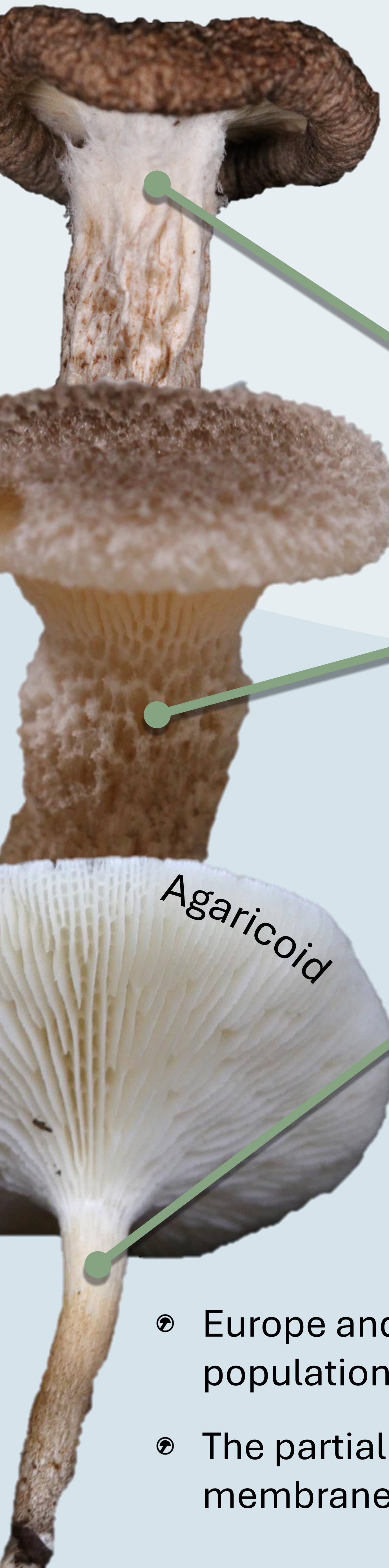


Lentinus tigrinus Genomes, Species Limits, and Development

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Biological Species Concept



Incompatible with:

- ☉ *L. squarrosulus*
- ☉ *L. sajor-caju*
- ☉ *L. crinitus*

Europe

- ☉ Agaricoid
- ☉ Partial veil
- ☉ Grows near water

Europe x N. Amer.

- ☉ Cross with secotioid strain
- ☉ Agaricoid F1
- ☉ Incomplete partial veil in F1

North America

- ☉ Agaricoid or secotioid
- ☉ No partial veil
- ☉ Grows near water

- ☉ Europe and N. America *L. tigrinus* populations are mating compatible¹
- ☉ The partial veil and secotioid membrane are distinct structures

Improving Genome Assembly: In Progress

Existing Genomes

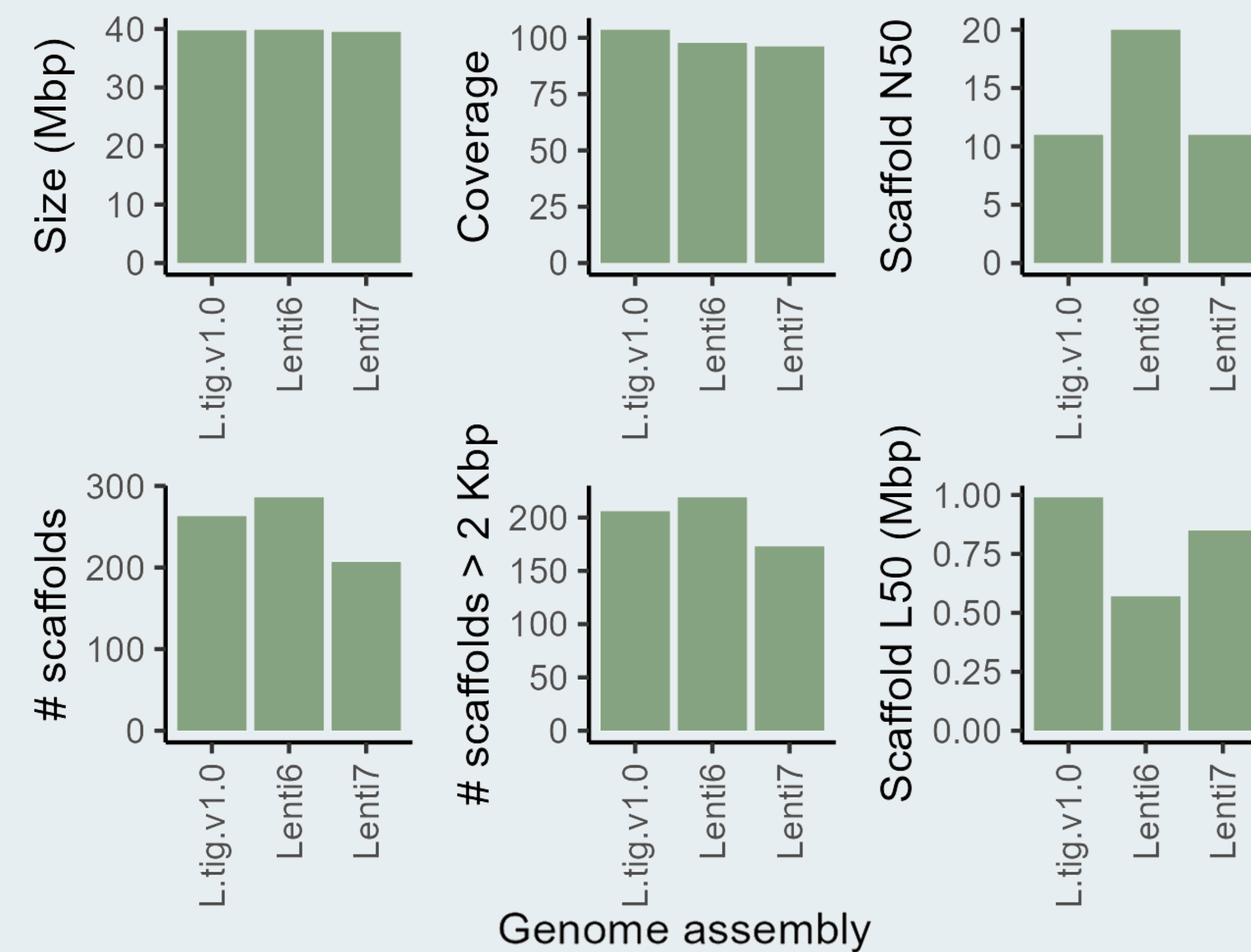


Fig. 1. Comparison of basic statistics of three *L. tigrinus* genomes available through MycoCosm.^{2,3}

Goals

- ☉ PacBio sequencing to improve genome assembly
- ☉ Conduct GWAS to identify locus (Q1)
- ☉ Assess population structure to determine species limits and selection (Q2/Q3)

Developmental Biology

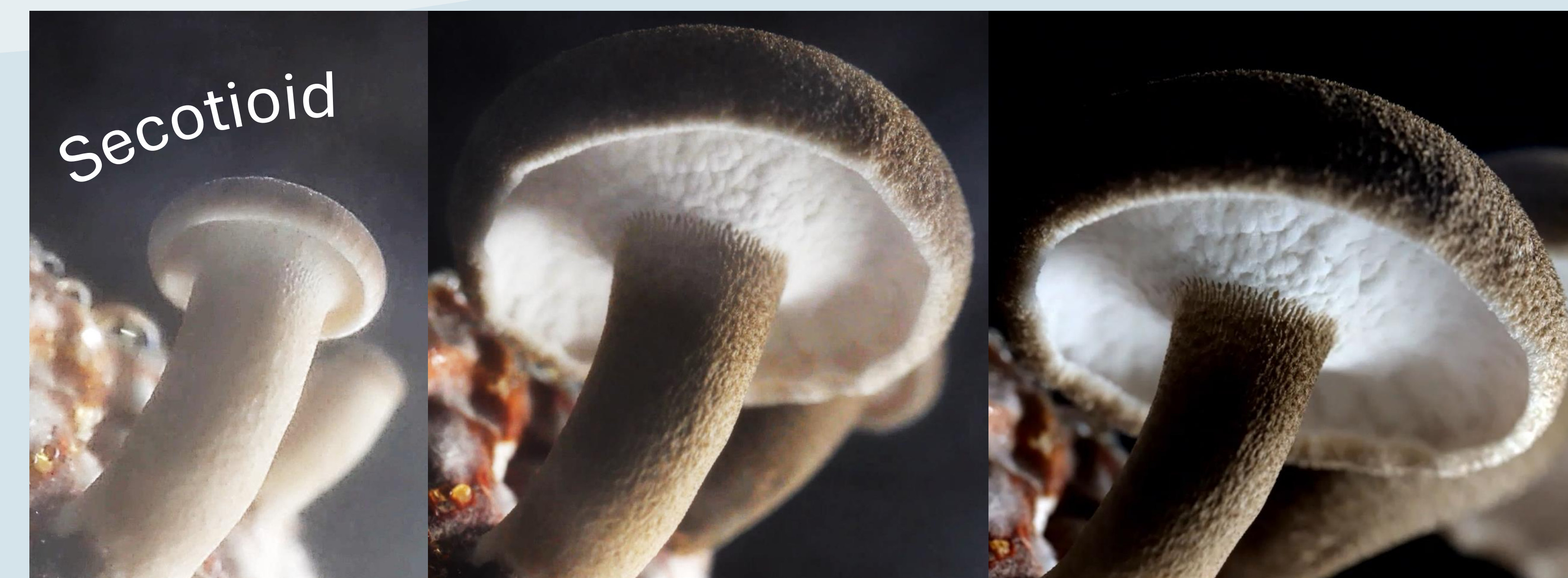
- ☉ Growth primarily through expansion
- ☉ Consistent with typical Agaricales model
- ☉ Inconsistent with typical Polyporales development
- ☉ Agaricales-like growth likely convergent



START

24 h

48 h



References

- Grand et al. Relationships within *Lentinus* subg. *Lentinus* (Polyporales, Agaricomycetes), with emphasis on sects. *Lentinus* and *Tigrini*. *Mycol Progress* 10, 399–413 (2011).
- Wu et al. Genomics and development of *Lentinus tigrinus*: A white-rot wood-decaying mushroom with dimorphic fruiting bodies. *Genome Biology and Evolution* 10, 3250–3261 (2018).
- Nordberg, H. et al. The genome portal of the Department of Energy Joint Genome Institute: 2014 updates. *Nucleic Acids Research* 42, D26–31 (2014).

Acknowledgements

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Questions

- What is the element controlling agaricoid/secotioid dimorphism?
 - ☉ Past work: region of 3 Mb, 4 scaffolds²
 - ☉ Likely a single locus in the genome²
- Are European and North American *L. tigrinus* specimens a single species?
 - ☉ Mating compatible but no overlap of morphological traits in the wild
 - ☉ Suggests lack of gene flow
- Is the secotioid form under selection?
 - ☉ Likely yes – preliminary MA population indicates heterozygote advantage