

The origin of alternation of generations in land plants: a focus on matrotrophy and hexose transport

Linda K. E. Graham and Lee W. Wilcox

Phil. Trans. R. Soc. Lond. B 2000 **355**, 757-767
doi: 10.1098/rstb.2000.0614

References

Article cited in:

<http://rstb.royalsocietypublishing.org/content/355/1398/757#related-urls>

Email alerting service

Receive free email alerts when new articles cite this article - sign up in the box at the top right-hand corner of the article or click [here](#)

To subscribe to *Phil. Trans. R. Soc. Lond. B* go to: <http://rstb.royalsocietypublishing.org/subscriptions>

The origin of alternation of generations in land plants: a focus on matrotrophy and hexose transport

Linda K. E. Graham and Lee W. Wilcox

Department of Botany, University of Wisconsin, 430 Lincoln Drive, Madison, WI 53706, USA (lkgraham@facstaff.wisc.edu)

A life history involving alternation of two developmentally associated, multicellular generations (sporophyte and gametophyte) is an autapomorphy of embryophytes (bryophytes + vascular plants). Microfossil data indicate that Mid–Late Ordovician land plants possessed such a life cycle, and that the origin of alternation of generations preceded this date. Molecular phylogenetic data unambiguously relate charophycean green algae to the ancestry of monophyletic embryophytes, and identify bryophytes as early-divergent land plants. Comparison of reproduction in charophyceans and bryophytes suggests that the following stages occurred during evolutionary origin of embryophytic alternation of generations: (i) origin of oogamy; (ii) retention of eggs and zygotes on the parental thallus; (iii) origin of matrotrophy (regulated transfer of nutritional and morphogenetic solutes from parental cells to the next generation); (iv) origin of a multicellular sporophyte generation; and (v) origin of non-flagellate, walled spores. Oogamy, egg/zygote retention and matrotrophy characterize at least some modern charophyceans, and are postulated to represent pre-adaptative features inherited by embryophytes from ancestral charophyceans. Matrotrophy is hypothesized to have preceded origin of the multicellular sporophytes of plants, and to represent a critical innovation. Molecular approaches to the study of the origins of matrotrophy include assessment of hexose transporter genes and protein family members and their expression patterns. The occurrence in modern charophyceans and bryophytes of chemically resistant tissues that exhibit distinctive morphology correlated with matrotrophy suggests that Early–Mid Ordovician or older microfossils relevant to the origin of land plant alternation of generations may be found.

Keywords: alternation of generations; embryophytes; bryophytes; charophycean green algae; hexose transporter genes and proteins

1. INTRODUCTION

Alternation of generations in autotrophs is generally defined as the occurrence of a life history in which there are at least two multicellular generations, the gametophyte and the sporophyte, linked by unicellular reproductive stages, namely gametes and spores (figure 1). Spores are generated by sporic meiosis, which is the type of meiosis associated with alternation of generations (Raven *et al.* 1999). This article does not address ‘alternation of generations’ that may occur in various autotrophic rotists that occur primarily as unicells (e.g. certain aptophyte algae), or in heterotrophs (such as foraminifera and fungi).

(a) *The occurrence of alternation of generations in autotrophs*

Life histories involving two or more alternating multicellular generations have evolved several times among photosynthetic protists (algae) (Graham & Wilcox 2000). For example, various bangiophycean red algae have a life history with two multicellular stages, and a three-stage life history appears to be a basic (plesiomorphic) feature of florideophycean red algae. Ancestors of phaeo-

phyceans (brown algae) independently acquired alternation of two multicellular generations; so far as is known, their closest extant relatives (tribophyceans and other ochrophyte/chromophyte/heterokont algae) lack such a life history. Among modern green algae, alternation of two multicellular generations occurs only in certain orders of the class Ulvophyceae, and is lacking in the three other green algal classes that include multicellular forms (namely Trebouxiophyceae, Chlorophyceae and Charophyceae *sensu* Graham & Wilcox (2000)). Evidence for independent evolution of alternation of generations in red, brown and green algae suggests that it is highly adaptive. Hypothetical adaptive aspects of life history variation in autotrophs are discussed by Bell & Koufopanou (1991), Otto & Goldstein (1992), Baillard (1997) and Bell (1997). A review of alternation of generations in land plants, with an emphasis on fossil branched gametophytes that are thought to be linked to the life histories of protracheophytes and early vascular plants, is provided by Kenrick (1994).

(b) *Ploidy change and alternation of generations*

It should be noted that while textbook depictions of alternation of generations in algae and land plants

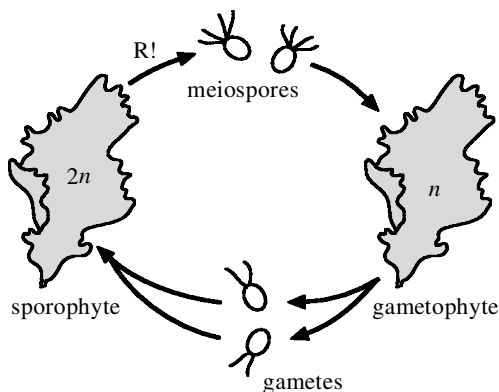


Figure 1. Diagram of alternation of multicellular generations. R! indicates the occurrence of meiosis. A life history involving spatially and temporally separate generations is characteristic of several groups of algae.

Typically describe the gametophytic generation as being haploid, and the sporophytic generation as diploid, there are many examples among the algae of life history phase change that are not correlated with change in chromosome number (Graham & Wilcox 2000). For example, the nuclei of sporophytes and gametophytes of the brown seaweed *Haplospora globosa* (Tilopteridales) possess the same number of chromosomes. However, the DNA level of sporophytic nuclei is twice that of gametophytic nuclei (Kuhlenkamp *et al.* 1993). In other algae, environmental factors are thought to be as, or perhaps more, important than chromosomal level in determining the direction of life history phase change. Environmental effects are regarded as possible explanations for cases of apogamy (transition to the sporophyte phase in the absence of gamete production and syngamy) and apospory (transition to the gametophytic phase in the absence of meiosis and spore production). In seedless plants, apogamy and apospory are also observed, but gene dosage effects are important. Maintenance of sporophytic growth depends on the presence of at least two sets of chromosomes, whereas gametophytic growth in culture does not continue when four or more sets of chromosomes are present (Bell 1991).

In higher plants, there are many examples of production of young sporophytes (embryos) from cells other than zygotes (e.g. microspore embryogenesis, somatic embryogenesis and apomixis) (Harada *et al.* 1998), and the genetic basis for such variants from the expected life history cycling is becoming clearer. For example, the *Arabidopsis* gene *LEAFY COTYLEDON 1 (LEC1)*, which encodes a transcription factor, is sufficient to induce embryo-like development from vegetative cells (Lotan *et al.* 1998).

(c) **Importance of sporophyte/gametophyte interactions**

The above variations having been noted, a life history involving alternation of multicellular gametophyte and sporophyte generations characterizes all groups of extant algae and plants, which comprise the Kingdom Plantae, as defined by Raven *et al.* (1999). Members of the extant plant kingdom constitute a monophyletic group that includes multiple lineages of early-divergent bryophytes

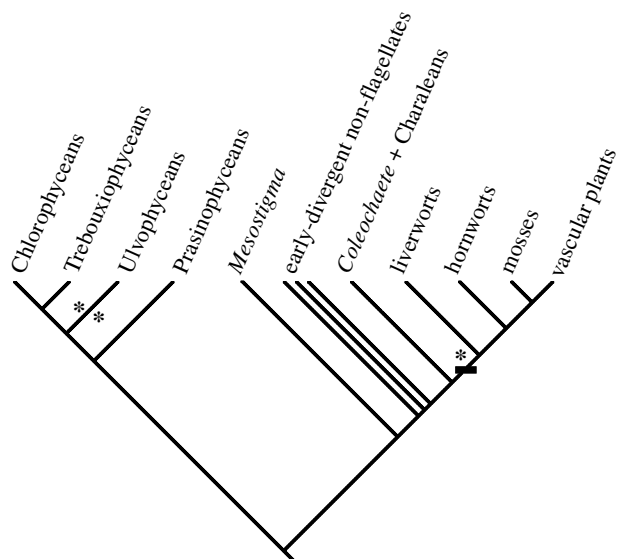


Figure 2. Diagrammatic representation of phylogenetic relationships among various groups of the green algae and embryophytes. Asterisks indicate cases of presumed independent origin of a life history involving alternation of two multicellular generations. The bar indicates the only known case among green autotrophs of the occurrence of a dependent, multicellular sporophyte (alternation of generations that are not separated temporally or spatially).

Table 1. *Matrotrophy and associated life history change have led to the origin of three high-diversity, long-lived clades*

clade	reproductive innovation	mechanism
florideophycean red algae	carposporophyte	$n/2n$ cell fusions
embryophytes	dependent embryo	placental transfer cells
eutherian mammals	viviparity	complex placenta

and later-divergent tracheophytes (Kenrick & Crane 1997). It is important to recognize that alternation of generations in the Kingdom Plantae is distinctive in that embryonic sporophytes occur in close spatial and temporal association with female (or bisexual) gametophytes. Plant embryos, including those of the simplest liverworts as well as derived angiosperms, seem generally to be nutritionally and developmentally dependent on parental gametophyte tissues for at least some period of time in early development. The presence of a dependent embryonic stage is the basis for the term embryophytes, commonly used as a synonym for Kingdom Plantae. The occurrence of alternation of multicellular generations coupled with dependent embryos in all groups of land plants suggests that these features are autapomorphic (unique and defining) features of embryophytes (figure 2). Multicellular sporophytes do not occur in the charophyceans, the green algal lineage most closely related to embryophytes (Graham 1993), and dependence of the embryonic sporophyte is lacking in most other algae that have alternation of generations. An exception to this generality is the floridophycean red algae, whose

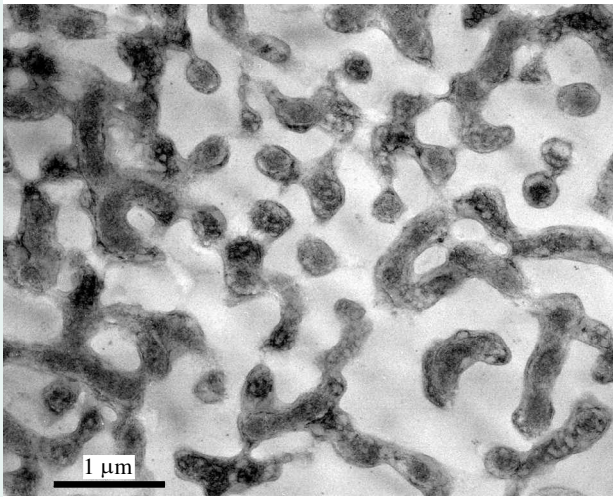


Figure 3. A transmission electron micrograph of a portion of placental transfer cell of the liverwort *Conocephalum conicum*, showing extensive development of wall ingrowths. The preparation was high-pressure frozen, a method that facilitates reservation of cell membranes.

carposporophyte generation is nutritionally and developmentally dependent on the female gametophyte.

In florideophyceans and land plants, the embryonic sporophyte's dependency has been described as matrotrophy (literally 'mother feeding'), by analogy to embryonic nutritional and developmental dependence in eutherians and other viviparous metazoans (Graham, 1996; E. 1996). Matrotrophy was a key innovation in the diversification of these three major clades (florideophyceans, embryophytes and eutherians), its adaptiveness in each case hypothetically related to increases in fecundity (table 1). In each case, there is a morphological correlate of matrotrophy—the placenta, a region of specialized cells (tissues in the cases of embryophytes and eutherians) that facilitates maternal to embryonic nutrient transfer.

In embryophytes (as in eutherians), there are no intercellular connections linking parental and embryonic tissues, hence transport of solutes occurs by means of cell membrane transporters (i.e. is apoplastic). Embryophytic placental cells typically possess elaborate systems of wall ingrowths, which greatly increase the surface area of the cell membrane across which facilitated diffusion or active transport must occur (figure 3). Such cells are known as placental transfer cells; these may occur on one or both sides of the generational gap. Placental transfer cells are well studied at the ultrastructural level in bryophytes (see reviews by Ligrone & Gambardella 1988; Ligrone *et al.* 1993) and in higher plants such as *Arabidopsis* (Murgia *et al.* 1993). Physiological studies provide substantial evidence that the placentae of bryophytes (Browning & Running 1979*a,b*; Renault *et al.* 1992) and flowering plants (Van Caesele *et al.* 1996) function in apoplastic transport.

Provision of the florideophycean embryo (carposporophyte) with nutrients from the maternal generation is suggested to be a strategy for amplifying the products of sexual recombination where fertilization rates are limited by the absence of flagella from male gametes. (It should

be noted, however, that few actual measurements of fertilization rates have been made for red algae.) A similar adaptive benefit has been hypothesized to accrue to seedless land plants, whose fertilization rates may be limited by availability of liquid water for transport of flagellate sperm (Searles 1980). The rest of this paper focuses on palaeontological, neontological and combined approaches to understanding the role of matrotrophy in the origin of alternation of generations and the dependent sporophytic embryo of land plants.

2. PALAEOLOGICAL EVIDENCE THAT EARLIEST-KNOWN (ORDOVICIAN) LAND PLANTS POSSESSED ALTERNATION OF GENERATIONS

Microfossils of the Mid–Late Ordovician age, described from Libyan deposits by Gray *et al.* (1982), Taylor (1995) and Strother *et al.* (1996), provide evidence that the earliest-known land plants possessed alternation of generations. These remains include spores arrayed in persistent tetrads. Persistence of the fossil tetrads suggests the presence of spore walls that were chemically resistant to the effects of microbial decomposition and diagenesis, as are the sporopollenin-impregnated spore walls of modern land plants, including those of bryophytes. Similar spore tetrads, known to have resulted from meiosis, are produced by the early-divergent (see Lewis *et al.* 1997) complex thalloid liverwort *Sphaerocarpos*. Such tetrads remained intact after high-temperature acid hydrolysis (Graham & Gray 2001), a procedure that to some extent mimics the degradative chemical and physical effects experienced by plant tissues during fossilization. Occurrence of the Ordovician spores in tetrads is strong evidence that they arose by sporic meiosis, which as noted earlier, is a hallmark of alternation of generations. So far as is known, meiosis is zygotic in all charophyceans exhibiting sexual reproduction (evidence reviewed by Graham 1993). The Ordovician spore evidence suggests that sporic meiosis was an innovation that occurred at the dawn of embryophytes.

Microfossil evidence that sporic meiosis coincided with origin of a multicellular sporophyte generation in earliest-known land plants was provided by Graham & Gray (2001), who demonstrated that the sporangial epidermis of *Sphaerocarpos*, on high-temperature acid treatment, falls apart into monostromatic cellular fragments resembling the most ancient multicellular fossils attributed to land plants. The latter, derived from Libyan deposits of Ordovician age, were associated with spores (Gray *et al.* 1982). On the basis of morphometric comparison with high-temperature acetolysed sporangial epidermis of the early-divergent moss *Sphagnum*, Kroken *et al.* (1996) suggested that the Ordovician cellular scraps represent sporangial epidermal remains, and that they are the earliest-known fossils of the sporophyte generation of land plants.

Hydrolysis-resistance of extant bryophyte sporangial epidermis was attributed to the presence of highly insoluble, wall-bound phenolic polymers, on the basis of specific autofluorescence properties (Kroken *et al.* 1996). Similar, resistant, autofluorescent materials occur in vegetative cell walls of various charophycean algae, suggesting that land plants inherited the capacity to

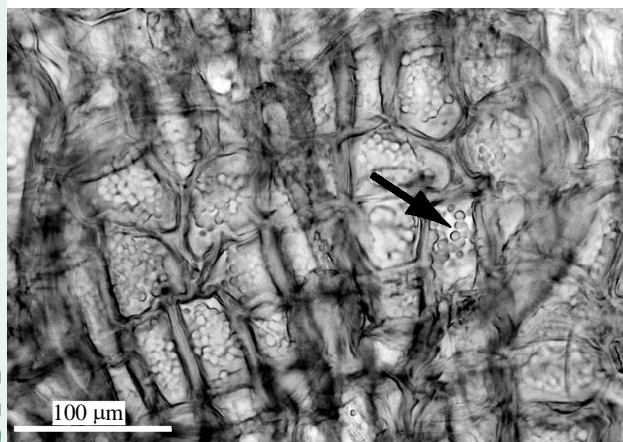


Figure 4. Light micrograph of a portion of a fertile *Coleochaete orbicularis* thallus that was subjected to high-temperature acid hydrolysis. Cell contents have been hydrolysed, but walls of the two large spherical zygotes and their enclosing cortical cells are resistant to hydrolysis. Extensive development of wall growths in cortical cells (see arrow, for example) is readily discernible.

produce such compounds from charophycean ancestors. Resistant, autofluorescent walls of the charophycean green alga *Coleochaete* that are associated with zygotes have Fourier transform infrared (FTIR) spectra that are similar to those of the decay-resistant walls of another charophycean, the desmid *Staurastrum* (Gunnison & Alexander 1975*a,b*), but different from typical plant sporopollenin (Delwiche *et al.* 1989).

Occurrence of similar, resistant compounds in walls of both charophyceans and bryophyte sporangial epidermis suggests the possibility that the Ordovician cellular remains cited earlier could be of charophycean origin. However, the only known example of resistant multicellular tissue in charophyceans is that of the thallus cells, which form a cortical layer that nearly completely encloses mature zygotes of *Coleochaete orbicularis* (Delwiche *et al.* 1989). In such an instance, the zygotes are quite conspicuous and much larger in diameter than the surrounding cells (figure 4); no such enlarged cells occur with the Ordovician cell scraps. Resistant materials in bryophyte sporangial epidermal cell walls may be adaptive in protecting developing spores from microbial attack; earliest land plants may also have derived such a benefit from their presence (Graham & Gray 2000). If Ordovician microfossil scraps do represent sporangial epidermis, the plants from which they were derived possessed a multicellular porophyte and, hence, alternation of generations. However, fossil remains of the placental region, representing evidence that plants had acquired matrotrophy by the Ordovician, are so far lacking.

3. HYPOTHETICAL STAGES IN THE ORIGIN OF ALTERNATION OF GENERATIONS IN EMBRYOPHYTES

Several stages (A–E) in the evolutionary origin of land plant alternation of generations (figure 5) can be deduced by comparison of the reproductive features of charophyceans and bryophytes.

(a) Stage A

An early step, occurring in charophyceans, is hypothesized to have been transition from isogamy to oogamy (production of flagellate sperm and larger, non-flagellate eggs), which also occurs in land plants. Determination of the direction of character transition is dependent on the existence of a robust phylogeny for charophyceans but, to date, phylogenies of charophyceans that are inferred from various nucleic acid sequence data have been highly incongruent. This state of affairs has most likely resulted from extinction, leaving us with a sparse extant representation of the group. Consequently, Graham, L. E. (1996) and Graham & Gray (2001) recommended reliance on molecular architectural data, such as intron insertion events, gene rearrangements or movement of genes between cellular genomes, as is also advocated by Qiu & Palmer (1999). Such data indicate that isogamous Zygnematales are earlier-divergent than oogamous *Coleochaete* and Charales (Graham & Gray 2001; Graham & Wilcox 2000).

However, Sluiman & Guihal (1999) reported that 18S rDNA sequence analysis suggests that the oogamous genus *Chaetosphaeridium*, which is frequently linked with *Coleochaete* on the basis of morphological similarities (see Graham & Wilcox 2000) and *rbcl* sequence data (Chapman *et al.* 1998), may not be closely related to *Coleochaete* but, rather, relatively early-divergent within charophyceans. Other small subunit rDNA sequence analyses indicate surprisingly early divergence of charaleans (Huss & Kranz 1997), this differing dramatically from *rbcl* data indicating that a clade including both Charales and *Coleochaete* is sister to embryophytes (McCourt *et al.* 1996). Further analysis of the evolutionary origin of oogamy in the charophycean–embryophyte lineage may depend on our ability to map isogamy to oogamy transition(s) on to a robust phylogeny.

Aspects of the transition to oogamy that require further study include: (i) origin of the complex, multicellular gametangia of Charales and transition from single-celled antheridia in some species of *Coleochaete* to multicellular aggregates of spermatangial and sterile cells that occur in other *Coleochaete* species (Graham 1993); (ii) changes in flagellate gamete anatomy that may be linked to transition to oogamy (Duncan *et al.* 1997); (iii) loss of flagellar development at the origin of egg cells; (iv) origin of the enlarged eggs, filled with food reserves, of charaleans; (v) origin of the trichogyne, a tubular protuberance of eggs, whose distal wall undergoes controlled hydrolysis, allowing release of sperm attractants and sperm entry in *Coleochaete* (Graham 1993); and (vi) the origin and chemical character of sperm attractants in oogamous charophyceans and bryophytes. A possible approach to the final issue might be to examine *Coleochaete* and charaleans for compounds that are known to coordinate mating in zygnemataleans (Sekimoto *et al.* 1993).

(b) Stage B

Retention of eggs (and zygotes that develop from them) on the maternal (or bisexual parental) thallus is the next hypothesized step (figure 5), as this is an essential precedent for the development of nutritional and developmental interactions between generations. Among charophyceans, examples of egg/zygote retention occur in

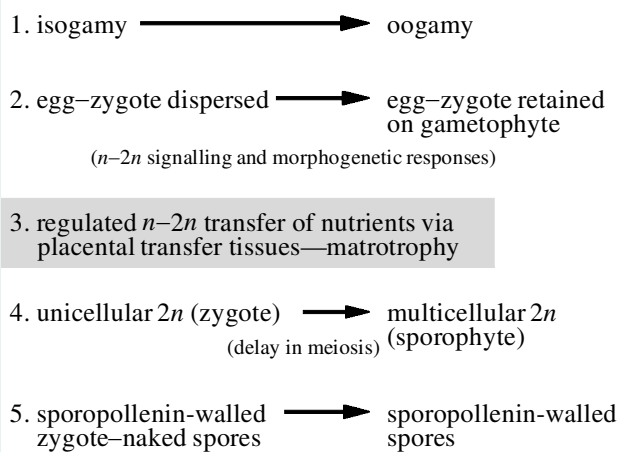


Figure 5. Hypothesized stages in the evolutionary origin of matrotrophic alternation of generations as it occurs in modern embryophytes. Some proposed mechanisms are shown in parentheses. Shading of no. 3 indicates the area of research that is described most fully in this paper.

Coleochaete and Charales, facilitated by mitotic production of a layer of corticating cells or elongation of spirally twisted tube cells, respectively. In both cases, the enclosing cells belong to the parental generation (Graham 1993). In Charales, tube cells undergo extension at the same time as the egg cell enlarges, such that mature eggs are thought to be near the maximal size reached by zygotes, and fully enclosed before fertilization occurs (Pickett-Heaps 1975). In contrast, in *Coleochaete*, the enclosing layer of parental cells does not develop until after fertilization, and zygotes (rather than eggs) undergo enlargement and storage accumulation. In both cases, as yet undefined cell–cell signalling processes probably coordinate development.

Only in *Coleochaete* could initial development of corticating cells be said to result from intergenerational zygote/parental thallus) communication, but zygote/parental cell signalling that influences maturation of the n/n cell complex may occur in both *Coleochaete* and Charales. Hence these are the taxa of choice for analysis of hypothesized communication systems, diffusible signalling molecules and their receptors. Some *Coleochaete* species produce zygotes that are less completely corticated than others (Szymńska 1989); comparison of expression patterns at critical developmental stages might reveal genetic differences relevant to the zygote retention issue. Eggs are reportedly not retained in *Chaetosphaeridium* (Thompson 1969); if *Chaetosphaeridium* is sister to *Coleochaete*, differential expression studies of these taxa could be very informative.

Molecules known to have signalling functions in higher plants that could also operate in charophyceans and early-divergent bryophytes include hydrogen peroxide, a diffusible and relatively long-lived molecule whose production is in part regulated by peroxisomal catalase (Karpinski *et al.* 1999); secreted peptides or small proteins that act as hormones (Gehring 1999; Fletcher *et al.* 1999); and sugars (Koch 1996; Graham, *et al.* 1996). The role of sugars is discussed below in more detail.

(c) Stage C

A third postulated step in origin of the life history of embryophytes (figure 5) is the origin of matrotrophy, transfer of nutrients and/or morphogenetic factors from parental tissues to zygotes. Such a step is dependent on the earlier evolution of retained eggs/zygotes (see above), because diffusive or degradative loss of parentally secreted nutrients or other compounds would greatly reduce their supply to progeny (zygotes) that are spatially separate, particularly in an aquatic environment. Contiguity of *Coleochaete* parental cells and zygotes, localized development of elaborate ingrowths on the walls of cortical cells that are in direct contact with zygotes, and conspicuous post-fertilization nutrient storage (in the form of starch and lipid) provide circumstantial evidence for occurrence of matrotrophy. Close spatial association of parental tissues whose cells possess typical placental transfer morphology (wall ingrowths) and physiological evidence for solute transport across the apoplastic, inter-generation junction (Browning & Gunning 1979a,b; Renault *et al.* 1992) provide evidence that matrotrophy occurs in bryophytes. These data suggest that embryophytes could have acquired matrotrophy from charophycean ancestors, and that matrotrophy preceded the origin of multicellular sporophytes. Molecular strategies for testing these possibilities are discussed in a later section of this paper.

(d) Stage D

Transition from a unicellular zygote (as produced by charophyceans) to production of a multicellular diploid sporophyte generation (as in embryophytes) by repeated mitotic division of the zygote is the next postulated step in the origin of land plant alternation of generations (figure 5). The mechanism most often hypothesized for this transformation is ‘delay in meiosis’. This means that a phase of mitotic proliferative growth would occur between zygote formation and spore production by meiosis. An alternative idea, that the land plant sporophyte originated in green algal ancestors that had first acquired alternation of multicellular generations, is currently less favoured, as it is not supported by modern phylogenetic analysis. Extant green algae that exhibit alternation of generations are not closely related to the ancestry of land plants, and charophycean green algae, which are closely related to embryophytes, lack alternation of generations, so far as is known. Hypothetical origin of embryophytes from green algae having temporally and spatially separate generations also does not provide a satisfactory explanation for the origin of matrotrophy.

The finding of an extant or fossil ‘charophycean’ that had intercalated even a single mitotic division between syngamy and meiosis would mean discovery of a possible transitional form, a descendant of such a transitional form or a close parallel to the simplest possible (two-celled) land plant sporophyte. The adaptive advantage of having a multicellular sporophyte is that greater numbers of genetically diverse meiospores could result, facilitating both colonization effectiveness and increase in population genetic variability, and hence greater evolutionary flexibility. Evidence that such adaptive advantage exists is provided by progressive increase in sporophyte size during the course of embryophyte evolution.

Reconstruction of the critical molecular events involved in the origin of a delay of meiosis may be possible by probing the genomes of extant charophyceans and early-divergent embryophytes for genes that are involved in the regulation of meiotic initiation. Such genes have been identified in metazoans (e.g. Barton & Kimble 1990; de Vries *et al.* 1999), yeasts (e.g. Englebrecht & Roeder 1990; Matsura *et al.* 1990; Printz *et al.* 1995; Iino *et al.* 1995; Colomina *et al.* 1999; de los Santos & Hollingsworth 1999; Edlmann *et al.* 1999; Ohno & Mattaj 1999; Sagee *et al.* 1999), and flowering plants (e.g. Bouchard 1990; Bai *et al.* 1999); a recent review of meiotic chromosome organization and segregation in plants is provided by Dawe (1998). Given such information, it may ultimately be possible to transform charophyceans so that their zygotes recapitulate the meiotic delay postulated to have occurred during the origin of embryophytes.

The origin of the quadripolar microtubular system (QMS) characteristic of embryophytic sporocytes, which provides an essential scaffolding for proper spatial segregation of plastids and nuclei into the four daughter products (Brown & Lemmon 1997), is unknown. Whether this structural component of meiotic cells is present in an earlier form in charophyceans is not known, and would require ultrastructural and fluorescence immunolocalization studies of germinating zygotes. Correlative analysis of nuclear DNA levels would also be valuable but has been difficult because large amounts of storage materials, autofluorescent sporopollenin-like zygote wall layers and autofluorescent cortical cell walls (in the case of *Coleochaete*) can interfere.

Microarray studies in yeast systems have revealed that ploidy change regulates the expression of genes involved in the cell cycle (such as G1 cyclins) and actin control of polarized growth. However, the mechanism by which ploidy change is sensed is not as yet understood (Galitski *et al.* 1999). Since homologous cell cycle and actin-regulation genes are likely to occur in plants, it is possible that ploidy changes occurring at the transitions between gametophytic and sporophytic phases of the life history are similarly important in determining differences in the morphology of the two stages.

(e) *Stage E*

A final step, from production of non-walled, flagellate meiospores, as occurs in modern *Coleochaete*, to production of nonflagellate, sporopollenin-walled meiospores, as occurs in all embryophyte groups, is also postulated to have been involved in the origin of plant alternation of generations (figure 5). This is because the relative efficiency of walled-spore dispersal and germination success on land may have driven evolutionary transition to increasingly larger multicellular sporophytes capable of producing many meiospores from a single fertilization event. The alternative, illustrated by charaleans, is production of many large, resistant zygotes (representing multiple fertilization events), each of which produces probably only one, non-resistant, meiotic product. Such a reproductive strategy would not be adaptive if fertilization frequency is limited by water availability. If spores lacking protective walls were unable to survive to germination following terrestrial dispersal, the multicellular sporophyte likewise would not be adaptive. The

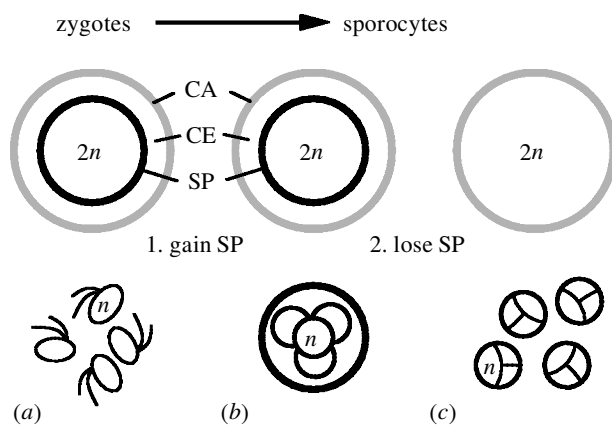


Figure 6. Hypothetical transformations involved in the origin of walled, non-flagellate spores, as they occur in modern embryophytes by gain of sporopollenin (SP) in spore walls (1), then loss of sporopollenin from sporocyte walls (2).

(a) The extant charophycean *Coleochaete* produces zygotes having callosic (CA), cellulosic (CE) and SP layers, within which develop flagellate, non-walled meiospores. (b) A hypothetical intermediate stage in which diploid sporocytes (that are homologous to charophycean zygotes) have resistant walls, and produce walled, non-flagellate meiospores. Persistence of the resistant sporocyte wall forms an envelope around meiospore tetrads, preventing their dissociation. (c) Sporopollenin-walled, dissociated meiospores as they occur in most embryophytes.

sporopollenin-coated wall of embryophytic spores has been regarded as having an adaptive function in the terrestrial environment, by providing structural stability and retarding microbial degradation during dispersal (Graham & Gray 2001).

It is as yet unclear whether multicellular sporophytes appeared prior to the origin of walled meiospores, or vice versa. It is possible that earliest embryophytes (defined by presence of a multicellular sporophyte, however small) might have lived in water or very moist environments in which selection did not operate heavily against unwalled, flagellate meiospores. Production of greater numbers of or more genetically diverse meiospores could have provided sufficient adaptive advantage for delay in meiosis to have occurred prior to origin of walled spores. Amplification (apparently by reduplication) of zygotic DNA levels and subsequent production of 8–32 meiospores in *Coleochaete* (Hopkins & McBride 1976) provides evidence of the adaptive value that a multicellular sporophyte could have in nearshore waters.

An inner wall layer of sporopollenin-like material is typically produced during zygote maturation in charophyceans. The transition to walled meiospores is postulated to have involved a change in the timing of sporopollenin production, such that it occurs at a later developmental stage, during spore maturation (figure 6). Transitional forms may have produced sporopollenin both at the spore mother cell stage (hypothesized to be homologous to charophycean zygotes by Graham 1990) and during spore maturation. Information regarding regulation of sporopollenin synthesis and deposition derived from analysis of higher plant mutants may be useful in deducing the genetic transformations that were involved in the production of sporopollenin-walled spores.

4. APPROACHES TO AN UNDERSTANDING OF THE EVOLUTIONARY ORIGIN OF MATROTROPHY AND PLACENTAL FUNCTION IN EMBRYOPHYTES

Combined neontological and palaeontological approaches, and comparative studies of extant charophyceans and bryophytes are recommended for elucidation of the evolutionary process surrounding the origin of embryophytic matrotrophy and placental function. An example of a combined neontological/palaeontological approach is the comparison of acid-hydrolysis resistant, autofluorescent mature placentae of *Coleochaete orbicularis* (the only charophycean known to produce placental transfer-like cells) (figure 4) and those of bryophytes (Kroken *et al.* 1996). Analogously, morphology of the placental interface of the extant pteridophyte *Tmesipteris elongata* was used by Frey *et al.* (1997) to postulate the occurrence of placentae in *Lorneophyton lignieri*, based on fossils that were originally published by Taylor & Taylor (1993).

Deposition of resistant polymers into the walls of placental cells occurs during late development of zygotes (*Coleochaete*) or sporophytes (bryophytes). These materials are not present (as judged by absence of specific wall auto-fluorescence and lack of resistance to high-temperature acid hydrolysis) in vegetative thalli or during early zygote development, including the period of presumed solute transport. Thus deposition appears to be under regulatory control. The function of resistant wall compounds is not proven but could include shutting off solute flow or protection from microbial attack (Kroken *et al.* 1996; Graham, L. E. 1996). Wall ingrowths of *Coleochaete* are also heavily impregnated and consequently resist acid hydrolysis. As a result, cell wall ingrowths, which represent a physical manifestation of matrotrophy, are readily visible at the light microscopic level, both in bright field (figure 4) and UV-V fluorescence excitation. This suggests that if ancestral charophyceans related to *Coleochaete*—organisms transitional between charophyceans and earliest embryophytes—or early embryophytes also possessed such features, as is likely, they should have left resistant remains in the form of distinctive microfossils. Thus it is possible that microfossils younger than those presently known may yet be found that will be informative regarding the earliest events in the history of embryophytes and their acquisition of alternation of generations. We are currently cataloguing high-temperature, acid-hydrolysed fertile thalli of various *coleochaete* species and placentae of modern bryophytes with the hope that they may provide useful search images for palaeontologists as they seek remains of earliest embryophytes or fossil evidence of matrotrophy.

Neontological approaches include comparative analyses of putative nutrient transport from maternal cells to zygotes of *Coleochaete* and between gametophytic and sporophytic tissues of early-divergent embryophytes. We have focused on comparative study of the hexose transporter gene family in charophyceans and early-divergent embryophytes (described in more detail below). Parallel efforts could be made to trace evolutionary change in amino acid transporters (Fischer *et al.* 1998) and H⁺-ATPases. Genetic analysis of the regulation and biosynthesis of wall-bound resistant polymers that characterize mature charophycean and bryophyte placentae is also

desirable. It may be possible to determine if parental imprinting affects allocation of resources at the plant placenta (reflecting intragenomic, i.e. paternal–maternal, conflict), as is believed to occur in eutherians (Haig 1996a,b, 1997) and, if so, when this might have evolved.

5. COMPARATIVE STUDY OF CHAROPHYCEAN/ EMBRYOPHYTE HEXOSE TRANSPORTER GENES AND PROTEINS

Physiological studies by Renault *et al.* (1992) in the early-divergent moss *Polytrichum* strongly implicate hexose sugars (rather than sucrose) as the major carbohydrate species that is moved from gametophyte to sporophyte. Because the intergenerational junction is devoid of symplastic connections (plasmodesmata), cell membrane hexose transporter proteins are implicated. Such transporters are thought to facilitate diffusion and/or actively cotransport sugars and protons; homologous proteins in bacteria, fungi, protists, higher plants and higher animals are members of a major family of transmembrane facilitators (Marger & Saier 1993).

It is postulated that glucose, once imported into zygotes/sporophytes, is transformed into starch or lipid storage, or used more immediately for embryo and sporophyte development and spore production. Both processes act as sinks that would tend to promote increased apoplastic transport from gametophytic sources. In higher plants, sugars not only serve as energy sources and building blocks for structural elements such as cellulose but are also potent regulatory molecules. Sugars function as signalling molecules that activate or repress genes involved in cell cycle regulation, photosynthesis and pigment production, glyoxylate metabolism, respiration, starch and sucrose synthesis and degradation, nitrogen metabolism, pathogen defence and wounding responses, and senescence (Ehness *et al.* 1997; Koch 1996; Graham, I. A. 1996; Truernit *et al.* 1996). For example, chlorophyll *a/b* binding protein genes and *rbcS* genes are sugar-repressible, whereas the gene encoding nitrate reductase is sugar-inducible. Further, genes essential for biosynthesis of ethylene are activated by sugars, and those for brassinolides are repressed. Receptors are hypothesized to be integral cell membrane hexose transporter proteins (Lalond *et al.* 1999).

Plants also possess sugar sensors. These are molecules that detect the presence of imported sugars and transduce the signalling responses. The sugar phosphorylation enzyme hexokinase (HXK) (which also functions as an early step of cytoplasmic glycolysis) is one example of a sugar sensor, because the sugar responses of higher plants require only that imported sugar is phosphorylated by this enzyme (Jang *et al.* 1997). It is thought that the signalling HXK is spatially associated with membrane hexose transporters (Smeekens & Rook 1997). HXK is highly conserved from bacteria to yeast and higher plants and animals, and is thus likely to occur in charophyceans and bryophytes, and to have similar sensing functions in these organisms.

We began our analysis of hexose transport by determining that at least some charophycean algae and bryophytes (e.g. *Coleochaete*, Graham *et al.* 1994; the charophycean desmid *Closterium*, Lewitus & Kana 1994;

and several *Sphagnum* moss species, P. Bilkey, unpublished data) are capable of importing and using exogenous hexose sugars, as deduced by growth experiments. This was necessary because a number of green algae are known to be incapable of utilizing hexose sugars. For example, all of the (chlorophycean) *Chlorococcum* species tested by Parker *et al.* (1961) were incapable of utilizing hexose sugars for growth. The model-system organism *Chlamydomonas reinhardtii* (also chlorophycean) is likewise unable to utilize exogenous sugars (Harris 1989). Forsberg (1965) concluded that axenic cultures of *Chara* were also unable to utilize exogenous sugars; however, Smith (1967) obtained evidence that *Nitella translucens* can take up and metabolize ¹⁴C-labelled glucose.

We found that hexose utilization by charophyceans and bryophytes increased under conditions in which dissolved inorganic carbon (DIC) was limiting to photosynthesis. This suggests a likely adaptive advantage of hexose-transport expression in vegetative cells or tissues. We then hypothesized that hexose transporter genes were brought under increased regulation in derived charophyceans having retained zygotes, such that hexose sugars can be moved from green cells of the parental generation to zygote storages, even under conditions when DIC is not limiting to vegetative growth. *Coleochaete* produces greater numbers of meiospores than any other green alga lacking alternation of generations, and we hypothesize that matrotrophy explains this difference. We suggest that matrotrophy was inherited by embryophytes from ancestral charophyceans and used as a means of supporting amplified spore production, and in aspects of sporophyte morphogenesis.

Because rapid diversification of tissue-specific members of gene families is involved in the origin of functional innovations and evolutionary radiation in other organismal groups (Iwabe *et al.* 1996; Henikoff *et al.* 1997), we are exploring the possibility that gene duplication, divergence and tissue-specific expression play important roles in the evolutionary origin of plant matrotrophy. We are currently testing these hypotheses by cataloguing sequences of hexose-transporter gene family members in selected charophyceans and bryophytes, then using the information to study expression patterns. We postulate that specific gene family members may occur only in charophyceans having retained zygotes, and that related, placenta-specific gene family members may occur in bryophytes and other embryophytes.

Hexose transporter genes of the green alga *Chlorella* Sauer & Tanner 1993) and *Arabidopsis* (as well as other flowering plants) have been well characterized at the molecular level, allowing identification of conserved regions. We have used such genetic database information to design primers for use in polymerase chain reaction amplification, cloning and sequencing of portions of several members of hexose transporter gene families in a number of charophyceans and bryophytes. We were successful in obtaining multiple, partial, putative hexose-transporter gene sequences from all three mosses investigated, *Sphagnum*, *Andreobryum* and *Mnium*, using the forward primer 5'-GATGGTACCGGATCCTTYTTYTARARYTIACIGGIATHAA-3' and the reverse primer 5'-GATCTGCAGTCGACTCDATIGGIAYICCYTTIGTYTCIGG-3'. 5' portions of the primers contain restriction sites to facilitate directional cloning.

Partial sequences of similar genes were also obtained from the hornwort *Megaceros*, the liverwort *Conocephalum* and several charophycean green algae with the same primers, but amplification product was not consistently obtained for liverworts and charophyceans. Evidence that our primers amplify portions of hexose transporter proteins includes high similarity in primary sequence data and in deduced hydropathy profiles of charophycean and bryophyte/amino acid sequences inferred from our DNA sequences and those of homologous regions of *Chlorella* HUPI (hexose uptake protein 1), and *Arabidopsis* STP 1 (sugar transporter protein 1).

In conclusion, it appears that the use of (i) neontological approaches such as molecular analysis of matrotrophy in charophyceans and bryophytes, and (ii) combined neontological/palaeontological approaches in which resistant morphology provides links between extant charophytes and bryophytes and microfossils, are likely to illuminate the evolutionary origin of the embryophytic life cycle to a degree previously thought impossible. Such studies provide not only insight into evolutionary mechanism, but also have implications for understanding the effects of earliest plants on Ordovician biogeochemistry and may be useful in deducing how the complex regulatory systems operating in higher plant reproductive development have been constructed over evolutionary time.

We are grateful to Colleen Lavin and the Integrated Microscopy Resource at the University of Wisconsin-Madison for assistance with high-pressure freezing techniques. John Raven and Dianne Edwards very kindly reviewed the manuscript and offered helpful suggestions. We also acknowledge National Science Foundation (USA) grant DEB 9628869, our colleagues Madeline Fisher, Evan Lau and Robin Kodner for assistance with the hexose transporter sequencing work, and Brent Mishler and Louis Lewis for providing moss and liverwort DNA.

REFERENCES

- Bai, X., Peirson, B. N., Dong, F., Xue, C. & Makaroff, C. 1999 Isolation and characterization of *SYN1*, a *RAD*-like gene essential for meiosis in *Arabidopsis*. *Pl. Cell* **11**, 417–430.
- Baillard, L. 1997 Is morphological alternation of generations in sexual reproduction a product of chronobiology? *Phytomorphology* **47**, 395–399.
- Barton, M. K. & Kimble, J. 1990 *Fog-1*, a regulatory gene required for specification of spermatogenesis in the germ line of *Caenorhabditis elegans*. *Genetics* **125**, 29–40.
- Bell, G. 1997 The evolution of the life cycle of brown seaweeds. *Biol. J. Linn. Soc.* **60**, 21–35.
- Bell, G. & Koufopanou, V. 1991 The architecture of the life cycle of small organisms. *Proc. R. Soc. Lond.* **B332**, 81–90.
- Bell, P. R. 1991 The life cycles of cryptogams. *Acta Bot. Malacitana* **16**, 5–18.
- Bouchard, R. A. 1990 Characterization of expressed meiotic prophase repeat transcript clones of *Lilium*: meiosis-specific expression, relatedness, and affinities to small heat shock protein genes. *Genome* **33**, 68–79.
- Brown, R. C. & Lemmon, B. E. 1997 The quadripolar microtubule system in lower land plants. *J. Pl. Res.* **110**, 93–106.
- Browning, A. J. & Gunning, B. E. S. 1979a Structure and function of transfer cells in the sporophyte haustorium of *Funaria hygrometrica* Hedw. I. The development and ultrastructure of the haustorium. *J. Exp. Bot.* **30**, 1233–1246.

- rowning, A. J. & Gunning B. E. S. 1979b Structure and function of transfer cells in the sporophyte haustorium of *Funaria hygrometrica* Hedw. II. Kinetics of uptake of labeled sugars and localization of absorbed products by freeze-substitution and autoradiography. *J. Exp. Bot.* **30**, 1247–1264.
- Shapman, R. L. (and 10 others) 1998 Molecular systematics of the green algae. In *Molecular systematics of plants. II* (ed. R. S. Soltis, D. E. Soltis & J. J. Doyle), pp. 508–540. Boston, MA: Kluwer Academic Publishers.
- Solomina, N., Gari, E., Gallego, C., Herrero, E. & Aldea, M. 1999 G1 cyclins block Imel pathway to make mitosis and meiosis incompatible in budding yeast. *EMBO J.* **18**, 320–329.
- Stawe, R. K. 1998 Meiotic chromosome organization and segregation in plants. *A. Rev. Pl. Physiol. Pl. Mol. Biol.* **49**, 371–395.
- de los Santos, T. & Hollingsworth, N. M. 1999 Red1p, a MEK1-dependent phosphoprotein that physically interacts with Hop1 during meiosis in yeast. *J. Biol. Chem.* **274**, 1783–1790.
- de Vries, S., Baart, E. B., Dekker, M., Siezen, A., de Rootig, D. G., de Boer, P. & te Riele, H. 1999 Mouse MutS-like protein Msh5 is required for proper chromosome synapsis in male and female meiosis. *Genes Dev.* **13**, 523–553.
- Delwiche, C. F., Graham, L. E. & Thomson, N. 1989 Lignin-like compounds and sporopollenin in *Coleochaete*, an algal model for land plant ancestry. *Science* **245**, 399–401.
- Duncan, T. M., Renzaglia, K. S. & Garbary, D. J. 1997 Ultrastructure and phylogeny of the spermatozoid of *Chara vulgaris* (Charophyceae). *Pl. Syst. Evol.* **204**, 125–140.
- Edelmann, E., Cohen, P. E., Kneitz, B., Winand, N., Lia, M., Heyer, J., Kolodner, R., Pollard, J. W. & Kucherlapati, R. 1999 Mammalian MutS homologue 5 is required for chromosome pairing in meiosis. *Nature Genet.* **21**, 123–127.
- Ehness, R., Ecker, M., Godt, D. R. & Roitsch, T. 1997 Glucose and stress independently regulate source and sink metabolism and defense mechanisms via signal transduction pathways involving protein phosphorylation. *Pl. Cell* **9**, 1825–1841.
- Engelbrecht, J. & Roeder, G. S. 1990 *MERI*, a yeast gene required for chromosome pairing and genetic recombination, is induced in meiosis. *Mol. Cell. Biol.* **10**, 2379–2389.
- Fischer, W.-N., André, B., Rentsch, D., Krolkiewicz, S., Tageder, M., Breitkraud, K. & Frommer, W. B. 1998 Amino acid transport in plants. *Trends Pl. Sci.* **3**, 188–195.
- Fletcher, J. C., Brand, U., Running, M. P., Simon, R. & Meyerowitz, E. M. 1999 Signaling of cell fate decisions by *CLAVATA3* in *Arabidopsis* shoot meristems. *Science* **283**, 1911–1914.
- Forsberg, C. 1965 Nutritional studies of *Chara* in axenic cultures. *Physiologia Pl.* **18**, 275–290.
- Frederick, W., Hofmann, M. & Hilger, H. H. 1997 Gametophyte-sporophyte junction in the Lower Devonian plant *Horneophyton lignieri*. *Nova Hedwigia* **64**, 549–552.
- Galitski, T., Saldanha, A. J., Styles, C. A., Lander, E. S. & Fink, G. R. 1999 Ploidy regulation of gene expression. *Science* **285**, 251–254.
- Gehring, G. 1999 Natriuretic peptides—a new class of plant hormone? *Ann. Bot.* **83**, 329–334.
- Graham, I. A. 1996 Carbohydrate control of gene expression in higher plants. *Res. Microbiol.* **147**, 572–580.
- Graham, L. E. 1990 Meiospore formation in charophycean algae. In *Microspores: evolution and ontogeny* (ed. S. Blackmore & R. B. Knox), pp. 43–54. London: Academic Press.
- Graham, L. E. 1993 *The origin of land plants*. New York: Wiley & Sons.
- Graham, L. E. 1996 Green algae to land plants: an evolutionary transition. *J. Pl. Res.* **109**, 241–251.
- Graham, L. E. & Gray, J. 2001 The origin, morphology and ecophysiology of early embryophytes: neontological and palaeontological perspectives. In *Plants invade the land: evolution and environmental perspectives* (ed. D. Edwards & P. Gensel). Columbia University Press. (In the press.)
- Graham, L. E. & Wilcox, L. W. 2000 *Algae*. Upper Saddle River, NJ: Prentice Hall.
- Graham, L. E., Graham, J. M., Russin, W. R. & Chesnick, J. M. 1994 Occurrence and phylogenetic significance of glucose utilization by charophycean algae: glucose enhancement of growth in *Coleochaete orbicularis*. *Am. J. Bot.* **81**, 423–432.
- Gray, J., Massa, D. & Boucot, A. J. 1982 Caradocian land plant microfossils from Libya. *Geology* **10**, 197–201.
- Gunnison, D. & Alexander, M. 1975a Resistance and susceptibility to decomposition by natural microbial communities. *Limnol. Oceanogr.* **20**, 64–70.
- Gunnison, D. & Alexander, M. 1975b Basis for the resistance of several algae to microbial decomposition. *Appl. Microbiol.* **29**, 729–738.
- Haig, D. 1996a Placental hormones, genomic imprinting, and maternal-fetal communication. *J. Evol. Biol.* **9**, 357–380.
- Haig, D. 1996b Gestational drive and the green-bearded placenta. *Proc. Natl Acad. Sci. USA* **93**, 6547–6551.
- Haig, D. 1997 Parental antagonism, relatedness asymmetries, and genomic imprinting. *Proc. R. Soc. Lond. B* **264**, 1657–1662.
- Harada, J. J., Lotan, T., Fischer, R. L. & Goldberg, R. B. 1998 Embryos without sex. *Trends Pl. Sci.* **3**, 452–453.
- Harris, E. 1989 *The Chlamydomonas handbook*. New York: Academic Press.
- Henikoff, S., Greene, E. A., Pietrokovski, S., Bork, R., Attwood, T. K. & Hood, L. 1997 Gene families: the taxonomy of protein paralogs and chimeras. *Science* **278**, 609–614.
- Hopkins, A. W. & McBride, G. E. 1976 The life history of *Coleochaete scutata* (Chlorophyceae) studied by a Feulgen microspectrophotometric analysis. *J. Phycol.* **12**, 29–35.
- Huss, V. A. R. & Kranz, H. D. 1997 Charophyte evolution and the origin of land plants. In *Origins of algae and their plastids* (ed. D. Bhattacharya), pp. 103–114. Vienna: Springer.
- Iino, Y., Hiramane, Y. & Yamamoto, M. 1995 The role of *cdc2* and other genes in meiosis in *Schizosaccharomyces pombe*. *Genetics* **140**, 1235–1245.
- Iwabe, N., Kumar, K. & Miyata, T. 1996 Evolution of gene families and relationship with organismal evolution. Rapid divergence of tissue-specific genes in the early evolution of chordates. *Mol. Biol. Evol.* **13**, 483–493.
- Jang, J. C., León, P., Zhou, L. & Sheen, J. 1997 Hexokinase as a sugar sensor in higher plants. *Pl. Cell* **9**, 5–19.
- Karpinski, S., Reynolds, H., Karpinska, B., Wingsle, G., Creissen, G. & Mullineaux, R. 1999 Systemic signaling and acclimation in response to excess excitation energy in *Arabidopsis*. *Science* **284**, 654–657.
- Kenrick, P. 1994 Alternation of generations in land plants: new phylogenetic information and palaeobotanical evidence. *Biol. Rev.* **69**, 293–330.
- Kenrick, P. & Crane, P. R. 1997 *The origin and early diversification of land plants: a cladistic study*. Smithsonian University Press.
- Koch, K. E. 1996 Carbohydrate-modulated gene expression in plants. *A. Rev. Pl. Physiol. Pl. Mol. Biol.* **47**, 509–540.
- Kroken, S. B., Graham, L. E. & Cook, M. E. 1996 Occurrence and evolutionary significance of resistant cell walls in charophytes and bryophytes. *Am. J. Bot.* **83**, 1241–1254.
- Kuhlenkamp, R., Mueller-Dieter, G. & Whittick, A. 1993 Genotypic variation and alternating DNA levels at constant chromosome numbers in the life history of the

- brown alga *Haplospora globosa* (Tilopteridales). *J. Phycol.* **29**, 377–380.
- alond, S., Boles, E., Hellman, H., Barker, L., Patrick, J. W., Frommer, W. B. & Ward, J. M. 1999 The dual function of sugar carriers: transport and sugar sensing. *Pl. Cell* **11**, 707–726.
- ewis, L. A., Mishler, B. D. & Vilgalys, R. 1997 Phylogenetic relationships of the liverworts (Hepaticae), a basal embryophyte lineage, inferred from nucleotide sequence data of the chloroplast gene *rbcL*. *Mol. Phylogenet. Evol.* **7**, 377–393.
- ewitus, A. J. & Kana, T. M. 1994 Responses of estuarine phytoplankton to exogenous glucose stimulation versus inhibition of photosynthesis and respiration. *Limnol. Oceanogr.* **39**, 182–189.
- igrone, R. & Gambardella, R. 1988 The sporophyte–gametophyte junction in bryophytes. *Adv. Bryol.* **3**, 225–274.
- igrone, R., Duckett, J. G. & Renzaglia, K. S. 1993 The gametophyte–sporophyte junction in land plants. *Adv. Bot. Res.* **19**, 232–317.
- otan, T., Ohto, M. A., Yee, K. M., West, M. A. L., Lo, R., Kwong, R. W., Yamagishi, K., Fischer, R. L., Goldberg, R. B., & Harada, J. J. 1998 *Arabidopsis* LEAFY/COTYLEDON 1 is sufficient to induce embryo development in vegetative cells. *Cell* **93**, 1195–1205.
- icCourt, R. M., Karol, K. G., Guerlesquin, M. & Feist, M. 1996 Phylogeny of extant genera in the family Characeae (Charales, Charophyceae) based on *rbcL* sequences and morphology. *Am. J. Bot.* **83**, 71–77.
- arger, M. D. & Saier Jr, M. H. 1993 A major superfamily of transmembrane facilitators that catalyze uniport, symport, and antiport. *Trends Biochem. Sci.* **18**, 13–20.
- atsura, A., Treinin, M., Mitsuzawa, H., Kassir, Y., Uno, I. & Simchen, G. 1990 The adenylate cyclase–protein kinase cascade regulates entry into meiosis in *Saccharomyces cerevisiae* through the gene *IME1*. *EMBO J.* **9**, 3225–3232.
- urgia, M., Hueng, B.-Q., Tucker, S. C. & Musgrave, M. E. 1993 Embryo sac lacking antipodal cells in *Arabidopsis thaliana* (Brassicaceae). *Am. J. Bot.* **80**, 824–838.
- hno, M., & Mattaj, I. N. 1999 Meiosis: MeiRNA hits the spot. *Curr. Biol.* **9**, R66–R69.
- otto, S. P. & Goldstein, B. 1992 Recombination and the evolution of diploidy. *Genetics* **131**, 745–751.
- arker, B. C., Bold, H. C. & Deason, T. R. 1961 Facultative heterotrophy in some chlorococcacean algae. *Science* **133**, 761–763.
- ickett-Heaps, J. D. 1975 *Green algae. Structure, reproduction and evolution in selected genera*. Sunderland, MA: Sinauer.
- rintz, S., Klein, F., Auer, H., Schweitzer, D. & Primig, M. 1995 DNA-binding factor (UBF) interacts with a positive regulatory element in the promoters of genes expressed during meiosis and vegetative growth in yeast. *Nucl. Acids Res.* **23**, 3449–3456.
- iu, Y.-L. & Palmer, J. D. 1999 Phylogeny of early land plants: Insights from genes and genomes. *Trends Pl. Sci.* **4**, 26–29.
- aven, P., Evert, R. F. & Eichhorn, S. E. 1999 *Biology of plants*. New York: Worth.
- enault, S., Bonnemain, J. L., Faye, L. & Gaudillere, J. P. 1992 Physiological aspects of sugar exchange between the gametophyte and the sporophyte of *Polytrichum formosum*. *Pl. Physiol.* **100**, 1815–1822.
- agee, S., Sherman, A., Shenhar, G., Robzyk, K., Ben, D. N., Simchen, G. & Kassir, Y. 1999 Multiple and distinct activation and repression sequences mediate the regulated transcription of *IME1*, a transcriptional activator of meiosis-specific genes in *Saccharomyces cerevisiae*. *Mol. Cell. Biol.* **18**, 1985–1995.
- Sauer, N. & Tanner, W. 1993 Molecular biology of sugar transporters in plants. *Bot. Acta* **196**, 277–286.
- Searles, R. B. 1980 The strategy of the red algal life history. *Am. Nat.* **115**, 113–120.
- Sekimoto, H., Inoki, Y. & Fuji, T. 1993 Detection and evaluation of an inducer of a diffusible mating pheromone of the heterothallic *Closterium peracerosum–strigosum–littorale* complex. *Pl. Cell Physiol.* **34**, 991–996.
- Sluiman, H. J. & Guihal, C. 1999 Phylogenetic position of *Chaetosphaeridium* (Chlorophyta), a basal lineage in the Charophyceae inferred from 18S rDNA sequence. *J. Phycol.* **35**, 395–402.
- Smeeckens, S. & Rook, F. 1997 Sugar sensing and sugar-mediated signal transduction in plants. *Pl. Physiol.* **115**, 7–13.
- Smith, F. A. 1967 Links between glucose uptake and metabolism in *Nitella translucens*. *J. Exp. Bot.* **18**, 348–358.
- Strother, P. K., Al Hajri, S. & Traverse, A. 1996 New evidence for land plants from the lower Middle Ordovician of Saudi Arabia. *Geology* **24**, 55–58.
- Szymánska, M. 1989 Three new *Coleochaete* species (Chlorophyta) from Poland. *Nova Hedwigia* **49**, 435–446.
- Taylor, T. N. & Taylor, E. L. 1993 *The biology and evolution of fossil plants*. Englewood Cliffs, NJ: Prentice Hall.
- Taylor, W. A. 1995 Ultrastructure of *Tetrahedralites medinesis* (Strother and Traverse) Wellman and Richardson, from the Upper Ordovician of southern Ohio. *Rev. Palaeobot. Palynol.* **85**, 183–187.
- Thompson, R. H. 1969 Sexual reproduction in *Chaetosphaeridium globosum* (Nordst.) Klebahn (Chlorophyceae) and description of a species new to science. *J. Phycol.* **5**, 285–290.
- Truernit, E., Schmid, J., Epple, P., Illig, J. & Sauer, N. 1996 The sink-specific and stress-regulated *Arabidopsis STP4* gene: enhanced expression of a gene encoding a monosaccharide transporter by wounding, elicitors, and pathogen challenge. *Pl. Cell* **8**, 2169–2182.
- Van Caesele, L., Klingner, B. & Sumner, M. J. 1996 The immunolocalization of plasma membrane H⁺-ATPase in the transfer cell region of *Brassica napus* (Brassicaceae) ovules. *Am. J. Bot.* **83**, 1386–1390.

Discussion

P. Kenrick (*Department of Palaeontology, Natural History Museum, London*). You suggest that fecundity was a major driving force behind the origin of the sporophyte generation in land plants: rarity of fertilization on land is compensated by the multiplication of the products of fertilization (spores) made possible by the development of sporophytes. Can similar arguments be made for the development of sporophytes in other algal groups (e.g. Ulvales)?

L. K. E. Graham. The most comparable situation occurs in red algae, where there is a parallel temporal–spatial association between the sporophyte (carposporophyte) and female gametophyte. At least theoretically, the frequency of red algal fertilization may be limited by the absence of flagella from male gametes. It has been argued (Searles 1980) that fecundity was the major driving force in the origin of the carposporophyte of red algae; origin of the land plant sporophyte was cited as a parallel example. Some physiological and anatomical work on red algal female gametophyte–carposporophyte associations (Hommersand & Fredericq 1990) seems to support Searles’s conjecture.

The hypothesis may reasonably be extended to Ulvales, though intimate association of sporophytes and gametophytes is lacking. Multicellular ulvacean sporophytes can clearly produce greater numbers of meiospores than can resumed ancestral forms (modelled by Ulotrichales whose diploid stage is unicellular), and this could be advantageous in the marine habitat (Graham & Wilcox 2000). It is possible that mating frequency in ulvaceans could be limited by the ability of gametes to locate each

other in a big ocean, but few studies have focused on mating success rates in algae. Such data could be helpful in testing the hypothesis.

Reference

- Hommersand, M. S. & Fredericq, S. 1990 Sexual reproduction and cystocarp development. In *Biology of the red algae* (ed. K. M. Cole & R. G. Sheath), pp. 305-346. Cambridge University Press.