

Development and physiology of the brown alga Ectocarpus siliculosus: two centuries of research

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Development and physiology of the brown alga Ectocarpus siliculosus: two centuries of research

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Summary

Brown algae share several important features with land plants, such as their photoautotrophic nature and their cellulose-containing wall, but the two groups are distantly related from an evolutionary point of view. The heterokont phylum, to which the brown algae belong, is a eukaryotic crown group that is phylogenetically distinct not only from the green lineage, but also from the red algae and the opisthokont phylum (fungi and animals). As a result of this independent evolutionary history, the brown algae exhibit many novel features and, moreover, have evolved complex multicellular development independently of the other major groups mentioned above. In 2004, a consortium of laboratories, including the Station Biologique in Roscoff and Genoscope, initiated a project to sequence the genome of *Ectocarpus siliculosus*, a small filamentous brown alga that is found in temperate, coastal environments throughout the globe. The *E. siliculosus* genome, which is currently being annotated, is expected to be the first completely characterised genome of a multicellular alga. In this review we look back over two centuries of work on this brown alga and highlight the advances that have led to the choice of *E. siliculosus* as a genomic and genetic model organism for the brown algae.

198 words

1	Keywords
2	Brown alga; phaeophyceae; <i>Ectocarpus</i> ; development; reproduction; pheromone;
3	metabolism; defense
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8	Abbreviations:
9	ARA, arachidonic acid; BL, blue light; CE: chloroplast envelope; CER: chloroplast
10	endoplasmic reticulum; Ci, inorganic carbon; EPA, eicosapentaenoic acid; DGDG,
11	digalastosyldiacylglycerol; DPG, diphosphatidylglycerol; DGTA
12	diacylglycerylhydroxymethyl- <i>N</i> , <i>N</i> , <i>N</i> -trimethyl-β-alanine; DGTS,
13	diacylglyceryltrimethylhomoserine; MGDG, monogalactosyldiacylglycerol; PC,
14	phosphatidylcholine; PE, phosphatidylethanolamine; PI, phosphatidylinositol; PG,
15	phospahtidylglycerol; SQGD, sulfoquinovosyldiacylglycerol.

I- Introduction

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The brown algae belong to the division Heterokonta and are therefore only very distantly related to the three most intensely studied eukaryotic groups, the animals, fungi and green plants (Baldauf, 2003; Davis, 2004; Fig. 1a). This independent evolutionary history has furnished brown algae with many novel metabolic, physiological, cellular and ecological characteristics including a complex halogen metabolism, cell walls containing many unusual polysaccharides and high resistance to osmotic stress. Developmental processes are particularly interesting in this group, which evolved complex multicellularity independently of the three other major groups listed above. From a more applied point of view, the evolutionary history of the brown algae also underlies the high commercial value of several members of the group in the sense that they have evolved novel biomolecules such as polysaccharides and defence elicitors that have a wide range of applications in industry (Klarzynski et al., 2000; McHugh, 2003). It is important to note, however, that whilst the independent evolutionary history of the brown algae is the source of much of the interest of this group, it can also be seen as a handicap because the well-developed model organisms from the plant and animal lineages are of limited relevance to brown algal biology. Specialised brown algal models have been developed in specific domains, for example members of the fucoids for cell biology approaches (see references in Corellou et al., 2005), but a polyvalent model organism that allows access to a wide range of questions at the molecular level has been lacking. This situation is changing with the development of genomic and genetic tools for the filamentous brown alga *Ectocarpus siliculosus*. The sequencing of the genome of this alga has recently been completed and the sequence is currently being annotated

1 (http://www.cns.fr/externe/English/Projets/Projet KY/organisme KY.html). It is therefore an 2 opportune moment to look back at the emergence of *Ectocarpus* as a model organism. Research on Ectocarpus began in the 19th century with descriptions of species and 3 taxonomy, followed by studies aimed at unravelling reproduction and life history. Other 4 5 major aspects that have been studied include the sexual pheromones and infection of 6 Ectocarpus by viruses. Research has also been carried out on ultrastructure, photosynthesis 7 and carbon uptake, gamete recognition and resistance to anti-fouling agents. Several 8 eukaryotic parasites of *Ectocarpus* have been described. A proposition to adopt *Ectocarpus* as 9 a general model organism for the brown algae was made in 2004 (Peters et al., 2004a). This 10 proposition was based partly on this alga's long history as an experimental organism but also took into account several features that make Ectocarpus an interesting model for genetic and 11 12 genomic approaches. These features include its small size, the fact that the entire life cycle 13 can be completed in Petri dishes in the laboratory (Müller et al., 1998), its high fertility and rapid growth (the life cycle can be completed in 3 months), the ease with which genetic 14 15 crosses can be carried out and the relatively small size of the genome (200 Mbp compared 16 with 1095 and 640 Mbp for Fucus serratus and Laminaria digitata respectively; Le Gall et 17 al., 1993 and Peters et al., 2004a). 18 Here we present an overview of the work that has been carried out on *Ectocarpus* over the 19 last two centuries and discuss how the availability of a number of genomic tools, in particular 20 the complete genome sequence, is expected to accelerate research in many domains of brown 21 algal biology in the coming years.

II- Diversity and taxonomy, distribution and ecology

1 - Diversity and taxonomy

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3 Dillwyn (1802-1809) published the first valid description of *Ectocarpus* (using the name 4 Conferva siliculosa) based on material collected by W.J. Hooker on "rocks in the sea at 5 Cromer and Hastings". These English localities lie in Norfolk and East Sussex, respectively. 6 Type material, collected by Hooker in 1807, is housed at BM (BM000685585 and 7 BM000685588) under the name C. confervoides. Lyngbye (1819) described the genus 8 Ectocarpus based on material from Denmark and cited C. siliculosa Dillwyn as basionym. 9 The correct nomenclature, therefore, is E. siliculosus (Dillwyn) Lyngbye (see Silva et al., 10 1996 for further details). E. siliculosus is the type species of the order Ectocarpales which 11 includes most of the smaller brown algae. Originally regarded as phylogenetically primitive, 12 molecular systematics has shown the Ectocarpales to belong to a group of brown algal orders 13 that evolved rather recently. They are closely related to the large and highly differentiated 14 Laminariales which are major components of coastal marine floras (Rousseau & de Reviers, 1999; Draisma et al., 2003; Cho et al., 2004; Kawai et al., 2007, Fig. 1b). Many species have 15 16 been described in *Ectocarpus* (www.algaebase.org, April 2007, lists 392 taxa of *Ectocarpus* 17 of whom 98 are flagged as "current"; numerous strains publicly available at the Culture 18 Collection of Algae and Protozoa (CCAP) in Oban, U-K, Gachon et al., 2007a). However, 19 only E. fasciculatus Harvey (1841) is currently recognized as a second well-defined species, 20 based on morphology (Russell 1966, 1967a), crossing studies and sequence analyses (Stache-21 Crain et al., 1997). Crossing experiments have shown that the taxon E. siliculosus may 22 represent a species complex (Stache-Crain et al., 1997) and ongoing, refined analyses are 23 expected to resolve this complex, increasing the number of recognised species. Identification 24 of different species of *Ectocarpus* based on morphology is difficult due to the plasticity of the 25 commonly examined features (habit, branching pattern, size of sporangia). In addition, the

1 two generations of a species may differ considerably (Müller, 1972a; Kornmann & Sahling,

2 1977).

2 – Distribution

E. siliculosus is distributed world-wide in temperate regions, but does not occur in the tropics and south of the Antarctic convergence (Stache, 1990; Wiencke & Clayton, 2002). It occurs in fully marine and in low-salinity habitats (e.g. 5 psu in Finland) and has even been recorded at a fresh-water site in Australia (West & Craft, 1996) and in a salt polluted river in Germany (Geissler, 1983). Records of *E. fasciculatus* are mainly from the North Atlantic but there are some from Korea, Chile, South Georgia and South Africa (www.algaebase.org, 2007). On the shore, *Ectocarpus* occurs from high inter-tidal pools to the sub-littoral. It is found on abiotic substrata (rocks, wood, plastic, ship hulls) and epiphytic on macrophytes or free-floating (Russell, 1967a,b; 1983a,b). As a result of its ability to grow on a range of abiotic substrates, *Ectocarpus* is a common fouling alga (Morris & Russell, 1974).

3 - Ecology

There have been a limited number of ecophysiological and ecological studies in *Ectocarpus*. Growth rate is dependent on temperature, and there is evidence that temperature also influences the life cycle, at least in some strains (Müller, 1963; Bolton, 1983; see the "sporophyte and gametophyte architecture" section). The thermo-sensitivity of different strains suggests that there is a genetic heterogeneity within the *Ectocarpus* genus (Bolton, 1983). A similar result was obtained for osmo-acclimation (Thomas & Kirst, 1991a,b; see the "Abiotic stress" section). In the field, *Ectocarpus* is a short-lived annual which may dominate the ectocarpoid flora on kelps (Russell, 1983a,b). Despite the commonness of *Ectocarpus*,

- there are few data concerning phenology in the field (mostly to be extracted from floristic
- works) and nothing precise on seasonality or habitats of the two generations.
- The availability of the *E. siliculosus* genome sequence is expected to facilitate the analysis
- 4 of the ecology of this species by providing a basis for the development of molecular markers.
- 5 Particularly important challenges in this respect will include the identification of the sex locus
- 6 and of genes specifically expressed in the two generations, as these will provide molecular
- 7 tools that can be used to investigate several aspects of the life cycle under field conditions.
- 8 Molecular markers will also allow the exploration of genetic polymorphism amongst
- 9 Ectocarpus species from multiple locations across the globe.

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III- Development

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1 – Life cycle and reproduction

Male and female gametes are morphologically identical in *Ectocarpus* (isogamy) but differ with respect to their physiology and their behaviour: female gametes settle sooner and produce a pheromone whilst male gametes swim for longer and are attracted to the pheromone produced by the female. Studies on the reproduction of *Ectocarpus* began with the observation of sexual fusions involving the attraction of male gametes to settled female gametes from field thalli of *E. siliculosus* at Naples, Italy (Berthold, 1881). These findings were hotly debated until Sauvageau (1896, 1897) and Oltmanns (1899) succeeded in repeating the experiment. Gamete fusions in *Ectocarpus* were later used by Hartmann (1934) to support his erroneous theory of relative sexuality (see Müller, 1976a for details). Knight (1929) identified the young, unilocular sporangium as the site of meiosis, Papenfuss (1935) and Kornmann (1956) published major contributions to the life history of *Ectocarpus*, and

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Boalch (1961) developed refined culture techniques. The entire life history of E. siliculosus from Naples was finally unravelled by Müller (1964, 1966, 1967, 1972b) using clonal cultures and chromosome counts. It is schematised in Fig. 2. The basic life history of E. siliculosus involves an alternation between the sporophyte and dioecious gametophytes, and sex determination is genotypic (Müller, 1967). Male and female gametophytes are morphologically indistinguishable. One of the problems for understanding the life history was that sporophytes and gametophytes are difficult to distinguish morphologically. Another problem was that zoids from plurilocular reproductive organs have different functions according to the generation forming them: on sporophytes they contain asexual zoospores that directly reproduce the sporophyte, on gametophytes they contain gametes. Further complications include the parthenogenesis of unfused gametes, which develop into haploid parthenogenetic sporophytes morphologically indistinguishable from diploid sporophytes, and heteroblasty (different fates) of spores from unilocular sporangia, developing either into gametophytes or into sporophytes. Furthermore, life cycle generation is not determined rigidly by ploidy (Müller, 1967). An important challenge for the future will be the characterisation of the genetic mechanisms that control life cycle progression in *Ectocarpus*. This will require the development of methodologies for positional cloning of mutated loci and of genome-wide methods to analyse gene expression throughout the life cycle. Work is currently ongoing in several groups to develop these techniques.

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2 – Sporophyte and gametophyte architecture

E. siliculosus is a small filamentous alga that grows to about 30 cm in length in nature, but it may become fertile in the laboratory at 1-3 cm.

1 Sporophyte development is initiated with the germination of the zygote. The first division 2 produces two cells of identical developmental fate (Peters et al., 2004b). Subsequent mitoses 3 lead to the formation of a basal (or prostrate) filamentous structure, defining the early 4 sporophyte (Fig. 3a). Phaeophycean hairs, i.e. hyaline filaments devoid of plastids developing 5 from a basal meristem, are absent in Ectocarpus but present in the sister genus Kuckuckia. 6 However, in *Ectocarpus* the distal end of filaments, or of plurilocular sporangia, may be less 7 pigmented and resemble a hair; such structures may be referred to as pseudo-hairs (Cardinal, 8 1964; Pedersen, 1989). If the growth conditions are favourable (Ravanko, 1970), erect 9 filaments (called "upright" filaments) emerge after a few days, contributing to the 10 establishment of an overall filamentous architecture (Fig. 3c). 11 The typical structure of a vegetative cell is illustrated in Fig. 4. Features common to all 12 brown algal cells include a chloroplast surrounded by four membranes, arranged as two 13 double-membraned envelopes (chloroplast envelope, CE). The second envelope is loosely associated with the chloroplast and forms part of the chloroplast endoplasmic reticulum 14 15 (CER). The lamellae of the chloroplast are composed of three thylakoids, which are absent 16 from the pyrenoid space (Bouck, 1965; Oliveira & Bisalputra, 1973). In Ectocarpus the 17 chloroplast is ribbon-shaped. The size and number of chloroplasts may vary within the same organism (Ravanko, 1970). Other typical features of *Ectocarpus* include several prominent 18 19 and pedunculated pyrenoids on the inner face of the chloroplast, which are used as a 20 taxonomic marker for the Ectocarpales (Evans, 1966; Rousseau & de Reviers 1999). CER and 21 CE envelopes also surround pyrenoids, this time being tightly adjacent. A third external 22 envelope, called the pyrenoid sac, surrounds the pyrenoid but has no connection with the 23 reticulum system (Bouck, 1965). The nuclear envelope is continuous with the CER, which, 24 itself is in close vicinity to the Golgi apparatus (Bouck, 1965; Oliveira & Bisalputra, 1973). It 25 has been hypothesised that these connections create a complex network of membranes

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allowing photosynthates to be efficiently transferred from the chloroplast to the Golgi apparatus, the latter being also in direct contact with the CER (Oliveira & Bisalputra, 1973). Cytoplasmic ER is dispersed throughout the cytoplasm and is mainly rough (Oliveira & Bisalputra, 1973). Osmiophilic bodies (OSB) which are thought to contain lipids, are dispersed throughout the cytoplasm and probably originate from the CER (Oliveira & Bisalputra, 1973). They have been observed within the cell wall, and also external to it. Vacuoles can be either large structures occupying peripheral locations (Oliveira & Bisalputra, 1973) in fixed material, or most of the cellular space (as in land plant cells; Knight, 1929, confirmed by data from our laboratory, after staining with cresyl blue or neutral red, on sporophytic filaments). The nuclear region encompasses two centrioles, which are considered as a microtubule organising centre (MTOC; Katsaros et al., 1991). The chromosome number in the haploid nucleus is estimated to be ca 25 (Peters et al., 2004a). Mitochondria are preferentially peripherally located. They are maternally inherited (Peters et al., 2004b). The cell wall consists of a fibrillar matrix (see the "Photosynthesis and carbohydrate metabolism" section) with several plasmodesmata distributed uniformly along the cross walls. Upon ageing, cell wall ingrowths occur, accompanied by a reduction in the size of the nucleus, mitochondria, ER and Golgi, followed by the disintegration of the chloroplasts and finally by autolysis of the cytoplasm (Oliveira & Bisalputra, 1977a,b). Most of the reproductive organs are carried by upright filaments (Müller, 1964). Two types of reproductive organs are produced by the sporophyte: plurilocular and unilocular sporangia. Plurilocular sporangia are cone-shaped three-dimensional structures of variable size (Knight, 1929), composed of a large number of locules with different shapes (Baker & Evans, 1973b; Fig. 3e). These locules are generated by several successive mitoses and each gives rise to a single zoospore, which is released through the apex of the sporangium. The mito-spores,

1 which are bi-flagellate and competent for swimming shortly after their release (Müller, 1980), 2 are a means of vegetative reproduction. Unilocular sporangia are born on the side of branches 3 (Fig. 3f, Baker & Evans, 1973a). A single meiosis occurs within the single thick-walled locule 4 (Baker & Evans, 1973a) and this is followed by several mitoses, which generate about a 5 hundred meio-spores, half of which are female and half male (Müller, 1980). Müller (1963) 6 and Ravanko (1970) reported that plurilocular sporangia were produced when the external 7 temperature was relatively high (~20°C or summer), whereas unilocular sporangia were 8 produced when the temperature was lowered to 13°C (mimicking winter conditions). 9 However, this temperature dependence is not observed in many strains of *Ectocarpus* (A.F. 10 Peters, unpublished results). Meio-spores germinate to produce haploid, dioecious 11 gametophytes (Fig. 3b,d). These are filamentous organisms similar to the sporophyte, but with 12 two important differences. Firstly, the meio-spore germinates asymmetrically to produce a 13 rhizoid and an upright filament, so no prostrate structure forms (A.F. Peters, unpublished 14 results). Secondly, the thallus is more ramified than that of the sporophyte (Müller, 1980). 15 Gametophytes produce only plurilocular gametangia and these are similar structurally to 16 plurilocular sporangia on sporophytes. Gametes resemble mito-spores in terms of their size 17 and motility. 18 The developmental patterning varies greatly across the *Ectocarpus* species complex and is 19 also dependent on the environment, growth conditions and even the age of the algae for some 20 features (Ravanko, 1970). This plasticity is observed for the branching frequency and the 21 number, shape, structure and positioning of the reproductive organs on filaments (Knight, 22 1929; Müller, 1980; Kim & Lee, 1992). Phytohormones, especially cytokinins, have also been 23 reported to influence the development of *Ectocarpus* sporophytes (Pedersén, 1968, 1973).

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As mentioned above, brown algae are interesting for developmental studies because they have independently evolved complex multicellularity. *Ectocarpus* is a relevant model to address this problem as it is closely related to complex algae such as the Laminariales and the Fucales. However, developmental processes in *Ectocarpus* are clearly simplified compared with its morphogenetically more complex sister families, which provides an advantage for the detailed dissection of these developmental processes. In particular the simple growth pattern of the uniseriate, branched filaments represents an ideal system for the combined application of genetic and mathematical modelling approaches to understanding developmental patterning and then addressing the issue of the evolution of development and multicellularity.

3 – Gametes and spores

reported (Baker & Evans 1973a,b; Lofthouse & Capon, 1975; Maier, 1997a,b). The typical structure of an *Ectocarpus* zoid is illustrated fig. 5.

**Ectocarpus* zoids correspond to the "primitive" type of brown algal zoid according to Kawai (1992). Gametes and spores typically contain a single chloroplast with a pyrenoid (Baker & Evans, 1973a; Lofthouse & Clayton, 1975; Maier, 1997a). As in the vegetative cells, lamella are composed of three thylakoids and the nuclear envelope is in continuity with the chloroplast endoplasmic reticulum (Maier, 1997a). The nucleus of the male gamete is rich in heterochromatin. Several dictyosomes are present in gametes and spores (Baker & Evans, 1973b; Maier, 1997a). The Golgi apparatus of mito-spores is very active both before and after release (Baker & Evans, 1973b). This secretory activity may have important functions in the biosynthesis of the adhesive required for gamete adhesion and for the synthesis of new cell wall compounds during germination.

Several electron microscopy studies of the motile cells (zoids) of *Ectocarpus* have been

1 Ectocarpus gametes and spores are characterized by two flagella with lateral insertion. One 2 is oriented forward and equipped with mastigonemes (hairs), and propels the cell with 3 meandering beats. The second is oriented obliquely backwards, and has no mastigonemes. 4 Most of the time this posterior flagella is passively dragged but occasional lateral beats induce 5 abrupt changes in direction of up to 180° (Geller & Müller, 1981). Gametes and spores have a 6 concave depression at the level of the eyespot into which the swelling of the proximal part of 7 the posterior flagellum fits (Baker & Evans, 1973b; Kreimer et al., 1991; Maier, 1997a). The 8 possible function of the eyespot is to reflect and focus incident light onto the site of 9 photoreception (Kawai et al., 1990; Kreimer et al., 1991). Zoids are capable of positive 10 phototaxis and their posterior flagellum shows strong autofluorescence when irradiated by 11 blue light (450 nm). The photoreceptor pigment is a flavoprotein, which is periodically 12 shaded by a carotenoid stigma (Müller et al., 1987; Kawai, 1988). The acronema (the 13 whiplash tip) is extremely sensitive to mechanical stress and plays an important role in 14 establishing the initial sexual contact between gametes. The details of the flagellar apparatus 15 of both female and male gametes have been studied by electron microscopy (Müller & Falk, 16 1973; Maier, 1997b). Despite their different behaviours, no difference in their fine structure 17 has been detected (Müller & Falk, 1973). 18 The pheromone that attracts male gametes to female gametes is an unsaturated hydrocarbon. 19 The substance initially identified was ectocarpene (all-cis-1-(cycloheptadiene-2',5'-vl)-20 butene-1) (Müller et al., 1971; Müller, 1976b; Müller, 1978; Müller & Schmid, 1988) but 21 more recently Boland et al. (1995) have shown that a thermally labile cyclopropyl precursor, 22 pre-ectocarpene, is more active by three orders of magnitude and is thus the actual 23 pheromone. 24 There is evidence that cell-to-cell recognition between *Ectocarpus* gametes is mediated by 25 N-acetyl glucosamine residues exposed on the plasma membrane of the female gametes and

1	that these residues are specifically recognized by a receptor on the male gamete (Schmid,
2	1993; Schmid et al., 1994a). In addition, the lectins concanavalin A (Con A) and Aleuria
3	aurantia agglutinin (AAA) bind specifically to the anterior flagella of Ectocarpus gametes
4	and the molecules with which they interact also may be involved in the gamete recognition
5	process (Maier & Schmid, 1995).
6	Zoids from plurilocular and unilocular Ectocarpus sporangia share similar overall
7	intracellular structures, but there are some important differences: plurilocular zoids are
8	smaller and swim faster with more rapid changes of direction (Baker & Evans, 1973a). Zoids
9	from unilocular sporangia are strikingly different from the other <i>Ectocarpus</i> cell types (both
10	plurilocular zoids and vegetative cells) in that the nucleus is physically separated from the
11	chloroplast. Moreover, secretory activity is lower than in zoids from plurilocular sporangia
12	(Baker & Evans, 1973b).
13	Gametes of <i>Ectocarpus</i> have also been used to study chemotaxis (Boland et al., 1983),
14	receptor modelling (Boland et al., 1989) and phototaxis (Kawai et al., 1990; Kreimer et al.,
15	1991). Protocols for isolation and biochemical characterisation of plasma membrane and CER
16	membrane have also been developed (Schmid et al., 1992). Together with more recently
17	developed molecular tools, these methods now offer access to a range of interesting
18	biochemical events that take place during gamete interaction at fertilization such as
19	chemoreception, cell-cell recognition and fusion processes.
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	IV- Metabolism
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24	1- Photosynthesis and carbohydrate metabolism

Marine environment constrains several aspects of photosynthesis in brown algae. First, carbon dioxide is not the main source of inorganic carbon (Ci). Indeed, seawater in equilibrium with air contains only 13 μM CO₂, but more than 2 mM anionic carbon, mainly in the form of HCO₃⁻ (Beer, 1994). Secondly, when seaweeds are immersed they do not receive the full light spectrum. Light absorption increases with water depth and varies according to the wavelength: red-light (650 nm) is first absorbed followed by purple (400 nm) and yellow-(550 nm) lights. In contrast green (500 nm) and blue (450 nm) lights display a strong penetration: at 10 m depth red light is almost fully absorbed whereas the absorption of green-and blue lights is not significant.

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Adapted to these conditions, brown algae differ from land plants by the pigment composition of their light harvesting complexes (LHCs). Chlorophyll c (chl c) and the carotenoid fucoxanthin are indeed the main light-haversting pigments of brown algae. Their presence broadens the absorption spectrum toward the green light relative to the chlorophytes. Brown algae features two different LHCs associated to photosystems I and II (PSI and PSII), but their pigment composition is controversial. Barrett and Anderson (1980) described a fucoxanthin-chl a/c protein and a violaxanthin enriched-chl a/c protein. Conversely Alberte et al. (1981) reported a chl a/c protein devoid of fucoxanthin and a second LHC containing chl a and fucoxanthin but not chlorophyll c. More recent analyses showed that the LHCs associated to PSI and PSII are in fact virtually identical with respect to their pigmentation and peptide composition. All complexes bound chl a, chl c and fucoxanthin in the proportion 6:2:7, but the LHC associated to PSI is significantly enriched in violaxanthin (De Martino et al., 2000). As in plants, these accessory pigments transfer energy to chlorophyll a within the photosynthetic reaction centres (Grossman et al., 1995). Violaxantin does not participate in light harvesting but is involved in photoprotection (Demmig-Adams & Adams, 1992; Lemoine et al., 1995).

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Under saturating red light, photosynthesis of E. siliculosus follows a circadian rhythm with maxima at about noon and can be stimulated by a pulse of blue light (BL) (Schmid & Dring, 1992). This stimulation is also observed in other Phaeophytes, but mostly absent in green or red algae (Schmid et al., 1994b). BL induces several responses: an acidification at the surface of E. siliculosus (Schmid & Dring, 1993), accompanied by bicarbonate uptake (Schmid, 1998). A plasma membrane H⁺-ATPase is thought to be activated, and the resulting acidification to increase the conversion of HCO₃⁻ to CO₂ in the extracellular space (Schmid & Dring, 1993). Recently, such BL-induced H⁺-ATPases have been identified in the brown alga Laminaria digitata (Klenell et al., 2002). BL also triggers the mobilization of an internal carbon source, since photosynthesis is stimulated even in the absence of external Ci (Schmid & Dring, 1996). As a result, a C4-like metabolism was initially proposed to exist in E. siliculosus (Schmid & Dring, 1996) but not all the enzymes necessary for a C4 cycle were detected (Bush & Schmid, 2001) and the pool of intermediates seems to be too small to act as an organic carbon stock (Hillrichs & Schmid, 2001). The sequestration of a pool of inorganic carbon in the vacuole and its movement to the cytosol in response to BL is now the favoured hypothesis (Schmid & Hillrichs, 2001). Analysis of the sequence of the E. siliculosus genome will help to confirm whether or not a C4 pathway exists in the brown algae. In contrast to the *Plantae* (Moreira et al., 2000), the *Phaeophyceae* do not store the carbon assimilated by photosynthesis as insoluble starch granules, but instead as the soluble 1,3-\u03b3glucan polymer (laminarin) localised in the cytosol (Craigie, 1974), and as mannitol, involved in osmo-acclimation (Davis et al., 2003; see the "Abiotic stresses" section). Brown algae also produce complex polysaccharides which constitute their cell wall. They synthesize some neutral polysaccharides in common with land plants, such as cellulose (Carpita & McCann, 2000), but also unique anionic polysaccharides, such as alginates and sulphated fucans

- 1 (Kloareg & Quatrano, 1988). Ectocarpus has not been well studied in this respect, but
- 2 preliminary analyses in our laboratory confirm that all of the polysaccharides typical of brown
- 3 algae are present in this genus (Estelle Deniaud, pers. comm.). The biosynthetic pathways of
- 4 these brown algal polysaccharides are essentially unknown and the genome of E. siliculosus
- 5 will be a much anticipated asset to investigate these crucial metabolisms.

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2- Lipid metabolism

Worldwide, at least 50 species of brown algae are used as human food. Their lipid content has therefore attracted considerable attention from the viewpoint of both nutrition and pharmacology. A number of studies have been conducted to profile and quantify the fatty acids and the different classes of lipids in these organisms, and to investigate whether the various lipid patterns correlate with the taxonomic position or any other characteristic of the brown algae. Brown algal polar lipids include several common glycolipids (MGDG, DGDG, SQGD, see list of abbreviations) and phospholipids (PC, PE, PI, PG and DPG). Interestingly, several reports have highlighted the high proportion of long chain polyunsaturated fatty acids (LC-PUFAs) (Eichenberger et al., 1993). Moreover, most of the Phaeophyceae contain the betaine lipids DGTA and DGTS, and either contain the common phosphatidylcholine phospholipids in surprising low amounts or do not produce them at all (Eichenberger et al., 1993). All Ectocarpales contain phosphatidylcholine, but exhibit interspecific variation in their DGTA content: E. fasciculatus strains contain this lipid while E. siliculosus strains do not (Müller & Eichenberger, 1995; Müller, 1995). DGTA may therefore be used as a taxonomic marker (Müller, 1995), although an E. fasciculatus strain deficient in DGTA biosynthesis has been described. Genetic analysis of such a strain identified an autosomal locus necessary for the biosynthesis of this lipid (Müller & Eichenberger, 1997).

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1 A novel phosphoglyceride, designated PX, was first isolated from E. siliculosus. It was 2 shown to account for 2-4% of total lipids (Schmid et al., 1994c), and to accumulate mostly in 3 the plasma membrane of gametes. Since PX is rich in 20:4n-6 (ARA) and 20:5n-3 (EPA), it 4 has been suggested that it could represent a potential reservoir for pheromone precursors. PX 5 was subsequently detected in other brown algae (Ectocarpales, Fucales and Sphacelariales; 6 Schmid et al., 1994c). 7 Lipid metabolism plays also major roles in the control of defense mechanisms. Therefore, 8 together with the genome sequence, it is likely that the efforts to screen mutants with altered 9 resistance to pathogen attacks, will lead to the phenotyping of plants impaired in lipid or fatty 10 acid metabolic pathways, thereby potentially revealing novel specific traits of this metabolism 11 in brown algae. 12 13 14 V- Interactions with the environment 15 16 1- Ectocarpus pathogens 17 Despite of their small size and ephemeral life stages, filamentous brown algae have been 18 frequently reported to be plagued by various pathogens, including viruses (Müller et al., 19 1998) and eukarvotic parasites of different phylogenetic lineages: oomycetes, chytrids and 20 hyphochytrids (Andrews 1976; Küpper & Müller 1999; Müller et al. 1999) and by parasites

related to the Plasmodiophorea (Karling, 1944; Maier et al., 2000). In addition, numerous

historical records described ectocarpoids with abnormal sporangia or vegetative cells

The oomycete Eurychasma dicksonii has been described mainly in wild populations of

Pylaiella littoralis (Küpper & Müller, 1999) but it displays a broad host range and infects

suspected to contain unknown parasites (Rattray, 1885; Müller et al., 1998).

1 various brown algae including Ectocarpus (Müller et al., 1999) in which it was initially 2 described by Wright (1879). There is a current effort to set up a defined pathosystem using E. 3 siliculosus and E. dicksonii, and Ectocarpus strains have been shown to exhibit differential 4 susceptibility to a defined Eurychasma strain. Conversely, several Eurychasma strains exhibit 5 different host specificities, suggesting co-evolution of the two species (Gachon et al., 2007b). 6 The molecular bases of resistance and virulence are under investigation. 7 Chytrids were described earlier by Petersen (1905) and the hyphochytrid Anisolpidium 8 ectocarpii was described by Karling (1943) and Johnson (1957). Like E. dicksonii, 9 Chytridium polysiphoniae (Chytridiomycota) is ubiquitous and can infect many hosts, including E. siliculosus and E. fasciculatus (Müller et al., 1999). Interestingly, its negative 10 11 effects on photosynthesis of its host was described at the cellular level in the related 12 ectocarpoid P. littoralis using fluorescence kinetic microscopy (Gachon et al., 2006). 13 Recently, the 18S rRNA genes of Chytridium polysiphoniae and Eurychasma dicksonii were sequenced and used to clarify their phylogenetic affiliations (Küpper et al., 2006). The 14 15 plasmodiophorean Maullinia ectocarpii is an obligate intracellular parasite of Ectocarpus spp. 16 (Maier et al., 2000). However, the extent to which this infection occurs in nature and its effect 17 on algal fitness are presently unknown. 18 Viral infections represent by far the most studied phenomenon in E. siliculosus (Müller, 19 1996; Müller & Knippers, 2001). Until the late eighties most reports of virus infections in 20 brown algal tissues were based on electron microscopy studies, which sporadically described 21 "virus-like particules" (VLP). Viruses were obtained in culture for the first time from a New 22 Zealand strain of E. siliculosus after lysis of host cells allowing evaluation of their infection 23 potential (Müller et al., 1990; Müller, 1991). Virus infections were found in about 50% of the 24 individuals of a given natural population (Dixon et al., 2000; Müller et al., 2000) and were

1 shown to occur worldwide in correlation with the cosmopolitan distribution of E. siliculosus 2 (Müller, 1991; Sengco et al., 1996). 3 The viruses that infect different ectocarpoid algae exhibit considerable variability in size 4 and diameter and, in general, they display a high level of host specificity (Müller et al., 1998). 5 However, several instances of trans-specific infection have been described, for example 6 between EsV-1 (Ectocarpus siliculosus virus-1) and Kuckuckia kylinii (Müller, 1992; Müller 7 & Schmid, 1996) and also between EfasV-1 (Ectocarpus fasiculatus virus-1) and E. 8 siliculosus (Müller et al., 1996; Sengco et al., 1996). Interestingly, EsV-1 and EfasV-1 are the 9 most similar of the brown algal viruses in terms of their genome size (Müller et al., 1996). 10 The EsV-1 virus specifically infects the single-celled gametes or spores, *i.e.* the only cells in 11 the life history that lack a cell wall (Maier & Müller, 1998). Following infection, a single 12 copy of the viral DNA appears to integrate into the host genome (Delaroque et al., 1999). The 13 viral DNA is then transmitted, via mitotic divisions, to all the cells of the developing alga. This has been confirmed by regenerating algae from protoplasts derived from virus-infected 14 15 gametophytes (Kuhlenkamp & Müller, 1994). Despite the fact that they carry the integrated 16 virus, vegetative cells do not produce viral particles (Müller et al., 1998). Viral particles are 17 only produced in reproductive organs (sporangia and gametangia) of mature algae from where 18 they are released to infect a new generation of zoids. In addition to these cycles of re-19 infection, the viral genome can be also transmitted to progeny through meiosis, in which case 20 it segregates as a Mendelian factor and is inherited by half of the progeny (Müller, 1991; 21 Bräutigam et al., 1995). The pathogenic character of viral infections has been unambiguously 22 confirmed, but this association's main impact is on reproductive success. Plant sterility varies 23 from partial (Müller et al., 1990) to total (Müller & Frenzer, 1993) but no significant 24 difference in photosynthesis, respiration and growth rate were observed in infected

1	gametophytes or sporophytes (Del Campo et al., 1997). This contrasts with the reduced
2	photosynthetic performance of Feldmannia species infected with FsV (Robledo et al., 1994).
3	The EsV-1 genome is a circular DNA molecule of a relatively large size (335 Kbp) for a
4	phycodnavirus (Van Etten & Meints, 1999; Van Etten et al., 2002) with double-stranded
5	regions interrupted by single-stranded regions (Lanka et al., 1993; Klein et al., 1994). Both
6	EsV-1 and the related Feldmania irregularis virus (FirrV-1) have been sequenced (Delaroque
7	et al., 2001; Delaroque et al., 2003). EsV-1 contains approximately 231 genes with a wide
8	range of predicted functions including DNA metabolism, signalling, transposition, DNA
9	integration and polysaccharide metabolism (Delaroque et al., 2000a,b; Delaroque et al.,
10	2003). It has also been proposed that the ability of the virus to integrate into its host's genome
11	could be exploited to develop a transformation vector for a wide range of brown algae,
12	including E. siliculosus (Henry & Meints, 1994, Delaroque et al., 1999). However, the
13	complex integration pattern of the virus into the algal genome will considerably complicate
14	this task (N. Delaroque, pers. comm.). A microarray has been constructed to analyse EsV-1
15	gene expression (Declan Schroeder, pers. comm.) and it will be particularly interesting in the
16	future to couple the analysis of viral and genome-wide host gene expression during viral
17	infection.
18	The development of genomic tools provides a new context to investigate the possible
19	genetic basis of the co-evolution between some pathogens and brown algae. The search for
20	inducers of defense responses and resistance against parasites is also still opened as, in
21	contrary with kelps, Ectocarpus does not react with an oxidative burst upon recognition of
22	alginate fragments (Küpper et al., 2002a).

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2- Abiotic stresses

1 E. siliculosus is able to exploit a wide range of habitats and environmental conditions (see 2 the "Distribution" section). This feature seems likely to be based at least as much as on a high 3 intrinsic genetic variability as on a general physiological toughness, as illustrated by work 4 carried out on copper and saline stress responses. 5 Interspecific variations in copper tolerance have been observed between different 6 strains of E. fasciculatus and E. siliculosus, with the latter being the most tolerant (Morris, 7 1974). Differences have also been observed among E. siliculosus strains that are differently 8 exposed to copper in their natural habitat (Russell & Morris, 1970; Hall, 1981). Cu²⁺ 9 interferes with the general process of photosynthesis in brown algae, and particularly in E. 10 siliculosus, by competing with magnesium for metal binding sites in the chlorophyll 11 molecules (Küpper et al., 2002b). A study of the mechanism of tolerance to copper and other 12 heavy metals suggested a co-tolerance to copper, cobalt and zinc, and provided evidence for 13 an exclusion mechanism to explain the particularly low sensitivity of E. siliculosus copper tolerant strains (Hall et al., 1979; Hall, 1980, 1981). However, as yet there is no clear 14 15 explanation for the intra-specific variation with respect to this trait within this species. 16 Ability of some E. siliculosus strains to tolerate copper has been considered to 17 develop bioassays in which this alga is used for monitoring marine antifouling characteristics 18 of copper-based materials (Hall and Baker, 1985, 1986). In addition, copper chloride was 19 used to inhibit *E. siliculosus* infestations in tank cultures of *Gracilaria gracilis* (Van Heerden 20 et al., 1997). 21 22 Russell and Bolton (1975) reported the occurrence of salinity ecotypes within *E. siliculosus*. 23 This study was extended by Thomas and Kirst (1991a,b) who showed that large differences in 24 photosynthesis, accumulation of osmotically active compounds (mannitol; Davis et al., 2003)

and vitality occur between E. siliculosus isolates from different geographic locations

- 1 following changes in salinity. They also observed that sporophytes were more salt tolerant
- 2 than gametophytes, irrespective of their level of ploidy.
- 3 Detailed investigations are necessary to decipher the physiological and cellular bases of salt
- 4 and heavy metal tolerance in *E. siliculosus*. Mutagenesis and transcriptomic approaches will
- 5 thus help to better understand the mechanisms involved in osmotic and oxidative adaptation,
- and to explain how these algae can cope with such a wide range of environmental conditions.

VI- Conclusion

model organism.

Taken together, the above sections illustrate the broad range of phenomena that have been studied in *Ectocarpus* and provide an indication of the domains that could be further explored in the future. Notably, a large proportion of this past work, covering many diverse aspects of *Ectocarpus* biology, was carried out in Dieter Müller's laboratory in Konstanz, and the efforts of this group have therefore laid the foundations for the development of *Ectocarpus* as a

The *Ectocarpus* genome project has federated a consortium of laboratories with an interest in this organism and these laboratories are currently developing several molecular tools. These include mutant screens, genetic transformation and genome-scale analysis of gene expression. Several developmental mutants have been isolated and positional cloning of some of the affected genes should be feasible in the near future. The availability of the genome sequence together with the ability to analyse gene function by forward and reverse genetic approaches will make it possible to address additional questions, many of which have been evoked in this review. Examples include the biosynthesis of diverse, brown-algal-specific metabolites, such as lipids and complex cell wall components, and the genetic basis of

1	resistance to biotic and abiotic aggression. The genome will also be an invaluable aide for the
2	study of the ecology of <i>Ectocarpus</i> , by serving as the base for the development of neutral and
3	selected molecular markers for the analysis of field isolates. In conclusion, the Ectocarpus
4	genome sequence and the tool development associated with this project are providing access
5	to a relatively unexplored branch of the eukaryotic tree and some exciting discoveries can be
6	expected in this domain in the coming years.
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16	
17	Figure legends:
18	
19	Figure 1: Phylogeny of brown algae and Ectocarpales.
20	a: Position of brown algae within the eukaryotes (adapted from Baldauf, 2003). Brown algae
21	belong to the heterokont phylum, which is phylogenetically distant from land plants and the
22	green and red algae. Photosynthetic organisms are framed. b : Position of the Ectocarpales (in
23	bold) within the brown algae (adapted from Kawai et al., 2007).
24	

1 Figure 2: Life cycle of <i>Ectocarpus siliculo</i>	sus.
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- 2 Diploid sporophytes produce meio-spores (by meiosis) in unilocular sporangia (UL). Meio-
- 3 spores grow into male or female gametophytes (dioecism). Gametophytes produce gametes in
- 4 plurilocular gametangia (PL). Fusion of gametes produces a zygote that grows into a diploid
- 5 sporophyte, completing the sexual cycle. Unfused gametes may grow parthenogenetically and
- 6 form a parthenosporophyte, which is indistinguishable from the diploid sporophyte. Both
- 7 sporophytes and parthenosporophytes can reproduce themselves asexually by the production
- 8 of mito-spores in plurilocular sporangia.

9

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Figure 3: Morphology of Ectocarpus siliculosus.

- Photographies of one week old vegetative sporophyte (a) and gametophyte (b) and schemes
- representing the whole body of the mature sporophyte (c) and gametophyte (d) after six
- weeks of growth, are shown. e: Plurilocular sporangium or gametangium (occurring on the
- sporophyte and the gametophyte, respectively) before (left) and after (right) release of zoids.
- 15 **f**: Unilocular sporangium from sporophyte. Sporangia and gametangia can be either sessile or
- pedicellate (Kim & Lee, 1992).

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Figure 4: General ultrastructure of a vegetative cell of *Ectocarpus siliculosus*.

- 19 The general ultrastructure of a vegetative cell is similar in both prostrate and erect filaments
- 20 (Oliveira & Bisalputra, 1973). The different compartments of the cell are illustrated (see text
- 21 for details). Lines represent membranes and define sub-cellular compartments, except for
- 22 thylakoids, drawn as a thick black line. Depending on their type and age, vegetative cell size
- varies from 10 to 35 µm in length, and 5 to 15 µm in width (in laboratory culture conditions).

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25

Figure 5: General intracellular structure of *Ectocarpus siliculosus* zoids.

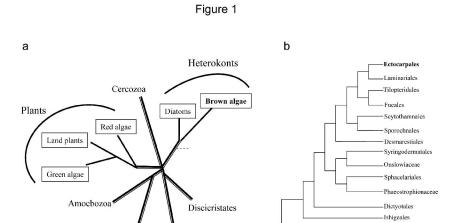
- 1 A scheme representing the different compartments of an *Ectocarpus siliculosus* zoid cell from
- 2 a plurilocular sporangium or gametangium is presented (see text for details). Legends are the
- 3 same as in fig. 4.

4

5

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- 9 the EsV-1 virus.



Opisthokonts

Excavates

Figure 1: Phylogeny of brown algae and Ectocarpales. a: Position of brown algae within the eukaryotes (adapted from Baldauf, 2003). Brown algae belong to the heterokont phylum, which is phylogenetically distant from land plants and the green and red algae. Photosynthetic organisms are framed. b: Position of the Ectocarpales (in bold) within the brown algae (adapted from Kawai et al., 2007).

275x190mm (300 x 300 DPI)

Choristocarpaceae

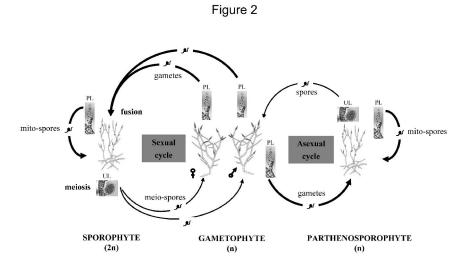


Figure 2: Life cycle of Ectocarpus siliculosus. Diploid sporophytes produce meio-spores (by meiosis) in unilocular sporangia (UL). Meio-spores grow into male or female gametophytes (dioecism). Gametophytes produce gametes in plurilocular gametangia (PL). Fusion of gametes produces a zygote that grows into a diploid sporophyte, completing the sexual cycle. Unfused gametes may grow parthenogenetically and form a parthenosporophyte, which is indistinguishable from the diploid sporophyte. Both sporophytes and parthenosporophytes can reproduce themselves asexually by the production of mito-spores in plurilocular sporangia.



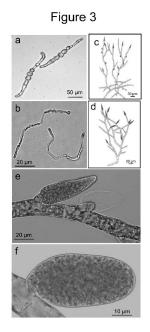


Figure 3: Morphology of Ectocarpus siliculosus. Photographies of one week old vegetative sporophyte (a) and gametophyte (b) and schemes representing the whole body of the mature sporophyte (c) and gametophyte (d) after six weeks of growth, are shown. e:

Plurilocular sporangium or gametangium (occurring on the sporophyte and the gametophyte, respectively) before (left) and after (right) release of zoids. f: Unilocular sporangium from sporophyte. Sporangia and gametangia can be either sessile or pedicellate (Kim & Lee, 1992).

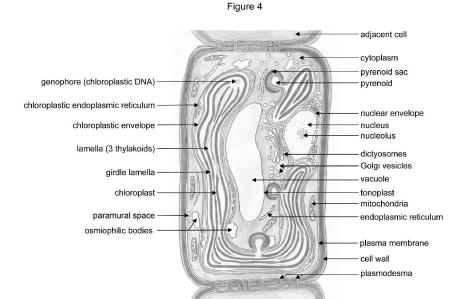


Figure 4: General ultrastructure of a vegetative cell of Ectocarpus siliculosus. The general ultrastructure of a vegetative cell is similar in both prostrate and erect filaments (Oliveira & Bisalputra, 1973). The different compartments of the cell are illustrated (see text for details). Lines represent membranes and define sub-cellular compartments, except for thylakoids, drawn as a thick black line. Depending on their type and age, vegetative cell size varies from 10 to 35 μ m in length, and 5 to 15 μ m in width (in laboratory culture conditions).

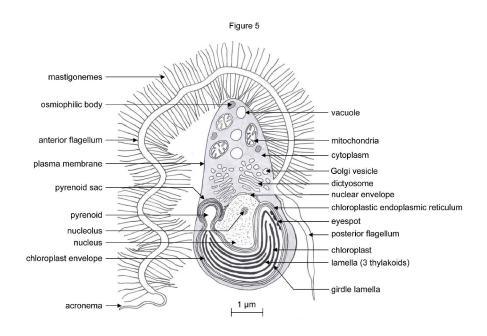


Figure 5: General intracellular structure of Ectocarpus siliculosus zoids. A scheme representing the different compartments of an Ectocarpus siliculosus zoid cell from a plurilocular sporangium or gametangium is presented (see text for details). Legends are the same as in Fig. 4.