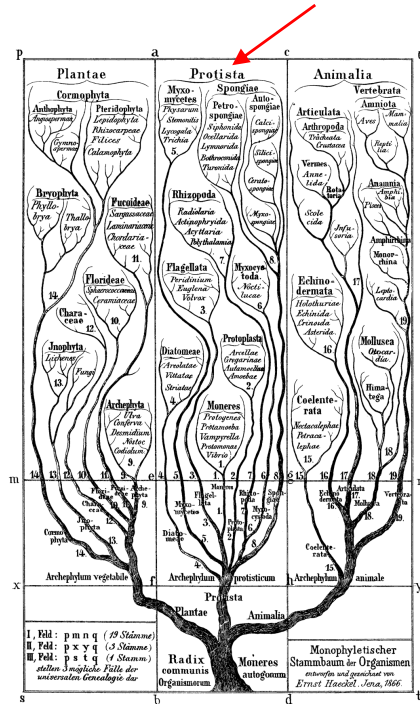


„secondary“ endosymbiosis
algae with rhodophyte-derived plastids

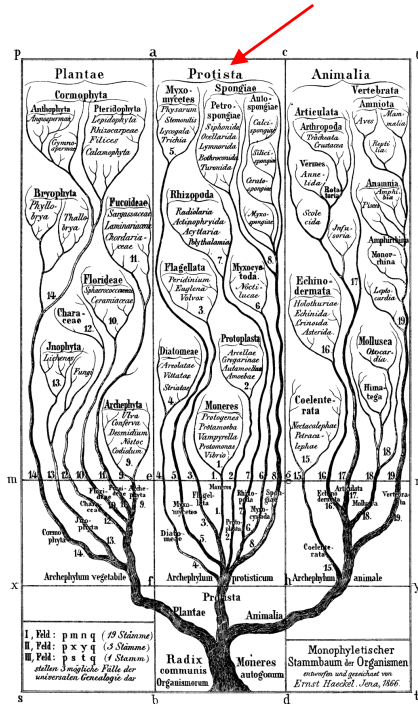
Eukaryotic tree of life



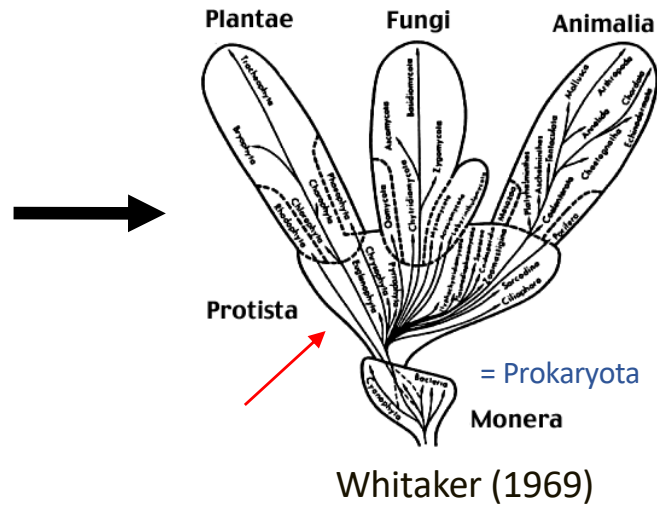
Haeckel (1866)

Classification: kingdoms

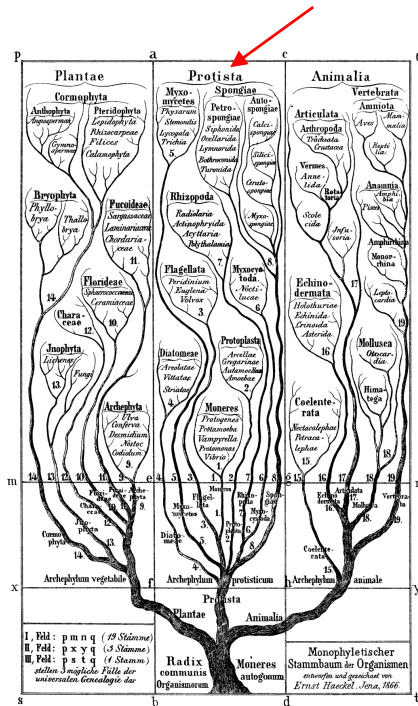
Eukaryotic tree of life



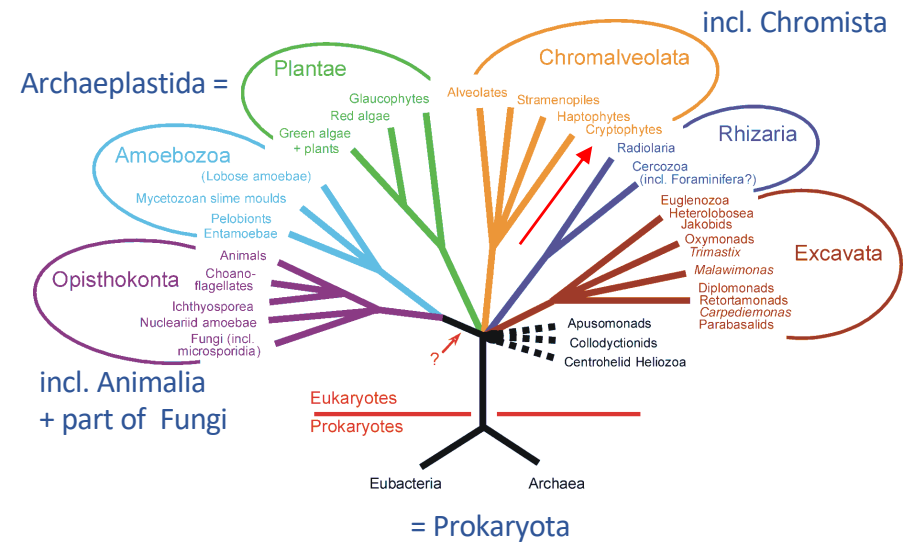
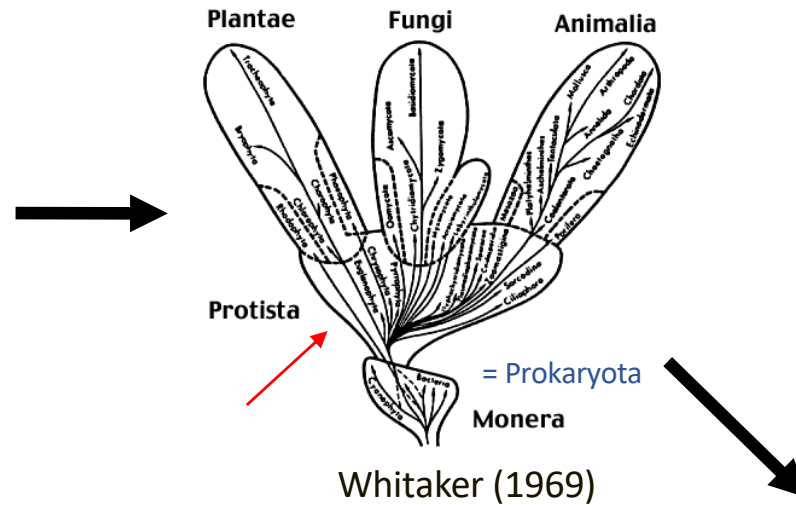
Haeckel (1866)



Eukaryotic tree of life

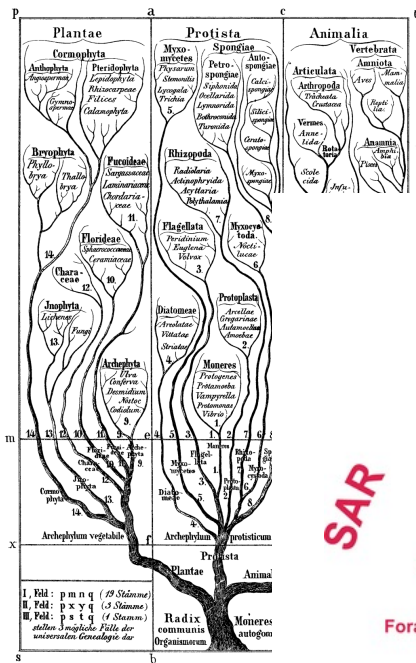


Haeckel (1866)

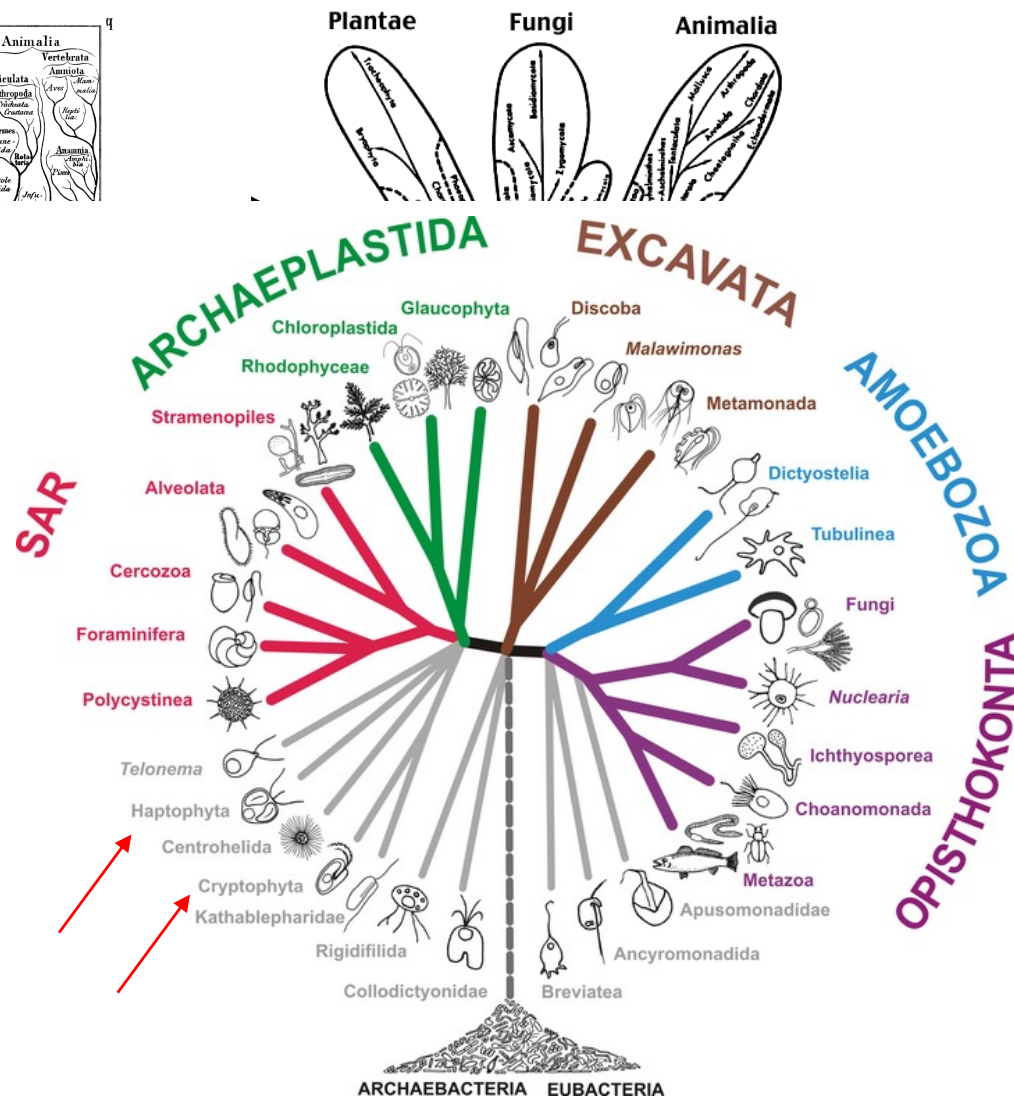


Classification: kingdoms

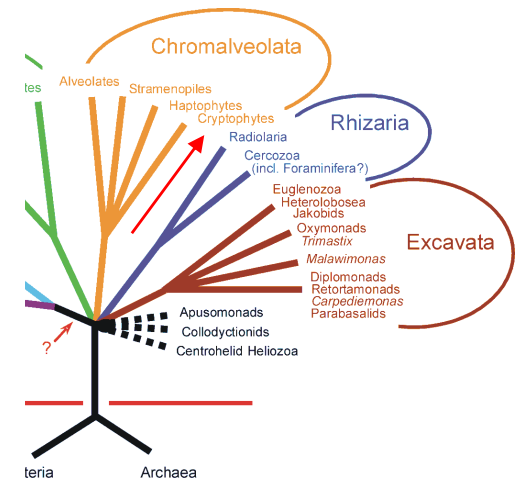
Eukaryotic tree of life



Haeckel (1866)

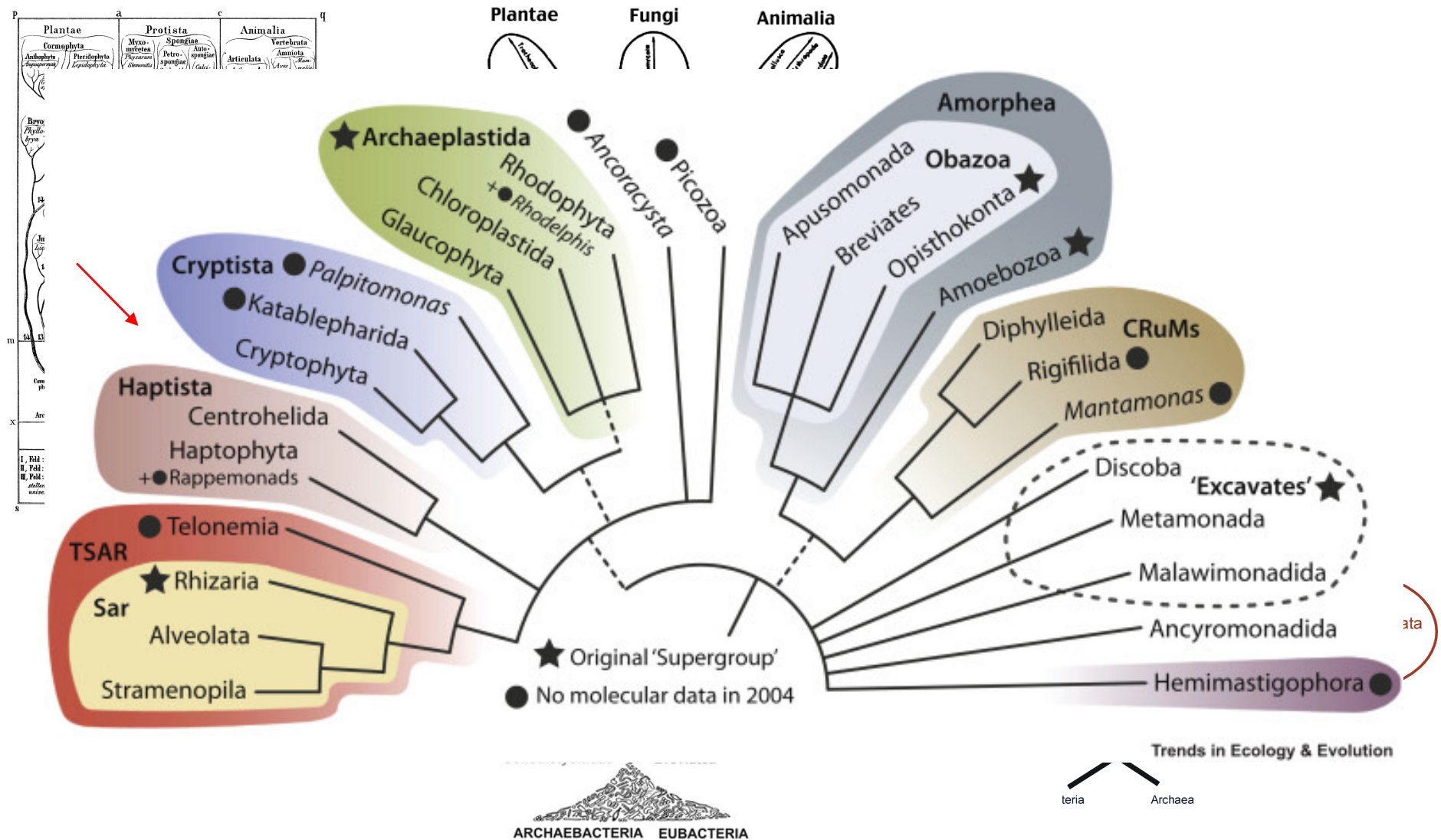


Adl et al. (2012)



Classification: kingdoms

Eukaryotic tree of life

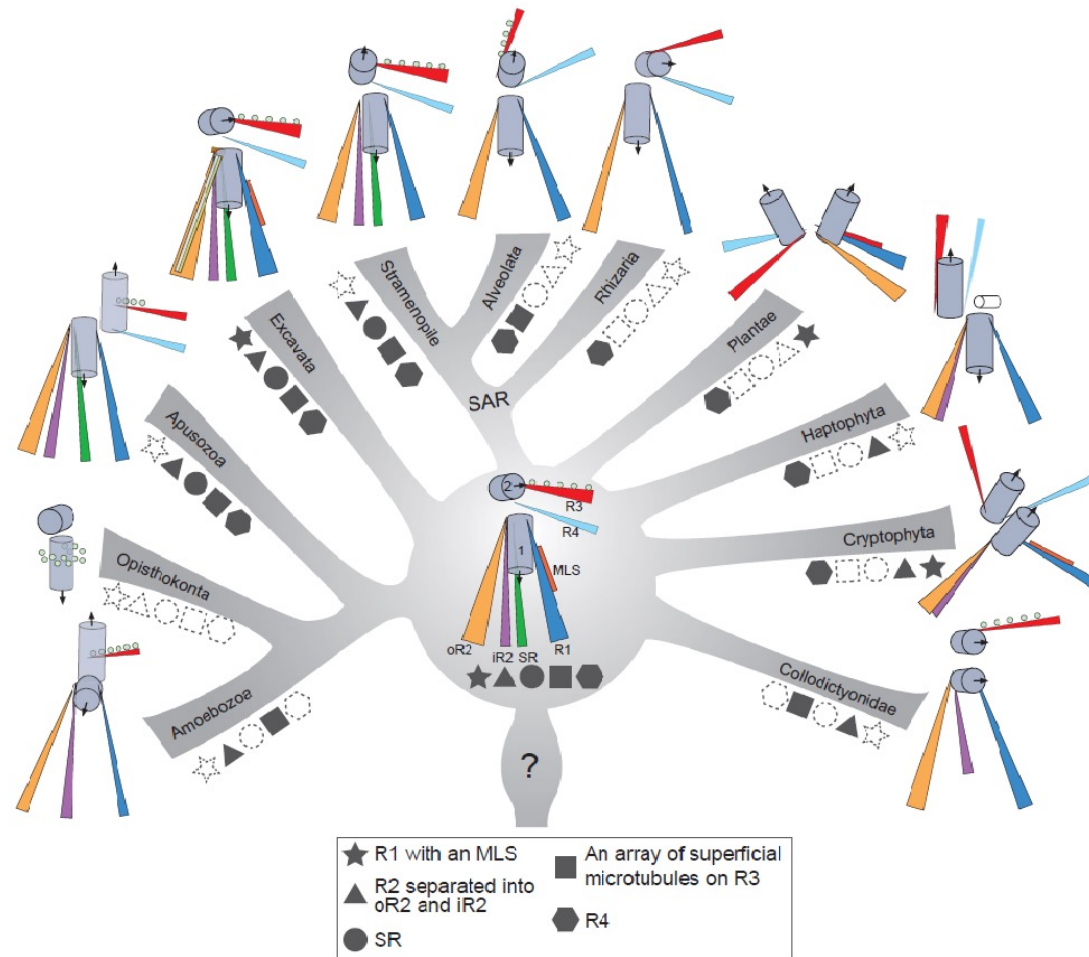


Burki et al. (2020)

Classification: kingdoms

Eukaryotic tree of life

- inferred ancestral traits for the flagellar apparatus
(variations in the structure of cytoskeletal system (MTOCs) reflect phylogenetic relationships)
- LECA (last Eukaryotic common ancestor) = best represented by typical Excavate



MTOC = microtubule organizing center, with few exceptions MTOCs consist of two basal bodies that anchor flagellar axonemes and different configurations of microtubular roots

MLS = multilayered structure

R1 = microtubule root 1

R2 = root involved with phagotrophy

R3 = superficial microtubules

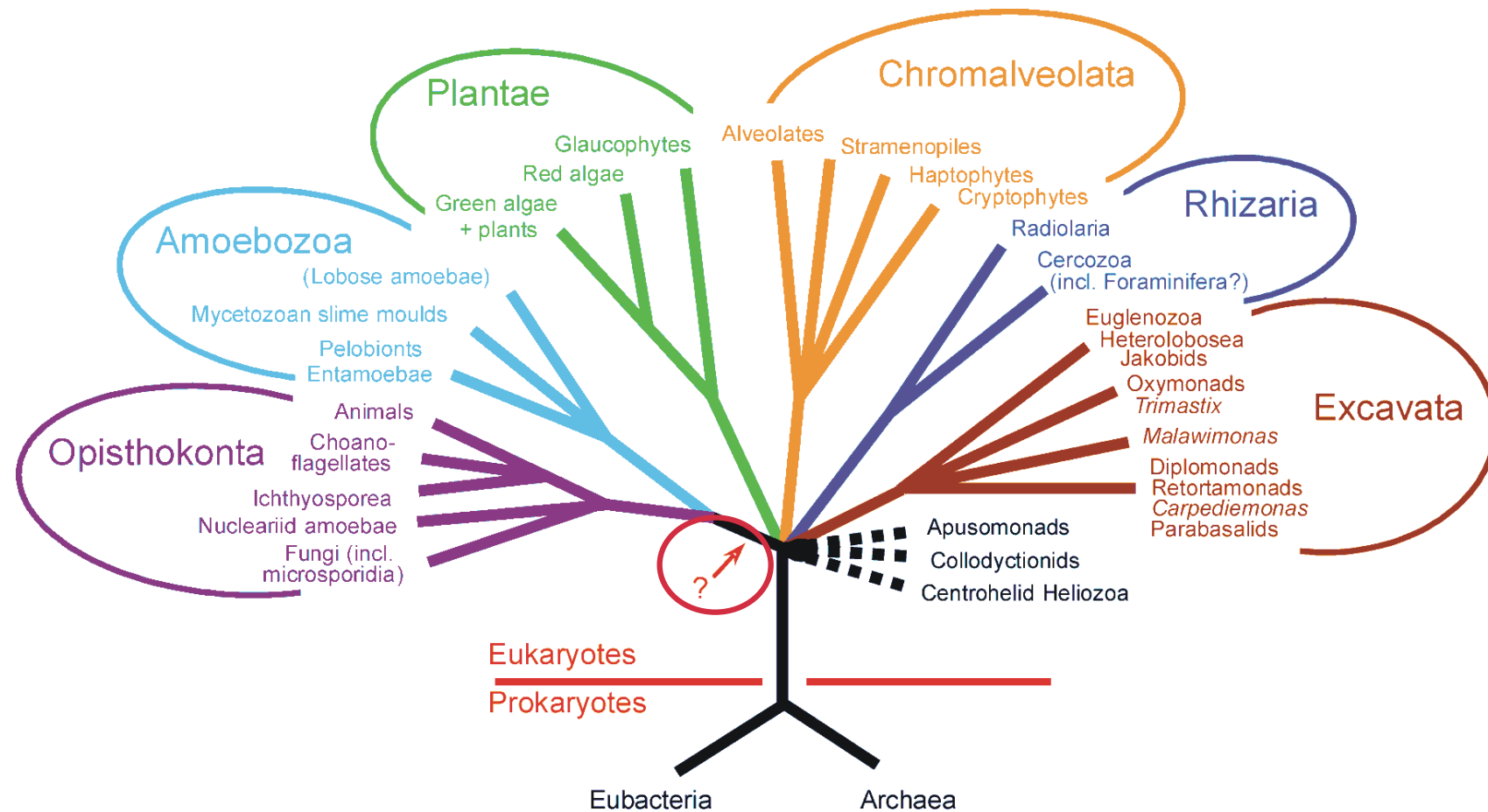
SR = a singlet root (feeding groove)

R4 = microtubule root 4

solid lines = presence of trait

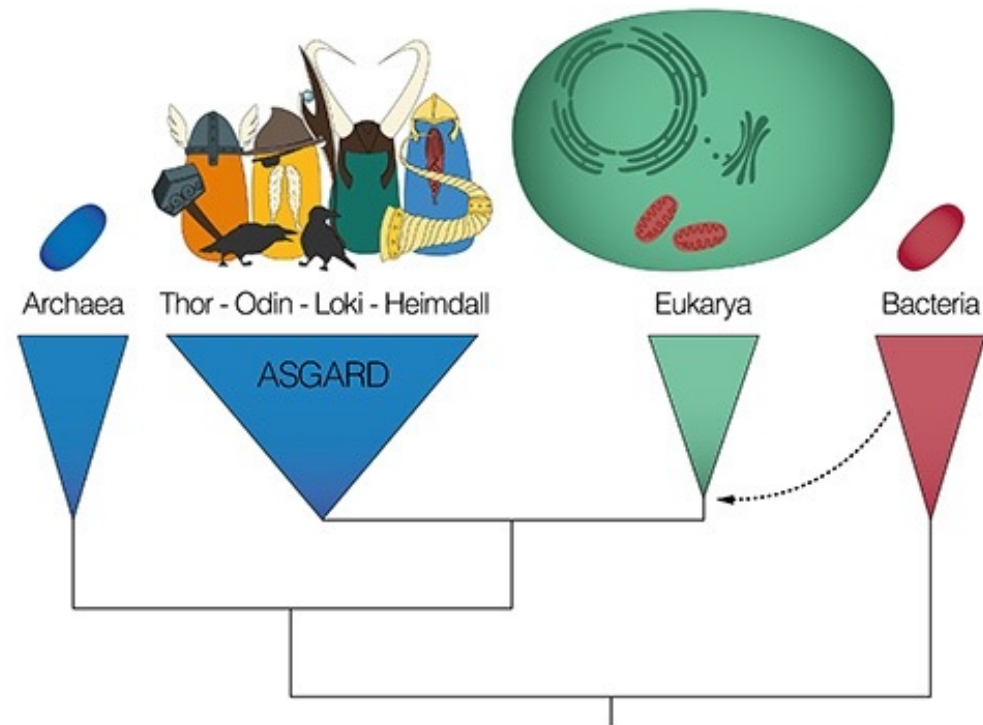
dashed outlines = loss of trait

Eukaryotic tree of life

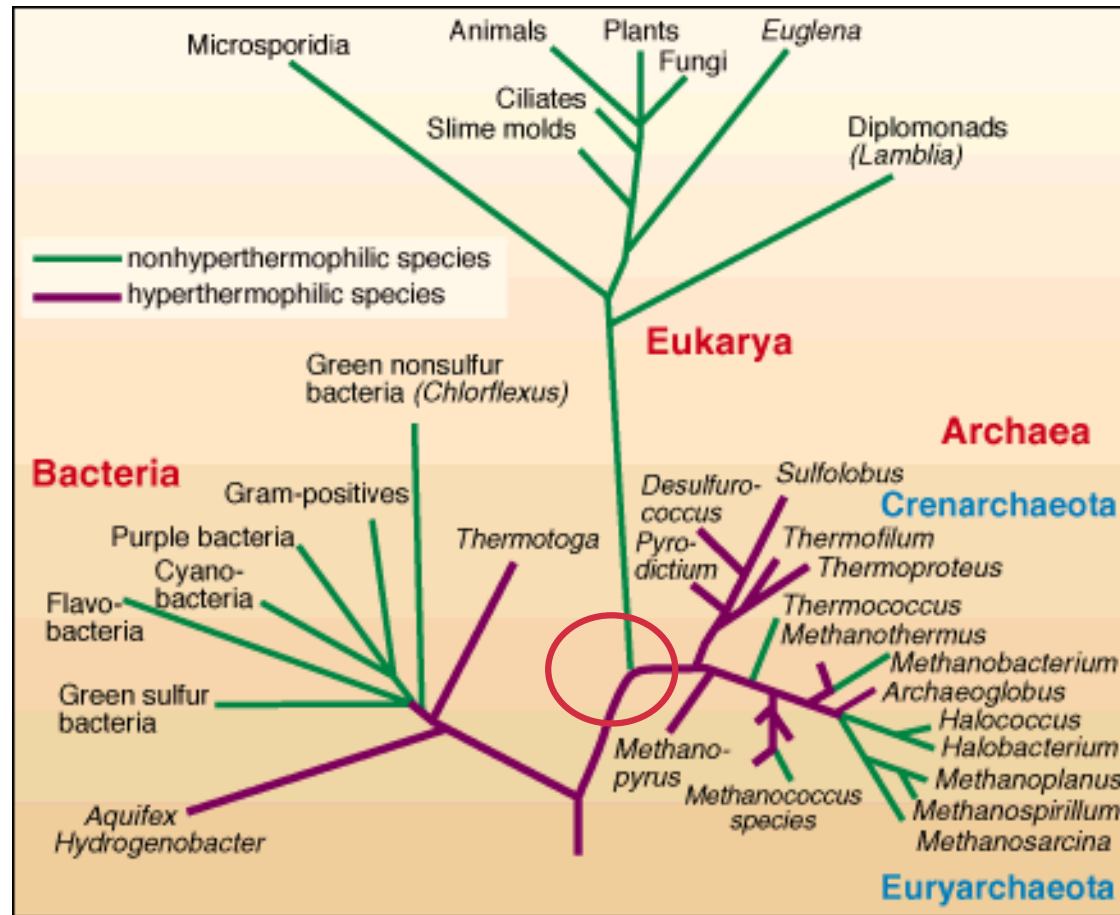


Eukaryotic tree of life

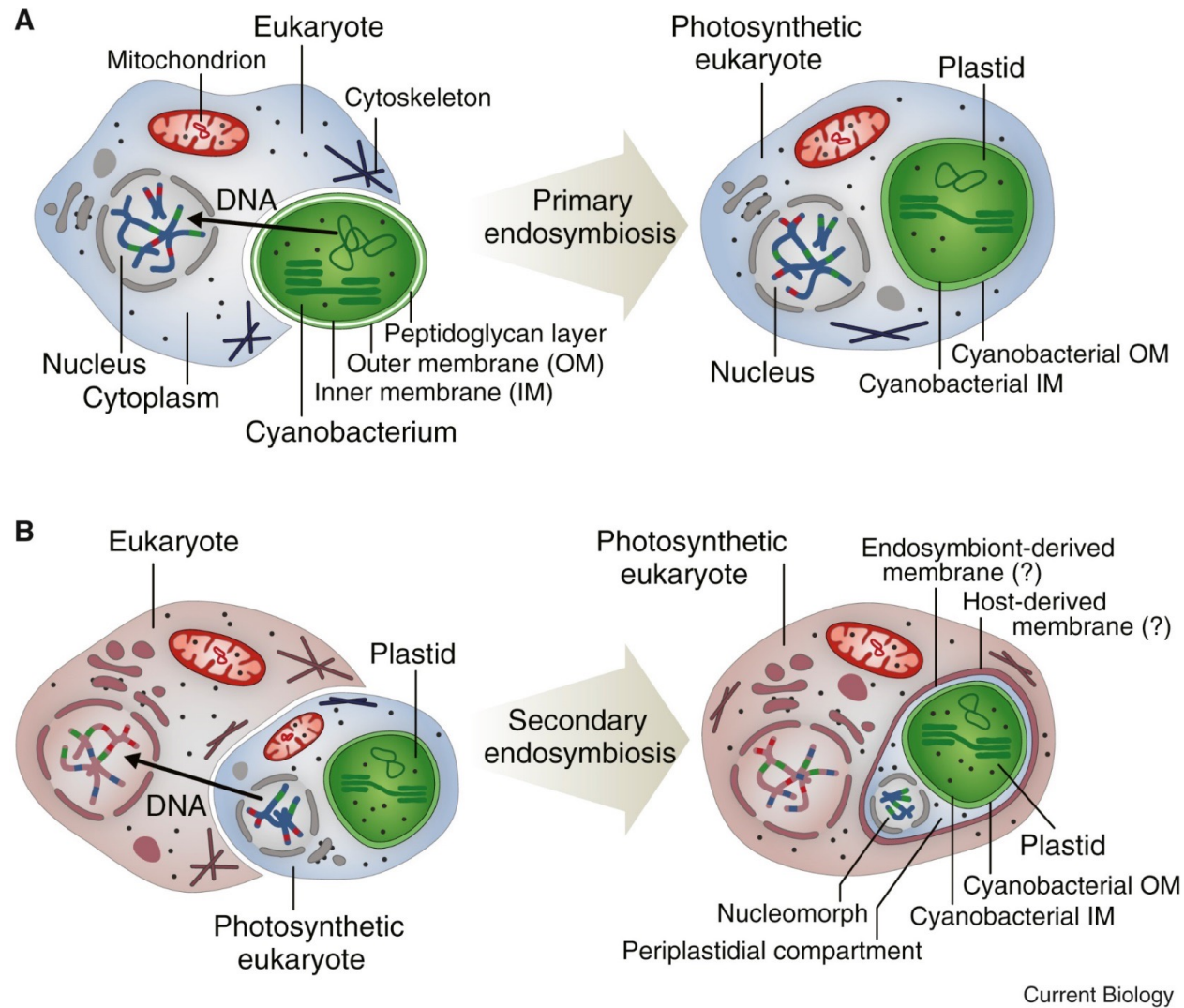
- Eukaryota nested in Archaea
- the Asgard superphylum: the closest prokaryotic relatives of eukaryotes
- their genomes are enriched for proteins formerly considered specific to eukaryotes -> the archaeal host cell already contained many key components that govern eukaryotic cellular complexity



Eukaryotic tree of life



Endosymbioses



Endosymbioses

Lineage	Belong to ^a	Plastid Membranes ^b	Putative Plastid Origin ^c	
Glaucophytes	Archaeplastida	2 (with peptidoglycan)	Cyanobacteria	primary endosymbiosis
Red algae	Archaeplastida	2	Cyanobacteria	
Green algae	Archaeplastida	2	Cyanobacteria	
<i>Paulinella chromatophora</i>	Rhizaria	2 (with peptidoglycan)	Cyanobacteria	
Chlorarachniophytes	Rhizaria	4	Green algae	higher order endosymbioses
Euglenids	Excavates	3	Green algae	
Cryptophytes	Cryptista	4 (with ER connection) ^d	Red algae	
Haptophytes	Haptista	4 (with ER connection) ^d	Red algae	
Ochrophytes	Stramenopiles	4 (with ER connection) ^d	Red algae	
Myxozoans	Alveolates	2–5	Red algae	

^aMain assemblage (no specific taxonomic rank) containing the plastid lineage.

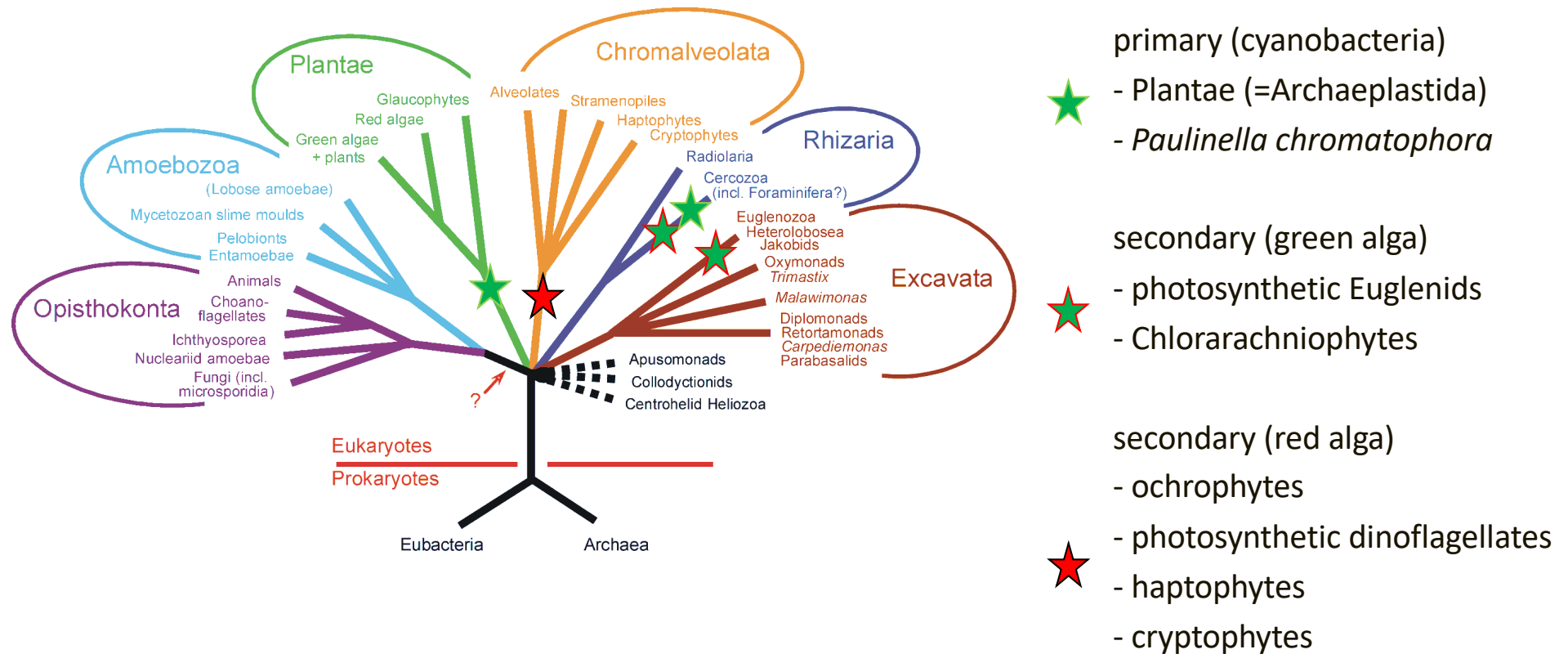
^bNumber of membranes dividing the host cytosol from the plastid stroma.

^cAll plastids ultimately trace back to cyanobacteria, so this column only refers to the direct source of plastids in the lineage under consideration.

^dThe plastid outermost membrane is continuous with the ER in these three lineages.

Endosymbioses

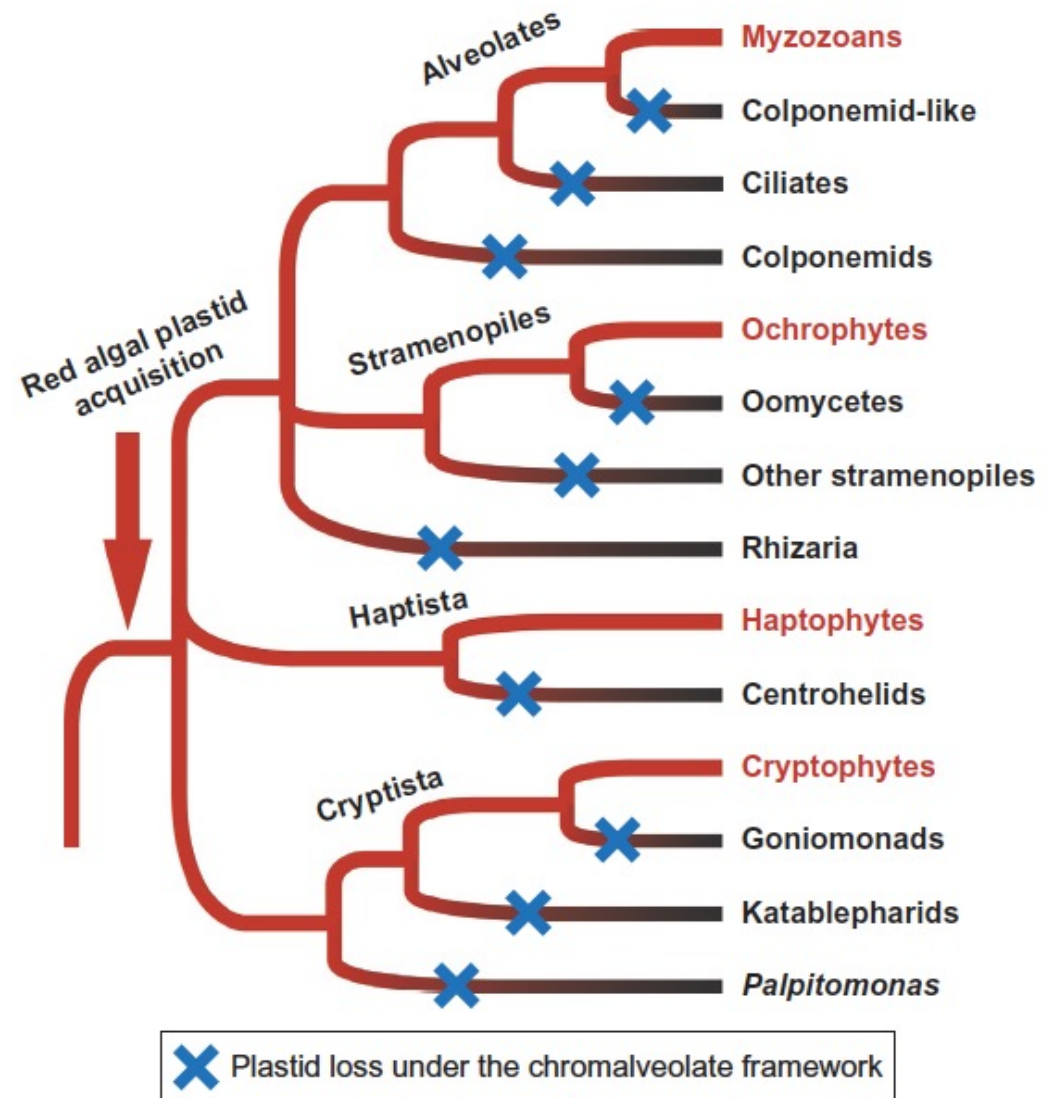
The Chromalveolate hypothesis (Cavalier-Smith 1999)



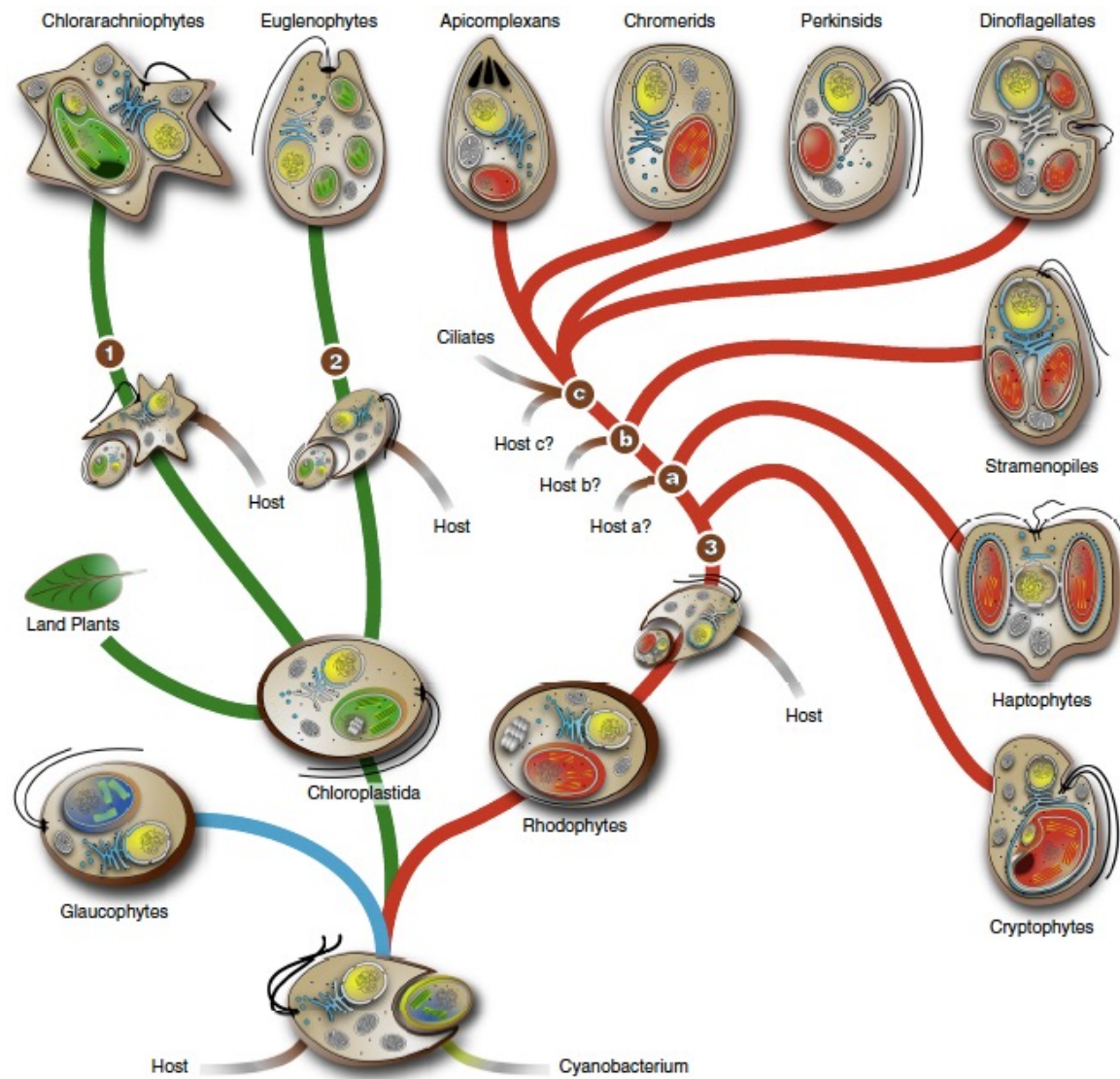
Endosymbioses

Hypothetical tree best fitting **chromalveolate hypothesis**

- the minimum count of plastid losses required is 10 (but in reality much more)
- an alternative better fitting model is needed



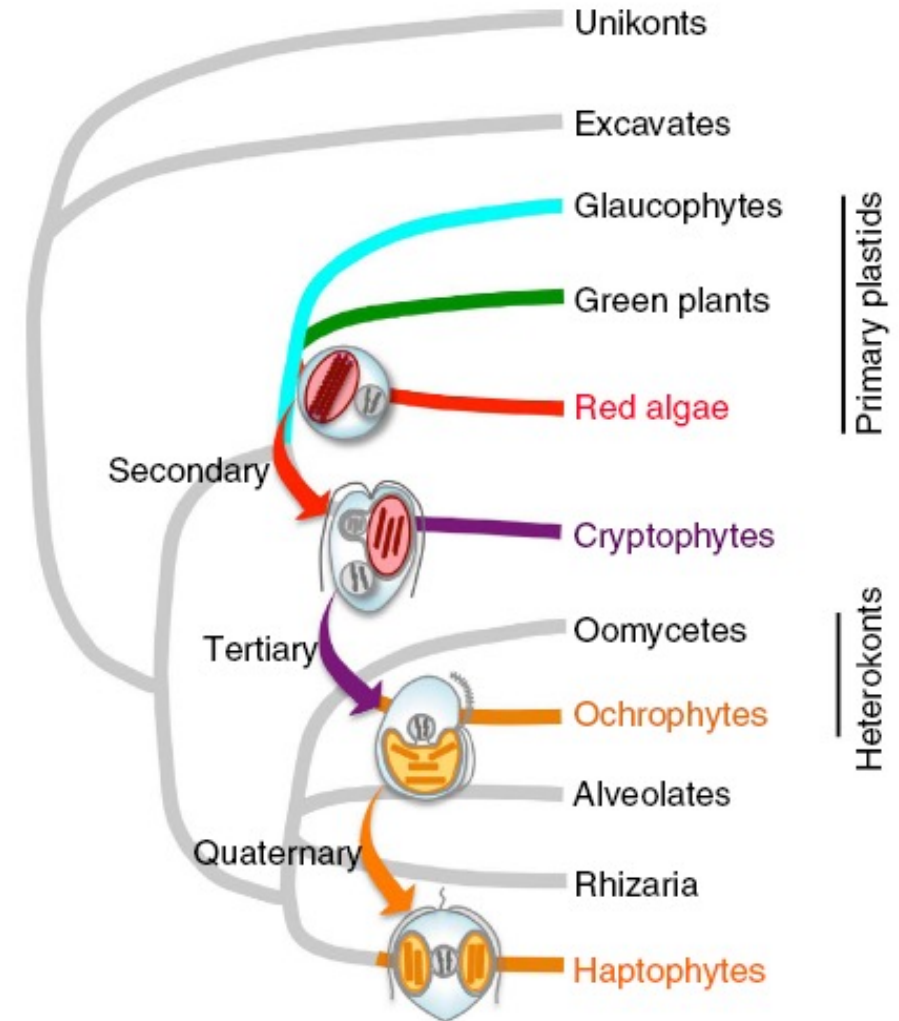
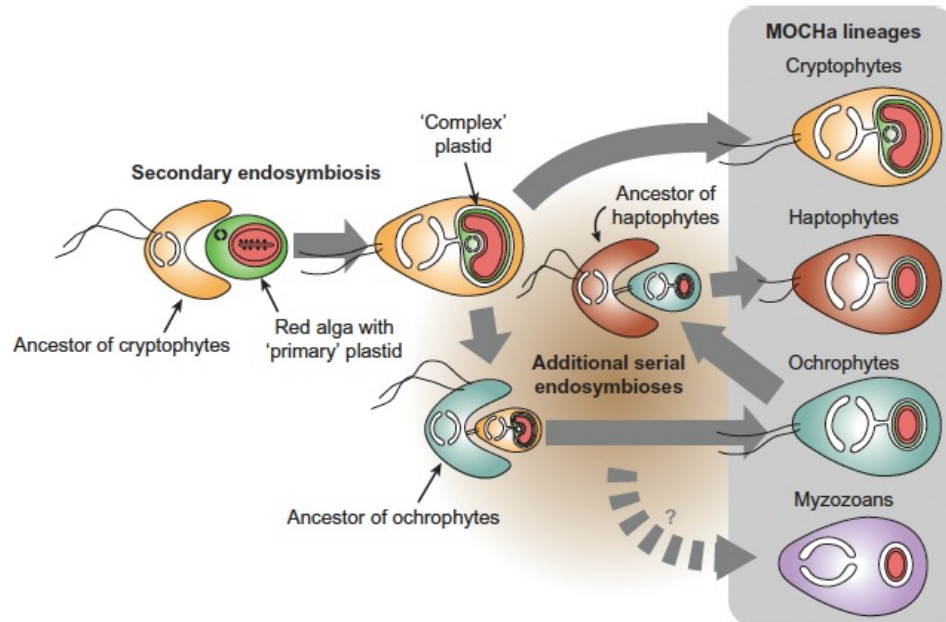
Endosymbioses



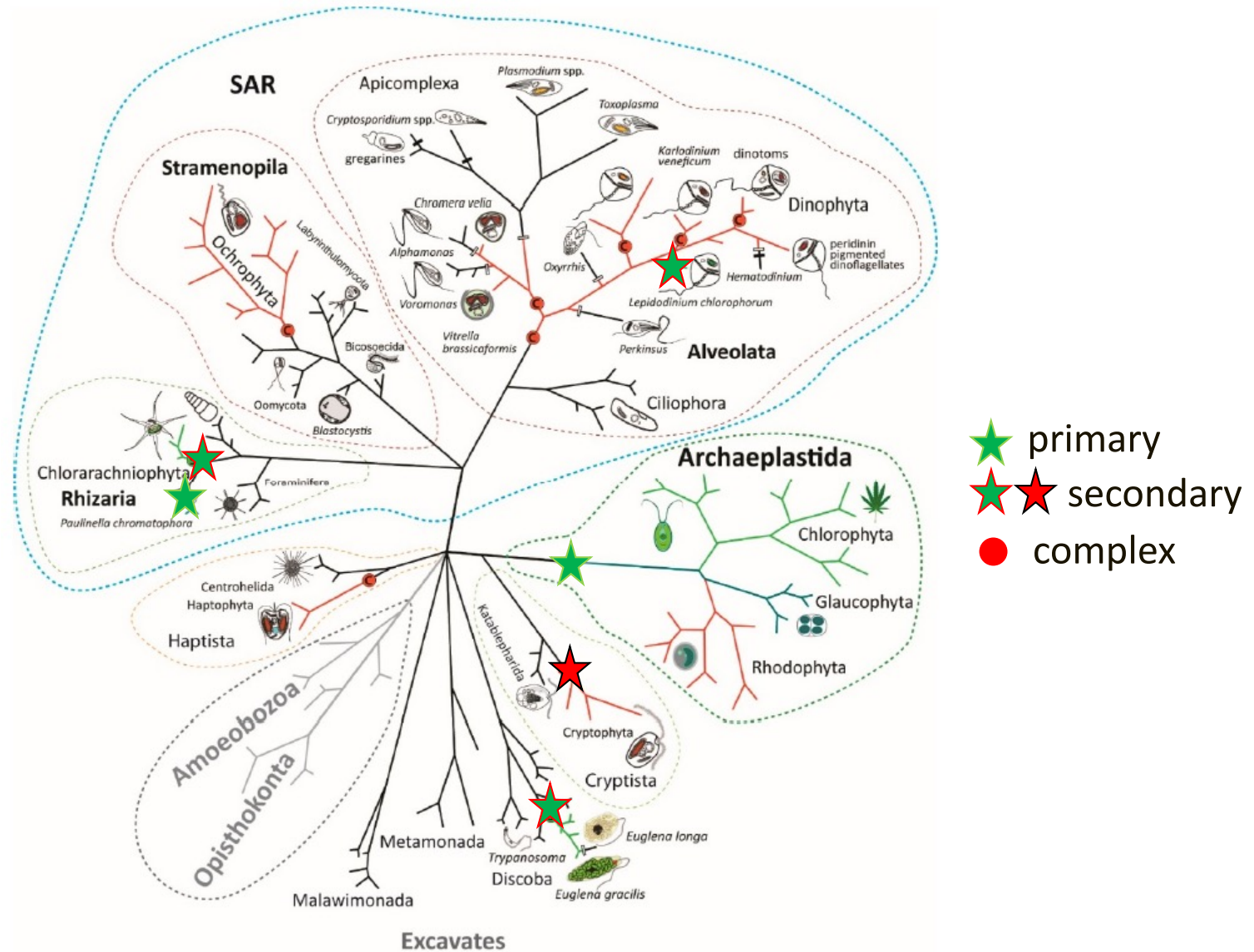
Current Opinion in Microbiology

Endosymbioses

Serial eukaryote-to-eukaryote
endosymbiosis



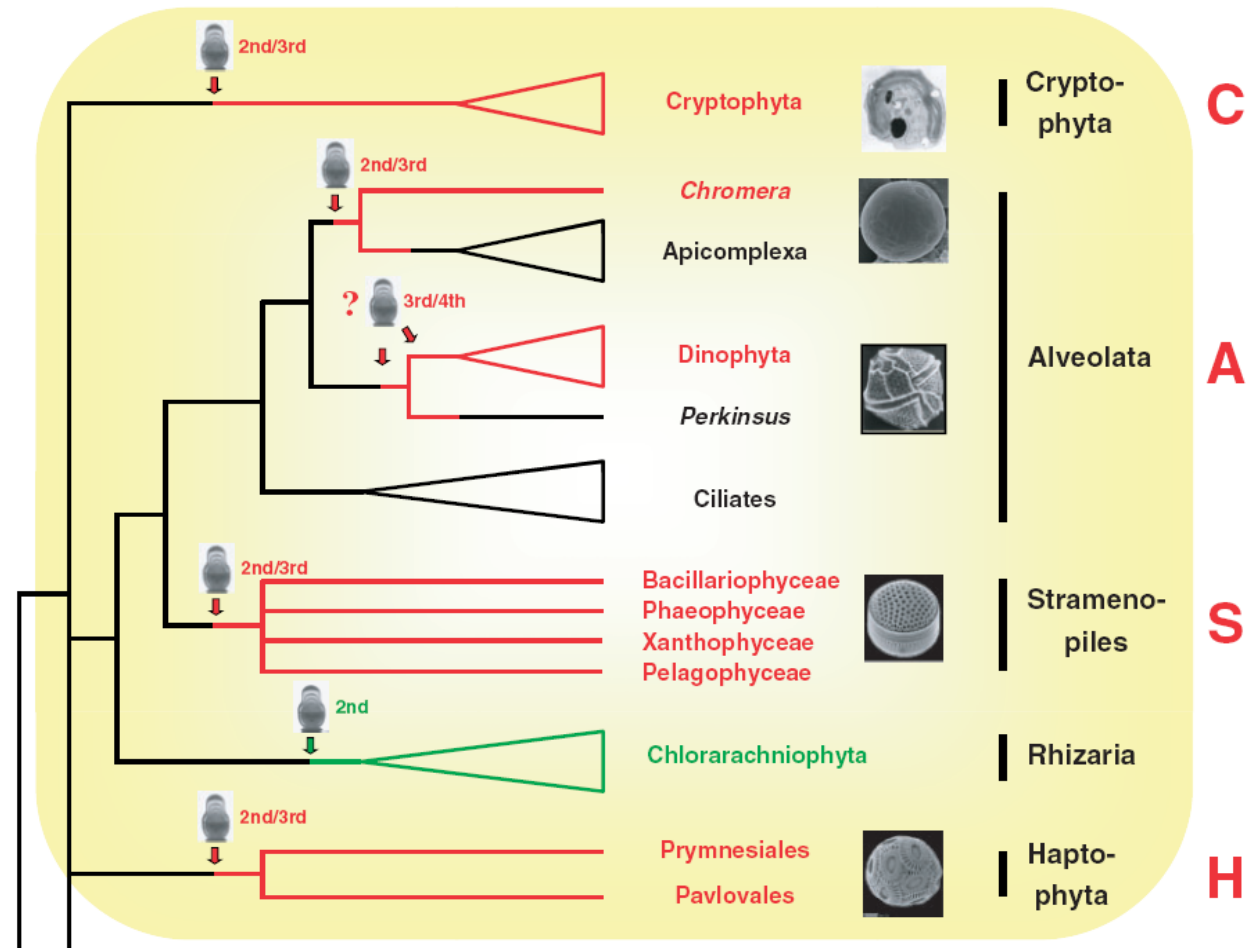
Endosymbioses



Endosymbioses

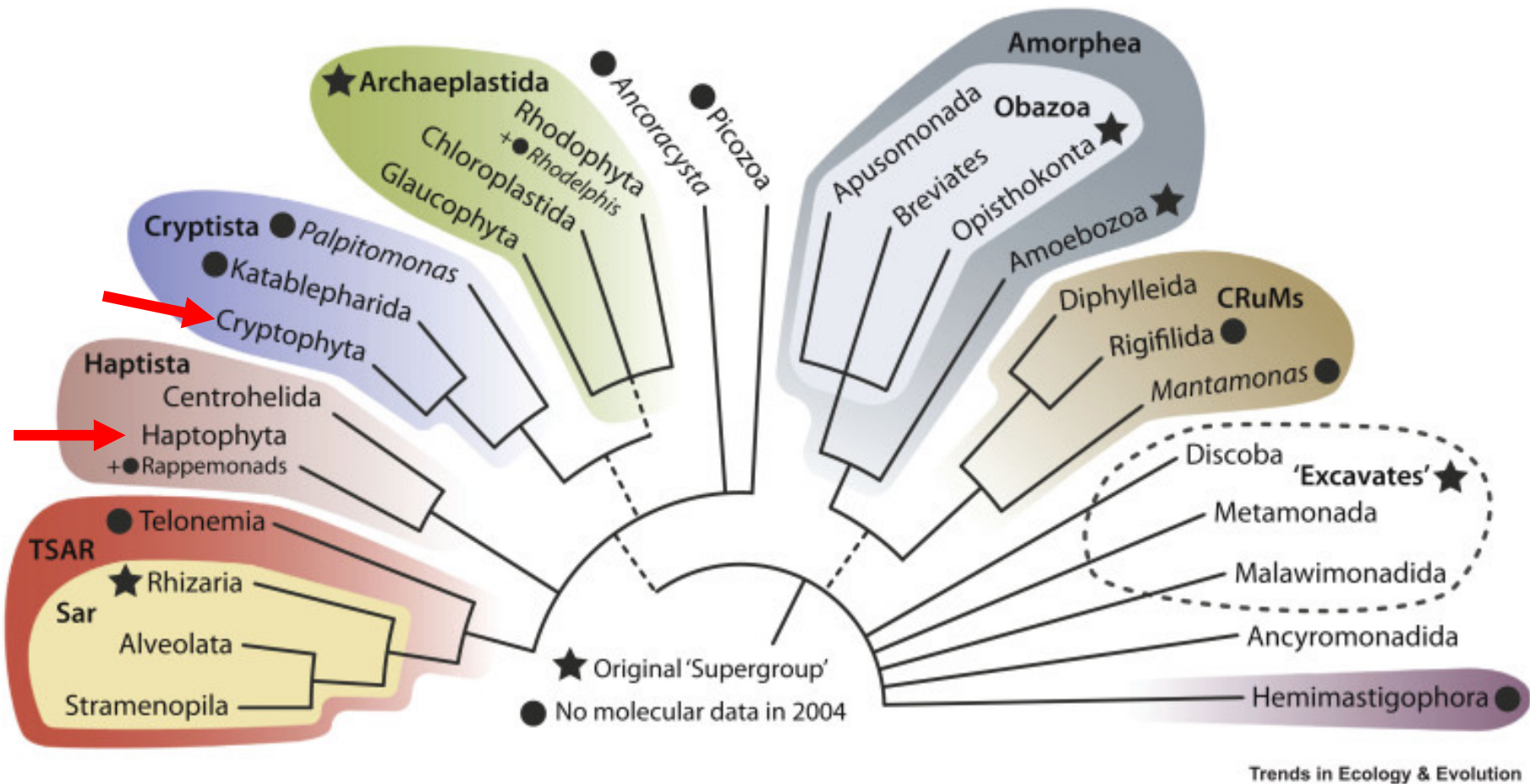
The rhodoplex hypothesis

- the complex plastids from CASH originate from a single secondary endosymbiosis with a rhodophyte
- subsequent eukaryote-to-eukaryote endosymbioses are needed to explain the incongruence observed between plastid-encoded and nuclear-encoded markers

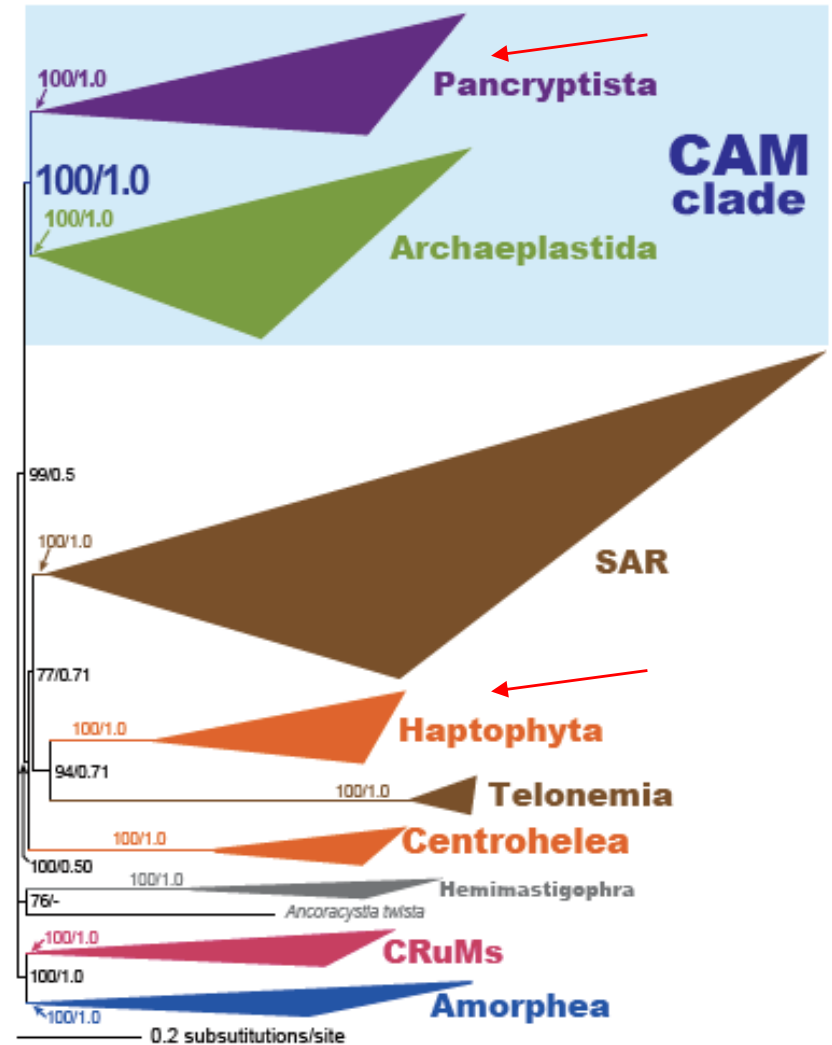
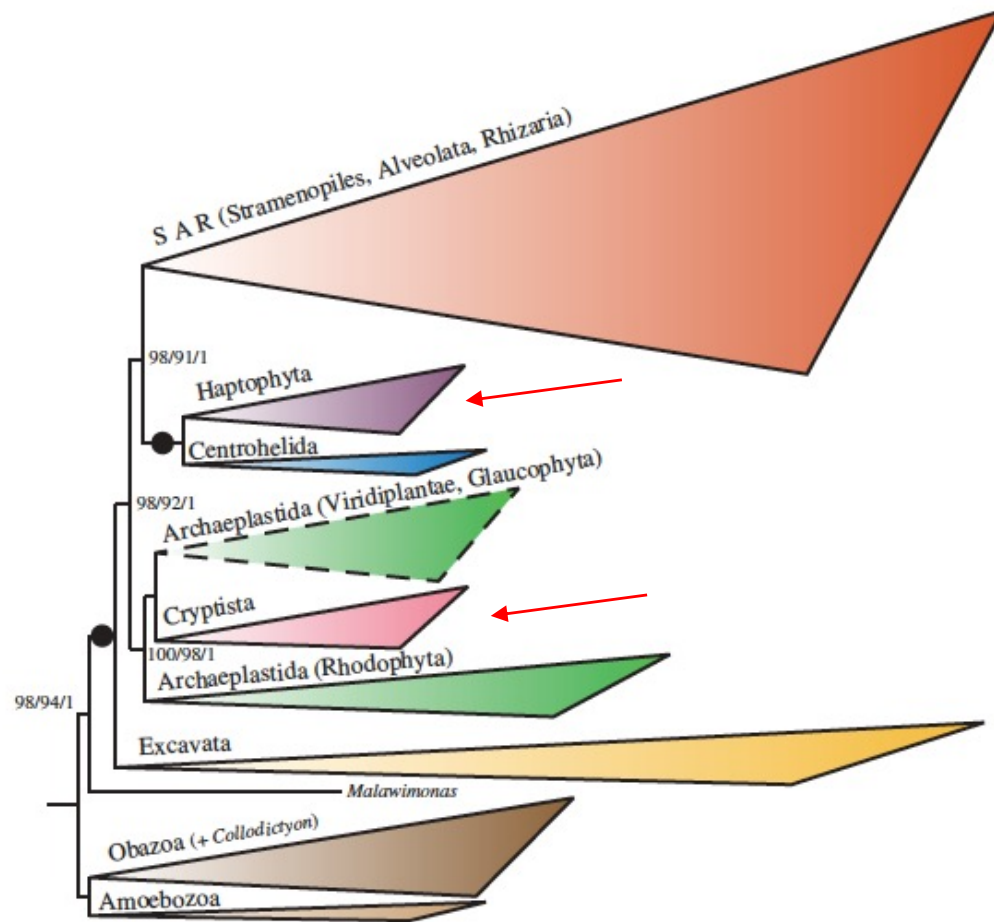


CASH = Cryptophytes, Alveolates, Stramenopiles, and Haptophytes; = complex algae with red plastids; = evolutionary independent groups, i.e. not chromalveolates

Haptophyta and Cryptophyta



Haptophyta and Cryptophyta



Haptophyta and Cryptophyta

<i>Algal group (chapter number)</i>	<i>Photosynthetic pigments^a</i>	<i>Chloroplast outer membranes</i>	<i>Thylakoid associations</i>	<i>Starch-like reserve^b</i>	<i>External covering^c</i>	<i>Flagella</i>
Cyanobacteria (3 & 4)	chl ^a , PE, PC, APC	0	0	Cyanophycean	Pepitoglycan matrices or walls	0
Red algae (5)	chl ^a , PE, PC, APC	2	0	Floridean	Walls with a galactose polymer matrix	0
Green algae (6–9)	chl ^{a, b}	2	2–6	True	Cellulosic walls, scales	0 – many
Euglenoid Algae (10)	chl ^{a, b}	3	3	Paramylon	Pellicle	1–2 emergent
Yellow-green and related algae (11)	chl ^{a, c}	4	3	Chrysolaminarin	Mostly cellulosic walls	2 unequal if present
Chrysophyte algae (12)	chl ^{a, c} fucoxanthin	4	3	Chrysolaminarin	None, scales, lorica	2 unequal
Haptophyte algae (13)	chl ^{a, c} fucoxanthin	4	3	Chrysolaminarin	Nonsiliceous scales	2 equal + haptoneema
Synurophyte algae (14)	chl ^{a, c} fucoxanthin	4	3	Chrysolaminarin	Siliceous scales	2 unequal
Diatoms (15–19)	chl ^{a, c} fucoxanthin	4	4	Chrysolaminarin	Siliceous frustule	1, reproductive cells only
Dinoflagellates (20)	chl ^{a, c} peridinin	3	3	True	Theca	2 unequal
Cryptomonads (21)	chl ^{a, c} PC or PE	4	2	True	Periplast	2 equal
Brown algae (22)	chl ^{a, c} fucoxanthin	4	3	Laminarin	Walls with alginate matrices	2 unequal

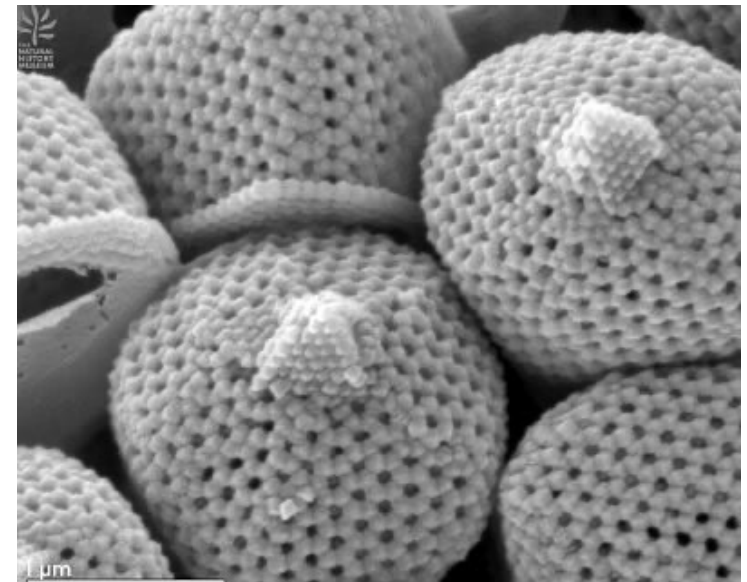
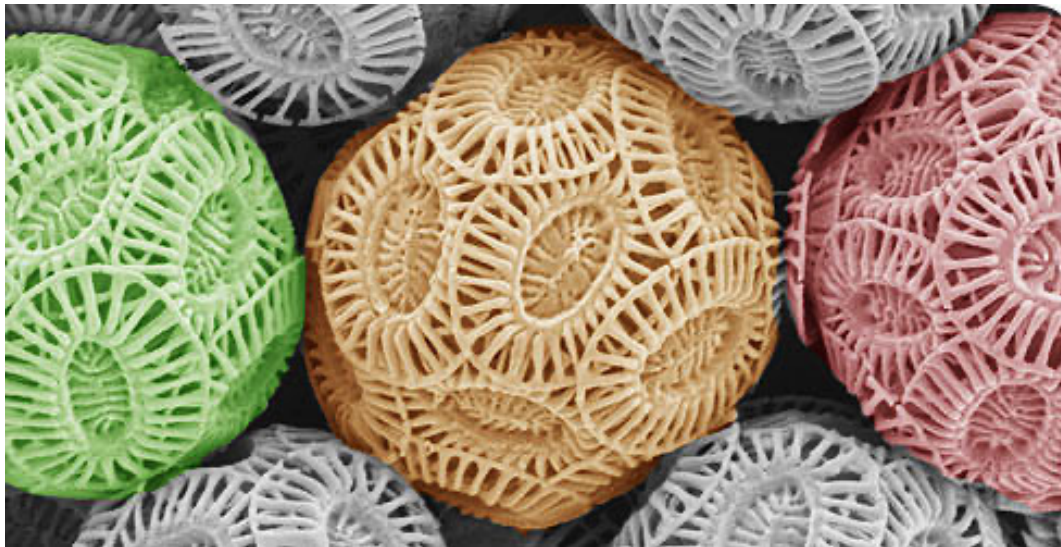
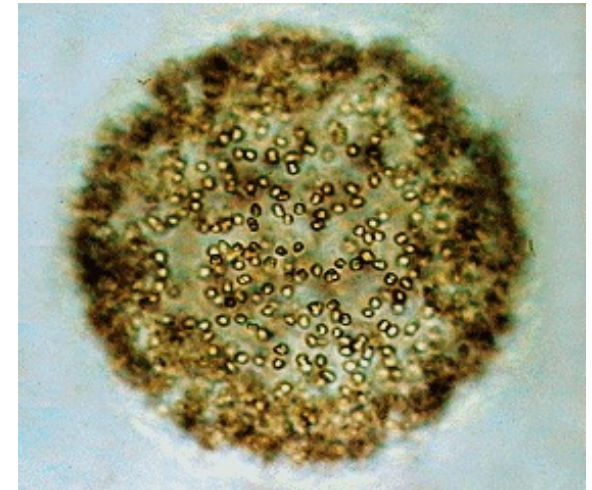
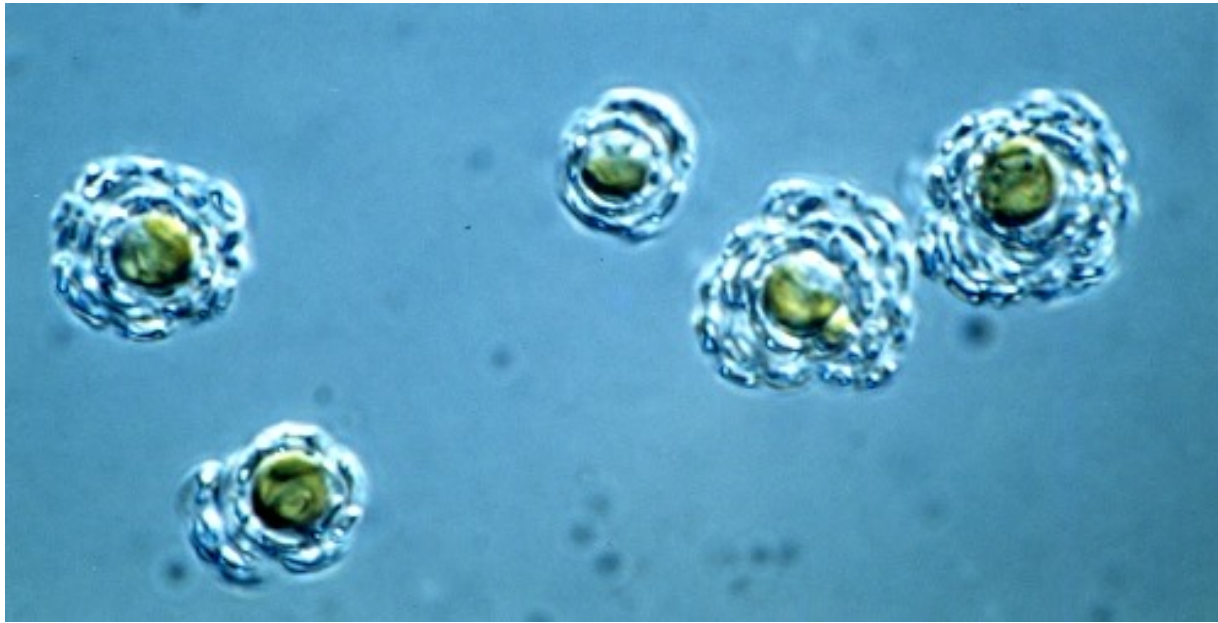
Source: Various phycology textbooks (e.g., Sze, 1998; and Graham and Wilcox, 2000).

^a chl = chlorophyll (green); PE = phycoerythrin (red); PC = phycocyanin (blue); APC = allophycocyanin (blue); fucoxanthin and peridinin (golden to brown).

^b All of the reserves are polymers of glucose. They differ by their linkages: cyanophycean and floridean α 1, 4 and α 1, 6 branches; true starch with amylose α 1, 4 and amylopectin α 1, 4 and α 1, 6 branches; paramylon β 1, 3; chrysolamin and laminarin β 1, 3 and β 1, 6 branches. Only true starch stains positively with iodine (purple to black).

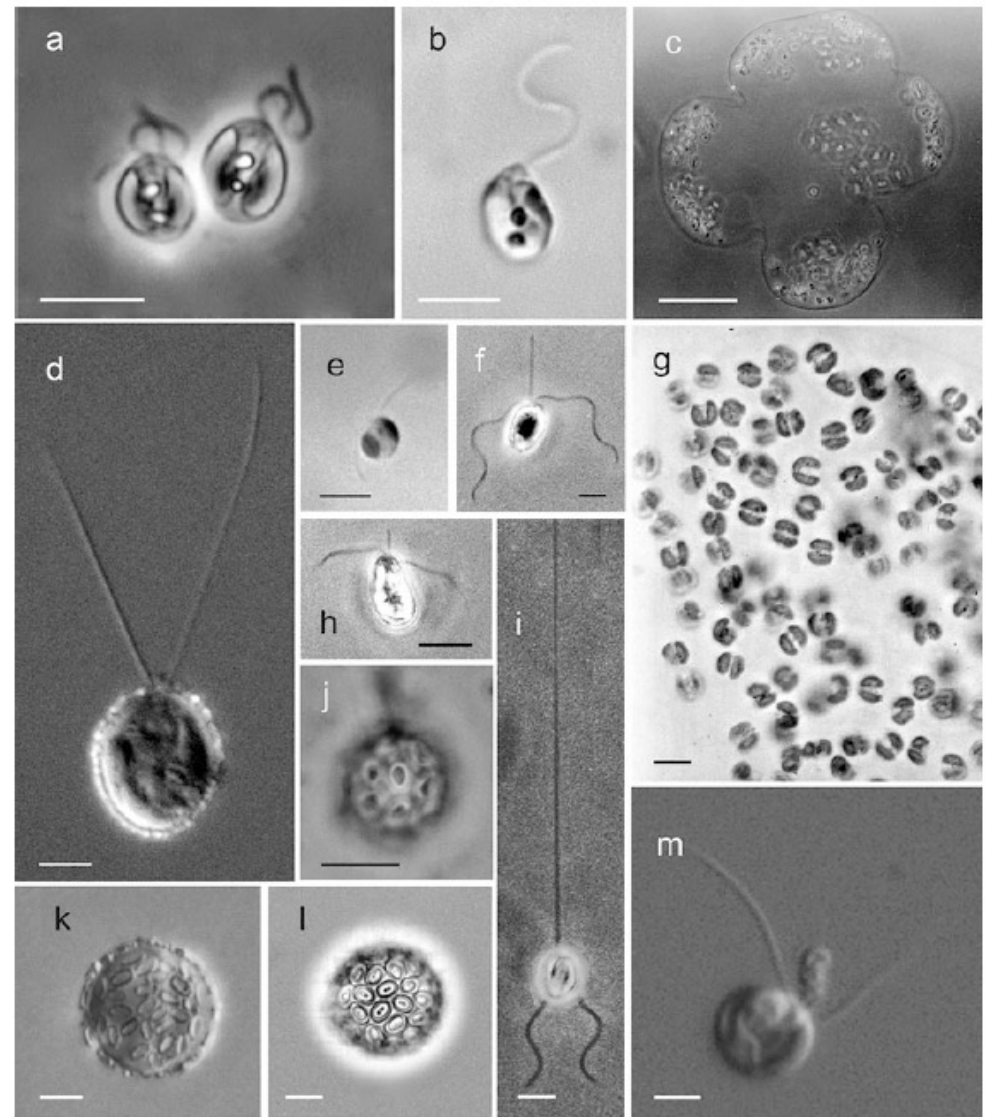
^c Pellicle and periplast within plasma membrane; the rest are external to it.

Haptophyta / Prymnesiophyta



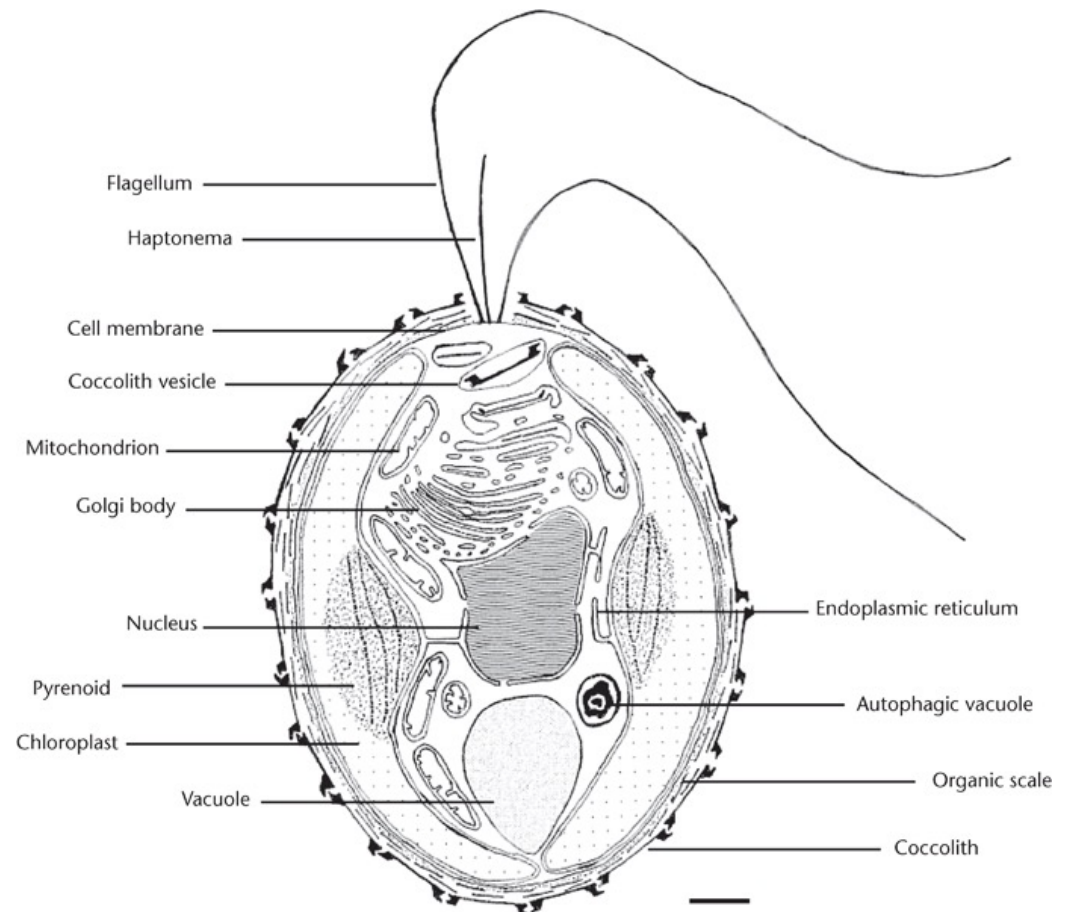
Haptophyta

- unicellular free-living protists
- predominantly planktonic
(picoplankton: $< 3 \mu\text{m}$ and nanoplankton: $2\text{-}20 \mu\text{m}$) and photoautotrophic [but many exhibit phagotrophy, some exclusively heterotrophic]
- main distribution in marine environments worldwide
- some form massive blooms (e.g., *Gephyrocapsa*/*Emiliania huxleyi*, *Phaeocystis* spp.) and some are toxic (e.g., *Prymnesium parvum*)
- important in global carbonate cycle (photosynthesis – 30-50% of total chlorophyll *a* biomass in oceans; calcification of scales - coccolithophores)



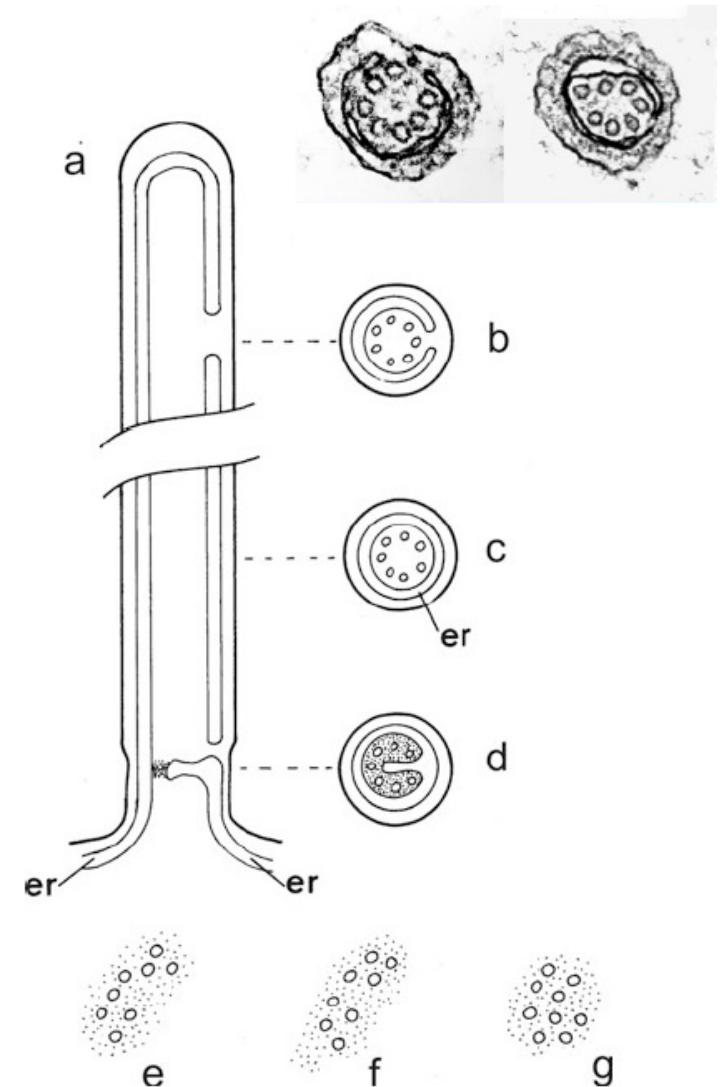
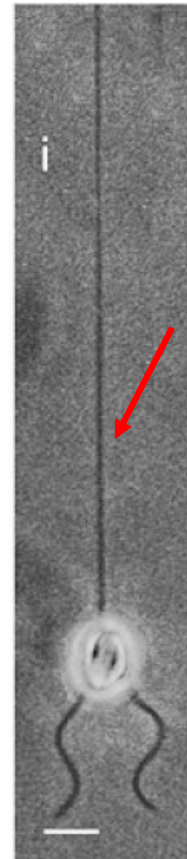
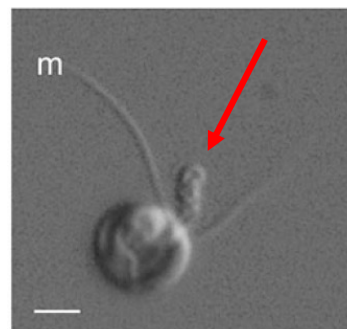
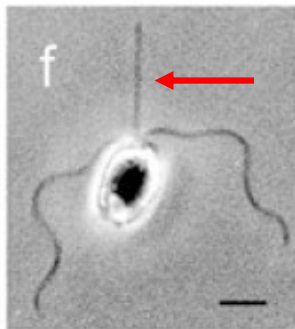
Haptophyta

- two similar naked **flagella**
[markedly unequal flagella with coverings in Pavlovophyceae, or missing in Coccolithophyceae]
- homo- or heterodynamic **flagella**
(9+2 arrangement of microtubules)
- **haptonema**: thin organelle resembling the flagellum [can be reduced in many Coccolithophyceae]; (Greek “*hapto*” = to touch)
- **haptonema**: avoidance response, capturing and gathering food particles, adhering to the substratum



Haptophyta

- **haptonema**: different inner structure;
cross-section: 6-7 microtubules in the free part of the organelle and 8-9 MT at the base; MT are surrounded by the ER
- differ in length among taxa [degraded in some Isochrysidales species, without hapt. MT]
- behavior coordinated with flagella
- cannot beat like a flagellum does but can be bent or coil



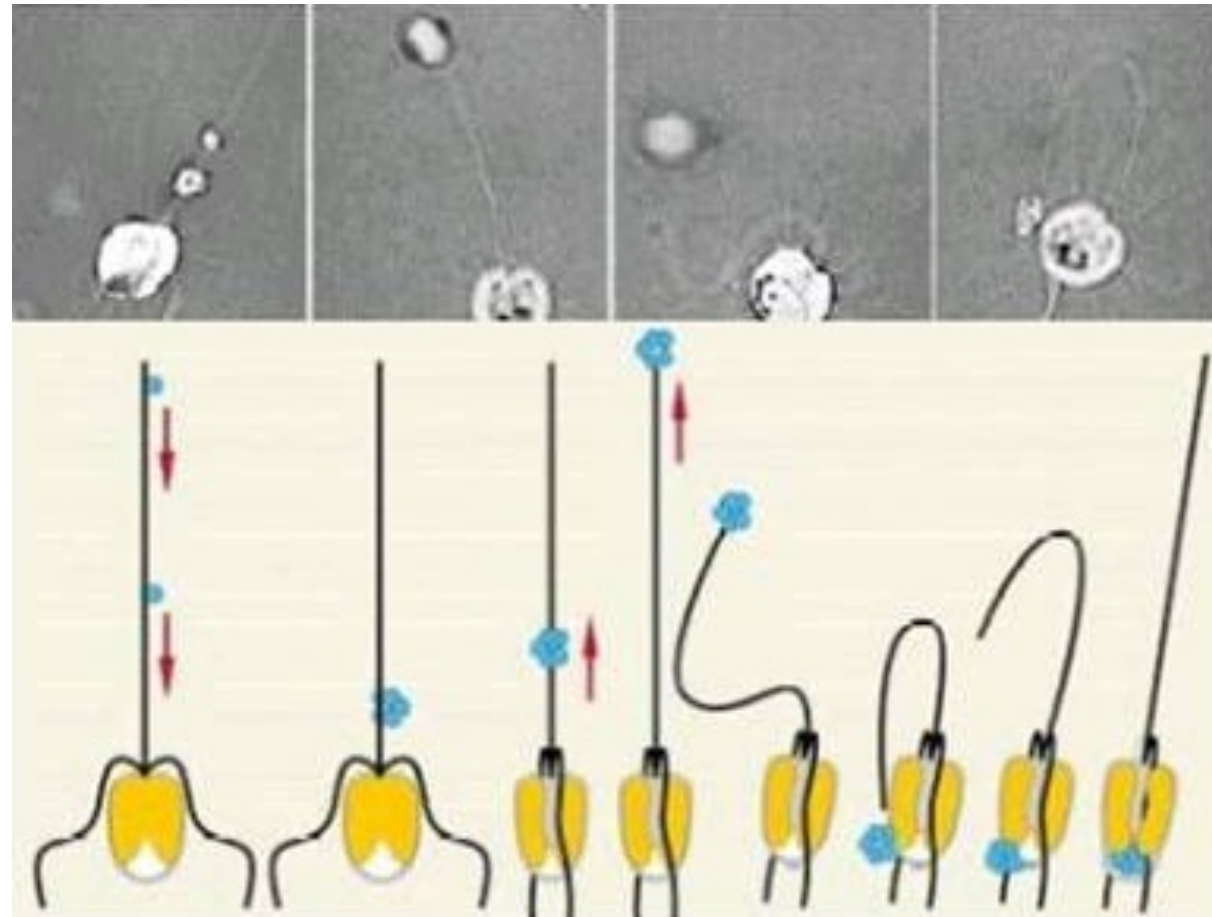
(a) longitudinal section with endoplasmic reticulum (er), microtubuli not shown; (b–d) transverse section at indicated positions showing endoplasmatic reticulum and microtubuli; (e–g) microtubuli within the cell.

Haptophyta

- **ingestion** of bacteria and smaller eukaryotes (phagotrophy)
- **obstacle avoidance:** coiling in extremely short time + the beating of flagella changes to the opposite direction

<https://bit.ly/3TsGZVX>

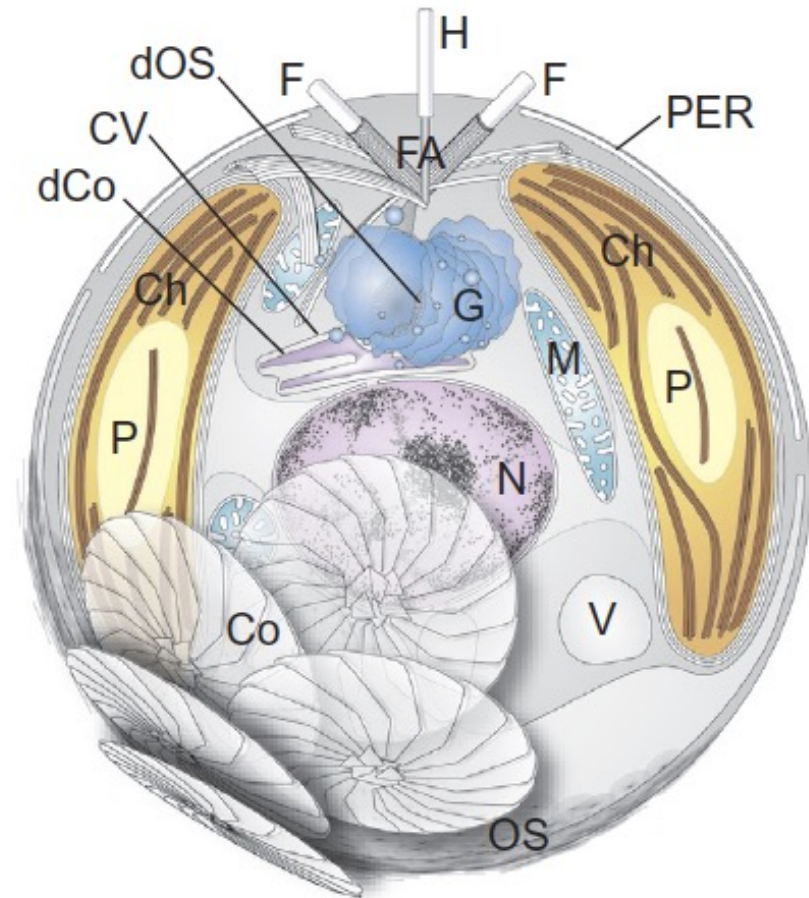
<https://bit.ly/3Tt60jT>



Chrysochromulina. Bacterial particles stick on the haptonema. They are transported to the aggregation center. The clump of bacteria is then transported to the end of haptonema, and delivered to the posterior cell end, where it is engulfed.

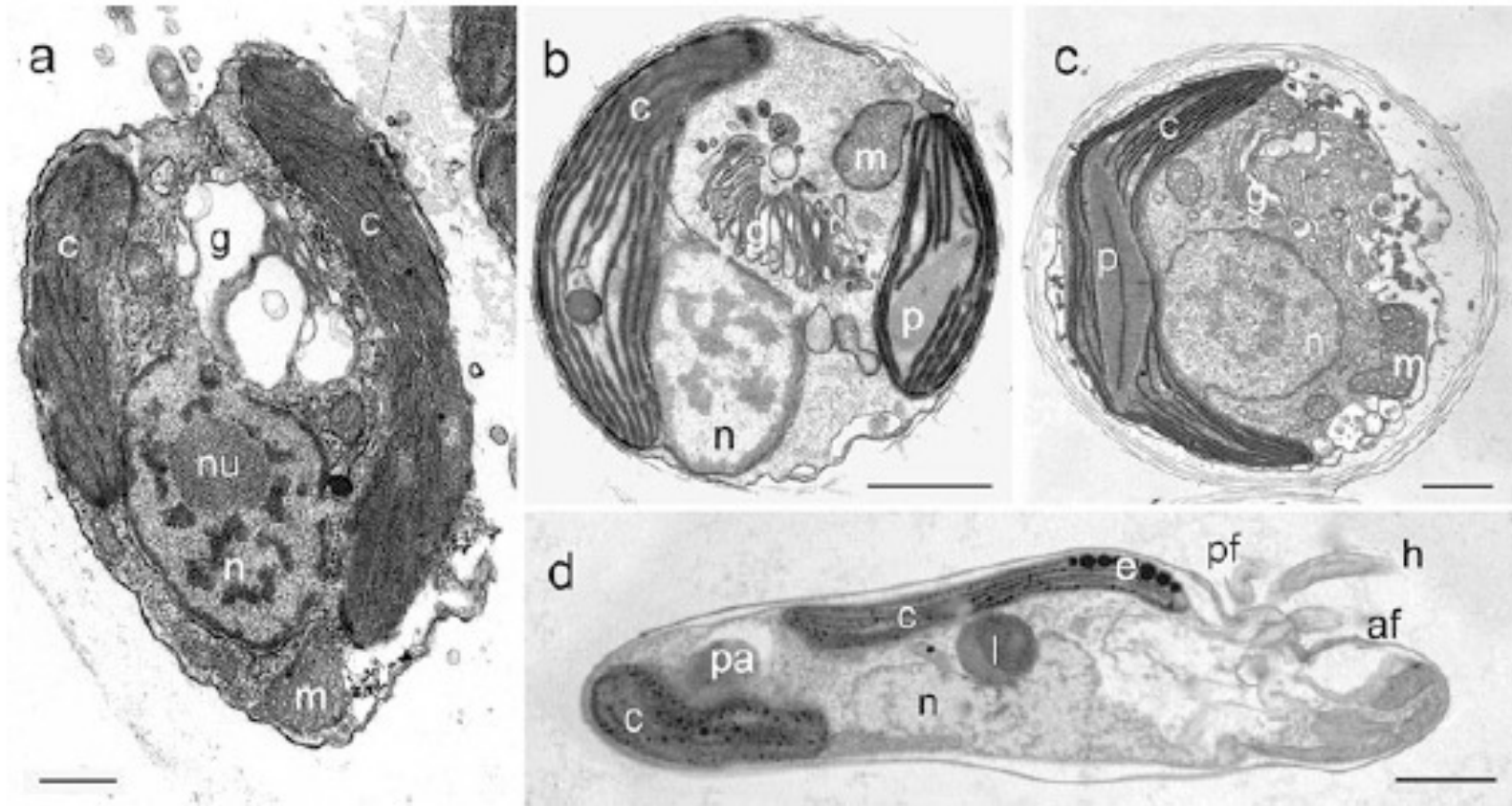
Haptophyta

- cells are surrounded by a cytoplasmatic membrane and peripheral ER [no cell wall]
- one or two yellow/golden-brown red-algal derived chloroplasts (4 membranes, outermost membrane from ER)
- plastids: with pyrenoid; lamellae comprised of 3 thylakoids; no stigma [but occurs in many Pavlovaphyceae]; no girdle lamella
- chlorophyll α and c_2 + carotenoids (fucoxanthin derivatives, beta-carotene)
- storage product: chrysolaminarin [paramylon in Pavlova], lipid bodies
- mitochondria has tubular cristae



Ch - chloroplast; **Co** - coccolith; **CV** - coccolith forming vesicle; **dCo** - developing heterococcolith; **dOS** - developing organic scale; **F** - flagellum; **FA** - flagellar apparatus; **G** - Golgi apparatus; **H** - haptonema; **M** - mitochondrion; **N** - nucleus; **OS** - organic scale; **P** - pyrenoid; **PER** - peripheral endoplasmic reticulum; **V** - vacuole.

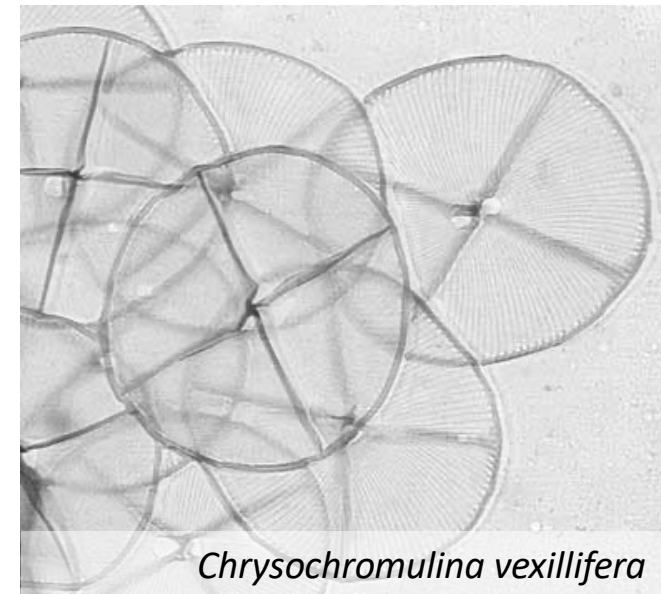
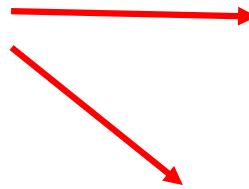
Haptophyta



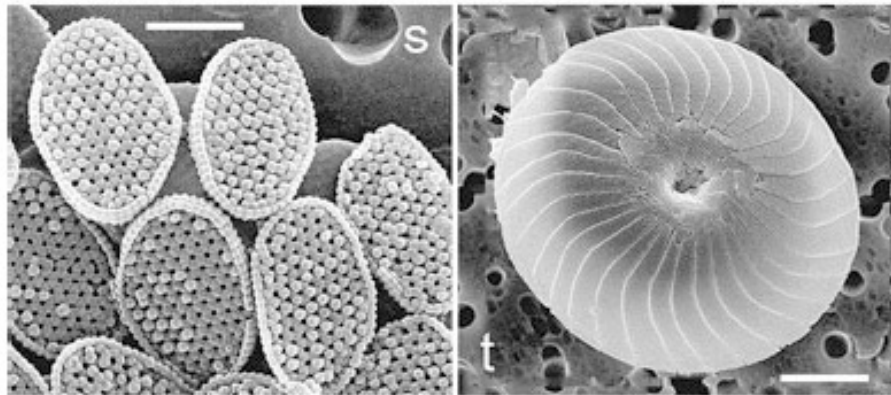
c - chloroplast; **n** - nucleus; **nu** - nucleolus; **pf** - posterior flagellum; **af** - anterior flagellum; **h** - haptonema; **p** - pyrenoid; **l** - lipid droplet; **m** - mitochondrion

Haptophyta

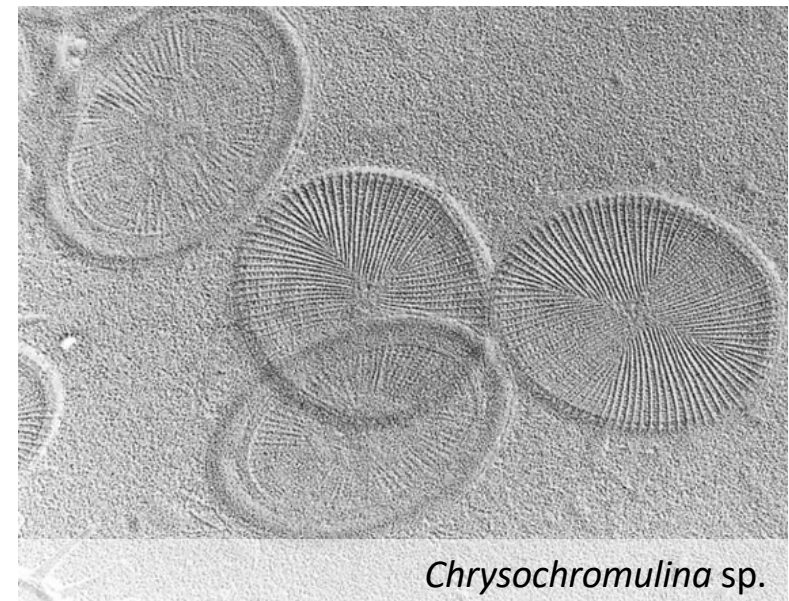
- usually covered with scales (production associated with Golgi vesicles)
- organic nonmineralized body scales (from carbohydrates; spiral microfibrils contain protein and cellulose)
- above body scales silicified (some Prymnesiales) or calcified scales = coccoliths (coccolithophores)
- scales: a diagnostic taxonomic character



Chrysochromulina vexillifera



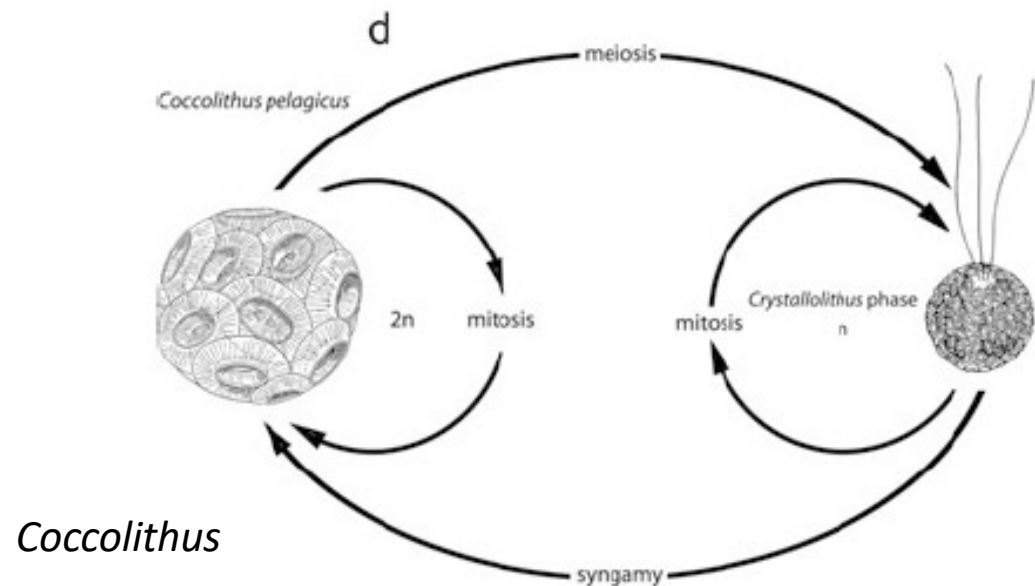
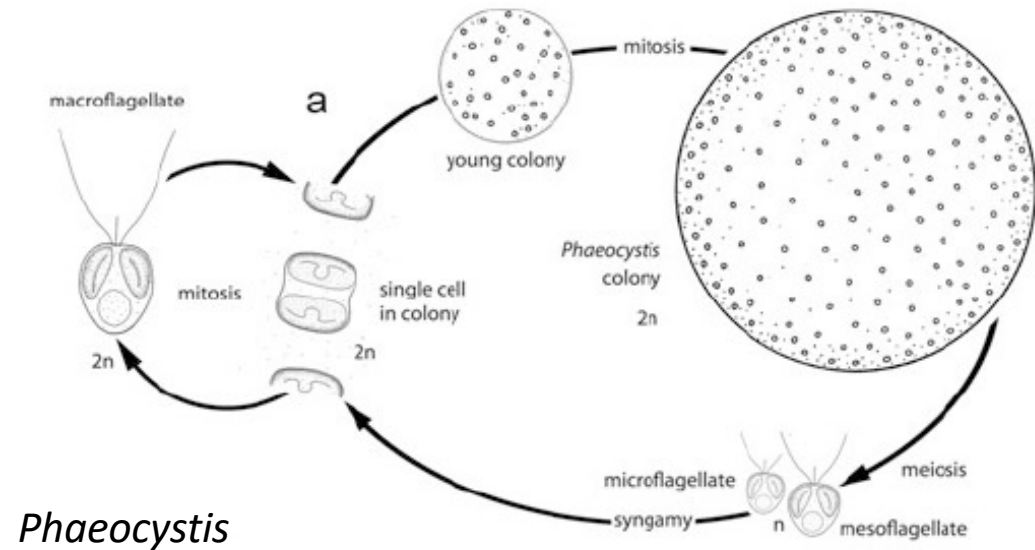
holococcoliths and heterococcolith



Chrysochromulina sp.

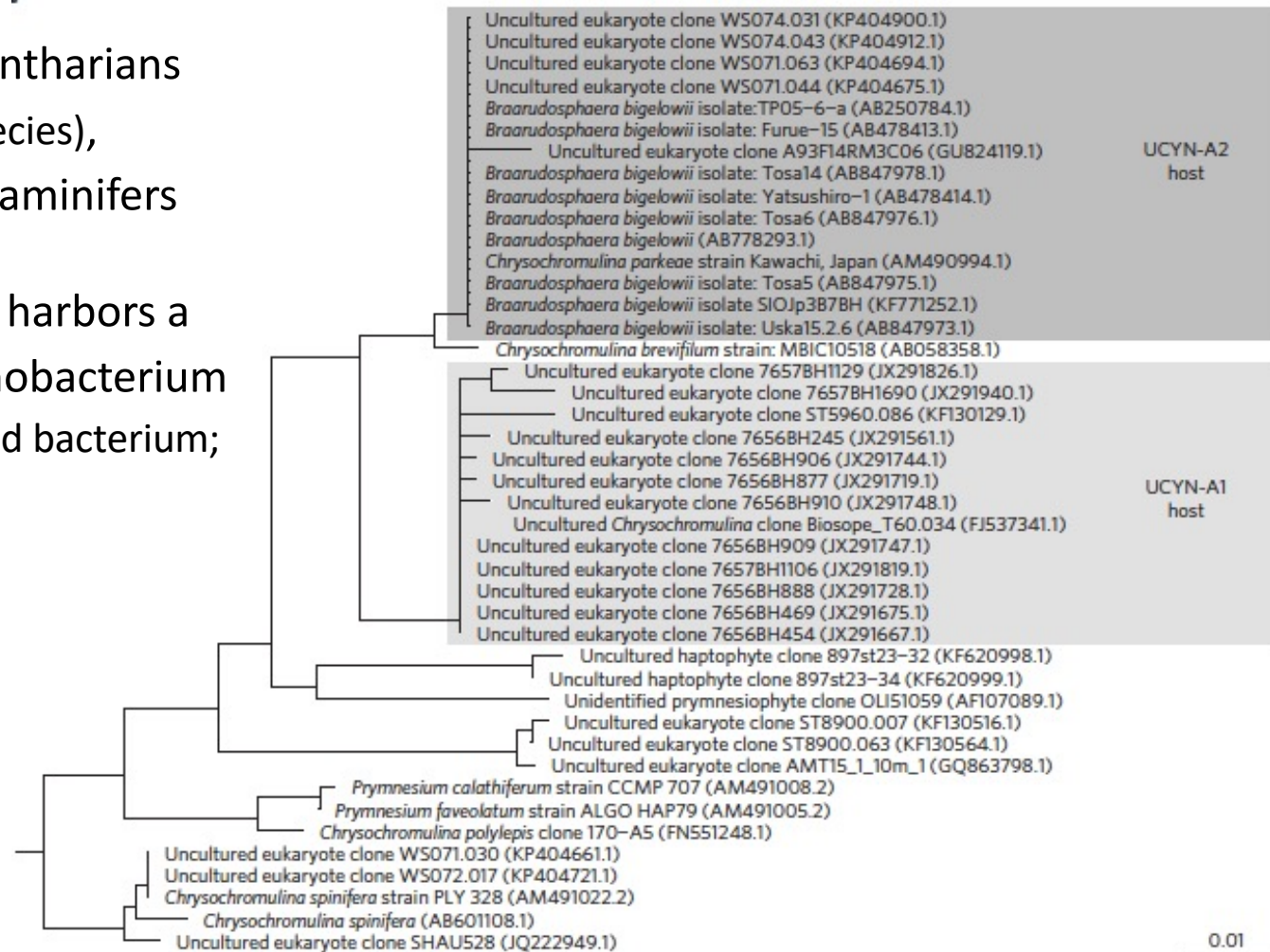
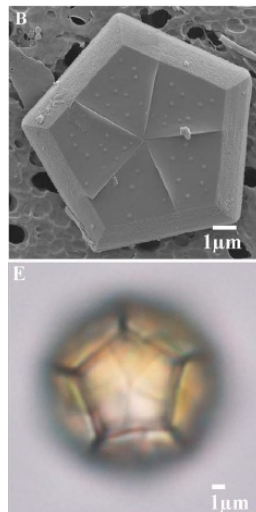
Haptophyta

- mitotic division: cell coverings are equally shared by daughter cells (new scales supply a deficiency after division)
- heteromorphic phases, alteration between motile and nonmotile forms
- alteration of a haploid stage with a diploid stage [so far not observed in Pavlovophyceae]
- haplo-diplontic life cycle: distinct genetics, morphology, and metabolism -> different ecological niche



Haptophyta

- as symbionts of acantharians (some *Phaeocystis* species), radiolarians and foraminifers
- haptophyte lineage harbors a nitrogen-fixing cyanobacterium (UCYN-A = uncultivated bacterium; in spheroid bodies)



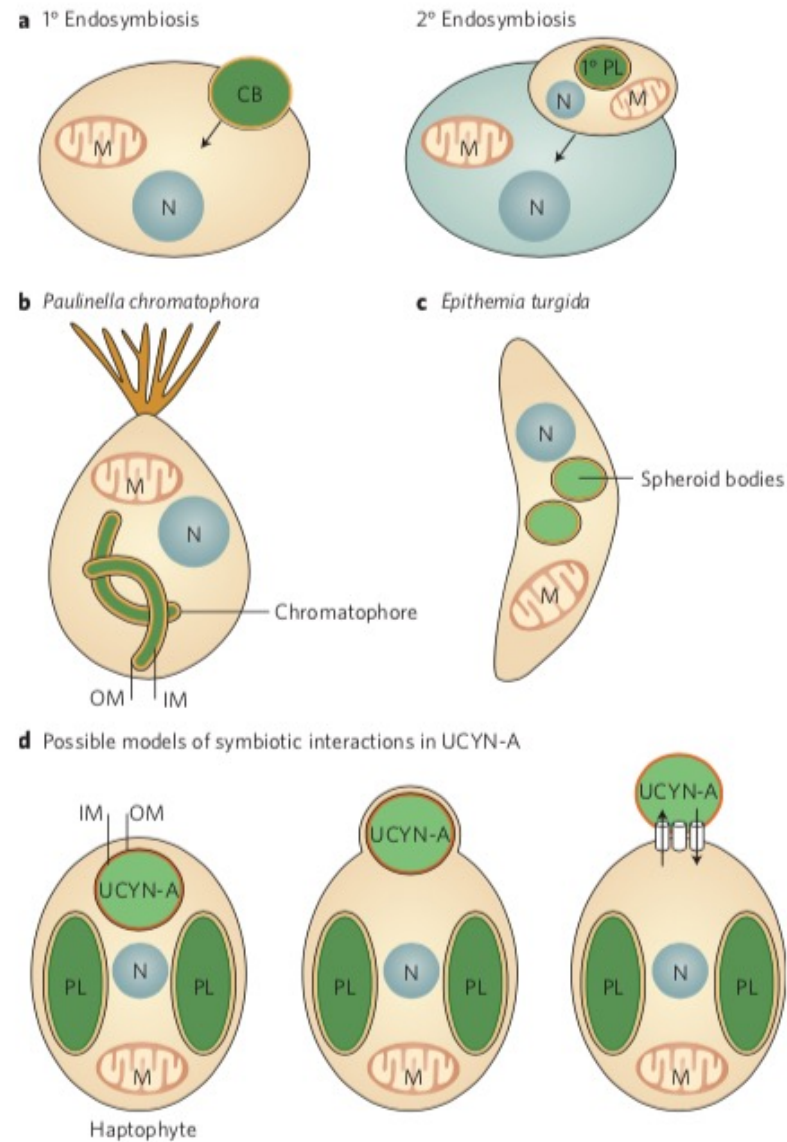
Braarudosphaera bigelowii

Haptophyta

- symbiotic interactions between N_2 -fixing prokaryotes and photosynthetic eukaryotes

d: Possible models of symbiotic interactions between UCYN-A and its haptophyte host.

CB, cyanobacterium; N, host nucleus; PL, plastids; M, mitochondrion; OM, outer membrane; IM, inner membrane.

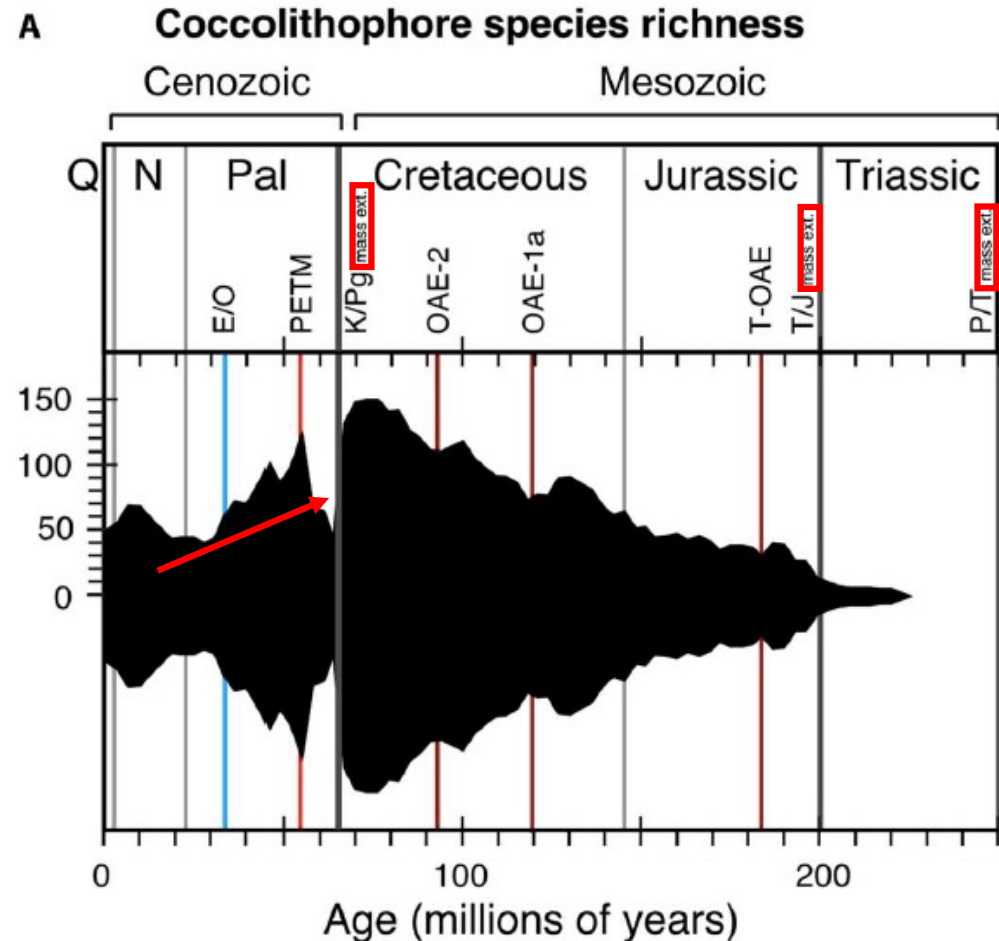


Haptophyta

- first appear in the fossil of the Late Triassic (~ 220 million years ago)
- the higher abundance during the Late Cretaceous (~95 mya)
- ~80 % of all coccolithophorids went extinct during the Cretaceous-Paleogene (K-Pg /K-T) event at the end of the Cretaceous



white chalk cliffs, England



Q - Quaternary; **N** - Neogene; **Pal** - Paleogene; **E/O** - Eocene/Oligocene glacial onset event; **PETM** - Paleocene/Eocene thermal maximum warming event; **K/Pg** - Cretaceous/Paleogene; **OAE** - oceanic anoxic event; **T-OAE** - Toarcian oceanic anoxic event; **T/J** - Triassic/Jurassic; **P/T** - Permian/ Triassic

REPORT

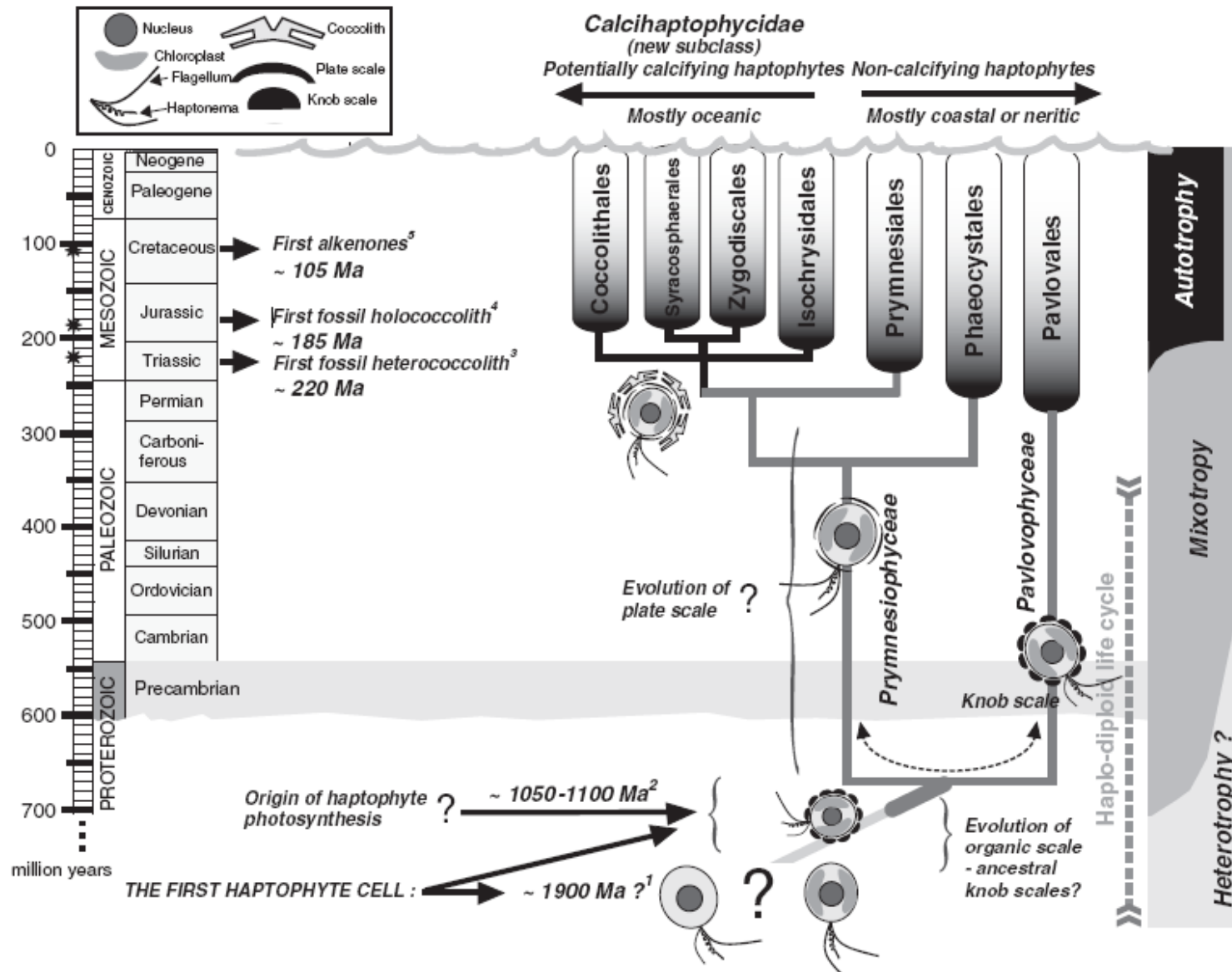
FOSSIL RECORD

Global record of “ghost” nannofossils reveals plankton resilience to high CO₂ and warming

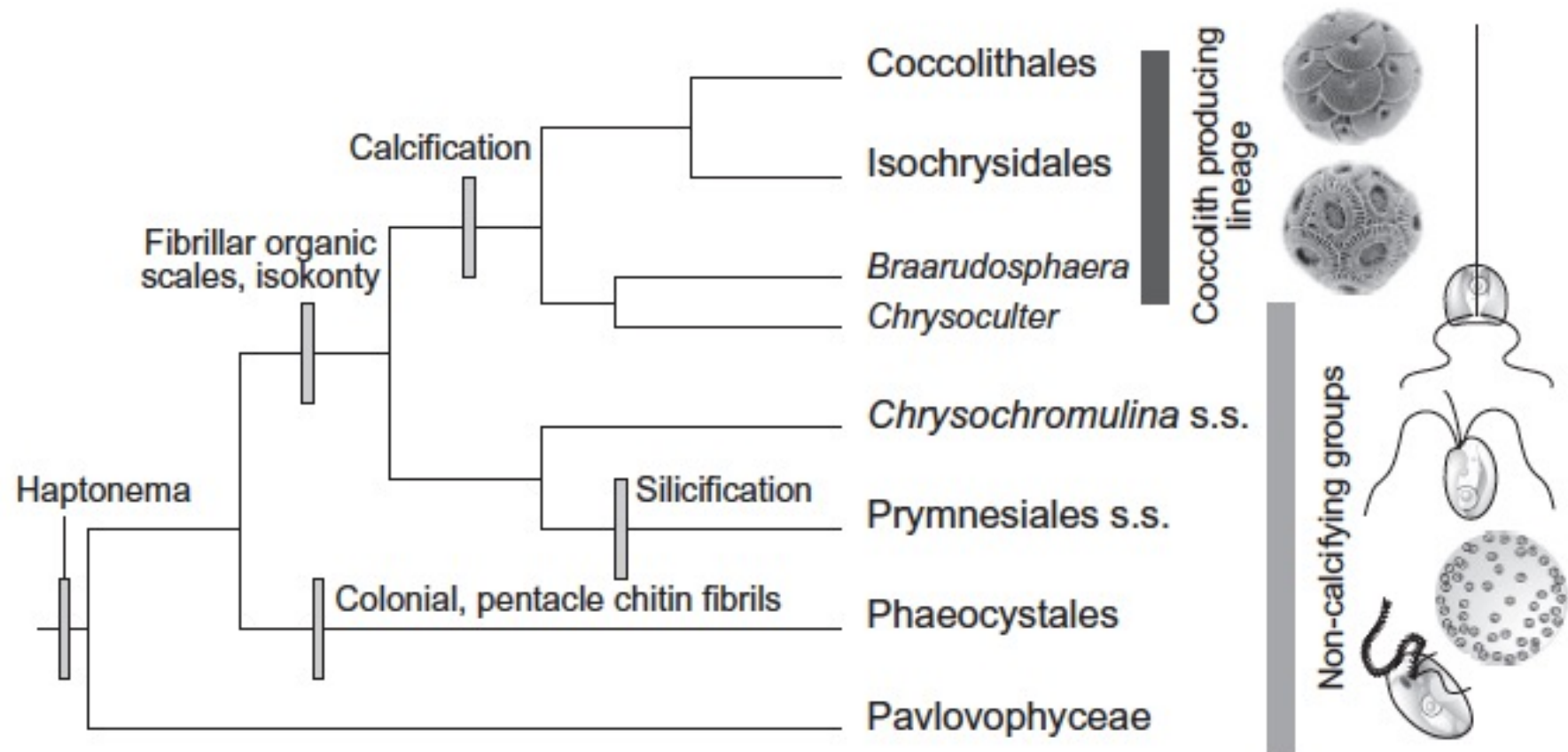
Sam M. Slater^{1*}, Paul Bown², Richard J. Twitchett³, Silvia Danise⁴, Vivi Vajda¹

Predictions of how marine calcifying organisms will respond to climate change rely heavily on the fossil record of nannoplankton. Declines in calcium carbonate (CaCO₃) and nannofossil abundance through several past global warming events have been interpreted as biocalcification crises caused by ocean acidification and related factors. We present a global record of imprint—or “ghost”—nannofossils that contradicts this view, revealing exquisitely preserved nannoplankton throughout an inferred Jurassic biocalcification crisis. Imprints from two further Cretaceous warming events confirm that the fossil records of these intervals have been strongly distorted by CaCO₃ dissolution. Although the rapidity of present-day climate change exceeds the temporal resolution of most fossil records, complicating direct comparison with past warming events, our findings demonstrate that nannoplankton were more resilient to past events than traditional fossil evidence suggests.

Haptophyta



- Haptophyta
- Pavlovophyceae
- Pavlovales (e.g. *Pavlova*, *Rebecca*)
- Coccolithophyceae (= *Prymnesiophyceae*)
- Phaeocystales (*Phaeocystis*)
- Isochrysidales (e.g. *Isochrysis*, *Emiliana*)
- Coccolithales (e.g. *Coccolithus*, *Syracosphaera*)
- Prymnesiales (e.g. *Prymnesium*, *Chrysochromulina*)

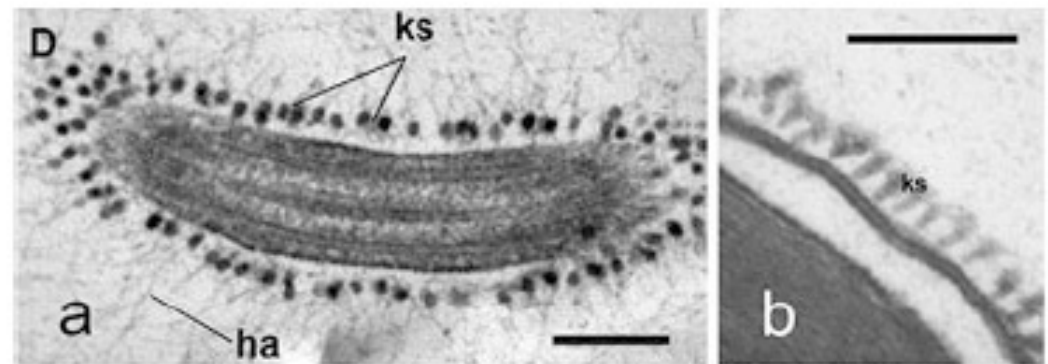
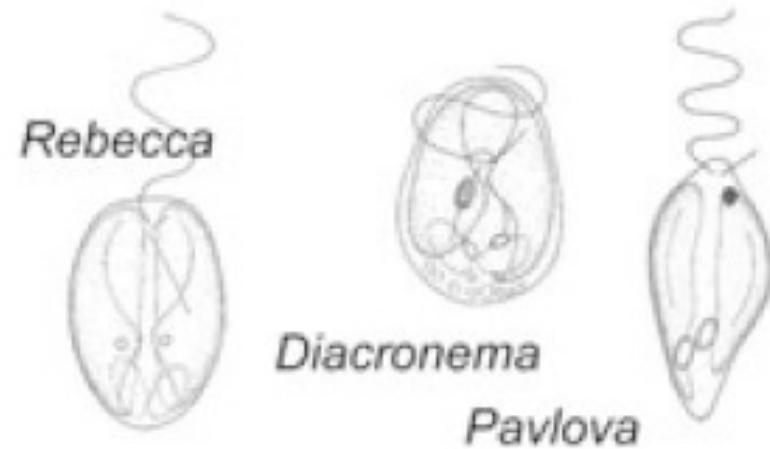


Haptophyta

Phylum Haptophyta	Cells with haptonema
Class: Pavlovophyceae	Cells with knob scales
Order: Pavloales	Motile cells with two unequal flagella, the longer with an investment of small knob scales and fine hairs and the shorter sometimes vestigial. Haptonema present but reduced (e.g., <i>Diacronema</i> , <i>Pavlova</i>)
Class: Coccolithophyceae (Prymnesiophyceae)	Cells with organic scales, with or without coccoliths
Order: Isochrysidales	Motile cells with two equal or subequal flagella, haptonema reduced or absent (e.g., <i>Ruttnera</i> , <i>Emiliana</i> , <i>Isochrysis</i>)
Order: Coccolithales	Cells coccolith bearing many genera with a polymorphic life cycle. Haptonemata recorded in several genera (e.g., <i>Acanthoica</i> , <i>Braarudosphaera</i> , <i>Calyptosphaera</i> , <i>Chrysotila</i> , <i>Coccolithus</i> , <i>Corisphaera</i> , <i>Crenalithus</i> , <i>Cyclolithella</i> , <i>Discosphaera</i> , <i>Helicosphaera</i> , <i>Laminolithus</i> , <i>Rhabdosphaera</i> , <i>Syracosphaera</i> , <i>Umbellosphaera</i> , <i>Umbilicosphaera</i>)
Order: Prymnesiales	Motile cells with two (rarely four) equal or subequal flagella and usually an obvious haptonema which may be very long and coiling (e.g., <i>Chrysochromulina</i> , <i>Haptolina</i> , <i>Prymnesium</i>)
Order: Phaeocystales	Motile cells with two flagella and short haptonema. Flagellate alternating with colonial stage (<i>Phaeocystis</i>)

Pavlovales

- flagella are markedly different in length (the more posteriorly inserted one is much shorter than the anterior flagellum)
- the longer flagellum usually carries a covering of fine hairs and knob-like scales [in some species scales may also be found also on the cell body]
- nonmotile cells may be invested with layers of mucilage
- a stigma or “eyespot” consisting of a single layer of lipid globules
- 4 genera and 13 species (Bendif et al. 2011)



(a–b) Pavlovophyceae: (a) *Pavlova pinguis*, knob scales on flagella; (b) cell surface with knob scales in *Rebecca salina*.

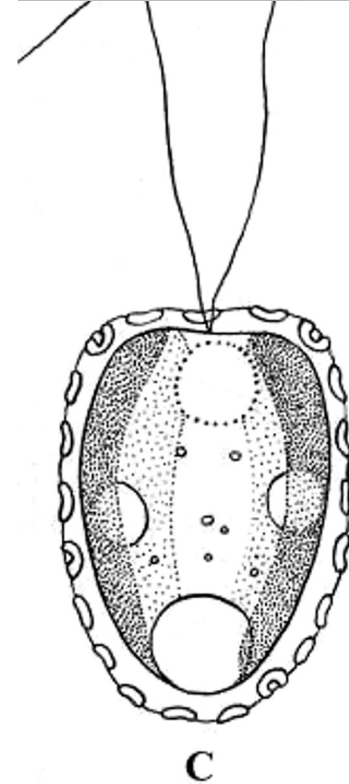
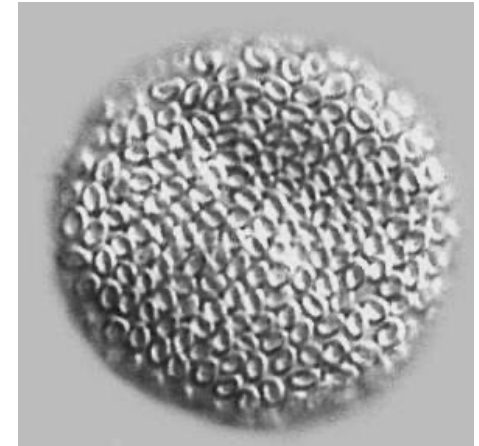
Pavlovales and Prymnesiales

- Pavlovales: common in nearshore coastal environments
- only a few species of Haptophyta have been reported from freshwater environments (lineages in blue) -> representatives of Pavlovales (*Diacronema*/*Exanthemachrysis*, *Pavlova*) and Prymnesiales (*Hymenomonas*, *Prymnesium*, *Chrysochromulina*)



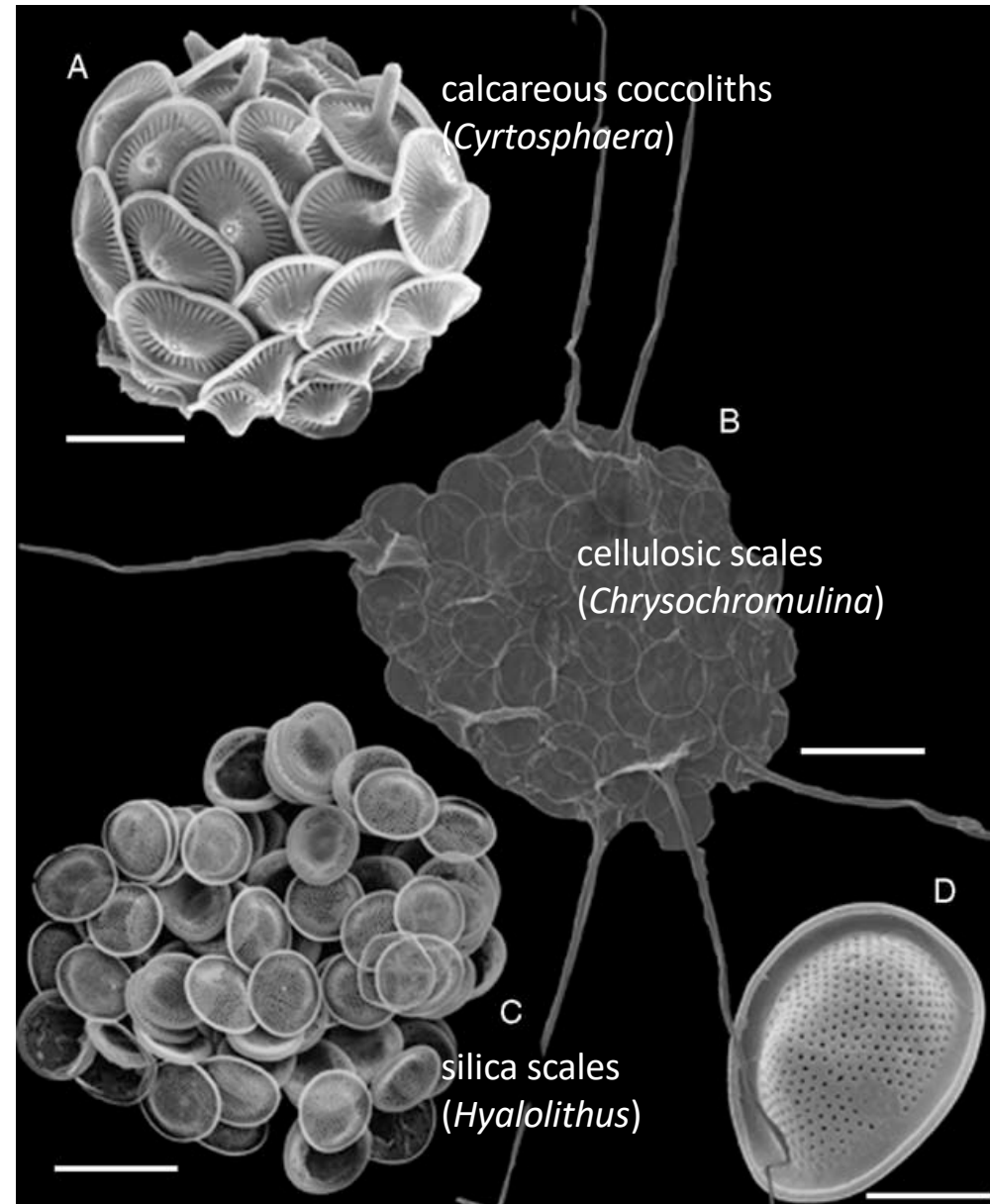
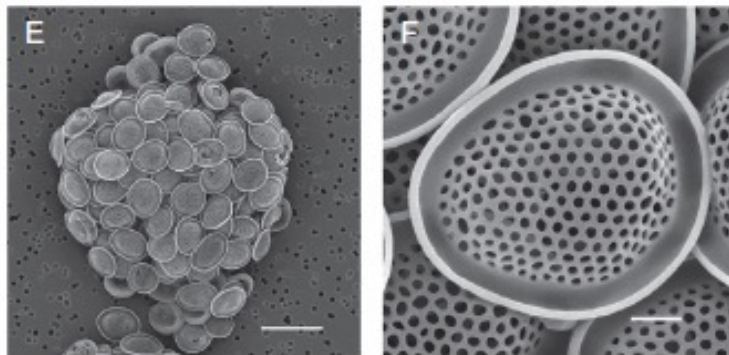
Prymnesiales

- *Hymenomonas roseola*, calcified scales
- freshwater in metaphyton of clear, oligotrophic waters



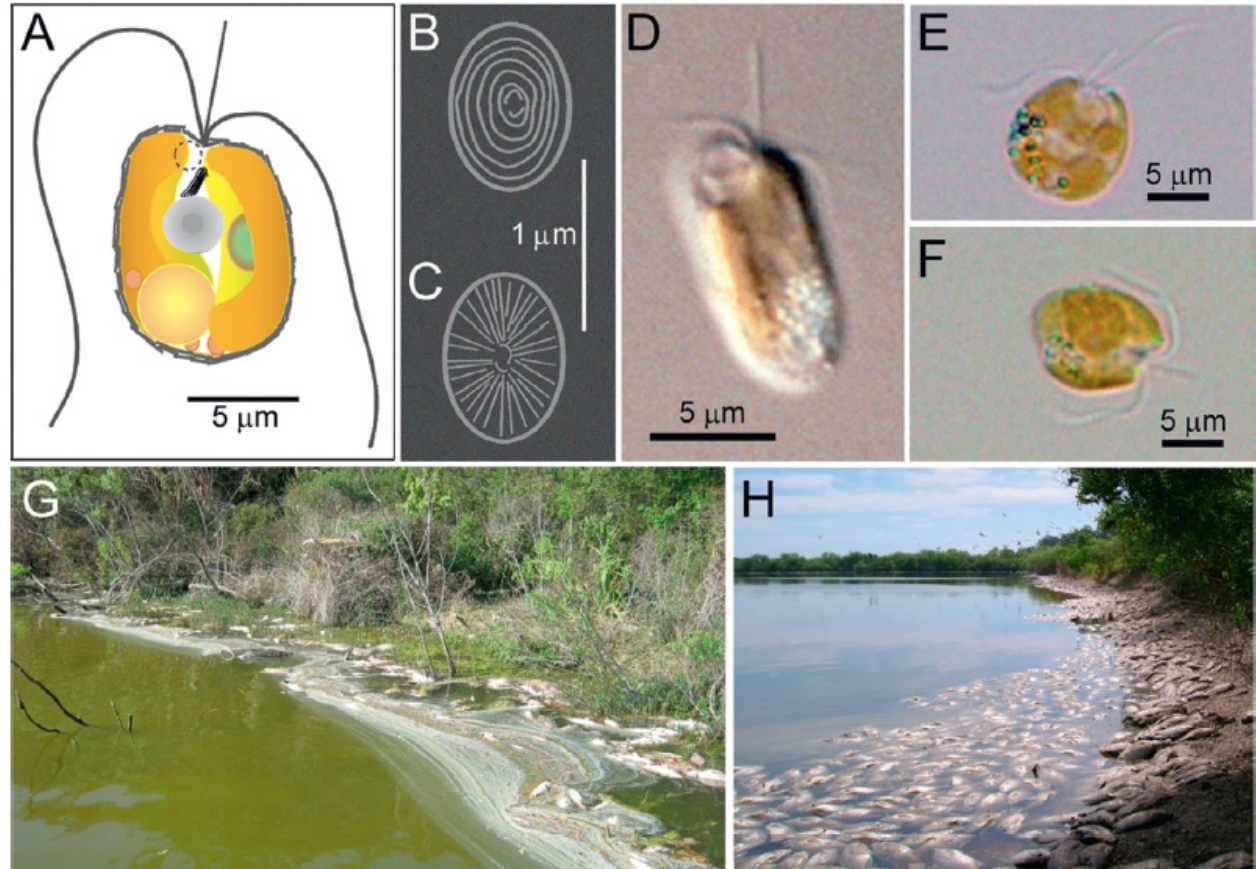
Prymnesiales

- very few species in Prymnesiales use silica to mineralize their scales
- production in SDV (silica deposition vesicles)
- small pores, deposited on cells in several layers
- *Prymnesium parvum* cysts are composed of layers of scales with siliceous material



Prymnesiales

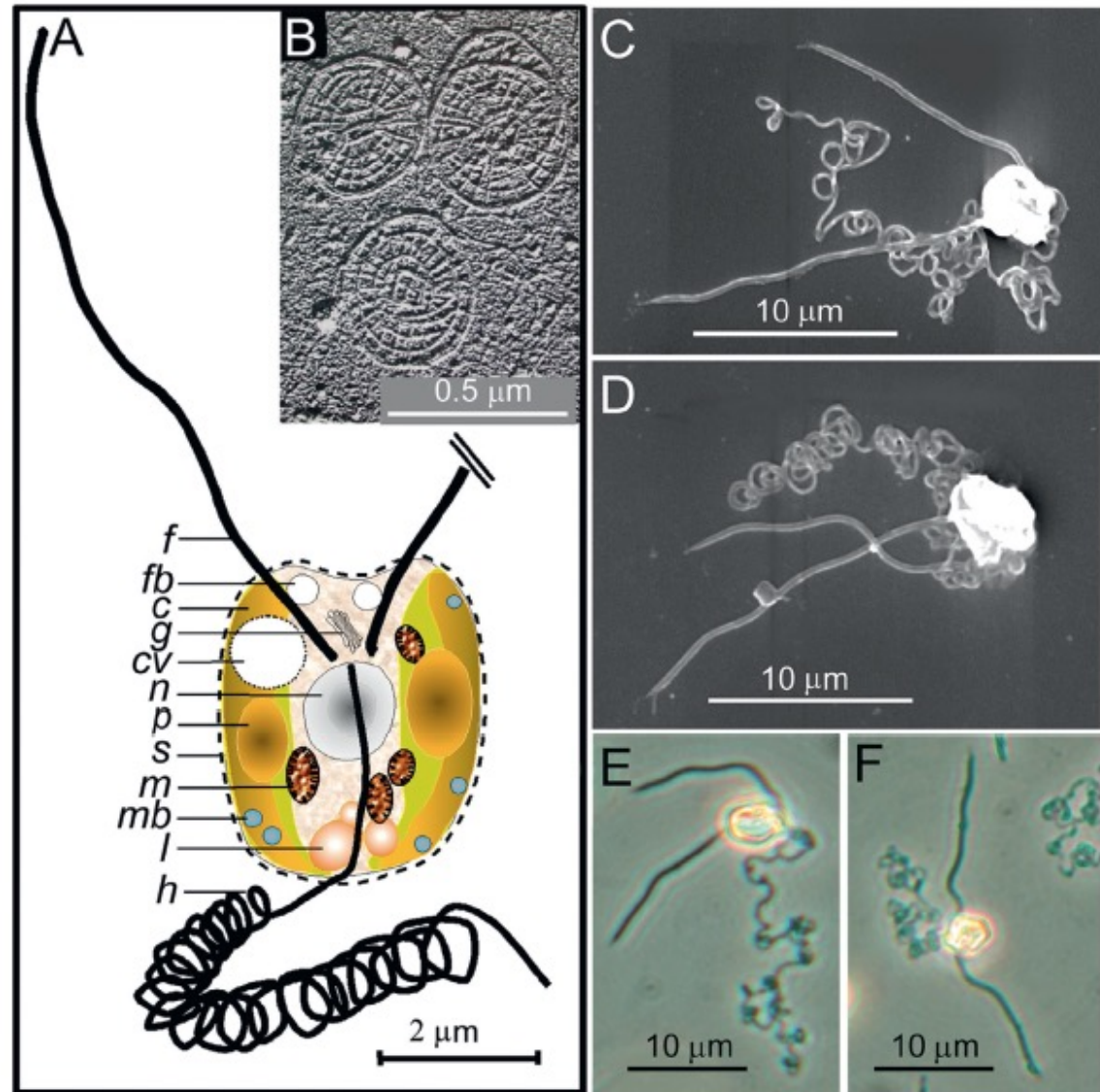
- *Prymnesium parvum*, mixotrophic, flagellates with organic scales
- toxic episodes in coastal area and freshwaters (toxins prymnesins, massive fish kills - ichthyotoxin)
- salinity appears to have been the primary factor controlling distribution of this species



Prymnesiales

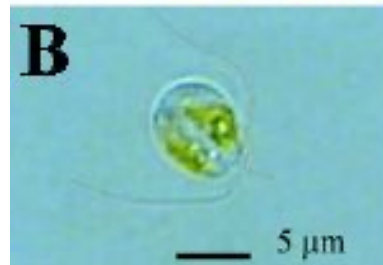
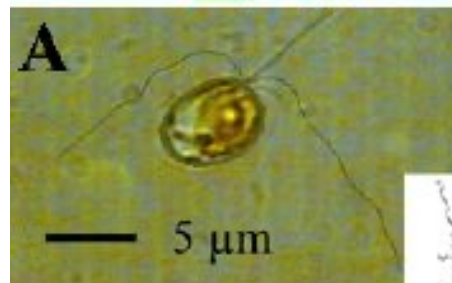
- *Chrysochromulina parva*, only organic scales
- widely distributed in freshwaters
- from eutrophic small ponds and rivers to large, deep, and cold lakes

f - flagellum; **fb** - lipid globule; **c** - chloroplast; **g** - Golgi body; **cv** - contractile vacuole; **n** - nucleus; **p** - pyrenoid; **s** - scale layer; **m** - mitochondrion; **mb** - muciferous body; **l** - leucosin vesicle; **h** - haptonema



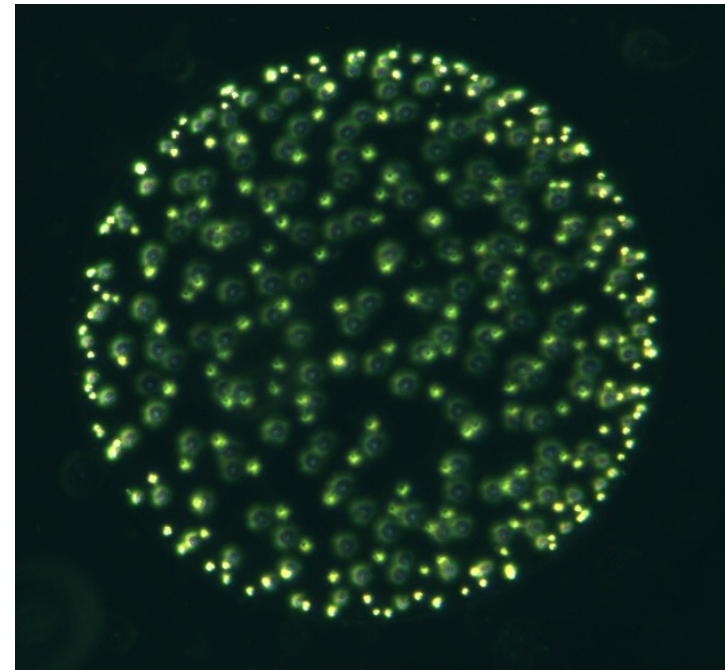
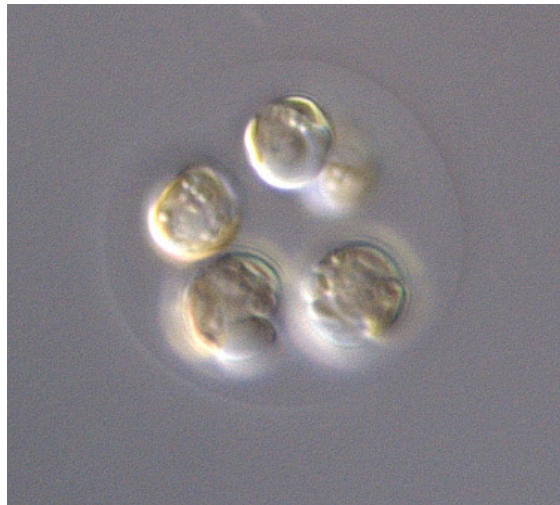
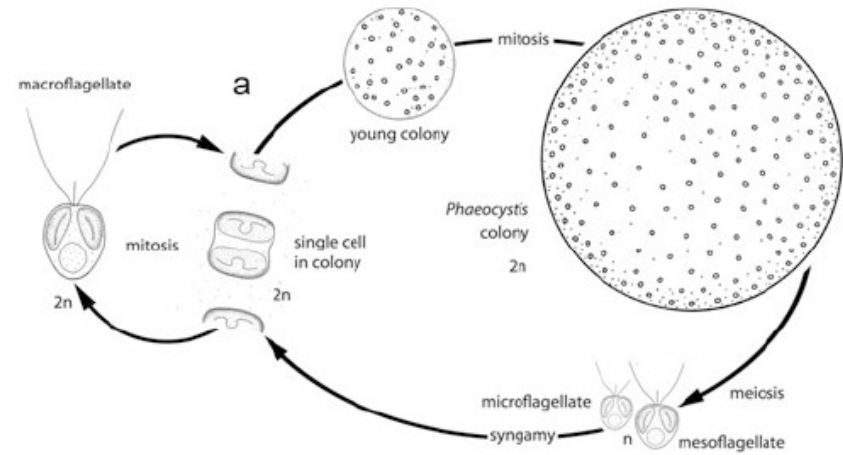
Prymnesiales

- *Chrysochromulina polylepis*
- production of ichthyotoxins



Phaeocystales

- *Phaeocystis pouchetii*: flagellates (3-8 μm) in large (up to a few mm or cm) mucilaginous colonies (pallmeloid stage)



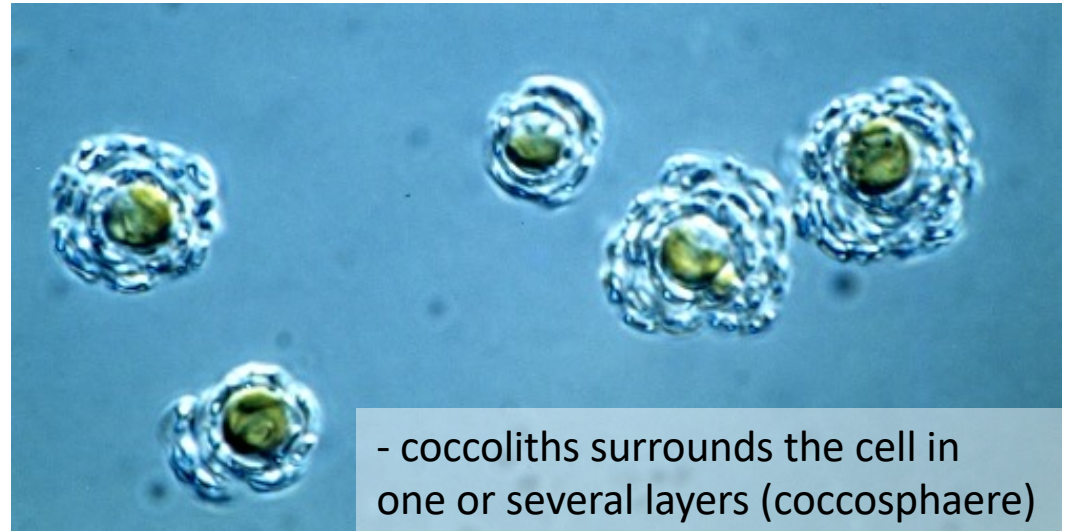
Phaeocystales

- *Phaeocystis* nuisance and ?toxic bloom, foam accumulated on beaches

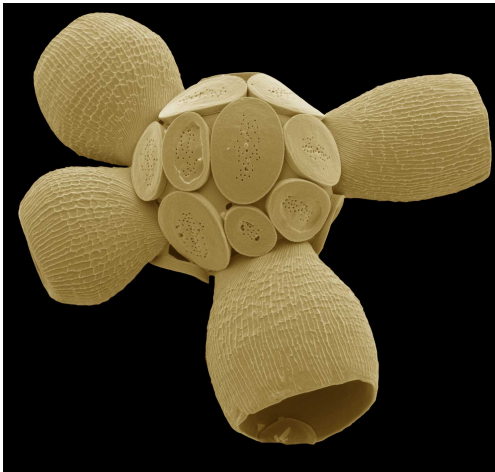


coccolithophores

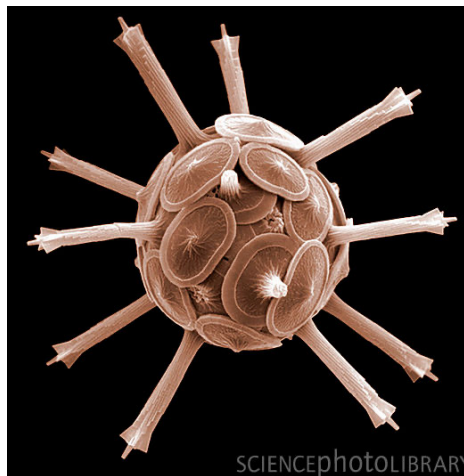
- coccolith-bearing haptophytes
- *kokkos* = berry, grain
lithos = stone (Huxley 1858)
- coccoliths especially in
Isochrysidales and **Coccolithales**



Scyphosphaera



Rhabdosphaera clavigera

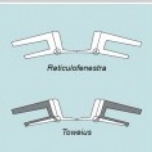
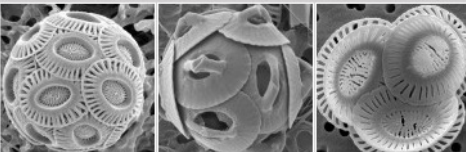
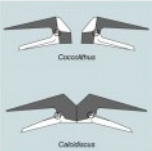
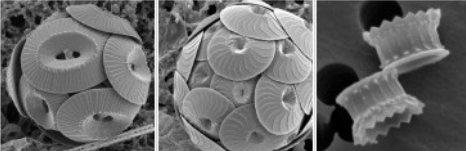
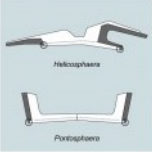
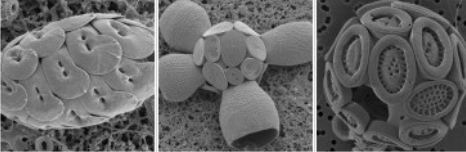
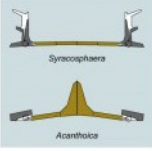
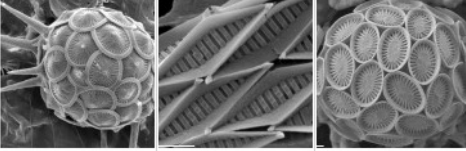
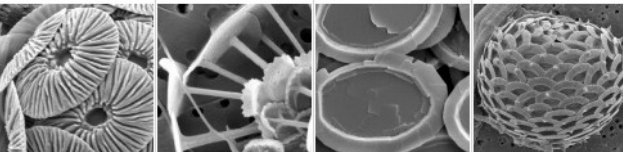


Syracosphaera anthos



- a huge variety of ornamented calcified scales

Coccolithophores

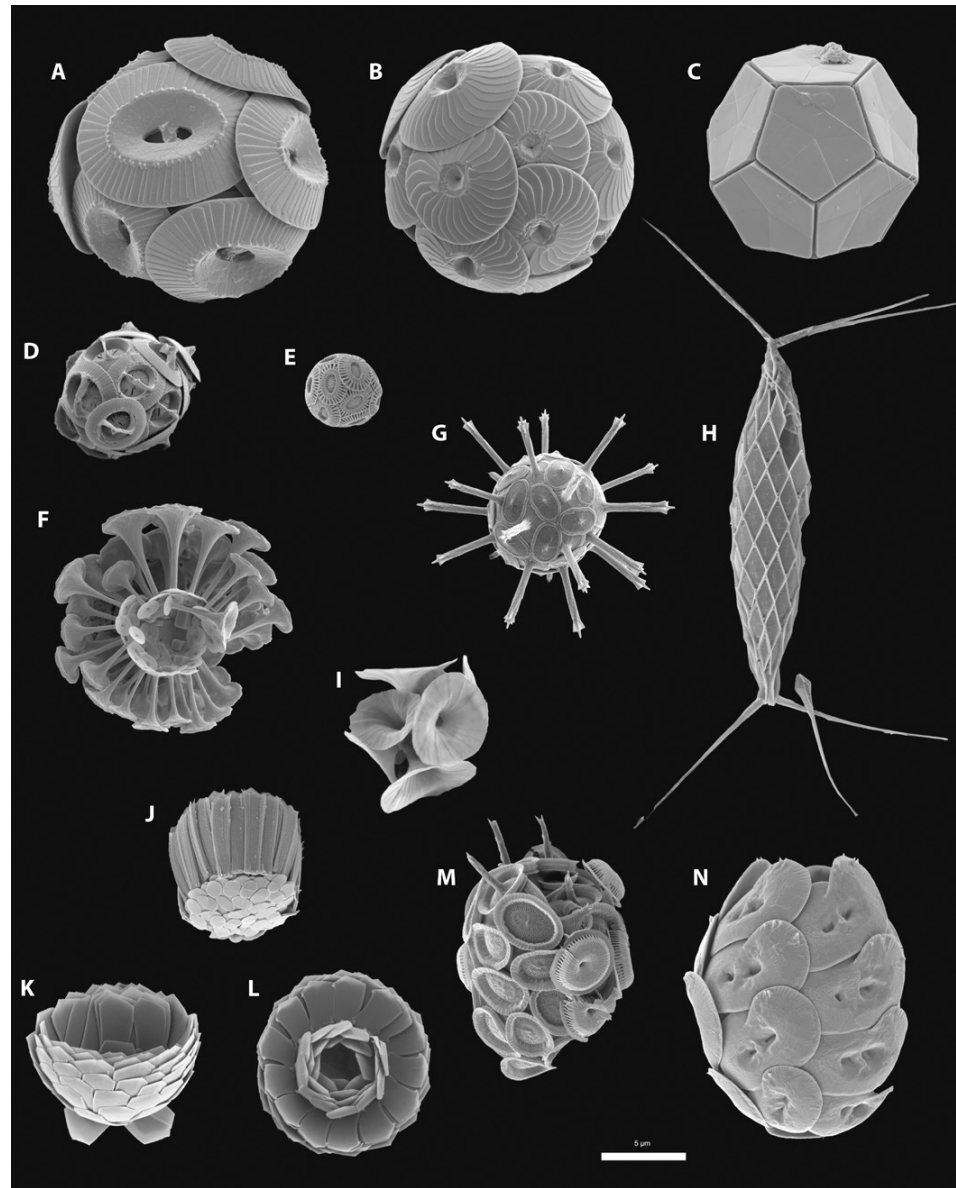
Daughter taxa: (blue => in age window 0-800Ma)				Granddaughter taxa	
Heterococcoliths					
		ISOCHRYSIDALES Heterococcoliths mostly placoliths with R-unit dominant. Motile phases with vestigial haptonema.		NOELAERHABDACEAE PRINSIACEAE ISOCHRYSIDACEAE	
		COCCOLITHALES Mostly placolith heterococcoliths, with V-unit forming the distal shield; R-unit the proximal shield.		COCCOLITHACEAE CALCIDISCACEAE HYMENOMONADACEAE PLEUROCHRYSIDACEAE	
		ZYGODISCALES Heterococcoliths with V-units forming upper/outer cycle of imbricated elements and R-units forming basal plate and central mass of irregular elements		HELICOSPHAERACEAE PONTOSPHAERACEAE ZYGODISCACEAE	
		SYRACOSPHAERALES Coccoliths with radial lath cycle of T-units, and disjunct, often complex, axial structure, coccospheres often polymorphic		CALCIOSOLENIACEAE RHABDOSPHAERACEAE SYRACOSPHAERACEAE	
		COCCOLITH FAMILIES INC SED Various heterococcolith groups not obviously related to the well-defined orders		ALISPHAERACEAE PAPPOSPHAERACEAE UMBELLOSPHAERACEAE NARROW RIMMED PLACOLITHS COCCOLITH GENERA INC SED UNIDENTIFIED	

Cenozoic and Modern Coccolithophores

<http://ina.tmsoc.org/Nannotax3/index.php?dir=Coccolithophores&top=34&base=300>

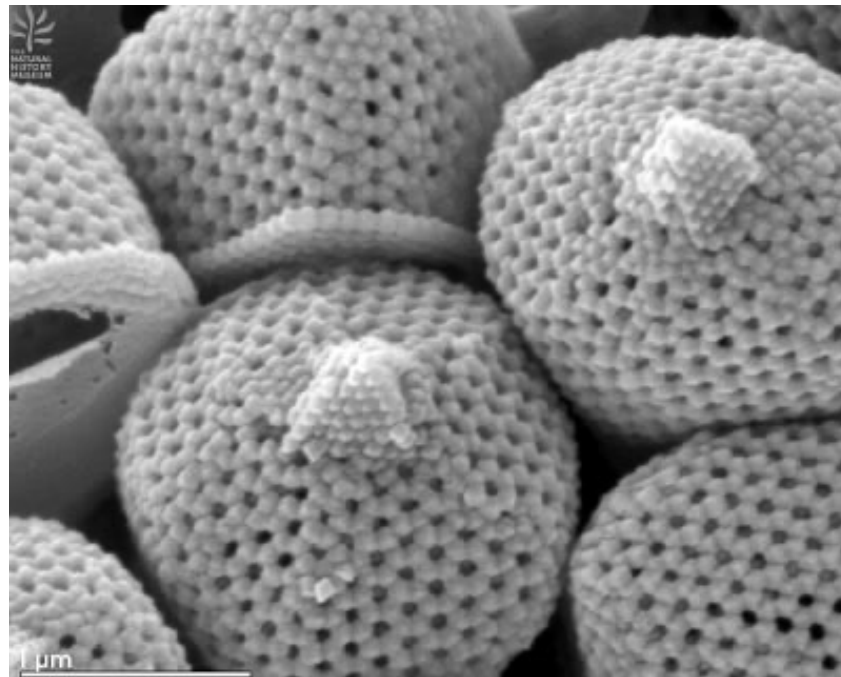
Coccolithophores

(A) *Coccolithus pelagicus*, (B) *Calcidiscus leptoporus*, (C) *Braarudosphaera bigelowii*, (D) *Gephyrocapsa oceanica*, (E) *E. huxleyi*, (F) *Discosphaera tubifera*, (G) *Rhabdosphaera clavigera*, (H) *Calciosolenia murrayi*, (I) *Umbellosphaera irregularis*, (J) *Gladiolithus flabellatus*, (K and L) *Florisphaera profunda*, (M) *Syracosphaera pulchra*, and (N) *Helicosphaera carteri*.
Scale bar, 5 μm .

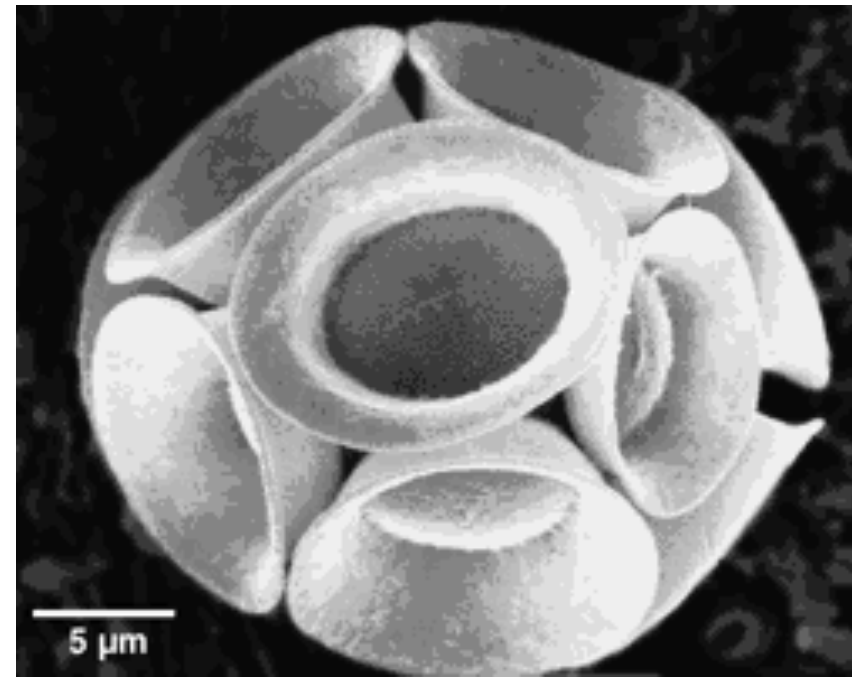


Coccolithophores

Holococcoliths: haploid cells;
individual crystals have simple
morphologies, no distinction between
the rim and central area

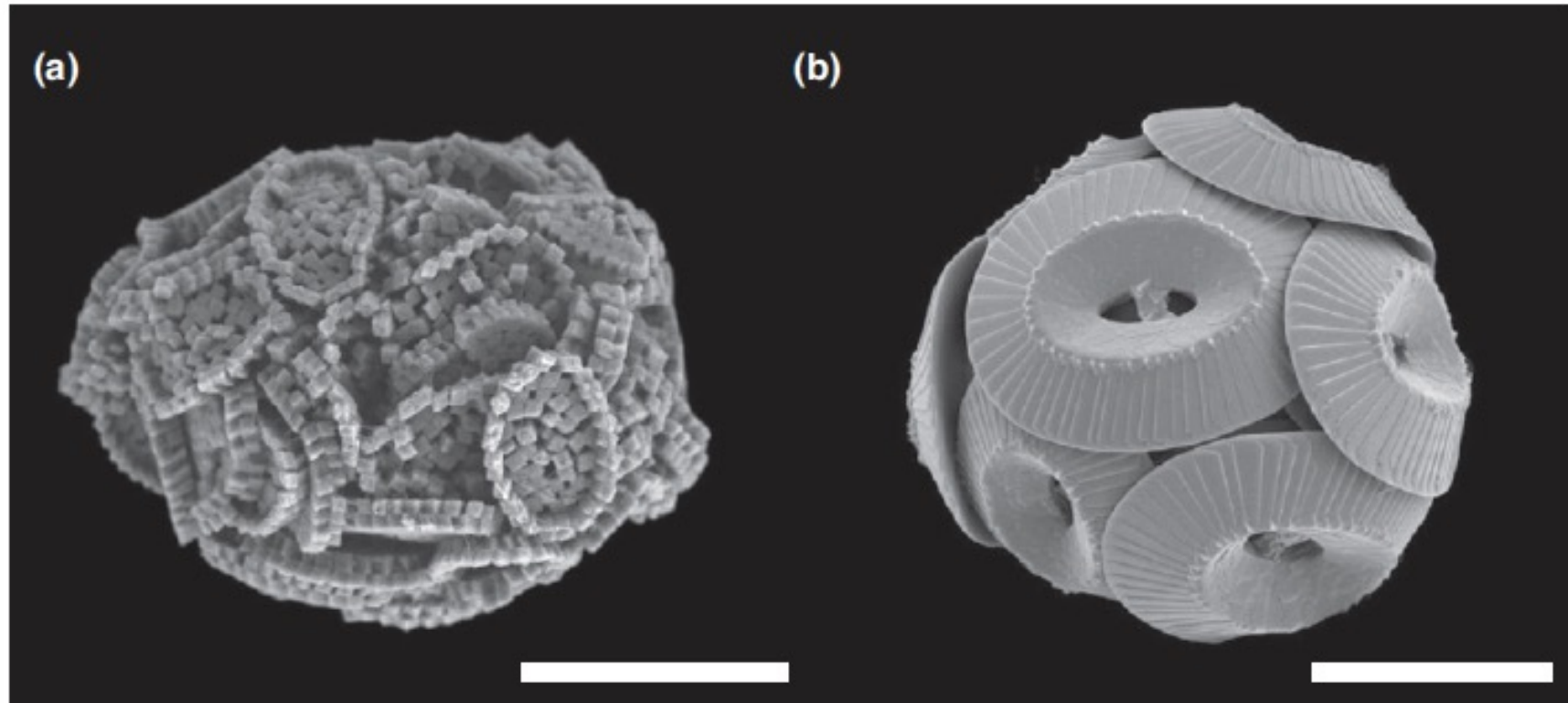


Heterococcoliths: diploid cells;
complex morphologies, a rim of
radial crystal units surrounding the
central area



Pontosphaera syracusana

Coccolithophores

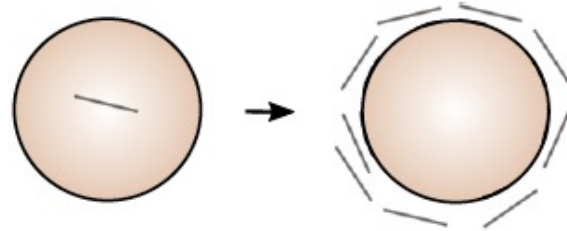


Coccolithus braarudii

Holococcoliths are comprised of identical rhombic crystals, whereas heterococcoliths are comprised of a variety of crystal sizes and shapes.

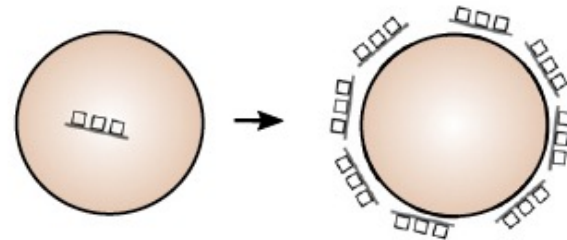
Coccolithophores

1. Noncalcified ancestor



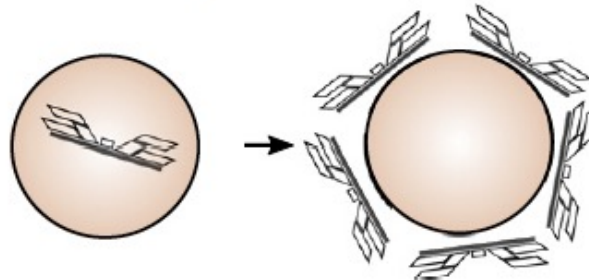
- Organic scales formed intracellularly

2. Emergence of calcified ancestor



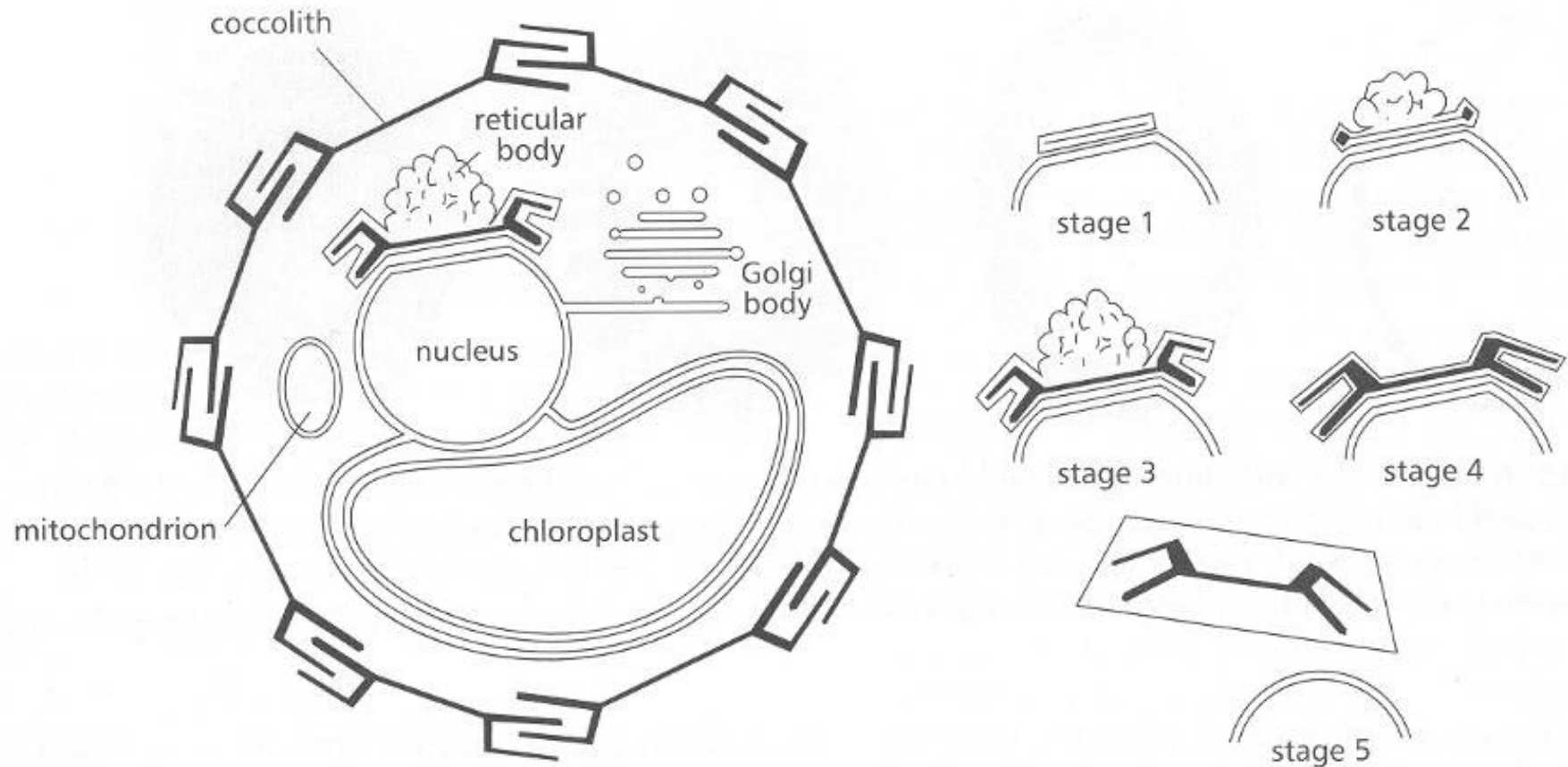
- Intracellular formation of calcite on organic scales
- Simple rhombic crystal morphology

3. Development of heterococcoliths



- Intracellular formation of calcite on specialised organic baseplate
- Recruitment of Si-dependent mechanism allows complex crystal morphology

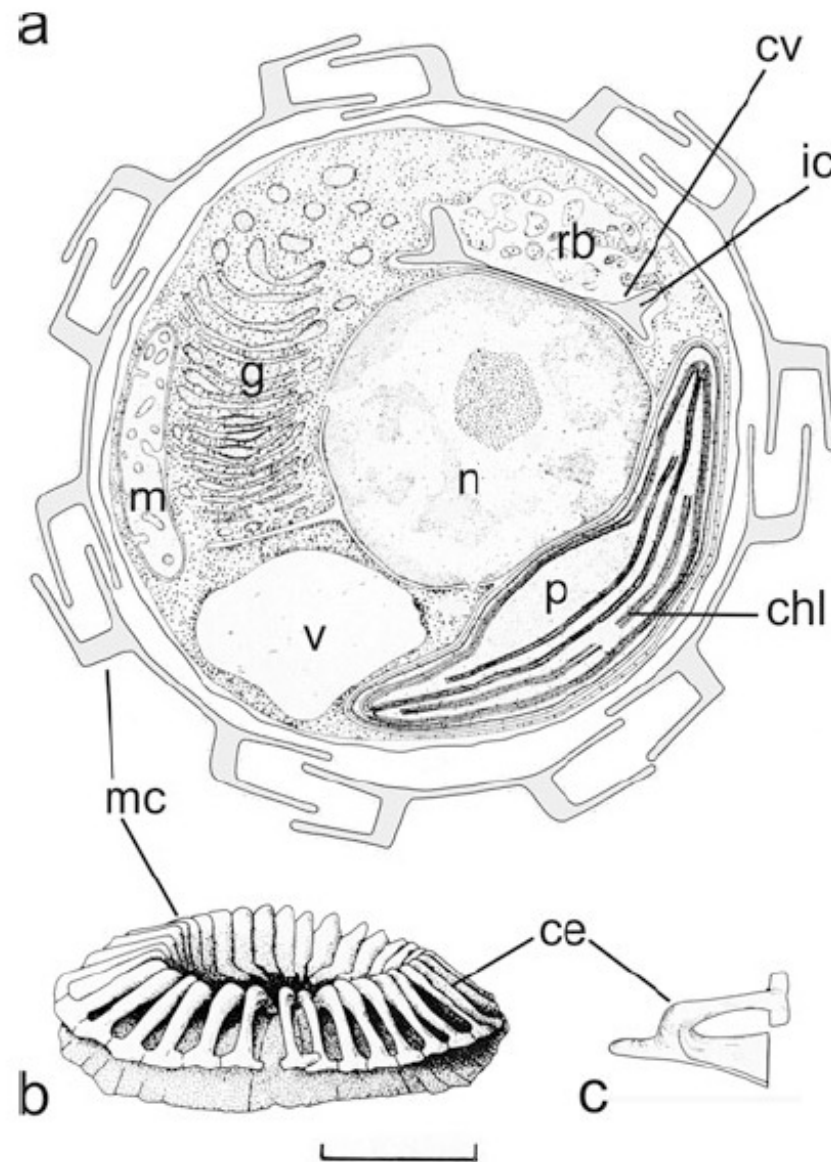
Coccolithophores



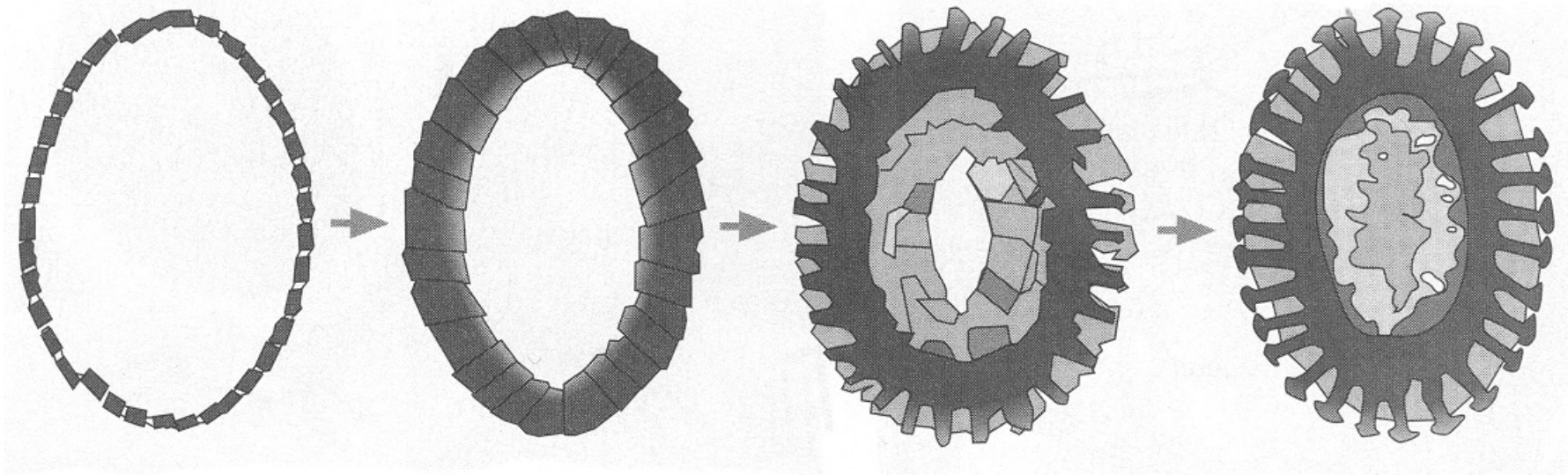
Biosynthesis of heterococcoliths: (1) Golgi vesicles fuse to form one big vesicle attached to the nuclear membrane, (2) thin organic lamella is formed inside the vesicle, reticular body (microtubular net) is attached outside, (3) calcification, (4) disappearing of reticular body, (5) complete coccolith is transferred to the cell surface and released outside.

Coccolithophores

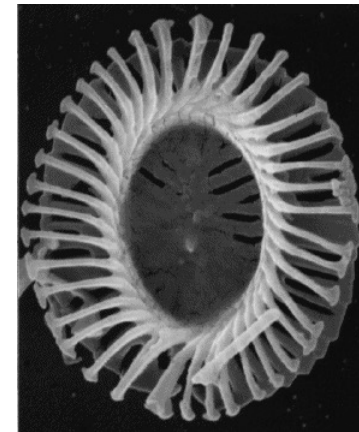
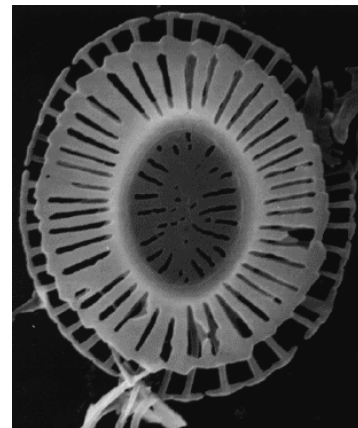
Schematic drawing *Emiliania huxleyi*: (a) transverse section of whole cell showing the chloroplast (chl), coccolith vesicle (cv), Golgi body (g), immature coccolith (ic), mature coccolith (mc), mitochondrion (m), nucleus (n), pyrenoid (p), reticulate body (rb), vacuole (v), (b) single mature coccolith showing elements arranged in an oval ring, (c) calcite element (ce) of a coccolith. Scalebar 1 μm



Coccolithophores

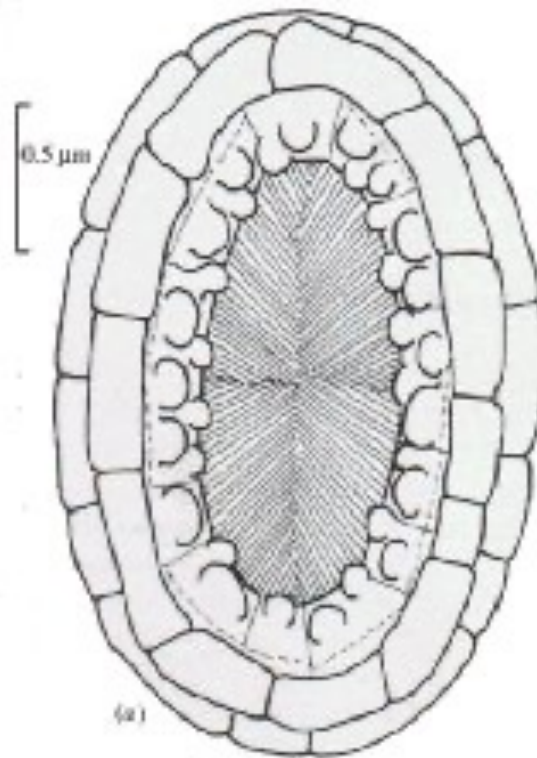


Biosynthesis of heterococcoliths
(*Emiliana huxleyi*)



Coccolithophores

- calcified scales are usually formed in addition to underlying organic scales

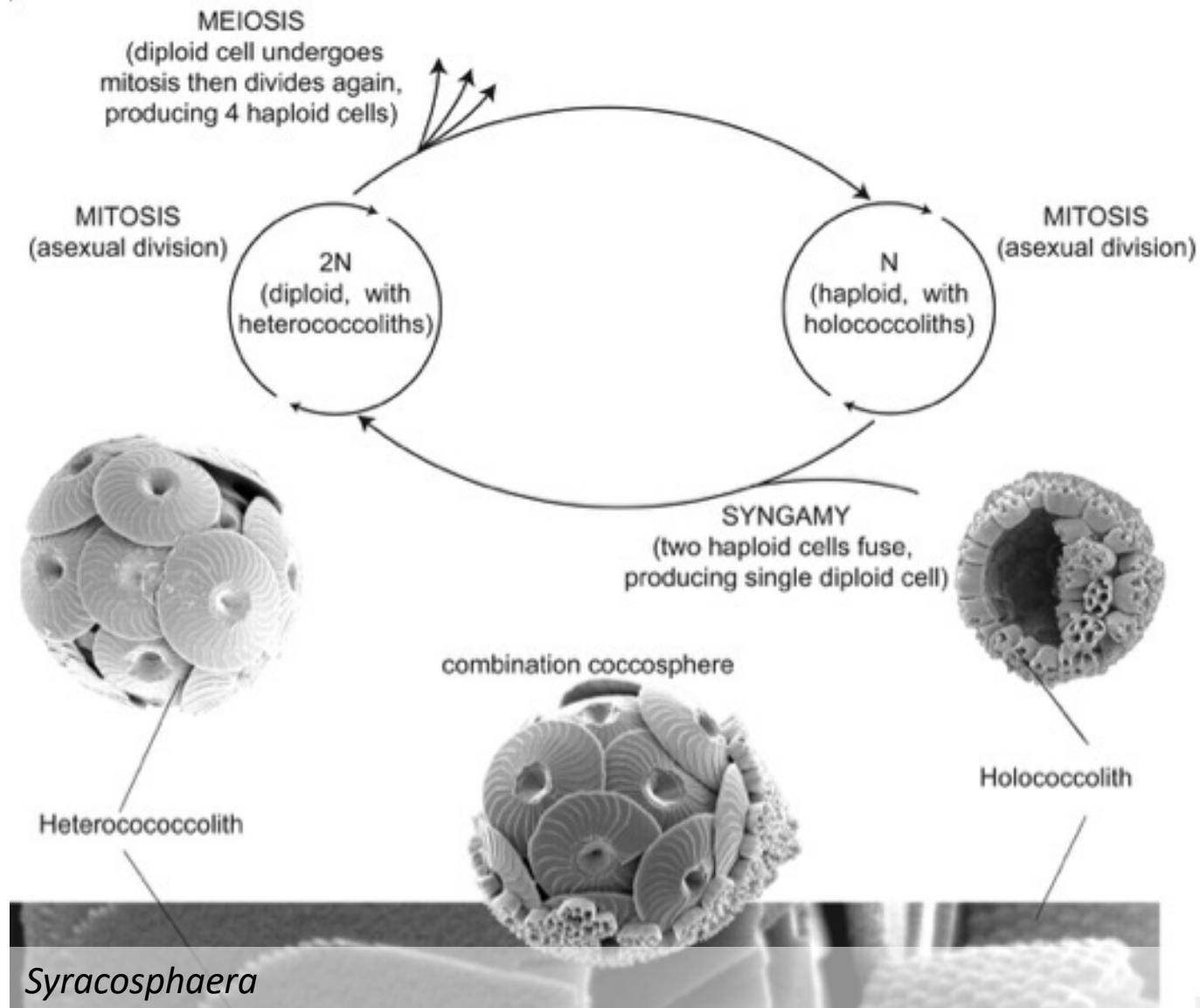


two calcite layers surround
the polysaccharide scale



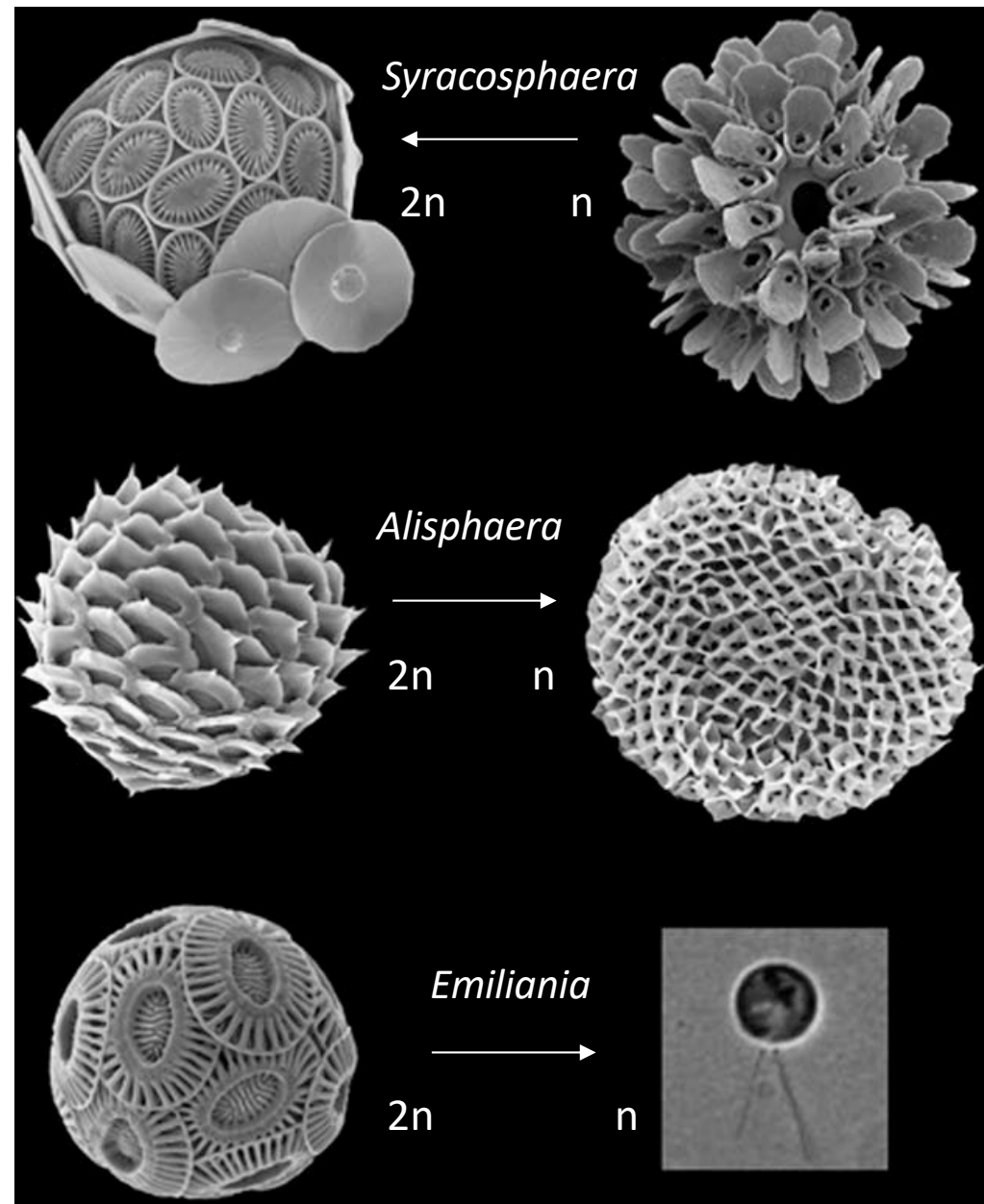
unmineralised scales are
composed of microfibrils

Coccolithophores



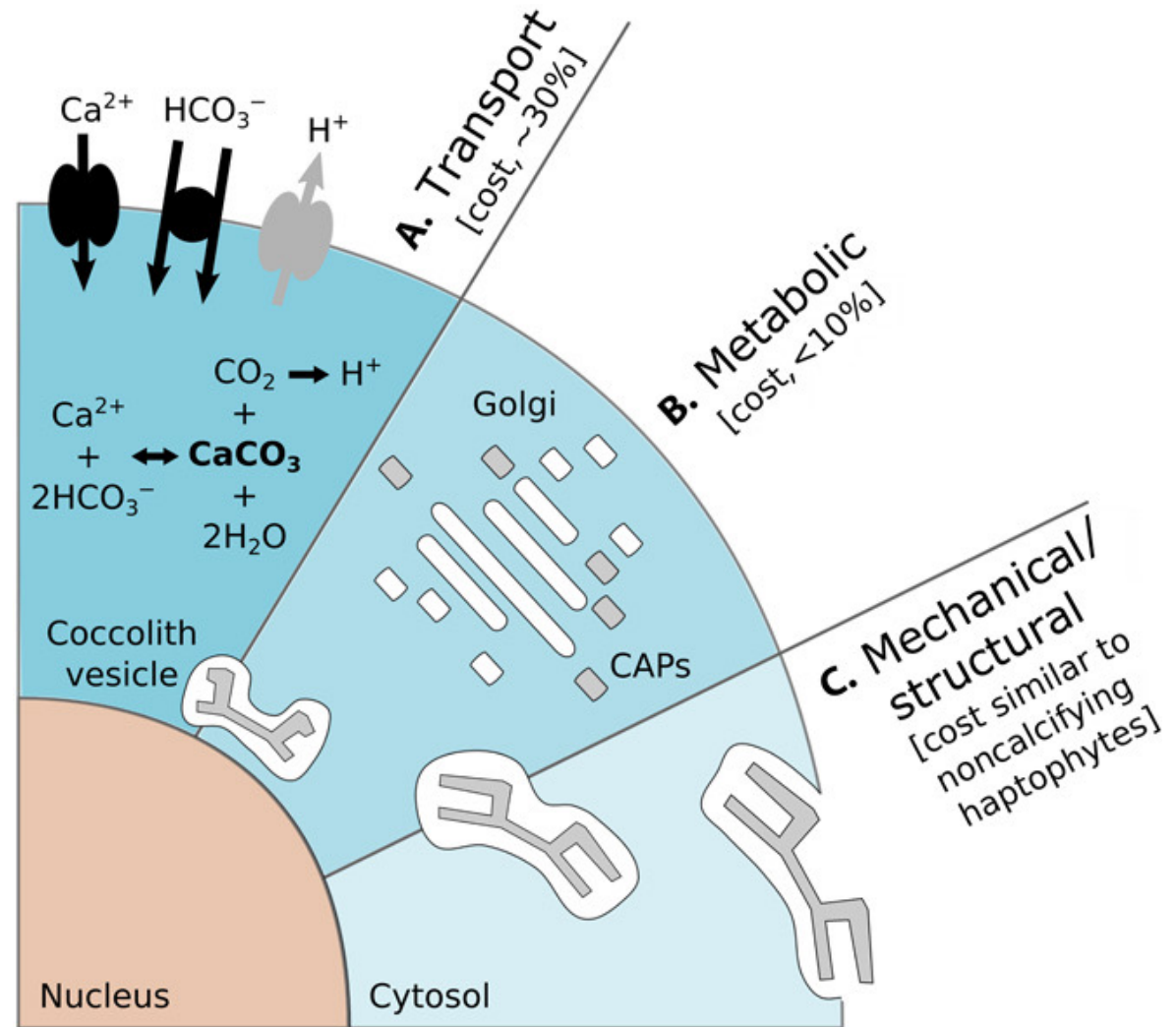
Coccolithophores

- holococcoliths and heterococcoliths occur on alternate phases of the life-cycle of single species
- it reflects a haplodiplontic life-cycle, with holococcoliths consistently occurring in the haploid phase
- nomenclature problems: a single species was occasionally described as separate species

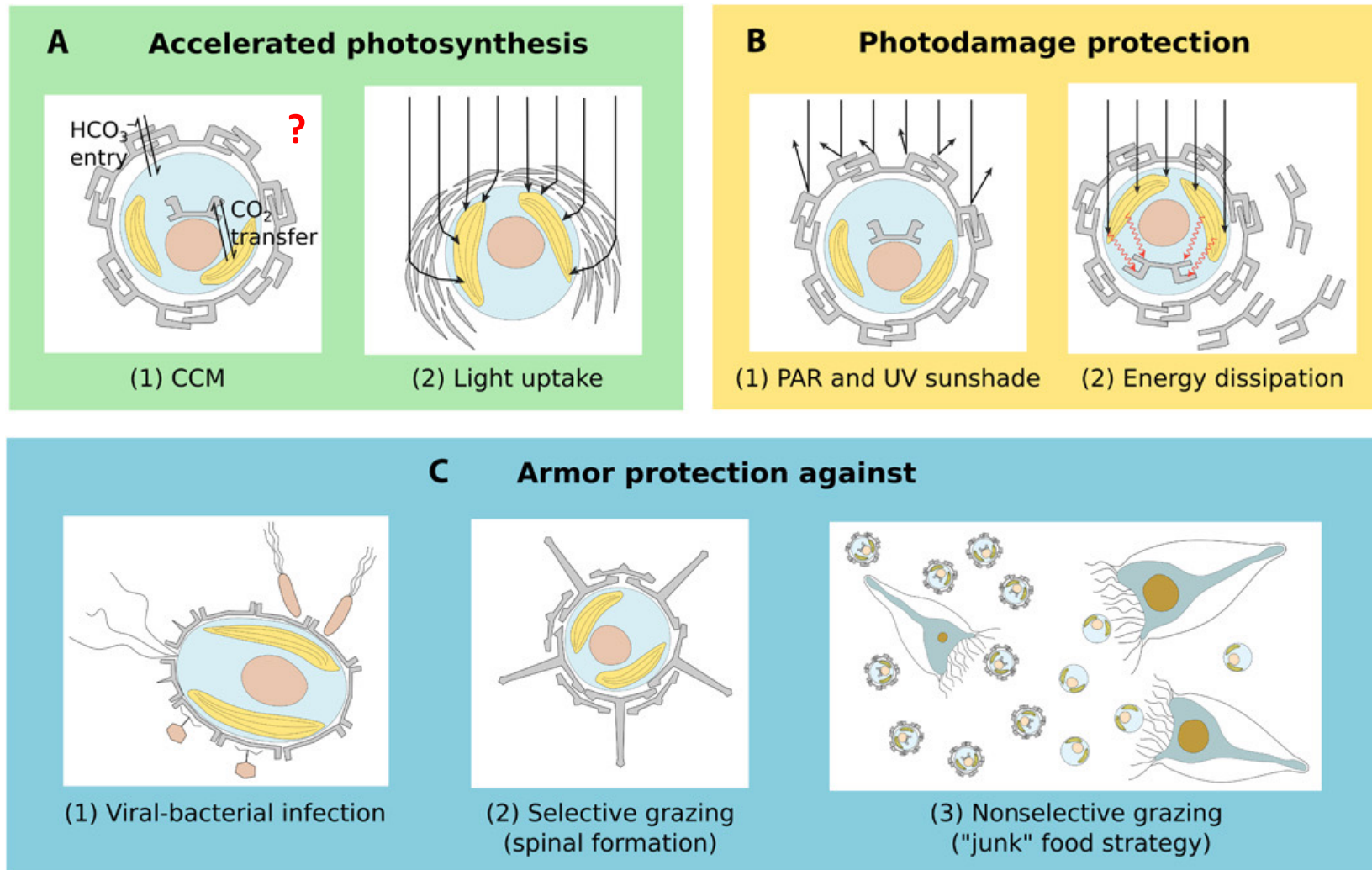


Coccolithophores

Schematic of the cellular processes associated with calcification and the approximate energetic costs of a coccolithophore cell.



Coccolithophores



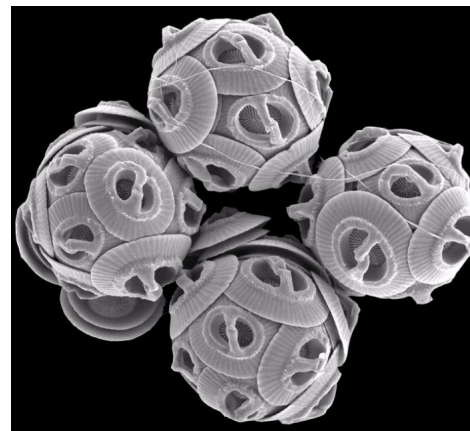
Proposed main benefits of calcification in coccolithophores.

Coccolithophores

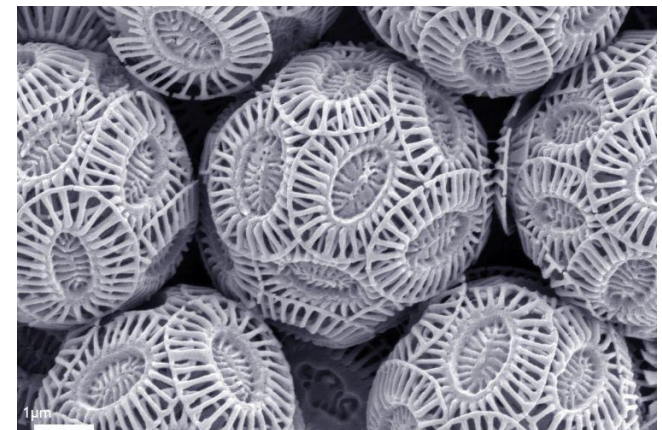
- global carbon cycle: fixing dissolved inorganic carbon (DIC) through photosynthesis and calcification
- sequester atmospheric CO₂ as limestone (Ca₂CO₃)
- white tides: blooms that frequently cover >100,000 km² in temperate and boreal regions



Satellite image of a bloom in the English Channel



Gephyrocapsa oceanica



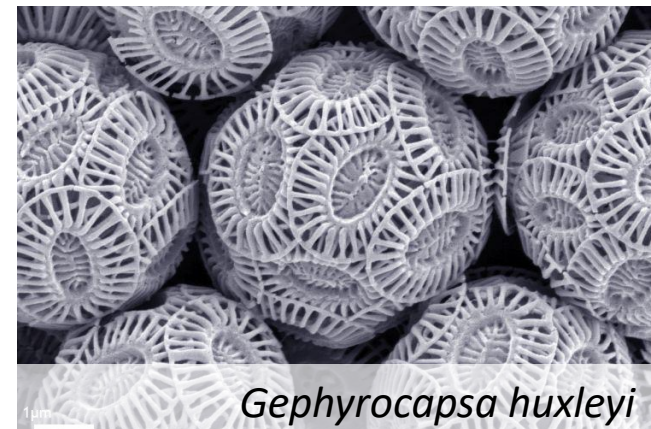
Gephyrocapsa huxleyi

Coccolithophores

- *E. huxleyi*: photo- and phagotrophy, multilayers and monomorphic coccoliths; no trace of haptonema
- tend to bloom in the stratified surface layers in the Earth's oceans (not inhibited by high irradiances)
- white tide blooms follow seasonal succession cycles: after diatom bloom (low nutrient and silicate concentrations)
- *E. huxleyi*: tolerates nitrogen deprivation much better than diatoms; able to substitute Fe-containing enzymes
- mature blooms are typically lysed by specific viruses (phyco-dna-viruses)

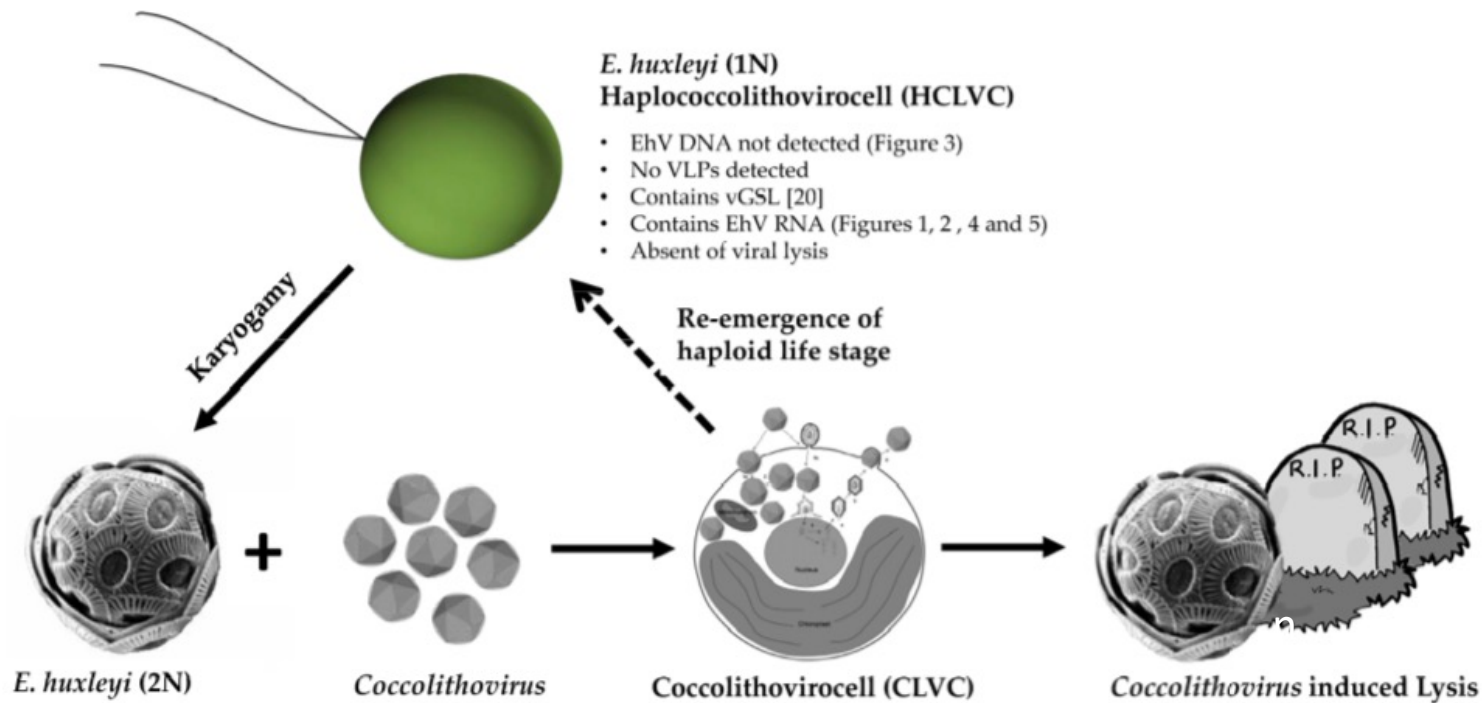


- blooms change optical properties: coccoliths or entire coccospheres are shed and scatter (by the CaCO_3 crystals) the incoming light (turning surface waters turbid and milky)

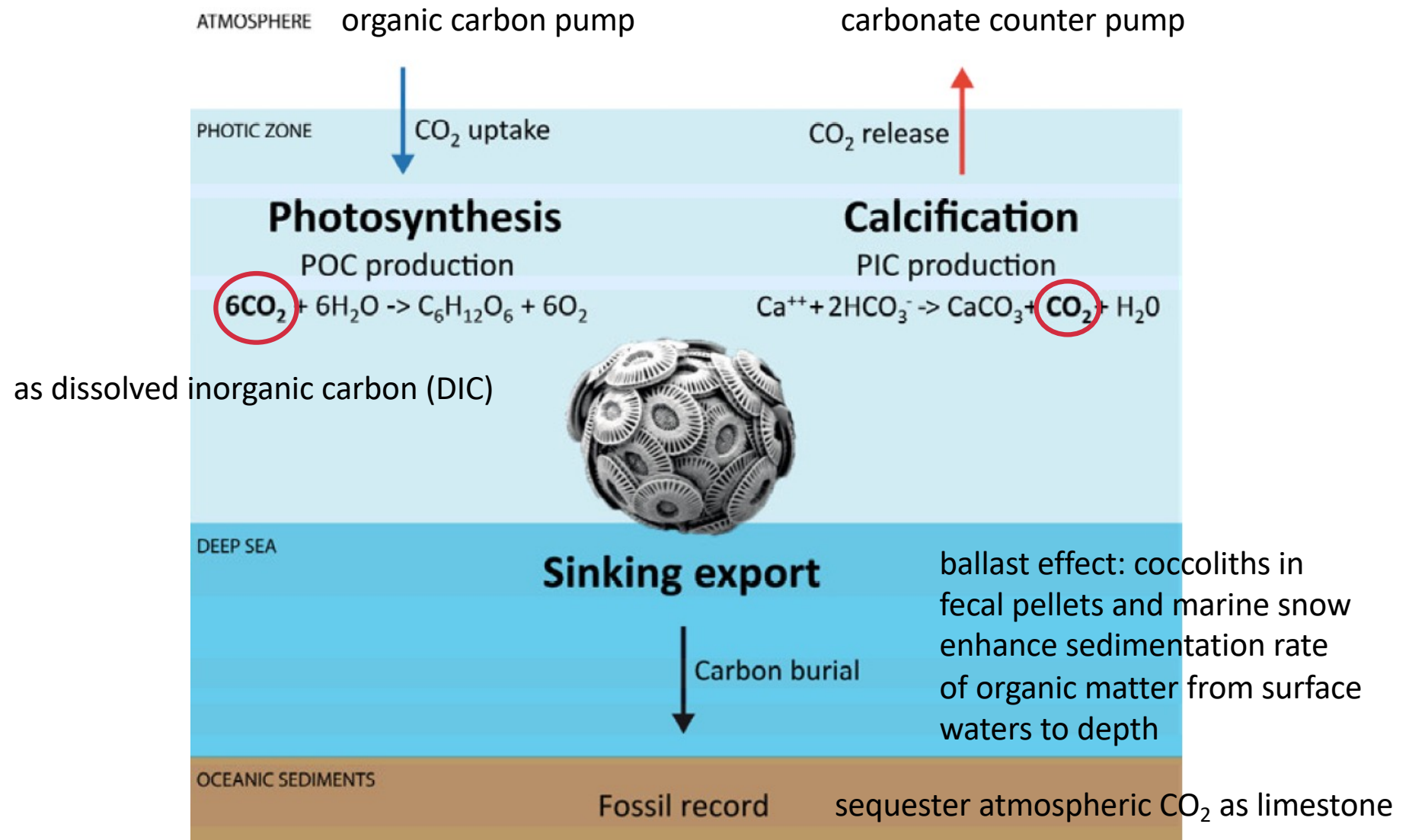


Coccolithophores

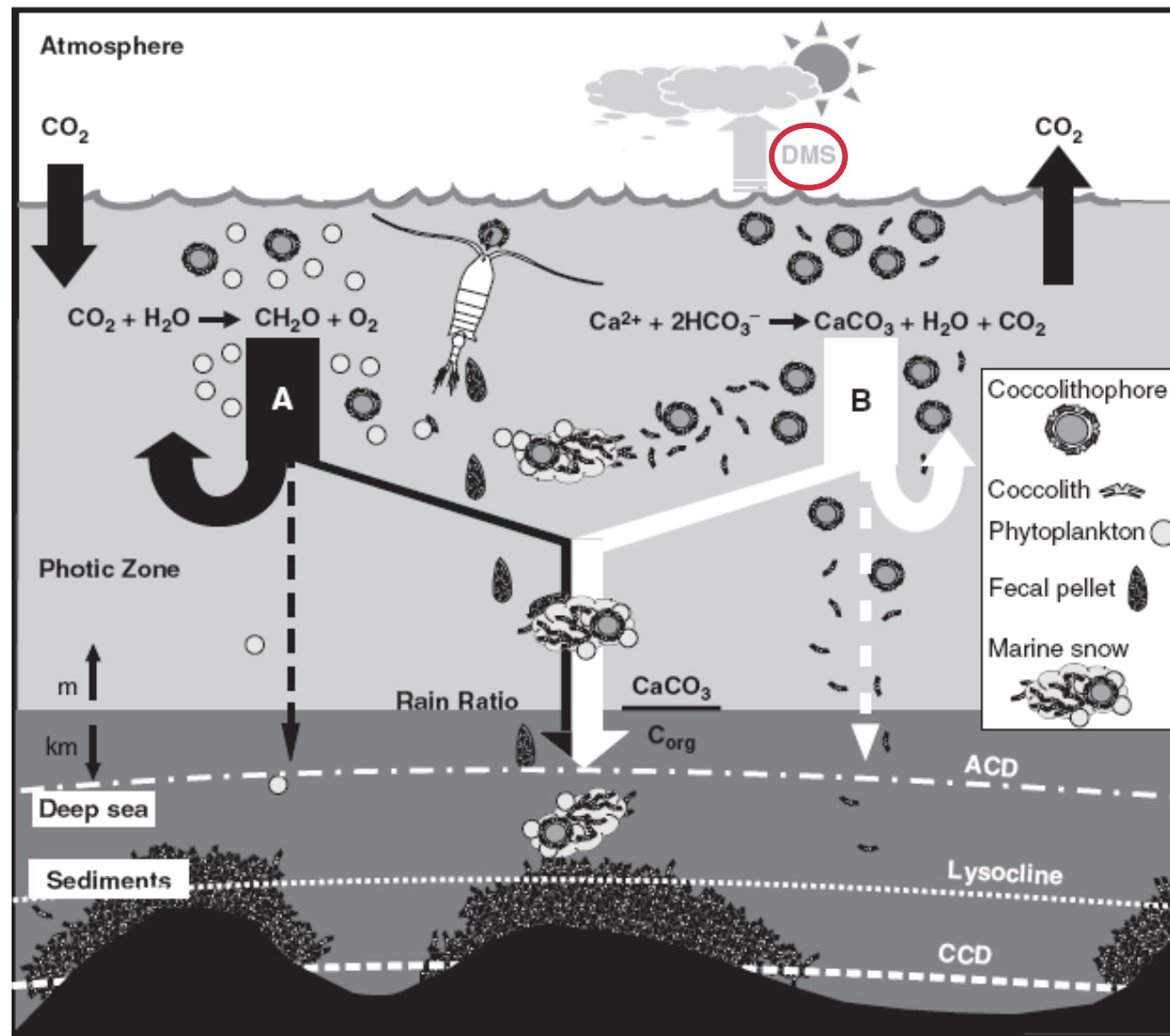
- *Emiliana huxleyi*: resistance to viral attack
- Cheshire cat escape strategy: haploid cells are thought to resist infection -> population as a new founding population
- Schrödinger's Cheshire Cat: the presence of viral RNA in the haploid cells -> an alternative stage of the virus life cycle, no evidence of viral cell lysis



Coccolithophores



Coccolithophores



DMS(P)

- marine phytoplankton: producers of dimethylsulfoniopropionate (DMSP)
- the most likely function of DMSP is related to the osmotic protection of the cell + antioxidant function + lysis of DMSP may be an effective chemical defense against grazing
- DMSP: degraded into dimethyl sulfide (DMS)
- DMS oxidation products act as (CCN) cloud condensation nuclei (atmospheric cloud formation)

TABLE 2. Intracellular concentration of DMSP in marine phytoplankton

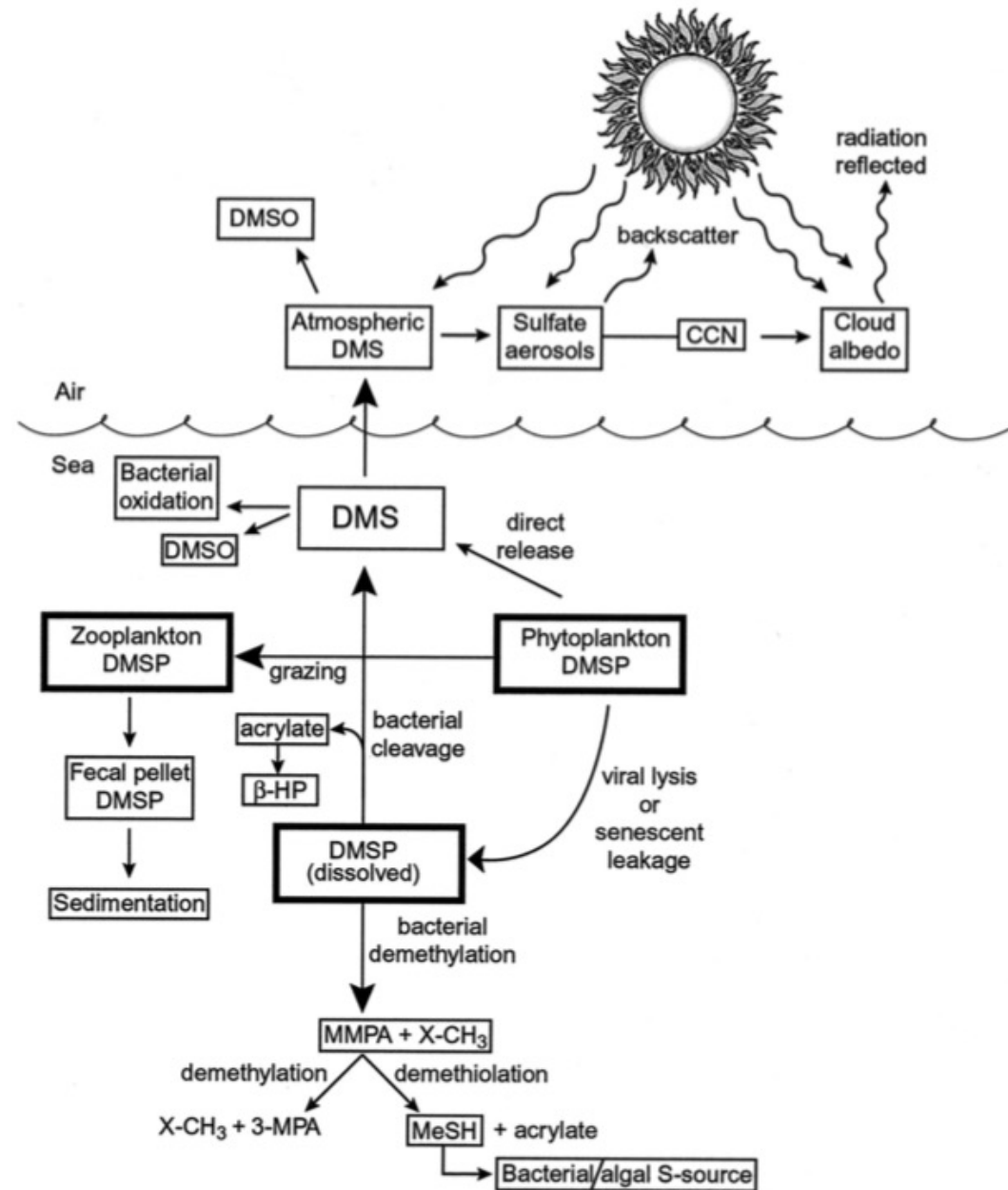
Organism	Cytosolic DMSP ^a (mM)	Reference
<i>Hymenomonas carterae</i>	120	Vairavamurthey et al. (120)
<i>Gymnodinium nelsoni</i>	280	Dacey and Wakeham (30)
<i>Platymonas subcordiformis</i>	170	Dickson and Kirst (40)
<i>Phaeocystis</i> sp.	71–169	Stefels and van Boekel (109)
<i>Melosira numuloides</i> (Diatoms)	264	Keller et al. (55)
<i>Chrysamoeba</i> sp. (chrysophyte)	596	
<i>Ochromonas</i> sp. (chrysophyte)	529	
<i>Prorocentrum</i> sp. strain IIB ₂ b ₁ (dinoflagellate)	1,082 ^b	
<i>Emiliania huxleyi</i> BT6 (coccolithophore)	166	
Diatoms (from sea water)	38 ± 18	Turner et al. (119)
<10-μm flagellates (from sea water)	125	Belviso et al. (15)
	650 ± 429	Turner et al. (119)
Dinoflagellates (from sea water)	640	Belviso et al. (15)
	70 ± 30	Turner et al. (119)

^a Per liter of cell volume.

^b A reviewer of this article pointed out that “seawater osmolarity is about 0.7 M, and DMSP is never the only osmolyte, so its intracellular concentrations can’t be >1 M.”

DMS(P)

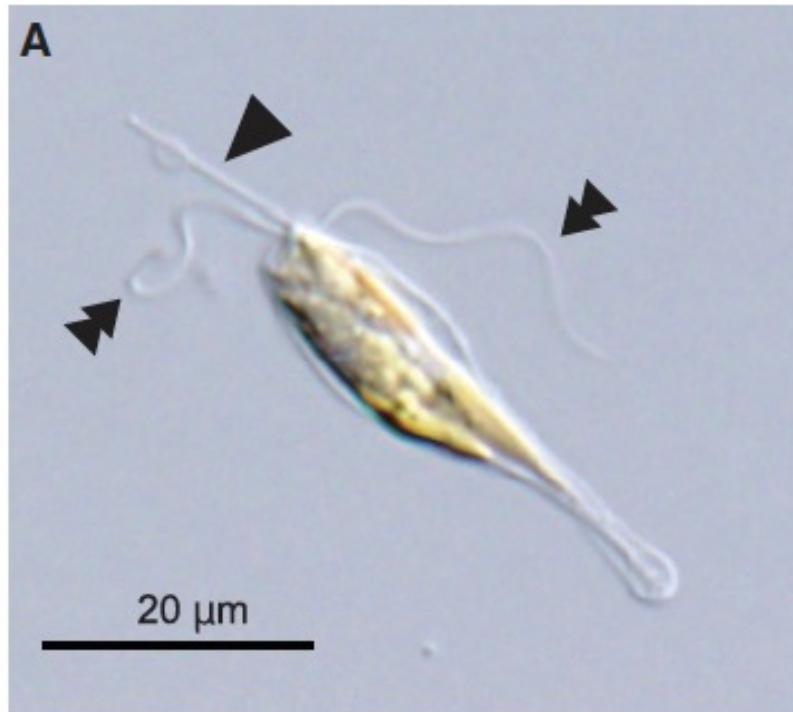
- DMSP cleavage to DMS plays an important biogeochemical role in the global sea-to-land transfer of sulfur
- it has a potential effect on remote oceanic weather patterns due to cloud formation -> reflection of solar radiation
- white tides: increasing of planetary albedo (i.e., increased light reflection) -> cooling down the Earth



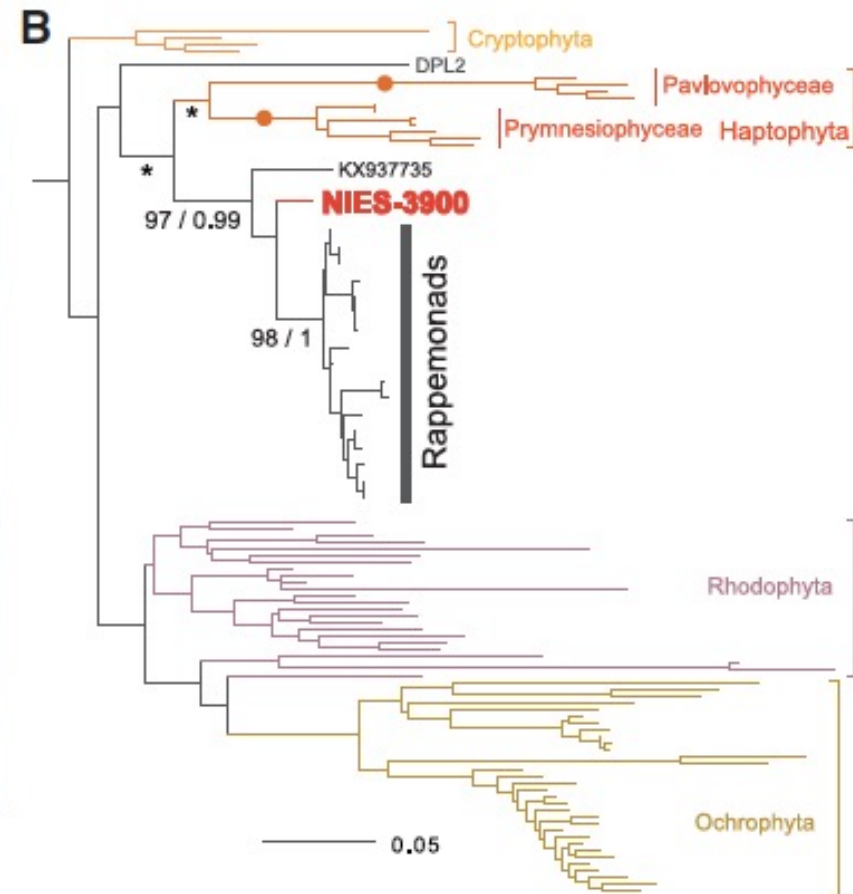
Haptophyta

- feeding of copepods, shellfish, aquaculture of bivalves, crustaceans, and fish
- nuisance or toxic blooms: serious economic losses
- paleolimnology: past climatic conditions, sedimentary rocks and ocean sediments composed of coccoliths
- climatic feedbacks? may affect planetary albedo and concentration of CO₂

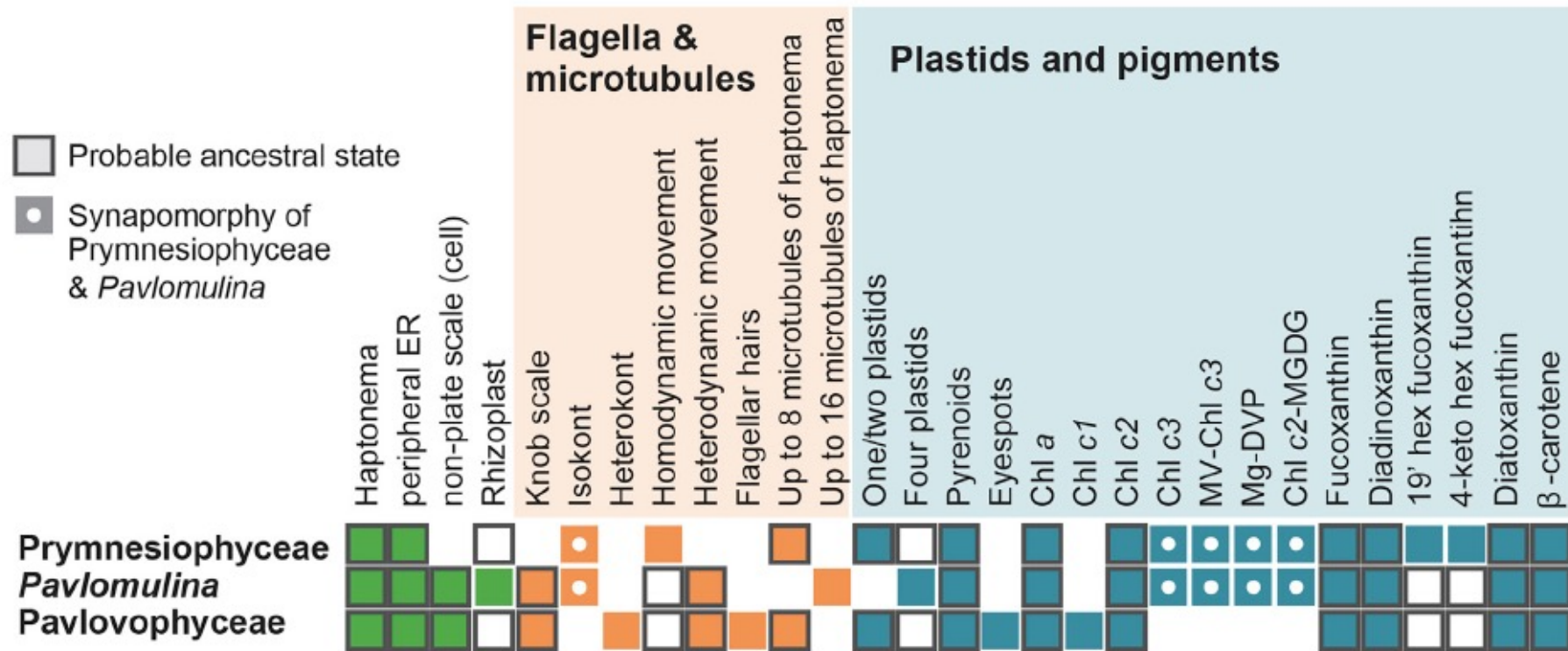
+ class Rappephyceae



NIES-3900 = *Pavlomulina ranunculiformis*



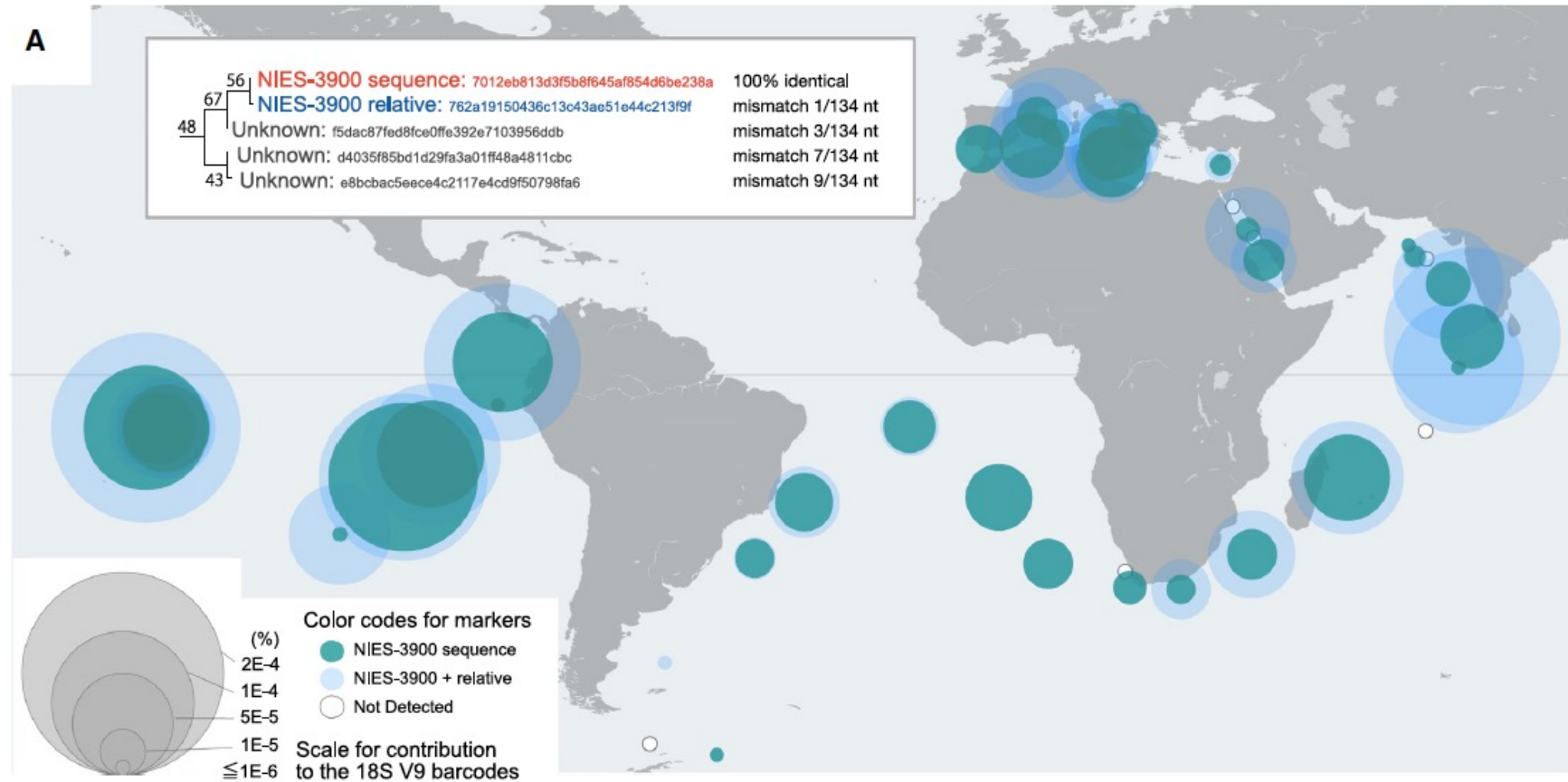
Rappephycea



-> class: Rappephyceae

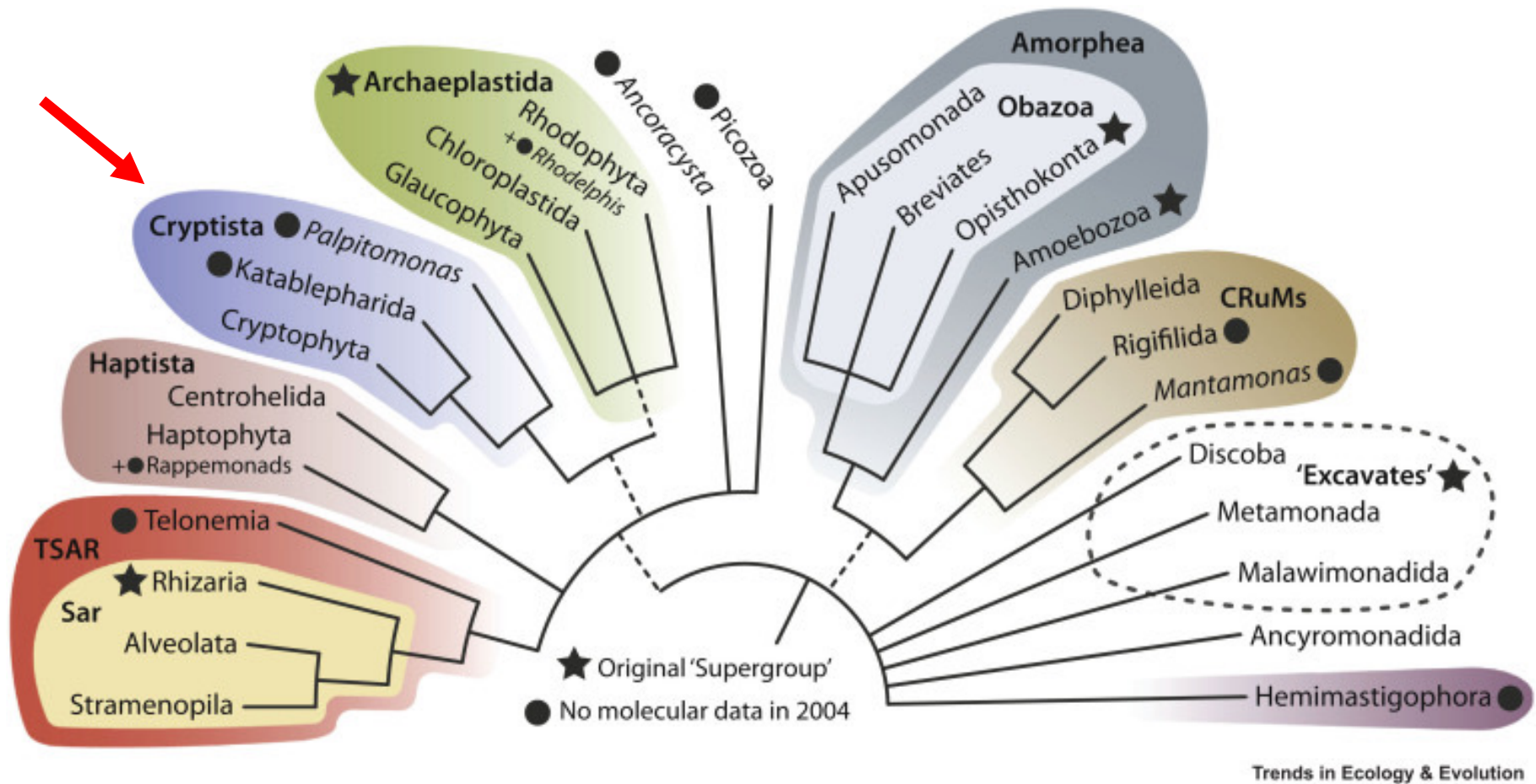
Rappephycea

NIES-3900 = *Pavlomulina ranunculiformis*

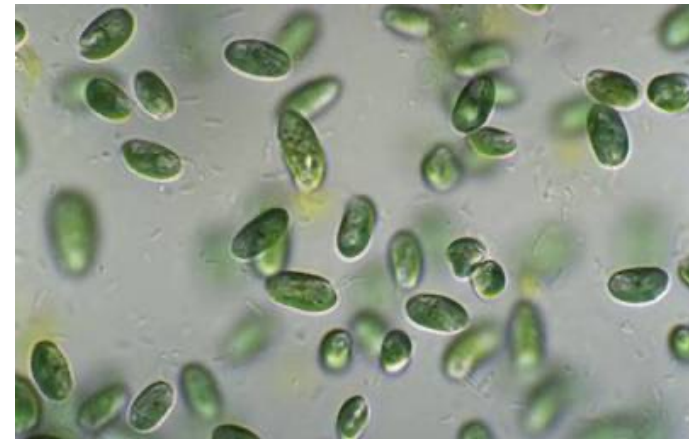
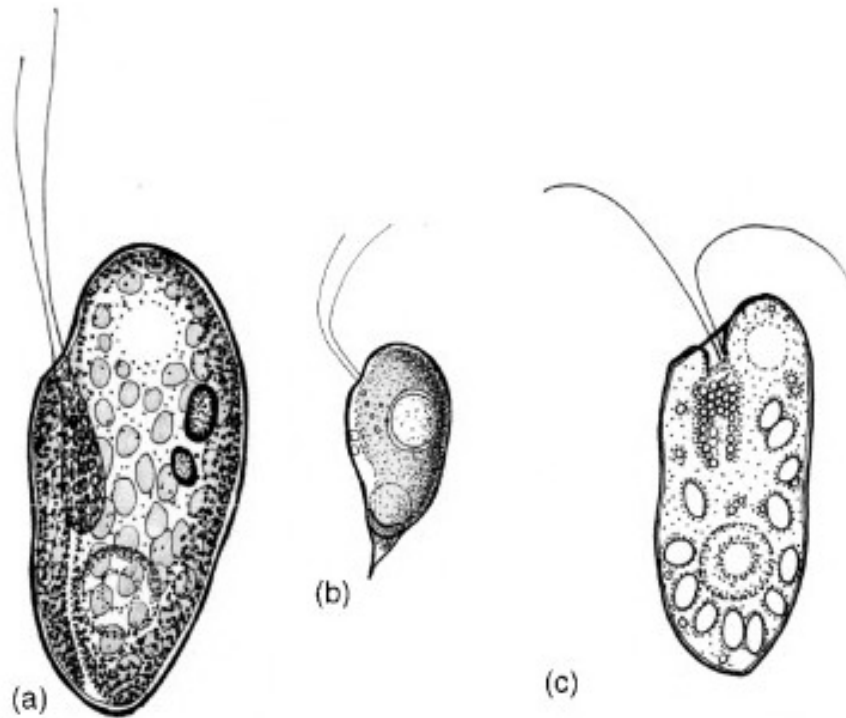


Rappephyceae have a cosmopolitan distribution, and an abundance comparable to or greater than the two previously known haptophyte classes.

Cryptista



Cryptomonads



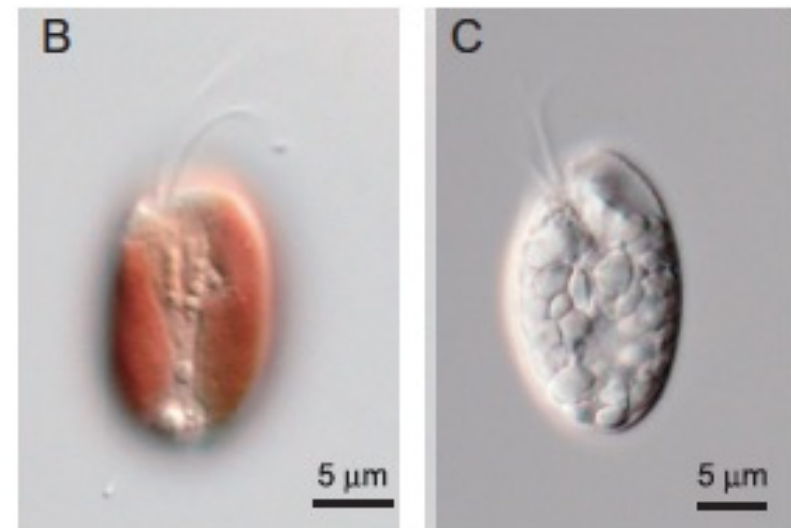
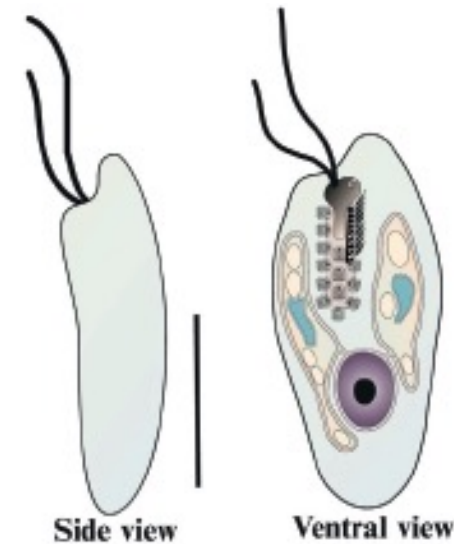
<https://m.youtube.com/watch?v=g6kXduiDrF0>

https://m.youtube.com/watch?v=iiMaUefa_3g

Cryptista – Cryptophyta - Cryptophyceae

Cryptophyta

- unicellular free-living flagellates
[palmeloid forms are also known]
- planktonic and photoautotrophic [may be mixotrophic, some exclusively heterotrophic with leucoplast], vertical migration, low-light specialists
- ubiquitous in the earth's hydrosphere: marine, brackish, and freshwater
- flattened asymmetric shape of cells, distinctive swimming motion (the cells rotate about its longitudinal axis), refractile ejectosomes (light microscopically visible)
- diverse coloration (chocolate brown, olive brown, bright brown, tomato red, dark orange, blue green, sky blue)

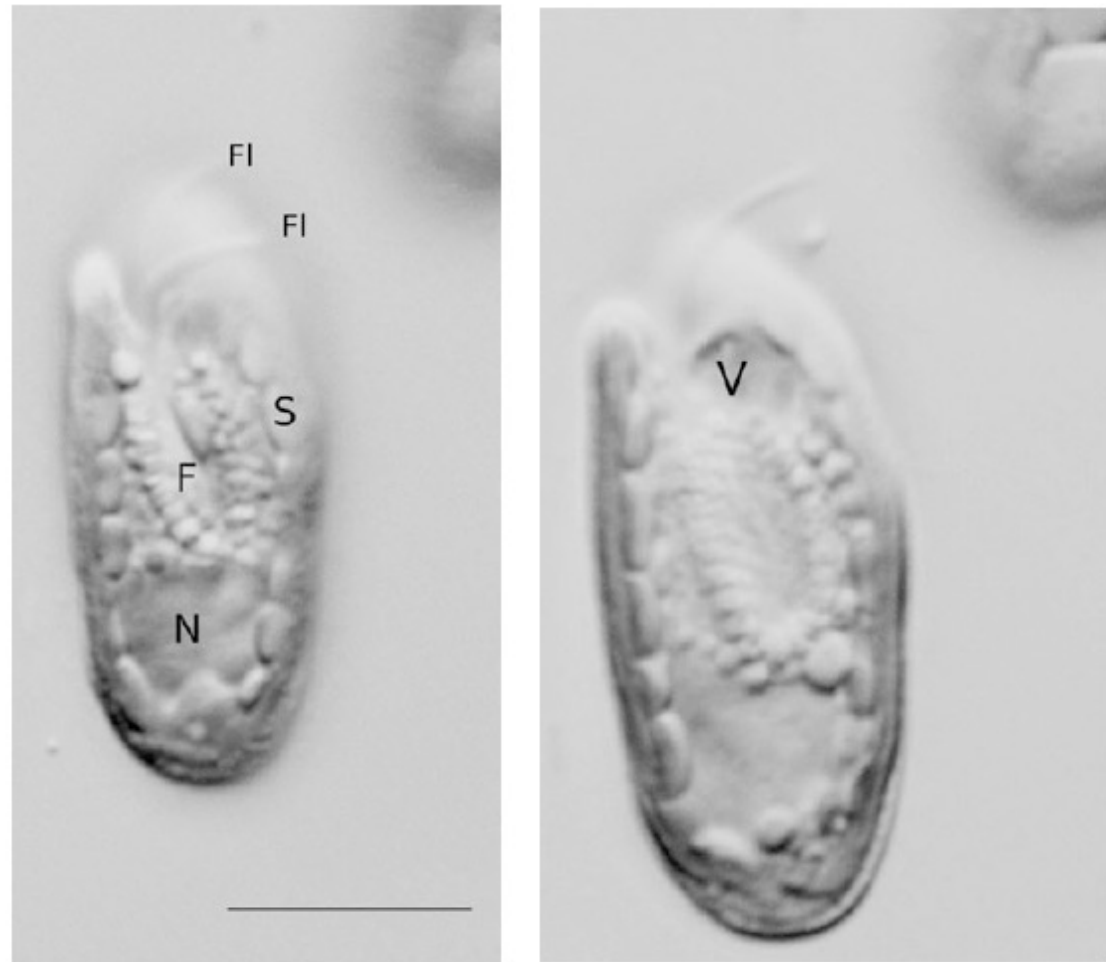


Cryptomonas curvata and *C. paramecium*

Cryptophyta

LEFT image: the ventral side of the cell, showing the elongate open furrow and approximate insertion site of the two flagella. The elliptical platelets lining the cell periphery are starch grains.

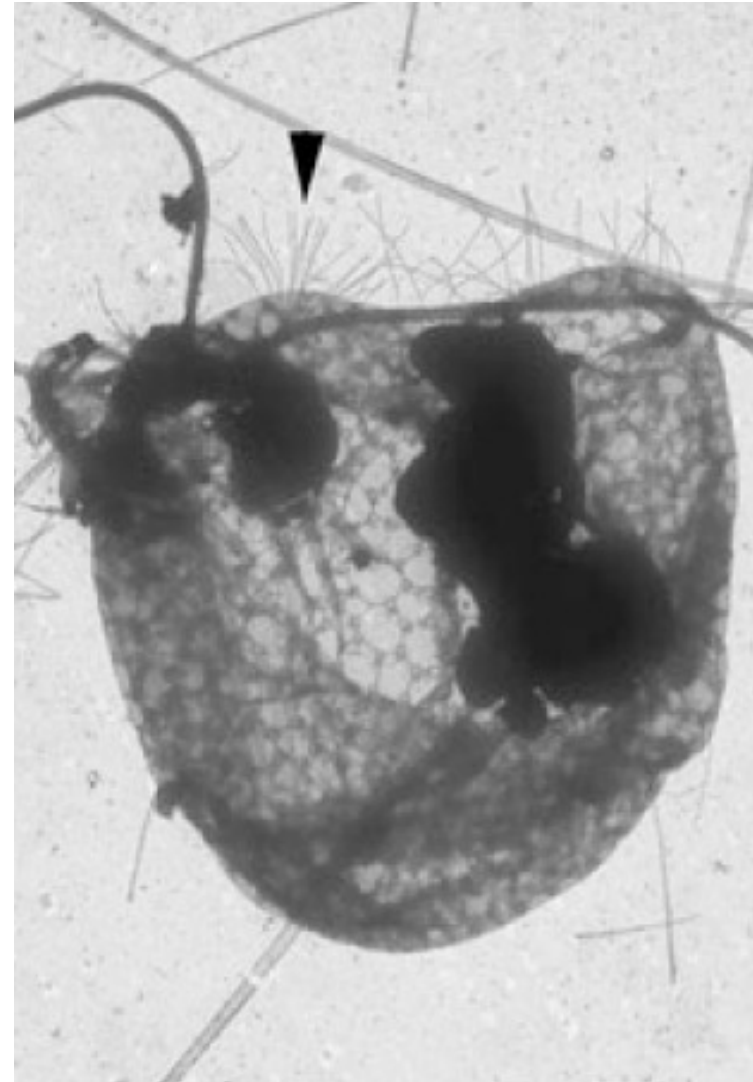
RIGHT image: plane of focus in the cell middle: the ejectosome-lined gullet.



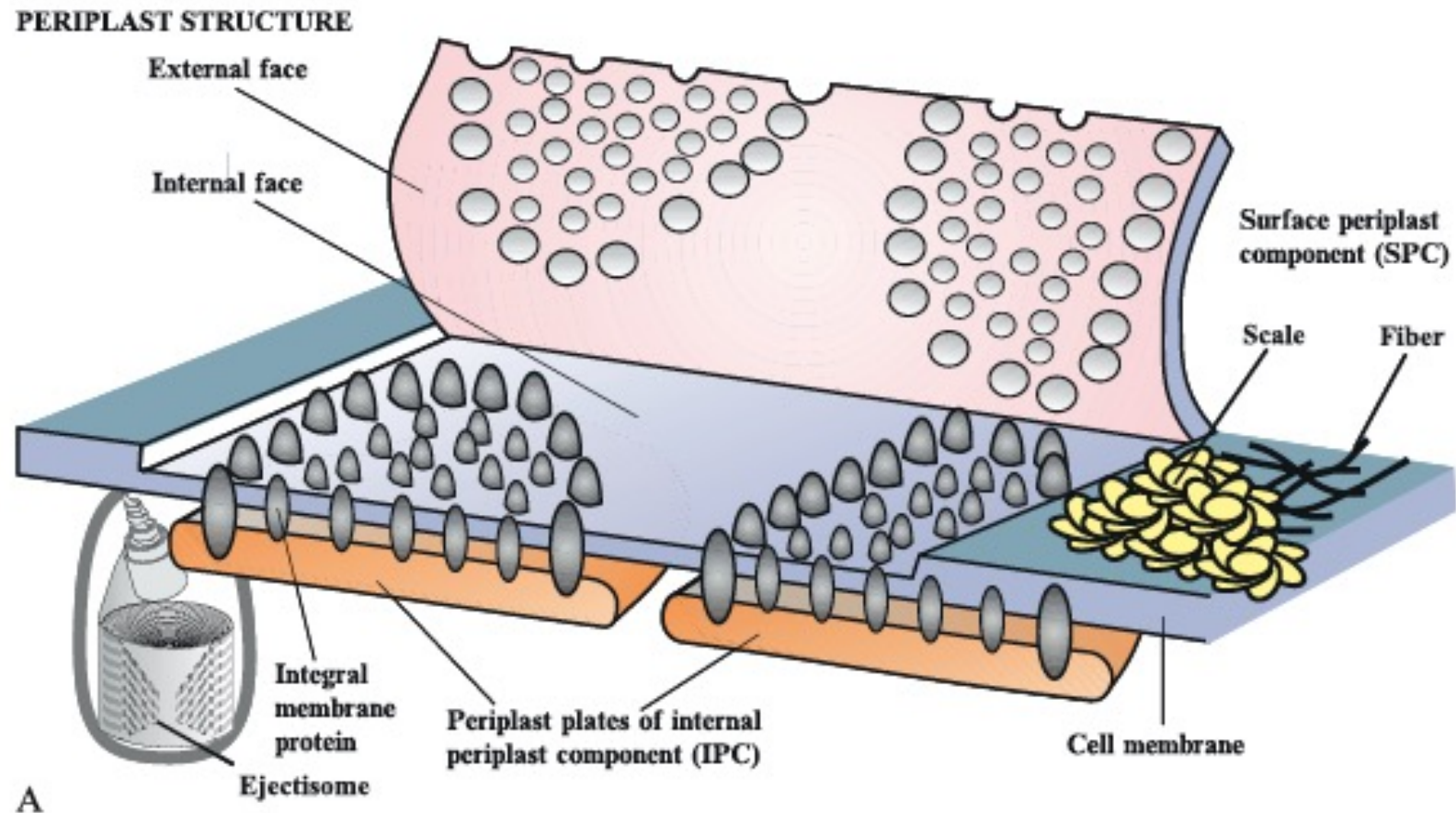
Cryptomonas borealis, the same cell: **FI** - flagella; **S** - starch grain; **F** – furrow-gullet system; **N** - nucleus; **V** - contractile vacuole

Cryptophyta

- **periplast:** composed of proteinaceous inner (IPC) and surface (SPC) layers that sandwich the plasma membrane (no true cell wall)
- the surface periplast component (SPC): covered by congruent plates, rosulate scales, and/or a fibrous coat
- the inner periplast component (IPC): hexagonal, polygonal, rectangular, or elongate plates; or may be continuous and sheetlike
- scales also in association with flagella

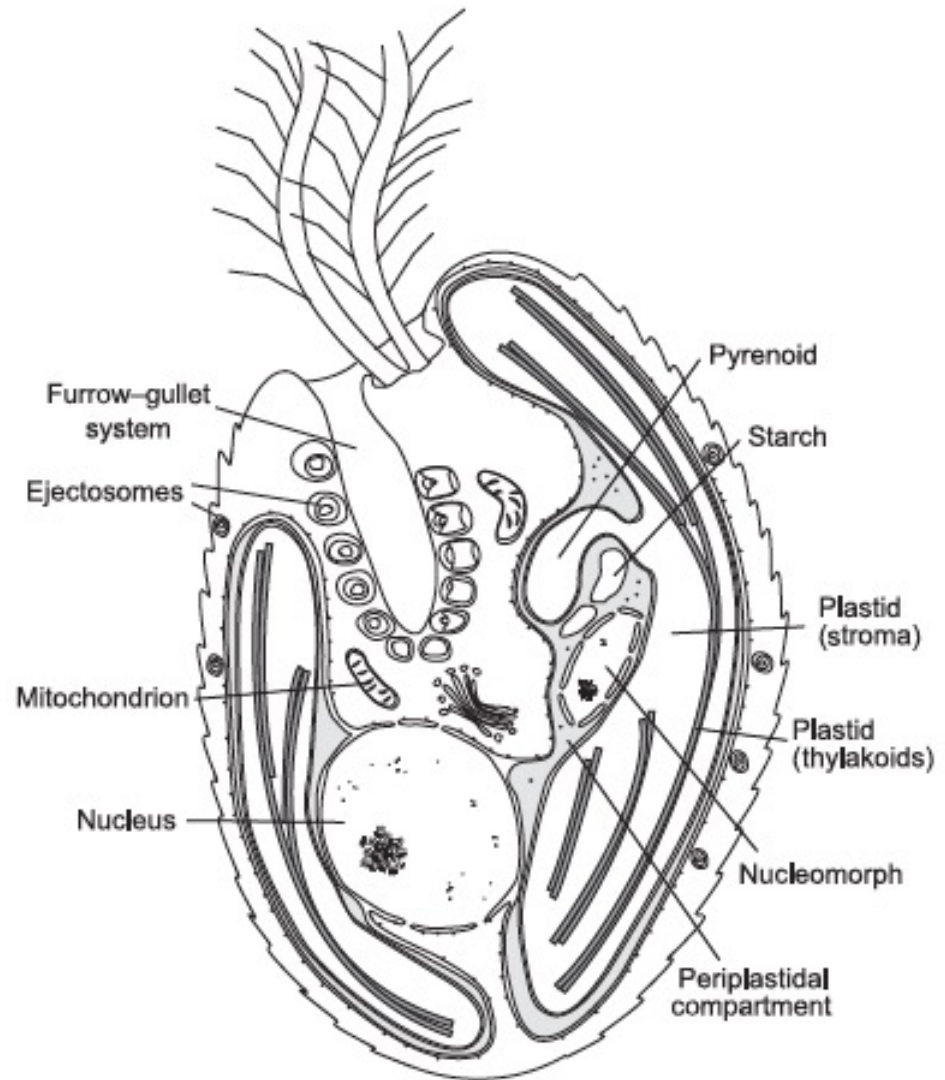


Cryptophyta



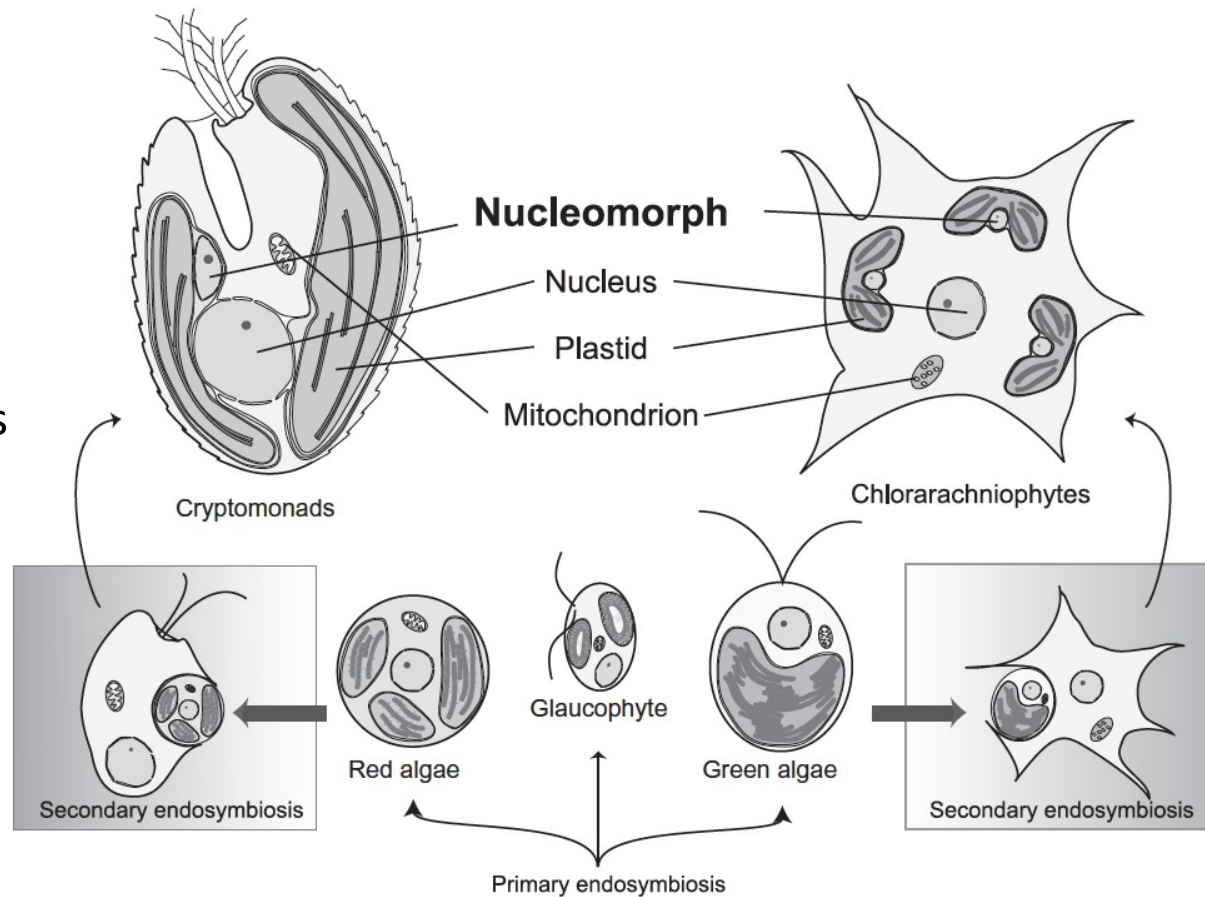
Cryptophyta

- two slightly unequal flagella in the vestibulum of an invagination = the furrow-gullet system; covered with bipartite flagellar hairs
- usually one H-shaped plastid per cell, a single pyrenoid [missing on some species]
- red-algal derived plastid; four membranes (the inner pair - stroma and the outer pair - the plastid ER, which is continuous with the nuclear envelope)
- expanded space = periplastidial compartment (inside: starch grains, pyrenoid, 80S ribosomes, nucleomorph)



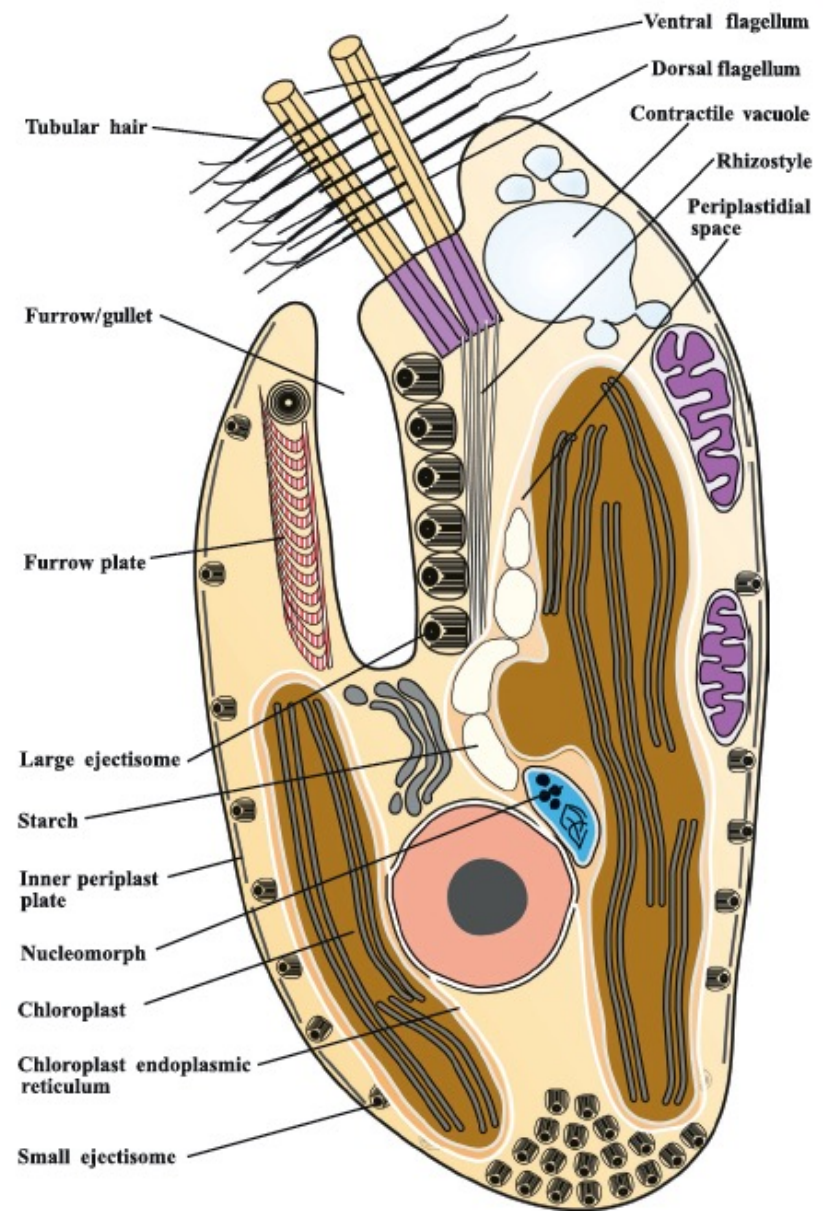
Cryptophyta

- **nucleomorph**: a residual nucleus belonging to a red algal endosymbiont
- often attached to pyrenoids
- two membranes
- presence of the nucleomorph: slow evolution or recent acquisition?



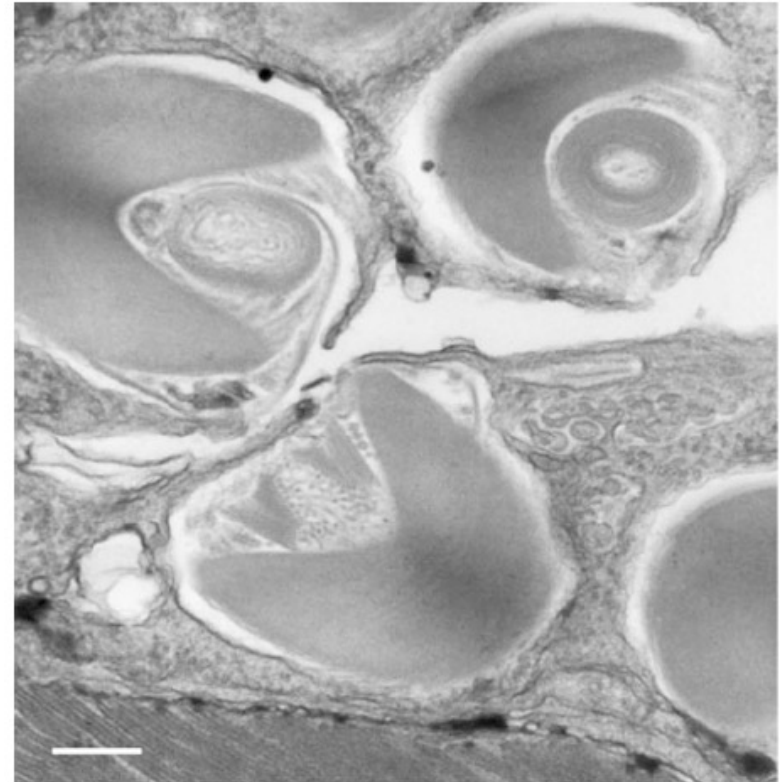
Cryptophyta

- plastids with lamellae comprised of 3 thylakoids, no girdle lamella
- diverse pigments: chlorophyll *a* and *c*₂ + carotenoid alloxanthin + one type of phycobiliprotein per cell: blue = phycocyanin or red = phycoerythrin (both evolved from red algal phycoerythrin)
- phycobiliproteins in thylakoid lumen (not aggregated in phycobilisomes as in Cyanobacteria, red algae, Glaucophyta)
- an eyespot consisting of carotene globules in the plastid (some *Chroomonas* species)
- mitochondria: fingerlike cristae
- starch and lipids as storage material



Cryptophyta

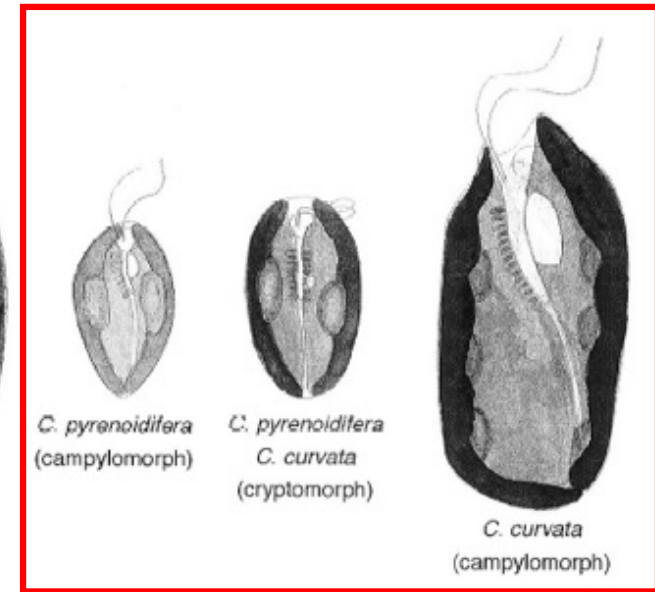
- **ejectosomes**: explosive organelles; large along furrow-gullet system and small ejectosomes near the periplast
- undischarged ejectosome is tightly coiled ribbon-like structure; enclosed within vesicles
- two connected ribbons of different sizes, that are rolled up and under tension
- discharged upon mechanical or chemical stress irritating the cells



large ejectosomes along furrow-gullet system (transmission electron microscopy)

Cryptophyta

- mitosis and cytokinesis (schizotomy) + indirect evidence for sexual reproduction: cell fusion (isogamy), four-flagella zygotes (planozygotes)
- cellular dimorphism: most likely haploid and diploid stage
- some form thick-walled cysts as resting stages (no ejectosomes, no furrow-gullet system)



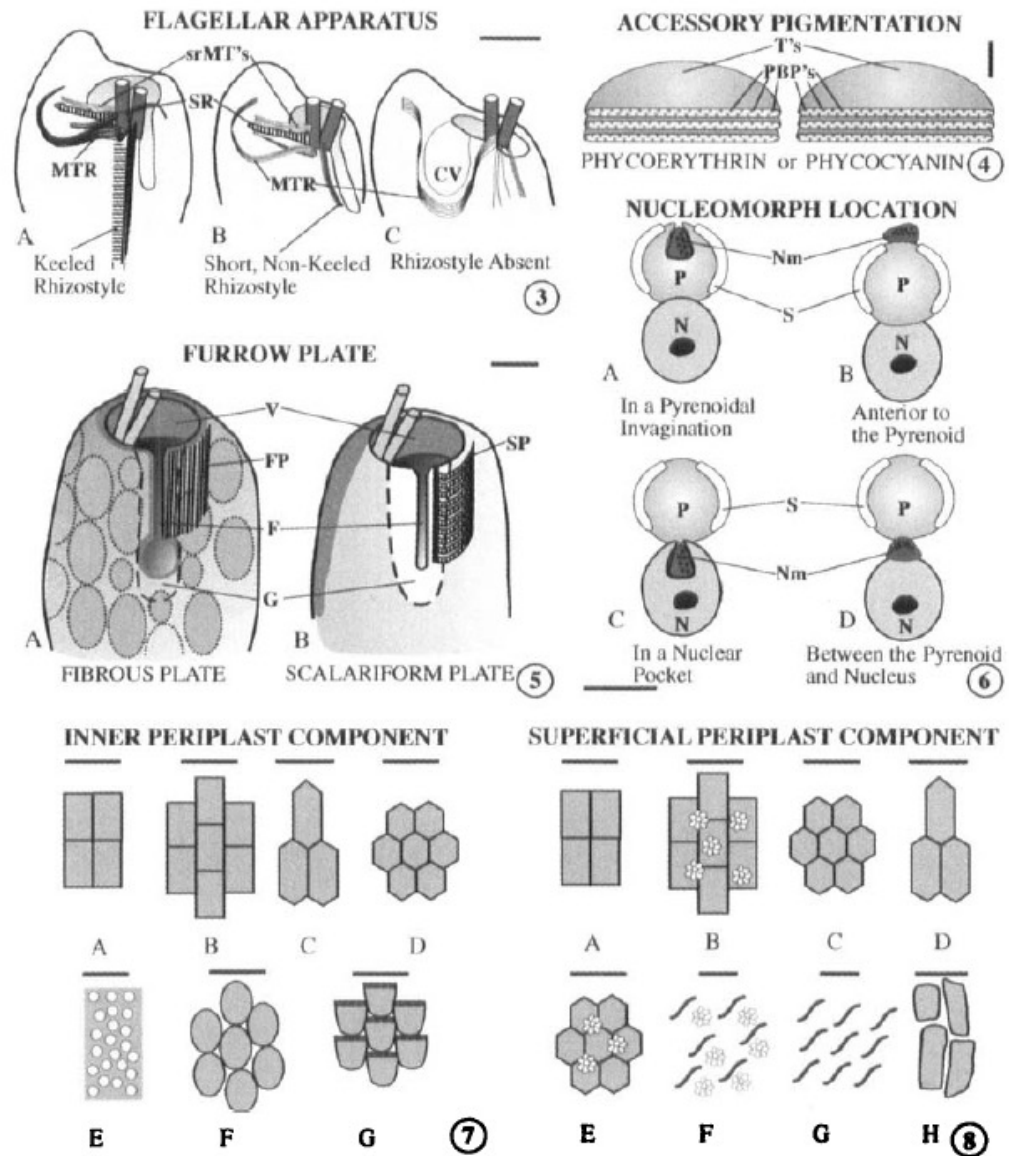
Cryptomonas spp.



resting cyst,
Cryptomonas curvata

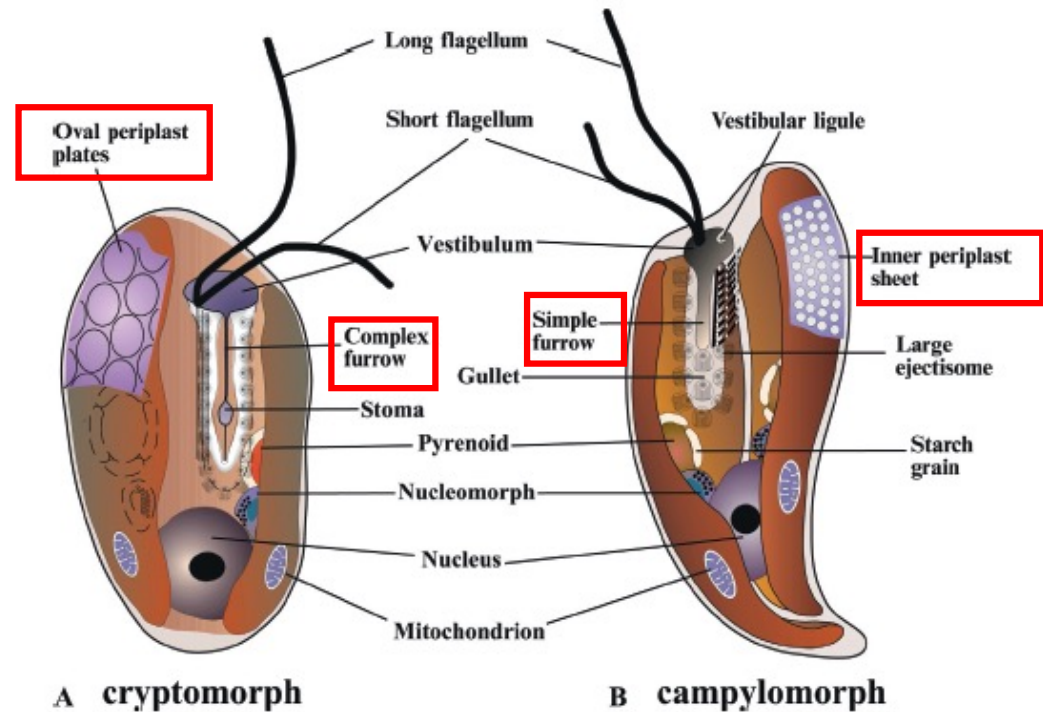
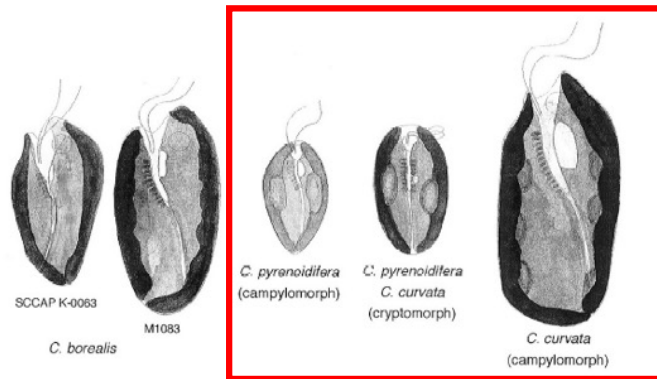
Cryptophyta

- fine-scale features used for classification: biliprotein type, plastid shape and number, position of nucleomorph, pyrenoid structure, flagellar apparatus, morphology of the furrow-gullet system, and ultrastructure of the periplast



Cryptophyta

- some characters are not stable within strains, problems with cell dimorphism



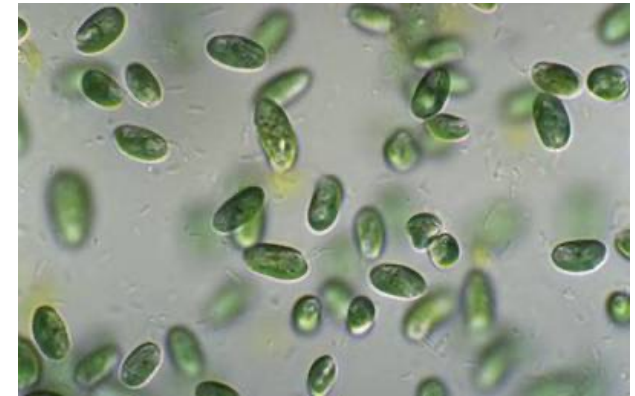
cell dimorphism (haplo/diplontic life cycle?)

Cryptophyta

- unreliability of color classification, because cell's color can change depending on the cell's condition
- a spectrogram analysis of pigment composition is helpful to recognize genera: eight types of distinct phycobiliproteins



Cryptomonas, Campylomonas (brown)



Chroomonas (blue)



Chilomonas (colourless)

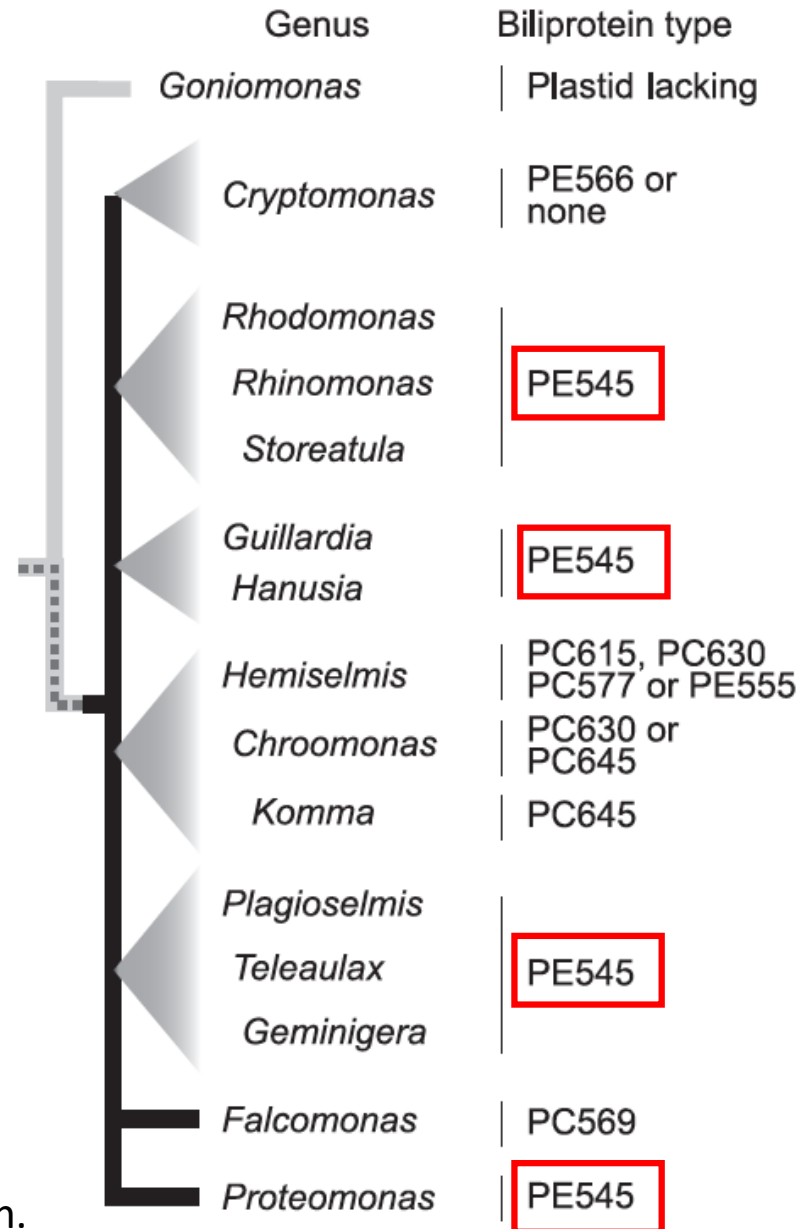


Rhodomonas (red)

Cryptophyta

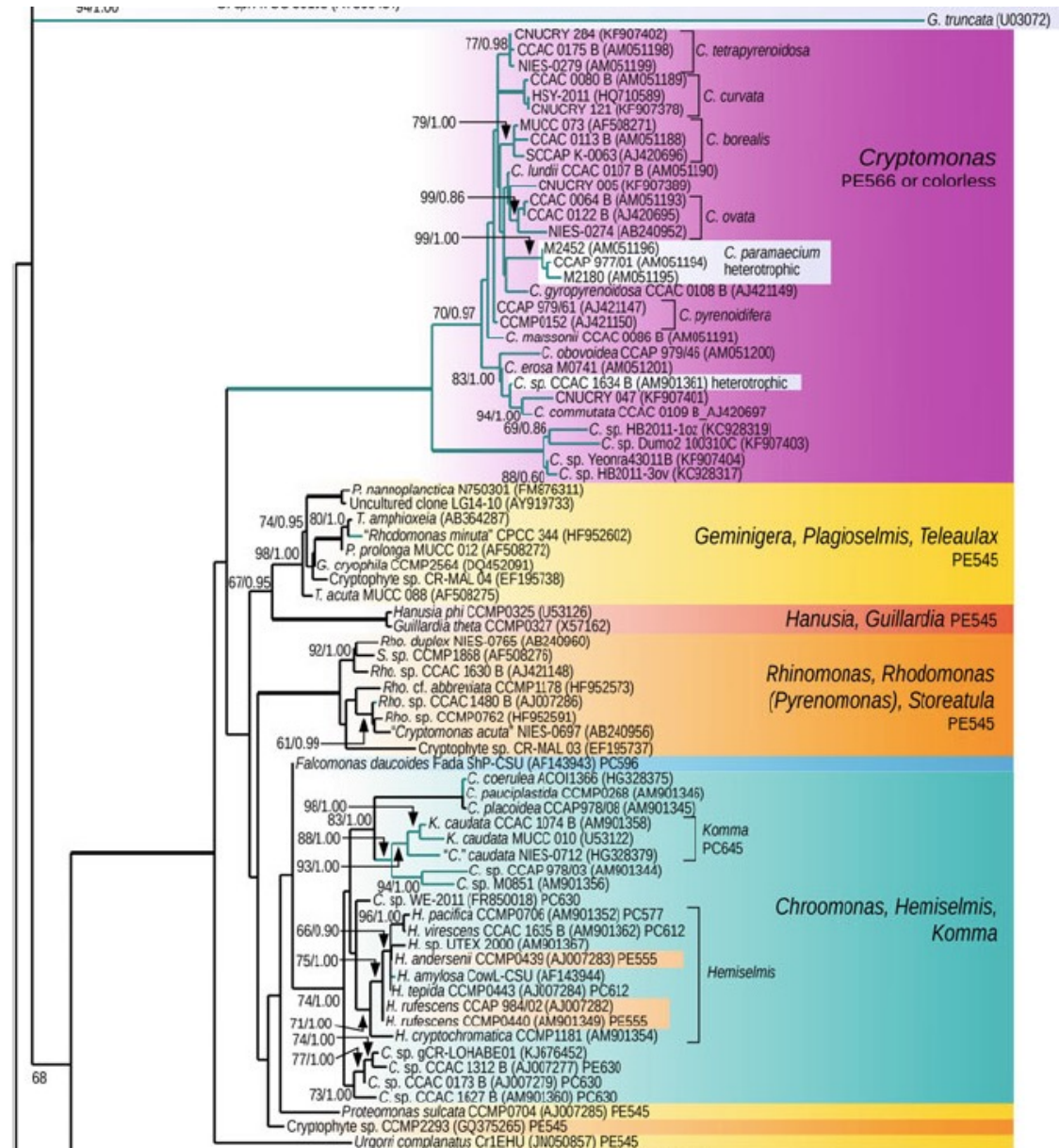
- a spectrogram analysis of phycobiliproteins
- different wavelength of their absorption maxima (e.g. phycoerythrin 545-PE545 = orange-red; phycocyanin 630-PC630)
- but some different genera share the same type of biliprotein or close relatives contain different types

Schematic phylogeny (18S rDNA) of the cryptomonads with pigment information.



Cryptophyta

- morphological classification incongruent with reconstruction of phylogeny (polyphyletic or paraphyletic genera, genetically (almost) identical strains classified as different genera)
- *Cryptomonas* – PE566 or colorless, exclusively freshwater
- lack of species specific morphological characters
- revisiting the classification of cryptomonads is necessary

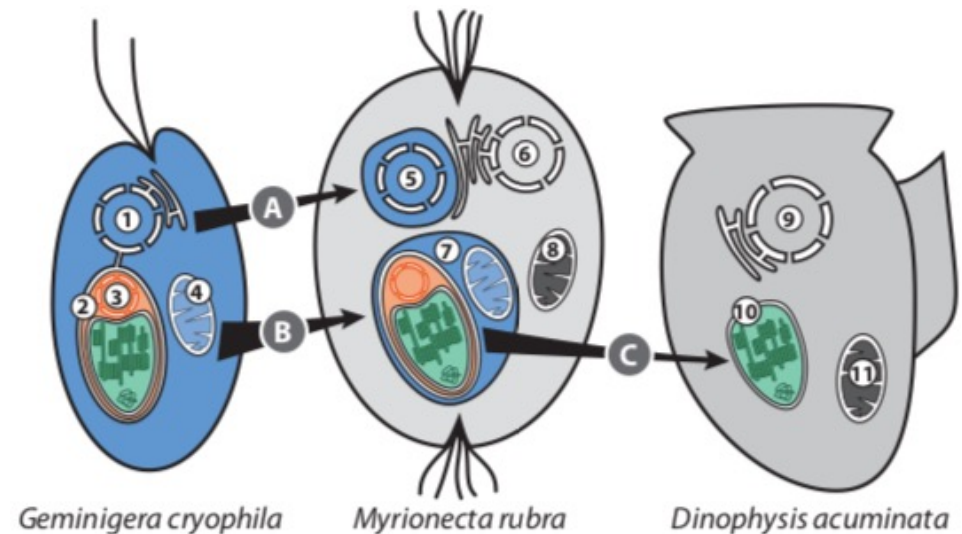


Cryptophyta

- nontoxic [*Chilomonas* may produce ichthyotoxin], easy to mass-cultivate + contain polyunsaturated fatty acid: as food for copepods in fish farming
- biliprotein pigments may be used as fluorescent dyes
- understanding the process of plastid acquisition: i) (non)photosynthetic plastids in cryptomonads; ii) apoplastic phagotrophic sister taxa (+ *Hatena arenicola* temporarily retain an engulfed green algal cell); iii) used as a temporary plastid = kleptoplastid (engulfed by dinophyte, ciliates)
- *Guillardia theta* – known plastid genome

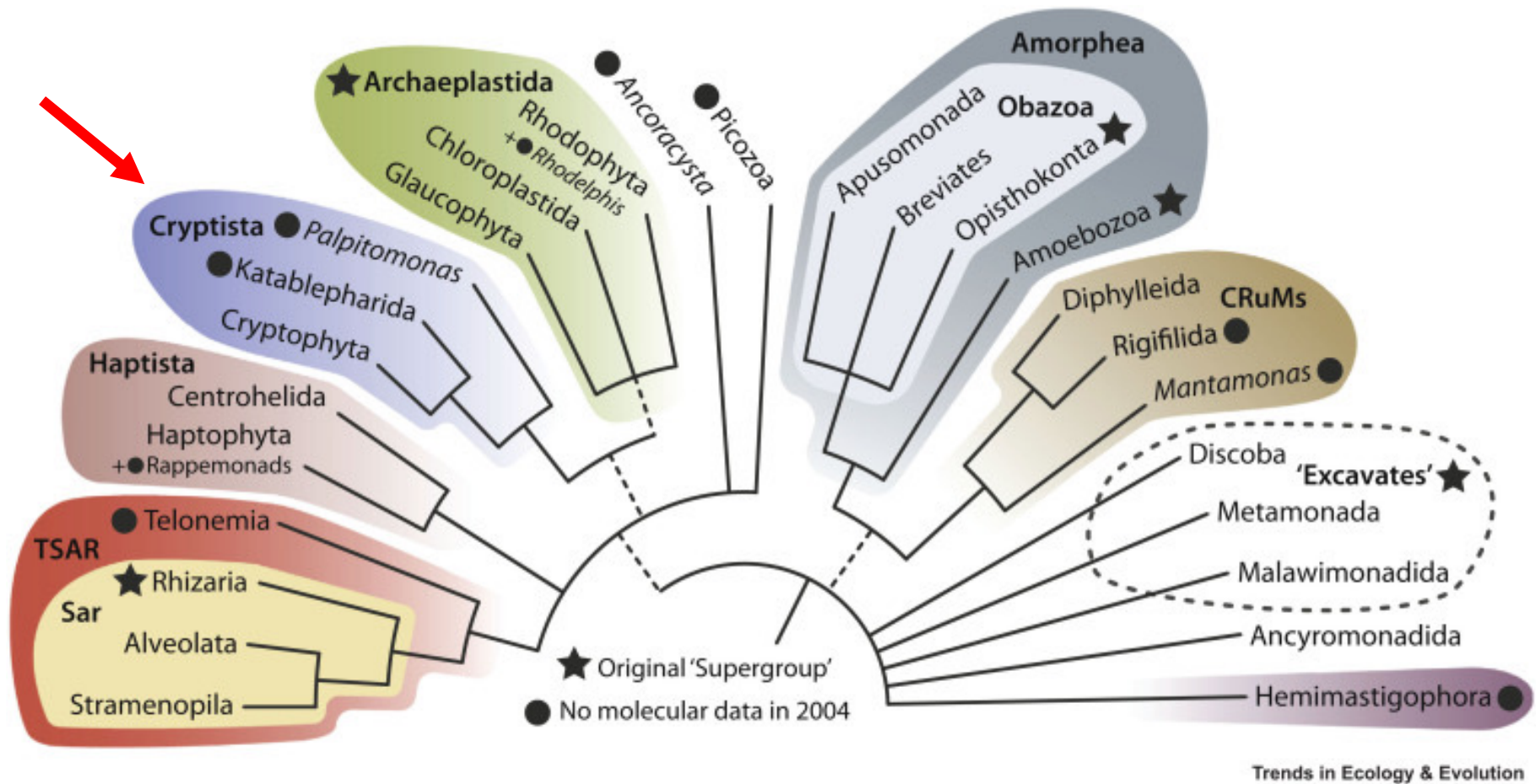


cryptophyte biliproteins, from left to right: PC630, PC577, PE555, PE566

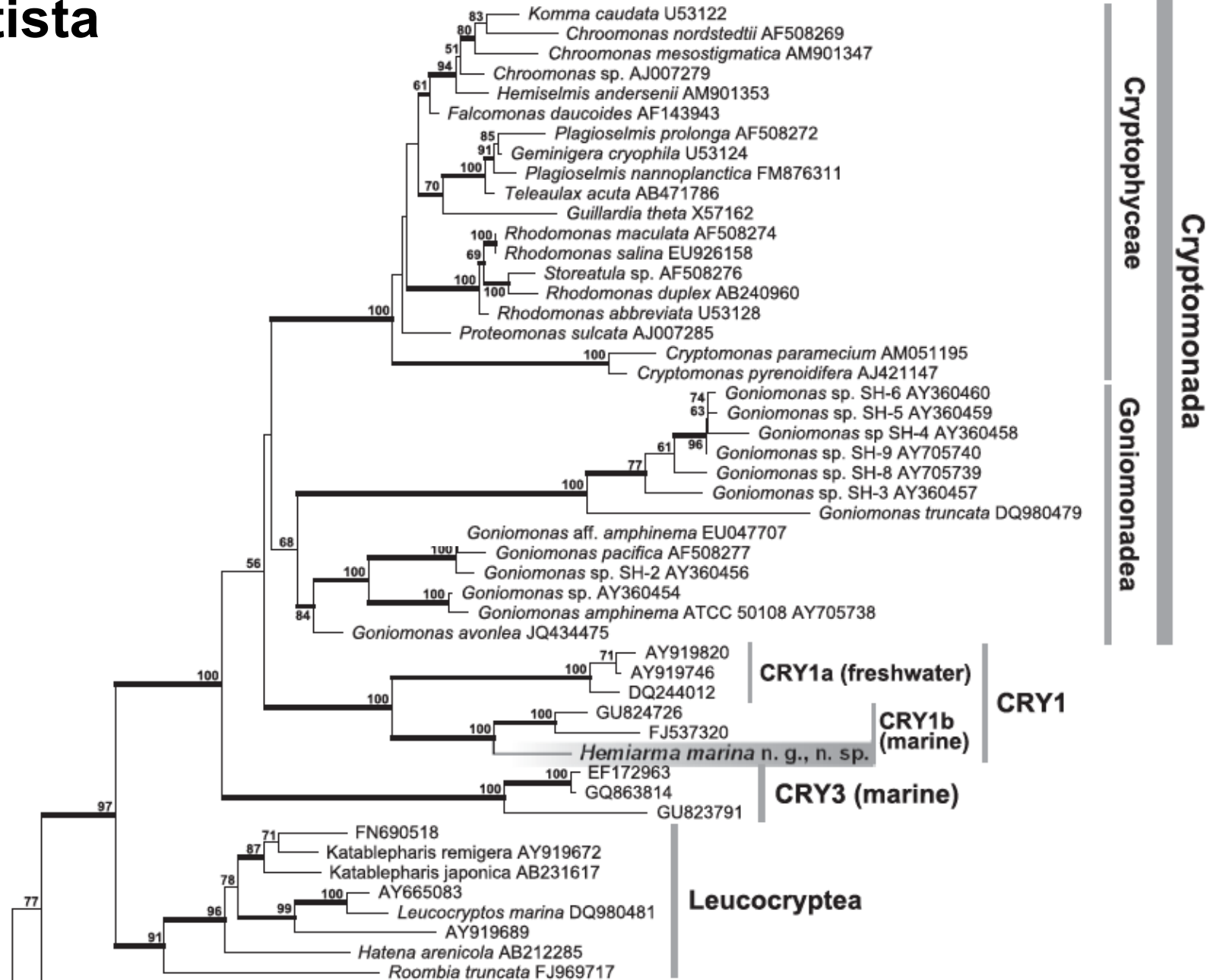


kleptoplast acquisition

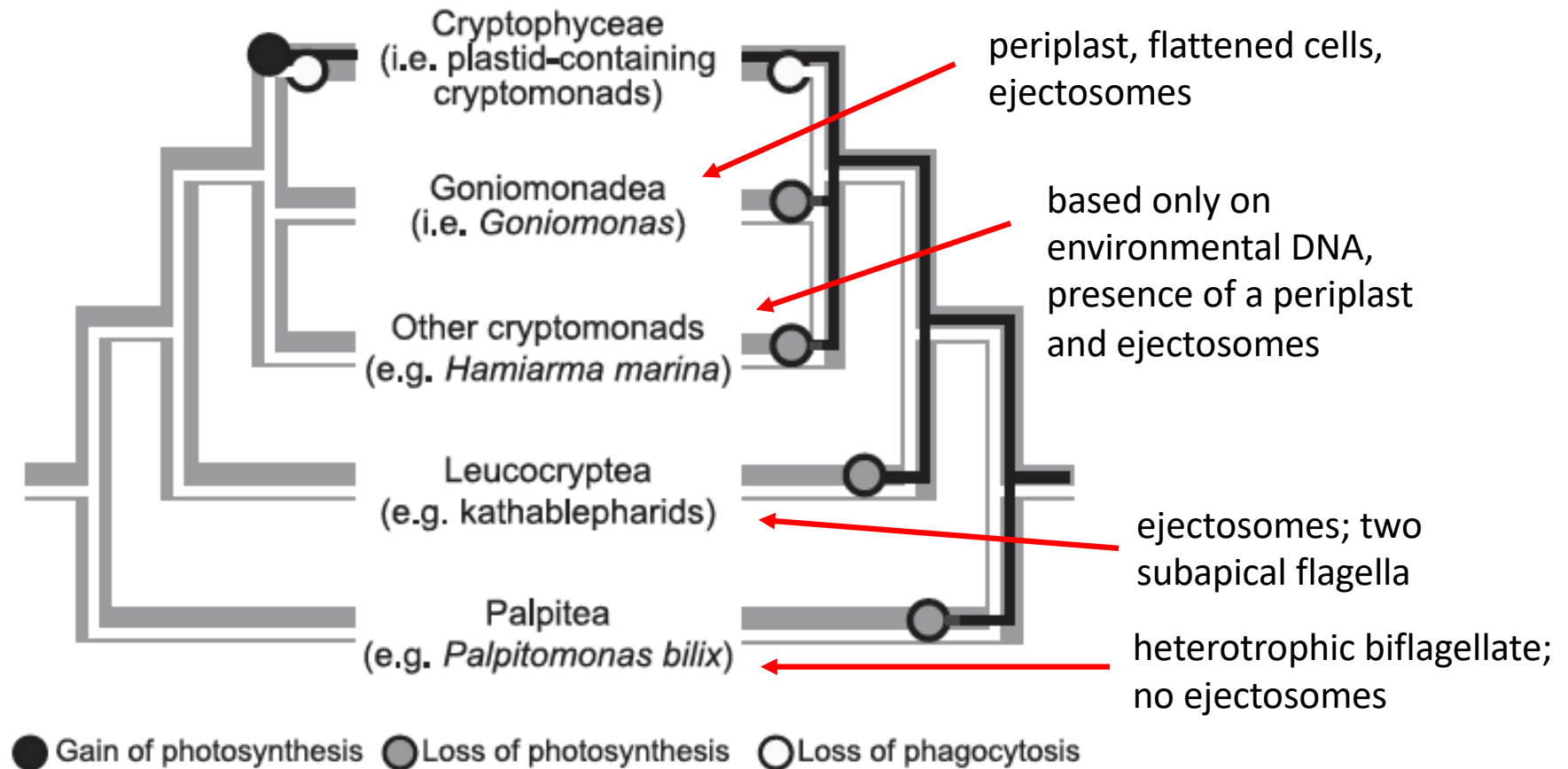
Cryptista



Cryptista



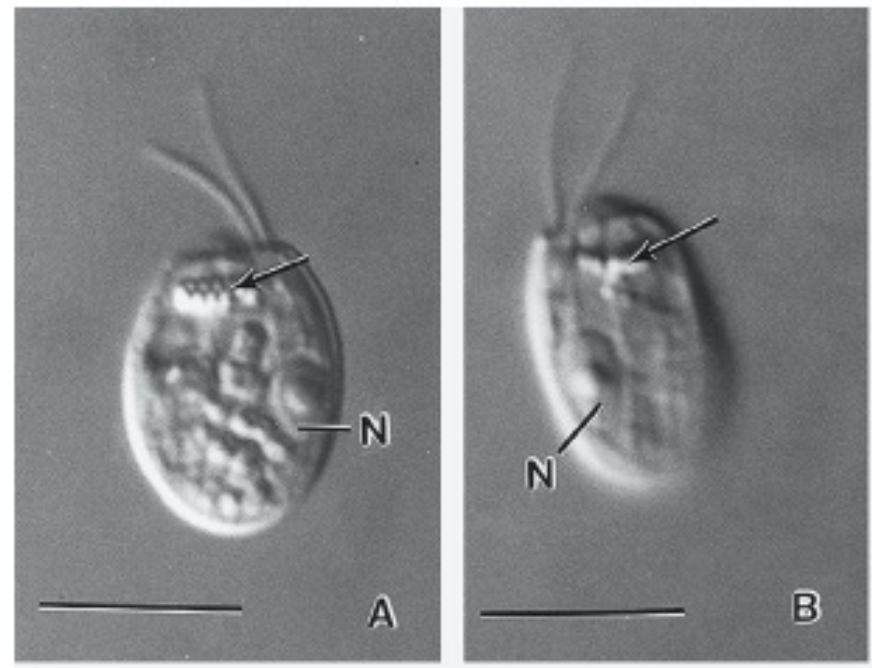
Cryptista



Two possible scenarios of the evolution of plastid acquisition or loss in Cryptista.

Goniomonadea

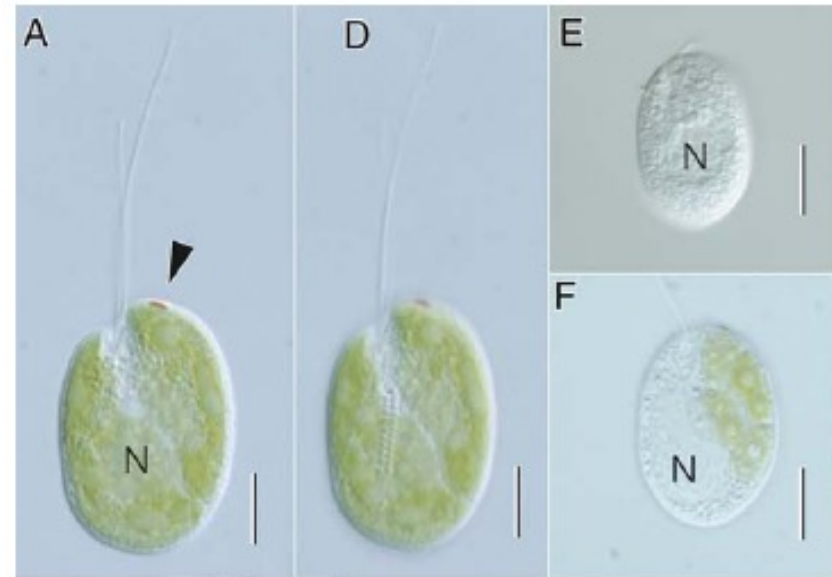
- freshwater habitats, seems to be benthic (prefer gliding along substrates)
- different in shape: markedly flattened with an obliquely truncate anterior
- apoplastic genus, phagotrophy (ingulf bacteria)
- different organization of ejectosomes (see black arrows in figures)
- light microscopically visible striation on the cell surface: large rectangular periplast plates
- long thin filaments cover both flagella



Goniomonas truncata

Katablepharids

- understudied group
- marine and freshwater habitats
- heterotrophic flagellates with ejectosomes
- *Hatena arenicola* harbors a *Nephroselmis* symbiont
- the symbiont plastid is selectively enlarged, while other structures such as the mitochondria, Golgi body, cytoskeleton, and endomembrane system are degraded



Hatena arenicola

Katablepharids

Half-plant, half-predator-hypothesis.

The solid line indicates a witnessed process, and the broken line indicates a hypothetical process. A-D: A green cell with the symbiont, lacking the feeding apparatus (A) divides (B) into one green (C) and one colorless cell (D). E-G: The colorless cell should form a feeding apparatus de novo and engulfs a *Nephroselmis* cell. G-H: The symbiont plastid selectively grows in the host cytoplasm. Because cell division of a colorless cell or a cell with an “immature” symbiont (H) has never been observed, uptake and the subsequent changes in both host and symbiont apparently occur within one generation.

