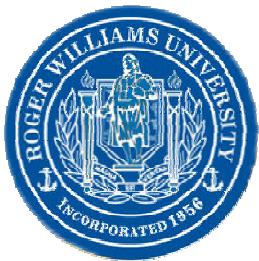




Introduction to the Classification of Green Algae



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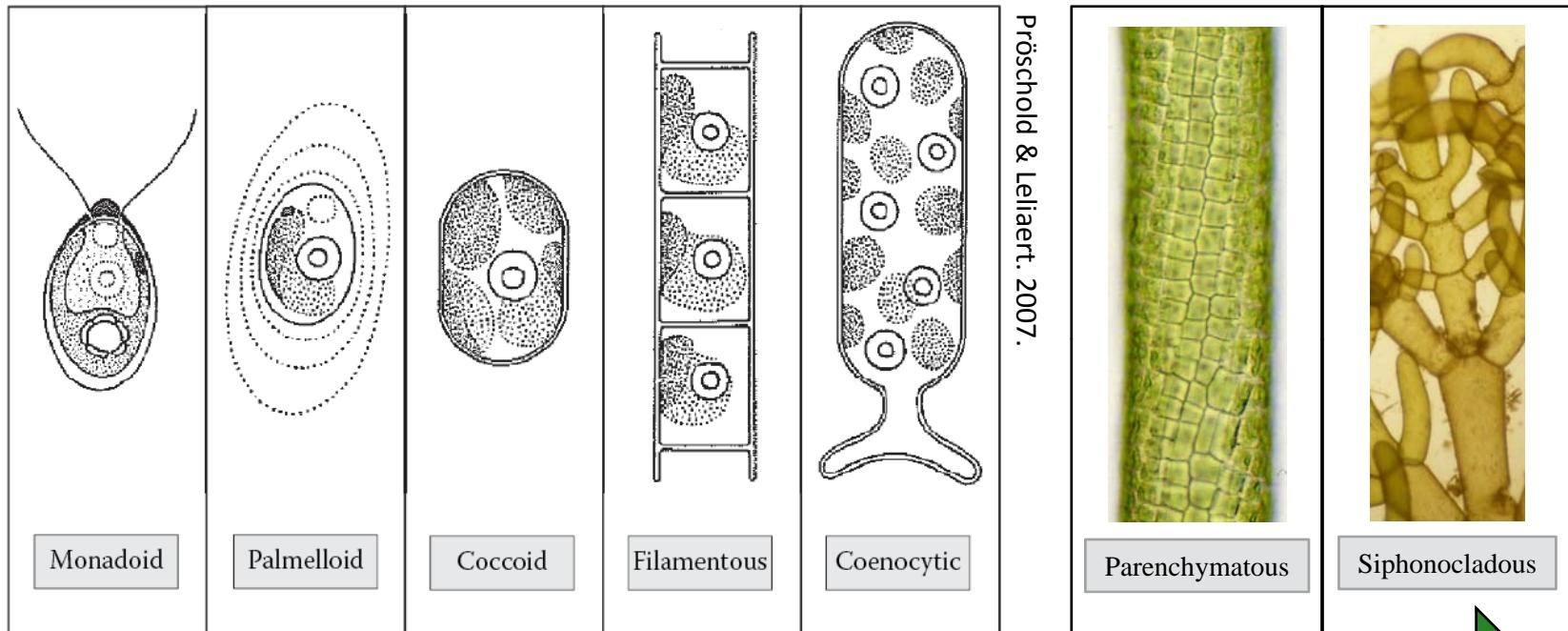


Classification

- 20+ classes of green algae have been described since Linnaeus
- Many criteria have been used to establish classes
 - Morphology
 - Ultrastructure
 - + mitosis, cell division, cytokinesis, flagellar apparatus
 - Life history patterns
 - Molecular
 - Primarily rRNA genes, *rbcL*
- Ambiguity remains, but consensus is emerging



Classification: Morphological Concept



Evolution of structural complexity

- Premise
 - Primitive unicells evolved structural complexity over time
 - Morphological divergence distinguishes groups

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Problems with Morphological Concept

- Selective environmental pressure may drive convergence
 - Biologically/genetically distinct species appear similar

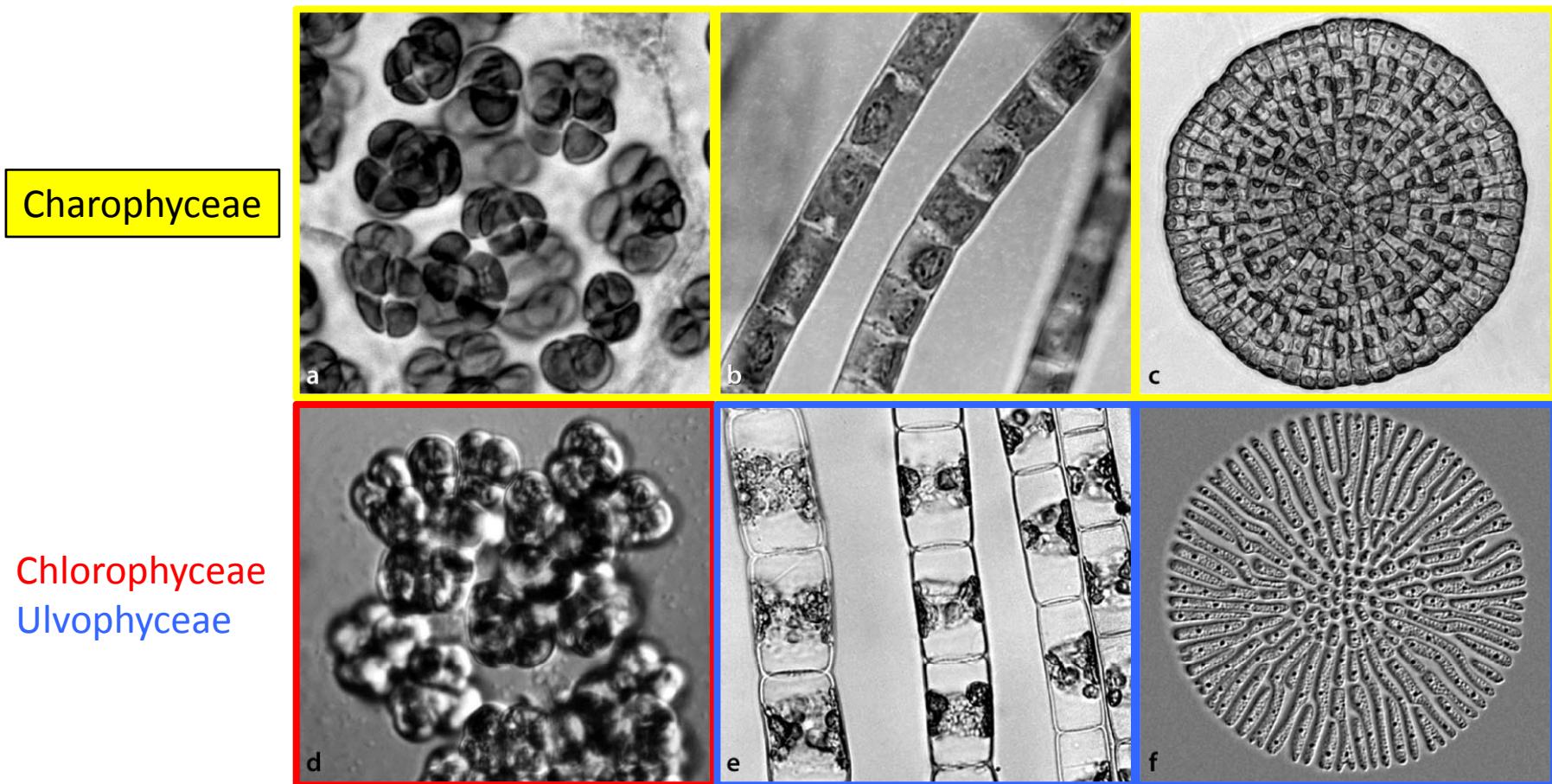


Fig. 16.18 Graham et al. 2008.

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Morphological Convergence w/in Class



Class: Ulvophyceae

Order: Ulvales

Genus: *Ulvaria*



Ulvophyceae

Ulotrichales

Monostroma

Chlorophyta: Morphological Convergence

- How can green algae be classified if they exhibit both divergent and convergent patterns of morphology?

7 Systematics of the green algae: conflict of classic and modern approaches

Thomas Pröschold and Frederik Leliaert

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ABSTRACT

Traditionally the green algae were classified in orders or classes according to the morphological species concept, but, for example, monoidial species (flagellates) were summarized in the order Volvocales, coccoids in the Chrysophycales, filaments in the Ulvophytales or Chlorophytales, and sphaeropleidians algae in the Chlamydomonadales or Siphonophytales. Later, a new classification was proposed based on ultrastructural investigations of the basal bodies in the flagellar apparatus and cell division. The species with basal bodies in clockwise (CW) or directly opposite (DO) orientation were classified in the class Chrysophyceae, the counter-clockwise (CCW) orientated species in the Chlophyceae and Trebouxiophyceae (= Plantastrophycaceae). Phylogenetic analyses of nuclear-encoded SSU and ITS rRNA sequences have basically confirmed the classification based on ultrastructural characters. However, most genera and orders are polyphyletic and the relationships between

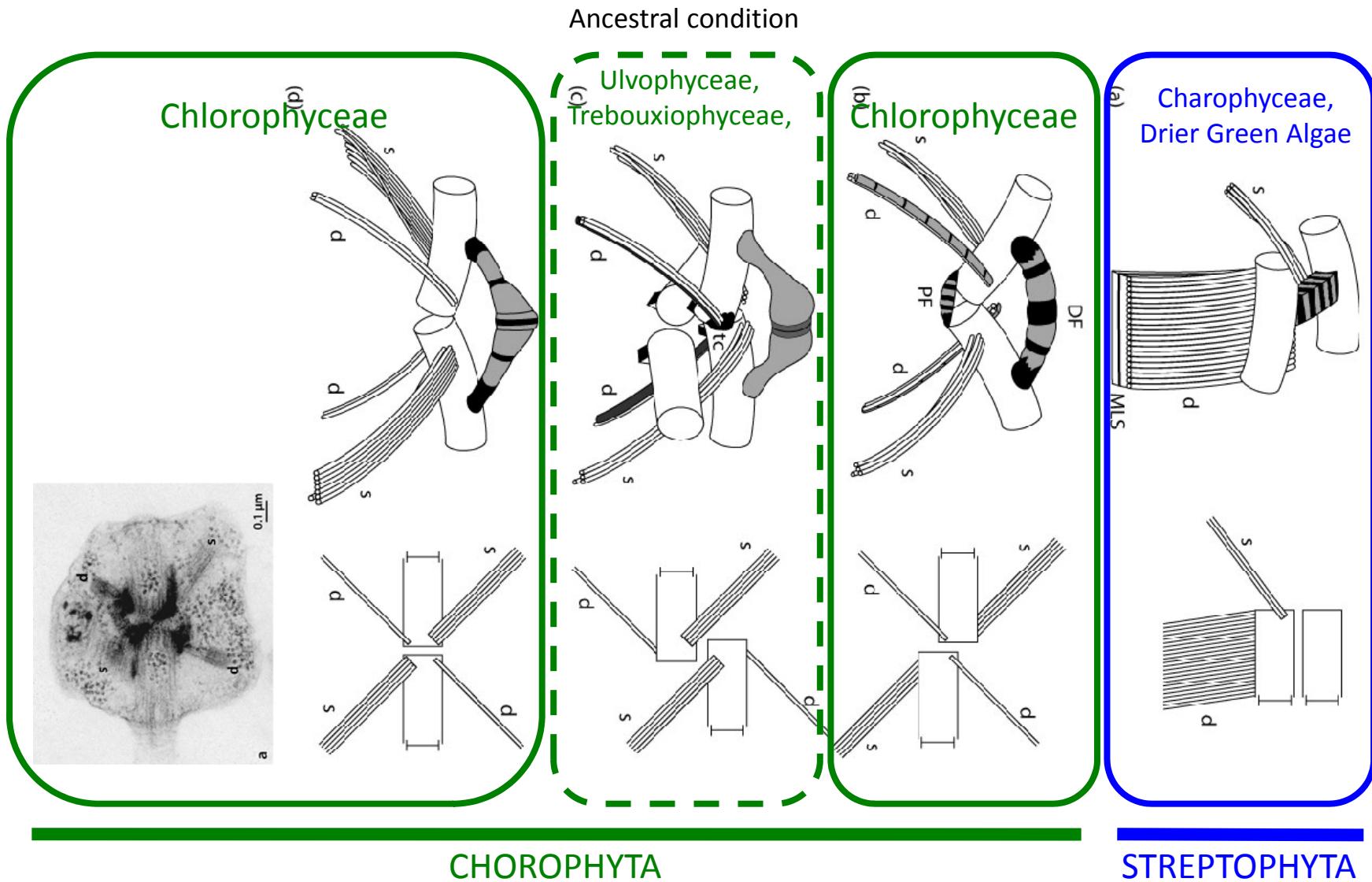


Ultrastructural Concept

- Mattox & Stewart (1984) proposed new Class-level taxonomy on the basis of basal body orientation
 - 1st radiation of green algae occurred among unicellular flagellates
 - Numerous ancient flagellate lineages diversified into contemporary forms along different evolutionary paths, but many with similar morphological results
- 4 different basal body patterns identified



Basal Body Orientations



Molecular Concept

- Elucidation of green algal phylogeny in molecular context commenced in early 1990s
- Early studies
 - confirmed distinction between Chlorophyta & Streptophyta already suggested by cellular and ultrastructural characters elucidated from EM
 - recognized 5 lineages:
 - Prasinophyceae, Chlorophyceae, Trebouxiophyceae, Ulvophyceae, Streptophyta + Land plants
 - Subsequent studies have confirmed these lineages, but relationships among & within the lineages remains unclear



Molecular Concept

(McCourt & Lewis 2004, Working Classification)

molecular data support the recognition of two phyla and a poorly resolve graded of early diverging flagellates informally known as Prasinophyceans

Chlorophyta

Streptophyta
(Charophyceae +
Drier Green Algae)

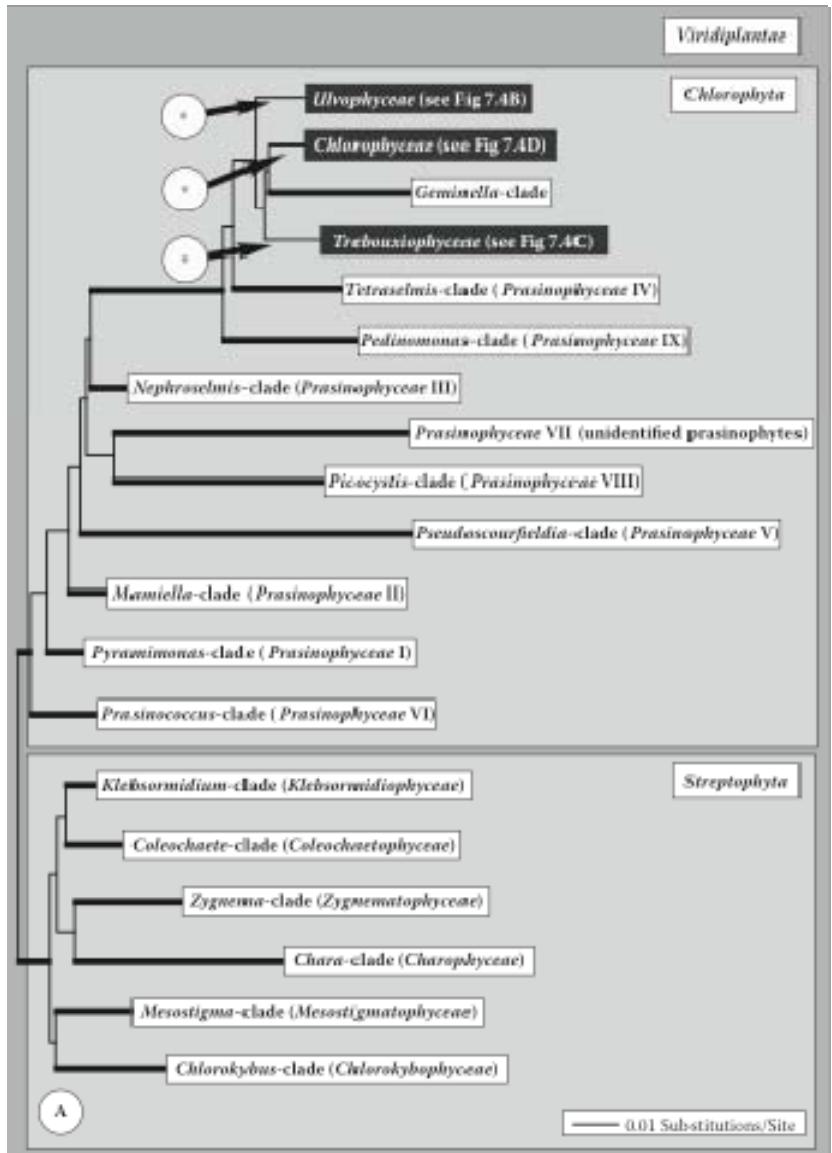
Viridoplantae or Chlorobionta

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SSU rDNA Molecular Concept

(Pröschold & Leliaert 2007)

- Chlorophyta + Streptophyta recovered
- UTC clade recovered with low BS support (no bold lines)
- Poorly resolved
Prasinophycean grade

FIGURE 7.4 Molecular phylogeny of the Viridiplantae based on SSU rDNA sequence comparisons. The phylogenetic tree shown was inferred by the neighbour-joining method based on distances of 1668 aligned positions of 498 taxa using PHYLIP 3.6 (Saitou and Nei 1987) and analysed with the dataset PART A (Figure 12) with confidence limits from 1000 BS replicates. Taxa and Node numbers by Mandelat et al. (Pröschold and Heckendorf 2002; Pröschold and Chaudhuri 2006). Bootstrap percentage values ($\pm 1\%$) are marked by bold brackets prior to node numbers in parentheses. Rooted tree of the Chlorophytidae and Streptophytidae clade. Taxa that are chlorophytes, Chlorophytidae, and Streptophytidae are represented by open circles in this analysis, the clade itself is blacked. In Node 10, G = 0.8. The clade containing species of the three so-called green algal clades marked by an asterisk were not included in bootstrap analyses, however, they were separately analysed in part B through D and marked with a black

Class Ulvophyceae

- Most macroscopic marine green algae belong to Class Ulvophyceae
- Ulvophyceae is circumscribed on the basis of a suite of plesiomorphic characters
 - Closed mitosis, persistent spindle, furrowing, CCW, flagellate cells with scales
- The absence of well-defined, shared-derived characters has resulted in uncertain relations within the class



Class Ulvophyceae

- Generally, 5 (or 6) orders are distinguished
 - flagellar apparatus
 - zoosporangial & gametangial structures
 - life history
- Orders are generally supported by molecular analysis, but relationships among orders are largely uncertain

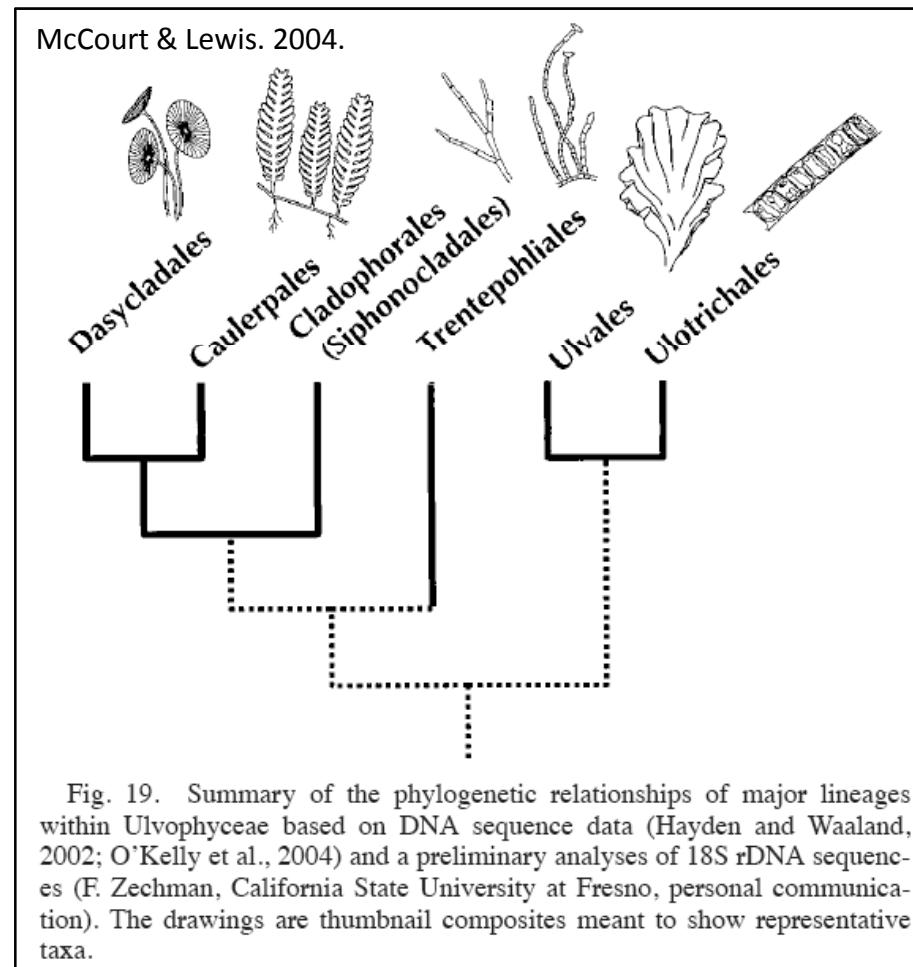
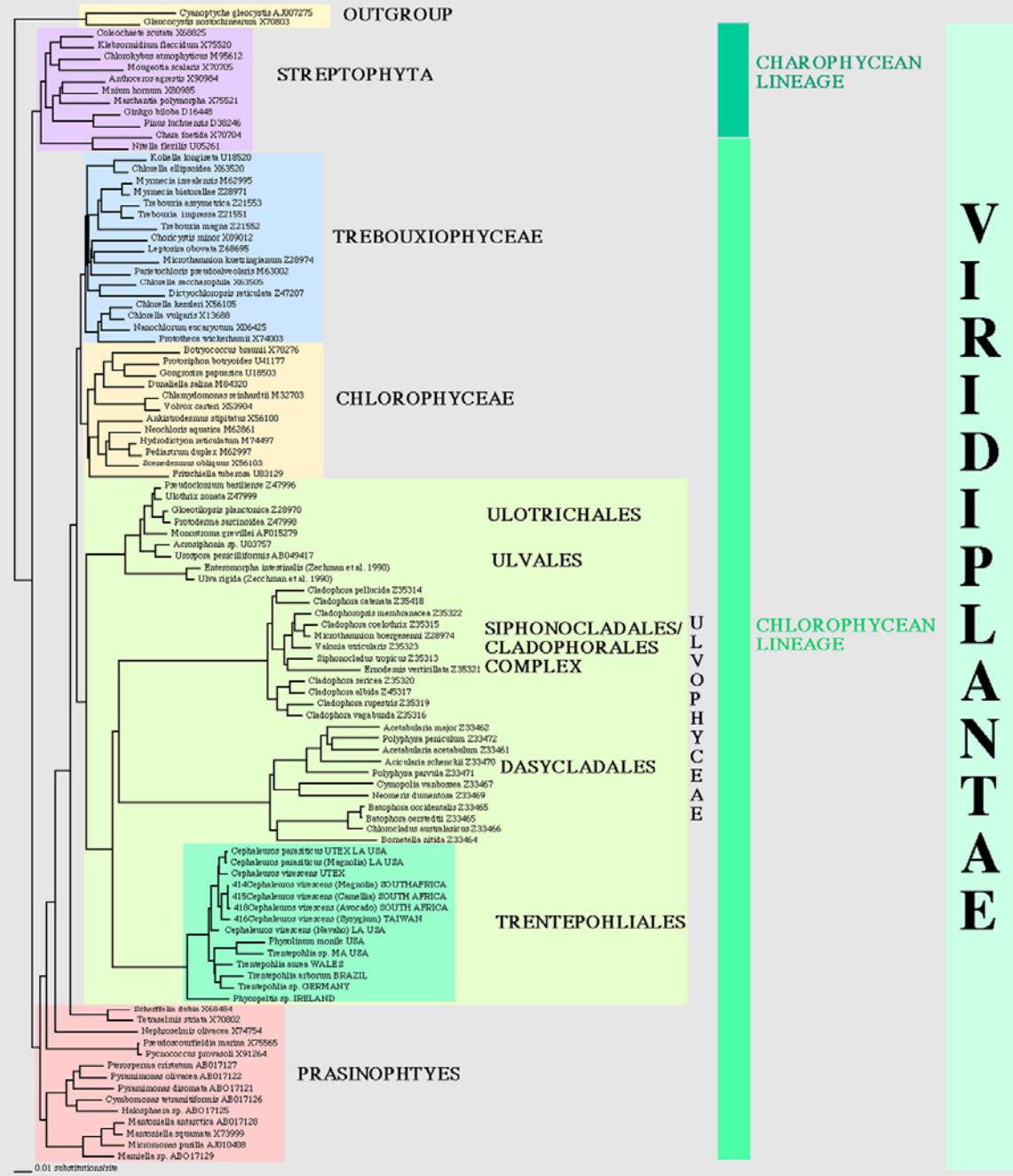


Fig. 19. Summary of the phylogenetic relationships of major lineages within Ulvophyceae based on DNA sequence data (Hayden and Waaland, 2002; O'Kelly et al., 2004) and a preliminary analyses of 18S rDNA sequences (F. Zechman, California State University at Fresno, personal communication). The drawings are thumbnail composites meant to show representative taxa.

Molecular Concept

(Lopez-Bautista et al. 2002)

Figure 22. Maximum likelihood tree, estimated from Kimura two-parameter model of sequence evolution using Modeltest parameters. $-\ln L = 22774.1914$. A total of 102 taxa with 1720 positions are represented in the phylogram. Genbank accession numbers are given after the scientific name.



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Sub-ordinal classification

- Generic and species level classification is frequently based on morphology which can be problematic
 - Phenotypic plasticity
 - Morphological convergence
- Integrated approaches are required to resolve these problems
- DNA sequencing (and distance clustering) is a start that can lead to the recognition of novel morphological, physiological or biological features

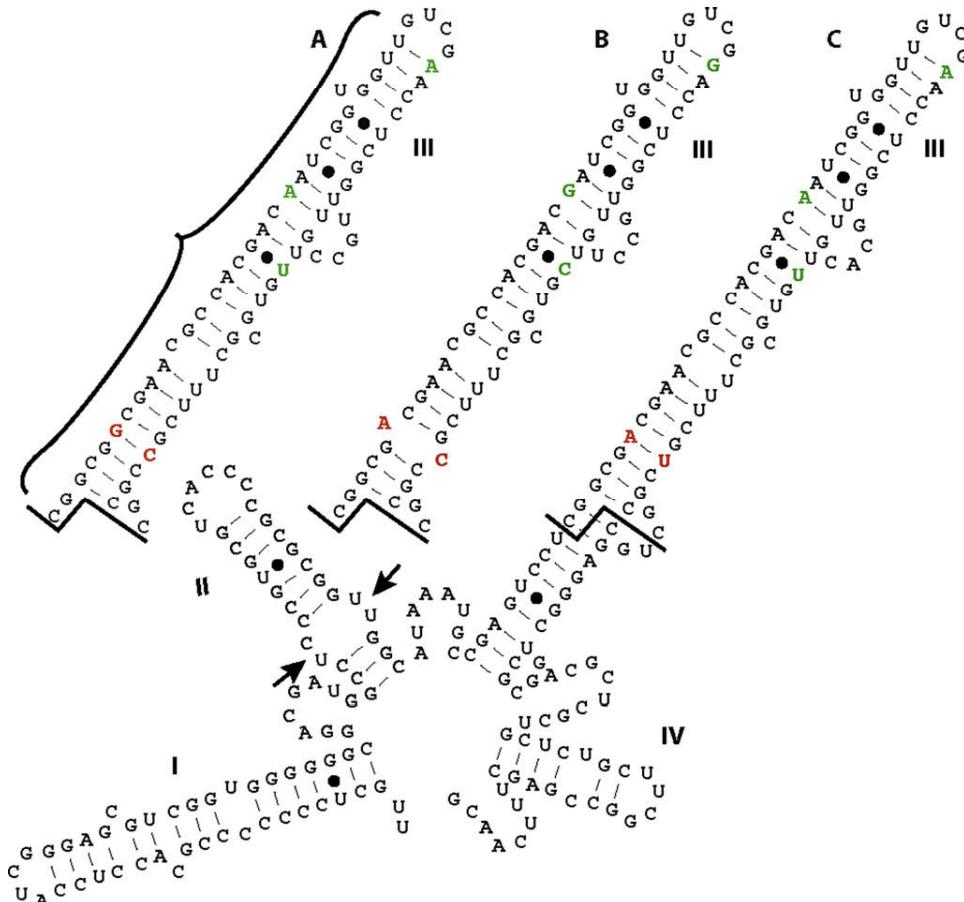
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Z-clades & Biological Species inferred from ITS2 2°structure (see Coleman 2009)



- As ITS2 differences between potential mates increase, sexual compatibility and zygote productivity decrease
- Organisms that differ by even one CBC in helix III are completely unable to cross
- Identity for the entire ITS2 correlates with significant interbreeding potential.

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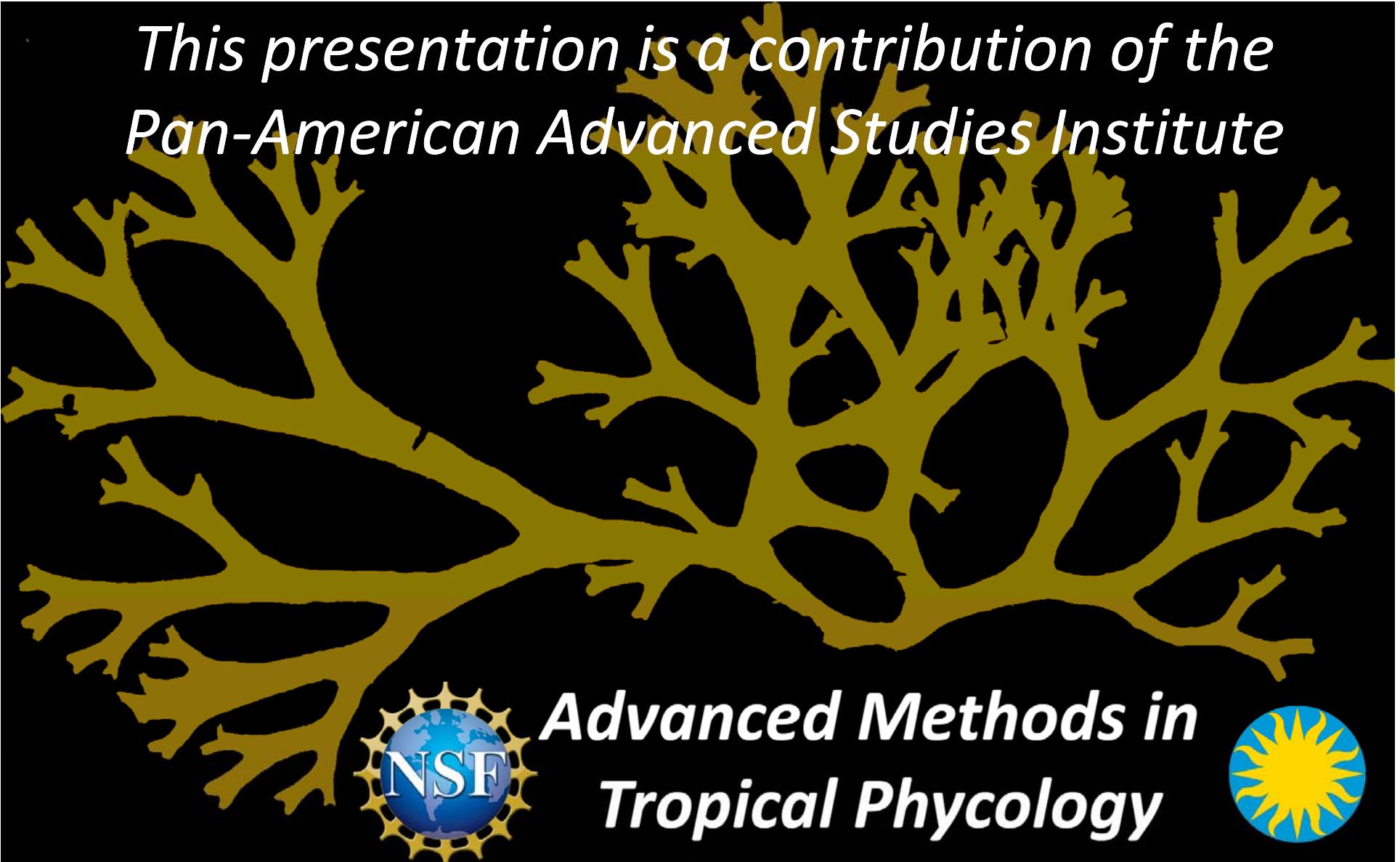


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