

Green Tree of Life: Resolving green plant phylogeny, from morphology to genomics

Presented by:

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Bigelow Laboratory for Ocean
Sciences

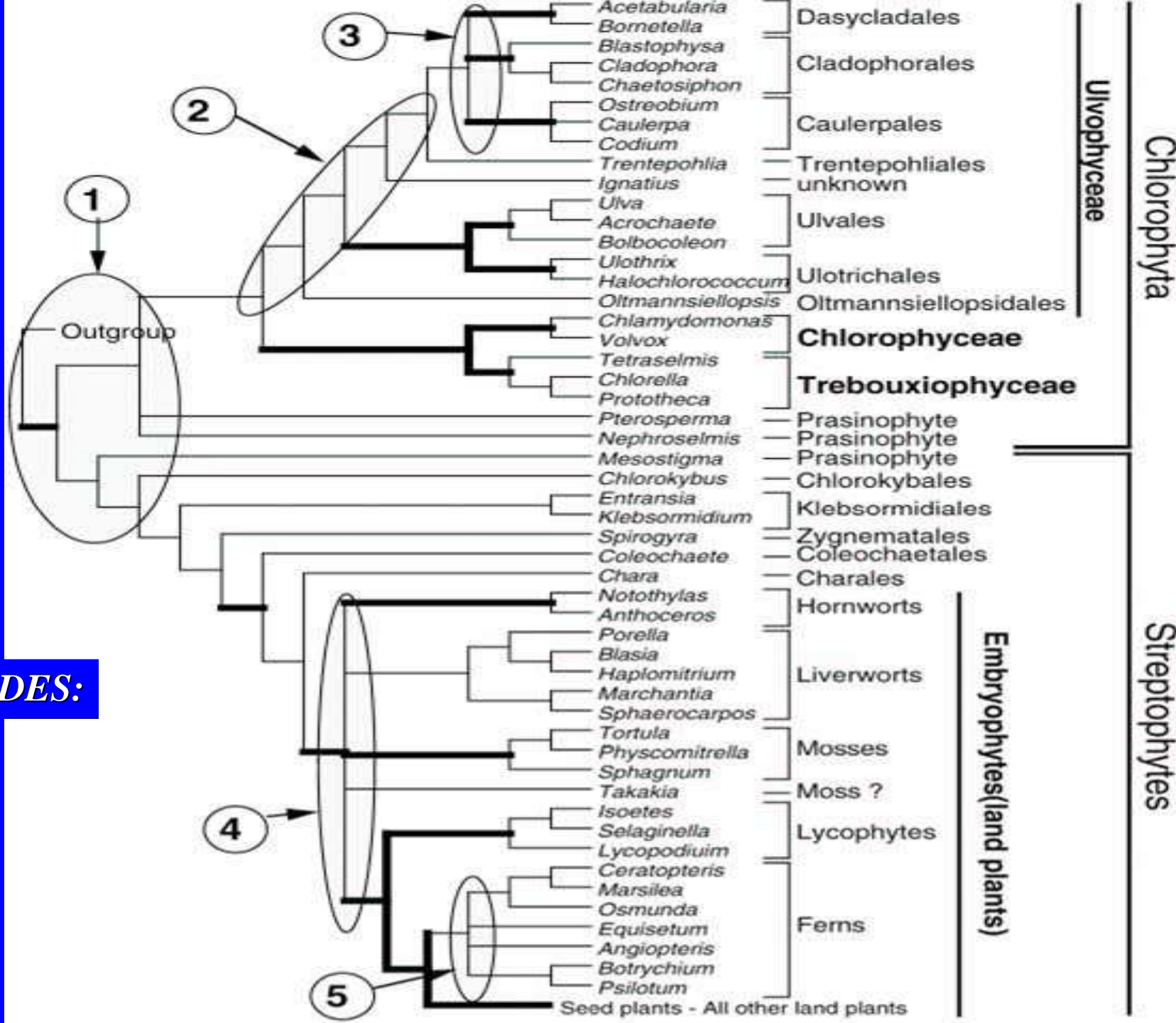
West Boothbay Harbor, Maine





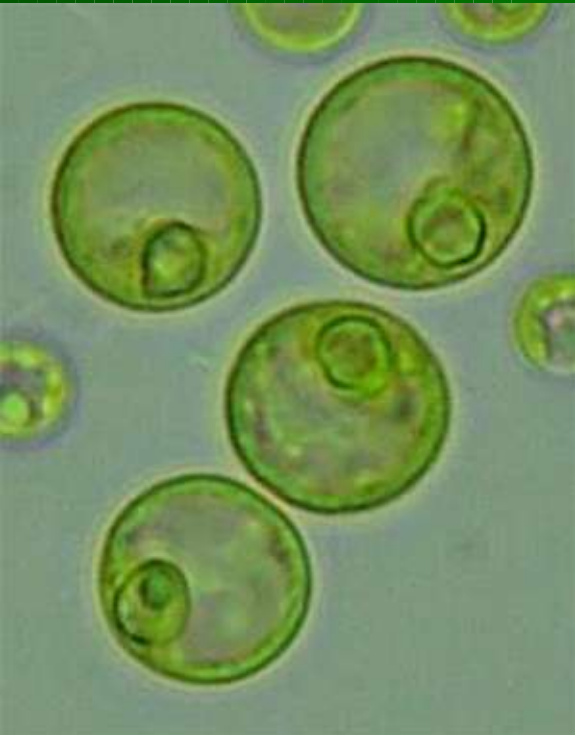
the green tree of life

UNRESOLVED NODES:



- CCMP 1293

- Coccoidal morphology, resembles *Chlorocystis*
- Ulotrichales ultrastructure
- Disjunct gene sequences (18s rRNA, *tufA*)



Ulvophyceae: One class or two?

- *Blastophysa* spp.
 - Cladophorales morphology
 - Cladophorales ultrastructure
 - Disjunct 18s sequence
 1. Sister to 1293
 2. Basal to Cladophorales



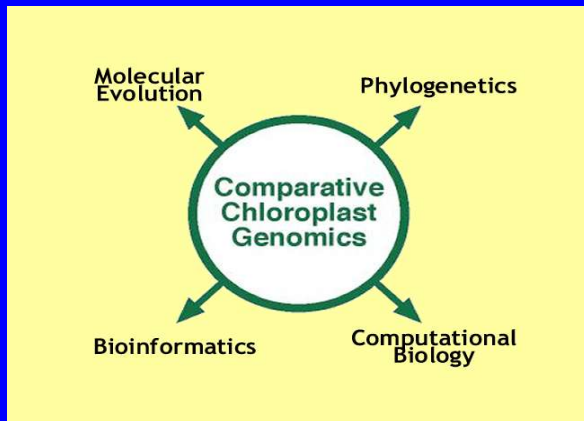
- Discontinuous morphological/molecular evolution

The Deep Green “family”

Some of the projects that have been funded as part of the “Deep Green” tradition of research cooperation.



DEEP TIME



deepestgreen





The Green Tree of Life (GToL):

“Deep Green Plant Phylogenetics: Novel Analytical Methods for Scaling from Genomics to Morphology”

TOWARD RESOLUTION OF GREEN PLANT PHYLOGENY



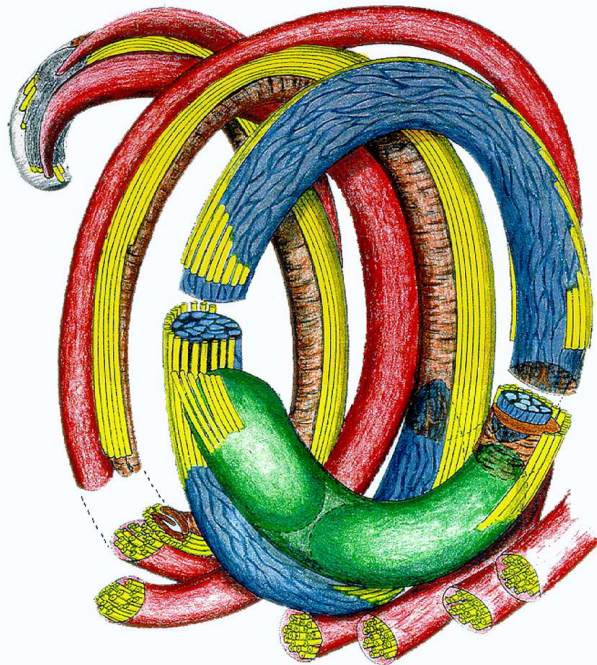
→ { O'Kelly 0228655
Donoghue 0228576
Mandoli, Olmstead 0228660
Mishler, Smith, Boore 0228729
Renzaglia 0228679
Wolf 0228432

Bigelow Lab for Ocean Sciences
Yale University
University of Washington
University of California, Berkeley
Southern Illinois University, Carbondale
Utah State Univ.

Goals:

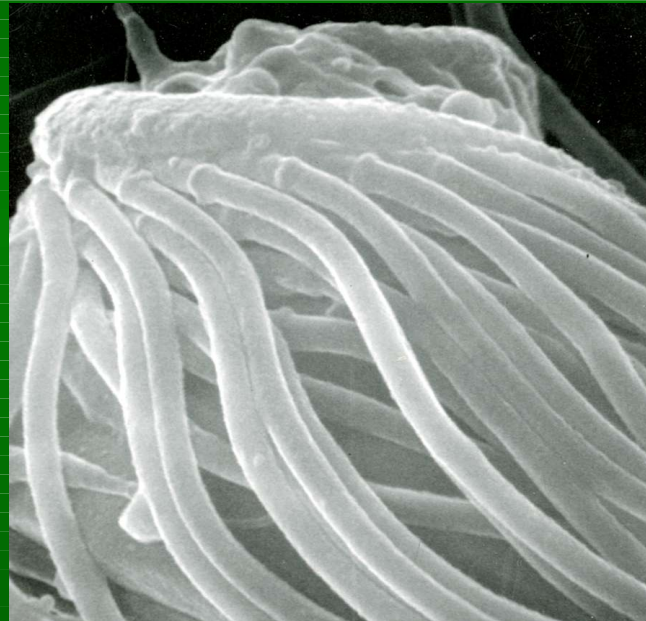
- * complete a matrix of whole genome sequences for chloroplasts and mitochondria and develop Bacterial Artificial Chromosome (BAC) nuclear genome libraries (where feasible given genome size) for ca. 50 representatives of the critical deep-branching lineages of green plants.
- * produce a comprehensive set of comparable morphological and ultrastructural data for these same taxa;
- incorporate inferences from across the phylogenetic hierarchy in green plants using methods designed to permit scaling across studies.

Morphological/ Ultrastructural Data: Spermatozoid Structure and Development in Embryophytes



Selaginella australiensis

Renzaglia



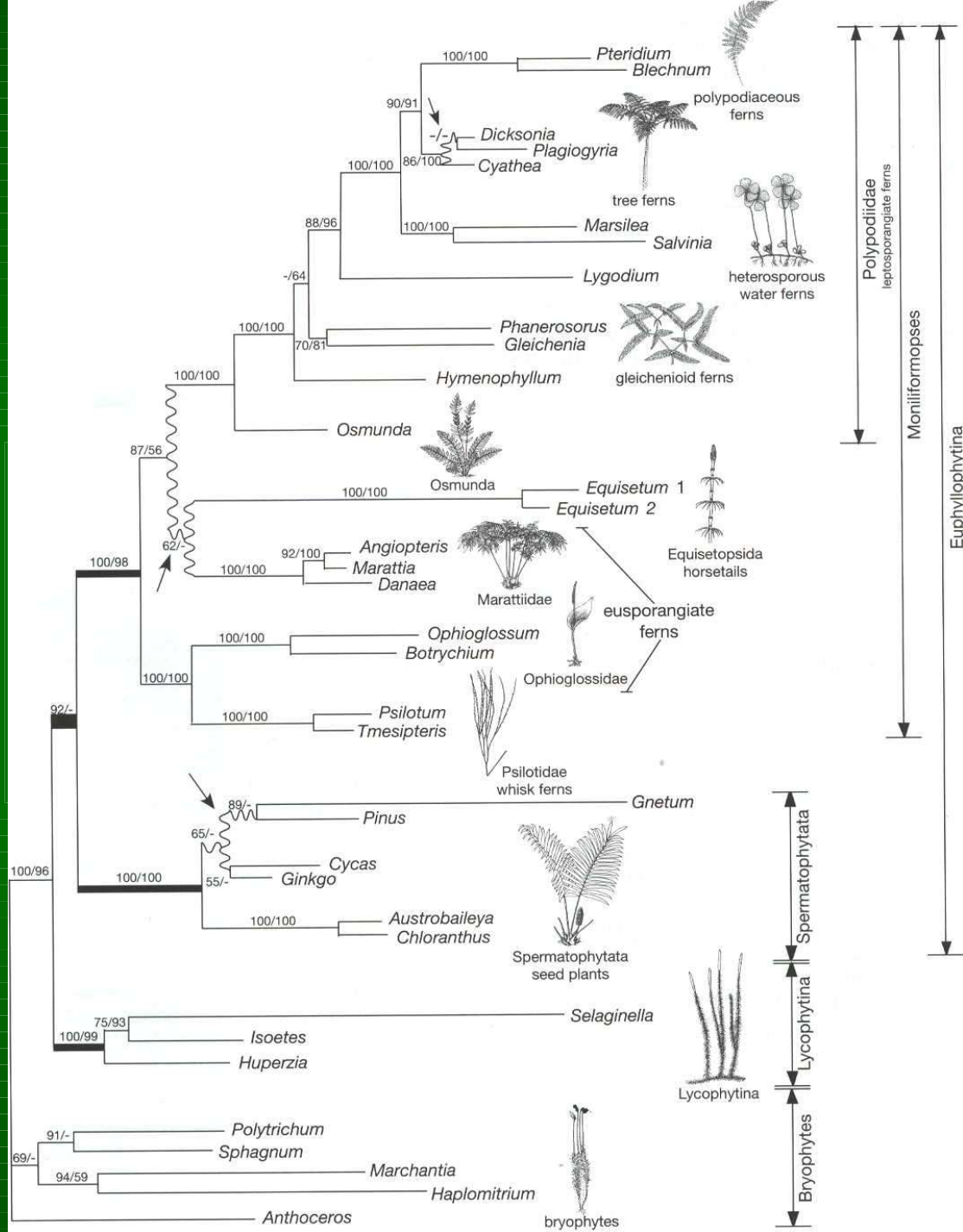
Bryophyte morphology: Teacher
Workshop (2003 BSA) and Teaching
Materials development.

Huperzia lucidula (*Lycopodium lucidulum*)



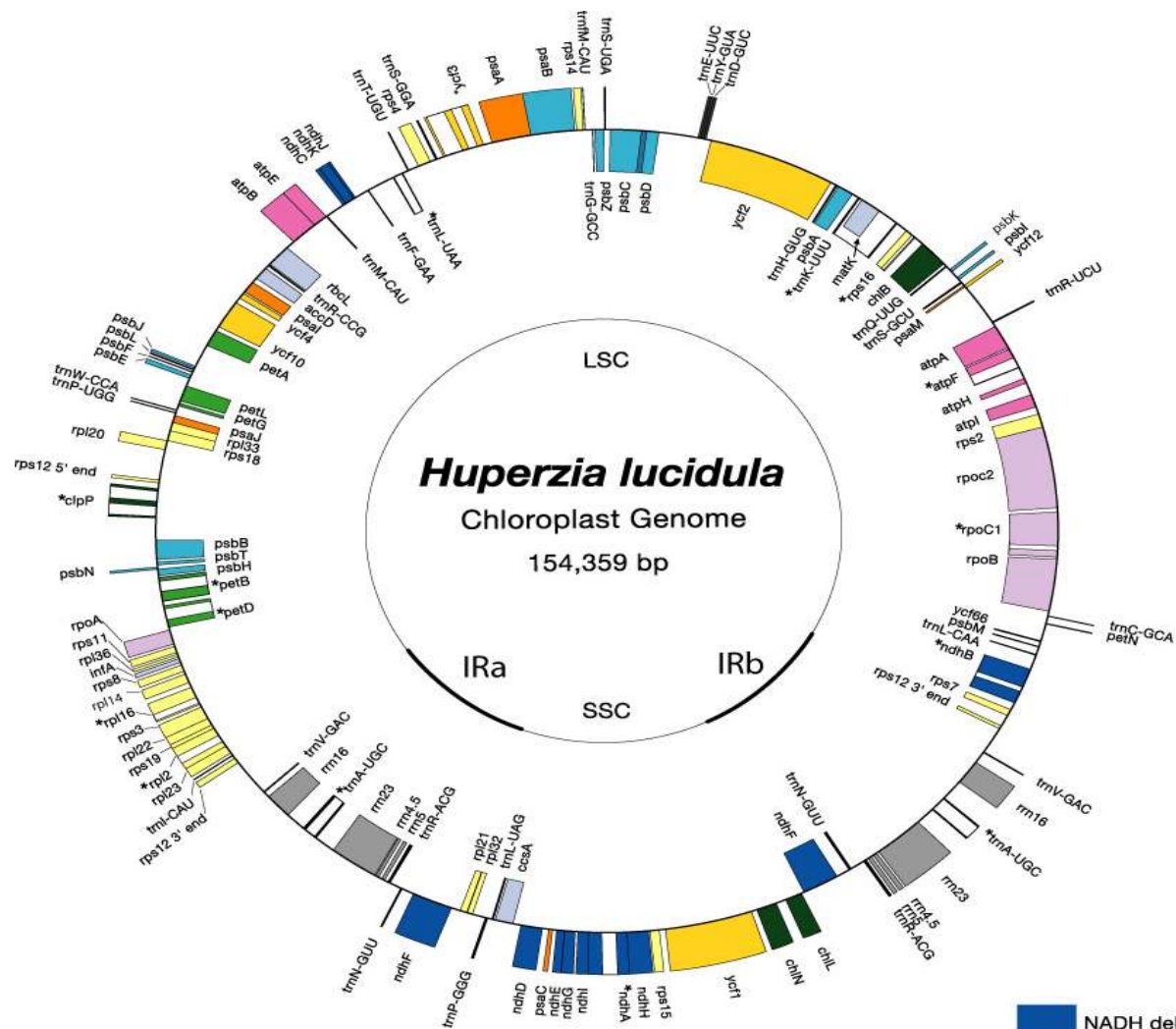
The dissection of *Huperzia*

- Specimen collected and identified (Wolf)
- Specimen vouchered (Mishler - Berkeley herbarium)
- Morphological and ultrastructural characterization (Renzaglia)
- Genomic analysis



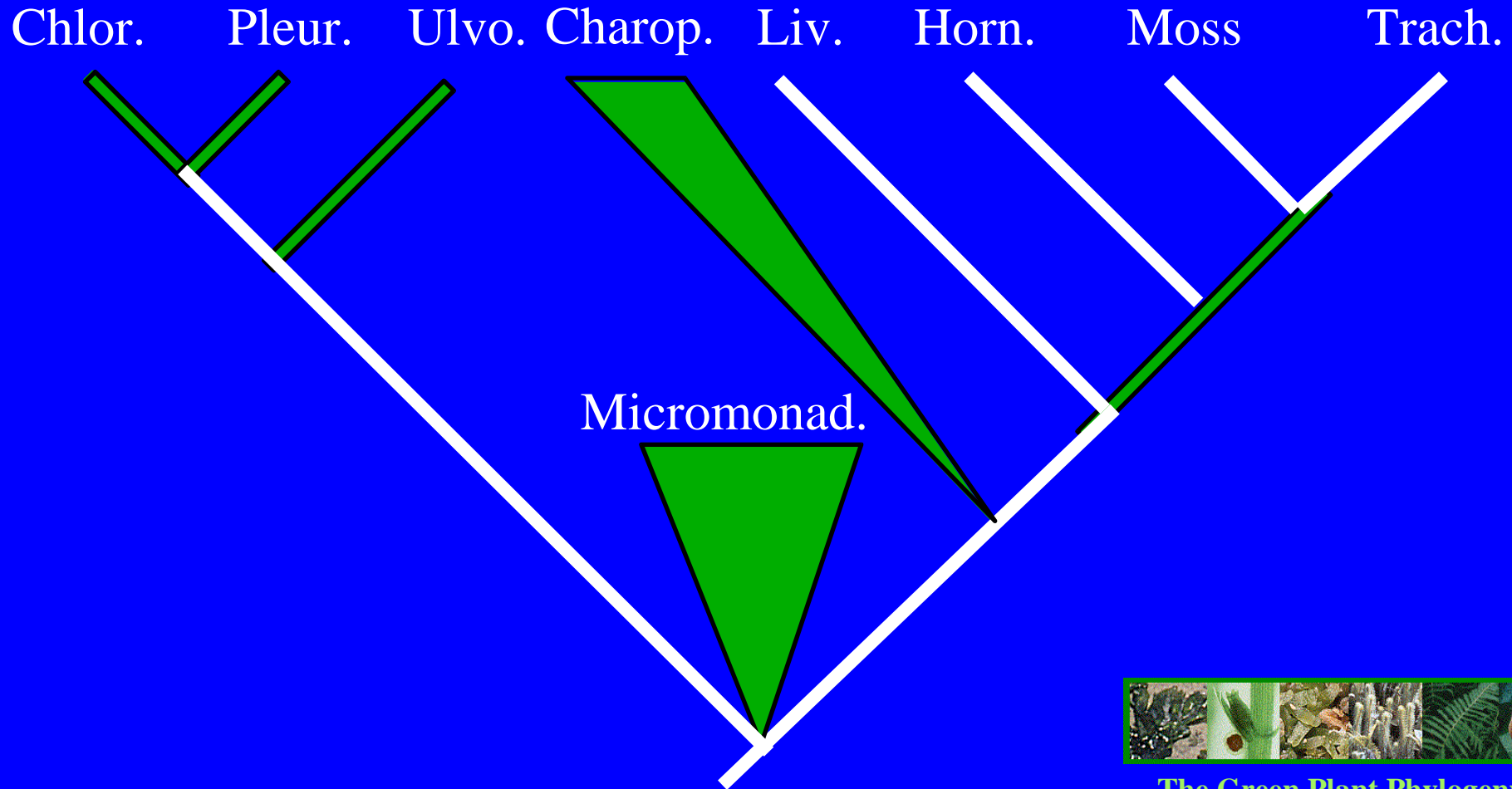
The story continues ...





- NADH dehydrogenases
- RNA polymerases
- Hypothetical proteins
- Chlorophyll biosynthesis
- Photosynthesis II
- Photosynthesis I
- Ribosomal RNAs
- Ribosomal proteins
- Cytochrome bb/f complex
- ATP synthases
- RNA polymerases

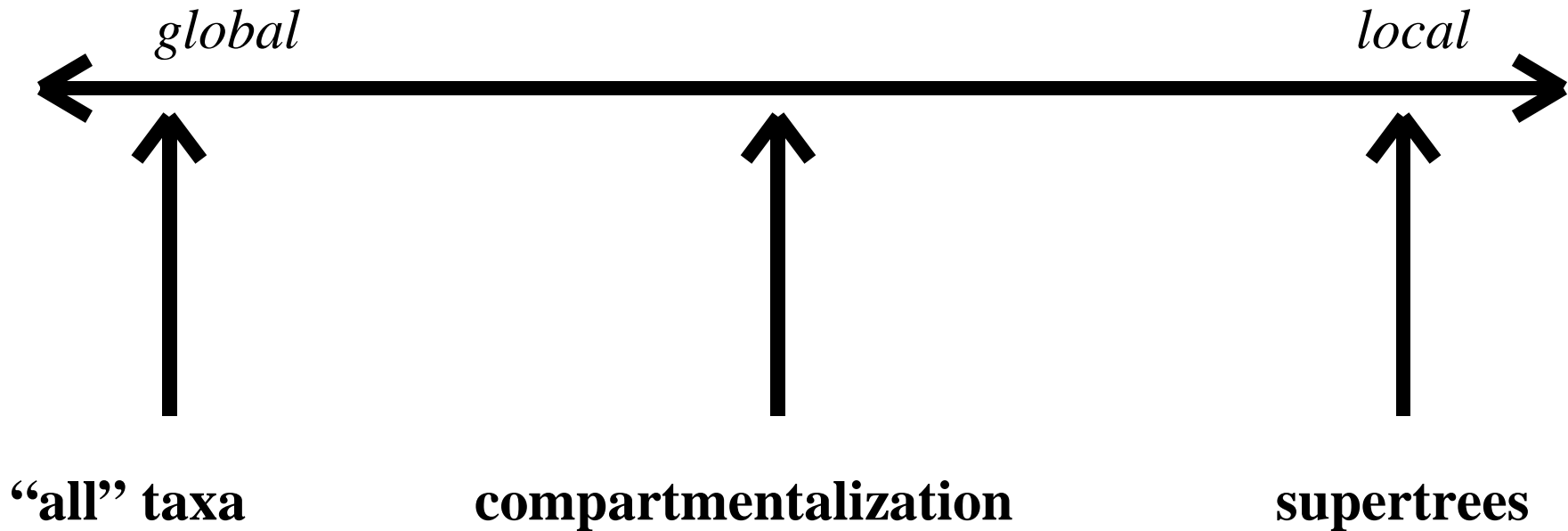
Outline of Green Plant relationships



The Green Plant Phylogeny
Research Coordination Group

<http://ucjeps.berkeley.edu/map2.html>

HOW TO CONCATENATE DIFFERENT ANALYSES TO BUILD THE TREE OF LIFE



HAPLOPHASE

DIPLOPHASE

