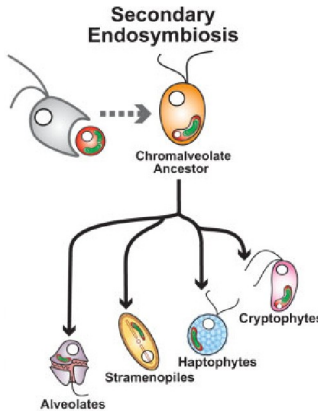


Trends in Ecology & Evolution

Figure 1. The New Tree of Eukaryotes.

This summary is based on a consensus of recent phylogenomic studies. The colored groupings correspond to the current 'supergroups'. Unresolved branching orders among lineages are shown as multifurcations. Broken lines reflect lesser uncertainties about the monophyly of certain groups. Star symbols denote taxa that were considered as supergroups in early versions of the supergroup model; thus, all original supergroups except Archaeplastida have either disappeared or been subsumed into new taxa. The circles show major lineages that had no molecular data when the supergroup model emerged, most often because they had not yet been discovered. Rappemonads (in parentheses) are placed on the basis of plastid rRNA data only. The putative new major lineages *Microheliella* and *Anaeramoeba* are not shown due to the limited evidence that they belong outside all existing groups shown here (Table 1).



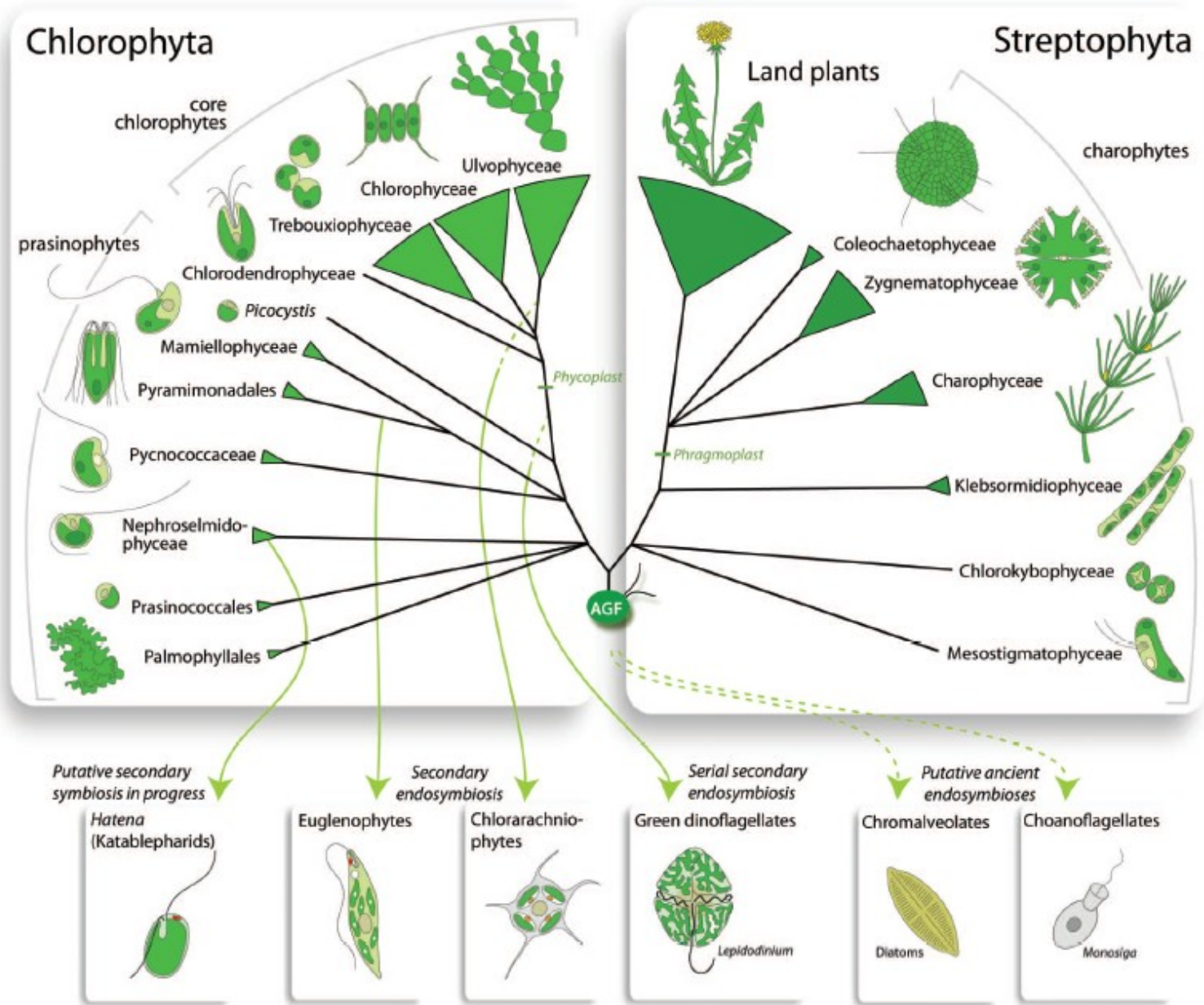
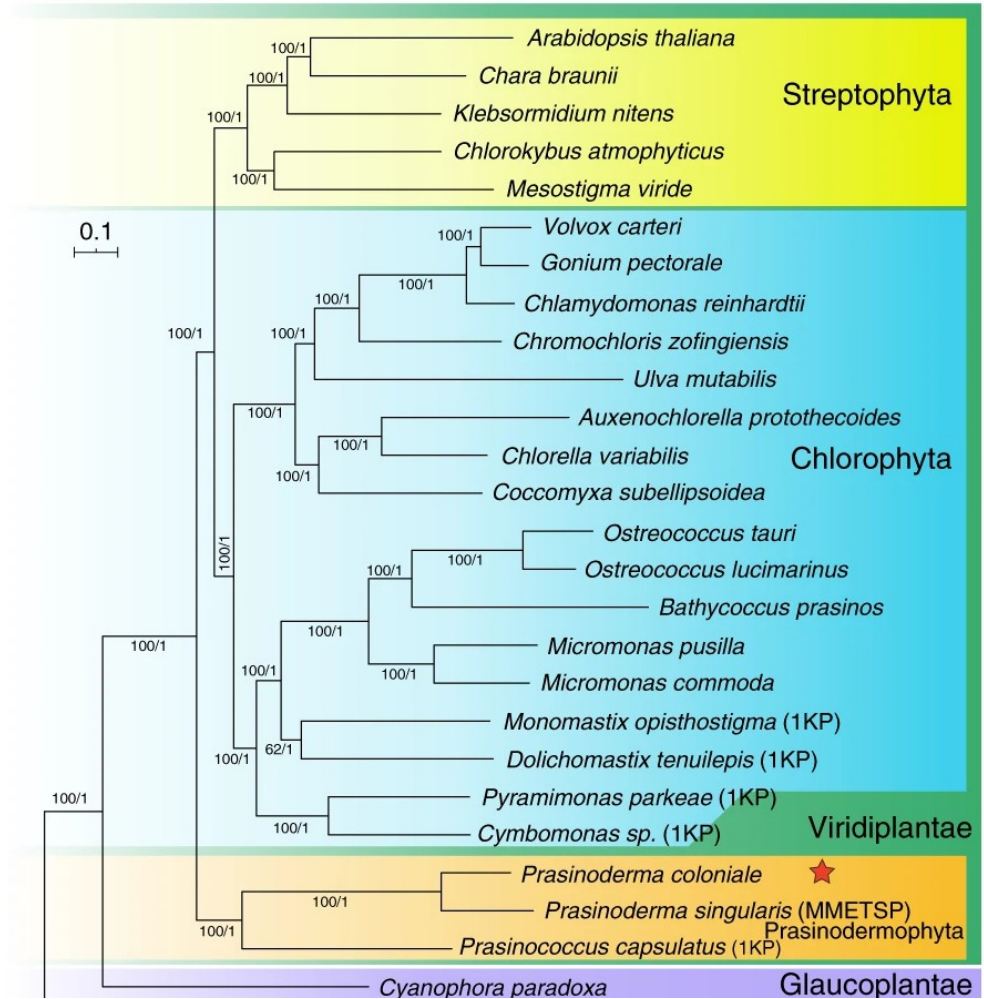
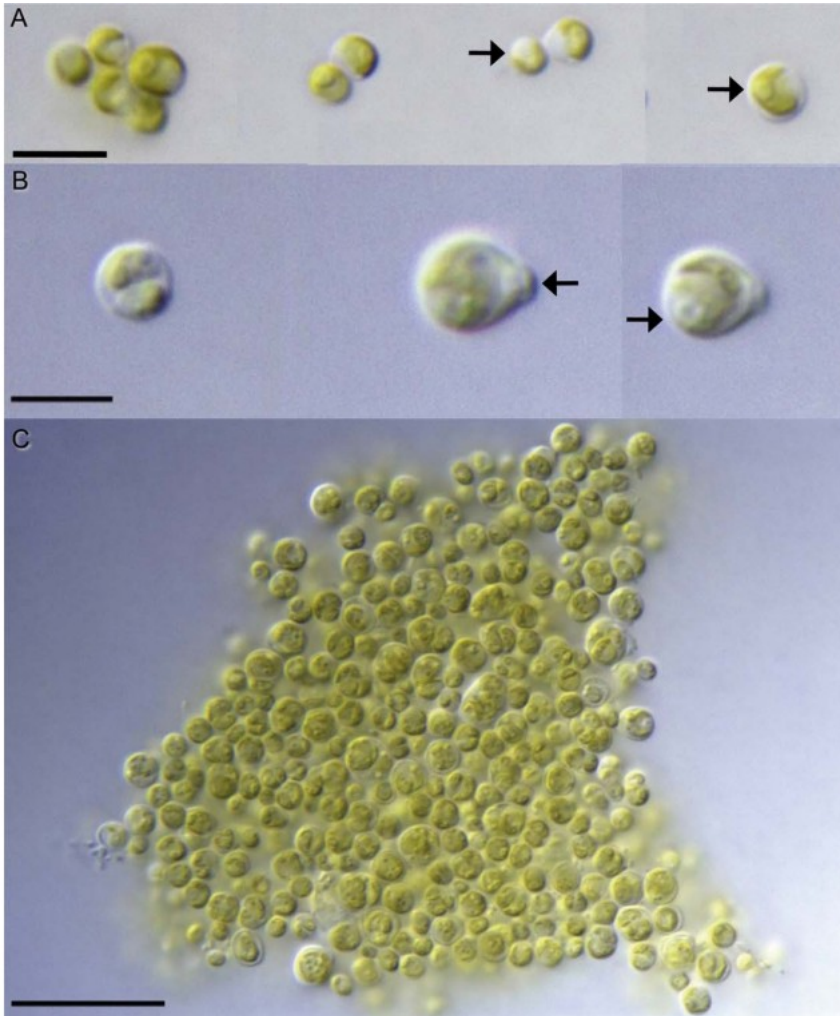


FIG. 2. Overview phylogeny of the green lineage (top) and spread of green genes in other eukaryotes (bottom). (Color figure available online.)

Prasinodermatophyta

the third lineage of Viridiplantae

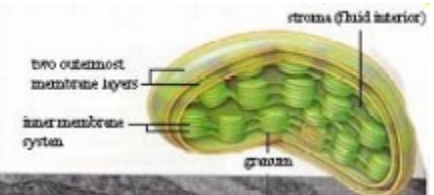


oceanic picoplankton (warm water regions)

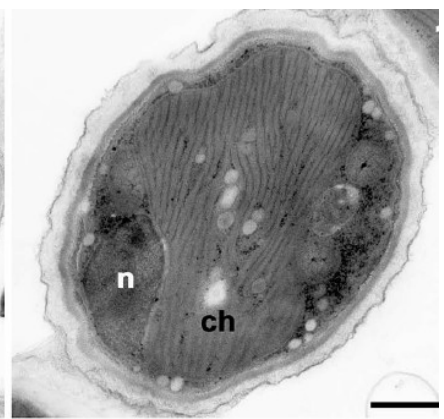
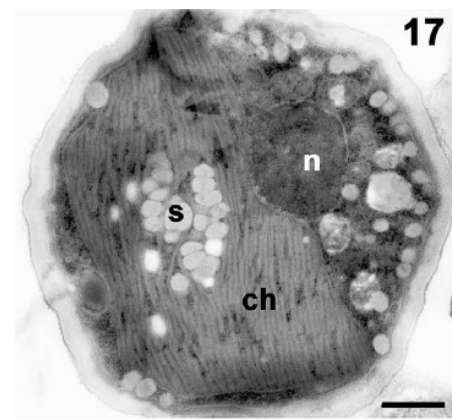
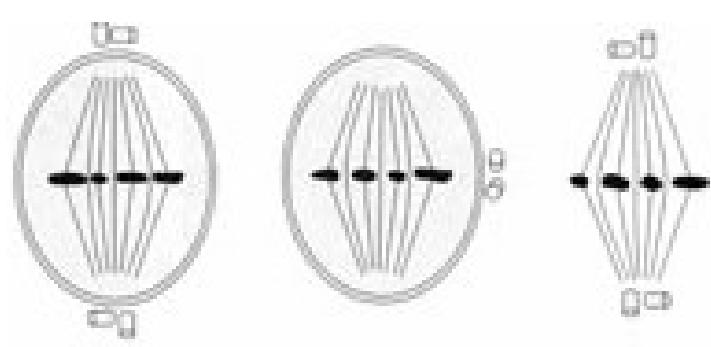
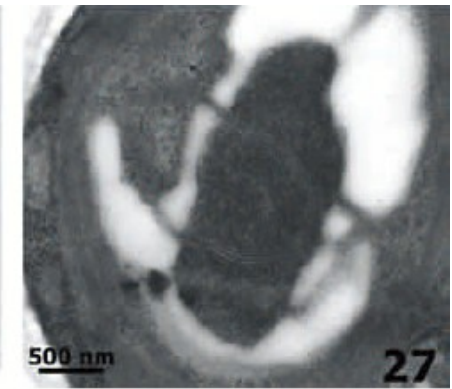
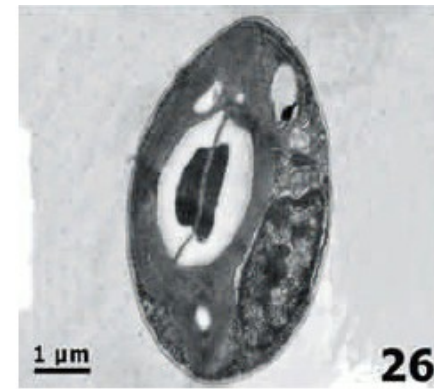
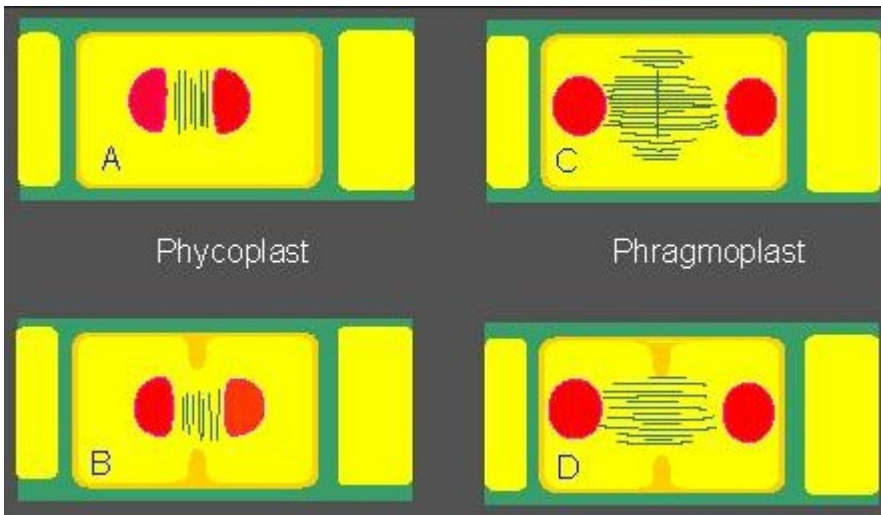
Li et al., 2020, Nat Ecol Evol

Jouenne et al., 2011, Protist

Kingdom: plants – *Plantae* (= *Archaeplastida*)
 Infrakingdom: ***Viridiplantae*** – green plants



two evolutionary lineages- *Chlorophyta* a *Streptophyta*



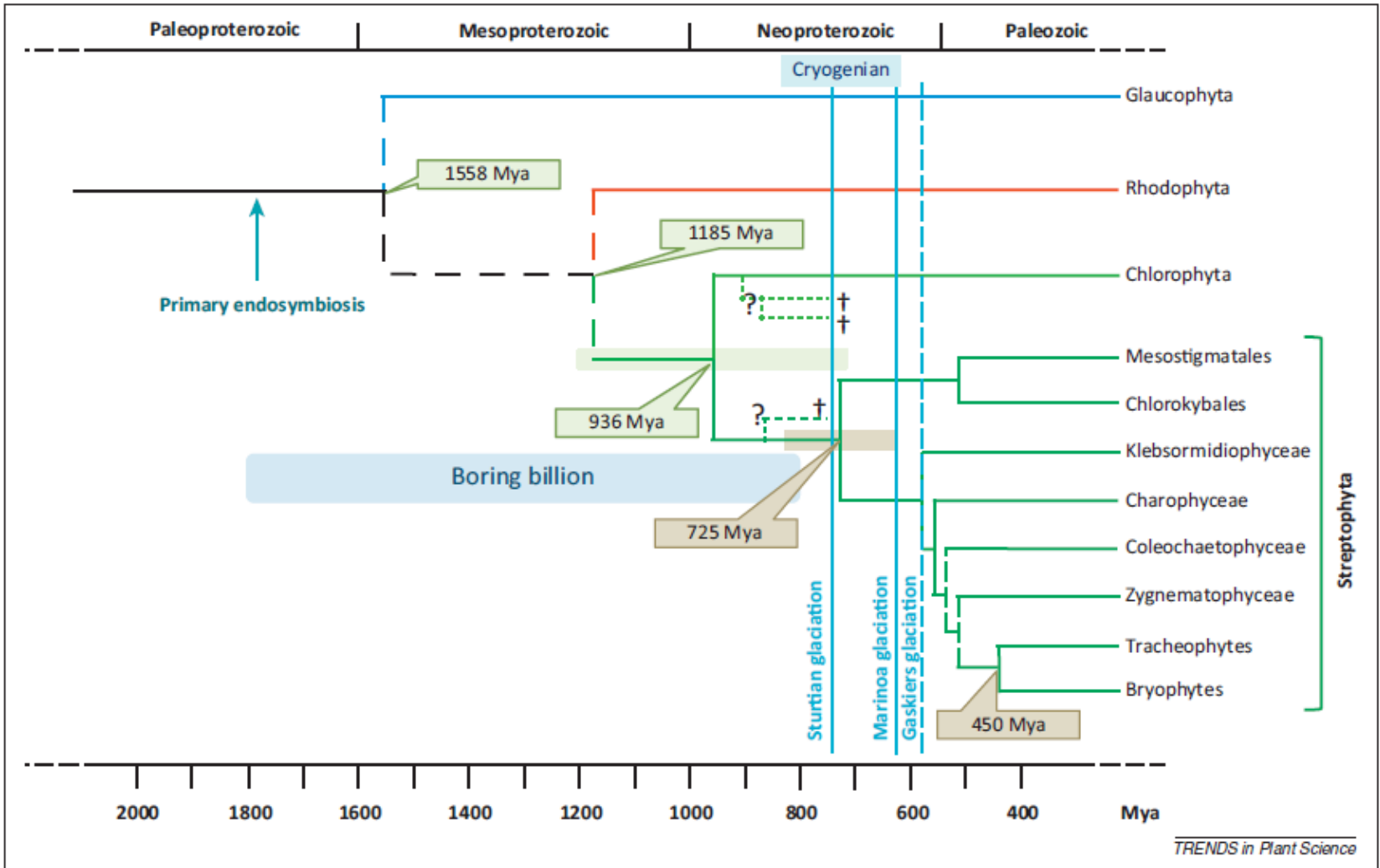


Figure 2. Plant evolution and major glaciation events. The scheme illustrates the evolution of plants. The streptophyte tree topology is based on [15,16]. Dashed lines indicate uncertain relationships. Nodes are dated according to TIMETREE (primary plastid groups, greenish boxed dates) or [25] (brownish boxed dates). The observed variation of divergence time estimates for chlorophytes and streptophytes is indicated with a greenish box, and the highest-probability density range given by [25] for the Mesostigma/other streptophytes split with a brownish box. The primary endosymbiosis is indicated by a vertical arrow. Today, extinct possible streptophyte and chlorophyte branches are marked with a question mark and terminate with a cross. Blue vertical lines indicate hard and soft (dashed line) snow ball states of the earth. The date for glaciation events and the boring billion period are based on [30]. The time frame for the Cryogenian period indicated is based on [36].

Evolution of green plant lineage

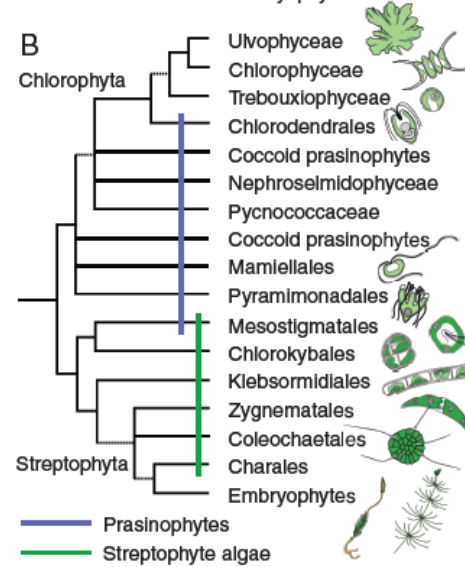
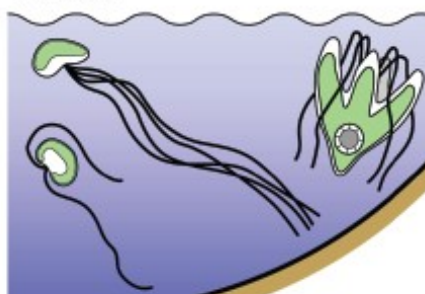


FIG. 1. Phylogenetic relationships among the major lineages of the Viridiplantae. Branches indicated by dotted lines are not well supported. (A) According to Lewis and McCourt (2004), and (B) based on unpublished, ongoing work by the authors. Some of the class names used by Lewis and McCourt (2004) have never been validly described, and for this reason we use order designations in (B) and throughout the text. The informal term 'prasinophytes' is commonly used for scaly green flagellates (Mesostigmatales and basal Chlorophyta).

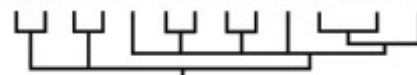
A. Neoproterozoic Era:

Cryogenian–Ediacaran Period
(approx. 850–540 MY ago)

Ocean



Ancient Chlorophyta:
marine and brackish
leiosphaerids and tasmanitids –
ancestors of extant 'prasinophytes'
(scaly green flagellates and coccoids)



CHLOROPHYTA

Freshwater



Ancient Streptophyta:
unknown ancestors of extant streptophyte lineages –
freshwater flagellates, coccoids,
sarcinoids and filaments



ancestors of *Mesostigma*
and *Chlorokybus* (scaly
freshwater flagellates)

STREPTOPHYTA

C. Paleozoic to Mesozoic Era:
Carboniferous–Cretaceous Period
 (approx. 360–65 MY ago)

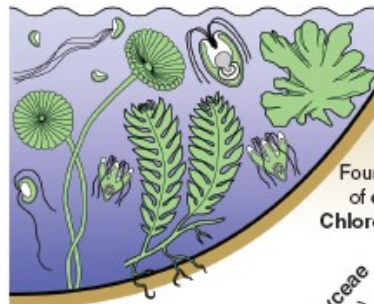
Ocean

- **Ulvophyceae** (e.g. Ulvales, Dasycladales, Bryopsidales, Cladophorales)
- **'Prasinophytes'** (e.g. Mamiellales, Pyramimonadales)

Freshwater

- **Chloro- and Trebouxiophyceae**
 (Chlorophyceae dominated freshwater phytoplankton communities after the Permian/Triassic mass extinction 250 MY ago)
- **Streptophyte algae** (mainly Charales and Zygnematales)
- **Embryophyte water plants** (first aquatic angiosperms: early Cretaceous Period)

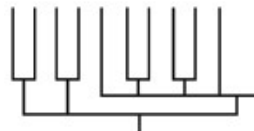
B. Paleozoic Era:
Ordovician–early Devonian Period
 (approx. 490–400 MY ago)



Four clades of derived Chlorophyta:



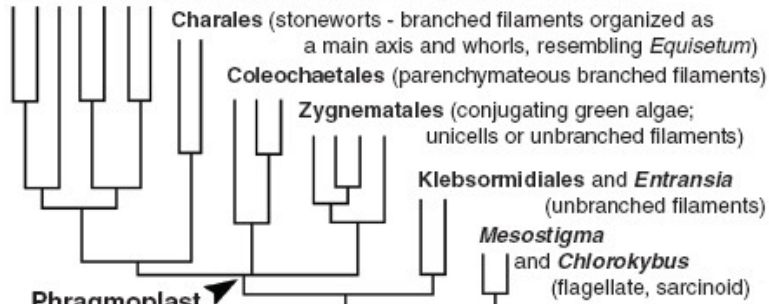
Ancient Chlorophyta:
 several clades of marine 'prasinophyte' flagellates and/or coccoids



CHLOROPHYTA

- **Rapid adaptive radiation of Chloro- and Trebouxiophyceae**
- **Gradual reduction in diversity and abundance of streptophyte green algae.**
 (exception: evolutionary diversification and speciation of the Zygnematales during the Jurassic and Cretaceous Period)

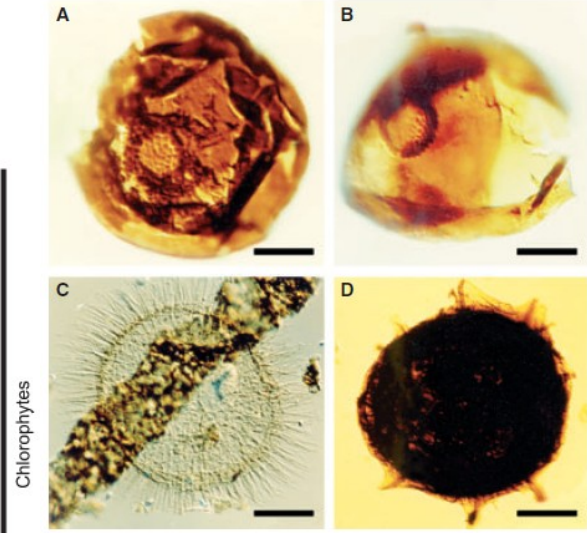
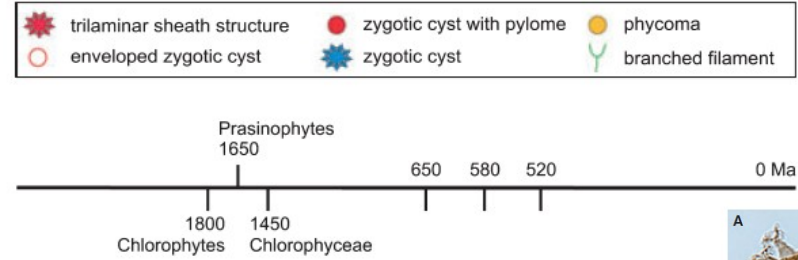
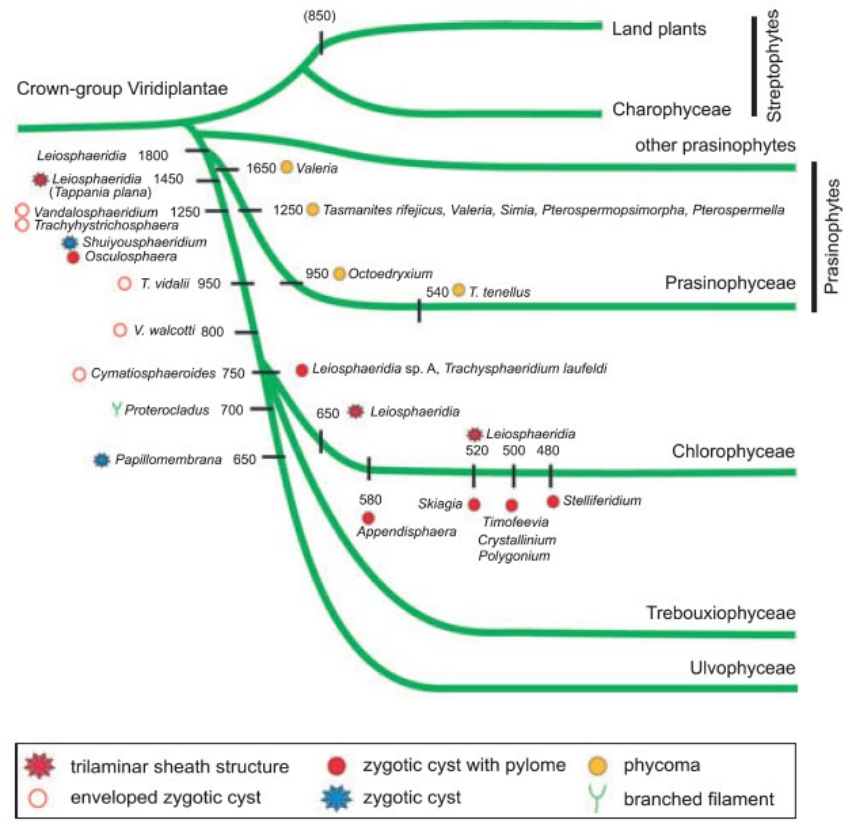
EMBRYOPHYTA (bryophytes, lycophytes, ferns, spermatophytes)



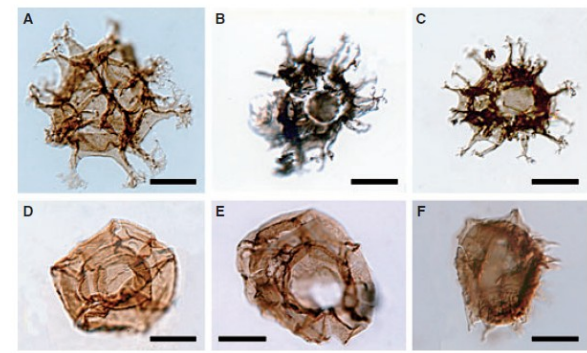
Phragmoplast

STREPTOPHYTA

TEXT-FIG. 4. Phylogeny of the Viridiplantae (modified from O’Kelly 2007 and Moustafa *et al.* 2009). The chronologically arranged, earliest appearance of microfossils interpreted to be green microalgae constrains the origin of the classes Prasinophyceae, Chlorophyceae and Ulvophyceae. The first appearance data of the microfossils are compiled from sources cited in the text. Terrestrial expansion of biota at c. 850 Ma is according to Knauth and Kennedy (2009). The time axis is drawn not in scale and shows the time of the origins of the Chlorophytes prior to c. 1800 Ma, Prasinophyceae c. 1650 Ma, Chlorophyceae c. 1450 Ma, and major radiations of phytoplankton at c. 650, 580 and 520 Ma.



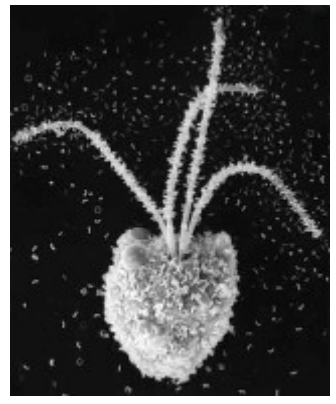
TEXT-FIG. 1. Neoproterozoic microfossils. A, B. *Leiosphaeridia* sp. A, with pylome. Chuar Group, the Great Canyon, northern Arizona (Cryogenian). A, Specimen LO 5658. B, Specimen LO 5659. C, *Appendisphaera grandis*. Khamaka Fromation, eastern Siberia (Ediacaran). Specimen PMU-Sib.1-L/27/1. D, *Trachyhystrichosphaera vidalii*. Khajpakh Foramtion, eastern Siberia (Tonian). Specimen PMU-Sib.6-N/43/3. Scale bars represent 10 μ m for A, 7 μ m for B, 38 μ m for C and 46 μ m for D. All are light photomicrographs.



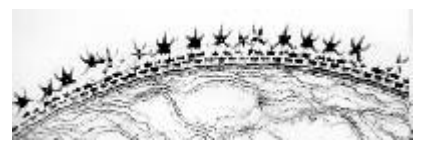
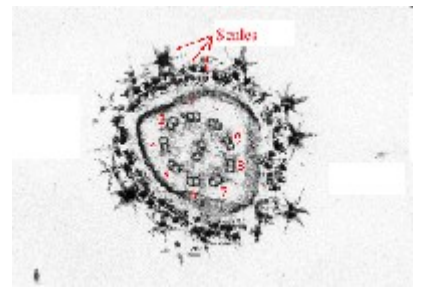
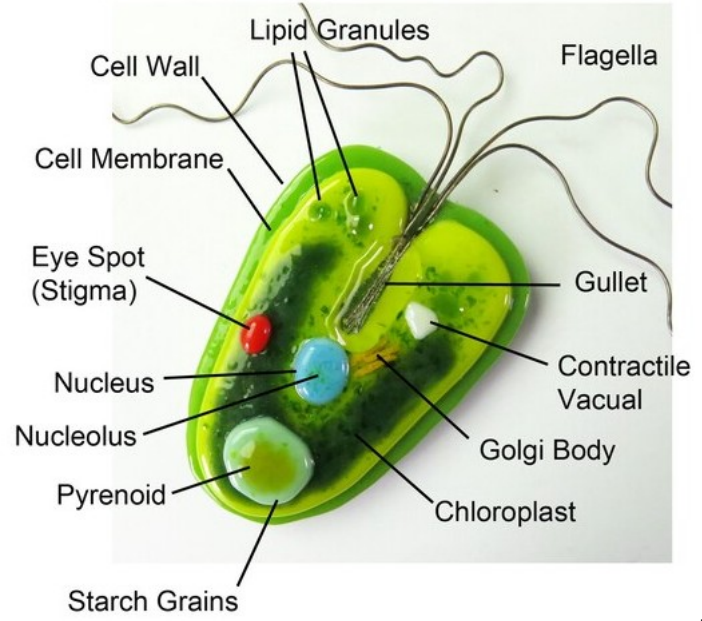
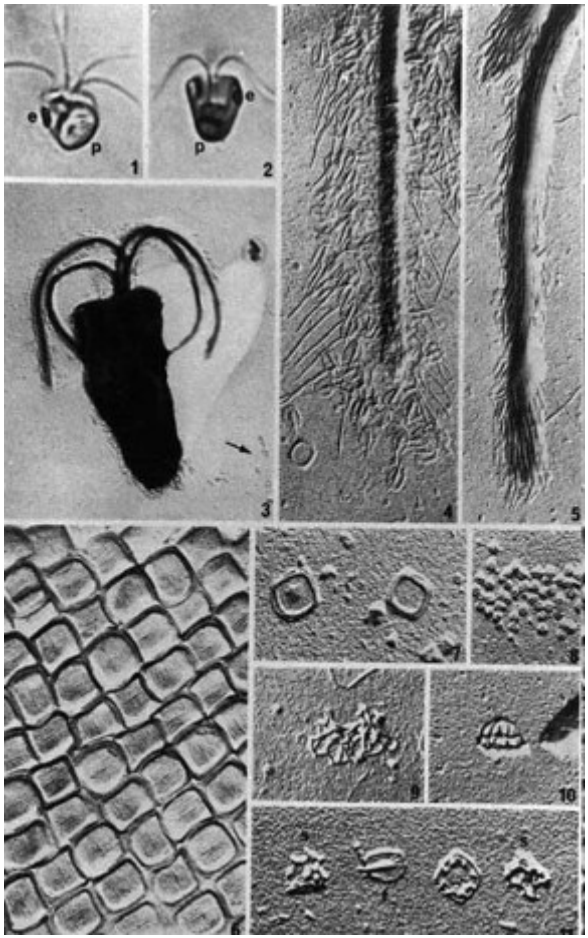
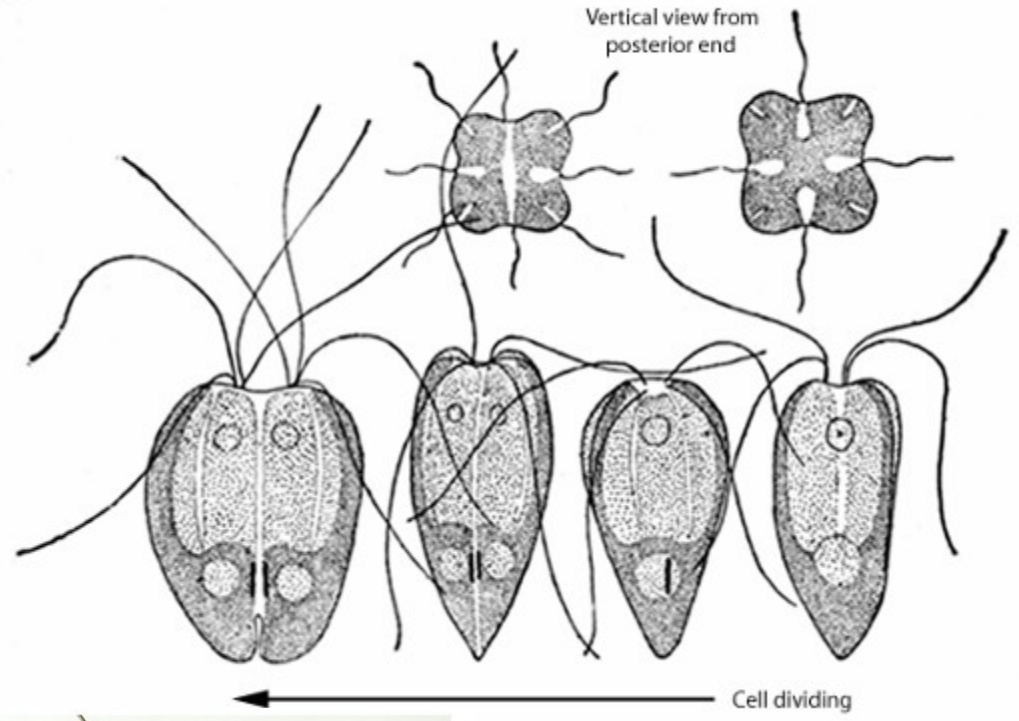
TEXT-FIG. 3. Cambrian microfossils. A–C. *Timofeevia lancare*. A, Immature stage, specimen SP-2009.10.A/10. B, Specimen with ontogenetically defined pylome, ZL05-13b-L/43. C, Specimen with opened pylome, ZL05-13-G/20/4. D–F. *Cristallinium camabriense*. D, Specimen in immature stage, SP-2009.11.C/15/2. E, Specimen with two open pylomes, SP-2009-12.V/16. F, *Polygonium varium* containing the endocyst, specimen PO106-11N2-A/15/4. Specimens A, D–F, Oville Formation, Cantabrian Mountains, northern Spain; specimens B, C, Playón Formation, Ossa-Morena Zone, central Spain. Scale bars represent 15 μ m for all specimens.

Prasinophyceae

Pyramimonas



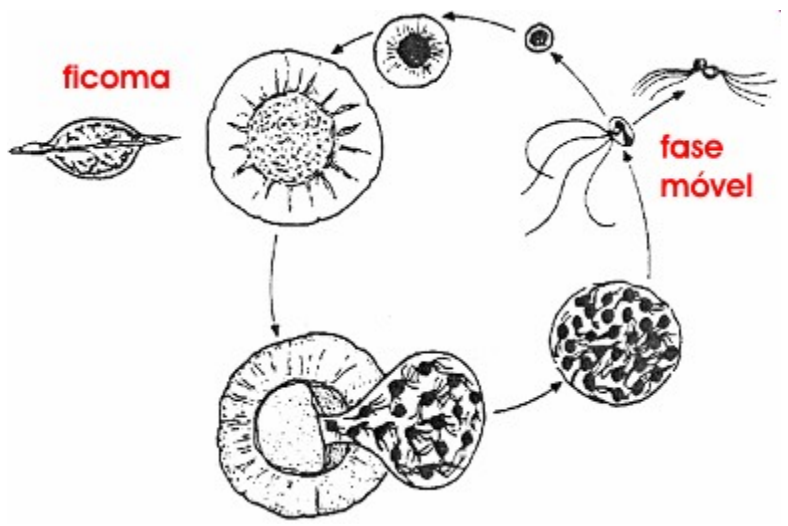
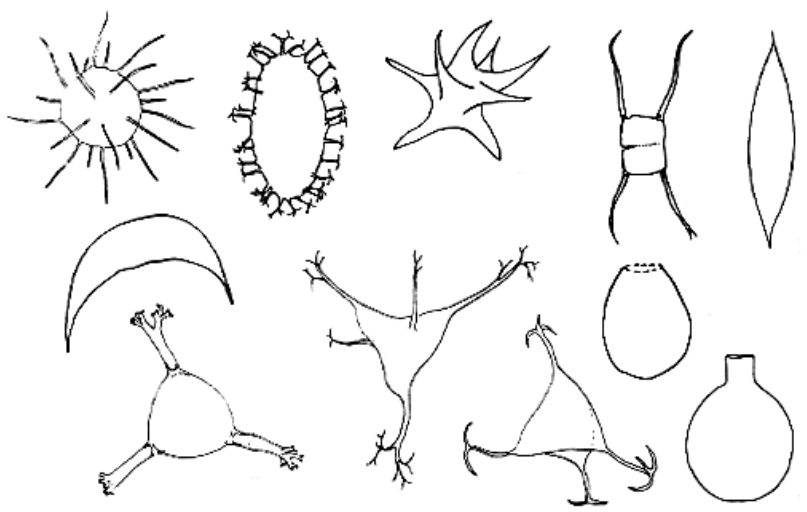
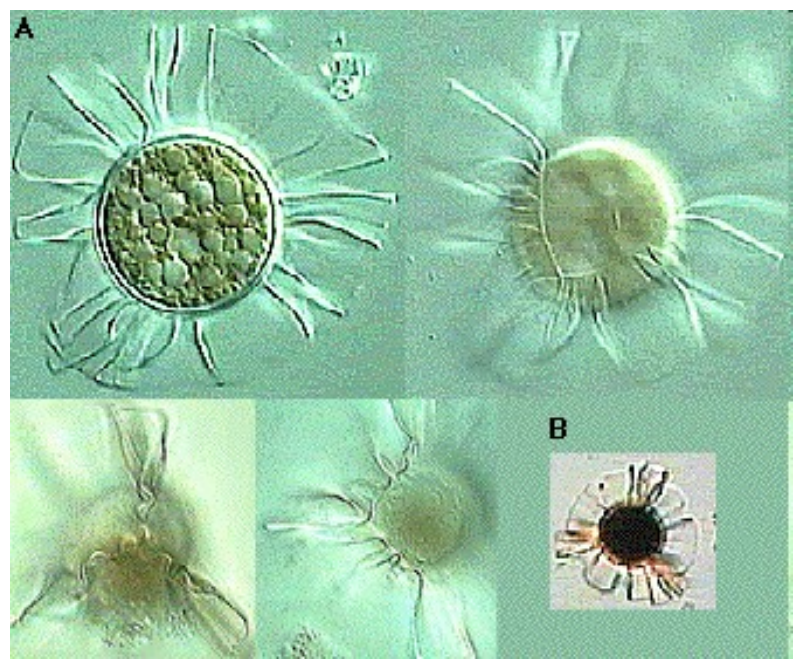
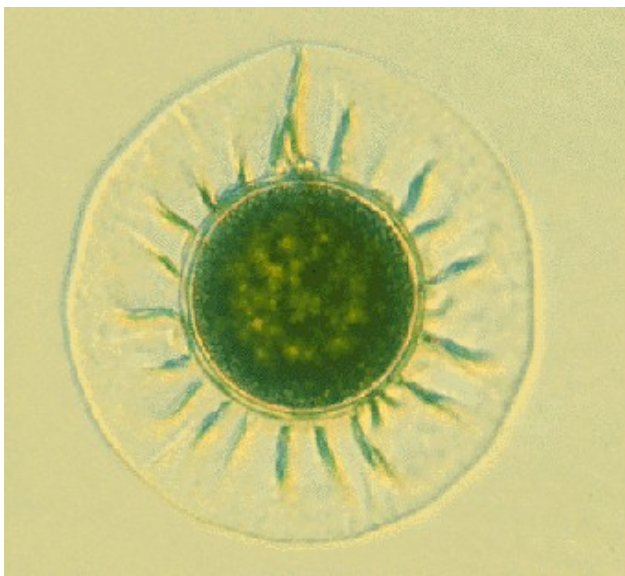
Pyramimonas



source: etsy.com

freshwater and marine flagellates

Prasinophyceae
Pterosperma

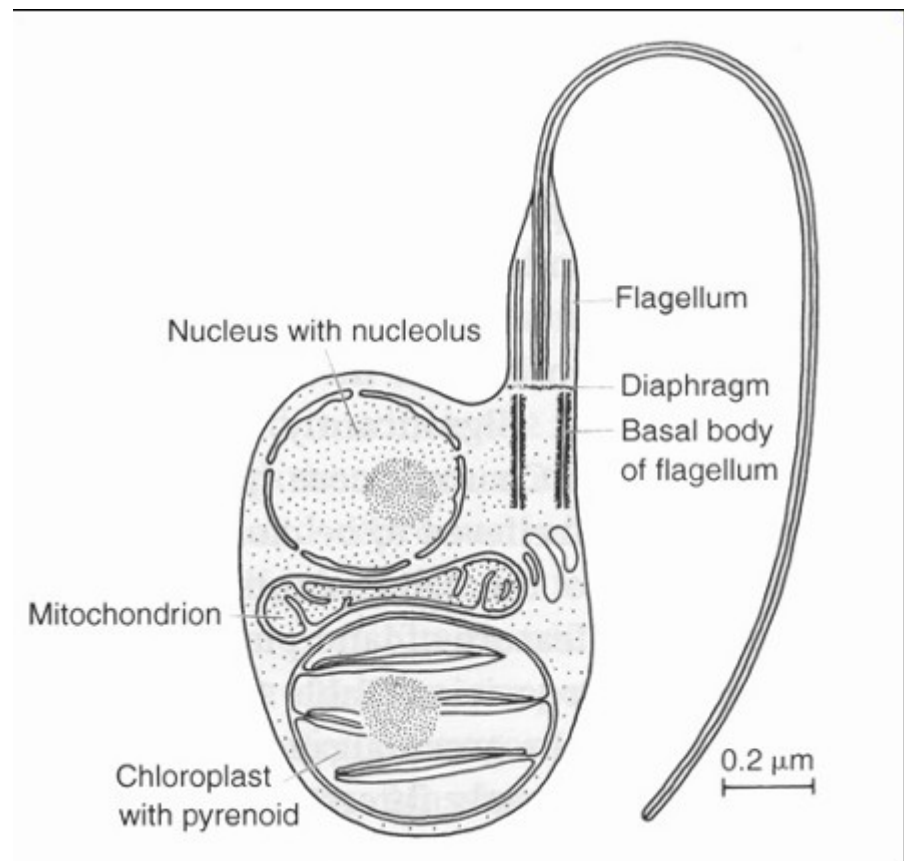
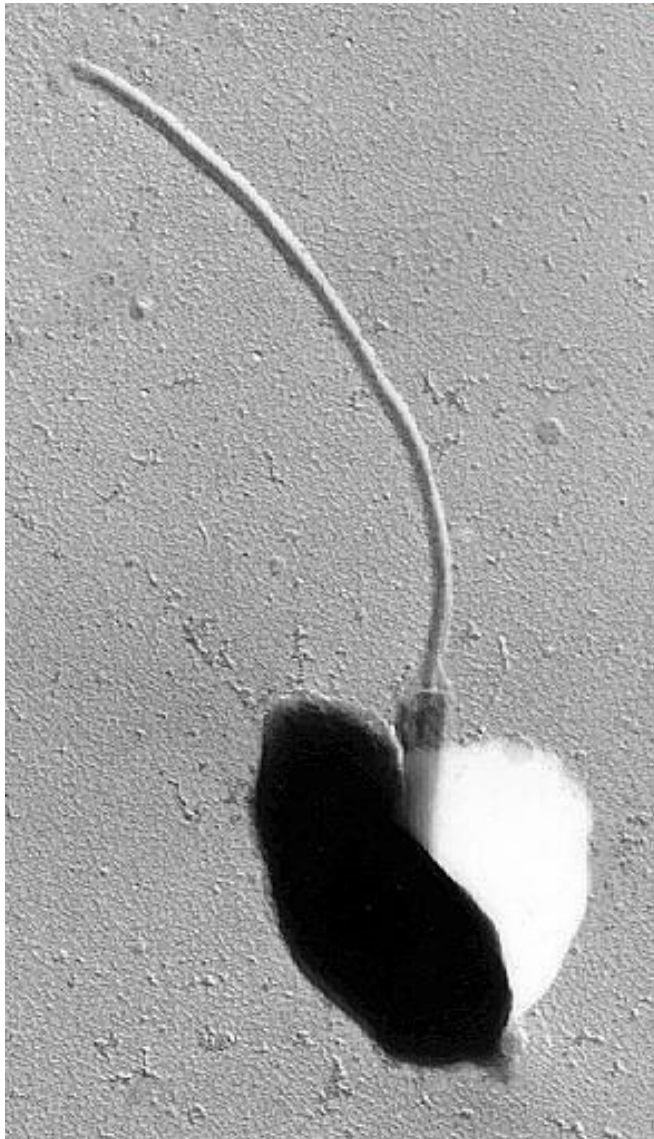


Acritarcha – very probably (partially) prasinophycean algae

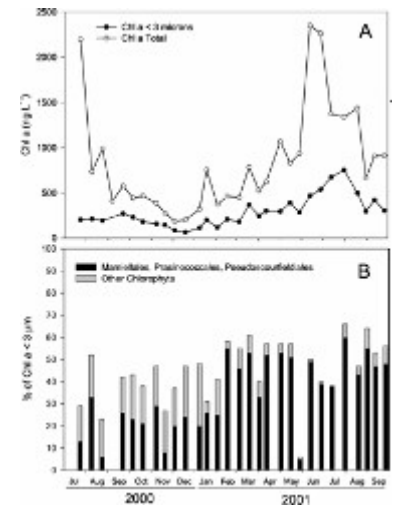
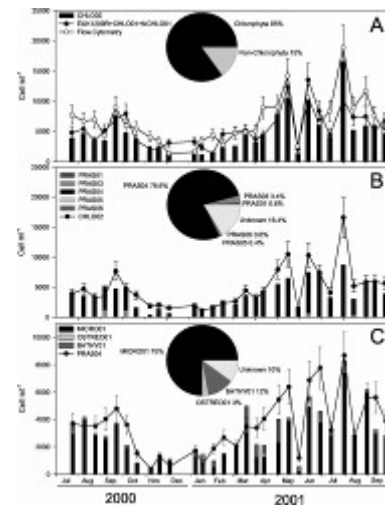
marine phytoplankton – cold seas

Mamiellophyceae

Micromonas



often as one of the dominants in picoplankton of cold shelf seas (i.e. North Sea)



Mamiellophyceae

Ostreococcus

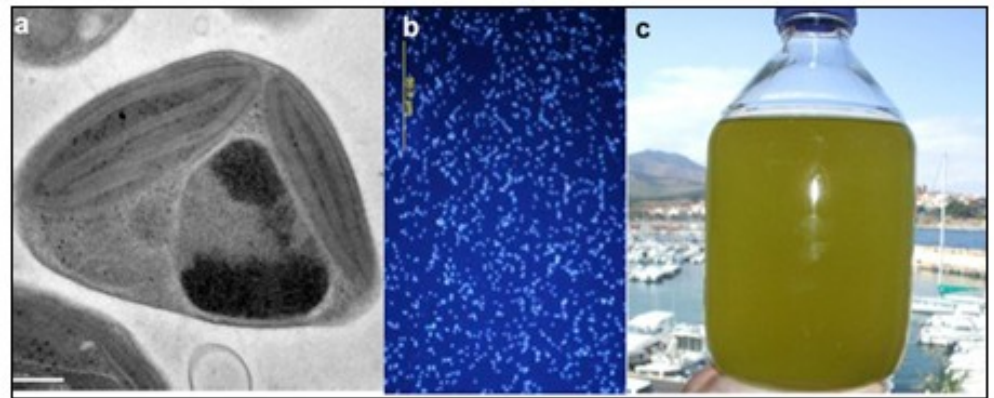
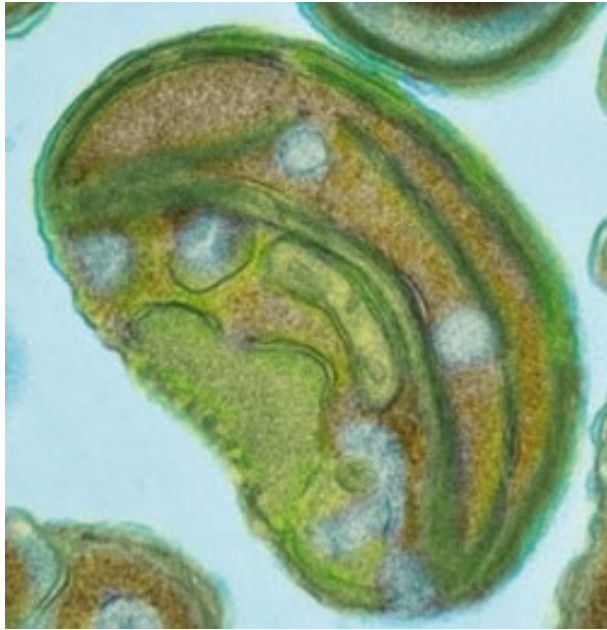
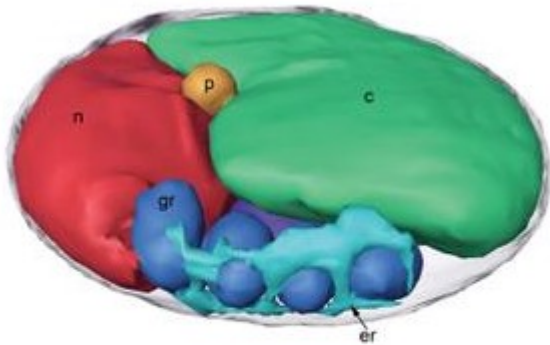


Photo: by Moreau lab, CNRS
a. TEM picture of *Ostreococcus tauri*
(ML Escande, Oceanological Observatory of Banyuls, France);
b. DAPI staining of *O. tauri* cells;
c. 2 Liter culture of *O. tauri* isolated from the bay of Banyuls seen in the background.



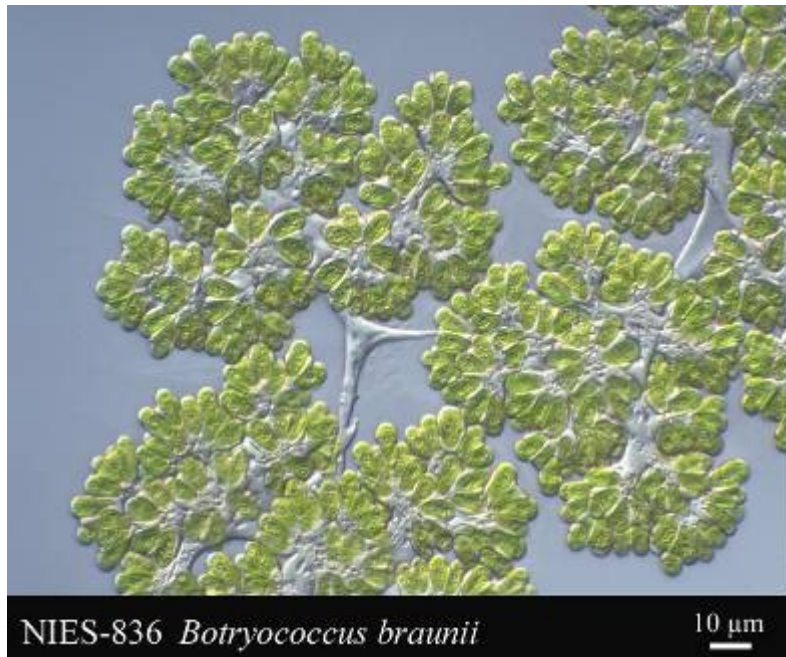
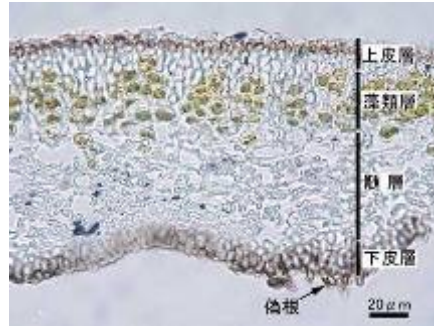
Electron cryotomographic reconstruction of an *O. tauri* cell. n = nucleus; c = chloroplast; p = peroxisome; er = endoplasmic reticulum. Source



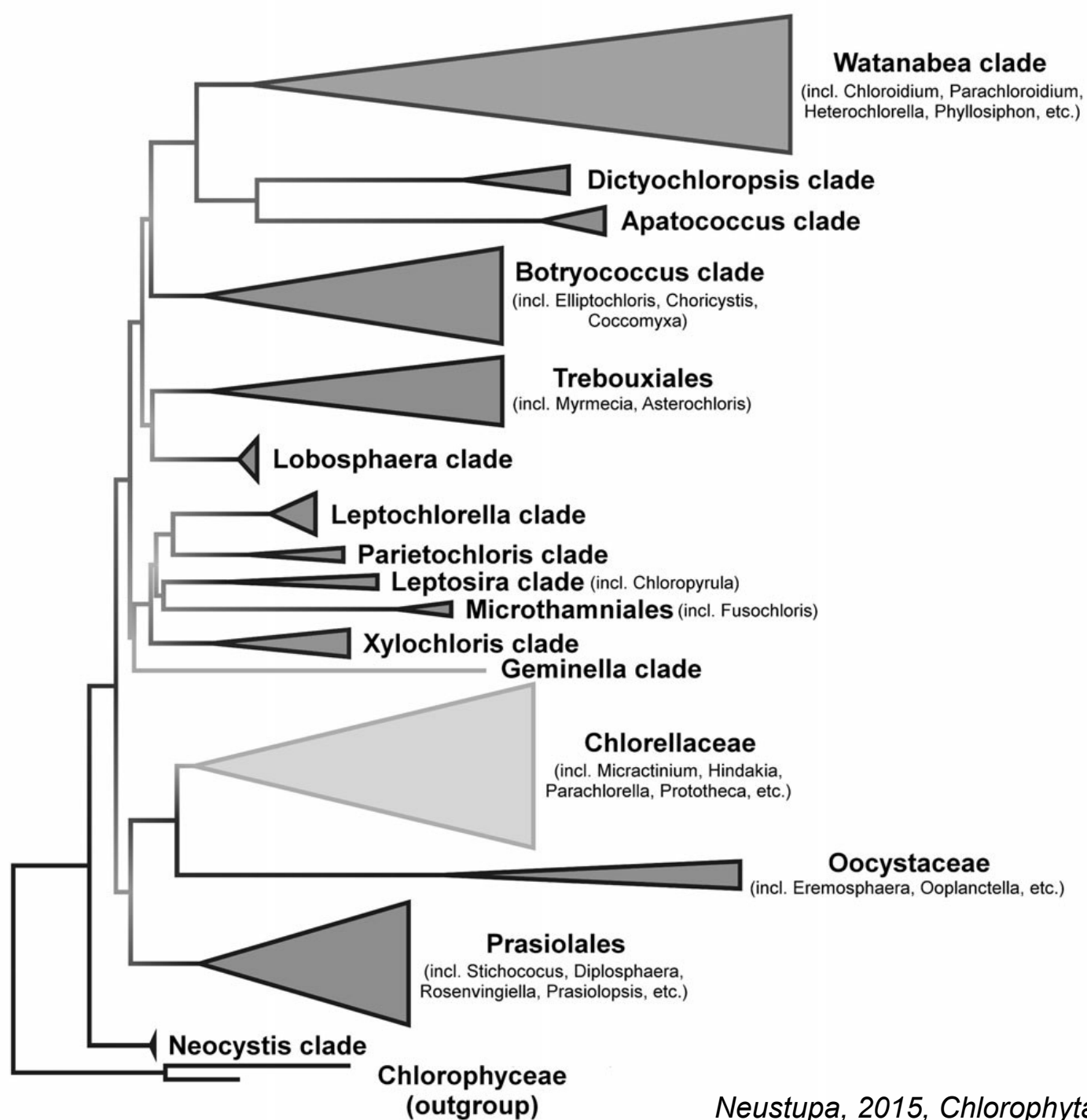
- described in 1994 – plankton in Étang de Thau
- the smallest known free-living eukaryotic organism

Třída: **Trebouxiophyceae**

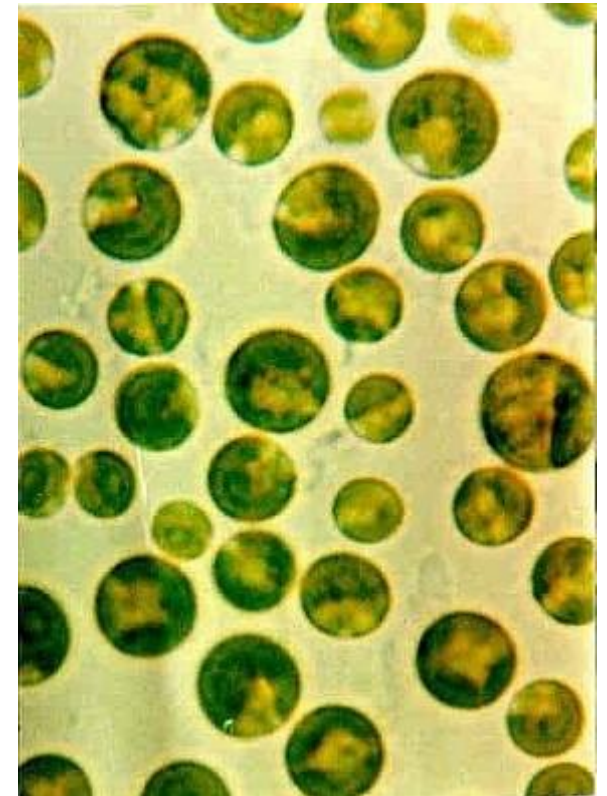
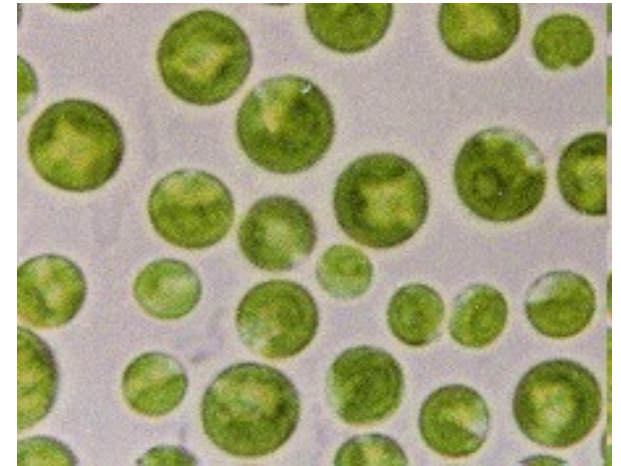
Trebouxia – the most frequent fotobiont



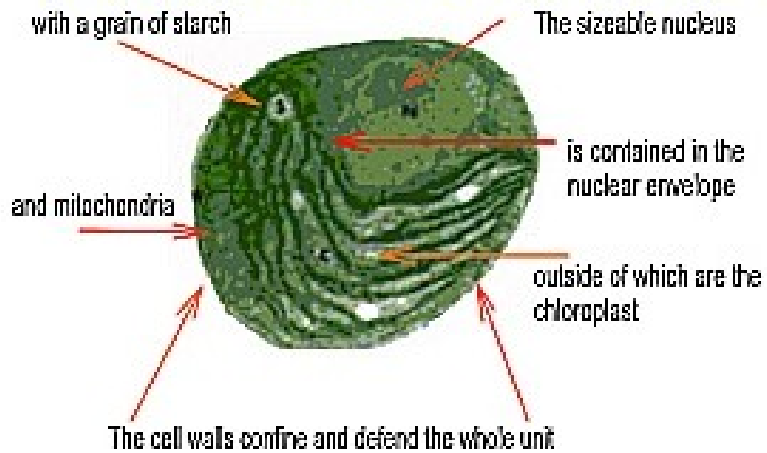
multiple successful transitions to terrestrial habitats but also several freshwater lineages

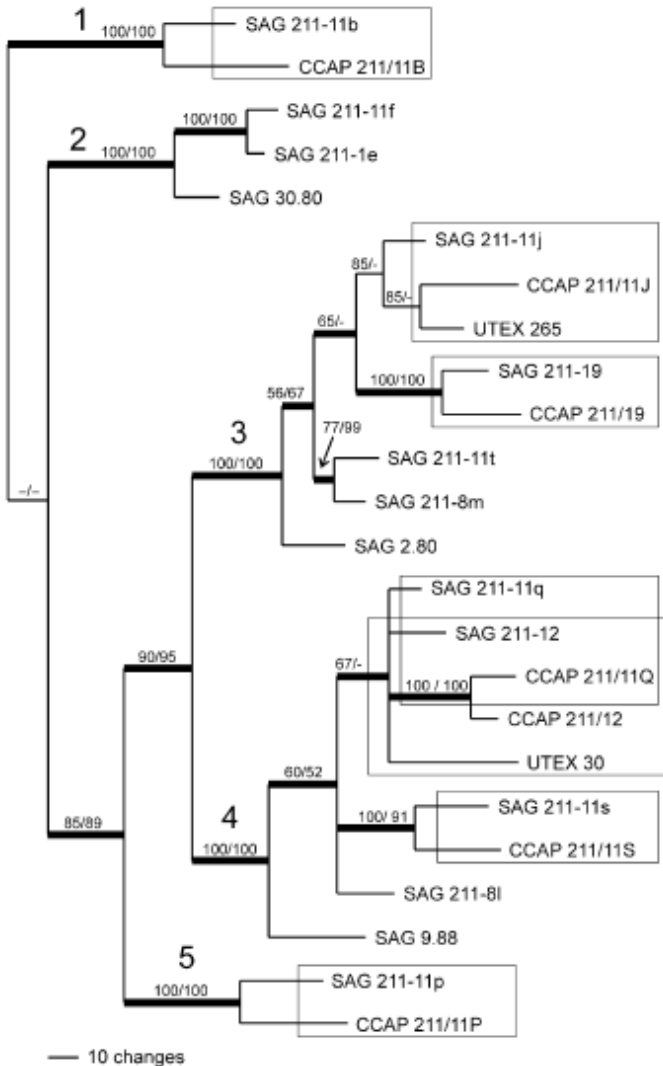
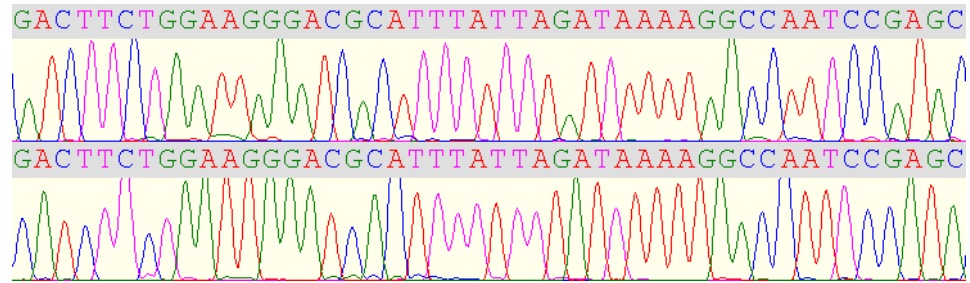


"Chlorella"

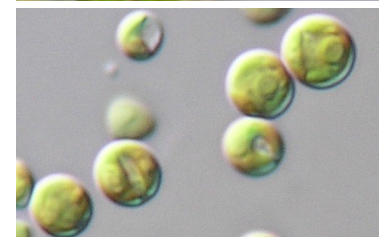
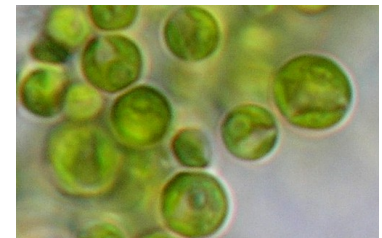
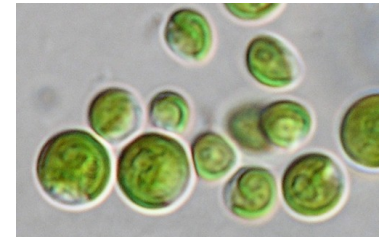
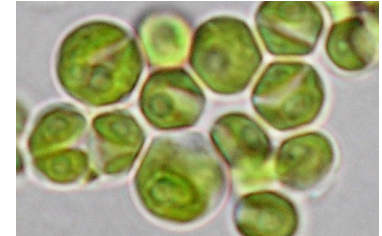
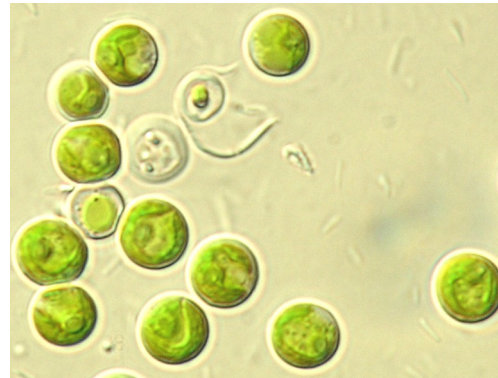


The Structure of a Chlorella Cell





Chlorella vulgaris



cryptic diversity hampers
defining species by
traditional (morphological) methods

Müller et al., 2005, *J. Phycol.* 41:1236-1247.

AFLP, MP phylogenetic tree

chlorella business...



Chlorella vs. Micractinium

herbivore-driven phenotypic plasticity

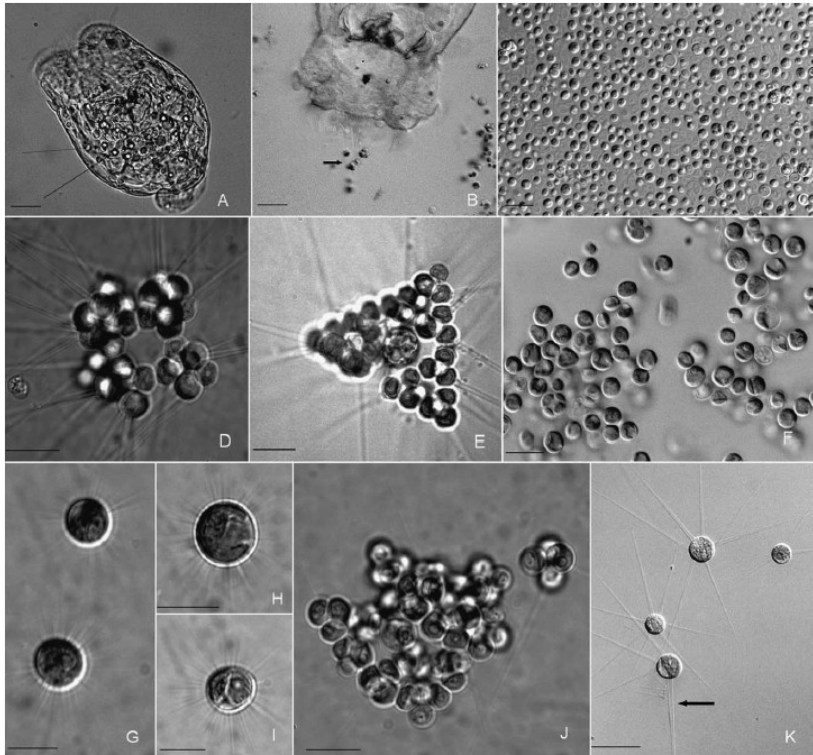


Figure 3. Organisms used in this study. Scale bars: A, B, 50 μ m; C, 25 μ m; D–F, J, K, 10 μ m; G–I, 5 μ m. A. *Brachionus calyciflorus* with ingested solitary, not-bristled cells of *Micractinium pusillum* (CCAP 248/5) (arrowhead). B. *Brachionus calyciflorus* ingesting coenobia of *Micractinium pusillum* (CCAP 248/5) (arrowhead). C. *Chlorella vulgaris* (CCAP 211/80). D–K. *Micractinium pusillum* under field and culture conditions. D. Cuboidal colony of eight spherical subcolonies; sampling site: Final Sewage Pond Nakuru, Kenya. E. Tetrahedral colony; sampling site: Speicher Radeburg, Germany. F. Strain CCAP 248/5, solitary cells without bristles in dense pre-cultures which can be used for the bristle induction experiments. G–J. Cultures from bristle induction experiments. G. Strain CCAP 248/7, tiny bristled solitary autospores. H. Strain SAG 13.81, bristled solitary vegetative cell. I. Strain CCAP 248/7, bristled mother cell with two autospores. J. Strain SAG 13.81, spherical colonies with or without bristles. K. Strain CCAP 248/5 after 24 h under grazing test conditions. The solitary cells developed strong bristles (arrow).

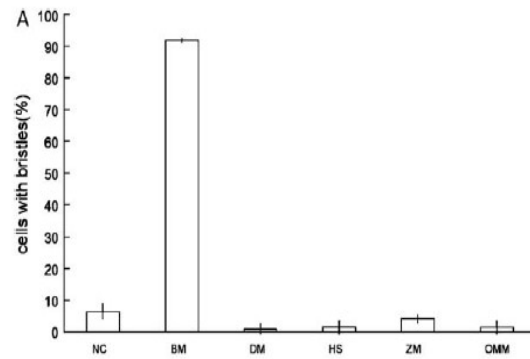
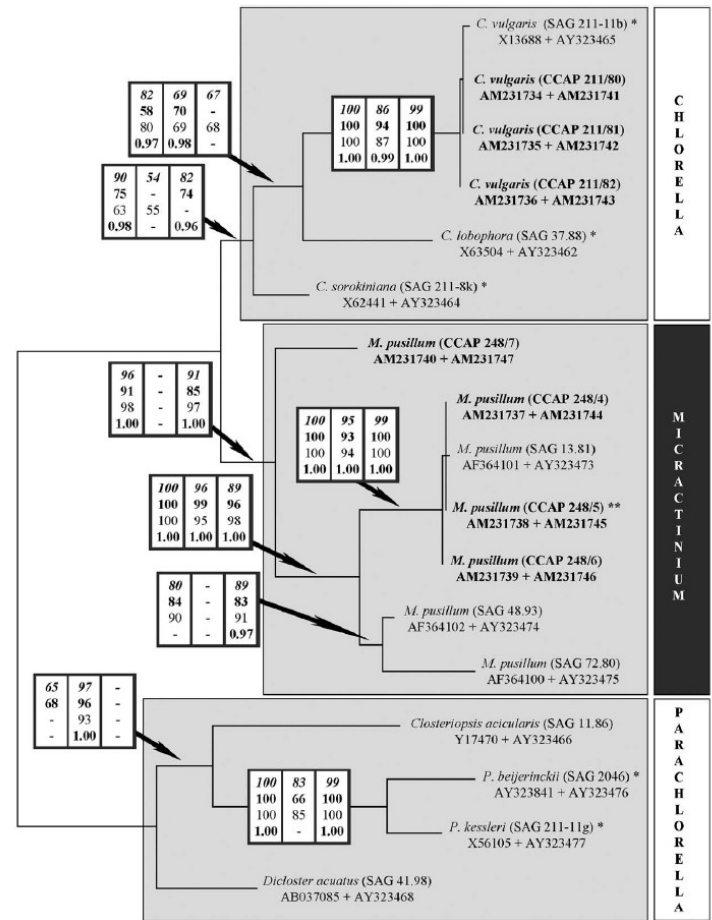
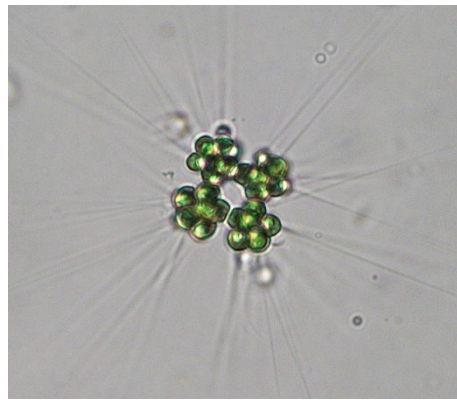
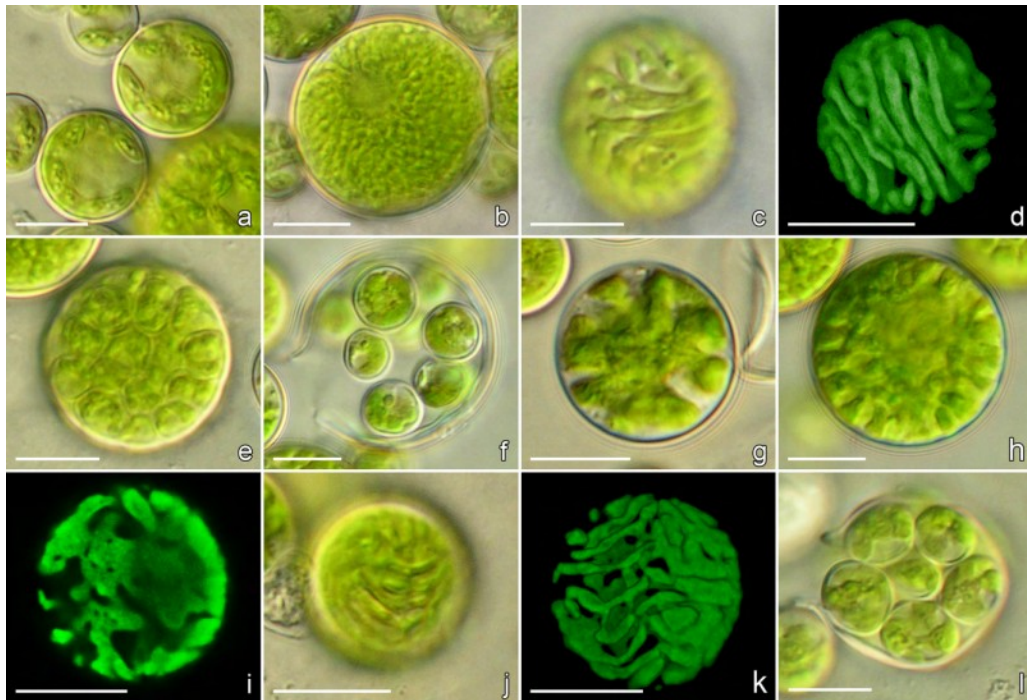
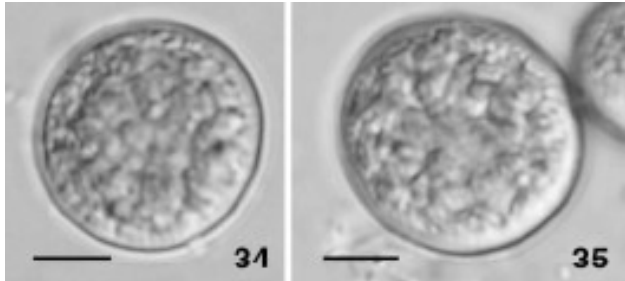
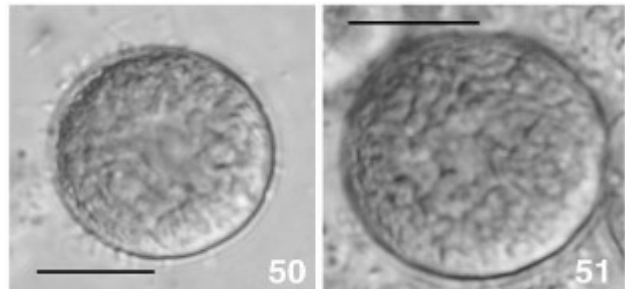
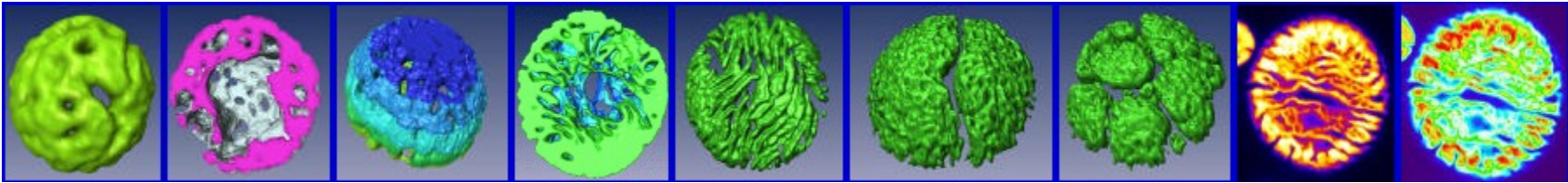


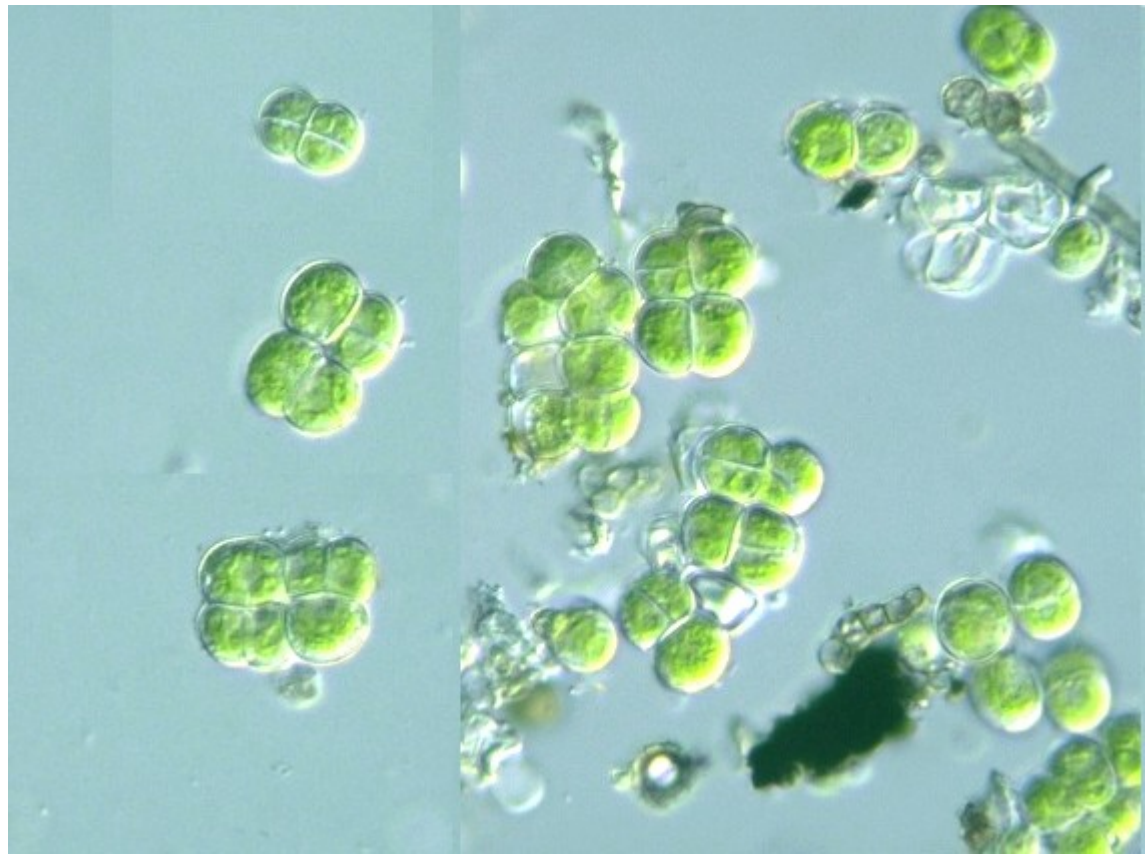
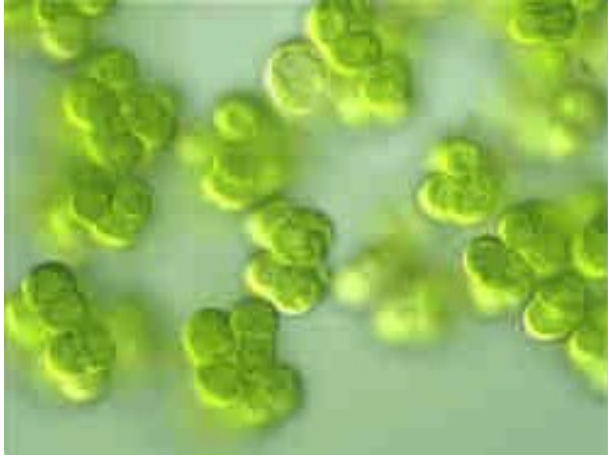
Figure 4. Influence of different test media on induction of bristle development in *Micractinium pusillum*. A. Percentage of bristled cells in strain CCAP 248/5 under the influence of different media, expressed as mean values (± 1 SD, $n = 3$). Abbreviations: NC = Negative control; BM = *Brachionus* culture medium filtrate; DM = *Daphnia* culture medium filtrate; HS = Humic substances; ZM = Z-medium; OMM = *Micractinium* culture medium from dense pre-culture. B. Percentage of bristled cells in

Genera *Symbiochloris* and *Dictyochloropsis* as examples of semicryptic evolutionary lineages



terrestrial biotopes – biofilms on bark and stones

Apatococcus

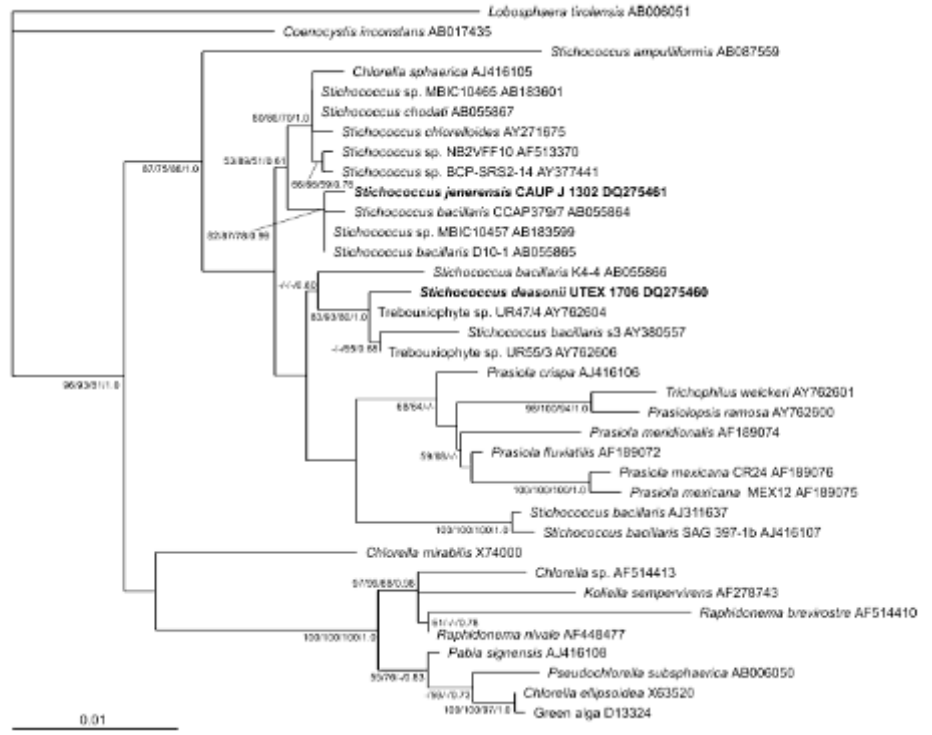
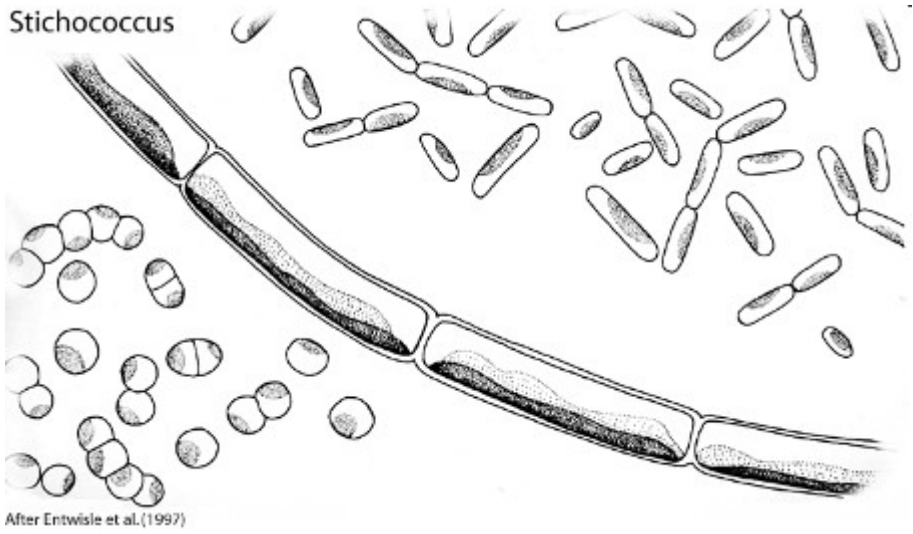
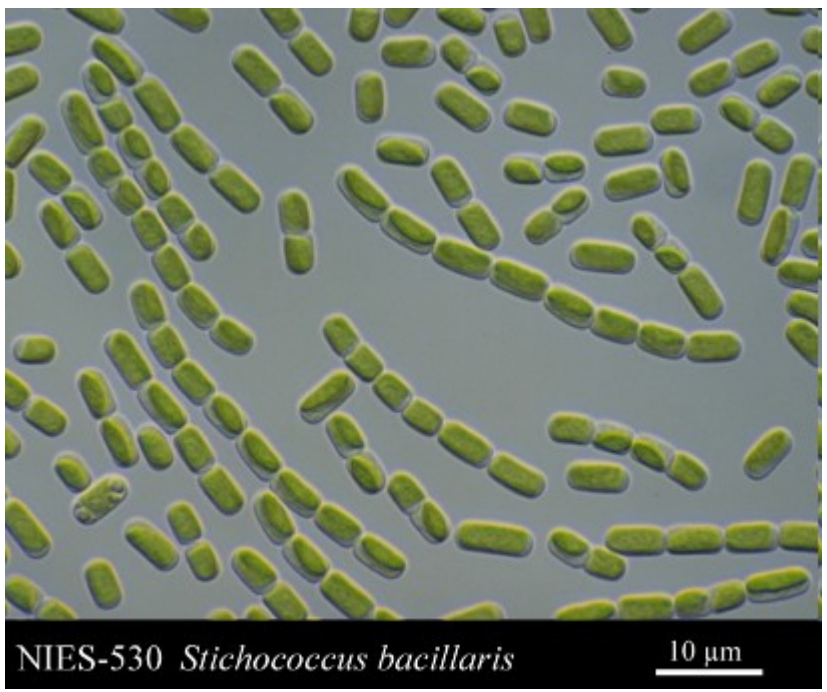


„the most frequent“ alga in the world?

sarcinoid, colonial thallus made by coccoid cells

mixotrophic nutrition → capability of surviving in extremely shaded microhabitats

Stichococcus (Prasiolales)



Prasiola (Prasiolales)



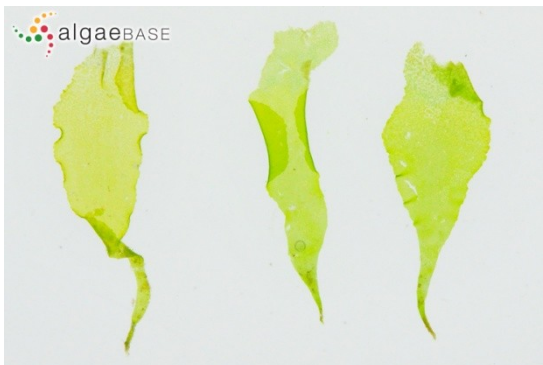
P. stipitata



P. crispa



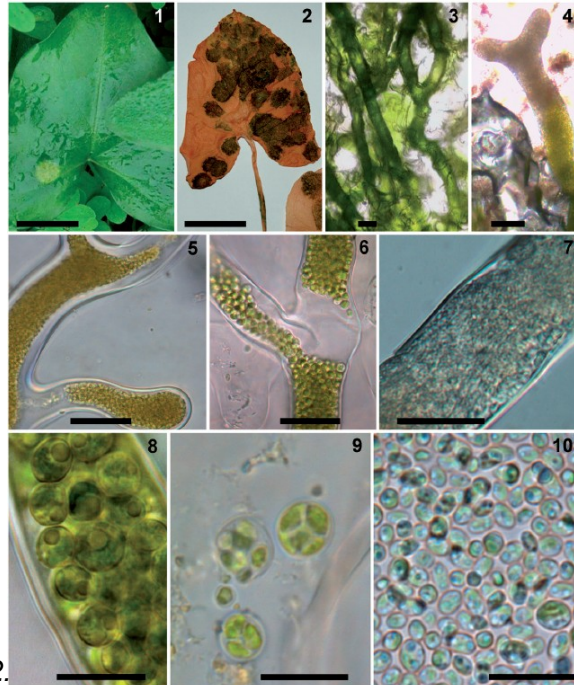
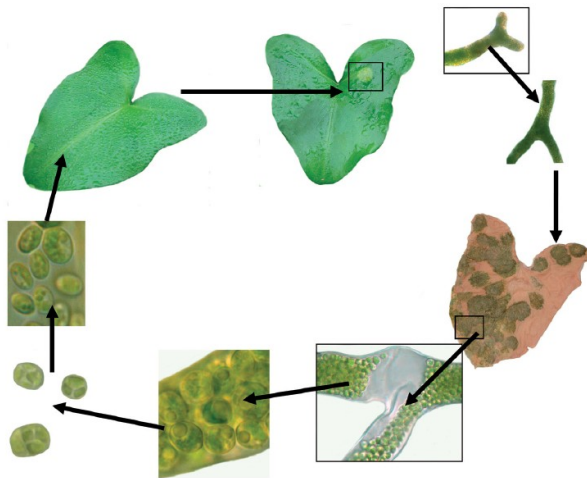
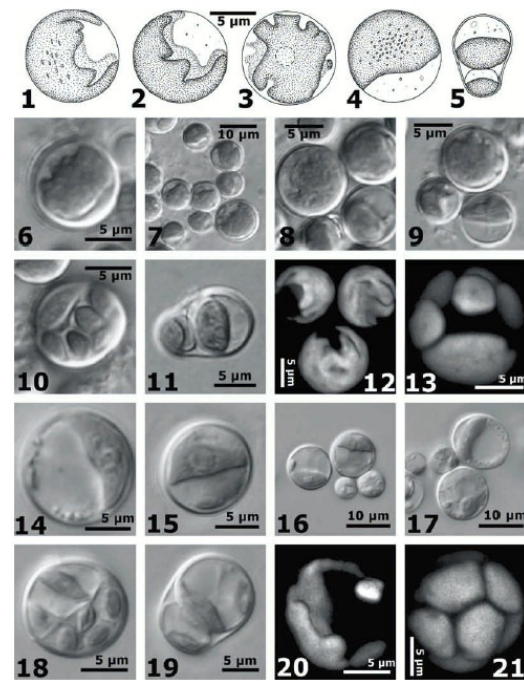
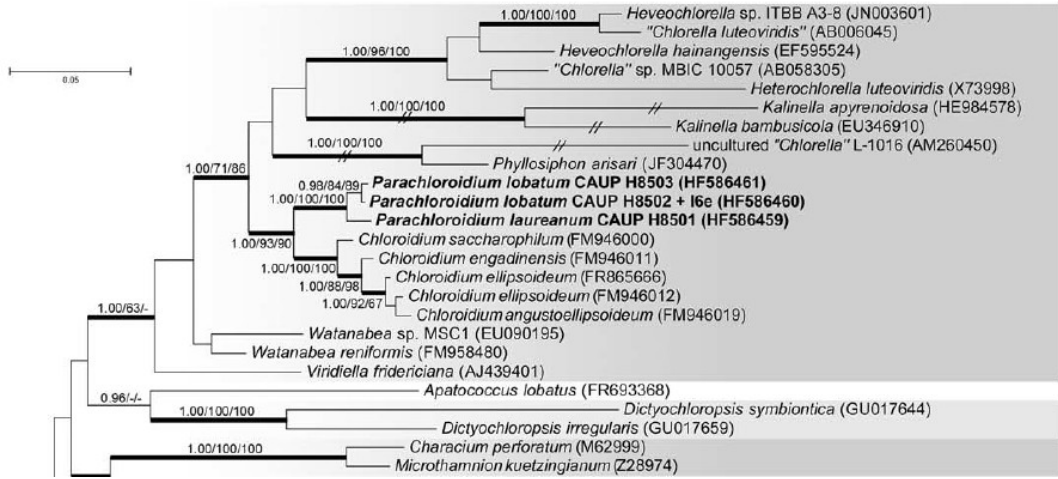
aquatic and terrestrial multicellular and macroscopic thalli in Trebouxiophyceae



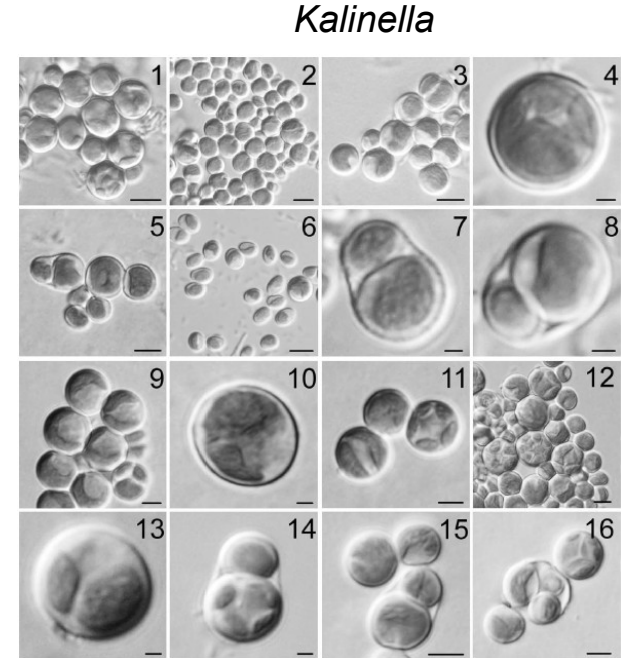
P. stipitata

Watanabea clade

(genera *Kalinella*, *Chloroidium*, *Parachloroidium*, *Phyllosiphon*, etc.)



Phyllosiphon



Parachloroidium

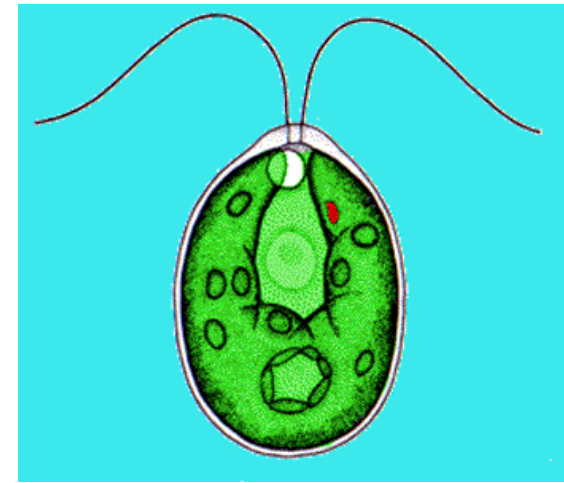
Neustupa et al. 2009, *Phycol. Res.* 57: 159-169.
 Aboal & Werner 2011, *Eur. J. Phycol.* 46: 181-192.
 Neustupa et al. 2013, *Phycologia* 52: 411-421.

Třída: ***Chlorophyceae***

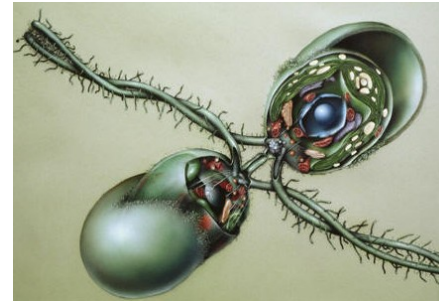
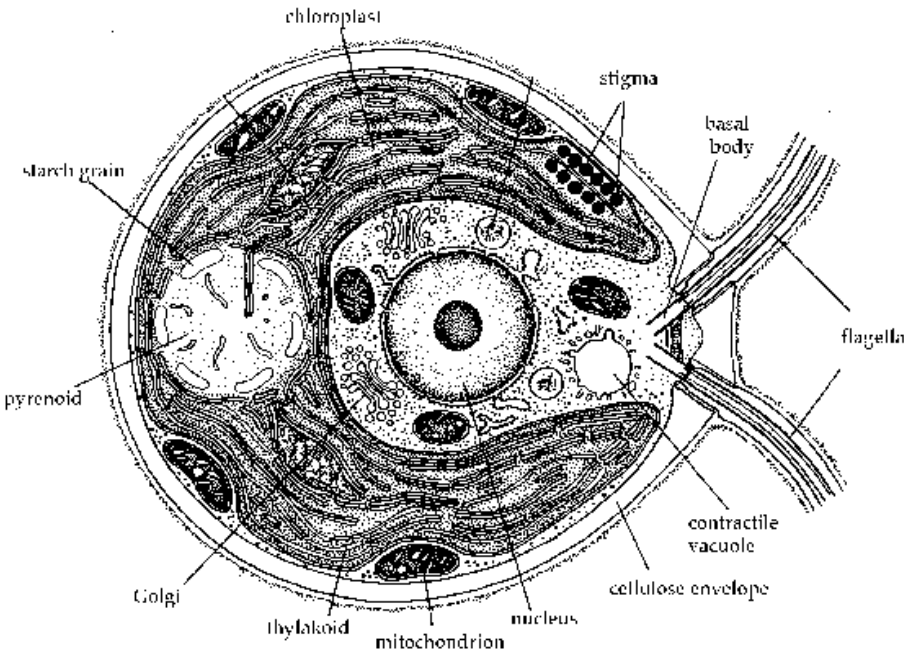
here belong most free-living „plant“ flagellates

coenobial or filamentous multicellularity

biotechnological importance – astaxanthin, beta-carotene production



Chlamydomonas - model flagellated organism in Viridiplantae
(*Volvox* - plant coenobial multicellularity based on flagellated cells)

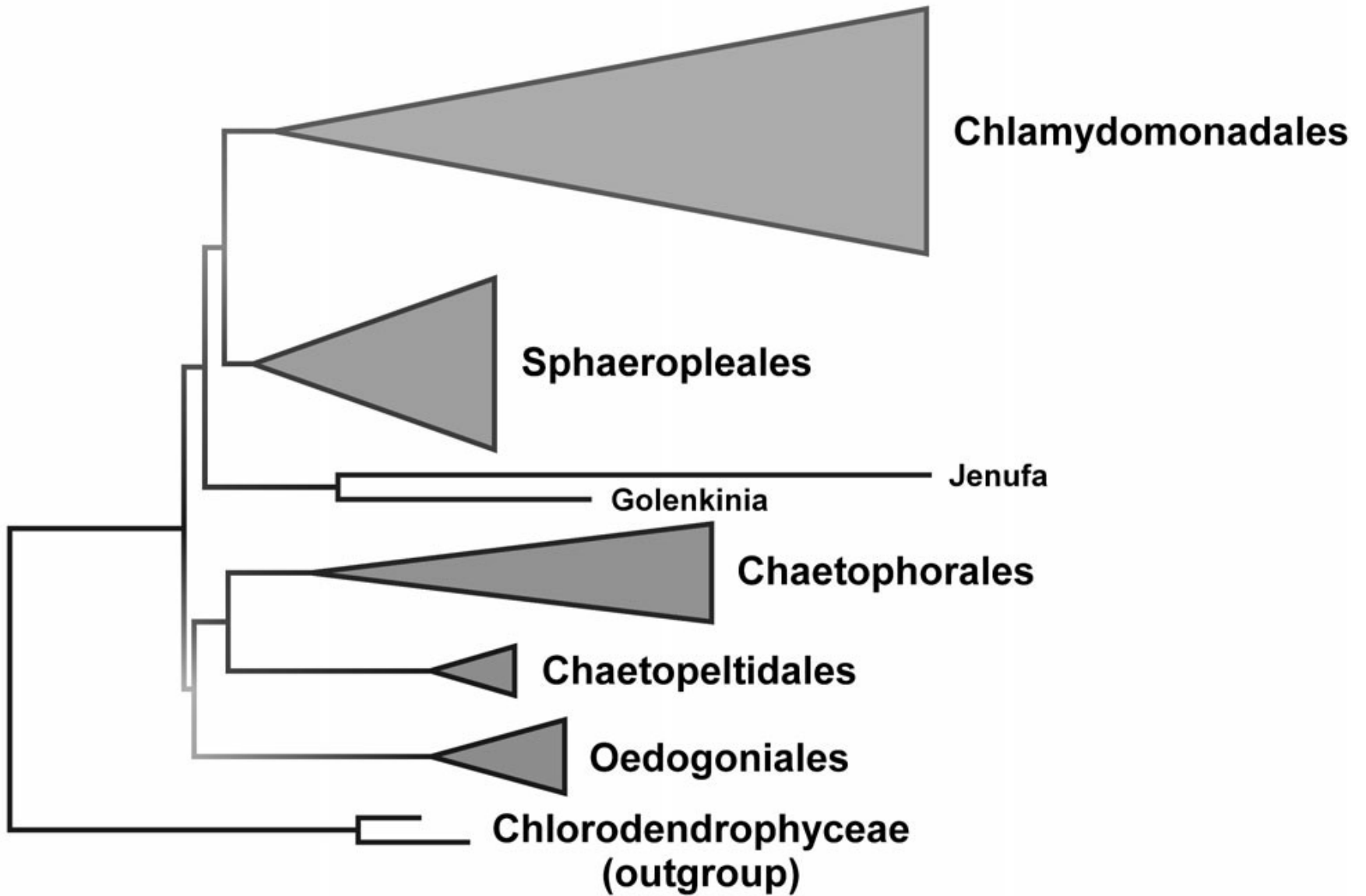


Chlamydomonas - izogamie
(http://bioweb.uwlax.edu/bio203/s2007/awowale_john/reproduction.htm)



Dunaliella, *Carteria*

Chlamydomonas



class: Chlorophyceae

order: **Chaetopeltidale**



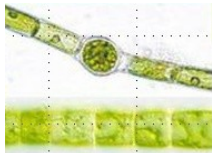
order: **Chaetophorales**

(*Chaetophora*, *Stigeoclonium*, *Dicranochaete*)



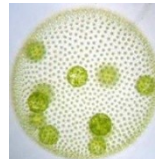
order: **Oedogoniales**

(*Oedogonium*, *Bulbochaete*, *Oedocladium*)



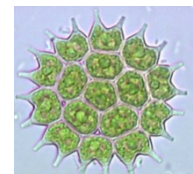
order: **Chlamydomonadales**

(*Carteria*, *Chlamydomonas*, *Dunaliella*, *Haematococcus*, *Volvox*, *Gonium*)

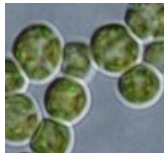
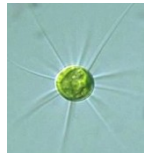


orer: **Sphaeropleales**

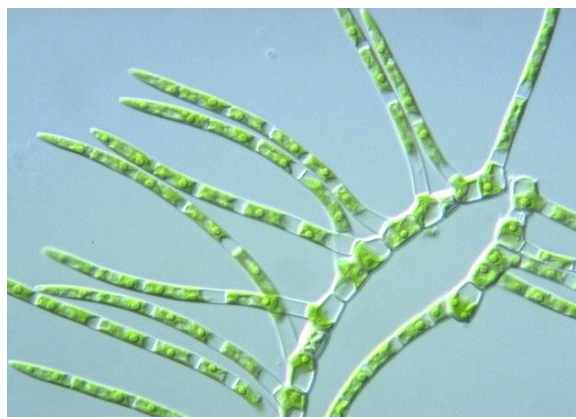
(*Scenedesmus*, *Desmodesmus*, *Pediastrum*, *Hydrodictyon*)



incertae sedis: *Golenkinia*, *Jenufa*



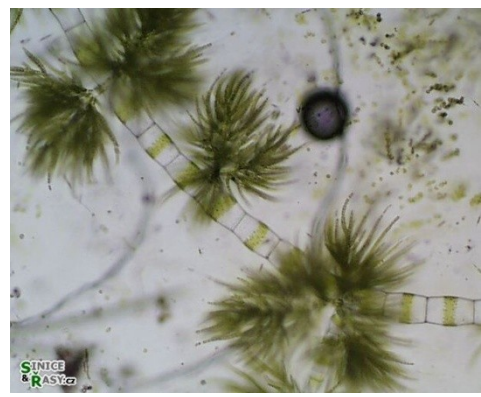
order: Chaetophorales



Stigeoclonium



Chaetophora

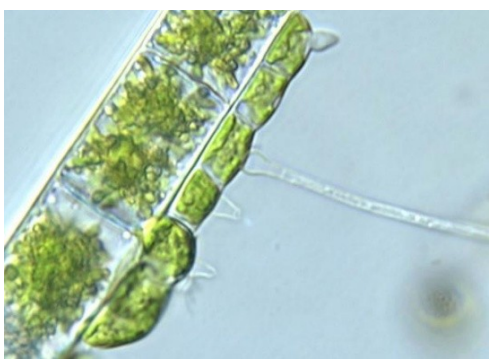


Draparnaldia

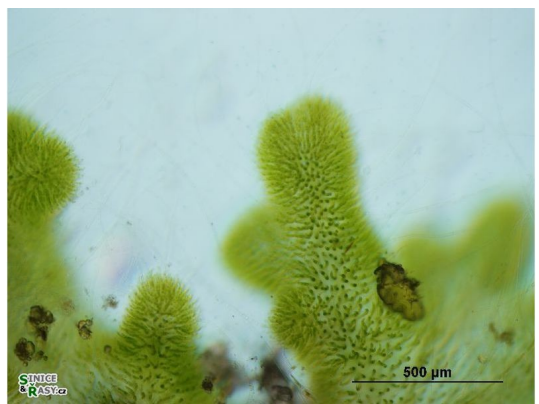


BioObs © Gilbert Billard

predominantly a freshwater lineage

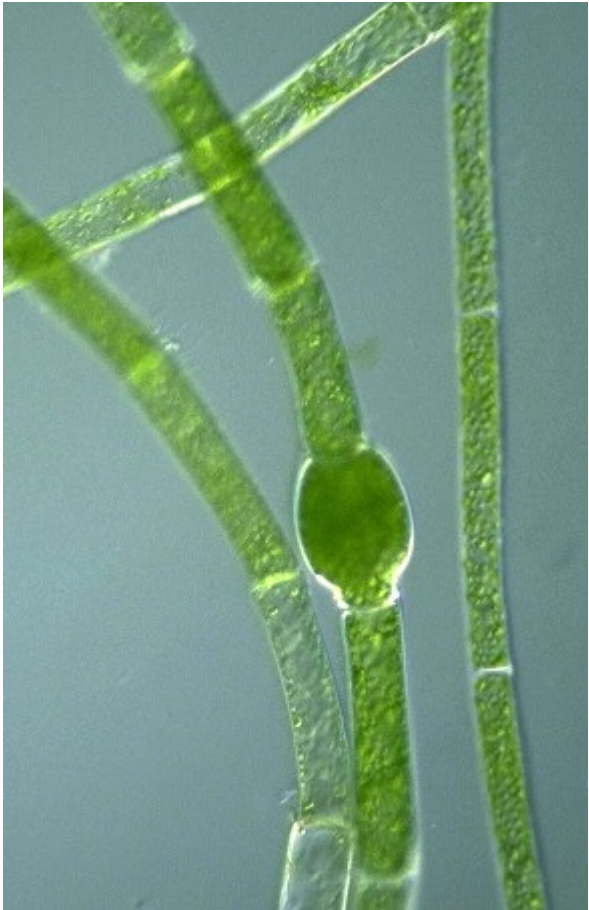


Aphanochaete

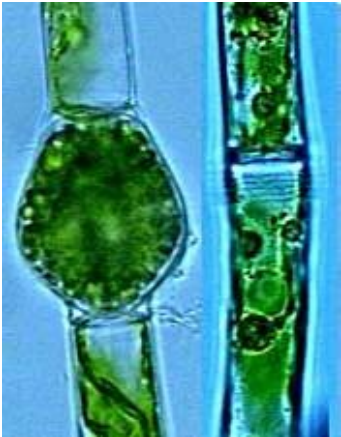


500 µm

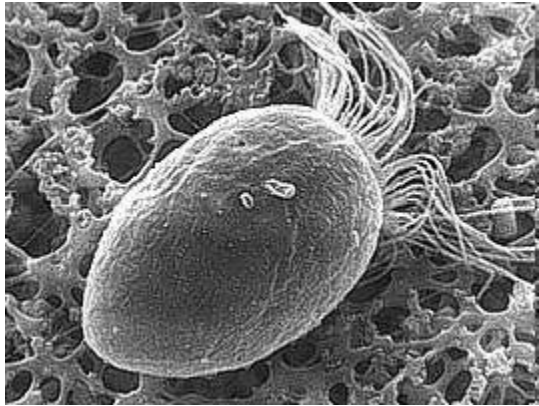
order: Oedogoniales



Bulbochaete



Oedogonium



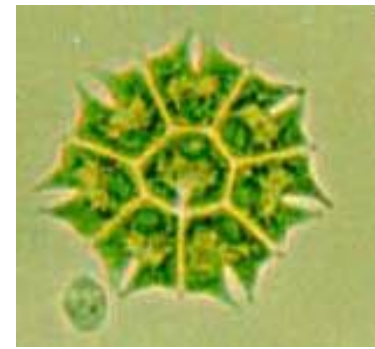
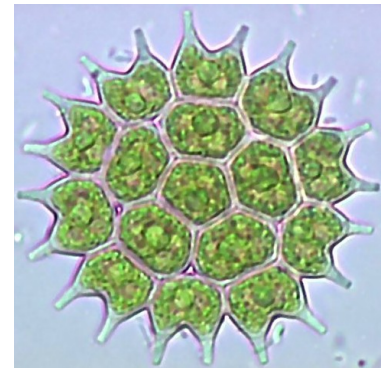
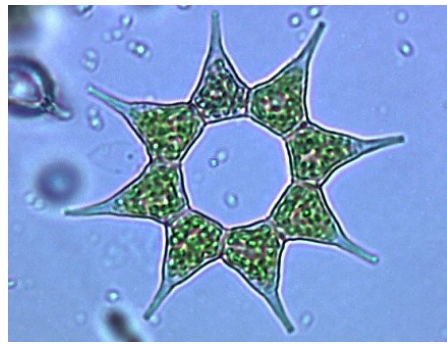
order: Sphaeropleales

Pediastrum and relatives

Scenedesmus and relatives

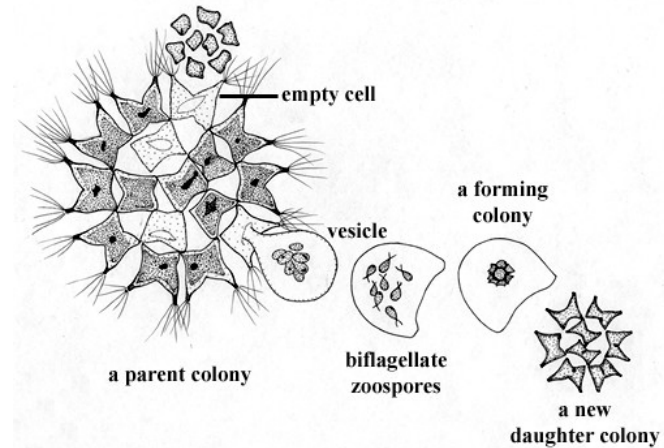
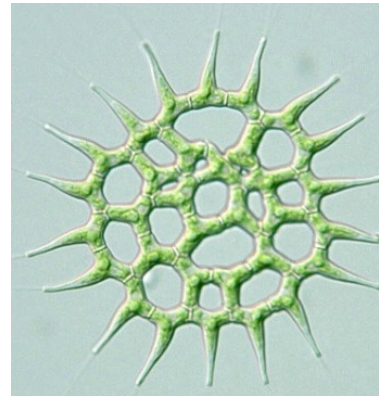
[incl. *Desmodesmus*, *Acutodesmus*,
Pectinodesmus, *Verrucodesmus*,
Chodatodesmus, etc...]

coccoid coenobial multicellularity

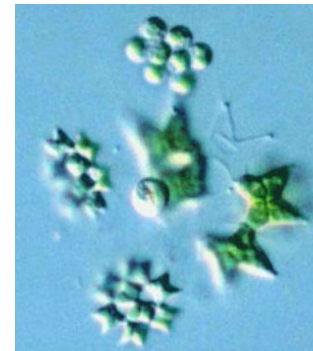


SCENEDESMUS

Pediastrum

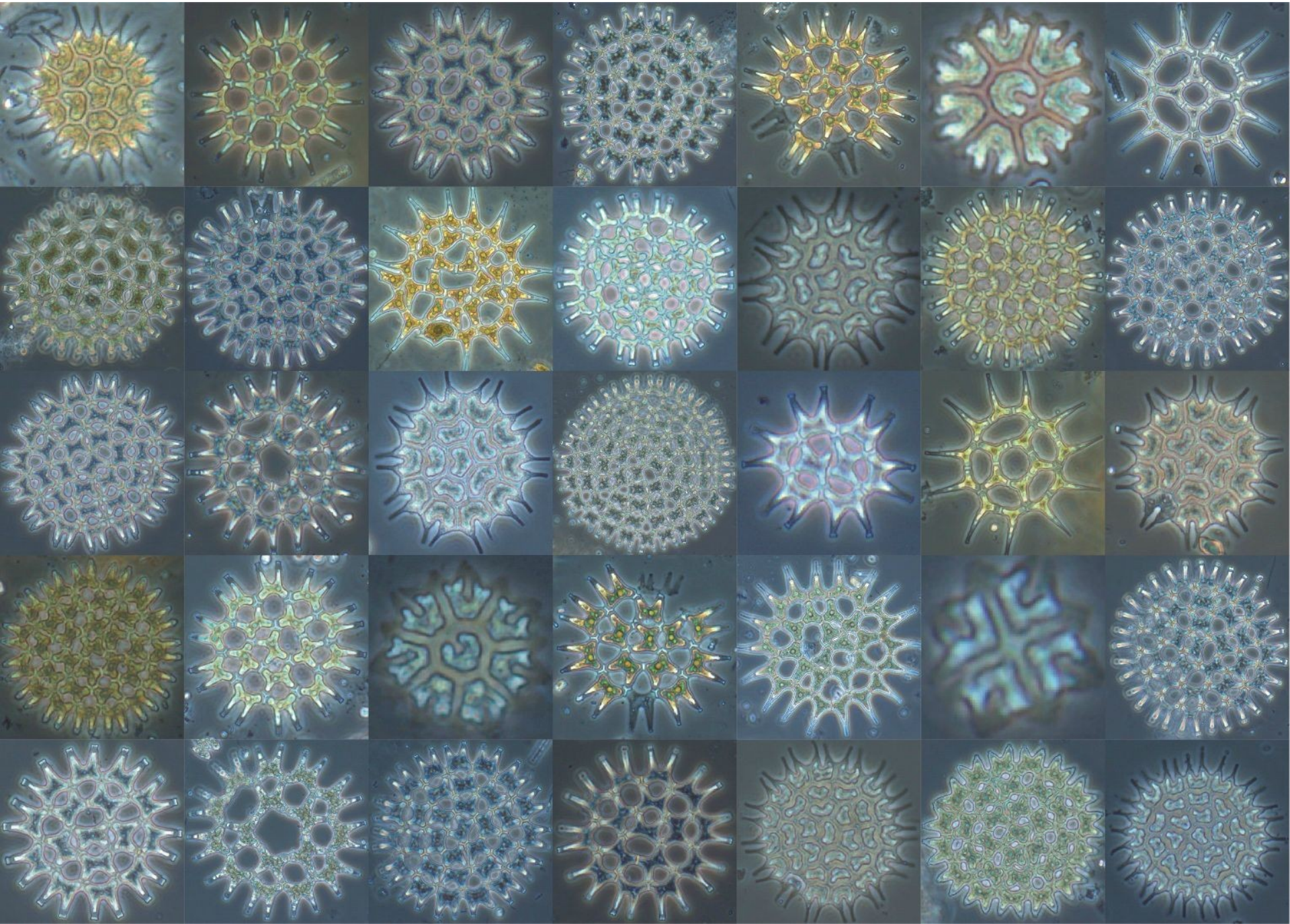


Scenedesmus, *Desmodesmus*



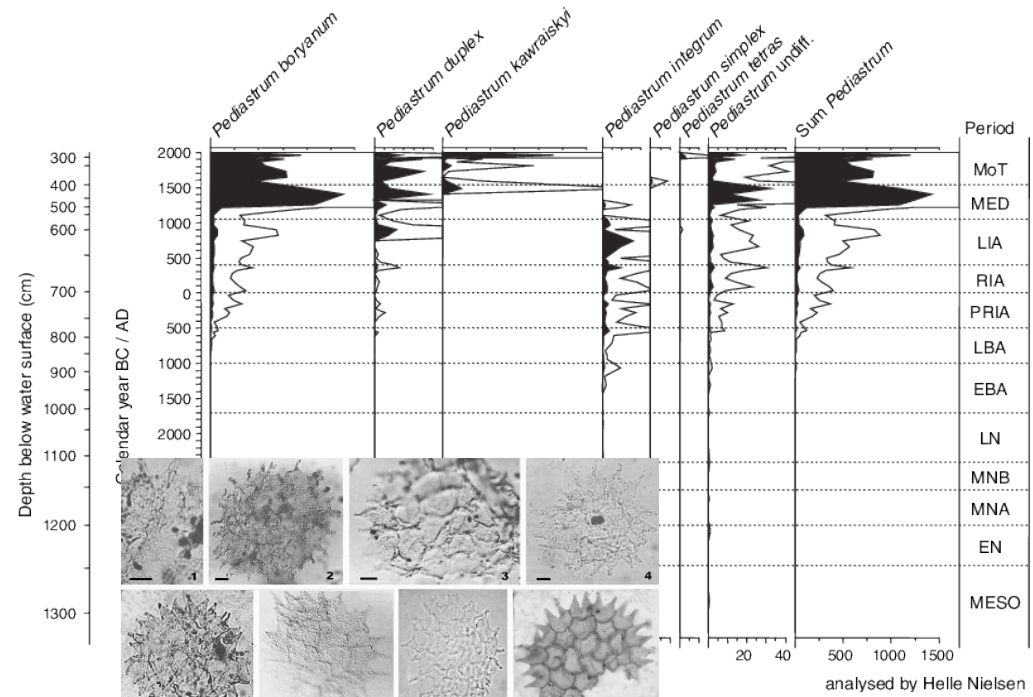
morphological plasticity (reaction norm of genotypes) – adaptation to herbivory and sinking stress

Pediastrum, Parapediastrum, Pseudopediastrum, Sorastrum, Monactinus, Lacunastrum, Stauridium, etc...



Bradshaw et al, 2005, *The Holocene*

subrecent eutrofication of a temperate lake



analysed by Helle Nielsen

Zamaloa & Tell, 2005, *Hydrobiologia*

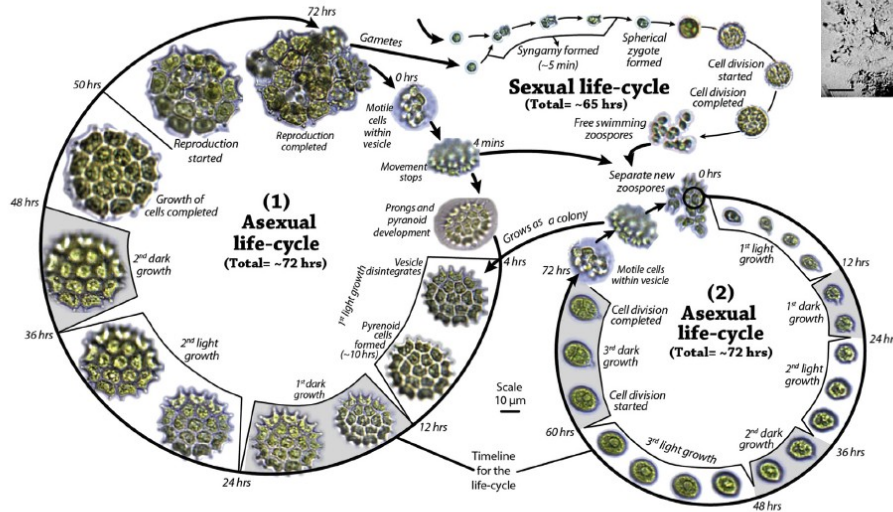
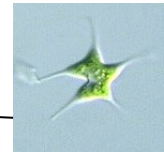
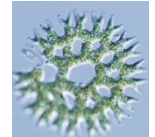
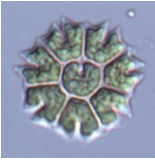
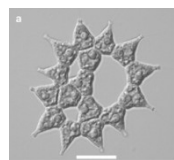
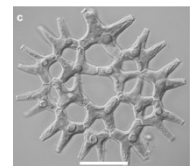
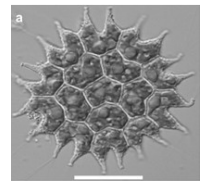
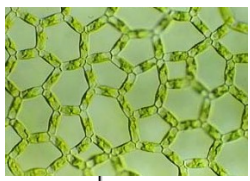
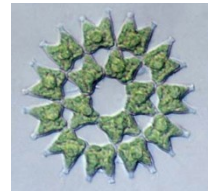
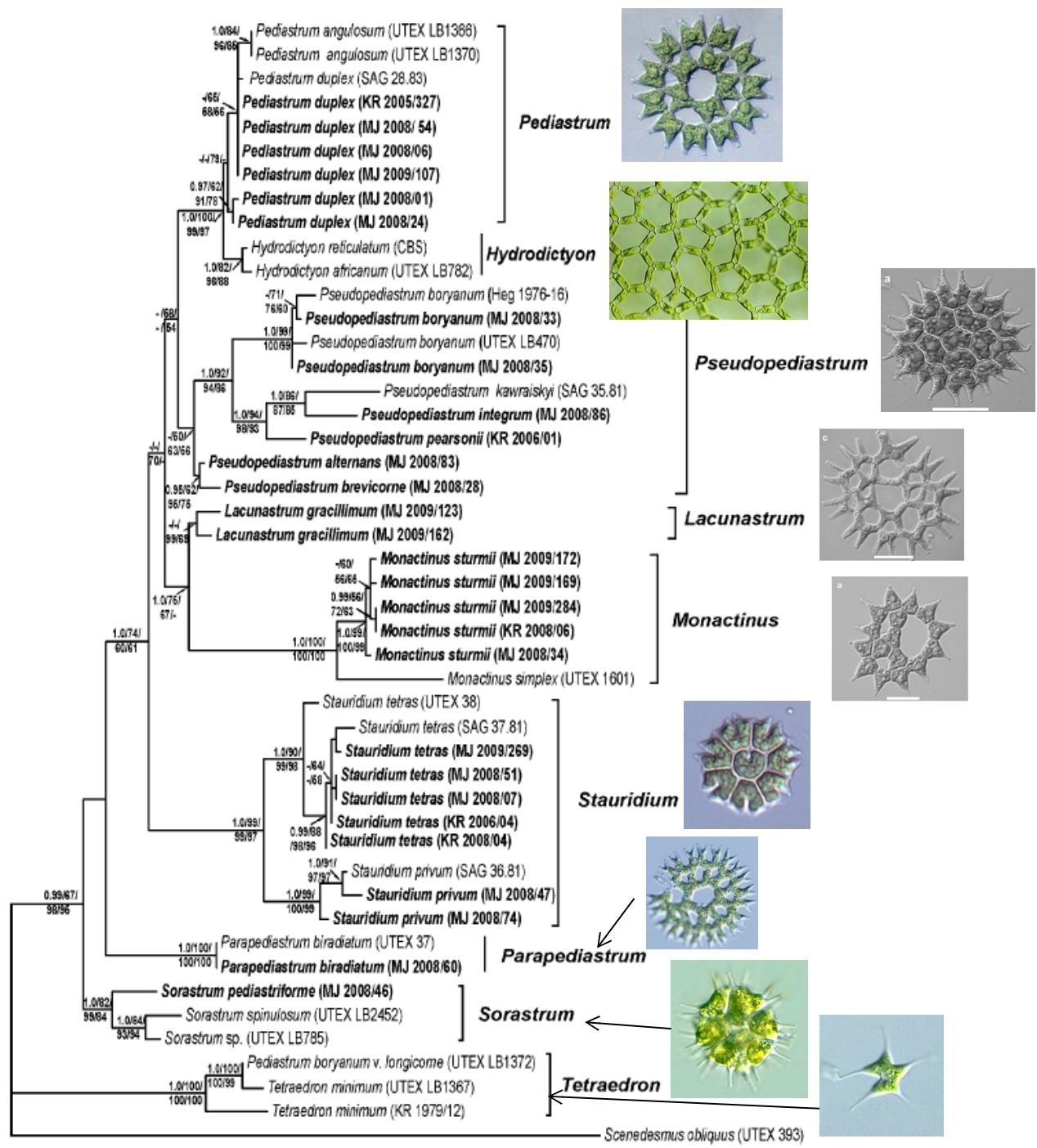
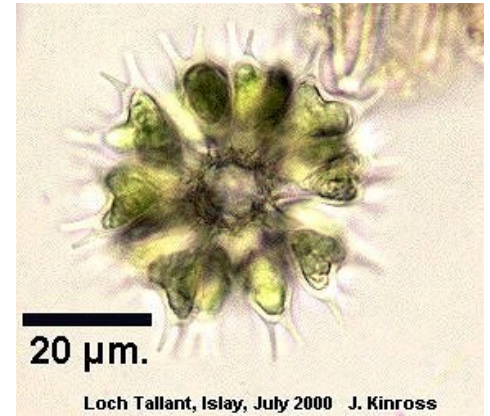
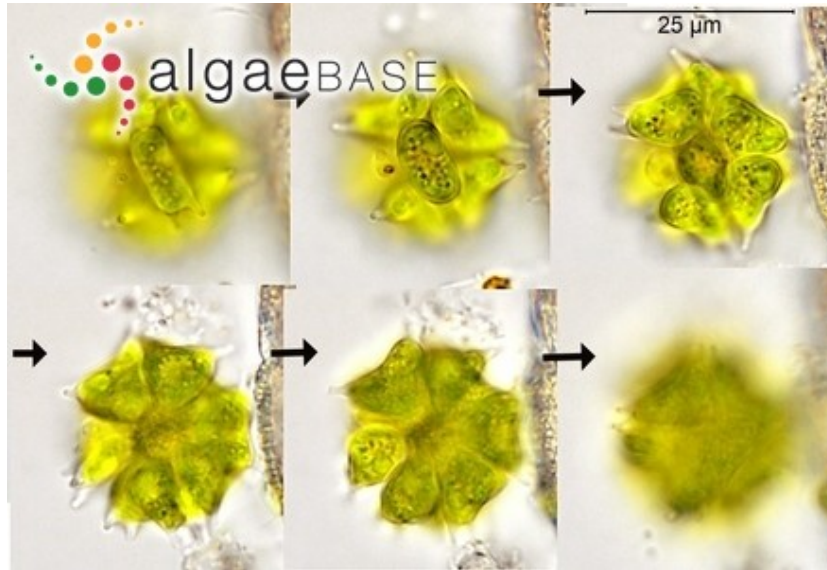
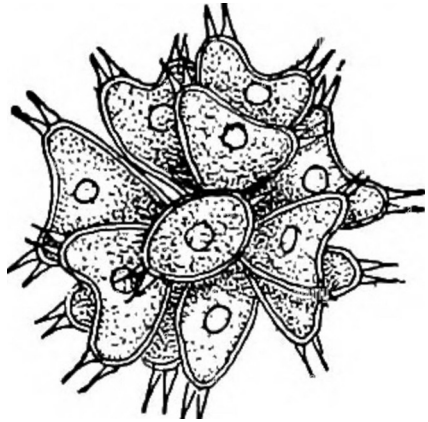


Fig. 3 – The asexual and sexual life-cycles of *Pediatrum boryanum* determined by observation of the growth of single cells/ colonies grown in a microcosm under 250 $\mu\text{Mol/m}^2\text{s}$ (12:12 h light and dark cycle) at 20 °C.

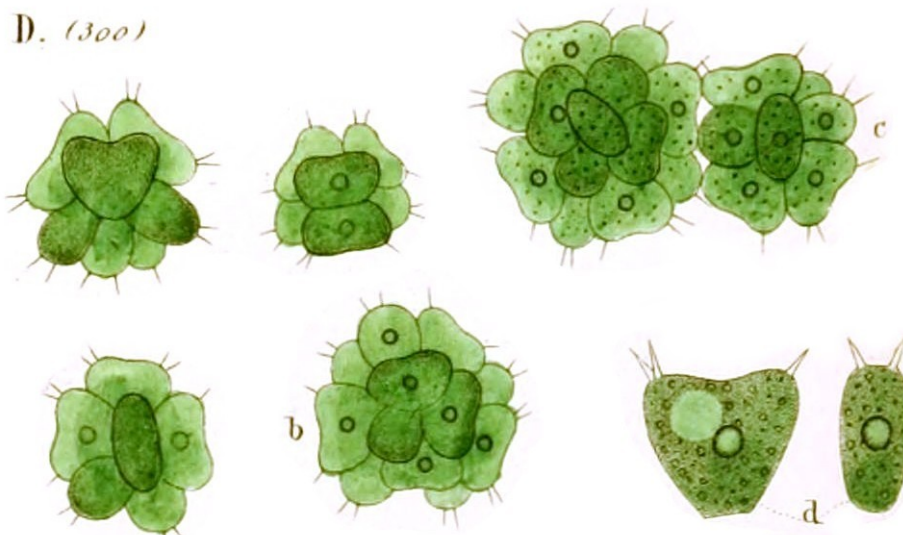
Park et al, 2014, *Water Research* 60



Sorastrum

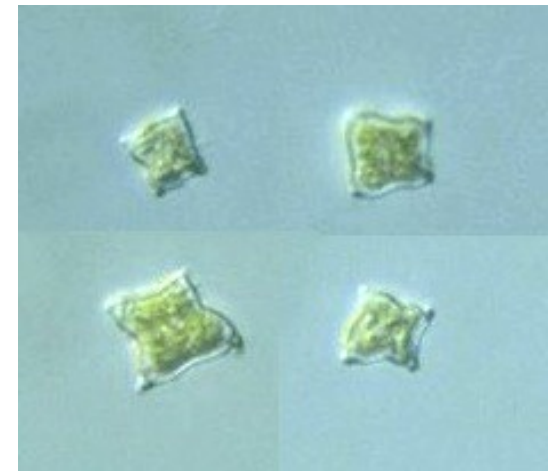


D. (300)



Sorastrum spinulosum (600)

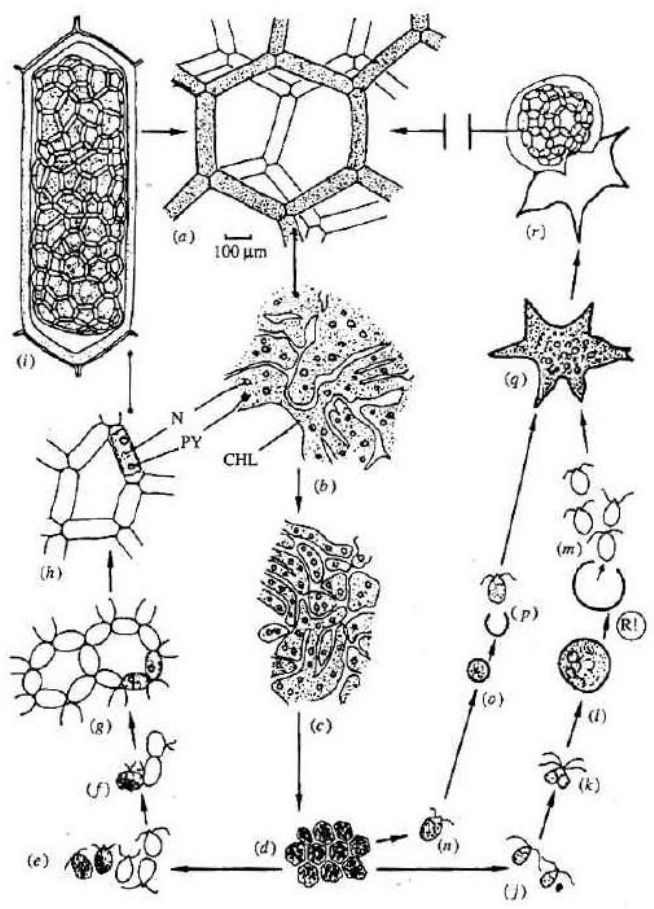
Tetraedron



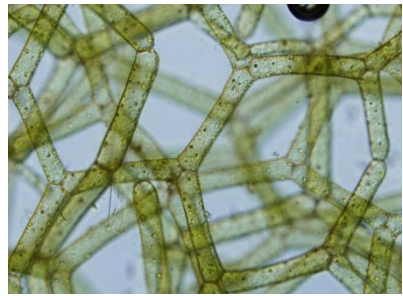
mostly (sub-)tropical eutrophic phytoplankton

unicellular lineage,
sister to coenobial taxa

Hydrodictyon



haplontic life cycle, meiospores from a zygote, develop into the single-celled stage which produces coenobial plant



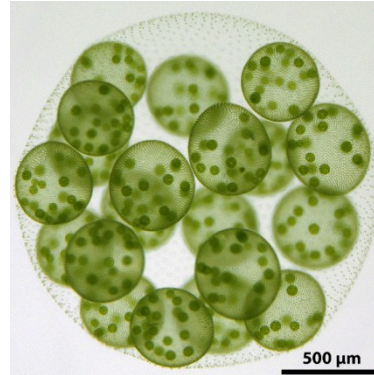
[Domin pond, Č.B.]

Chlamydomonadales

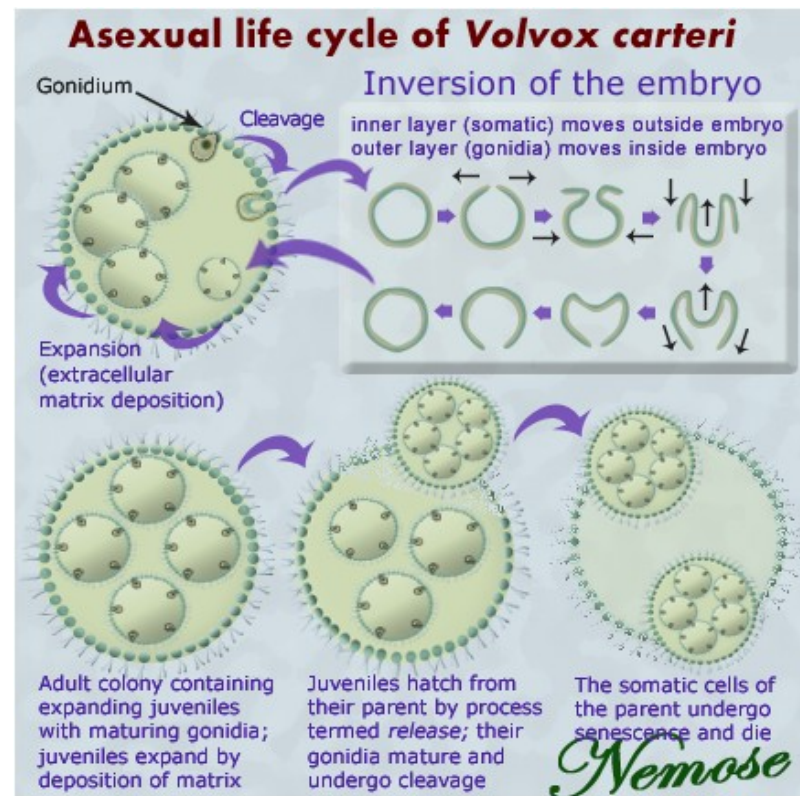
coenobial multicellularity of flagellates
(*Volvox*, *Gonium*, *Eudorina*)



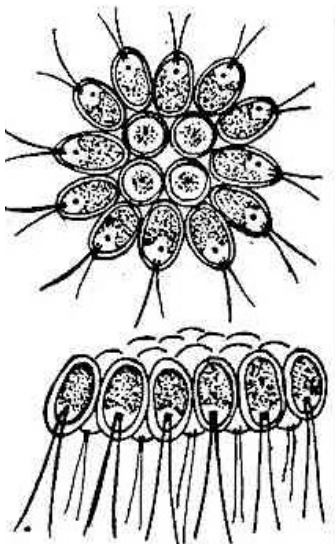
Pleodorina



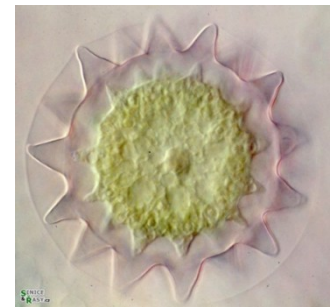
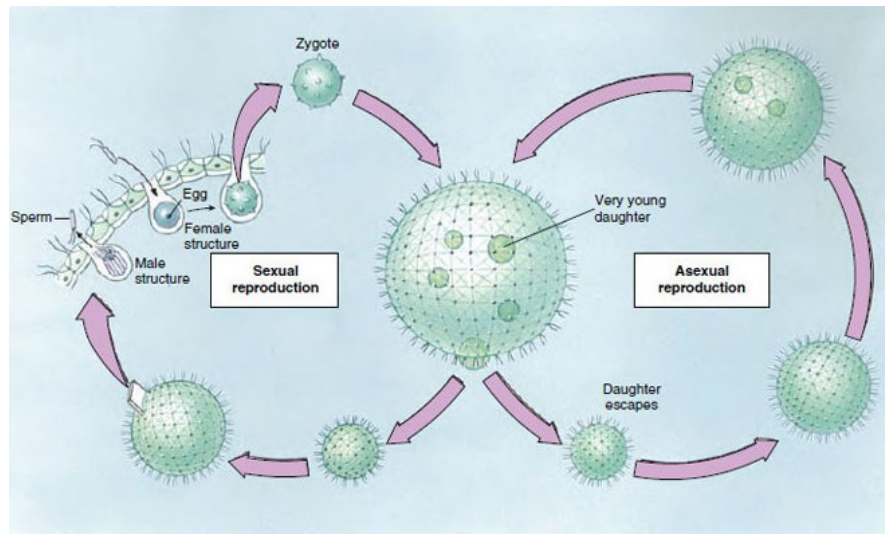
Volvox



<http://www.metamicrobe.com/volvox/>

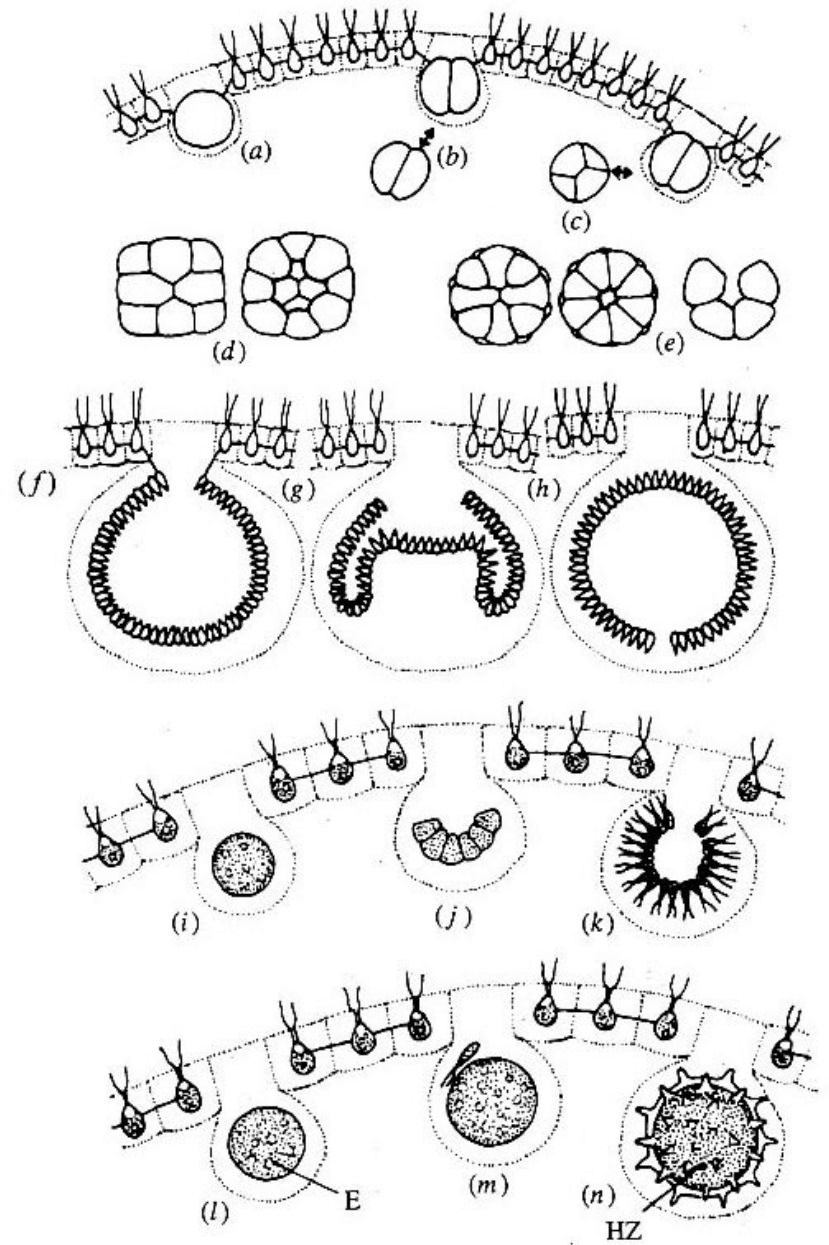
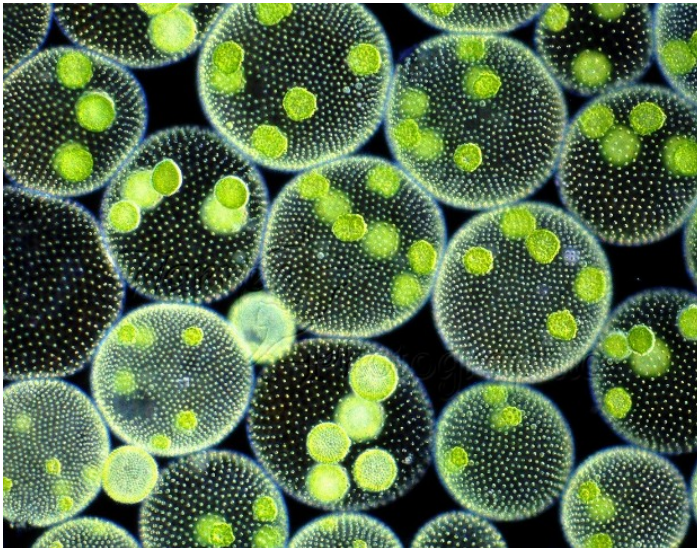
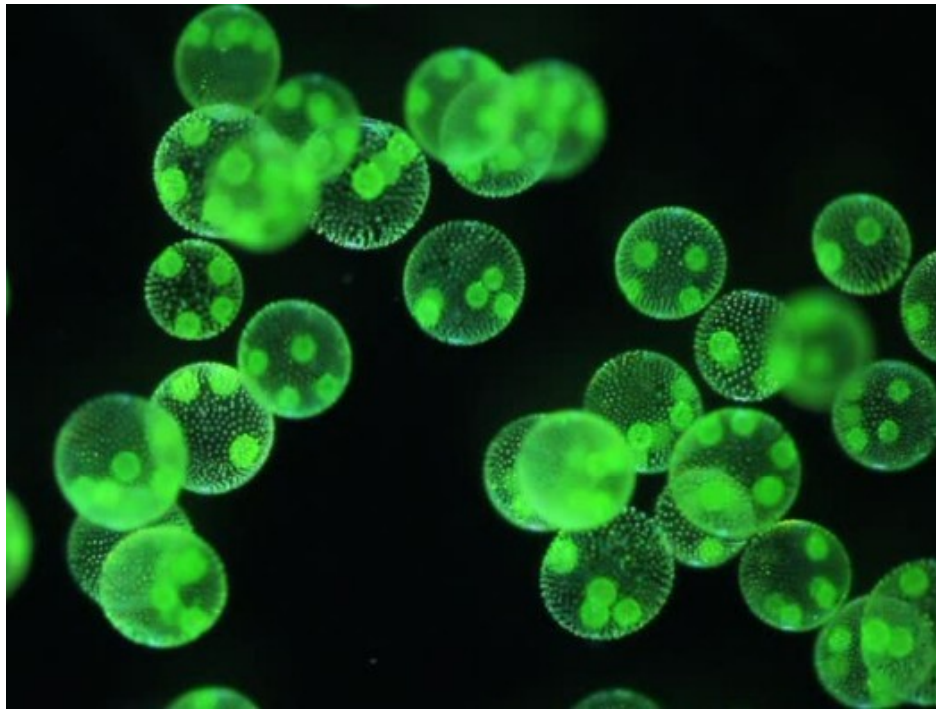


Gonium



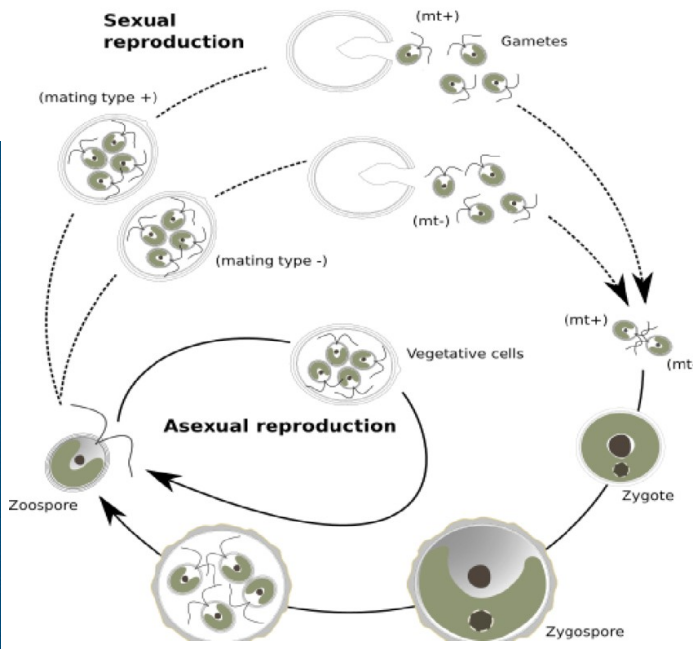
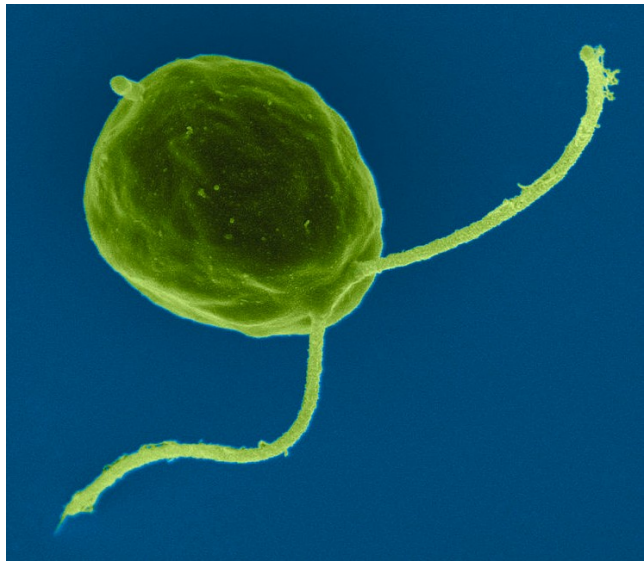
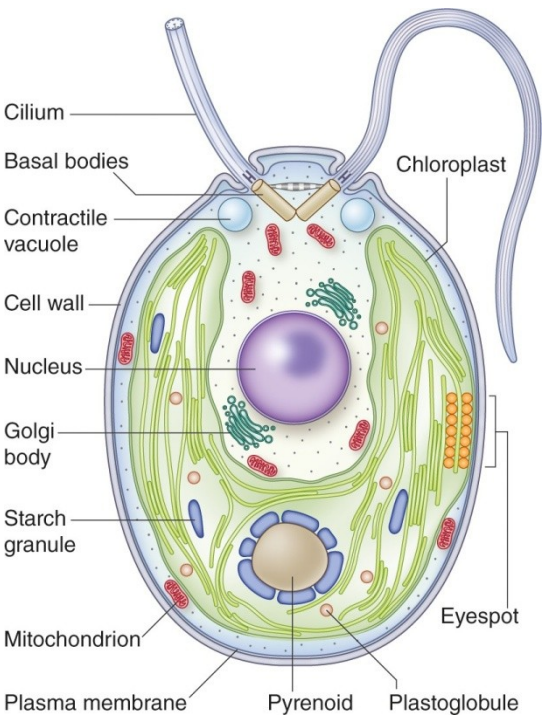
Volvox - zygospore

Volvox



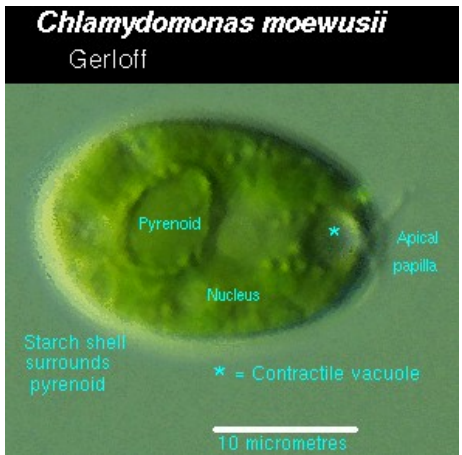
asexual: daughter colonies, sexual: oogamy

Chlamydomonas

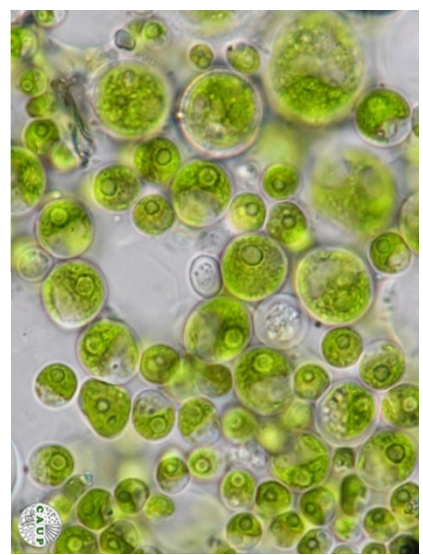


Baudelet et al., 2017, Algal Res.

the model free-living
flagellated plant organism
(*C. reinhardtii*)

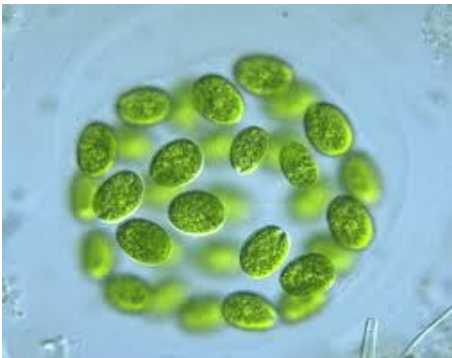
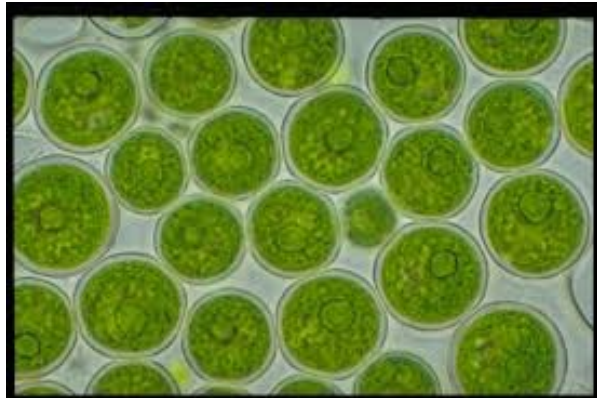
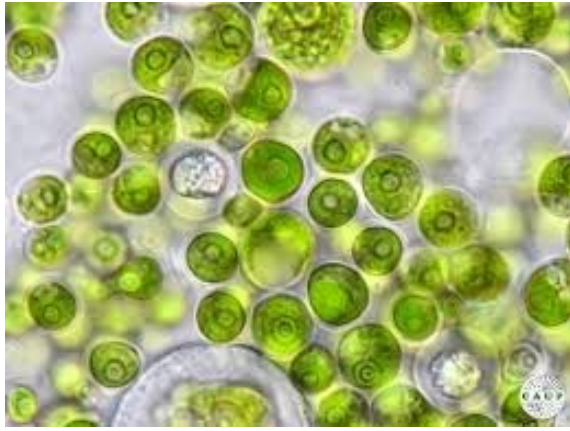


Chlorococcum

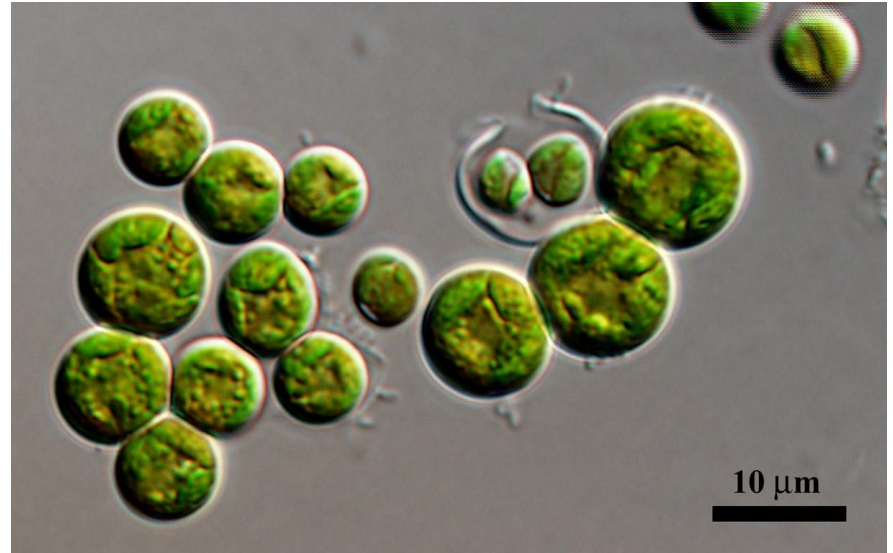


<http://megasun.bch.umontreal.ca/protists/chlamy/summary.html>

Chlorococcum



Bracteacoccus



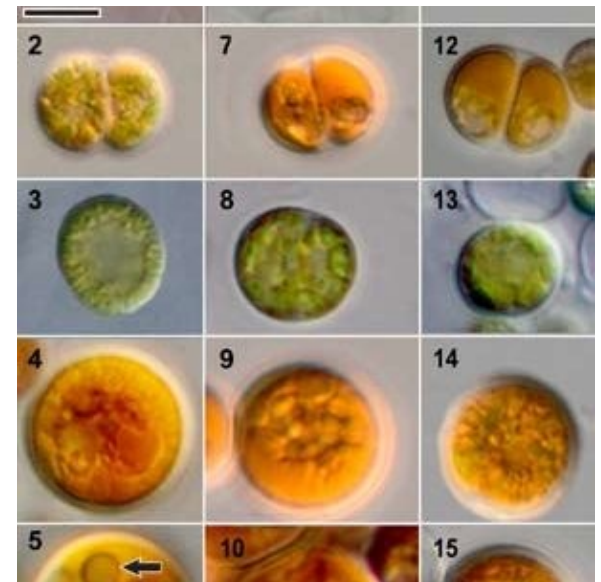
often occur in soil

simplified globular morphology

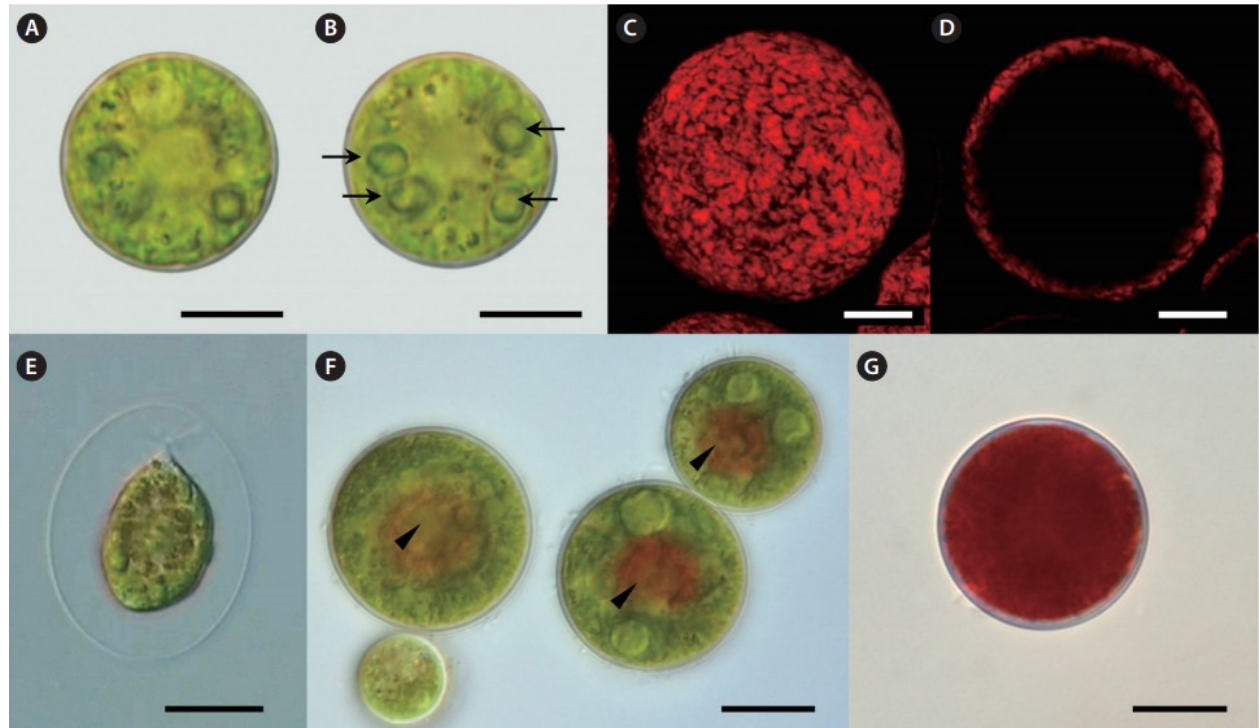
Chlamydomonas-like zoospores

lipidic storage compounds

high amounts of carotenoids

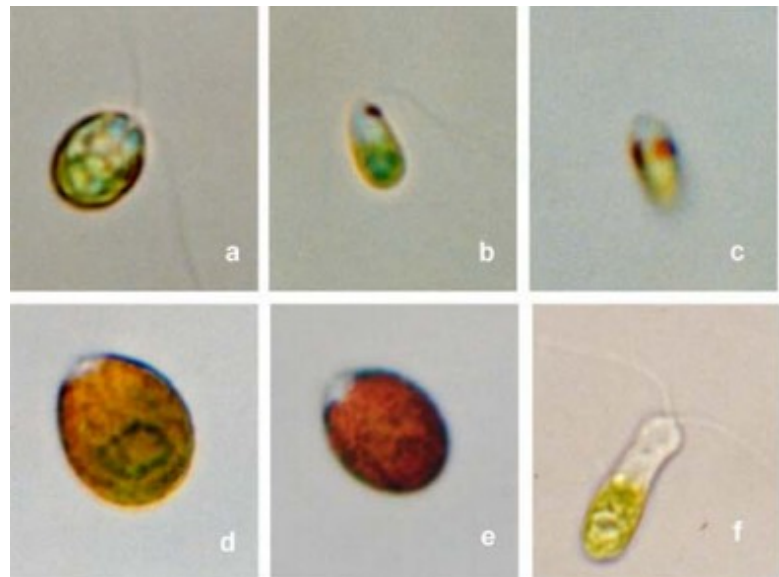


Haematococcus



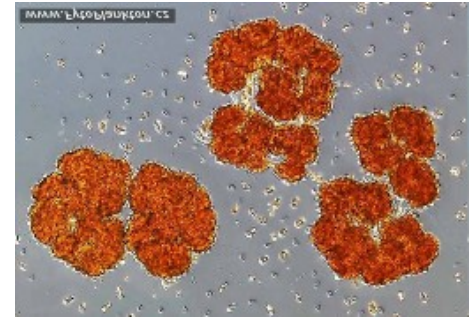
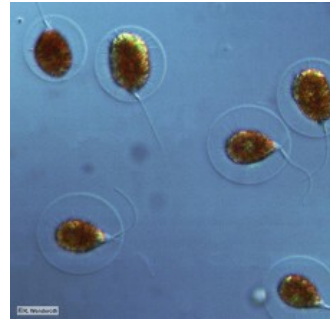
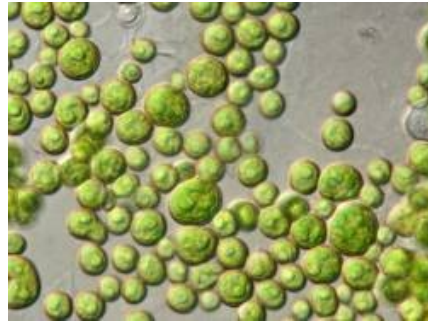
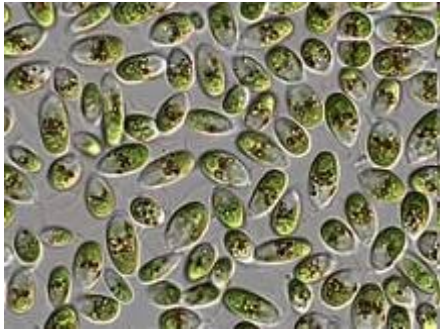
Klochko et al., 2013, Algae

Dunaliella

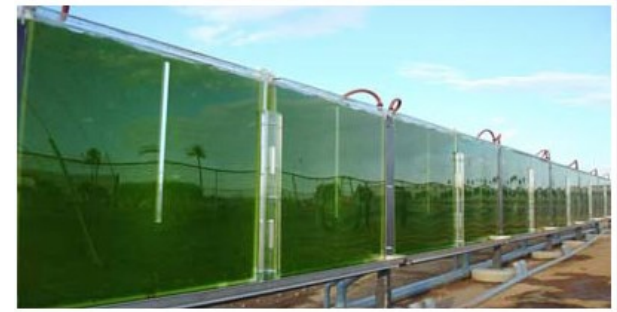
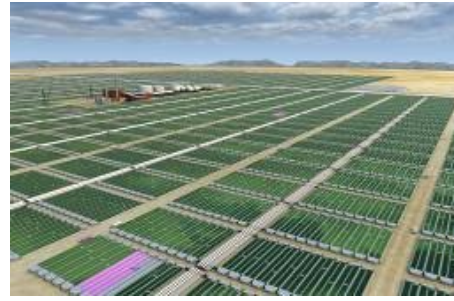
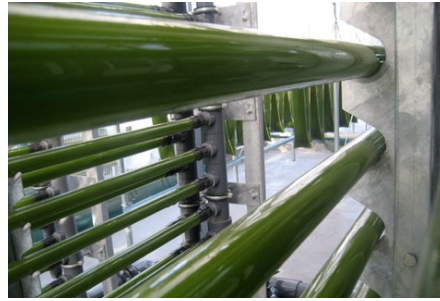


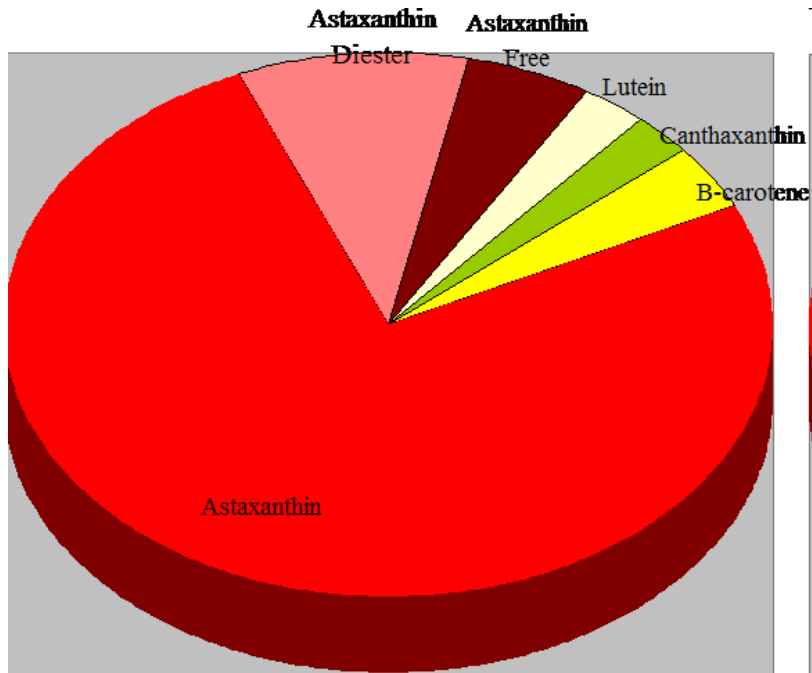
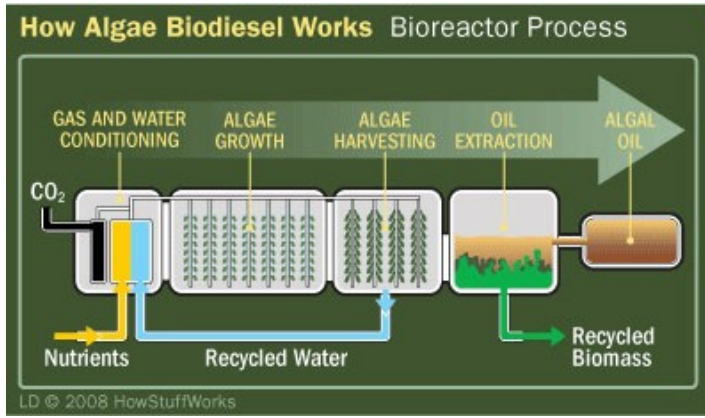
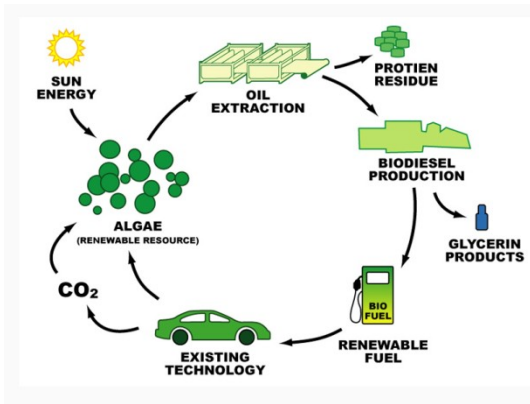
Borowitzka & Siva, 2007, J. Appl. Phycol.

photobioreactores and green algae



*Chlorophyceae (Muriellopsis, Dunaliella, Haematococcus),
Eustigmatophyceae (Nannochloropsis), Trebouxiophyceae (Botryococcus)*





■ B-carotene	■ Astaxanthin	■ Astaxanthin Diester
■ Astaxanthin Free	■ Lutein	■ Canthaxanthin

<http://www.igb.fraunhofer.de/en/competences/environmental-biotechnology/microalgae/astaxanthin.html>

<http://www.algae-biotech.com/>

Haematococcus – composition of carotenoids

Algenhaus, Hamburg