%), both of which had NW aspect.

offspring. There was typically a single tree at each site. Three to six branches on each tree were used for measurements of photosynthesis and transpiration. More intensive sampling and measurements of CO₂ and humidity response, as well as low temperature response (5–15°C) were performed on trees at the Morris Arboretum (seven trees) or on the campus of the University of Pennsylvania (two trees). Measurements were performed on days with full sun or slight overcast; photosynthetically-active photon flux density (PPFD) was greater than the light saturation value for this species as determined in preliminary experiments (and shown below). Dark respiration was determined by completely covering the cuvette with foil. Tests were performed both in the field and in the laboratory at 5, 15, 25, 35 and 40+°C. The lowest and highest temperatures were only tested in the laboratory.

Stand structure

Stand structure for the Napartulik site was inferred by analogy with modern plantations growing in Japan. We compared stem density and diameter with branching patterns from open- and closed-grown plantation trees. We analysed four stands at three experimental forests on Honshu Island, Japan, located in the warm-temperate climate zone. At the Tokyo University Forests, Tanashi Experimental Station, two small stands of differing age and initial planting density were analysed. One stand (hereafter referred to as 'Tanashi A') was established in 1953 from two parent trees. Trees were established on a 2 m square grid pattern and the original plot size was about 600 m². A second stand (hereafter referred to as 'Tanashi B') was established in 1980 and was planted on a 1 m square grid; the stand is about 420 m² in area. The third stand was located on the Izu Peninsula at the Aono Forest of the Arboricultural Research Institute, Minamiizu. The fourth stand analysed was located at the Kyoto University, Kamigamo Experimental Forest. All stands were derived from rooted cuttings.

In June 2000, 14 plantation trees across a range of stem diameters (minimum 8.9 cm, maximum 40.1 cm, measured 1.4 m above the soil; dbh) were selected and sampled using the methods of Vann *et al.* (1998) and Whittaker *et al.* (1974). Briefly, trees were felled and a measuring tape was laid along the stem axis from tip to base. Diameter of the stem was measured at intervals of 1–2 m. The distances of all living and dead branches were tallied from the tip downward along with branch diameter at the point at which it joined the stem. Measurements were made using standard dbh tapes for stem and large diameter branches (>10 cm). Smaller diameters were measured using digital calipers. Five to six live branches were selected from the canopy of each tree and the length, fresh foliage weight and fresh branch wood weight were measured. Field measurements of stem, branch and leaf components were made with a combination of spring scales (10, 25, 50, 125 kg capacity) and a portable electronic scale (5 kg capacity). Dry weight determination of these fractions was based on the fresh/dry weight ratios determined on oven-dried (65°C) subsamples. Stems were cut into logs (1–2 m long). Fresh weights and dimensions of the logs were recorded and sample discs about 20 cm in diameter were cut from the logs near the base, middle and top of each tree for dry weight determination. Dry weights of logs were calculated from the log fresh weights and disc fresh/dry ratios.

Three plantation trees and all open-grown trees from arboretea or academic insititutions were sampled using a non-destructive technique (see Vann *et al.*, 1998). This involved climbing each tree to a point as close to the apex of the tree as possible (typically within 3 m of the apex). This distance was determined based on the canopy density and safety of the climber. The area of the canopy not directly accessible to the climber was assessed

visually and branches were counted and diameters estimated by eye. Total tree height of open grown trees was determined using a distance tape in combination with a height pole raised by the climber. The diameter of each limb below the top was measured and recorded. The climber then moved down the length of the stem measuring stem diameter at 1–2 m intervals and the basal diameters of all branches.

The data for branch diameter, branch length and branch foliar biomass were natural log transformed prior to analysis. Branch wood and foliage dry weights ($n = 80$) were regressed against branch basal diameter using ordinary least-squares regression to predict branch and foliar biomass. These regressions were then applied to the basal diameters of all non-weighed branches and the results summed to estimate the dry weight of branch wood and foliage in the total canopy of each sample tree. Measured weights were related to diameters using ordinary least-squares regressions with the standard allometric formula $Y = \alpha x^\beta$, where $\alpha = e^{\gamma + \varepsilon}$ and $\gamma =$ the regression intercept and $\varepsilon =$ an error term equal to $\frac{1}{2}$ the mean squared error from the regression analysis. The error term compensates for a bias introduced during back-transformation of regression estimates (Bell *et al.*, 1984). Statistical analyses were completed using the statistical program JMP 4.03 (SAS Institute, Raleigh, NC).

Fossil evidence

Fossils examined in this study were recovered from sediments of the Buchanan Lake Formation, Eureka Sound Group on Axel Heiberg Island (Ricketts, 1991). The Buchanan Lake formation crops out sporadically throughout the Sverdrup Basin and is well exposed at Napartulik. Thirty-seven fossil stems were excavated from unconsolidated mud and siltstone layers. We located fossil stems that were protruding from the sedimentary layers and excavated the stem with shovels and pick axes until a complete sample was obtained or permafrost impeded further excavation. We preferentially sampled stems that were buried at shallow depths above the permafrost layer or were oriented in a direction parallel to the slope, which generally permitted greater lengths of stems to be exposed.

The length of each exposed log was measured with standard measuring tapes. Logs were typically fractured into several segments and any gap between each segment was measured to adjust the total length of each log. Diameter measurements were taken at 50 cm intervals along the stem using large metal calipers. We also mapped the location of exposed branch stubs and branch scars on the stem surface using the distance from the basal end of the log as a reference point, as well as an approximate location of the branch stub on the stem at that distance (e.g. top centre or bottom right). For each branch stub, two diameters were measured to the nearest millimetre (major and minor) using digital calipers. Diameters were estimated for missing branch stubs. We also assessed the external morphology of each branch stub to determine whether a callus of wood had formed around the stub. We used the presence of a callus as an indicator of whether the branch was alive or dead at the time of burial (e.g. Mattheck, 1991:34–36). In the course of excavating fossil logs we recovered the remains of seven stems that were characteristic of the upper portions of tree stems with numerous exposed small diameter branch stubs projecting from the surface of the fossil. We applied allometric equations developed from plantation grown *M. glyptostroboides* (see above) to estimate foliar and branch wood dry weights for the fossil tree tops.

Forest stand density for the Napartulik site has been described previously; the forest (designated 'N-layer') was originally surveyed by Francis (1991) and later resurveyed by T. Sweda of Ehime University, Japan. A preliminary analysis of Sweda's mapping data

was published by Basinger *et al.* (1994). Sweda's raw dataset includes measurements of stump diameter (base and top), location and type of soil in which the stump was rooted (i.e. mineral soil or organic soil). We re-analysed Sweda's original dataset of field measurements of *in situ* stumps to determine the stand density and basal area of the N-layer forest. Because forest biomass studies typically report results based on bole diameter at a standard stem height (dbh), we reconstructed fossil stem dbh (cm) using a regression equation relating stump basal diameter (D_b , cm) to dbh developed from plantation grown *M. glyptostroboides* trees ($\text{dbh} = 0.775 D_b^{0.990}$; $R^2 = 0.97$, $n = 444$). We reconstructed fossil stem height using a combination of techniques. Kuser *et al.* (1997) summarized the measurements of 52 *M. glyptostroboides* growing within its native range in China. Using their original dataset we developed the following allometric regression equation expressing the relationship between tree height (m) and dbh (m): $\text{Tree ht} = 42.26 \text{ dbh}^{0.678}$ ($R^2 = 0.63$). We also used sequential measurements made along the length of the log sections to determine a rate of stem taper, from which we derived an estimate of tree height using standard parabolic volume curves and the basal diameters of the fossil stumps.

Light measurements

Measurements of PPFD were determined using a quantum sensor (Li-Cor Corp., Lawrence, KS, USA) attached to a datalogger (LI-1000, Li-Cor). Multiple values (100–200) were recorded for each of the three plantations in Japan; these were obtained via a random walk through the plantation at approximately 2 m intervals or some 4–10 m outside of the plantation in the open. Values for the Arctic site were continuously recorded as hourly averages and total integrated sums during July 1999 and 2000.

Seedling germination study

In the spring of 2001 we established a series of greenhouse experiments to examine the influence of seedbed soil type and the influence of leaf litter on seed germination success of *M. glyptostroboides* seedlings under well-watered conditions. We germinated commercially available seeds (Sheffield Seed Company, Locke, NY) on three types of substrate; Pro Mix Bx (Premier Horticulture, Dorval Quebec, Canada), a *Sphagnum* L. peat based potting soil; pure washed beach sand; and a mineral soil (Earthgro Topsoil, Scotts Company, OH, USA). A second experiment evaluated the effect of two levels of litter layer thickness on germination success. We collected senescent *M. glyptostroboides* litter from a 45 year-old stand in Princeton, NJ. Litter was air dried and manually sorted to remove any seeds. Three treatments included a no litter control, a thin litter treatment (15 g dried litter, equivalent to 2 mm thickness) and a thick litter treatment (35 g dry litter equivalent to 5 mm thickness). Dried litter was spread to a uniform thickness across the surface of each tray.

Both experimental treatments were carried out using an unblocked design (four replicates of three soil types in the case of the substrate treatment; and eight replicates of three litter treatments in the litter thickness experiment). Germination trays were moved to a new position at random every 7 days to avoid any variation in environmental conditions within the experimental area. We germinated 2000 seeds in the soil treatment and 1000 seeds in the litter addition treatment. We then tracked percentage germination in each treatment over a period of 6 weeks. All germination percentages were arcsine transformed prior to statistical analysis. We used ANOVA to analyse the final percentage germination values for each treatment to test the hypothesis that substrate type or presence of litter had no effect on germination success.

Results

Light environment

Average light intensity in the modern Arctic is moderate compared with the values seen at lower latitudes (Table 21.3). Peak values are in excess of that which saturates the photosystem in *M. glyptostroboides* (Figure 21.1B), so light would not limit the photosynthetic rate on clear days. The range of values seen on overcast days are about one-third or less of the clear sky values, but would not be expected to limit significantly carbon gain in this species. On clear days, during the polar summer of continuous light, the total PPF received in the Arctic is about twice that received at 40°N latitude for the same period.

Palaeolatitude for the Fossil Forest site was about 78°N (Irving and Wynne, 1991); the small difference between this value and the current latitude is not expected to result in significant differences in the light environment at these high latitudes.

Gas exchange

Temperature, light, humidity and CO₂ response did not differ significantly between individuals or leaves at either a single location or between locations (data not shown). Consequently, all trees were pooled for the data presented in the accompanying figures. The data shown in Figure 21.1A and B represent all measurements from all trees at all locations; CO₂ and humidity data were determined only from seven trees at the Morris Arboretum. The CO₂ uptake rate shows a fairly broad response to temperature, with the maximal values occurring between 15° and 25°C (Figure 21.1A). Based on the general shape of the curve, it appears that the optimal temperature for photosynthesis in this species is about 20°C. This peak is about 2.5 times the rate seen at 5°C. The rate of CO₂ uptake drops rapidly above 25°C, with one half the maximum rate at about 30°C; above 35°C the rate drops very sharply. Leaves exposed to temperatures above 40°C did not return to pre-exposure rates within one hour, suggesting damage to the photosynthetic apparatus (data not shown).

The CO₂ uptake rate of *M. glyptostroboides* shows a very rapid response to low light intensities, saturating at a relatively low light intensity, about 700 μE (Figure 21.1B), less than one-half the typical clear-day intensity seen in the Arctic (Table 21.4). The leaves

Table 21.3 Summary of light intensity (PPFD) measurements made at several sites

| <i>Light received (PPFD)</i> | | | <i>Canopy light penetration</i> | | |
|------------------------------|-----------|----|---------------------------------|------------|-----------|
| | | | Above | Below | |
| <i>Arctic (summer peak)</i> | | | | | |
| Full sun | 1500–1700 | μE | Tanashi | 458 ± 75 | 9 ± 3 μE |
| Overcast | 300–500 | μE | Izu | 1415 ± 123 | 15 ± 6 μE |
| Total per day | 130 | E | Kyoto | 476 ± 113 | 16 ± 9 μE |
| <i>Low latitude</i> | | | | | |
| Full sun | 2200–2400 | μE | | | |
| Total per day | 60 | E | | | |

Values are average peak intensities except total per day, which is the total integrated input for 24 hours (Arctic) or 10 hours (Philadelphia). Arctic is the Napartulik site on Axel Heiberg Island., Nunavut, Canada; low latitude is Philadelphia, Pennsylvania, USA. Canopy light measurements were made in Japan, at or near the cities listed. *Note:* measurements at Tanashi and Kyoto were performed on overcast days. Units: E = Einsteins, or moles of photons m⁻²s⁻¹ (for wavelengths active in stimulating photosynthesis).

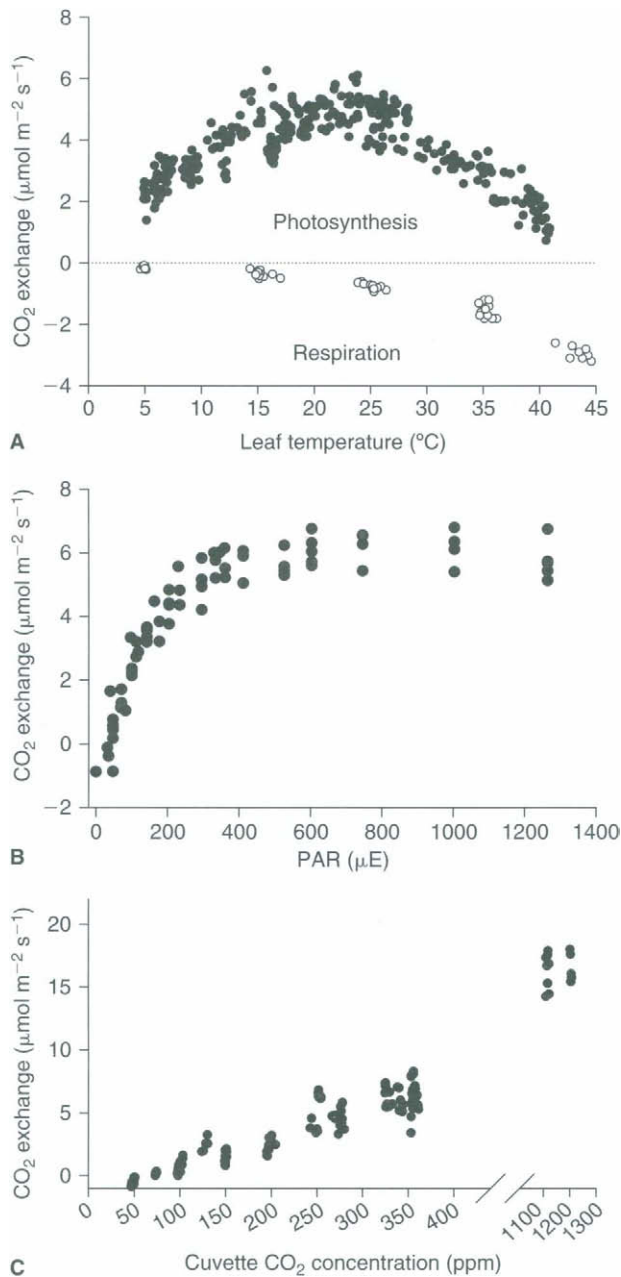


Figure 21.1 Carbon dioxide exchange in *Metasequoia glyptostroboides* in response to several environmental parameters. A. Response to temperature; closed circles represent net photosynthesis; open circles show 'dark' respiration. B. Response to light intensity, measured as photosynthetically active radiation ($\mu\text{mol photons m}^{-2} \text{s}^{-1}$). C. Response to CO₂ concentration of the input air stream.

maintain one-half the saturated rate at a very low 150 μE . However, below 100 μE , carbon gain is trivial. The low light saturation value indicates that the photosystem is unable to exploit the higher light intensities seen at lower latitudes. Following Leverenz (1987), we calculated the convexity of the light response curve (θ). The value, 0.9, is high compared

Table 21.4 Metrics from tree, plantation and fossil forest analysis

| | | <i>Extant</i> | | | | | | | | | <i>Fossil</i> |
|--------------------------------------|------------|--------------------|--------------|--------------|-------------|-------------------------|----------|------|--------|------|-----------------------|
| | | <i>Plantations</i> | | | | <i>Open grown trees</i> | | | | | |
| Site | | Tanashi | Tanashi | Izu | Kyoto | Kyoto #8 | Kyoto #1 | NCSU | Tulane | Duke | Napartulik |
| Age | (years) | 20 | 48 | 48 | 48 | 48 | 48 | 50 | 50 | 50 | >250 |
| Original planting density | (trees/ha) | 10000 | 2500 | 3086 | 3500 | na | na | na | na | na | unknown |
| Tree density | (trees/ha) | 3219 | 816 | 2000 | 1000 | na | na | na | na | na | 480/1275 ^c |
| Height | (m) | 18 | 30.7 | 31 | 28.5 | 37.6 | 37.1 | 22 | 27.4 | 19 | 39, 42 ^d |
| Length of canopy | (m) | 6.9 | 10 | 9.5 | 8.3 | 35.3 | 32.8 | 21.5 | 26.4 | 18 | ca. 9 |
| Average dbh | (cm) | 12.1 | 35.8 | 25.9 | 24.4 | 76.8 | 62.1 | 67.4 | 66 | 103 | 41/25 |
| dbh range (cm) | (min, max) | (1.5, 40.1) | (19.8, 50.6) | (10.0, 65.7) | (6.2, 52.0) | na | na | na | na | na | 20, 131 |
| Longest branchless span ^a | (m) | 8.4 | 7.5 | 24.3 | 14 | 4 | 2.3 | 1.5 | 1.3 | 0.8 | 6.5 ^b |
| Average branch diameter | (cm) | 0.88 | 3.4 | 2.1 | 1.6 | 5.1 | 4.8 | 7 | 5.8 | 11 | 2.5 |
| Minimum | | <0.5 | <0.5 | 0.5 | <0.5 | 1.8 | 1.1 | 1.9 | 1.1 | 1.1 | <0.5 |
| Maximum | | 3 | 7.6 | 5.8 | 4.1 | 10.49 | 9.2 | 12.8 | 12.6 | 16.2 | 7.6 |
| Average tree biomass | | | | | | | | | | | |
| Trunk | (kg) | 50 | 338 | 204 | 170 | | | | | | 660 ^e |
| Canopy foliage | (g) | 2000 | 11639 | 7248 | 6231 | | | | | | 6582 |
| Canopy branch | (g) | 5064 | 42606 | 24672 | 19692 | | | | | | 27105 |

^a without live branches in living trees; without emergent branch remnants in fossils.

^b from an 8.4 m fossil log.

^c values represent stems with dbh > 15 cm/stems of all recorded sizes.

^d first value based on parabolic taper, second value based on regression.

^e reconstructed values.

with many needle-leaved conifers (0.44–0.58; Leverenz, 1996), but is consistent with the planar nature of the leaves and the observed shade intolerance.

Short-term exposure to elevated CO₂ resulted in an equivalent increase in net photosynthesis (Figure 21.1C). Carbon gain decreased rapidly with declining CO₂ concentrations, reaching the compensation point between 50 and 100 ppm, consistent with the general pattern seen in C₃ plants. There was little response to changes in humidity over a range of 20–80% RH. Measurements at higher RH values were confounded by temperature effects. Below 20% RH, carbon exchange became erratic and unstable. Based on transpiration data (not shown), it appeared that the stomata were opening and closing irregularly, possibly in an attempt to conserve water.

Dark respiration measurements were performed at five different temperatures; values at the highest temperature ranged from 1.6 to 3.2 $\mu\text{mol CO}_2\text{m}^{-2}\text{s}^{-1}$ (mean 2.4, $n = 8$). At this setting ($>40^\circ\text{C}$), the gas analyser had difficulty in maintaining a steady temperature. At 35°C , the CO₂ efflux rate varied from 1.2 to 1.8 $\mu\text{mol m}^{-2}\text{s}^{-1}$ (mean 1.6, $n = 12$); at 25°C the rate ranged from 0.6 to 0.93 (mean 0.76, $n = 12$); at 15°C the mean value was 0.35 (range 0.18–0.51, $n = 12$); and at the lowest temperature, 5°C , the range was 0.09–0.22 (mean 0.15, $n = 8$). This is summarized in Figure 21.1A.

Stand structure

Plantation grown *M. glyptostroboides* exhibited rapid height growth, attaining relatively uniform canopy heights within each plantation. Average height of canopy dominant trees in the 20-year-old stand was 18 m. Canopy dominant trees in all of the 48-year-old plantations attained similar maximum tree heights (28.5–31 m). The height corresponded to vertical growth rates of approximately 0.9m year^{-1} in the youngest plantation and between 0.6 to 0.65m year^{-1} in the older plantations.

Average stem diameter increased with decreasing original planting density (see Table 21.4). Despite relatively uniform canopy heights, we found that stem diameter varied considerably within a given stand (see Table 21.4). This is reflected by an overall poor relationship (e.g. Tanashi B, $R^2 = 0.19$) between bole diameter and total tree height of the canopy dominants in a given plantation.

On average, the youngest trees possessed the smallest branches and expressed the least variability in branch diameter (mean 0.88 cm, range $<0.5\text{--}3.0$, $n = 372$). In contrast, trees in the older, low density plantation (Tanashi B) had the largest average branch diameter (mean 3.4 cm) and the widest range ($<0.5\text{--}7.6$, $n = 390$). As expected, we found that the stands of trees with the largest diameter branches also had the largest foliar and branch biomass. Length of the live canopy differed among plantations (see Table 21.4). However, when expressed as a function of total tree height the proportion of live canopy to total tree height was similar among the older plantations (29 to 31%) but slightly higher in the youngest plantation where 38% of the total tree height was covered by live canopy.

Open grown solitary trees differed greatly from plantation grown trees in their overall structure. Total tree height varied considerably among solitary trees (range 19–37.6 m) and solitary trees had significantly larger stem diameters than plantation grown trees of similar age. The canopy structure of solitary *M. glyptostroboides* differed considerably from plantation grown trees. Maximum branch diameters were observed on the Duke Gardens tree (mean 11.0 cm, range 1.1–16.2, $n = 104$); whereas Kyoto #1 was observed to have the smallest branches and least variability of the open grown trees (mean 4.8 cm, range 1.1–9.2, $n = 129$). We found no sign of self-pruning occurring on open grown trees. The canopy of open grown *M. glyptostroboides* was more extensive on solitary as

opposed to plantation grown trees (see Table 21.4). This corresponds to a smaller amount of branch-free stem wood (expressed as the longest span between live branches) on open grown trees than on plantation grown trees. In contrast, plantation trees exhibited signs of strong self-pruning and long lengths of bole lacking live branches.

Fossils recovered from the Napartulik locality indicate a large stature, low-density forest of moderate to high biomass (see Table 21.4). We estimated fossil tree height to be 39 m using the tapers of excavated logs. Using reconstructed dbh of the ten largest well-preserved fossil stumps and the height to diameter relationship of modern *M. glyptostroboides* from China we estimated a fossil tree height of 42 m. Of the 37 fossil logs recovered from the sediments, the longest log with no evidence of exposed branches was 8.4 m in length; 6.5 m of the log was clear wood with no branches. A 16.7 m log was also exposed, which had 6 m of branch-free wood. We estimated the live canopy of the fossil *Metasequoia* trees was approximately 9 m on the basis of the length of fossil tree tops we recovered. Branch diameters measured on exposed branch stubs from seven nearly complete upper stem (i.e. canopy) segments averaged 2.5 cm; range <0.5–7.6, $n = 565$). We estimated the average fossil canopy consisted of about 6580 g dw of foliage and 27.1 kg dw branch wood using allometric equations derived from plantation grown trees.

Seedling germination

The majority of seeds germinated 7 days after sowing and no new seedlings emerged after 28 days in any treatment. Seedling germination was low in all treatments, with approximately 5% maximum germination (Table 21.5). Soil type did not affect the germination rate significantly ($F_{2,9} = 1.1439$, $P > 0.3$), although it is possible that it would have had an effect on survival, which we did not follow beyond 4 weeks. Depth of burial, however, did have an effect with decreasing germination as the litter layer increased in depth; the thick litter layer reduced germination significantly ($F_{2,9} = 48.67$, $P < 0.0001$). Similarly, we observed no germination within the plantations in Japan, although seedlings were observed on an open road cut across from a plantation and in light shade under trees at the road's edge in the Kyoto plantation (personal observation).

Discussion

The Napartulik site has been reconstructed as a deltaic environment, either a braid- or meander-plain (Ricketts, 1991; McIver and Basinger, 1999). The remnant forests in

Table 21.5 Results of germination trials of *Metasequoia glyptostroboides* germinated on differing substrates and in the presence or absence of *Metasequoia* litter. Values followed by the same letter are not significantly different ($P > 0.05$)

| Substrate type | Germination percentage (mean followed by s.e.) | |
|------------------------------------|--|---|
| Potting soil | 5.1 ± 0.4 | a |
| Mineral soil | 4.6 ± 0.4 | a |
| Beach sand | 4.4 ± 0.2 | a |
| Burial depth (all on potting soil) | | |
| No litter | 5.0 ± 0.4 | a |
| ~2 mm depth (thin layer) | 4.2 ± 0.2 | a |
| ~5 mm depth (thick layer) | 1.3 ± 0.3 | b |

China exist in a dissimilar habitat (Yang and Jin, 2000); it is montane and, although the species tends to be concentrated near riparian channels, it is lacking a permanent water source. Consequently, a modern analogue of the Napartulik site does not exist. However, several lines of evidence suggest that the ancient forests followed a riparian model, becoming established on open flood sediments as even-aged cohorts (*sensu* Oliver and Larson, 1996:145–167). First, the seeds are small and winged, similar to *Sequoia sempervirens* [D. Don (Endl.)], whose seeds do not fall far from the tree (Olson *et al.*, 1990); however, they do float. Second, the seedling germination study indicates that the best germination rate occurs on open soil (see Table 21.5), as opposed to beneath forest floor litter. This is consistent with results from *Sequoiadendron giganteum* [(Lindl.) Buchholz], whose seedlings rarely become established on undisturbed forest floor, apparently resulting from rapid desiccation of the thick litter (Olson *et al.*, 1990). Moreover, Falder *et al.* (1999) describe *in situ* fossil *Metasequoia* seedlings from a Palaeocene floodplain environment and interpret *Metasequoia* as a colonizer of marshy, mineral soil habitats. Finally, the species attains its best growth in the absence of water stress (Kuser, 1982). Considering the low light levels within the forest stands (see Table 21.3), it seems reasonably evident that the establishment of individuals within the stand is unlikely, consequently, multi-cohort stands would not result. Rather, the simultaneous germination of individuals on open flood soils would produce single-cohort stands. Although the existence of multiple stump diameters has been taken as evidence of a multi-aged stand (e.g. Nobori *et al.*, 1997), the data from plantations in Japan demonstrate that single-cohort stands of similar canopy height can have a wide range of trunk diameters (see Table 21.4). We therefore contend that seedling physiology as reflected in germination and light tolerance is consistent with single-cohort stands subject to infrequent major disruptions or ‘patch’-scale disturbances. However, seasonal flooding would produce new areas which could be rapidly colonized. This is suggestive of an evolutionary trend toward an invasive habit, exploiting riparian habitats where competition may have been lower in the past.

The thermotolerance of CO₂ uptake in *M. glyptostroboides* is fairly broad (see Figure 21.1A), and is consistent with the patterns of growth seen in planted specimens throughout the world (Kuser, 1982). This range of temperature is also consistent with the geographical extent of the fossil *Metasequoia* and the presumed range of palaeoclimate. This suggests that, at least for temperature, *M. glyptostroboides* is an adequate NLR. However, it also indicates that *M. glyptostroboides* is of limited use in predicting palaeoclimate. Certainly, it would be expected to perform poorly in subtropical and tropical environments. Water availability appears to be the primary determinant of habitat for this species (Kuser, 1982). In warm-temperate regions with adequate water supply, the tree performs remarkably well; for example, the tree on the NCSU campus (MAT = 15.4°C) is increasing in diameter by about 1.2 cm year⁻¹ and the average rate at the Tanashi plantation (MAT = 13.7°C) is 5.8 mm year⁻¹ (C.J. Williams, unpublished data). Presumably, carbon gain during the cooler spring and fall months offsets the losses associated with the peak summer temperatures in these locations. In contrast, the Napartulik trees grew only about 1.38 ± 0.52 mm year⁻¹ (C.J. Williams, unpublished data). Since water was unlikely to be limiting in the riparian environment at Napartulik, it seems reasonable to assume that the slow growth rates were a consequence of low temperatures.

The photosynthetic apparatus of *M. glyptostroboides* saturates at a relatively low level of light, approximately 1700 μE (see Figure 21.1B). Indeed, chlorophyll content declines with increasing light intensity (Ida, 1981b) and the pigment rhodoxanthin increases (Ida, 1981a; Czczuga, 1987a), leading to a reddish cast in the upper leaves of the trees (personal

observation). This pigment is normally an autumnal coloration (Ida, 1981a; Czezcuga, 1987b); it probably serves in a protective capacity during the summer, shading the chloroplast pigments. The low light saturation level is consistent with the evolution of the photosynthetic system under the reduced light levels above the Arctic circle; full-sun measurements in the Arctic are relatively low (see Table 21.3), but exceed the saturation value. A light saturation value greater than that actually experienced by the plant would represent an investment in unused photosynthetic materials and would likely be selected against as an unnecessary energetic cost. The performance under low light conditions is intriguing; it is conceivable that *Metasequoia* evolved in a foggy riparian environment. However, the high shoot convexity value (θ) is seemingly inconsistent with this scenario, as lower convexity values are thought to be related to more efficient absorbance of diffuse light (Leverenz, 1987) within the conifer shoot. The planar nature of the *Metasequoia* leaf, compared with the needle display of conifers with low convexity values, undoubtedly explains the θ value. Interestingly, the fact that the light saturation level is low, in spite of the modern species' distribution at low latitudes, suggests that the species has not experienced a strong selective pressure to adapt physiologically to alternate light regimes and may not have evolved much in other aspects either. It is worth noting that, in high light environments and at the top of trees, the leaf morphology changes, with the leaflets becoming substantially narrower (D.R. Vann, personal observation; see also Chapter 20). This response appears to be quite plastic in the modern species. Although there is an extensive, well-preserved litter layer at the Napartulik site, the fossil leaves seldom show such a narrow leaflet morphology (B. LePage, personal observation). This implies low light levels in the Eocene Arctic and that the modern species' primary response to light intensities at low latitudes is morphological, rather than physiological. Specific leaf area varies such that mass-based photosynthetic rates are similar in sun and shade leaves (see Chapter 20).

Alternately, the roughly three-fold increase in CO₂ uptake accompanying a three-fold increase in the CO₂ concentration of the cuvette input air stream (see Figure 21.1C) suggests that the plateau in the light response curve is not primarily a limitation in the carboxylation capacity of the cell. Instead, it implies a diffusional limitation. If this is true, it suggests that stomatal density may limit carboxylation efficiency (on a quantum basis). This may have arisen as a compromise between water loss and carbon uptake. Xie and co-workers reported that *M. glyptostroboides* has low drought tolerance and poor xylem hydraulic conductivity (Xie *et al.*, 1999a,b). Further experimentation is required to resolve this question. Presumably, with an adequate CO₂ supply, the light saturation value would increase, indicating that the photosystem's capacity is not saturated at current ambient CO₂ levels.

Plants that shed their leaves simultaneously on an annual basis are considered 'deciduous'; generally deciduous species are responding to a seasonal stress such as cold or drought. As the late Tertiary and much of the Quaternary are considered to have had relatively mild climates, without the large temperature fluctuations seen today at the higher latitudes, cold seems an unlikely candidate as the driving force behind deciduousness in *Metasequoia*, particularly as its evergreen relatives all tolerate frost (Sakai and Okada, 1971; Sakai, 1983). At higher altitudes and in cold lowland regions, evergreen taxa survive the winter in a state of suspended physiology. It has been argued that warmer temperatures could lead to substantial respiratory losses in the winter (Axelrod, 1984). If these losses cannot be replaced, as would occur during the polar night, the plants might not survive. This has not been explored extensively, although the results of Read and Francis (1992) were consistent with this idea.

Metasequoia apparently evolved near the Arctic circle during a period when lowland winter temperatures probably seldom went below freezing (Yang and Jin, 2000). It is therefore reasonable to ask whether the polar light seasonality could have been a driving force for the evolution of deciduousness in this species. To test this, we used the foliar biomass estimates derived for the Napartulik site (see Table 21.4) and the gas-exchange data to derive a coarse carbon budget for the canopy under various temperature scenarios. An 'average' tree at the Napartulik site was calculated to hold approximately 6580 g of foliage (see Table 21.4). Using an average foliage density of $60 \text{ g m}^{-2} \text{ dw}$ (D.R. Vann, unpublished data; see also Chapter 20 who provide a range from 30 to $90 \text{ g m}^{-2} \text{ dw}$), there is some 108 m^2 in the canopy foliage. Based on the photosynthetic rates determined above and using 3780 total hours of daylight (at 65°N latitude, for days with daylength longer than 5 h), a canopy of this size can fix a maximum of perhaps $8.8 \text{ kmol CO}_2 \text{ year}^{-1}$. This figure assumes that the entire canopy is sunlit and fully functional. The actual value is likely considerably lower, as we do not consider a number of factors, such as the loss of photosynthetic capacity during senescence or the effect of rain or overcast weather. Carbon content of foliage is about 49%, so the amount of carbon in the foliage is 3.2 kg. Construction cost is some 30–50% (e.g. Baruch and Goldstein, 1999), therefore, the cost of rebuilding the foliage annually is around 0.4 kmol CO_2 .

Based on the values given in the Results section, maintenance respiration rates for the leaf during mean growing season temperatures of 10, 15, 20 and 25°C would be 0.42, 0.51, 0.84 and 1.1 kmol CO_2 respectively. Dark season respiration at temperatures of 2, 5 and 10°C would amount to 0.11, 0.29 and 0.55 kmol CO_2 respectively. Based on these estimates, and using 0.4 kmol CO_2 as the canopy replacement cost, the 'break-even' temperature is somewhere around 7°C . This figure is a coarse estimate. It is reasonable, however, to expect that temperatures between 5 and 10°C would favour a loss of foliage during the dark months. This also assumes that the reserves are present in the plant. The total amount of carbohydrate required to support this respiration rate amounts to some 12% of the dry weight of the leaf, an unrealistic figure. Storage in other parts of the plant, such as the twigs, is possible, but translocation in the dark, without transpiration to assist, would require an additional energy expenditure. In any case, this analysis is also sensitive to latitude; during the portion of the year with 5 or more hours of daylight, total dark hours decrease with increasing latitudes; these hours accrue prior to and after the polar summer of continuous light. At the latitude where the sum of spring and autumn dark and light hours are equal, the total respiration requirements decline to the point where the foliage could presumably survive a warmer winter (about 8°C). This is an important result for evaluating the evolution of deciduousness as a physiological trait, as it constrains the environment under which this trait emerged. Winter temperatures below 5°C would appear to allow the retention of an evergreen habit, whereas warmer temperatures accompanied by prolonged dark periods represent a selection pressure towards deciduousness. This analysis does not examine whole-tree carbon balance; it is likely that competition for photosynthate from cambial respiration, below-ground growth and reserves for spring budbreak would reduce the balance to favour lower temperatures.

The logic of this result can be generalized to other species. It is consistent with the hypothesis that deciduous plants could have greater success at surviving and/or dominating in warm, dark winters. We cannot establish that dark winters initiated the deciduous trait; it could have existed prior to the migration of these species above the Arctic circle, where it provided a competitive edge over evergreen types. Applying the details of the rather naive analysis above to other species would not be appropriate. There is a large variation in the relationship

between specific leaf area and photosynthesis, resulting in a spread of 'break-even' temperatures. Nonetheless, the general approach could work for other species. However, in all cases, the metabolic rates are so low below about 2°C, that it appears most species could survive, so cold, dark winters would not appear to be an important selective force. This is interesting in light of the fact that the majority of modern clades, in which deciduousness is present, evolved prior to the global cooling trends producing the modern temperate world.

As intimated above, it is possible that *Metasequoia* had already evolved the deciduous trait in response to drought (see Chapter 20). In this case, drought deciduousness could have been important in aiding the colonizing of Arctic habitats by *Metasequoia*. Different cues trigger drought-related leaf-fall compared with dark-deciduousness; based on the phenology of *Metasequoia*, it is responding to a seasonal light cue, probably short days. When it experiences drought, the plant generally responds with a loss of green shoots, rather than the normal senescent process. Considering the species' association with hydric environments and its large water storage capacity within the trunk, it seems unlikely that drought drove the development of deciduousness. Conceivably, *Metasequoia*'s ancestor might have become deciduous in response to a seasonally predictable dry period, triggered by a seasonal light cue. The true origin is unresolved at this time, but it has bearing on the other deciduous members of this clade, *Taxodium* and *Glyptostrobus*, as it is likely the trait evolved in the common ancestor. Each of these is strongly associated with wetland habitats and drought sensitivity.

In conclusion, the variables measured to date support the idea that *M. glyptostroboides* is very likely an example of evolutionary stasis and, as such, it is a good NLR. However, its rather broad climatic tolerance means it is not useful in reconstructing past environments, with the probable exception of indicating a moist habitat. In terms of the evolution of physiological parameters, there seems to be little evidence that this has occurred, as the plasticity of the modern plant would allow its success in all known fossil locations. This does not preclude ecotypic variations (*sensu* Fryer and Ledig, 1972) in the history of the genus, however, the modern form is most likely little changed from the dominant fossil form (*M. occidentalis*). The similarity of fossil and modern stand structure as affected by physiological constraints suggests that the past forests functioned in a similar manner and that sunlight duration does not affect this process. Finally, we provide physiological evidence supporting the hypothesis that evolution of deciduousness could have been driven by warm, dark winters at high (although not polar) latitudes.

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Adaptive ancientness of vascular plants to exploitation of low-nutrient substrates – a neobotanical overview

Christopher N Page

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Introduction

Several living families of ferns have taxonomic fossil lineages which date back at least into the Lower Mesozoic (e.g. Osmundaceae, Dicksoniaceae, Matoniaceae, Dipteridaceae) or even into the Palaeozoic (Marattiaceae, Schizaeceae) (Cleal, 1993; Collinson, 1996; Rothwell, 1996a,b; van Konijnenburg-van Cittert, 1999). Various conifer groups (Rothwell, 1982; Page, 1990) as well as clubmosses of the living families Lycopodiaceae and Selaginellaceae (Thomas, 1992) also date back to the Palaeozoic, as do the Equisetaceae, whose sole living genus *Equisetum* has been suggested to have even changed little since that time (Page, 1972a). Indeed, some whole extant fern floras are ancient: that

of the Canary Islands is an intact fern flora of Miocene age (Page, 1973), that of southern Siberia, though modified by later glacial influences, contains elements which appear to originate from the Cretaceous (Gureyeva, 2001). Among living pteridophytes, ancient elements as well as modern elements often coexist (e.g. Stewart and Rothwell, 1993). In the 1970s, I had the honour (so far as I can work out) of first appropriating the word 'diversity' into biology (Page, 1979a). Similarly, for convenience of reference, a new single phrase is needed for all of these ancient living vascular plants (from pteridophytes to conifers) – as there is no comprehensive taxonomic term. These are here collectively christened Ancient Living Vascular Plants or ALVPs for short (the 'OAPs' of the plant world).

The operation and dynamics of ecological strategies by which these ALVPs have succeeded in achieving their long-term survival, as well as widespread adaptive diversity, have been identified using evidence gained from the living members of these ancient groups, combining observations from field, whole-plant experimental cultural and laboratory approaches (Page, 1979a,b, 1990, 1997b, 2002a and b). This chapter outlines fundamental but little-researched aspects of the edaphic adaptations of ALVP clubmosses, horsetails, ferns and conifers, concentrating on the potential of many, on a global basis, for colonization of substrates characterized by usually direct mineral surfaces of permanently low-nutrient availability and often also of acute nutrient disequilibrium.

These adaptations enable many ALVPs today to continue to exist in sites in which they show exceptional abilities at 'doing without', reflecting extremely efficient standards of nutrient management. In consequence, these abilities enable these ALVPs to occupy diverse edaphically marginal habitats for which today most other modern plant competition is necessarily low. They effectively avoid encountering the many more vigorous competitors that would be more dangerously confronted within almost all more nutrient-rich, more mesic habitats, and the presence to this day of so many ALVPs in such low-nutrient sites remains a stringent test of the efficacy of this strategy for long-term survival.

The modern array of low-nutrient habitats

The term 'low-nutrient habitats' here encompasses all habitats which are, by their very nature, far from being nutrient-rich. Many, but not all, are the direct results of the dynamics of Earth Science processes and involve exposures of largely mineral terrains, with little previous amelioration by climatic or biological agents. Others are added to by biological activity, such as nutrient-poor peat surfaces and bark-surface epiphytic habitats. Among the total of nutrient-poor habitats arising, two broad regimes of tolerance types exist widely among ALVPs. These are tolerance of:

1. regimes of low-nutrient levels *per se*, in which the great majority of mineral elements are either lacking or in very short supply (e.g. the majority of exposed mineral surfaces, peats and plant bark surfaces)
2. regimes of low-nutrient levels of essential mineral elements, but in which non-essential mineral elements may additionally occur in excess, some of which may be regarded as generally toxic (e.g. mineral surfaces containing heavy metal loadings).

Within each of these regimes, field evidence suggests that there are many variations of mineral availability, combination and quantity, which overall determine the habitats to which suites of ALVP species are differentially able to tolerate, while availability of moisture

regimes and habitat accessibility to propagules (e.g. epiphytic sites) further prescribes the groups of plants differentially exploiting them.

The suites of habitats of low-nutrient levels *per se* are especially widespread. Beside the epiphytic habitats (noted above), they include such sites as lowland sand-plains and inundated sand, clay and gravel stream-sides and basins, intrinsically nutrient-poor sandy heathlands and savannahs, and leached soils such as laterites. Some of these habitats are ones also associated with wildfire regimes. In upland areas, widely occurring sites include a great variety of direct erosion-surface mineral exposures of most rocks of all types in virtually all climates, forming cliffs, scree, slump, slide and landslip surfaces, newly formed volcanic outpourings, ash and ejecta surfaces and, additionally, in the wet tropics, especially high mountain rocky summits, ridges and saddles, where gleyed clay ridges result. Ecologically, many of these seem remarkably little studied. For example, mountain summit ridges and saddles form habitats which contain (typically dominantly, Page, 1979a) special suites of ALVPs suggesting these to be among the most extreme in mineral deficiency anywhere.

The suites of habitats in which low-nutrient levels of essential mineral elements combine with toxic non-essential mineral element availability (mineral overload habitats) require further physiological abilities for toleration of such excesses. Such habitats occur where potentially toxic rock types of high mineral-yielding ability occur (and especially where metalliferous elements are yielded). In wild sites at all altitudes, these can include cliffs, major landslip surfaces, landslide debris fans, scree slopes and appropriate post-volcanic disturbance sites. Particularly toxic are the minerals of naturally surface-exposed ultramafic peridotite (volcanic) and serpentine (metamorphic) rock types, in which manganese, cobalt, nickel and chromium typically occur and whose mineral excesses of these elements often combine with simultaneous deficiencies of calcium. Other less toxic habitats, but in which mineral excesses play a particularly species-selective role, are those where calcium or magnesium are particularly high, such as in carboniferous or dolomitic limestones respectively. Man has locally added greatly to the surface occurrence of some of these in the form of discarded minespoil tailings, particularly where workable metal-containing ores occur (e.g. zinc, lead, copper, tin, cadmium and strontium), with discarded mine-tailing exposures still with toxic amounts of metals present also often containing arsenic and sulphurous compounds. In all, the exact mineral excesses involved clearly depend on the rock type itself, its chemical constitution, levels of mineral overload and the rate at which such excesses of minerals are yielded under the occurring climatic conditions.

Whether the sites are merely especially low-nutrient *per se*, or are ones of potentially toxic high mineral overload levels, and whether the habitats are wild or man-made, all these sites present surfaces exposed to potential colonization by arriving ALVP propagules, including wind-borne seeds of conifers and, especially, the highly mobile airborne spores of pteridophytes. Such immigrant seeds and spores have the potential to arrive from the earliest stages of new rock-surface exposures following disturbance events (such as landslides, but also smaller disturbance-event mosaics), when exposure surfaces are particularly fresh. These immigrant propagules include chance arrival of 'unadapted' species arrays from potentially long-distance sources (notably so in Pteridophyta), or from other existing 'adapted' species already occurring on other nearby rock exposures. In each, as mineral substrates vary, local selection for particularly closely-adapted populations undoubtedly applies critical selection pressures, and close experimental investigation of this across an array of ALVPs and their habitats is much needed.

The modern diversity of ALVP colonists of low-nutrient habitats

Taxonomic distribution

Wide arrays of ALVP species from diverse families are today specialist colonists of many low-nutrient sites under both temperate and tropical conditions. There can be considerable specializations of life-form between different ALVPs exploiting these relatively demanding situations and often different taxonomic groups target specific suites of habitats.

On the basis of personal verification in the field (and, for most genera, experience too of experimental cultivation of the same material), the taxonomic spectrum of extant low-nutrient-tolerant ALVPs by genera is presented in Table 22.1.

From Table 22.1 it can be seen that *at least 85 genera of pteridophytes* (19% of 414 total pteridophyte genera recognized by Crabbe *et al.*, 1975) and *at least 19 genera of conifers* (27% of 68 total conifer genera recognized by Page, 1990) are identified here as possessing species which are tolerant of growth on low-nutrient substrates. *This is a total of 104 of 482 genera of extant pteridophytes and conifers together (= 21%)*. In the list, Cycadaceae (s.l.), Gnetales and Ephedraceae (although they are ALVPs) are excluded on the basis of lesser direct personal experience. Were these to be included also, my estimate is that it is likely that all cycad genera with the exceptions of *Bowenia* (tropical Australia) and *Stangeria* (tropical South Africa) would additionally qualify to be included, as would virtually all Ephedraceae. This would enhance still further the trend already identified among pteridophyte and conifer ALVPs for which I have more comprehensive field evidence.

From these wild habitat indicators, if I were to look for the greatest extremist tolerators of low-mineral nutrition availability in this list, I would single out the following: *Lycopodium* (Lycopodiaceae), *Lycopodiella* (Lycopodiaceae), *Huperzia* (Lycopodiaceae), *Equisetum* (Equisetaceae), *Schizaea* (Schizaeaceae), *Gleichenia* (Gleicheniaceae), *Sticherus* (Gleicheniaceae), *Dicranopteris* (Gleicheniaceae), *Platyzoma* (Platyzomataceae), *Stromatopteris* (Stromatopteridaceae), *Lindsaea* (Lindsaeaceae), *Sphenomeris* (Lindsaeaceae), *Schizoloma* (Lindsaeaceae), *Syngamma* (Lindsaeaceae), *Matonia* (Matoniaceae), *Dipteris* (Polypodiaceae), *Widdringtonia* (Cupressaceae), *Cupressus* (Cupressaceae), *Actinostrobus* (Cupressaceae), *Callitris* (Cupressaceae), *Neocallitropsis* (Cupressaceae), *Austrocedrus* (Cupressaceae) and *Pinus* (Pinaceae).

Habitat distribution

Not all ALVPs occur today in low-nutrient habitats, for many have adopted life-forms which enable them to avoid excesses of vegetative competition in other ways. However, the above significant numbers of ALVPs do, and both the taxonomic and habitat diversity of these reflected in the above section, is considerable. In terms of overall habitat trends among ALVPs which do have members which occur in low-nutrient habitats, pteridophytes usually predominate today where such low-nutrient habitats occur in moister climates at all latitudes, but with special diversity in the wet tropics, often reaching their zenith at mid-mountain altitudes (Page, 1979a,b). There are, however, some ALVP fern genera (notably the monotypic *Platyzoma* in Australia and *Stromatopteris* in New Caledonia) which occupy particularly sub-desert terrains. More widely, however, conifers and then cycads occur in progressively drier low-nutrient sites, with conifers in both temperate and tropical conditions (and especially in tropical montane habitats, but also in dry Mediterranean ones) and cycads in mainly dry, high-light tropical ones.

Epiphytic sites are chiefly dominated, among ALVPs, by many ferns and a few club-mosses, which gain rapid access through the high mobility of their airborne spores. All (and

Table 22.1 The taxonomic spectrum of ancient living vascular plant (ALVP) genera each with at least some species able to occupy low-nutrient sites

| | |
|---|---------------------------------------|
| Clubmosses | Ferns (Continued) |
| <i>Selaginella</i> (Selaginellaceae) | <i>Asplenium</i> (Aspleniaceae) |
| <i>Huperzia</i> (Lycopodiaceae) | <i>Pleurosorus</i> (Aspleniaceae) |
| <i>Lycopodium</i> (Lycopodiaceae)* | <i>Matonia</i> (Matoniaceae)** |
| <i>Lycopodiella</i> (Lycopodiaceae) | <i>Phanerosorus</i> (Matoniaceae) |
| <i>Phylloglossum</i> (Lycopodiaceae) | <i>Dipteris</i> (Polypodiaceae)** |
| Horsetails | <i>Cheiropleuria</i> (Polypodiaceae) |
| <i>Equisetum</i> (Equisetaceae) | <i>Platynerium</i> (Polypodiaceae)* |
| Ferns | <i>Polypodium</i> (Polypodiaceae)* |
| <i>Ophioglossum</i> (Ophioglossaceae)* | <i>Goniophlebium</i> (Polypodiaceae)* |
| <i>Osmunda</i> (Osmundaceae) | <i>Dictymia</i> (Polypodiaceae)* |
| <i>Todea</i> (Osmundaceae) | <i>Syngamma</i> (Polypodiaceae)* |
| <i>Leptopteris</i> (Osmundaceae) | <i>Pleopeltis</i> (Polypodiaceae)* |
| <i>Schizaea</i> (Schizaeaceae) | <i>Microgramma</i> (Polypodiaceae)* |
| <i>Lygodium</i> (Schizaeaceae) | <i>Phlebodium</i> (Polypodiaceae)* |
| <i>Gleichenia</i> (Gleicheniaceae)** | <i>Lemmaphyllum</i> (Polypodiaceae)* |
| <i>Sticherus</i> (Gleicheniaceae) | <i>Belvisia</i> (Polypodiaceae)* |
| <i>Dicranopteris</i> (Gleicheniaceae)** | <i>Pyrrhosia</i> (Polypodiaceae)* |
| <i>Platyzoma</i> (Platyzomataceae) | <i>Microsorium</i> (Polypodiaceae)* |
| <i>Stromatopteris</i> (Stromatopteridaceae) | <i>Leptochilus</i> (Polypodiaceae)* |
| <i>Hymenophyllum</i> (Hymenophyllaceae)* | <i>Colysis</i> (Polypodiaceae)* |
| <i>Trichomanes</i> (Hymenophyllaceae)* | <i>Aglaomorpha</i> (Polypodiaceae)* |
| <i>Didymoglossum</i> (Hymenophyllaceae)* | <i>Merinthosorus</i> (Polypodiaceae)* |
| <i>Culcita</i> (Dicksoniaceae) | <i>Drynaria</i> (Polypodiaceae)* |
| <i>Dennstaedtia</i> (Dennstaedtiaceae) | <i>Lecanopteris</i> (Polypodiaceae)* |
| <i>Microlepia</i> (Dennstaedtiaceae) | <i>Crypsinus</i> (Polypodiaceae)* |
| <i>Lindsaea</i> (Lindsaeaceae) | <i>Selliguea</i> (Polypodiaceae)* |
| <i>Sphenomeris</i> (Lindsaeaceae) | <i>Grammitis</i> (Grammitidaceae)* |
| <i>Schizoloma</i> (Lindsaeaceae) | <i>Antrophyum</i> (Vittariaceae) |
| <i>Syngamma</i> (Lindsaeaceae) | <i>Vittaria</i> (Vittariaceae)* |
| <i>Hypolepis</i> (Hypolepidaceae) | Conifers |
| <i>Pteridium</i> (Hypolepidaceae) | <i>Araucaria</i> (Araucariaceae) |
| <i>Histiopteris</i> (Pteridaceae) | <i>Pinus</i> (Pinaceae) |
| <i>Pellaea</i> (Pteridaceae) | <i>Larix</i> (Pinaceae) |
| <i>Cheilanthes</i> (Sinopteridaceae) | <i>Actinostrobus</i> (Cupressaceae) |
| <i>Actiniopteris</i> (Cryptogrammeaceae) | <i>Cupressus</i> (Cupressaceae) |
| <i>Cryptogramma</i> (Cryptogrammeaceae) | <i>Austrocedrus</i> (Cupressaceae) |
| <i>Onychium</i> (Cryptogrammeaceae) | <i>Diselma</i> (Cupressaceae) |
| <i>Hemionitis</i> (Gymnogrammeaceae) | <i>Callitris</i> (Cupressaceae) |
| <i>Pityrogramma</i> (Gymnogrammeaceae) | <i>Juniperus</i> (Cupressaceae) |
| <i>Davallia</i> (Davalliaceae)* | <i>Microbiota</i> (Cupressaceae) |
| <i>Scyphularia</i> (Davalliaceae)* | <i>Neocallitropsis</i> (Cupressaceae) |
| <i>Humata</i> (Davalliaceae)* | <i>Tetraclinis</i> (Cupressaceae) |
| <i>Oleandra</i> (Oleandraceae)* | <i>Dacrydium</i> (Podocarpaceae) |
| <i>Nephrolepis</i> (Oleandraceae)* | <i>Manoao</i> (Podocarpaceae) |
| <i>Arthropteris</i> (Oleandraceae)* | <i>Lepidothamnus</i> (Podocarpaceae) |
| <i>Rumohra</i> (Aspidiaceae)* | <i>Microcachrys</i> (Podocarpaceae) |
| <i>Elaphoglossum</i> (Aspidiaceae)* | <i>Microstrobos</i> (Podocarpaceae) |
| <i>Doodia</i> (Blechnaceae) | <i>Taxus</i> (Taxaceae) |
| <i>Blechnum</i> (Blechnaceae) | <i>Pseudotaxus</i> (Taxaceae) |

Those genera marked with an asterisk (*) include members with notable epiphytic tolerances. Those marked with paired asterisks (**) include extreme terrestrial specialists of tropical high mountain summit and ridge environments.

especially ferns) predominate under conditions of frequent rainfall and diminish in species numbers in progressively drier epiphytic sites. The genera containing epiphytic species of clubmosses or ferns are those most likely to include the widest arrays of individual taxa most tolerant of overall low mineral levels, but not necessarily tolerant of high availability of other mineral elements. In the wild, subtle differences among tolerances of epiphytic taxa, especially in the tropics, are reflected in the tendency for different pteridophyte epiphytes either to be generalist or more narrowly specialist species as far as preferred host-tree habitat selection is concerned and complex arboreal species-habitat niche mosaics exist (Gardette, 1996; Page, 2002b). Some epiphytic genera additionally show unusual, sometimes bizarre, structural adaptations to modest natural enhancement of arboreal mineral scavenging, such as production of morphological adaptations to intercept and subsequently compost falling leaves (e.g. *Platyserium*, some *Asplenium*), or to form associations with animals and receive detritus (notably from ants, for which specialist myrmecophilous fern genera such as *Lecanopteris* (extant Old World tropics) and *Solanopteris* (extant New World tropics) represent extreme adaptations).

In exposed rocky habitats, many ALVPs exist. These include especially ferns and conifers, each with suites of unrelated genera adapted to tolerance not only of the rock types outcropping, but also of the physical constraints of these habitats which include exposure to frequent and irregular droughts. Within both ferns and conifers, further taxonomic specialists occur with adaptations to rock types in which other great mineral excesses additionally occur. These are usually adaptee species from within existing rock-inhabiting specialist genera and related to ones tolerant of other mineral-poor habitats but with less toxic mineral overloads. Unusual rock types, including metal-loaded ones, such as ultramafic rock sites, provide habitats for specialist species of many ALVP genera either of pteridophytes or conifers (e.g. Wherry, 1920; Ishizuka, 1961; Krukeberg, 1964; Yoshioka, 1974; Jaffre, 1980, 1995; Sleep, 1985; Jaffre *et al.*, 1987; Jaffre and Veillon, 1990; Page and Hollands, 1987; Page, 1999 and in press a,b). These extraordinary nutrient-poor mineral disequilibrium sites clearly require unusual degrees of tolerance which can be highly taxon-specific.

In free-draining almost purely sandy soils, sometimes with acidic peat horizons, as well as in essentially rocky habitats, conifers often predominate. Such sandy soils today mostly carry members of the families Pinaceae and Cupressaceae in the northern hemisphere and Cupressaceae and Podocarpaceae in the southern hemisphere, sometimes with specialist Schizaceae and Lycopodiaceae. Additionally, dry rocky habitats globally are occupied extensively by members of virtually all conifer families including Araucariaceae, and there may be also a number of specialist ferns. One of the most extensive of such areas of such habitats is across the south-eastern and eastern lowlands of the USA, where substantial pine-dominated savannahs occur. Here, soils and rocky substrates, which are acid and low in nutrients (Little, 1979), are especially widespread. Extensive areas are dominated by Loblolly Pine (*Pinus taeda* L.) and Shortleaf Pine (*P. echinata* Mill.), while some of the most apparently nutrient-poor extremes I have seen anywhere are the habitats of Sand Pine (*Pinus clausa* Vasey) in coastal white sands of the Florida panhandle and of Hard Pine (*P. rigida* Mill.) dominating the 'Pine Barrens' of the New Jersey coastal plain.

Fireburn habitats, often themselves occurring in already low-nutrient terrains which are also dry, such as sand plains and rocky surfaces (above), are sites to which many conifers and various pteridophytes characteristically contain locally-adapted species (often evolved from other non fire-adapted local species groups). Among conifers, these especially include *Pinus* (with other genera occurring more locally) and *Araucaria* and *Callitris* (in the southern hemisphere). Among pteridophytes there are a number of genera recorded in fire-climax

vegetation, including, in sub-Saharan Africa especially, species of *Actiniopteris*, *Adiantum*, *Anemia*, *Aspidotis*, *Cheilanthes*, *Mohria*, *Pellaea* and *Lycopodium* (Kornas, 1978) and in Queensland especially species of *Adiantum*, *Blechnum*, *Doodia*, *Cheilanthes*, *Cyclosorus*, *Dicranopteris*, *Gleichenia*, *Lastreopsis* and *Schizaea* (Page, personal observations). Globally *Pteridium* occurs especially widely in such fireburn habitats and, unusually, has the ability to pioneer ash-burn surfaces of high potash levels and highly alkaline pH, metamorphosing to become an acidophilous plant once its rhizomes penetrate beyond the initial ash surface layers (Page, 1986).

Low-nutrient poor clay and gravel sites, especially in temperate zones, often typical of areas of glacial retreat, can form habitats for ALVP clubmosses (especially where eventually overlain by acidic peats) or for horsetails (usually on direct mineral surface exposures). Today the former group often extends in such sites abundantly to nutrient-poor mountain summit habitats, both in high temperate latitudes and on mountain ranges in the tropics, such as those of the Andes (Olgaard, personal communication), while ALVP horsetails are most abundant in such sites in lowland areas, where they can be especially rapid colonizers of such newly-exposed mineral surfaces (Page, 1967). For horsetails, the mineral significance of such nutrient-poor sites is further complicated by the over-riding requirements of all members of this plant group for sites which also contain both adequate available bases (usually calcium) as well as abundant available silica – an element integral in the structuring of all aerial parts (Page, 1972b). No horsetails succeed (nor probably ever have) as true epiphytes (though they might theoretically have existed as scandent climbers), since the necessity for this unusual mineral combination clearly requires maintenance of permanent root contact with the ground.

On directly nutrient poor edaphic sites characterized by erosion-regimes and constant surface moisture downwash such as today on high-mountain saddles of tropical mountains, pteridophytes, but not usually conifers, gain the necessary up-slope access and so these nutrient-poor habitats are typically dominated by ALVP ferns. Of these, unrelated plants of surprisingly large size and often surface-creeping rhizomes with notably sprawling and often rampant frond habits are especially characteristic, with tough growth structure forming ridge-top fern thickets typically yielding much outwash debris. Characteristic genera are the ALVP ferns *Gleichenia*, *Dicranopteris*, *Dipteris*, *Matonia* and *Christensenia*. Species of *Lycopodium*, especially *L. cernuum*, are occasionally present.

Surfaces of volcanic rocks have their own, usually peculiar, suites of colonizing ALVPs, especially ferns and sometimes horsetails. To what degree volcanic lava, ash and other ejecta surfaces represent low-nutrient ones clearly depends on many factors, since such volcanic outpourings have the repute of often eventually giving rise to highly fertile soils. Yet it is, however, at the earliest stages of post-cessation activity by the volcanic source itself that diverse ALVP components today, most especially, achieve pioneering vegetational roles in the arising low-competition habitats. At these stages, the substrates have weathered relatively little, and most nutrients would appear to be highly unavailable. Examples of such pioneer ALVPs thus occur in many geographically-scattered volcanogenic habitats. These can include both conifers and pteridophytes, the proportions dictated by colonization intervals and, especially, by geographic location of continentality versus oceanicity. For example, I have recorded the conifers *Thujopsis dolabrata* Sieb. & Zucc. and *Picea polita* Carr. colonizing such sites in Honshu, Japan, where these trees eventually re-form substantial forests. But where volcanogenic disturbances occur far from such seed sources, and especially so on oceanic islands, then promoted by the high access-mobility of airborne spores, contingents of unrelated pteridophyte genera are usually the colonists. Fern genera I have seen so-behaving

in the field (Canary Islands, Mauritius, Hokkaido, Yakushima, Philippines, New Zealand, Western Samoa, Hawaii, Cascades, Chile) are: *Arthropteris*, *Asplenium*, *Blechnum*, *Cheilanthes*, *Cibotium*, *Davallia*, *Dryopteris*, *Elaphoglossum*, *Humata*, *Hypolepis*, *Lygodium*, *Microgramma*, *Microsorium*, *Nephrolepis*, *Notholaena*, *Odontoloma*, *Onychium*, *Pellaea*, *Phymatodes*, *Phymatosorus*, *Pityrogramma*, *Platycerium*, *Polypodium*, *Psilotum*, *Pteridium*, *Pteris*, *Pyrrosia*, *Sadleria*, *Sphenomeris* and the fern allies *Selaginella*, *Lycopodium* and *Equisetum* (Page, 1979a *et seq.*, but only those with multiple or extensive occurrences are included within the main list here). Further, *Dicranopteris*, *Gleichenia* and *Sticherus* usually occur on more degraded lavas and some of these and other genera (e.g. *Christella*, *Cyclosorus*, *Doodia*, *Histiopteris*, *Hypolepis*) colonize and may persist in the inclement (poor and often sulphur-laden) terrain around hot-spring sites (e.g. Parris, 1976). Similar observations recording ferns to be early colonizers of lava flows and other volcanogenic surfaces have been made by a number of authors in other localities (e.g. Gates, 1914 (Philippines); McCaugherty, 1917 (Hawaii); Hartt and Neal, 1940 (Hawaii); Beard, 1945 (W. Indies); Crookes, 1960 (Rangitoto); Fosberg, 1967 (Hawaii and Galapagos); Eggler, 1971 (Hawaii); Uhe, 1974a–c (Kermadec Is., Savai'i and Tonga); Yoshioka, 1974 (Miyakejima and Sakurajima Is.); Benl, 1976 (Cameroons); Spicer *et al.*, 1985 (Chiapas, Mexico). Some of these colonists (e.g. *Nephrolepis biserrata* (Sw.) Schott in Western Samoa) I found freely colonizing lava surfaces of as little as 30 years of age, and *Equisetum arvense* L. (Spicer, personal communication) established on bare ash-fall deposits within the Mt St Helens blast zone in a considerably shorter period than this. On the island of Krakatau where the post-eruptive succession has been particularly well documented (Treub, 1888; Ernst, 1908; Turrill, 1935; Docters van Leeuwen, 1936), eleven species of ferns were firmly established on explosion breccia within as little as three years of its August 1883 cataclysmic eruption.

Within each of these general habitat types, further factors then govern the actual species of ALVPs which are present. A very few species of ALVPs, almost exclusively ferns, seem able occasionally to exchange between some (albeit usually a limited array) of these habitats and these usually occur between certain epiphytic and rock-exposure faces of relatively low mineral-yielding rocks. Mostly, however, suites of species are characteristic of one type of site and not found in another. The frequency of occurrence and taxonomic diversity of ALVPs in all of these low-nutrient habitat-roles today, with subtle differences between sometimes related taxa and their adaptations and niches within suites, and major taxonomic, habit and consequent differences in taphronomic potential between suites, almost certainly indicates that there have been similar occupations with similar differences in pteridophytes in many palaeo-settings throughout land-plant evolutionary time.

It is significant that several terrestrial groups of clubmoss and ancient fern species present in the palaeo-record survive today especially as residents of extreme tropical mountain ridge habitats, of which on personal experience, *Gleichenia*, *Dicranopteris*, *Matonia*, *Dipteris* fall closely within this most edaphically exacting modern habitat category.

Responses to cultivation

Experiences of exposure of a great array of globally wild-sourced low-nutrient tolerant ALVPs to experimental cultivation not only endorse that observed ecological abilities have a physiological basis, but that species, though sometimes flexible in response, are more often especially closely attuned to both tolerance and often optimization of growth under regimes of the low levels of mineral availabilities actually occurring in their wild habitats. If artificially exposed to high-nutrient availability, most fail to show significant growth-enhancement

response, failing completely to accommodate to enhanced nutrient availability, even at levels which would be regarded as mesic by general flowering plant standards. Under such conditions, rapid death usually ensues, even when, under experimental conditions, they are protected from the effects of competition for these habitats from other plants more demanding of these conditions. One can only conclude that, in the great majority of cases, their physiology is specifically attuned to high-nutrient deficiency and exposure to anything better presumably causes what I can only describe as 'nutrient shock'. (It would be rather analogous to putting high-octane aviation fuel into an old Landrover in the hope of going faster cross-country, and finding the engine merely explodes as a result.)

In fact in ALVPs, the responses tend to vary subtly (but with apparently substantial ecological consequences) between suites of species (whether closely interrelated or not), but often showing close similarities of response when from similar originating habitats. For example, the genera containing epiphytic species of clubmosses or ferns are those most likely to include individual taxa most tolerant of overall low mineral levels, but not necessarily tolerance of any more than extremely modest levels of further nutritional application. Such plants respond successfully in cultivation to culture usually only on low nutrient substrates and, once established, can be horticulturally nutrient-fed (e.g. N-P-K) only at *extremely modest levels*. Greater feeding concentrations kill them rapidly. In the case of epiphytes, such modest tolerated nutrient levels in experimental cultivation presumably approximate to equivalents of available levels of such substances in the wild, through leachates, through leaf-composting strategies of large 'tank' epiphytes (Page, 1979a,b), and incoming animal debris and droppings, such as those of colonizing ants (e.g. *Azteca* spp., Janzen, 1974).

At an even greater extreme, ground-rooted genera from tropical high mountain-ridge habitats (e.g. the fern genera *Gleichenia*, *Matonia*, *Christensenia*, *Dipteris*, which can be large and deceptively luxuriant plants in the wild), are especially sensitive to virtually any imposed feeding regimes in cultivation, to the extent that it is difficult to devise edaphic conditions which are sufficiently poor on which to grow them (most feed-free composts as well as bracken-peat have been tried as substrates with varying degrees of success and perennation, and even expanded polystyrene has been tried!).

These observations suggest the existence of underlying physiological mechanisms in many ALVPs (and especially pteridophytes) of appropriate habitats which are:

1. exceptionally efficient in nutrient management by general flowering plant standards, and which
2. become effectively 'saturated' at quite low, sometimes extremely low, levels of nutrient availability.

Further, the additional habitat category of low-nutrient tolerant plants from toxic mineral rock sites clearly must have other physiological mechanisms for tolerance of the excesses of toxic mineral overload. Cultural experience of the relatively few ALVPs from these sources which have been tried, has so-far produced varied results with different genera, suggesting that here wide differences in potentials and responses may exist. For example, some temperate conifer species exclusively from wild ultramafic soils prove difficult if not impossible to grow in cultivation. These include *Pinus sabiniana* Dougl. which I seed-sampled from ultramafic rocks in California (when samples of nearby non-ultramafic *Pinus coulteri* D. Don. from the same climate grew successfully) and *Picea glehnii* Voss. which I seed-sampled from ultramafic rocks in Hokkaido (when nearby non-ultramafic *Picea jezoensis* Carr. seed-sampled from non-ultramafic rocks in the same climate grew vigorously). Further,

even within-species there appear to be populational contrasts: preliminary experiments showed that *Asplenium adiantum-nigrum* L. ferns spore-sampled from ultramafic (serpentine) sources on the Lizard, Cornwall, grew well in cultivation on their native rocks but less successfully on non-ultramafic rocks, while cultures from parental populations on nearby non-ultramafic (shale) rock sources grew well on their own rock types but not at all on the ultramafic ones. These data help to explain the exclusive confinement of these ALVPs to their respective rock types in the wild, and the evolution of obligate physiological adaptations seem indicated in these cases.

However, by contrast, several ALVP genera of tropical conifers (e.g. *Neocallitropsis araucarioides* (R.H. Compton) Florin, *Dacrydium araucarioides* Brongn. & Gris, *Dacrydium guillauminii* Buchholz, *Retrophyllum minor* (Carr.) C.N. Page, *Agathis ovata* Warburg) which I seed-sampled from wild trees growing on peridotite rock exposures in New Caledonia (Page, 1999, in press b), proved to grow successfully in cultivation on regular horticultural compost substrates, where they also tended to show far more rapid rates of growth than in the wild. Here, at least for these plants, the evolution of physiological adaptations to these relatively exacting terrains would seem to be facultative. However, by contrast, the ALVP fern *Stromatopteris*, growing in association with *Dacrydium araucarioides*, failed to respond at all to attempts to grow it in cultivation. This seems to suggest that different evolutionary pathways of evolution of low-mineral nutrition levels and tolerance of other elements in excess have been involved, even between different ALVP members co-associated today in the same habitat!

Discussion

The success and diversity which pteridophytes and conifers have evolutionarily achieved in all habitats through time has been especially indicated from a palaeobotanic perspective by Seward (1933), Harris (1976), Miller (1976, 1977, 1982), Edwards (1980), Taylor (1981), Galtier and Scott (1985), Scott and Galtier (1985), Thomas (1985, 1986), Thomas and Spicer (1987), Rothwell (1982, 1987, 1996a,b), Stockey (1989), Collinson (1990, 1996, 2002), Stewart and Rothwell (1993), Taylor and Taylor (1993), Bateman (1994), Galtier and Phillips (1996) and DiMichele and Philips (2002). The surviving diversity of extant pteridophytes, including many aspects of their ecology, has been presented especially by Bower (1923), Verdoorn (1938), Holttum (1938, 1968), Copeland (1907, 1947), Jermy *et al.* (1973), Lovis (1977), Dyer (1979), Page (1979a,b, 1987), Tryon and Tryon (1982), Dyer and Page (1985), Camus *et al.* (1996), and that of conifers especially by Florin (1940, 1963), Page (1990), Enright and Hill (1995), Farjon (1998) and Richardson (1998). Attempts to analyse the ecological strategies by which this long-term survival has been achieved have been made for pteridophytes and conifers from a neobotanical perspective (Page, 2002b and a respectively). From the latter evidence it is clear that the living plants can provide information about the strengths of the abilities of these plants, many unique to them and to their life cycles, life styles and natural history, which would not have been easy to deduce or necessarily even possible to perceive, from examination of fossil material alone. On the other hand, the fossil material provides a factual basis of morphology in time and space and (often) some indications of habitat, against which to compare ideas formulated from extant material. Collinson (1996: 350), appropriately described the role of fossils as 'providing a time dimension for almost all aspects of interest in modern material'. This clearly includes setting a time-frame to the

inferences which can be made from the living material in relation to when and where adaptations demonstrated would have been of ecological benefit in the geological past, and thus, by further implication, when the necessary supporting physiologies to achieve such adaptations would likely have come into play.

Other theoretical aspects also help to establish a logical background for such neobotanical interpretations. For example, the evolution of several aspects of plant architecture relevant to ALVPs has been presented by Kurmann and Hemsley (1999). As a habitat background to ALVP success, vegetational consequences of angiosperm radiation have been presented by Crane (1987) and summarized with respect to fossil ferns by Collinson (1996). Contrasts between gymnosperm and angiosperm neo-ecology have been drawn by Bond (1989), and the importance of considering long time-frames in understanding slowly operating biotic processes in palaeoecology has been stressed by Schoonmaker and Foster (1991).

Against this background, observations on the ecology of the whole of the diversity of living members particularly allows collective inferences to be drawn about the advantages of the whole diversity of the abilities of ALVPs in relation to long-term survival of these ancient plants, the ancientness of the traits indicated and the role which adaptation to these habitats has played in vascular plant evolution. Further, the importance of consideration of habitat in relation to strategies has been stressed by Southwood (1977), albeit for animals, but I adopt strong similarities of approach in the marrying of ALVP neo- and palaeo-ecological botanical data. For example, in probably the great majority of habitats in which ALVPs occur, there are likely to be a constantly reiterating processes of ecological dynamics of colonization and recolonization, producing an overall 'pattern and process' (*sensu* Watt, 1947) to the ALVP community as a whole, with strategic advantages of pioneering among plants able to tolerate the substrates exposed recurrently regained. Linking closely with this, the importance of such disturbance regimes in activating colonization opportunity for pteridophytes (and probably actively important for regeneration of very many ALVPs in the field) has been stressed in relation to the achievement of modern conservation goals for these groups (Page, 2002a).

Advantages of low-nutrient toleration for survival of ALVPs

Tolerance of low-nutrient regimes opens colonization opportunities which are not necessarily available to competing species. Advantages gained are specifically:

1. A time-frame advantage – opportunity for access to habitats in which newly-exposed mineral-poor surfaces become suddenly available
2. A spatial advantage – opportunity for access to habitats which are widely available
3. A strategic biotic advantage – avoidance of vigorously evolving competition for more nutrient-rich habitats.

Clearly, the more generalized the low-nutrient terrain, the more colonists will be able to pioneer it. Experiences in experimental cultivation indicate that for the edaphic low-nutrient generalists, a wide array of opened habitats are accessible for those species which are able to tolerate the ruling nutrient disequilibria. For those of most extreme mineral-loaded habitats, high and exactly-limited tolerances and limitations are involved. Further, such experimental evidence suggests that although, especially on the more extreme terrains, ALVPs with such adaptations are also relatively slow-growing in the wild by most flowering-plant standards, such rates of growth are not necessarily disadvantageous in such

marginal wild low-nutrient habitats in the absence of more vigorous competition (most ALVPs are thus superb exploiters of the 'tortoise and hare' syndrome).

Direct colonization of each of these sites is most likely to follow in the immediate wake of disturbance regimes which have opened temporarily competition-free sites (Page, 2002a). Access factors then often determine to which colonists the habitats are available. A short-lived rocky site, if not moisture-limited, can be rapidly pioneered by pteridophytes because of the high mobility of their airborne spores. It often takes only a little longer for conifers to arrive, depending on the source and type of seed. A poor sandy habitat can be easily available to pteridophyte spores and conifer seeds, given modest time for the latter, though moisture availability may limit the pteridophyte component. Epiphytic sites are an important low-nutrient target for pteridophyte spores but are seldom successfully accessed (except on fallen logs) by conifer seeds, even though they may be well-able to tolerate the habitats. Consequently, pteridophyte epiphytic diversity is abundant (and I suspect has been through the fossil record, wherever potential habitats have been present) but, despite their potentials for low-nutrition tolerance, recurrent poor access is almost certainly the main reason for the absence through evolution of epiphytic herbaceous conifers.

Of further biological significance is that newly-opened habitats of all types, because of their freedom from initial competition, are also sites in which enhanced opportunities exist for simultaneous colonization by randomly-derived individuals growing closely together. This is particularly significant for pteridophytes for which enhanced opportunities for occasional outcrossing across species boundaries can be a consequence. Such a role of newly-opened habitats is undoubtedly significant in hybrid formation in many pteridophytes today (Page, 1967, 1987, 1997a,b, 2000) and has been especially widely recorded in *Equisetum* (e.g. Page and Barker, 1985; Page, 1987). Such hybrids, in modern Pteridophyta, form the building blocks of evolution through allopolyploidy and such habitats have likely been additionally significant in this respect throughout land-plant evolution (Page, 2002b).

Ancientness of low-nutrient toleration

Today, specialist ALVPs able to exploit low-nutrient habitats come from a very wide range of taxonomic groups, the palaeo-elements of which have appeared (where known) at many different times in the fossil record. The adaptations and ecologies which they show today demonstrate the taxonomic diversity and effectiveness with which ALVPs are able to respond to the challenge of occupancy of such sites, and it has been independently suggested in relation to conifers (Lusk and Matus, 2000) that traits other than growth are important on low fertility sites. Mycotrophy is probably an important further element of this whole nutrition issue, which is beyond the immediate scope of this chapter, but which I am aware has itself almost certainly assisted with many of the uptake processes of these groups throughout geological time, and also probably has a history back to the first land colonizers (e.g. Taylor and Taylor, 1993, 1997; Taylor *et al.*, 1995). We need to find out much more about its role in relation to ALVPs and especially in relation to those of the habitats discussed here.

Low-nutrient habitats of the types we see today must have occurred and recurred at least as widely on Earth throughout the geological record, varying mainly only in location, emphasis and extent between different Epochs and Periods, responding to both minor and more major geo-tectonic, orogenic and occasional catastrophic events. (Witness, for example,

the 'fern-spike' immediately following the dynamic events at least of the K/T boundary interpreted as representing the first Tertiary vegetation – e.g. Spicer, 1989: 295; Nichols *et al.*, 1992; Lerbekmo *et al.*, 1999.) Indeed, disturbance of the environment associated with many episodes of extinction in the fossil record (e.g. Boulter, 1997) may have regularly provided extensive exposures of those nutrient, mineral-dominated substrates, in which ferns have subsequently expanded widely. Is it the case that only at these times (and possibly other boundaries) had ALVP (and especially fern) colonists evolved that could widely utilize the arising terrains? Alternatively, although it cannot be ruled out that there have also been other members of the same families and even genera that succeeded in more mesic terrains in the past, evidence from the living ALVPs suggests that low-nutrient toleration is likely to be a strategy which very probably similarly occurred and recurred through appreciable geological timescales, with cohorts of such species similarly existing wherever and whenever opportunity offered with at least as much efficiency and diversity as do their survivors today.

Other palaeo-evidence of colonization of low-nutrient sites is mainly by analogy. Conifers with many members able to colonize a variety of low-nutrient sites today are, as genera, often far more geographically widespread as fossils than they are today (Florin, 1940, 1963). Fireburn sites are known since at least the Early Carboniferous (Scott and Jones, 1994; Falcon-Lang, 1998, 1999), and volcanic ash habitats from at least similar times (e.g. Scott and Galtier, 1985; Brousmitche *et al.*, 1992; Falcon-Lang and Cantrill, 2002). Indeed, that the ancientness of this strategy probably provided opportunism for colonization even among early filicaleans is indicated by Galtier and Phillips (1996). Epiphytism among fossil ferns, although yet very fragmentarily known (e.g. Sanhi, 1931; Rothwell, 1991; Poole and Page, 2000) probably too has a very long fossil history. Some ferns of extremely nutrient-poor white-sand habitats with a long evolutionary history (e.g. *Platyzoma*, Tryon and Vida, 1976) grow in a curious colonial 'fairy-ring' formation today (Page and Clifford, 1981) providing one potential fossil 'signal' of such habitats. Also notable here is that so many terrestrial ferns as far back as the Early Carboniferous have been described, in habit, as 'sprawling' or 'rampant' (e.g. DiMichele and Phillips, 2002) and one cannot help but link this habit today with that so characteristic of several surviving pteridophytes (e.g. *Gleichenia*, *Dicranopteris*, *Dennstaedtia*) of the poorest edaphic terrains, themselves also so often mineral dominated sites associated with disturbance events and, from time to time, fireburn. The 'fern thickets' of Cretaceous Antarctica (Cantrill, 1996) suggest somewhat similar habitats much later in the fossil record and habitats which include such genera as *Blechnum* today, while the 'fern prairies' in the mid-Cretaceous probably occurring over sites of volcanogenic disturbance events (Collinson, 1996, 2002) may also indicate extensive low nutrient habitats at the time.

I have not included within this account the additional array of ALVPs which are largely swamp or open-water species (both pteridophytes and conifers, see Collinson, 2001, 2002 for recent reviews of the occurrence of fossil Cainozoic members of these) since I am not convinced that such swamp habitats are necessarily as completely nutrient-poor to the degree that are the habitats discussed here. Further, the response in cultivation of many of their members to enhanced nutritional levels tends more often to be a positive one, suggesting that different physiological mechanisms characterize these suites of genera and that there are other reasons for the survival of these ALVPs in the habitats which they occupy. In comparison with plants from the habitats discussed, swamp species are relatively well-represented in the fossil record and a separate study of these relating them close to the *ecological* experience of the modern plants would be desirable.

Theoretical perspectives

Throughout the geological record of pteridophytes and conifers, I have the impression of an overall sparsity of evidence about the ecology and precise habitat types of so many of the plants whose morphology is often discussed at length. This may well be partly due to lack of knowledge of such papers on my part, although I find that Collinson and van Konijnenburg-van Cittert (2002) have also drawn attention to seeking more direct evidence of ancient fern ecology as an avenue for future research. However, recent valuable contributions in relation to palaeo-ecology of ferns have been made, for example, by Watson and Alvin (1996), Bateman *et al.* (2000), Collinson (2002), Deng (2002), DiMichele and Phillips (2002) and Van Konijnenburg-van Cittert (2002). Palaeo co-associations between specific conifers types, ferns types and lycopod types may well be overlooked elements in supporting conclusions about nutrient richness or otherwise of habitats here.

Of evidence further back in time, of that known to me, there is some direct evidence of the mineraliferous nature of habitats of at least some early plants such as *Rhynia* (Trewin, 1994; Rice 1995; Powell *et al.*, 2000; Rice *et al.*, 2002; DiMichele and Phillips, 2002). Although it may be merely by chance that in such hot-spring habitats this material became most extensively preserved, this evidence at least indicates the close association of these plants and probably mineral-dominated terrains dating back at least to the early Devonian.

I thus contend that the types of habitat adaptations discussed here among ALVPs are likely to be ancient and probably actually primitive within land plants, and have a history which is likely to be a continuous one back to the ecology of the earliest land colonizers. For at these times the earliest land colonizers must have once faced habitats virtually everywhere which were extensively nutrient-poor but exceptionally mineral-rich terrains. It was the ability to manage efficiently the exploitation of these hostile terrains wherever water was not limiting, that so enabled these colonizers to be so successful in achieving the breakthrough of utilization and pioneering of these early land surfaces.

As all of the above conditions are geologically enduring, they have thus almost certainly operated to the advantage of the survival of ancient forebears of our modern ALVPs through time in exactly the ways in which they did originally, and still do today. Indeed, it is likely that seldom, if ever, on a global scale, have ALVPs needed to face such overall biotic competition pressures as are presented by modern biota, and especially by the flowering plants. The persistence of our ALVPs still today is thus high testament and stringent test of the efficacy and success for survival of ALVPs enduring strategy of low-nutrient tolerance.

Plants of these habitats, perhaps partly because they are likely to be always under-represented throughout the fossil record, tend to be little-appreciated components of land plant evolutionary success. The extant ALVPs of these habitats may well, however, often represent most closely the ecological cores from which others, and especially many of those of mesic habitats, have tended to evolve. There has probably always been a slow, sometimes very slow, rate of evolutionary turnover among extremist species at the periphery of nutrient availability, while, by contrast, higher, and often much higher, evolutionary turnover rates have probably always tended to dominate the mesic ones.

The displays of slow growth and considerable tolerances of exacting and quite extreme edaphic conditions which many ALVPs continue to show today nevertheless echo the existence of significant evolutionary niches through time, their basic parameters of which must have changed remarkably little through long time-spans. I suggest it is rarely, in the course of evolution, that diversification 'throws-up' new plants better able to exploit these more marginal conditions than do the ancient ones already holding them.

Conclusions

The wide taxonomic range of ALVPs exploiting today's nutrient-poor terrains reflects a wide distribution of the underlying physiological abilities, suggesting that they are unlikely to be new, but are ones which these plant groups have had the potentials to call upon through their past history. In the past, a wide variety of low-nutrient habitats of direct mineral exposure will have also repeatedly existed and continuously recurred on land throughout the geological record, as an immediate result of the dynamism of Earth Science processes. At various times, some of these will have been more extensive than at others. Adaptations similar to those seen today among ALVPs, must have occurred through much of land-plant evolutionary time and that occupation of such marginal low-competition habitats has often led to long-term survival among the taxa able to do so. The enabling physiologies are probably themselves ancient and may well be more primitive within land plants than are necessarily those of the occupants of more mesic habitats today, and reflect more closely those of the earliest land colonists.

Such habitats and their surviving ALVP occupants, may thus well actually represent, in their exacting adaptations, our nearest living equivalents to the adaptations of even the earliest vascular land colonizers. These, I suspect, were not only equally efficient in nutrient management of their Spartan resources, but as a result, like many extant ALVPs, I strongly suspect, may have grown large, were individually often long-lived and usually did everything slowly!

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23

The evolution of aluminium accumulation in angiosperms

Steven Jansen, Toshihiro Watanabe, Steven Dessein,
Elmar Robbrecht and Erik Smets

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Aluminium in the environment and its toxicity for plants

Aluminium (Al) is ubiquitous in our environment, as it is the most abundant metal and the third most common element in the earth's crust. Rocks and soils consist of primary and secondary aluminosilicate minerals. Although it occurs as a constituent element in the silicon tetrahedra and aluminium octahedra in all soils and as oxides and hydroxides in highly weathered tropical soils, its solubility in soil solutions is very limited and varies according to soil pH. The naturally occurring Al forms are usually non-toxic and stable, but soluble Al ions (Al^{3+} , $\text{Al}(\text{OH})^{2+}$, and $\text{Al}(\text{OH})_2^+$) become available in soil solutions when pH is below 5.5 and may be detrimental to many plant species, especially some widely cultivated crop plants such as wheat (*Triticum aestivum* L.), corn (*Zea mays* L.), or soybean (*Glycine max* (L.) Merr.) (Roy *et al.*, 1988; Marschner, 1995; Kochian, 1995). For this reason, Al toxicity is a well-studied topic in agricultural research all over the world.

The principal symptom of Al toxicity in plants is a rapid inhibition of the root growth, which results in a reduced and damaged root system and can lead to mineral deficiencies (e.g. Ca, Mg) and water stress (Foy *et al.*, 1978; Roy *et al.*, 1988). The task to unravel the toxic properties of Al in plants is complex and a number of different mechanisms have been proposed including Al apoplastic lesions, interactions with the cell wall, the plasma membrane, or the root symplasm (Taylor, 1988a; Lüttge and Clarkson, 1992; Kochian, 1995).

Most of the native plant species on acid soils do not suffer from Al toxicity since they exhibit a wide range of resistance mechanisms. There are two main strategies to deal with Al toxicity, namely exclusion from the root apex, which is the major site of Al toxicity, and development of one or more mechanisms to tolerate Al once it has been taken up by the plant. Most physiologists working in the field of metal stress also distinguish between internal and external mechanisms depending on the site of Al response. Internal mechanisms occur in the symplasm, while external mechanisms take place in the apoplasm. The hypotheses proposed for external resistance include: (1) binding or fixation of Al in the cell wall; (2) exudation of Al chelator ligands, especially organic acids, or possibly phosphate; (3) selective permeability of the plasma membrane; (4) plant-induced pH barrier in the rhizosphere; and (5) Al efflux (Taylor, 1991; Kochian, 1995; Ma, 2000; Taylor *et al.*, 2000; Ma *et al.*, 2001). Our understanding of symplastic detoxification of Al is even more fragmentary and focuses on the following hypotheses: (1) formation of Al chelates by organic acids, proteins, or other organic ligands; (2) compartmentalization in the vacuole; (3) synthesis of Al tolerant proteins; and (4) elevated enzyme activity (Taylor, 1991; Kochian, 1995; Ma, 2000; Ma *et al.*, 2001). Because of the complex interactions between Al and the plant, it is very likely that plants are able to use a number of different mechanisms to confer Al resistance. This is supported by studies focusing on genetic mechanisms, which have shown that Al resistance is a trait that can be controlled by one or more major genes and several minor genes (Aniol, 1990; Larsen *et al.*, 1998; Ma *et al.*, 2000).

What are Al accumulators?

Only a small fraction of Al is taken up in plants before it is returned to its mineral forms. Since the majority exhibits a variety of mechanisms that exclude high Al levels from the shoot, the mean content of Al in plant tissues is found to be 200 mg kg^{-1} or 0.02% of the dry matter (Hutchinson, 1943, 1945). Moreover, Al levels in the aerial plant tissues are generally below 300 mg kg^{-1} . Differential uptake and transport between root and shoot in these Al excluding plants leads to rather constant low shoot levels over a wide range of Al concentrations in the soil.

Contrary to Al excluders, a limited number of plant species can be termed Al accumulators, or Al plants, because they take up Al in above ground tissues in quantities far above the average level (Hutchinson and Wollack, 1943; Hutchinson, 1945; Robinson and Edgington, 1945; Chenery, 1948a). A frequency histogram of Al levels in related taxa or acid, tropical rainforest floras usually shows a clear bimodality of normal Al levels and very high values representing Al accumulators. The establishment of a 1000 mg kg^{-1} threshold for dried leaf tissue has been suggested by several authors and is followed in this chapter as it generally serves to differentiate Al accumulators from non-accumulators very well. Masunaga *et al.* (1998), however, proposed a criterion of 3000 mg kg^{-1} for the definition of Al plants based on the relationships between Al concentrations and five other elements (Ca, Mg, P, S, Si).

Al concentrations above 1000 mg kg^{-1} are not only limited to leaves, but may also be found in tissues of the wood, bark, fruit and seed (Haridasan, 1987; Silva, 1990; Masunaga *et al.*, 1998). Although the Al content of leaves is generally higher than in other aerial tissues, there appear to be quite a lot of plant species in which the Al bark content is above 1000 mg kg^{-1} , but lower in the leaves. It is therefore suggested that Al accumulators should be defined as plants in which an Al concentration of at least 1000 mg kg^{-1} has been recorded in the dry matter of leaves in at least one specimen growing in its natural habitat.

Although exact analytical techniques are preferable to determine whether or not plants show Al accumulation, most Al accumulators have been reported using semiquantitative tests. The 'Aluminon' (ammonium aurine tricarboxylate) test was first used by Chenery (1946, 1948b). This simple but adequate test can be applied to both living and dried material. The pinkish to orange reagent acquires a distinctive dark red to crimson colour when the Al content in the tissue tested exceeds 1000 mg kg^{-1} . Kukachka and Miller (1980) devised a similar chemical spot-test using a chrome azurol-S solution to detect Al accumulation in dried wood samples.

The distribution of Al accumulators in flowering plants

Our knowledge on the distribution of Al accumulators mainly builds on the substantial studies of Yoshii and Jimbo (1932), Chenery (1946, 1948a,b, 1949), Chenery and Sporne (1976), Webb (1954) and Kukachka and Miller (1980). A list of families with Al accumulating taxa is summarised in Table 23.1 based on these earlier studies. At present, the feature is found in approximately 55 angiosperm families, which largely belong to the eudicots. About 93% of all Al accumulators that have been recorded belong to the asterids and rosids. Within these groups, Al accumulators are mainly restricted to the orders Myrtales and Gentianales, which account for 42.5% and 35% respectively of the total number of Al accumulators known. It should be emphasized, however, that Table 23.1 can only be regarded as a preliminary compilation since it is likely that several, as yet unidentified Al accumulators growing on natural and man-made acid soils remain to be discovered by plant scientists.

Examples of families that include many Al accumulators are Hydrangeaceae, Melastomataceae, Rubiaceae, Theaceae, Symplocaceae and Vochysiaceae. The tea bush (*Camellia sinensis* (L.) Kuntze) is probably one of the most important economic Al accumulators (Chenery, 1955). Another frequently cited Al accumulator is *Hydrangea macrophylla* DC. The blue or pink flower colour of this ornamental species is found to be dependent on the Al concentration in the shoot (Allen, 1943; Chenery, 1946; Takeda *et al.*, 1985). The occurrence of Al accumulation in these families may illustrate the close connection of this feature with taxonomy.

Al accumulators are mainly woody, perennial taxa from the tropics. Noteworthy is the lack of Al accumulators in the monocotyledons. According to Chenery (1949), this lack is closely connected with the rarity of species with high cell sap acidities and with the herbaceous habit. Remarkable Al accumulators in the monocots, however, include at least six genera of Rapateaceae, the genus *Aletris* (Liliaceae) and a few grass genera (Poaceae) (Chenery, 1949). Furthermore, there are also data available on Al accumulation outside angiosperms, in particular ferns, clubmosses, liverworts and algae, but the feature appears to be entirely lacking in gymnosperms (Yoshii and Jimbo, 1932; Hutchinson, 1945; Chenery, 1949; Webb, 1954).

Table 23.1 List of Al accumulators in angiosperms summarized from Jansen *et al.* (2002); classification follows APG (1998); taxa in bold include strong and/or numerous Al accumulators; *Cardiopteridaceae *sensu* Karehead (2001)

| | <i>Order</i> | <i>Family</i> |
|---------------------|--------------|--|
| Basal angiosperms | | Amborellaceae, Illiciaceae, Trimeniaceae, Winteraceae |
| | Laurales | Lauraceae, Monimiaceae, Siparunaceae |
| Monocots | Asparagales | Orchidaceae |
| | Liliales | Liliaceae |
| Commelinoids | | Rapateaceae |
| | Poales | Poaceae |
| Basal eudicots | Proteales | Proteaceae (Grevilleoideae, Placospermum C.T.White and W.D.Francis) |
| | Ranunculales | Lardizabalaceae |
| Basal core eudicots | Santalales | Olacaceae |
| | Saxifragales | Daphniphyllaceae, Grossulariaceae, Saxifragaceae |
| Eurosids I | Cucurbitales | Anisophylleaceae |
| | Fabales | Polygalaceae (Moutabeae, Xanthophylleae) |
| | Fagales | Fagaceae, Juglandaceae |
| | Malpighiales | Euphorbiaceae (Aporuseae, Phyllanthae subtribe Uapaceae), Flacourtiaceae? (Soyauxia Oliv.), Goupiaceae, Lacistemataceae, Violaceae |
| | Oxalidales | Cunoniaceae |
| Eurosids II | Myrtales | Combretaceae, Crypteroniaceae, Melastomataceae, Memecylaceae, Myrtaceae, Rhynchoalycaceae, Vochysiaceae |
| Basal asterids | Cornales | Cornaceae, Hydrangeaceae |
| | Ericales | Diapensiaceae, Ebenaceae, Lecythydaceae, Myrsinaceae, Symplocaceae, Ternstroemiaceae, Theaceae |
| Euasterids I | Gentianales | Apocynaceae, Gentianaceae, Strychnaceae, Rubiaceae (Rubioideae) |
| | Lamiales | Lentibulariaceae |
| Euasterids II | | Icacinaceae s.str. ? (<i>Platea</i> Blume), Polyosmaceae |
| | Aquifoliales | Phyllonomaceae |
| Uncertain position | | Cardiopteridaceae* (including <i>Gonocaryum</i> Miq., <i>Leptaulus</i> Benth.), Geissolomataceae, Pentaphylacaceae, Peridiscaceae |

Al accumulation: a useful character in plant systematics

Al accumulation has attracted little taxonomic attention in earlier classification systems. Only a few authors believed that a high Al concentration might not only characterize certain species, genera and families, but to a certain extent also rather loosely defined groups of allied families (Yoshii and Jimbo, 1932; Hutchinson, 1943; Chenery, 1948b, 1949; Chenery and Sporne, 1976; Chenery and Metcalfe, 1983). Cronquist (1980) was not convinced that the feature proves systematic significance due to its scattered taxonomic distribution and the pervasive parallelisms among the angiosperms. It can be suggested that the following problems have hindered the use of this phytochemical character in systematics: (1) conflicts with morphological taxonomies; (2) difficulties in access to data on Al concentrations; (3) the poor phylogenetic knowledge of the angiosperms; and (4) more attention to ecological and physiological aspects of Al accumulators than its phylogenetic significance.

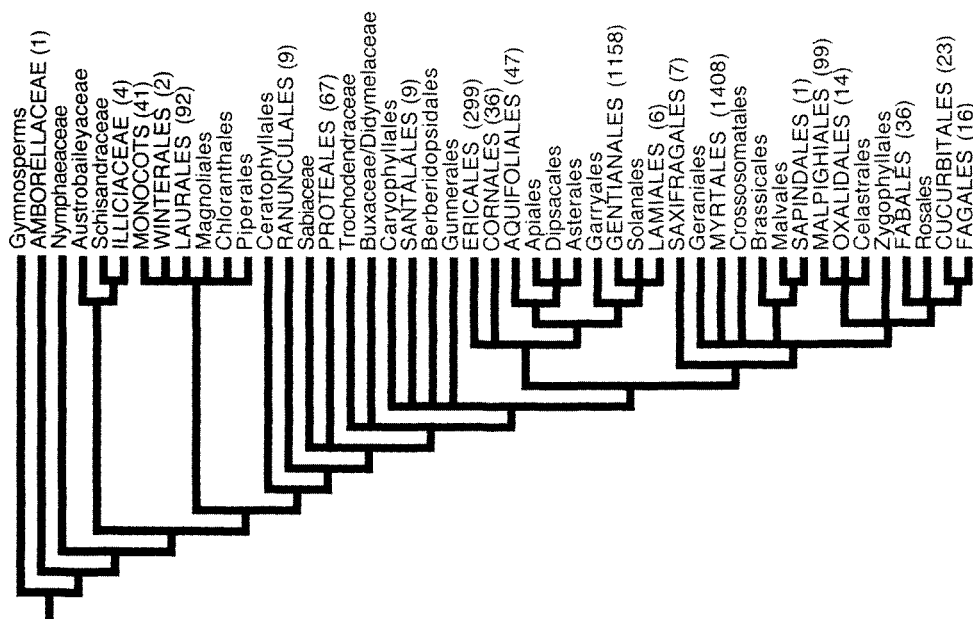


Figure 23.1 Angiosperm phylogeny inferred from 18S rDNA, *rbcl*, and *atpB* sequences (Soltis *et al.*, 2000); taxa including Al accumulators are printed in capitals; the number of Al accumulating plants known is indicated between brackets.

Recent phylogenetic analyses based on molecular sequence data allow us to evaluate the evolutionary trends of characters and character states in a more accurate way than in the past. With respect to Al accumulation, this character has independently been developed in several plant groups that are not necessarily closely related, but its occurrence is far from being randomly distributed. The primitive status of Al accumulation in angiosperms has previously been suggested on the basis of statistical correlations (Chenery and Sporne, 1976). Its primitive (plesiomorphic) or derived (apomorphic) nature, however, largely depends on the taxonomic level and should be evaluated for each taxonomic group separately (Jansen *et al.*, 2002). For instance, at the family level of Rubiaceae the presence of Al accumulation represents a synapomorphy that characterizes one of the three subfamilies, namely Rubioideae. Within the Rubioideae, however, the feature should be considered as a symplesiomorphy and the secondary loss of the feature as evolutionarily advanced or specialized in the herbaceous members (Jansen *et al.*, 2000a,b). In general, Al accumulation is most widespread at the base of major clades that belong to the higher groups of the eudicots, namely rosids (Malpighiales, Oxalidales, Myrtales) and asterids (Cornales, Ericales, Gentianales, Aquifoliales). On the other hand, Al accumulators are not common in the basal angiosperms, with the exception of *Amborella* Baill. (Amborellaceae) and some members within the Laurales (Figure. 23.1).

The evolutionary trend of Al accumulation, moreover, is complicated by the occurrence of numerous reversals (or losses) and parallel origins. Alternatively, the low incidence of Al accumulators in derived groups might also be correlated with their frequent herbaceous habit (see further). The two examples given below illustrate the potential significance of Al accumulation in plant systematics.

Al accumulators in the family Polygalaceae appear to be restricted to the following genera: *Barnhartia* Gleason, *Diclidanthera* Mart., *Eriandra* P.Royen & Steenis, *Moutabea*

Aubl. and *Xanthophyllum* Roxb. (Chenery, 1948b; Eriksen, 1993). The taxonomic position of these genera within the family supports the delimitation of the Xanthophylleae and Moutabeae, while the feature is lacking in all members of the Carpolobieae and Polygalae (Eriksen, 1993). Since Al accumulators take the most basal position within this family (Persson, 2001), it can be hypothesized that the feature has been lost in the more derived clades.

In contrast with the distribution of Al accumulation in Polygalaceae, a possible relationship between the Al accumulating families Anisophylleaceae and Cunoniaceae is in conflict with results from molecular data. Al accumulation is strongly present in all members of the Anisophylleaceae and clearly distinguishes this family from the Rhizophoraceae (Dahlgren, 1988). Several Al accumulating genera have been reported in the Cunoniaceae, namely in the genera *Acrophyllum*, *Anodopetalum*, *Aphanopetalum*, *Ceratopetalum*, *Gillbeea*, *Platylophus*, *Schizomeria* and *Spiraeanthemum* (Chenery, 1948b; Webb, 1954). A recent investigation of floral structures suggests that Anisophylleaceae and Cunoniaceae may be closely related, which is also supported by paleobotanical evidence (Matthews *et al.*, 2001; Schönemberger *et al.*, 2001). Nevertheless, Cunoniaceae appeared to have a position within Oxalidales according to molecular sequence data, while Anisophylleaceae were placed in the Cucurbitales (APG, 1998; Savolainen *et al.*, 2000; Soltis *et al.*, 2000). The presence of Al accumulation in both families may provide additional support for their potential close relationship. Moreover, this feature is not known in other families of the Cucurbitales or Oxalidales. It is clear, however, that more extensive phylogenetic studies based on molecular evidence are required to study critically the position of both families. Furthermore, the distribution of Al accumulators within Cunoniaceae needs further attention, since several non-accumulators have been reported as well. Preliminary data from literature indicate that Al accumulation characterizes the tribe Schizomerieae and the clade comprising *Acrophyllum* and *Gillbeea*, which take a relatively basal position within the family (Bradford and Barnes, 2001).

The ecology of Al accumulators

It has often been stated that Al levels in leaves are subject to ecological influences. According to Chenery and Metcalfe (1983) Al accumulation is considered to be 'a process that depends partly on the influence of heredity and partly on ecological influence'. Among ecological factors, the one which comes to mind most readily is soil acidity. Indeed, the soil pH comprises the most important factor in Al uptake because the solubility and bioavailability of Al increases with decreasing pH. Most Al accumulators grow in leached, acid soils and they occur particularly in tropical, humid rainforests and savannahs. Al accumulators are much less common in temperate regions or dry areas (Webb, 1954; Larcher, 1980; Lüttge, 1997). Hence, it is essential to incorporate ecological factors such as moisture, acidity and Al solubility in the soil when comparing Al levels of plants from different localities.

Another topic that merits attention is the relationship between Al accumulation and growth forms of angiosperms, especially the woody versus herbaceous habit. The number of soft stemmed herbs that show Al accumulation is very low. A remarkable example is for instance *Coccocypselum* (Rubiaceae), which is a small neotropical genus of creeping herbs with metallic blue berries. All specimens of *Coccocypselum* growing in its natural habitat are found to be strong Al accumulators (Jansen *et al.*, 2000a). Other herbaceous Al accumulators occur in Lentibulariaceae (*Genlisea*, *Utricularia*) and in a few monocots. One may speculate that the higher amount of secondary cell wall material in woody

plants is favourable for the binding or fixation of Al in the cell wall of 'dead' tissues. This idea may indicate that Al levels are usually higher in wood than in leaves, but the reverse has generally been found. Another aspect that may partly affect Al concentrations in plant tissues includes the difference in life span of the organs or the growth rate between woody and herbaceous plants. It has been demonstrated that the Al concentration in the xylem sap of buckwheat is not much lower than that of woody plants (Ma and Hiradate, 2000). This may suggest that mechanisms of Al uptake and transport from roots to shoots are not responsible for the absence of high Al levels in the shoot of herbaceous plants.

A study of different specimens of Al accumulators from various localities may allow us to distinguish between obligate and facultative Al accumulators. Obligate Al accumulators can be defined as species in which the Al content of the shoot is constantly above 1000 mg kg^{-1} , irrespective of the Al solubility in the soil, while the internal Al concentration in facultative accumulators (also termed 'indicator' species) more or less reflects the external soil level. Noteworthy is that Al concentrations in obligate accumulators do not vary during seasonal changes. Al levels in above ground organs of facultative accumulators, however, may vary depending on seasonal variations (de Medeiros and Haridasan, 1985; Mazorra *et al.*, 1987).

Al accumulators usually do not show symptoms of Al toxicity in acidic soils. There is even evidence that Al stimulates the growth of several Al accumulating and Al tolerant plants. Examples of Al accumulators that show a beneficial effect of Al are *Camellia sinensis* (L.) Kuntze, *Miconia albicans* (Sw.) Triana and *Melastoma malabathricum* L. (Figure. 23.2; Matsumoto *et al.*, 1976; Konishi *et al.*, 1985; Haridasan, 1988; Osaki *et al.*, 1997; Watanabe *et al.*, 1997, 1998; Watanabe and Osaki, 2001). A similar effect has been reported in *Eucalyptus gummiifera* Hochr. (Mullette, 1975). This species grows on acidic soils, but is not an Al accumulator. These results raise interesting physiological questions about the possible role of Al in the metabolism of plants with Al accumulation and



Figure 23.2 Al-induced growth enhancement in *Melastoma malabathricum* L. (Melastomataceae); seedlings were grown in a nutrient solution with Al (left) and without Al (right) for 3 weeks.

tolerance to Al toxicity. It can be suggested that Al is essential rather than beneficial for Al accumulators since healthy plants in acid, Al-rich soils may turn into sick plants with necrosis and leaf yellowing when growing on a non-acid, Al-poor soil. It is unknown whether this holds true for all Al accumulators.

We still know very little about the biological and evolutionary significance of metal accumulation in general. Hypotheses that have been addressed include the following: drought resistance, inadvertent uptake, tolerance or disposal of metal from plants and defence against herbivores or pathogens (Raskin *et al.*, 1994). With respect to Al, it is difficult to estimate the physiological cost that detoxification of Al in above ground plant tissues may imply in comparison with detoxification by exclusion mechanisms such as exudation of Al chelator ligands or impermeability of the endodermic cells to Al. It is likely that plants use different mechanisms effectively to guarantee safety and protection of vital tissues and organs. It can also be suggested that the tolerance limits to survive in a wide range of soil pH are higher for Al accumulators than for non-accumulators or Al excluders, which may indicate that Al accumulation is a more advantageous strategy for plants that grow in acidic soils.

After studying two forest communities in the cerrado region of central Brazil, Haridasan and Araújo (1988) concluded that Al accumulators growing on a strongly acid dysotrophic latosol account for 17.3% of the total importance value of the community. On the other hand, Al accumulators account for only 11.7% of the total importance value in a community on a mesotrophic but slightly acidic soil. This difference could be explained on the basis of a possible successful, but not essential, adaptive strategy of Al accumulators for survival and competition in strongly acidic and dysotrophic latosols. Al accumulators, however, are not absolutely restricted to this soil type. Hence, it is difficult to speculate any worthwhile hypothesis regarding the adaptive strategy of Al accumulators. Al accumulation is most likely to be just one of several options which allow plants to survive in an extremely acid, nutrient deficient soil environment (Haridasan, 1987; Haridasan and Araújo, 1988).

The evolution of heavy metal accumulation

Plant adaptation to high-metal soils and ecotypic variation with regard to resistance and accumulation of heavy metals (e.g. Ni, Zn, Pb, Cd, Co, Cu, Mn, Cr, Se) is a common phenomenon (Baker, 1987; Macnair, 1993). Often there is a single species exhibiting accumulation of a heavy metal within a genus, suggesting that differences in heavy metal content can be attributed to rather recent evolutionary processes (Reeves and Baker, 2000). For instance Ni accumulators in Rubiaceae are limited to a few genera and its taxonomic implications appear to be significant at the specific or generic level. In this way, the distribution of Ni accumulating species within the huge genus *Psychotria*, which includes both Ni accumulators and non-accumulators, may be interesting for classification purposes among species. The most unifying feature of heavy metal accumulators is probably their restricted distribution and high degree of endemism (Baker and Brooks, 1989). By contrast, Broadley *et al.* (2001) found significant variation in heavy metal accumulation at the taxonomic level of orders (especially Malpighiales, Brassicales and Asterales) and above. This may indicate that rather ancient evolutionary traits influence heavy metal accumulation in angiosperms.

In our opinion, Al accumulation in leaves shows stronger phylogenetic signals at a higher taxonomic level than heavy metal accumulation. This difference may be explained by the availability of the elements and their role in the metabolic processes of plants. Acid soils with

high levels of soluble Al comprise up to 30% of the world's ice-free land area (von Uexküll and Mutert, 1995), while metalliferous soils usually have a rather isolated and local distribution. Metalliferous soils show abnormally high concentrations of some elements that are normally present only at minor or trace levels. Unlike Al accumulation, the effect of metalliferous soils on plants may result in the development of a characteristic, local flora of metal-tolerant species (Reeves and Baker, 2000). Moreover, at least some heavy metals, such as Cu, Mn and Zn, have been shown to be essential for plant growth. Thus it is hypothesized that plant adaptation to metalliferous soils may have developed at the level of species or varieties, while the strategy of Al accumulation or exclusion appears to have affected larger taxonomic groups such as plant families or orders. It is also likely that Al accumulation is controlled by different physiological processes than heavy metal accumulation.

The wider relevance of studies on Al accumulation

Al accumulators have been used in traditional dying technologies for centuries. Although most of these plants contain no colouring substances, the Al in their bark and leaves operates as a mordant instead of alum and serves to set the colours furnished by other plants (Robinson and Edgington, 1945). The traditional use of a plant as a mordant has revealed many cases of Al accumulating plants, especially in South East Asia. For instance as early as 1743, Rumphius described a tree as *Arbor aluminosa* or 'Aluyn-Boom', which is a species of *Symplocos* (Symplocaceae) (Hutchinson, 1943).

Plant species show considerable genetic variation in their Al sensitivity. In general, species with low resistance to Al stress are only able to grow in soils with low Al levels, while the plants with moderate or high resistance can survive higher Al concentrations. Over recent years, numerous intensive research efforts have investigated resistance to Al stress, especially with respect to crop plants such as *Zea mays* L., *Triticum aestivum* L., *Secale cereale* L., *Glycine max* (L.) Merr. and *Hordeum vulgare* L. (e.g. Taylor, 1988b; Carver and Ownby, 1995; Kochian, 1995). These studies illustrate the agronomic importance of Al toxicity. Hence, further understanding of the distribution of Al accumulators and their physiological processes may help to develop more resistant crops or plants that can be used as forage for animals.

A better understanding of the nutritional strategies of plants adapted to different soils may also be essential for employing certain species in forest management, soil improvement and recuperation of degraded lands in tropical rainforests or savannahs. This may also provide useful applications for forestry in temperate areas of Europe and North America, where acid rain is thought to be one of the causes of forest decline (Godbold *et al.*, 1988). Ideally, studies on Al accumulating plants should be integrated in the whole understanding of Al in living organisms. They should also be considered in any assessment of the total dietary intake of Al in human beings, as Al is suggested to play a role in human neurodegenerative disorders such as Alzheimer's disease (Exley, 2001).

General conclusion

Al accumulation in plants is not only a very important and ecophysiological highly interesting phenomenon, but it also provides useful information for systematic purposes at relatively high taxonomic levels. Its primitive status in the derived groups of the

angiosperms is generally confirmed following recent molecular phylogenies. In particular, the feature appears to be largely restricted to woody representatives of the basal branches of eurosids and asterids.

Further fruitful research on Al plants should incorporate a multidisciplinary approach ranging from ecology, soil biology, evolutionary biology, taxonomy, physiology and phytochemistry. It may not be easy to achieve a synthesis between these diverging fields, but data on the systematic distribution, localization of Al in different plant tissues and ecological or habitat categories may offer an excellent start for collaborative efforts to understand the biology of Al accumulators.

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