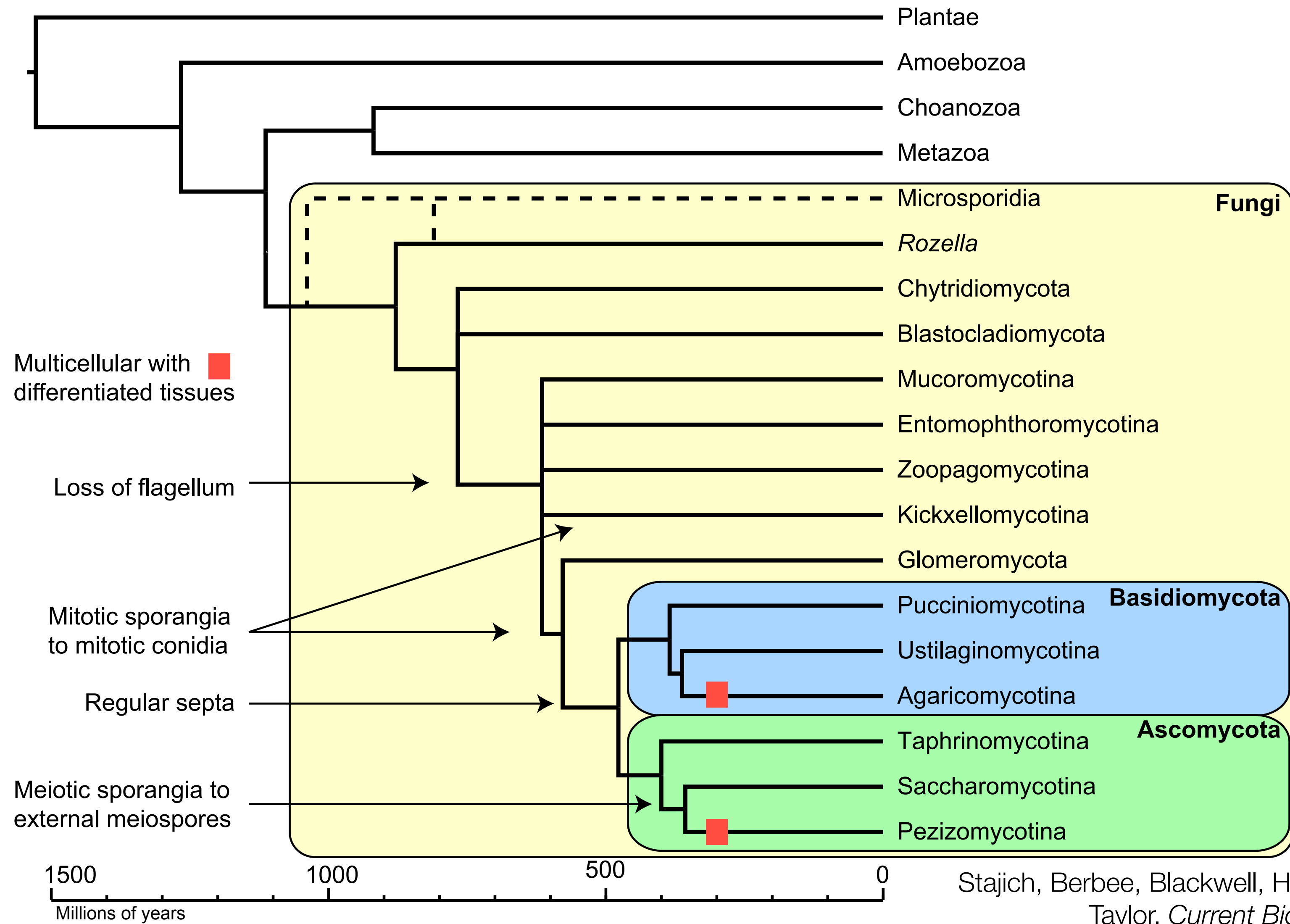


Connecting ultrastructure to molecules: Studying evolution of molecular components of cellular division in fungi from comparative genomics.

Jason Stajich
University of California, Riverside

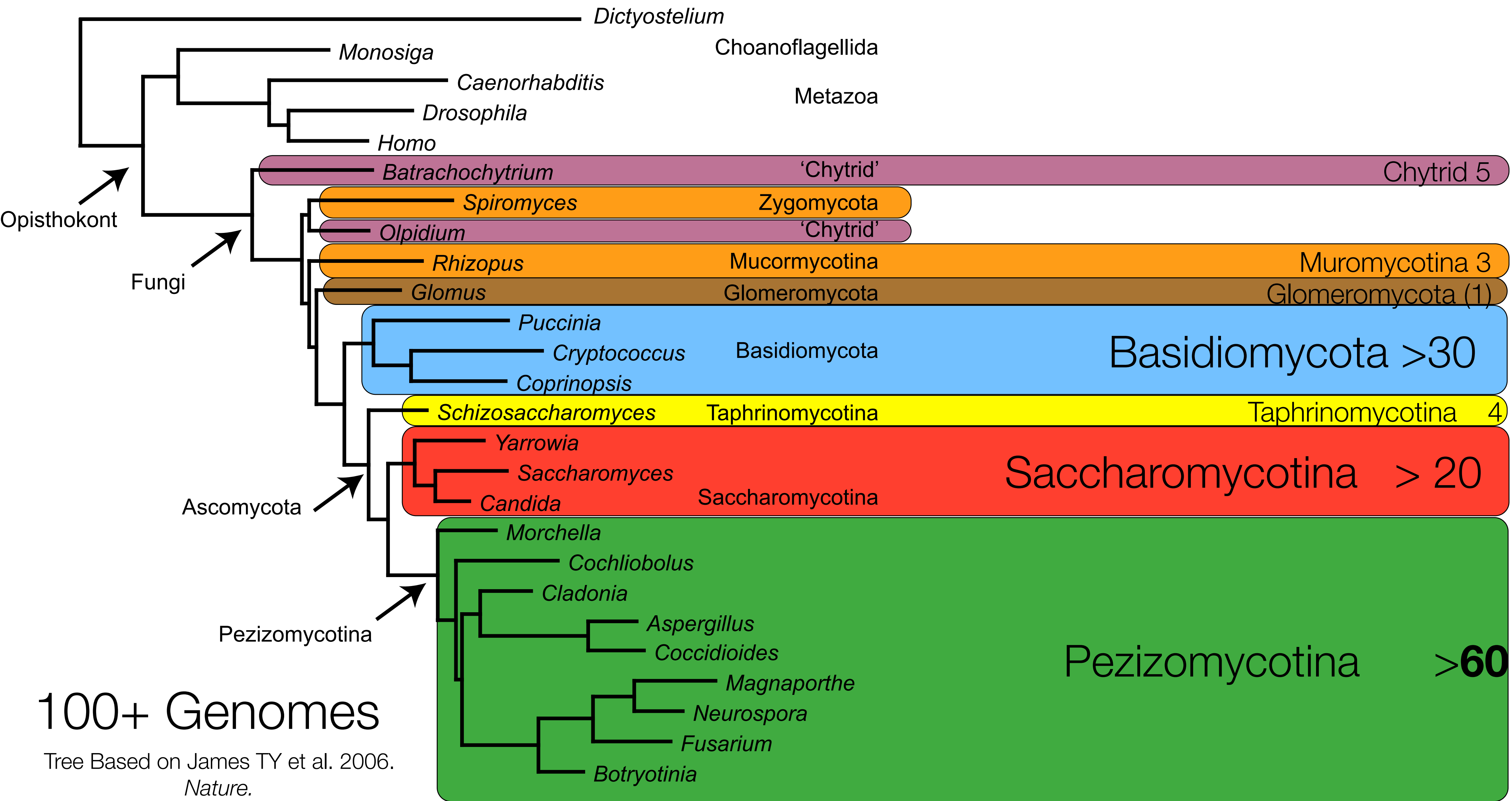
Evolutionary transitions in fungal cellular structure



Connecting evolutionary transitions to molecules

- Comparative genomics can help inventory the genes shared or missing between species
- Tracing the evolution from a flagellated, aquatic ancestor to filamentous and
- Using genome sequences from Chytridiomycota, Mucormycotina, and Dikarya fungi can compare genes that are found only in Dikarya fungi and linked to the clade-specific phenotypes
- 3 transitions illuminated
 - Cellular division
 - Cell wall composition
 - Septa

Genome samples from fungi

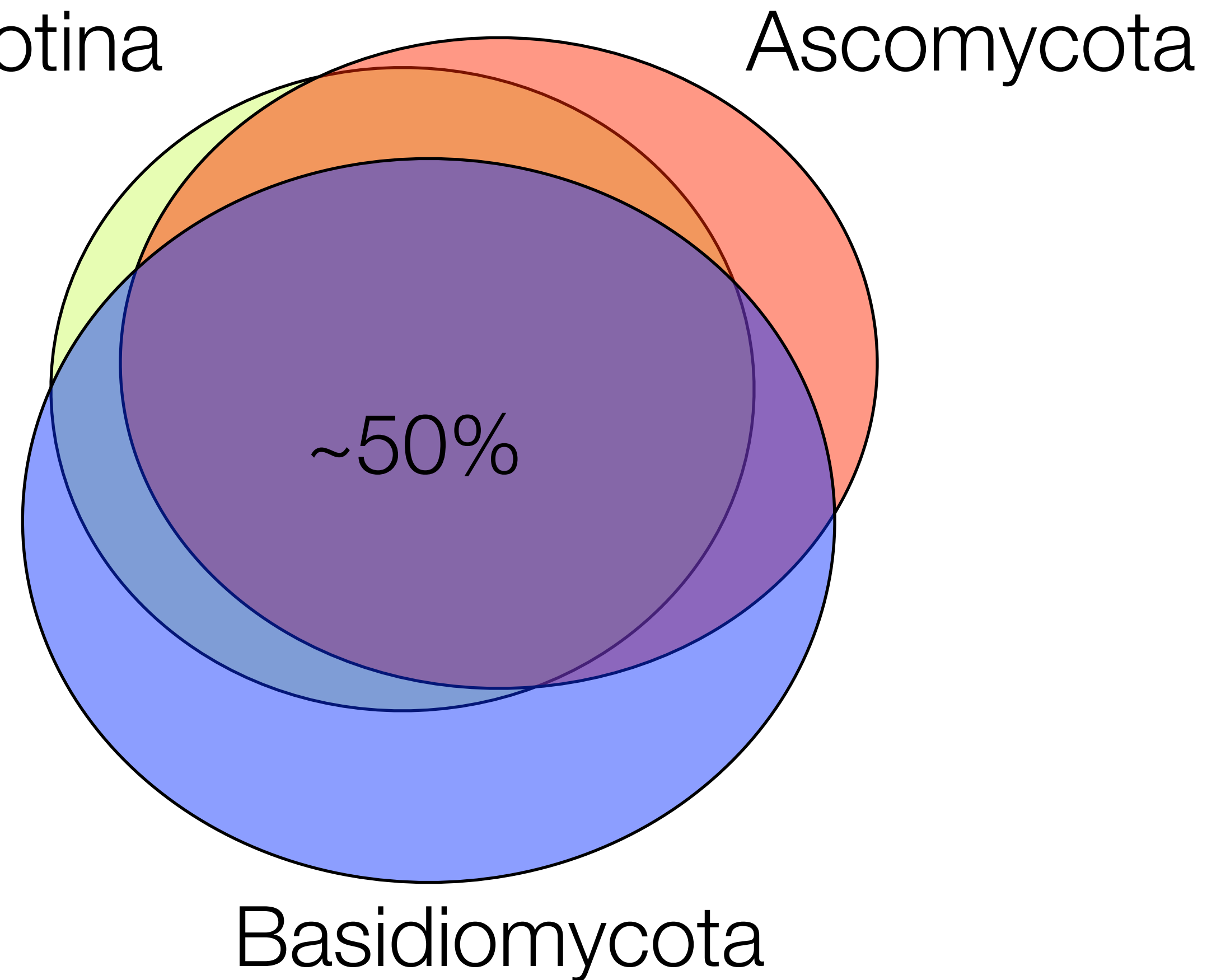


B. dendrobatidis genes shared with other fungi

Via all-vs-all similar searches using
FASTP and 1e-5 E-value cutoff

Using genomes from 30 fungi and the
B. dendrobatidis genome

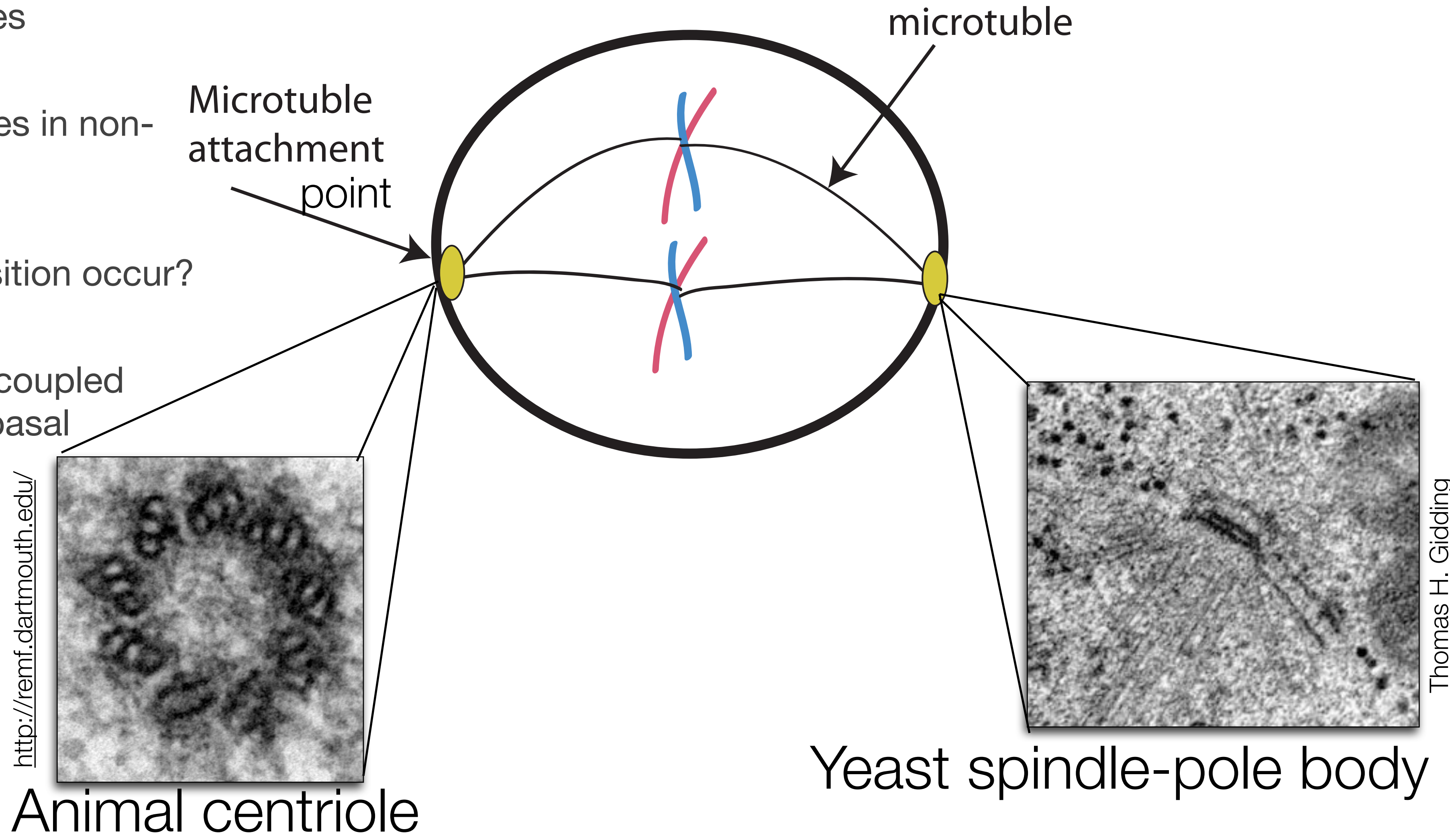
Identify the shared genes



~8800 genes in *B. dendrobatidis*

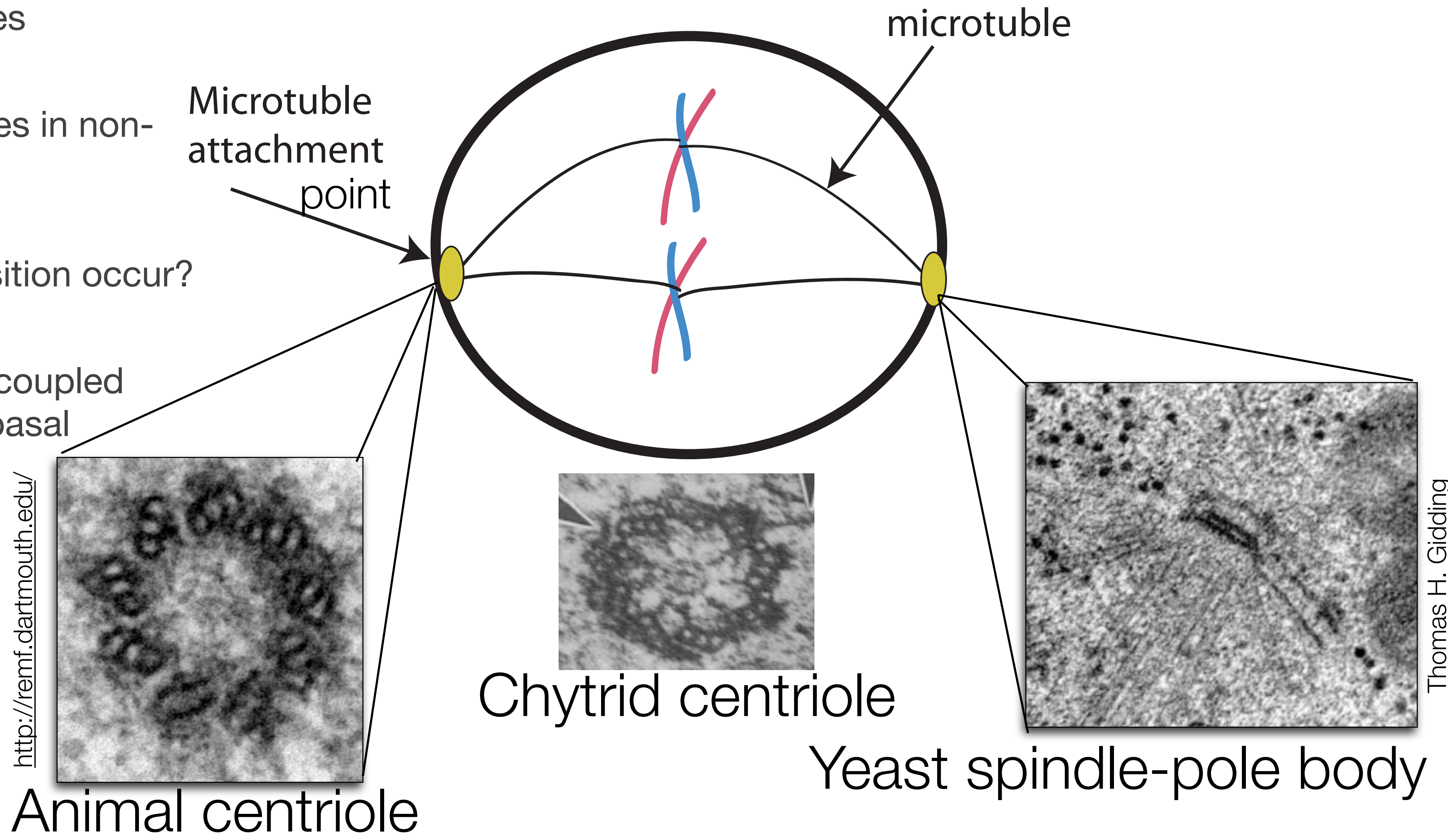
Differences between animal-like and fungal Microtubule attachment points

- Transition from flagellated cells with centrioles
- Spindle pole bodies in non-flagellated fungi
- How did this transition occur?
- Loss of flagellum coupled with transition in basal bodies?



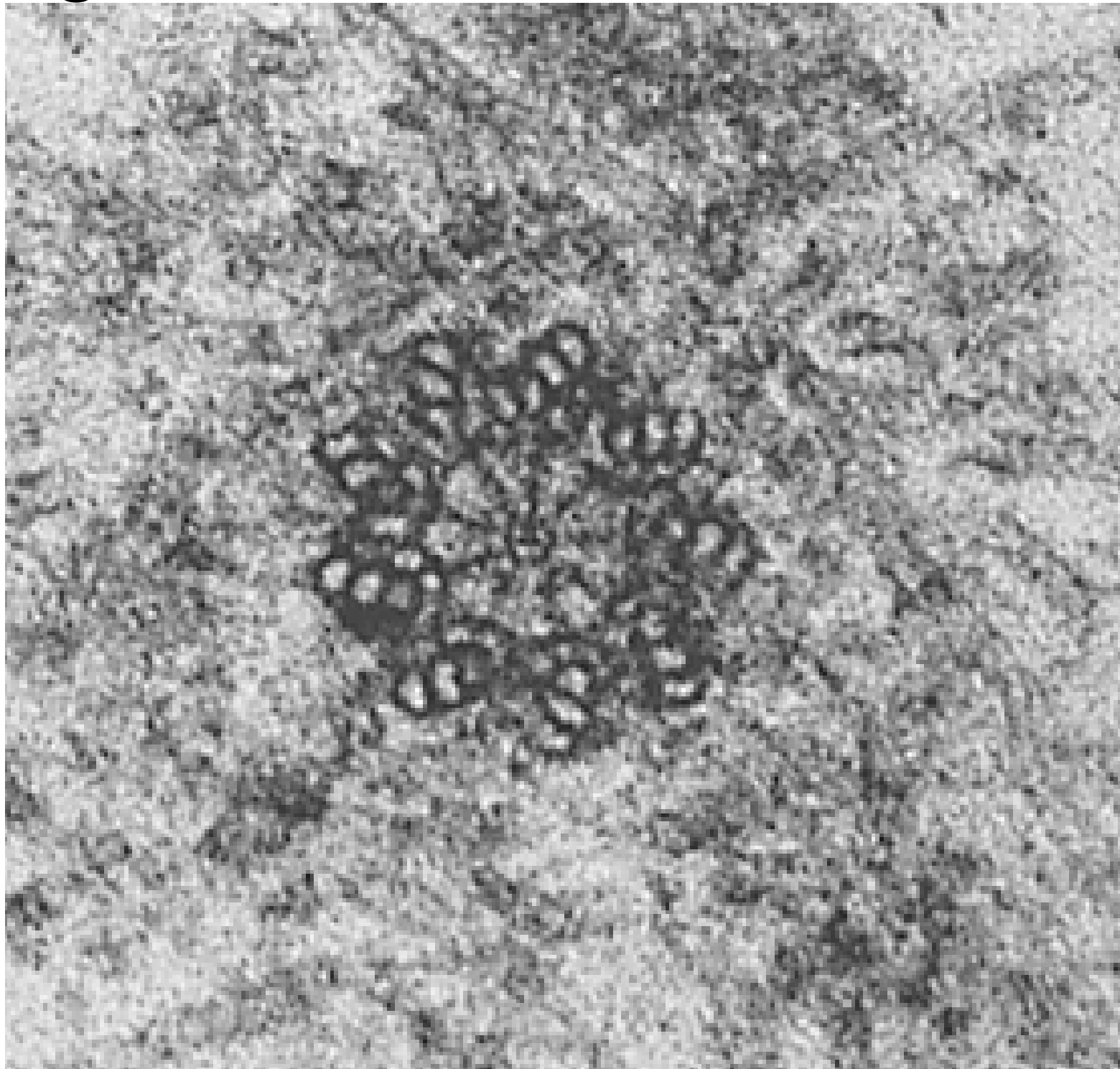
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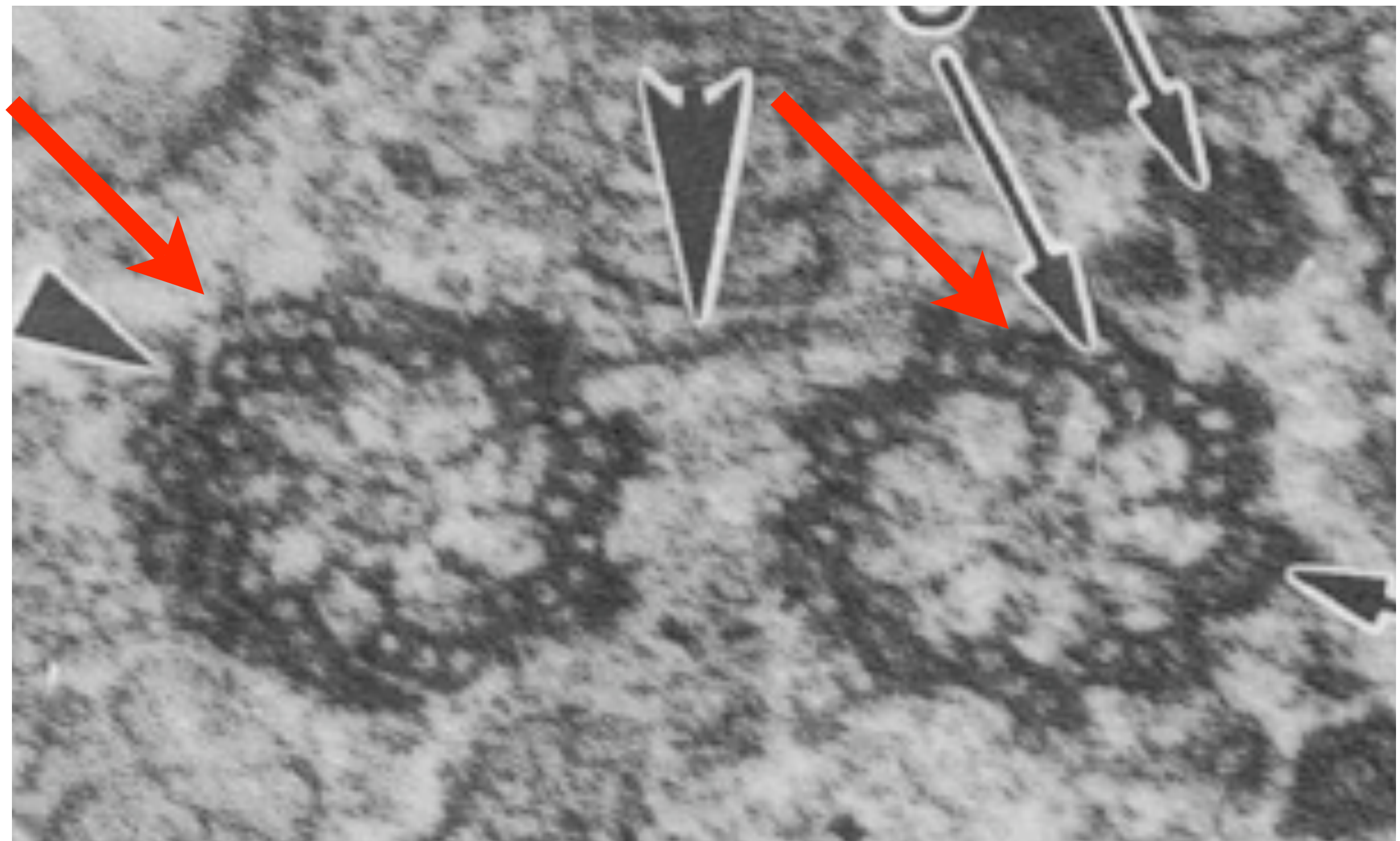
Centrioles in animals and chytrids are homologous structures

Cross-section of a rat kangaroo (Marsupial, Animalia) centriole region



Berns MW et al. *J Cell Biol* 1977

Chytrid zoospore cross section with 2 centrioles



Entophlyctis luteolus

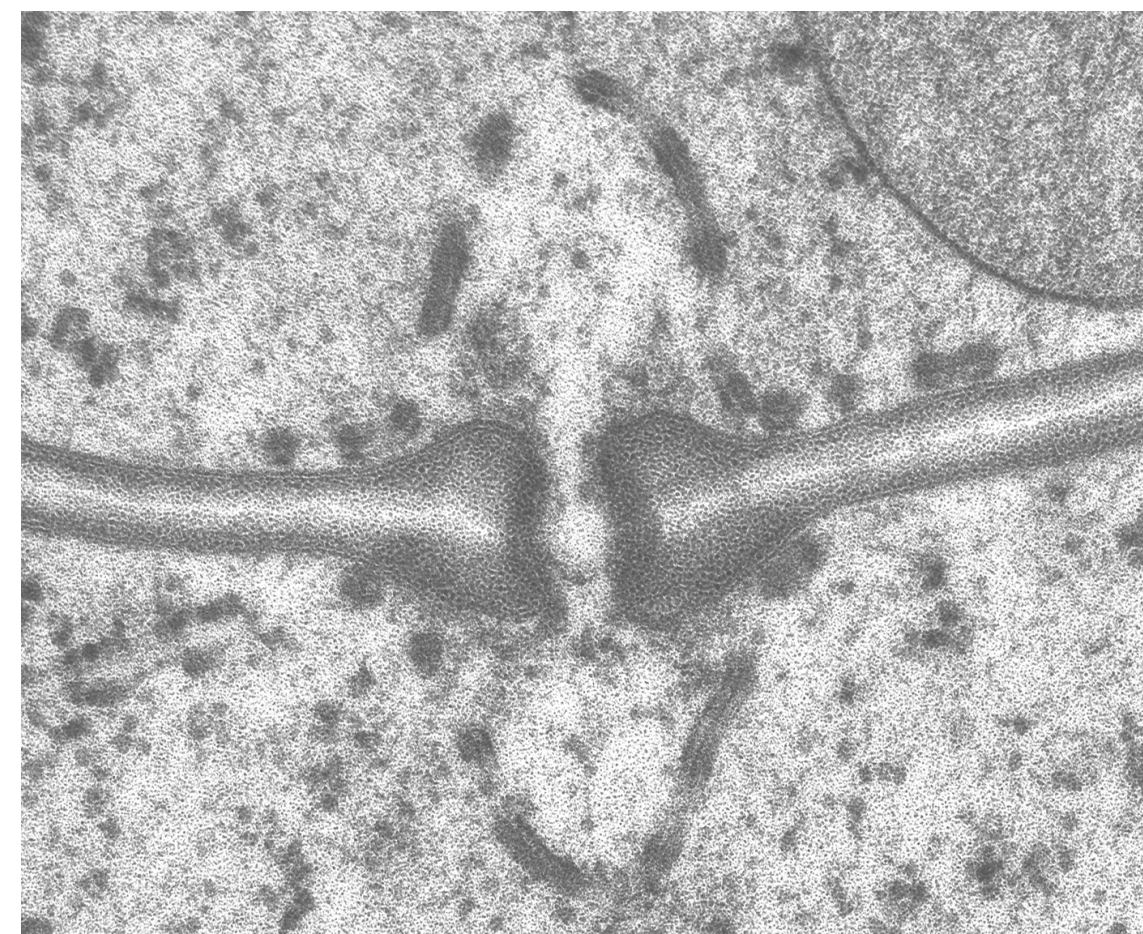
Longcore. *Mycologia* 1995

Chytrid genome is missing some chromosome segregation and mitosis related genes

- Missing genes in the *B. dendrobatidis* genome
- MSP3, KAR1, KAR2 for nuclear membrane fusion during karyogamy and Spindle-body duplication
- SPC42 - central plaque component of spindle pole bodies
- CEP3 - essential kinetochore protein
- CIN2 - Tubulin folding protein
- REC8 for sister chromatid cohesion
- DASH Complex for kinetochores coupling during mitosis (10 genes)

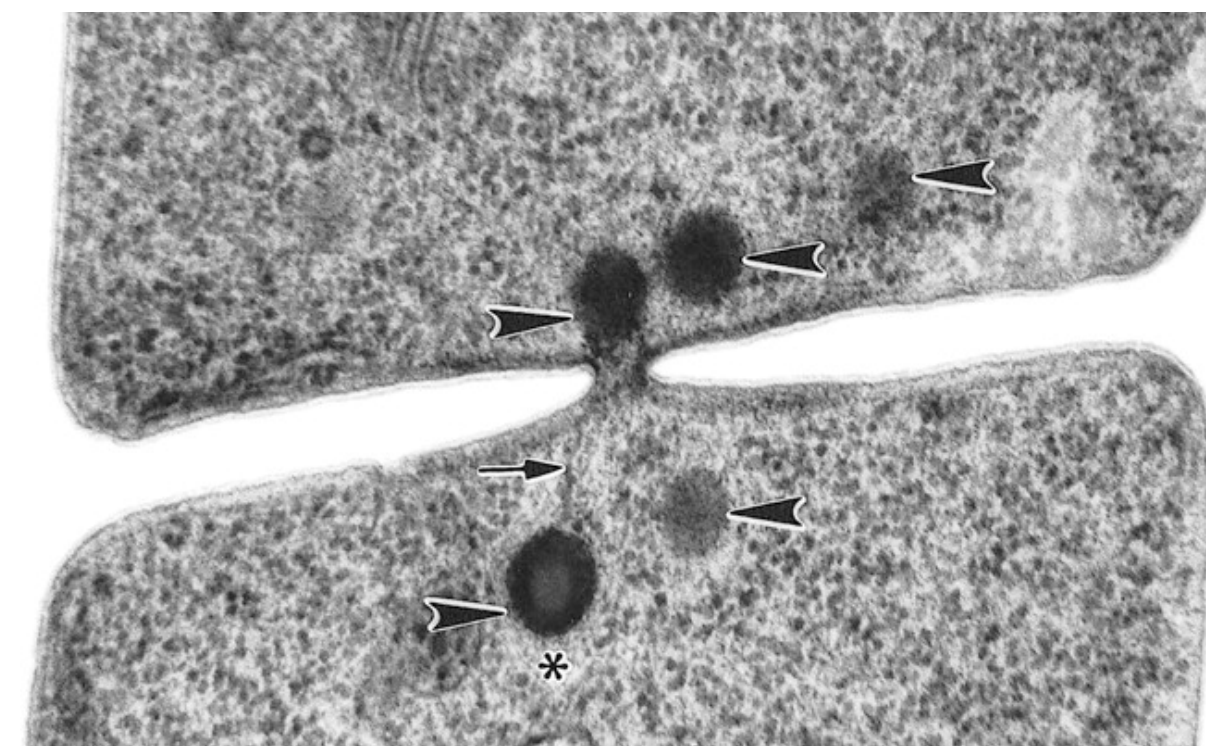
Septa

- Septa allow for separation of cells, isolate cytoplasm, allow for cellular differentiation.
- Regular septa in the Dikarya, but septa do exist in some pre-Dikarya
- Septa observed in the Kickxellomycotina (*Coemansia*)
- All 6 septin genes identified in *Aspergillus* are found in early fungi (*B. dendrobatidis*)



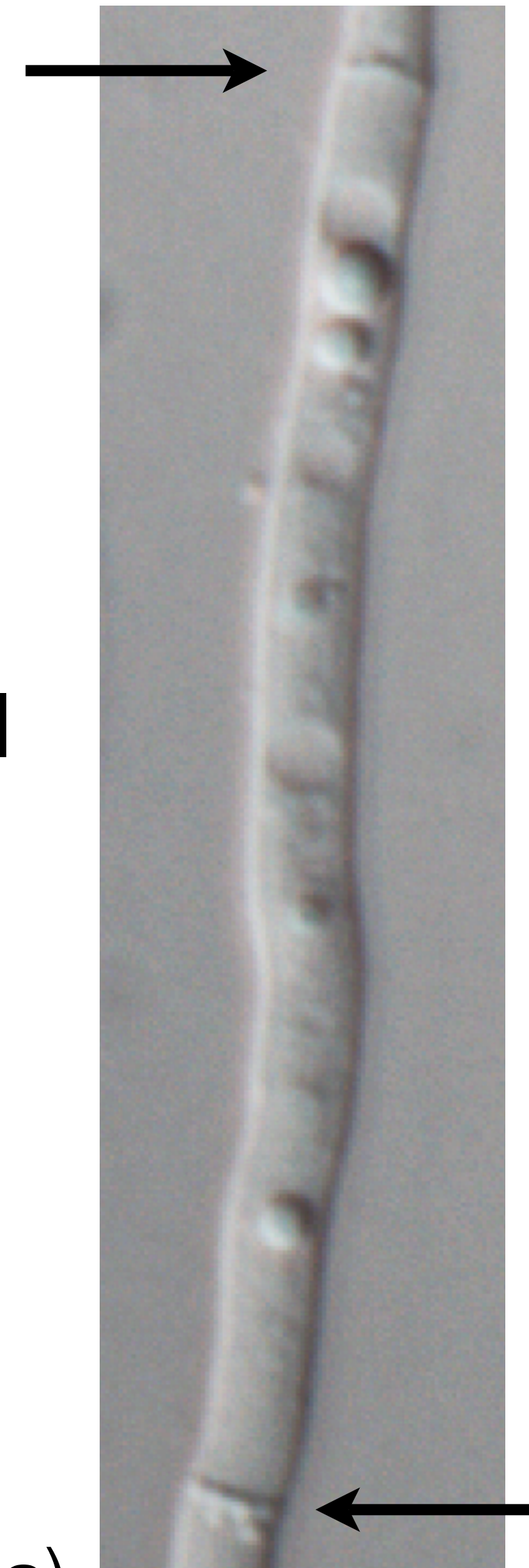
Auriscalpium vulgare
(Agaricomycotina) hyphal
septum.

Celio et al. Mycologia. 2007



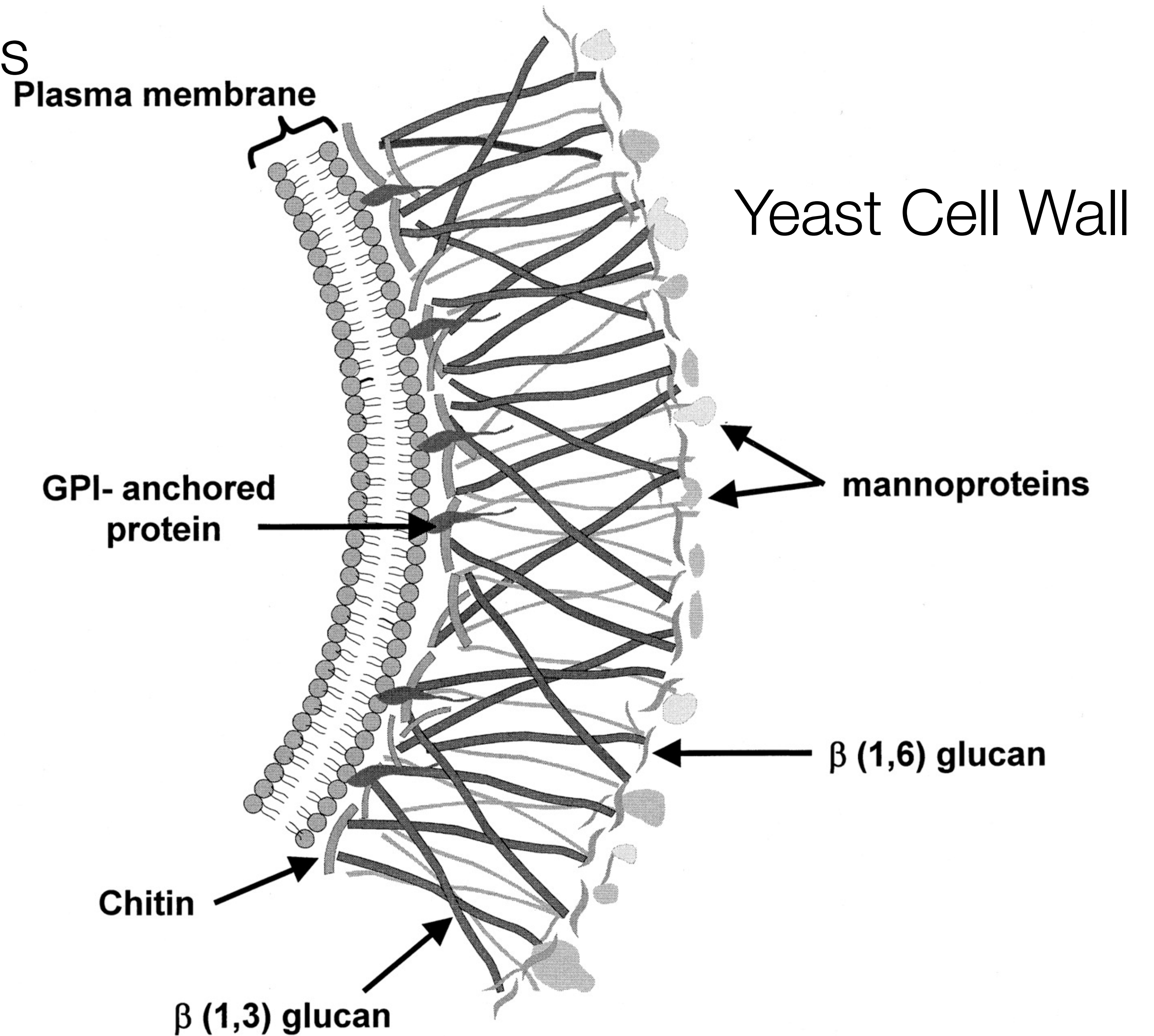
A. nidulans (Pezizomycotina)
septum and woronin body.

Momany et al. Mycologia. 2002



Coemansia

Fungal cell wall changes



Cell wall changes from Early fungi

- Looking at the proteins known to make up cell wall in well studied yeast *Saccharomyces cerevisiae*
- Identified series of genes missing in Bd including major components of the 1,3- β glucan biosynthesis pathway (FKS1) and many of the KTH transporters
- With collaborators (JP Latge) we showed through a simple cell wall composition assay
 - Missing $\beta(1,3)$ - & $\beta(1,6)$ -glucans but contain $\beta(1,4)$ - glucan: cellulose.
 - Few "higher" fungi have been documented to have cellulose in cell walls (Ascomycota)
- Bd is much less sensitive to Enchinocandin drugs which target FKS1
- Does gain of genes to make $\beta(1,3)$ -glucan and/or loss of cellulose represent one of the major transitions from aquatic to terrestrial lifestyles for fungi?

Connecting molecules to evolutionary transitions

- Nuclear division changes
 - Basal body, centriole loss coupled with loss of flagellum
 - What happened first? Where did spindle-pole body genes evolve from - duplication or novel gene creation?
Did some of flagellum & basal body genes evolve new functions?
- Multicellular development
 - Differentiation of multicellular tissue, septation, and signaling
- Cell wall changes
 - 1,3 Beta-glucan and 1,6 Beta-glucan synthesis evolved after Chytrid divergence
 - Was this a necessary transition with the aquatic to terrestrial life changes?

Identifying an ancient photoreceptor

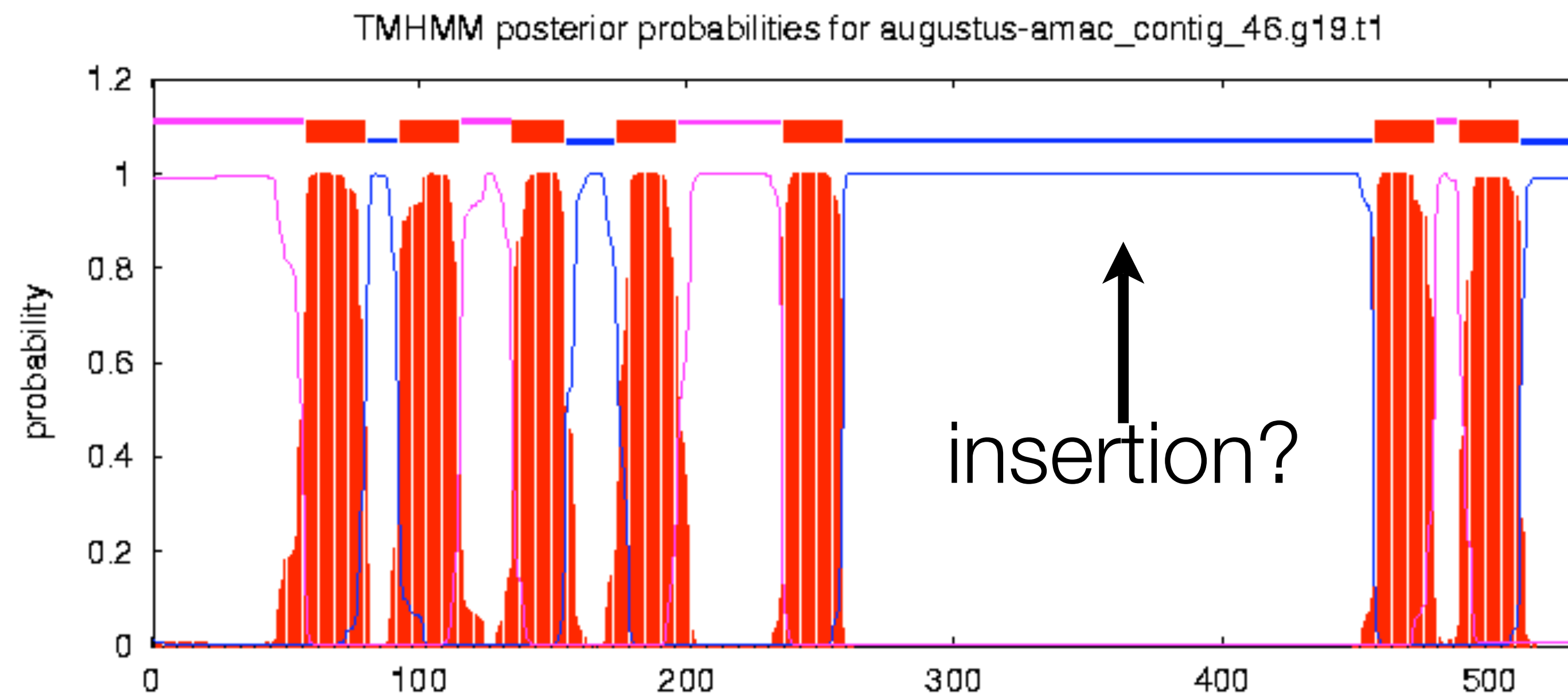
- Fungi have some light sensing molecules but no known rhodopsins have been cloned.
- Rhodopsin molecule implicated in zoospore phototaxis in *Allomyces* (by comparing wavelengths where cells responded)
- Sequence similarity identifies a candidate 7-transmembrane protein, but has insertion found in both *Bd* and *Allomyces* (draft genome).

Rhodopsin guides fungal phototaxis

Rhodopsins act as the visual pigments of multicellular animals and also guide the swimming of green algae¹ towards or away from light (phototaxis). If these rhodopsin molecules have a common evolutionary origin, fungi, the sister group of animals², would also be expected to use a rhodopsin-based photoreceptor. Here we show that rhodopsin guides the zoospores of the fungus *Allomyces reticulatus*³ towards the light, suggesting that the origin of vision might have been the phototaxis of their unicellular ancestors.

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Syracuse University, Syracuse,
New York 13244-1130, USA*

NATURE | VOL 387 | 29 MAY 1997



Transitions inferred from genome comparisons

AFTER CHYTRID SPLIT

Loss of:

- Flagella, centrioles
- **1,4 Beta-glucan synthesis**

Gain of:

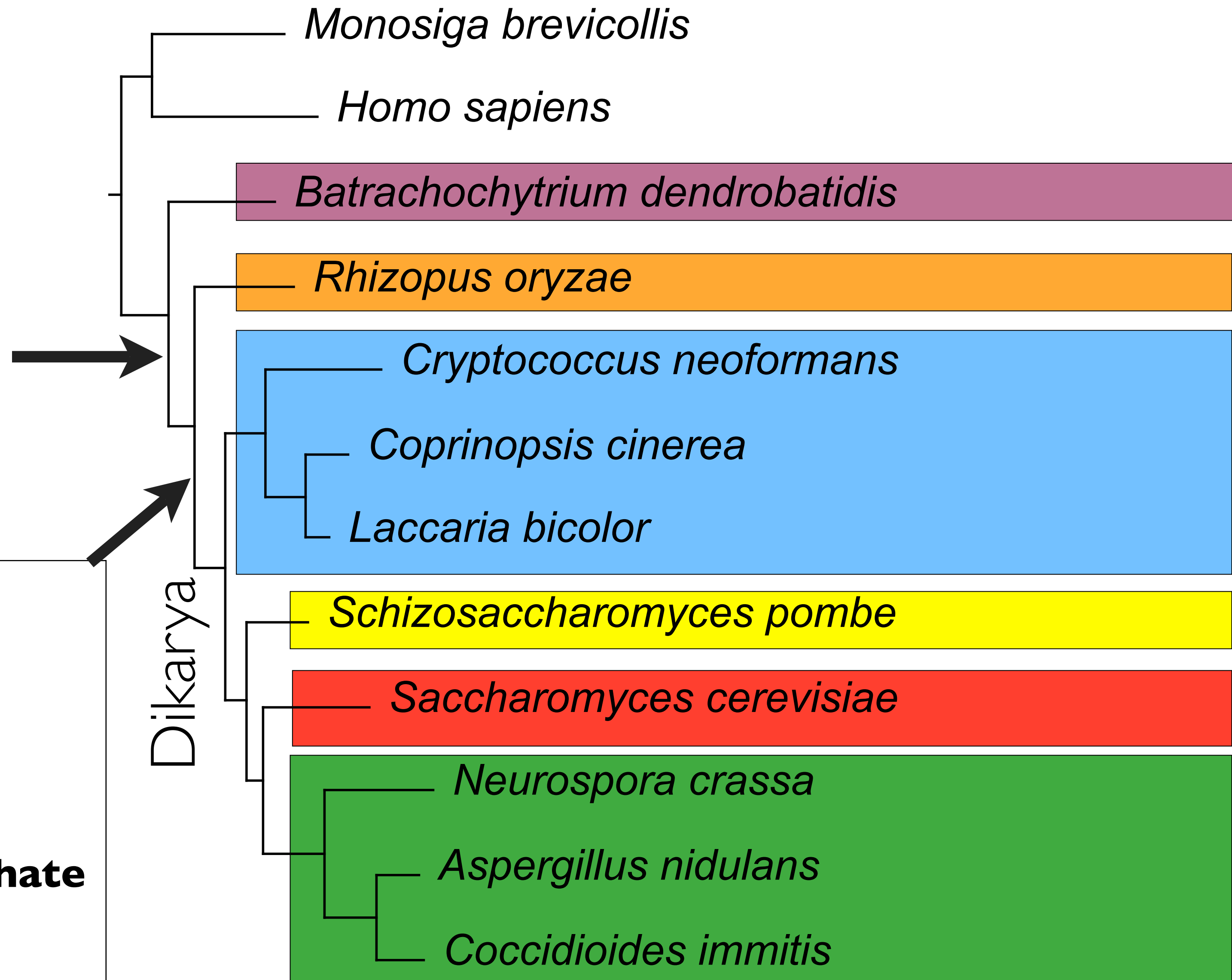
- STE50 adaptor
- Some glucan synthase and transferases
- **Rhodopsin-like 7TM**

IN DIKARYA, Gain of:

- Spindle-pole bodies
- 1,3 Beta glucan synthase
- Thiamine biosynthesis
- STE3 pheromone receptor
- **DASH complex**
- **KTR mannosylphosphate family**

Loss of:

- Chitosanase

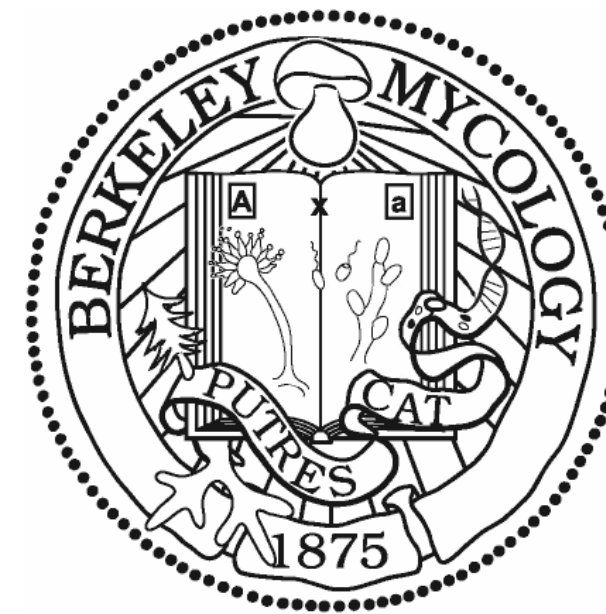


Thanks

John Taylor & Laboratory (Univ California, Berkeley)

Mary Berbee (Univ British Columbia)

UC Berkeley Imaging facility & Barbara Waaland



Miller Institute

B.dendrobatidis sequencing

Joint Genome Institute & Broad Institute

<http://fungalgenomes.org/blog>

New research starting at UC Riverside

- Interested in research in fungal genomics?
 - Evolution of fungal development
 - Post-transcriptional gene regulation and small RNAs
 - Fungal cell evolution in early diverging chytrid and 'zygomycete' fungi
- Bioinformatics and genome informatics of fungal comparative genomics

