

# Emergence and diversification of a highly invasive tree pathogen lineage

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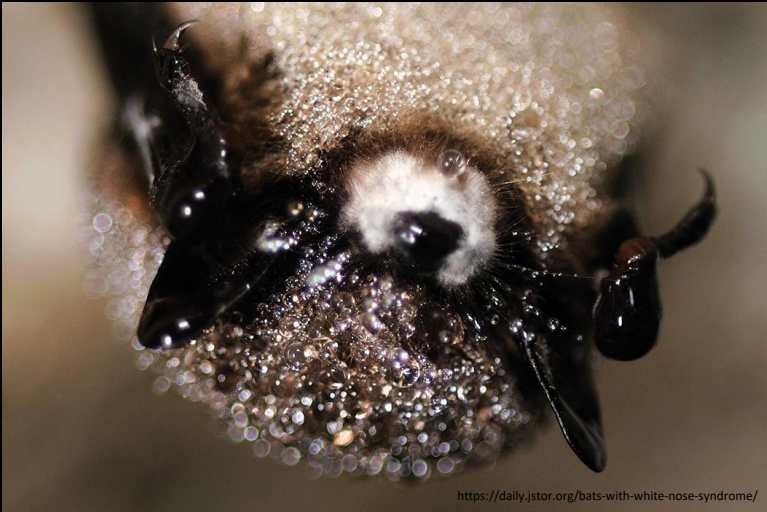
<sup>2</sup> Laboratory of Evolutionary Genetics, Institute of Biology, University of Neuchâtel (Switzerland)



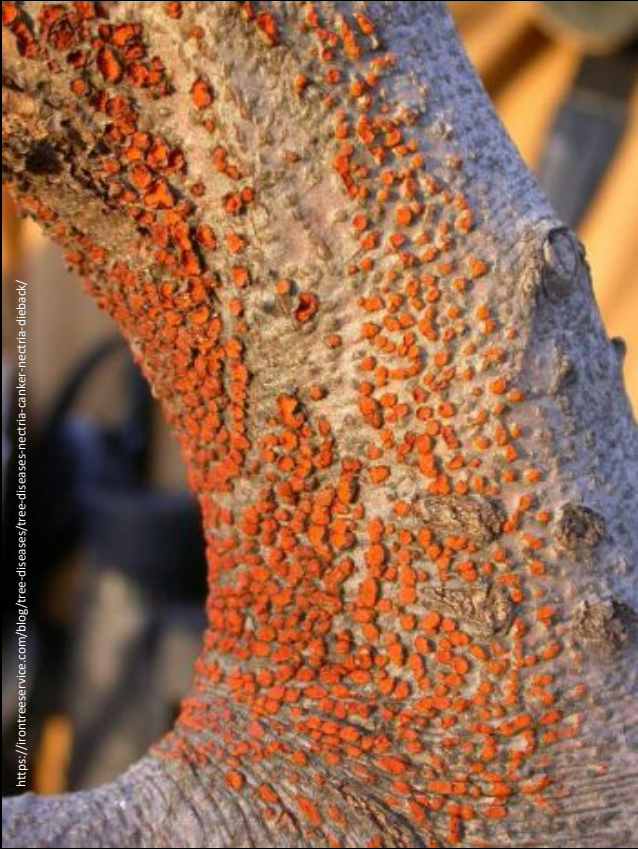
Nick Talbot, University of Exeter



<https://agrinfolbank.com.pk/>



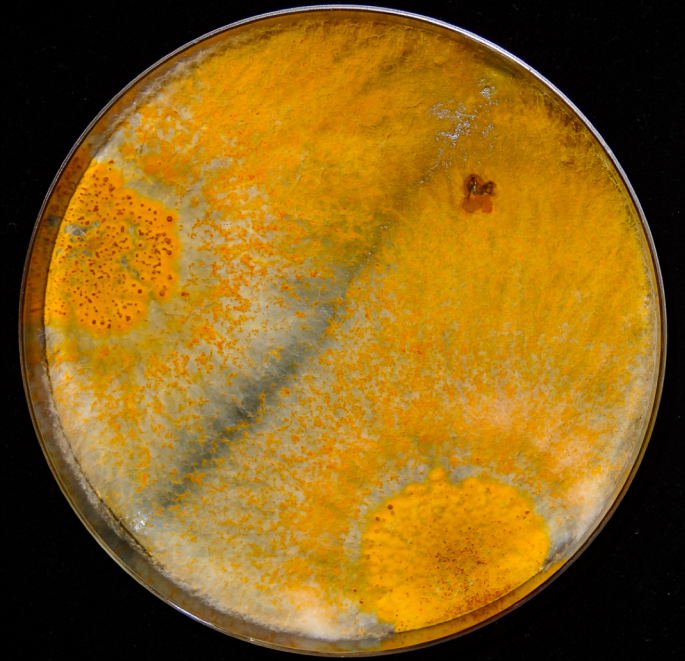
<https://daily.jstor.org/bats-with-white-nose-syndrome/>



<https://ronreservice.com/blog/tree-diseases-nectria-canker-nectria-dieback/>



# Chestnut blight – *Cryphonectria parasitica*



# Pathogen vs saprophyte



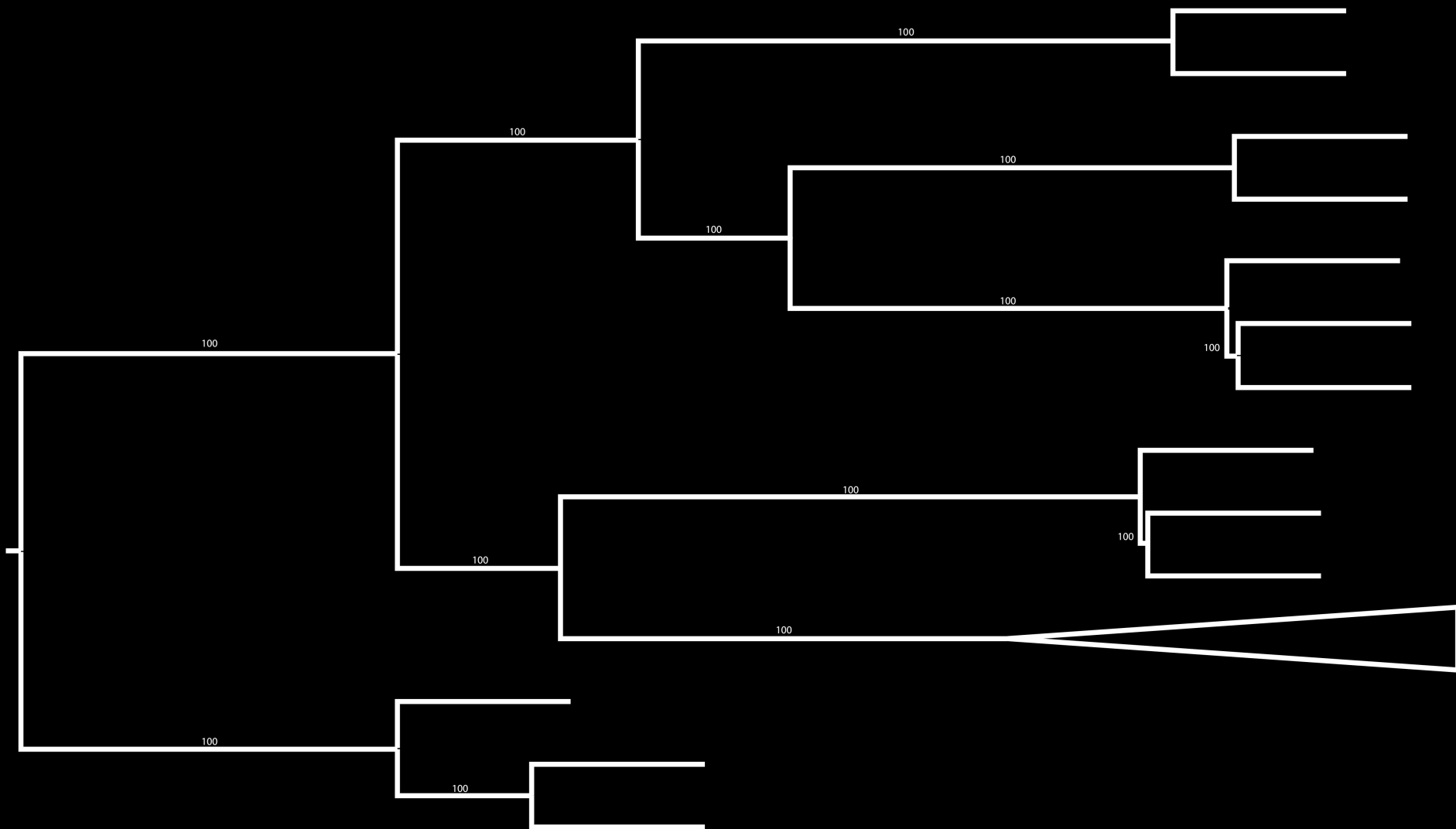
Pathogen vs saprophyte

Emergence & diversification of a  
highly invasive pathogen lineage

Pathogen vs saprophyte

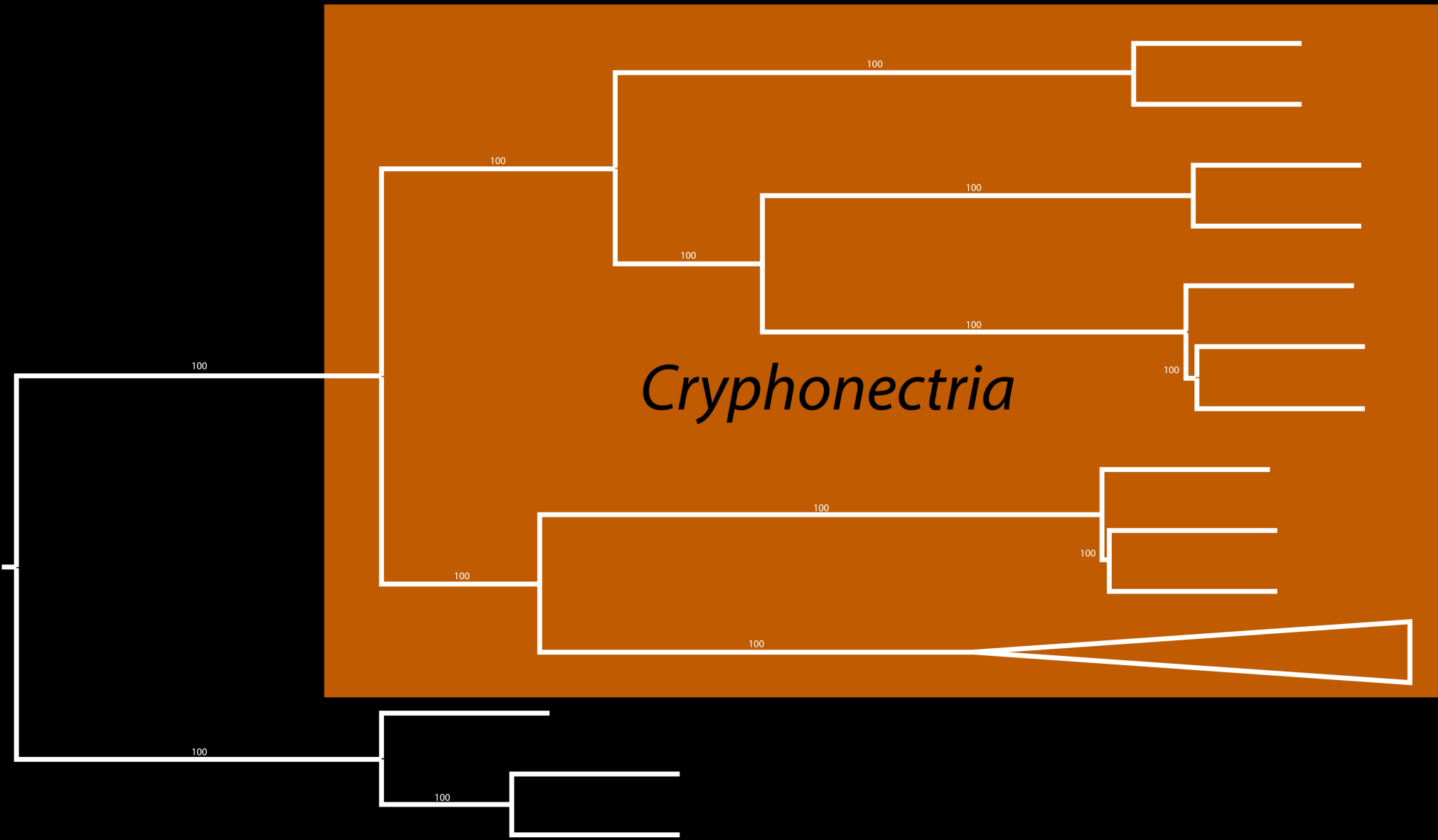
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Illumina WGS data (~300 isolates)

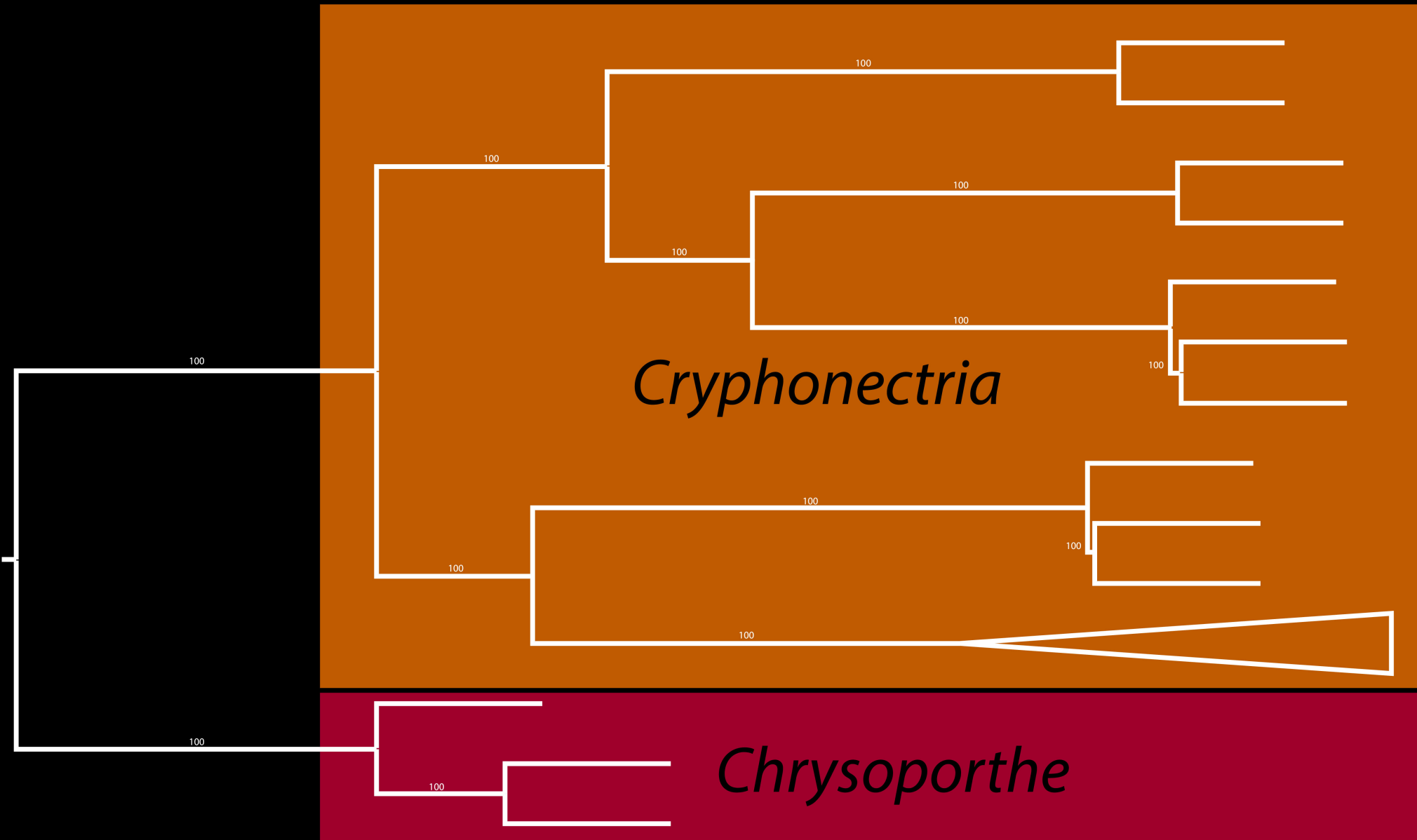


2.0

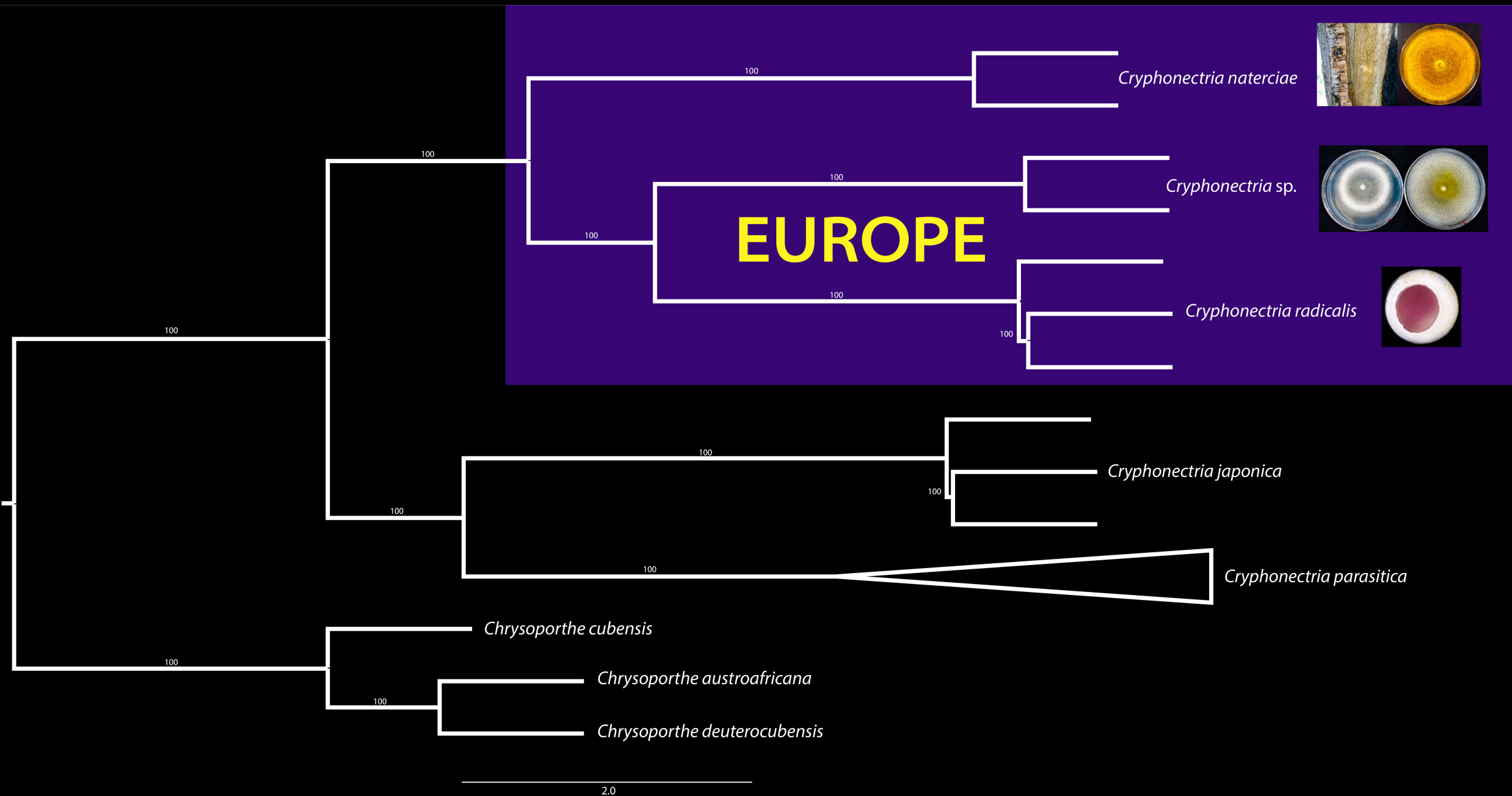




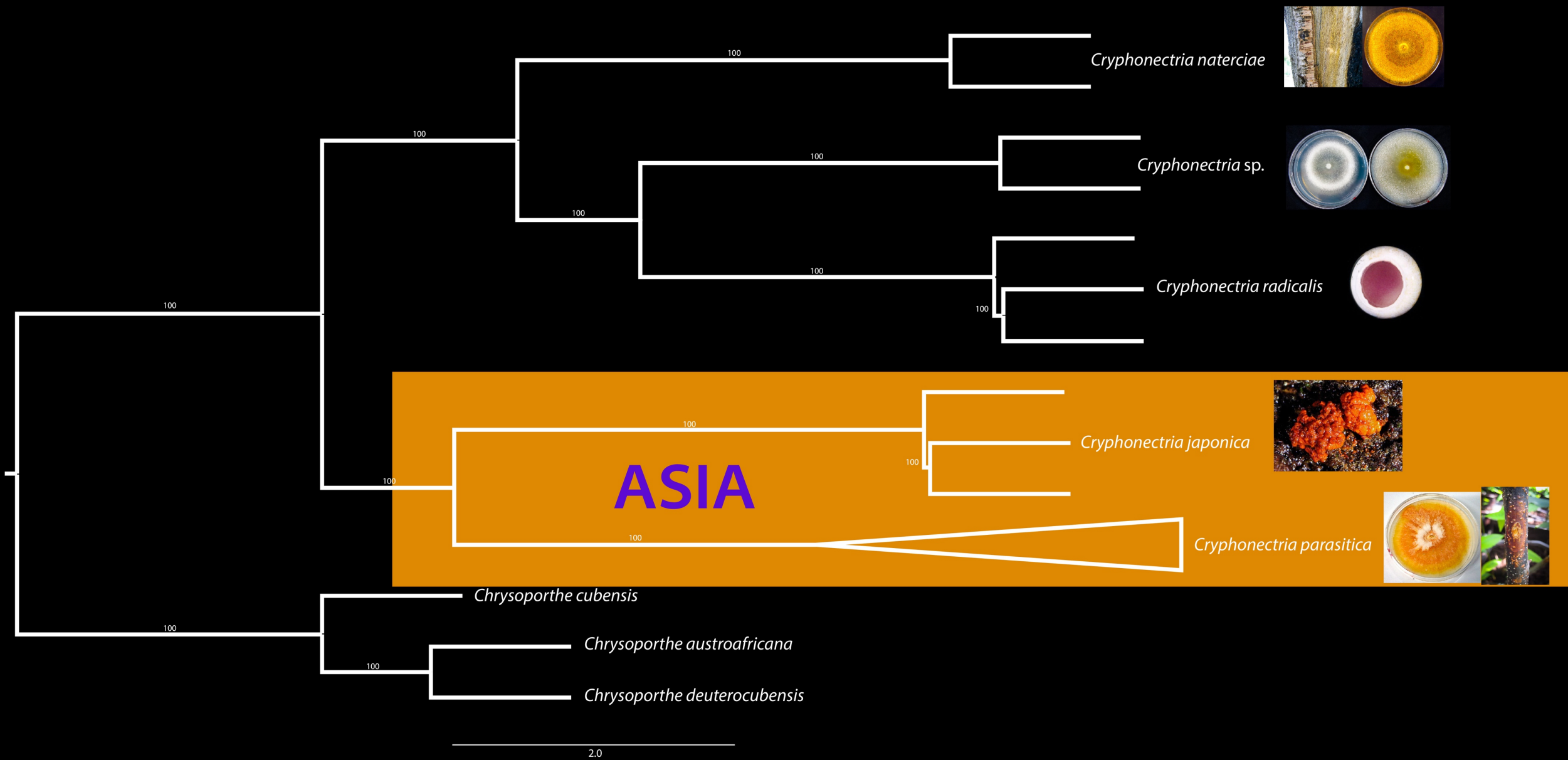
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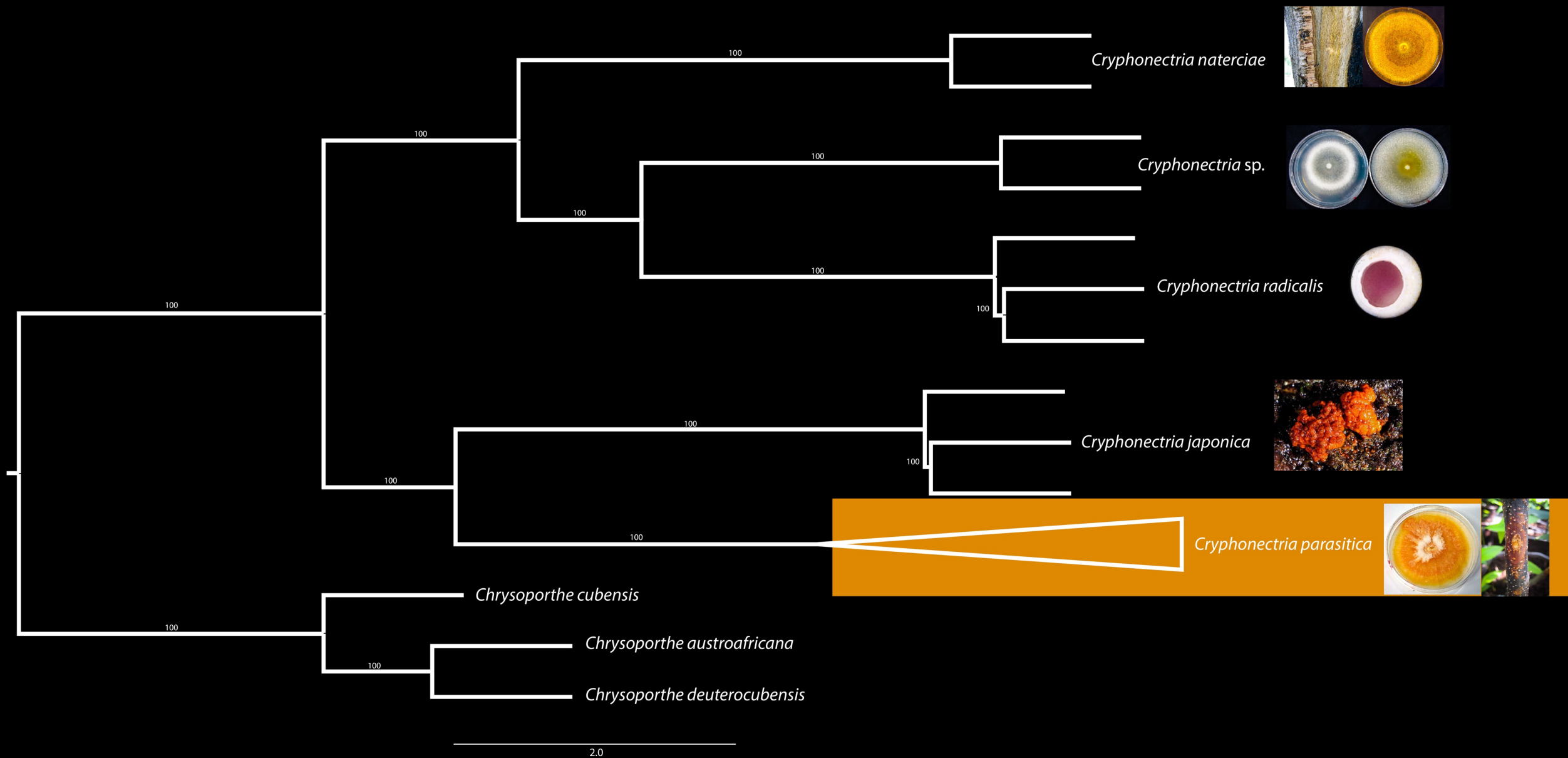


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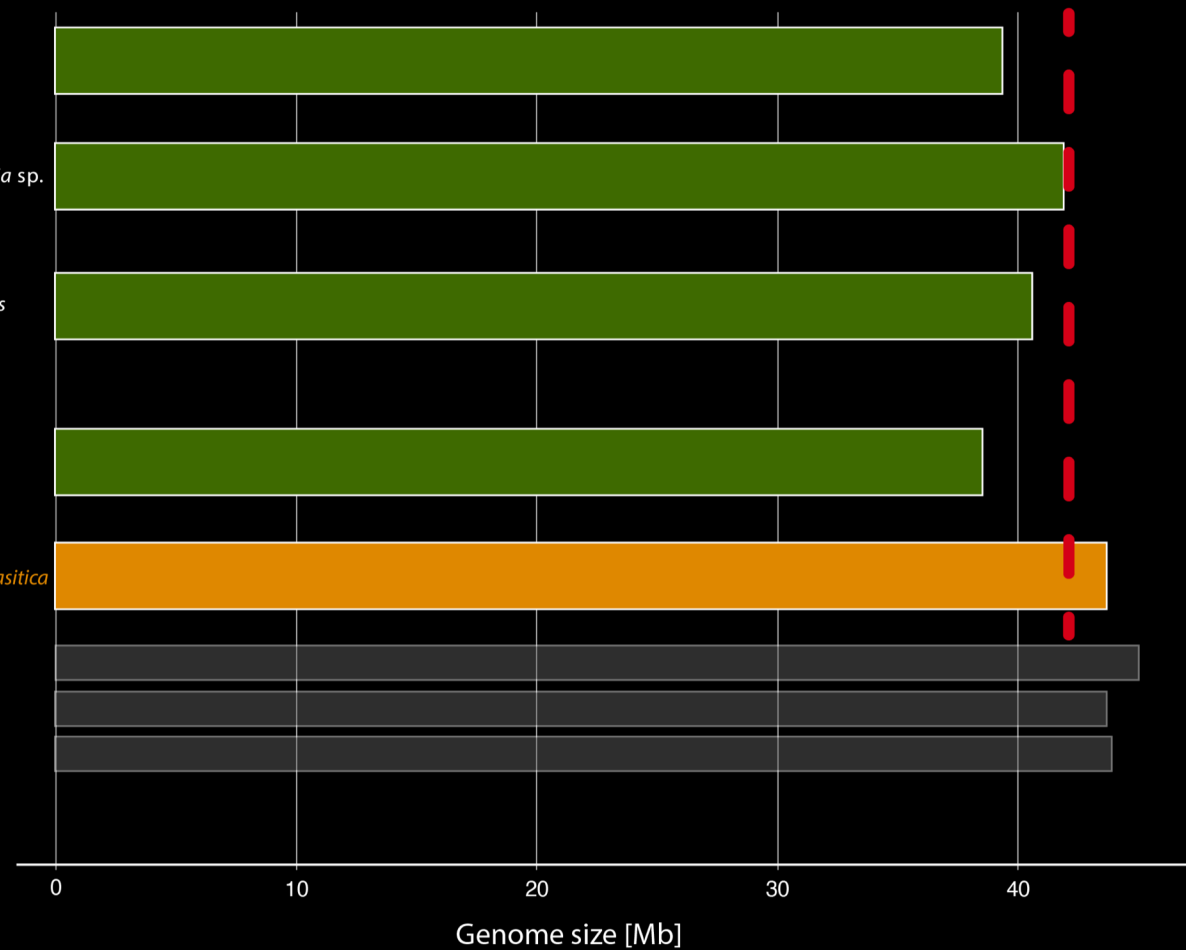
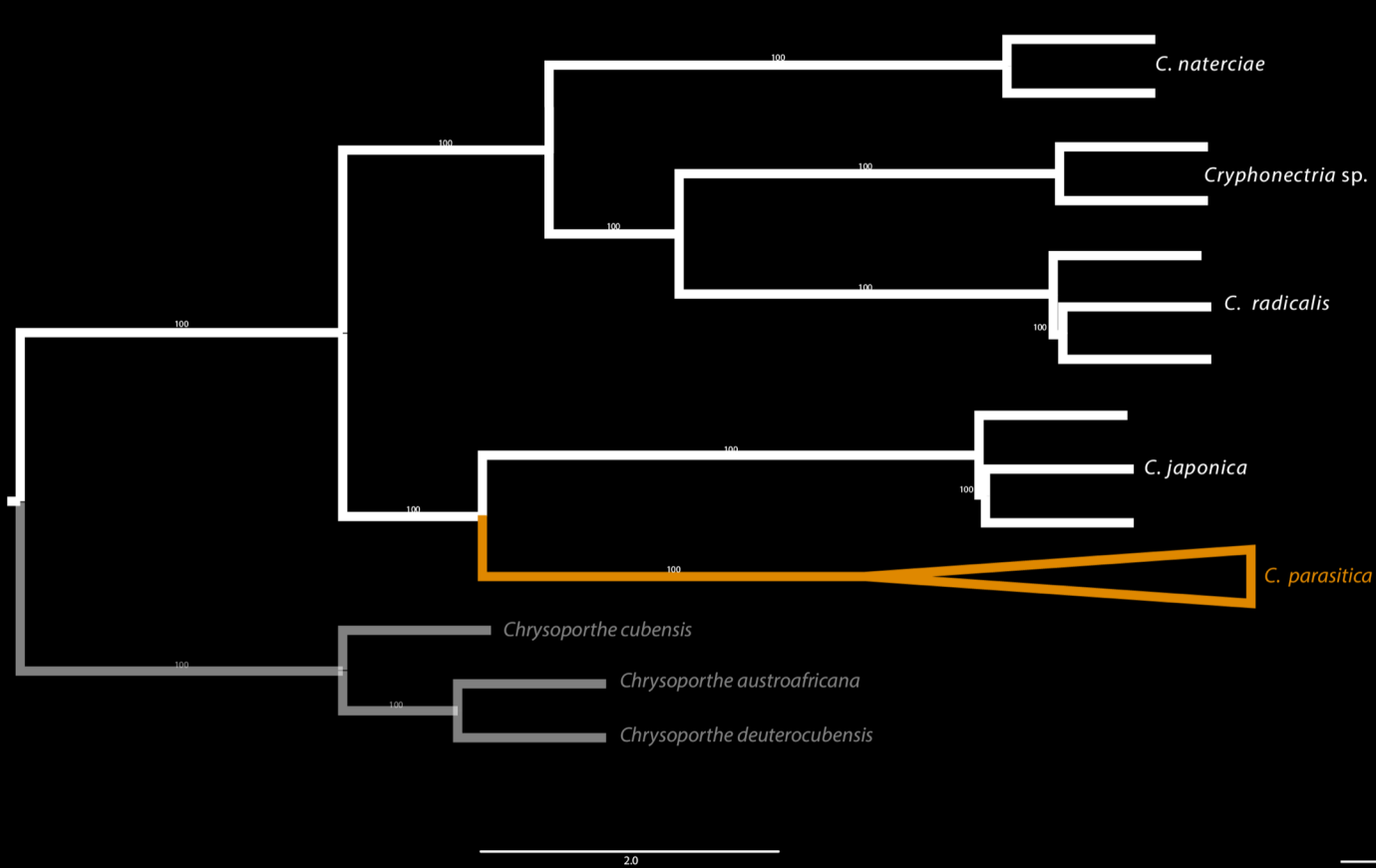






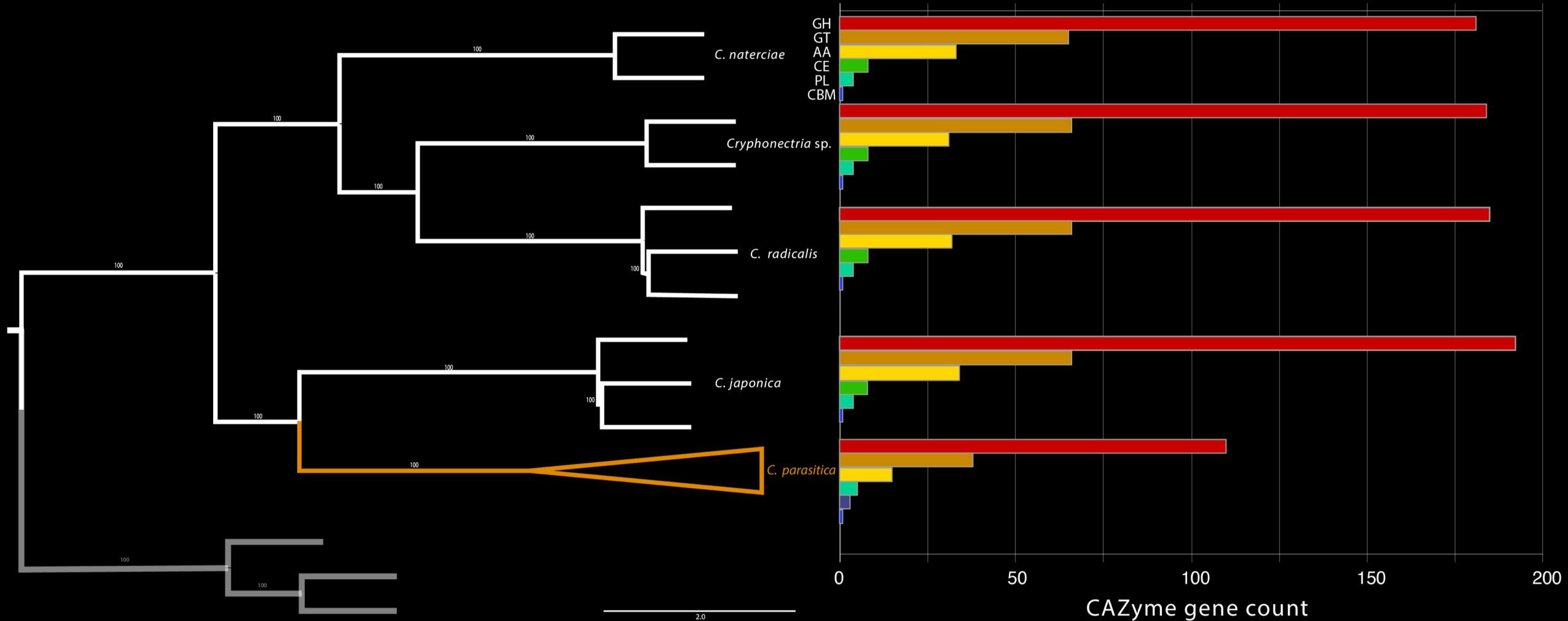


# Genome size

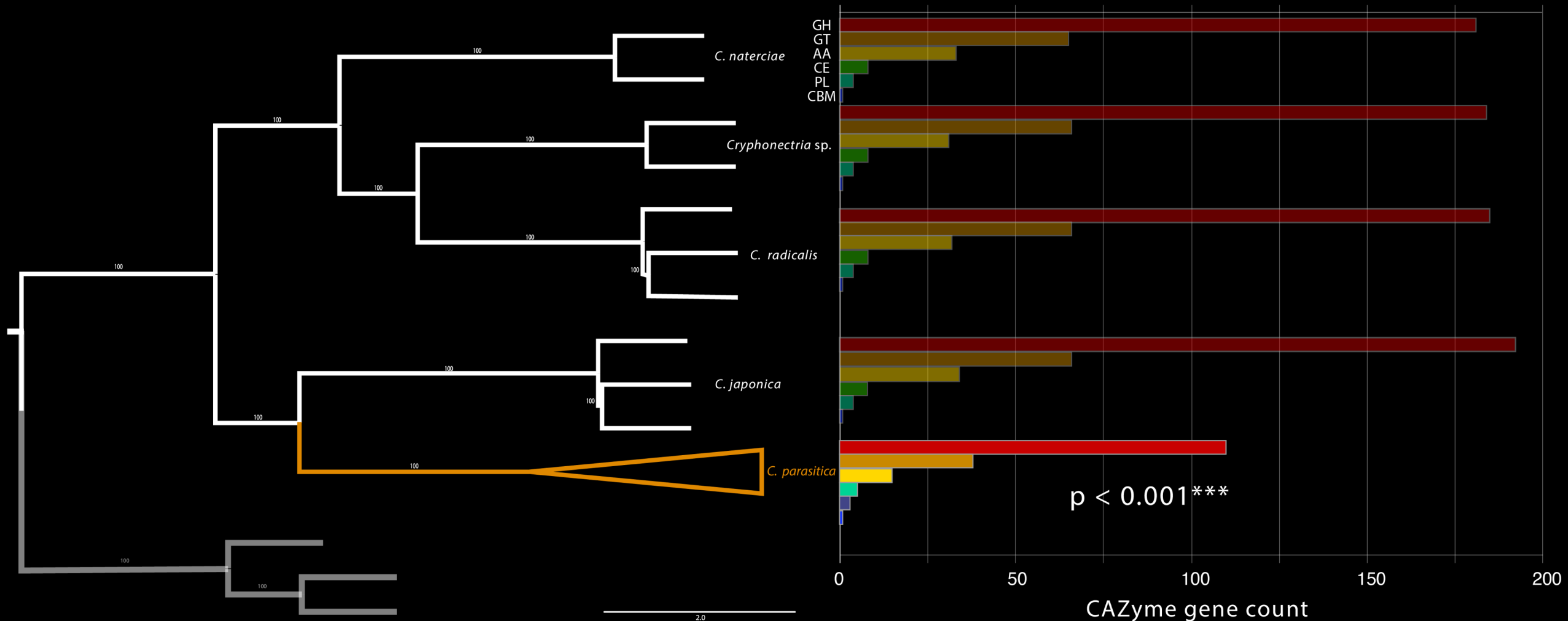




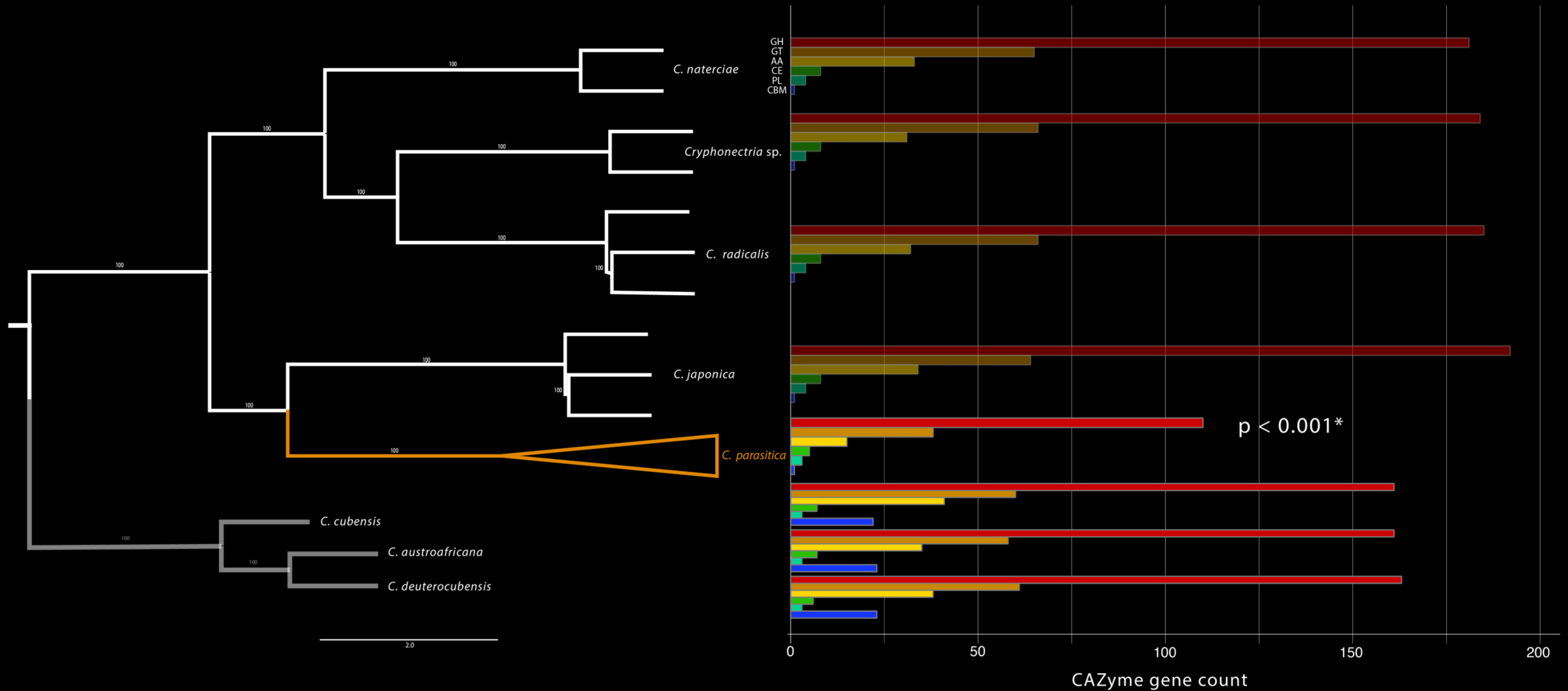
# Cell wall degrading enzymes



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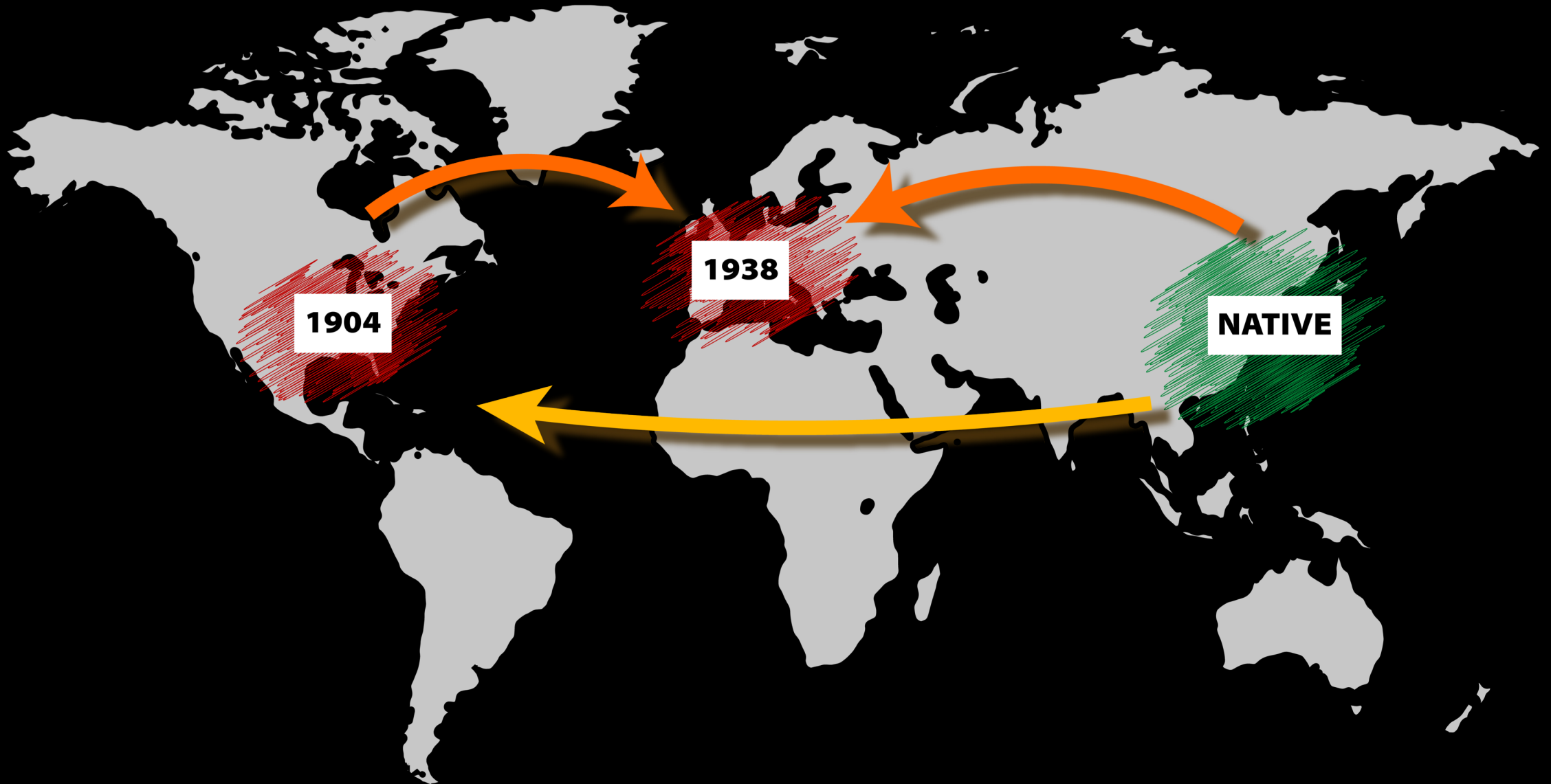


# Cell wall degrading enzymes



Emergence & diversification of a  
highly invasive lineage

# The chestnut blight invasion





# Low diversity in south-eastern Europe



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- vc type: EU-12

# Low diversity in south-eastern Europe



- vc type: EU-12
- Mating type: MAT-1

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- vc type: EU-12
- Mating type: MAT-1
- **S12 lineage = "clonal"**

# What we don't know about S12...



Origin?



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**Origin?**

**Diversity in S12?**

# What we don't know about S12...

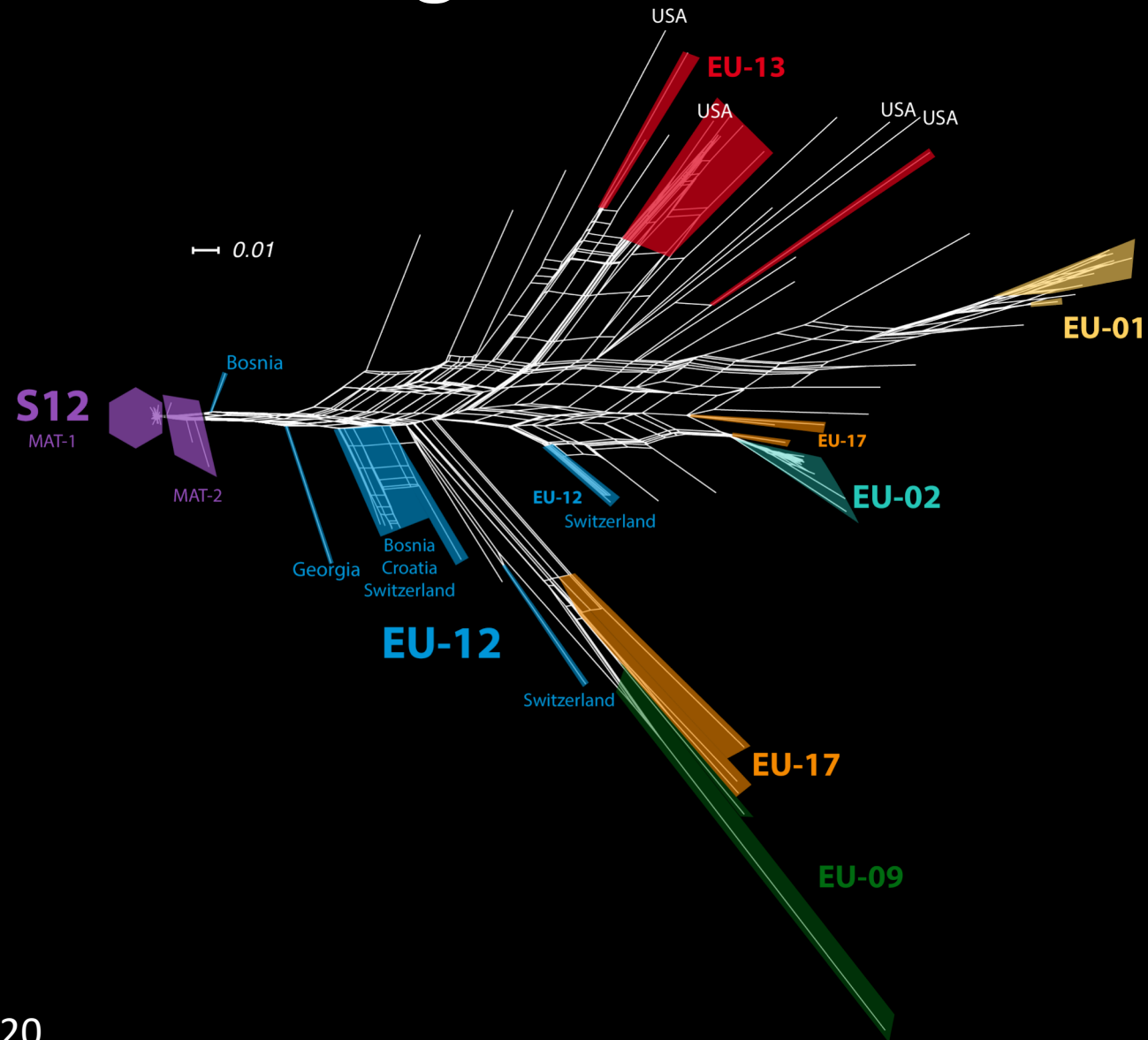


**Origin?**

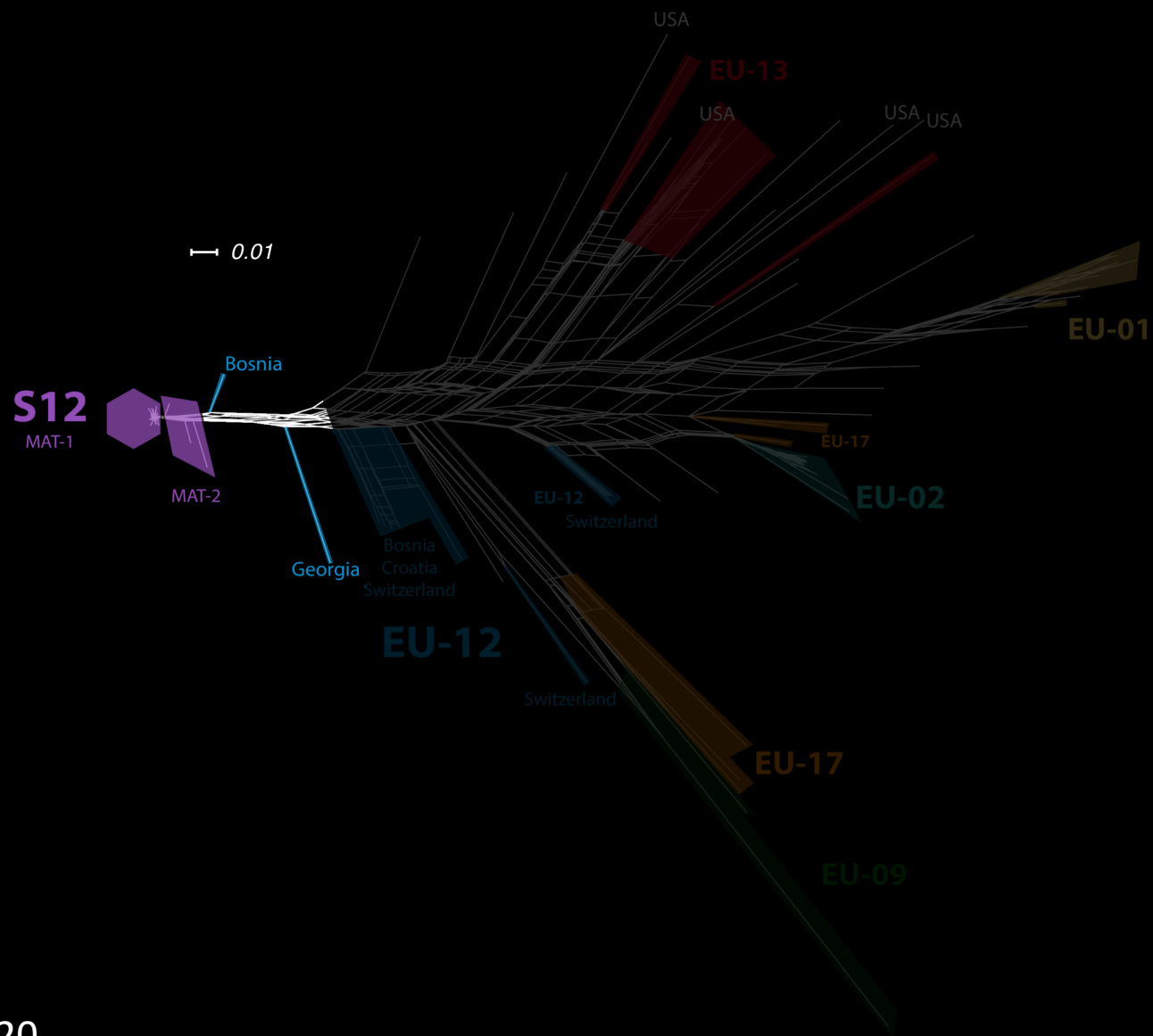
**Diversity in S12?**

**Evidence for adaptive evolution?**

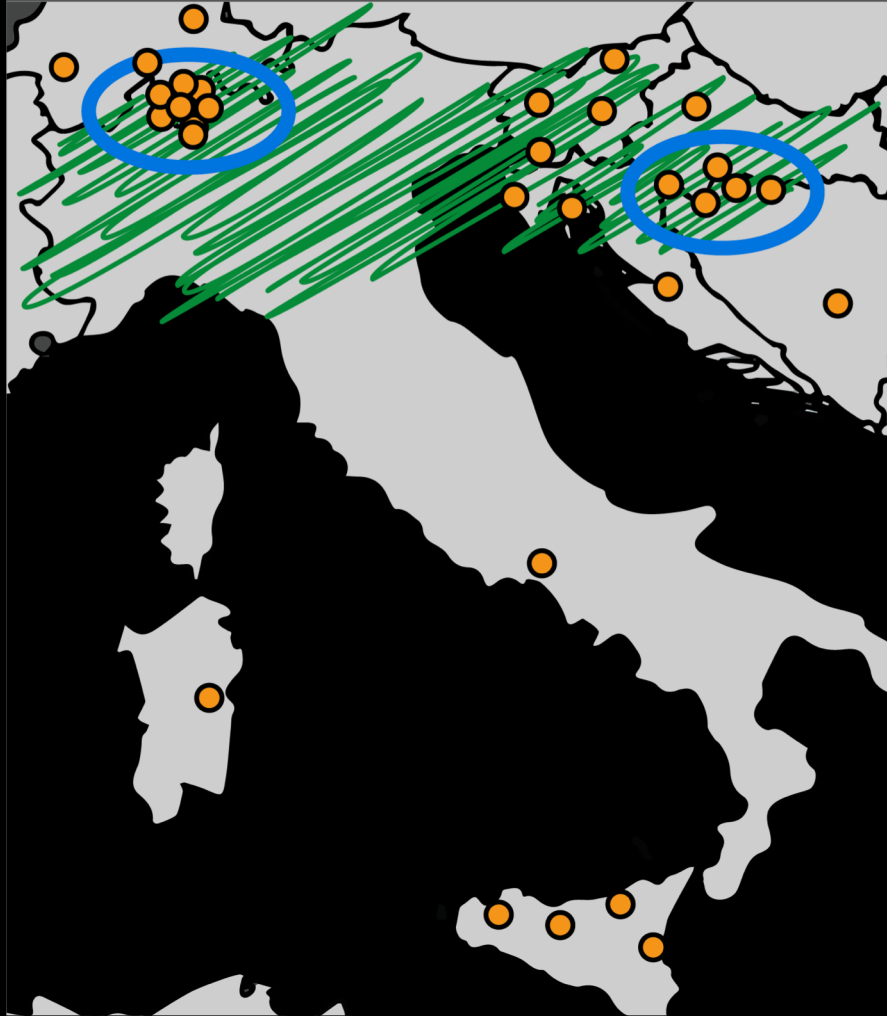
# Origin of S12



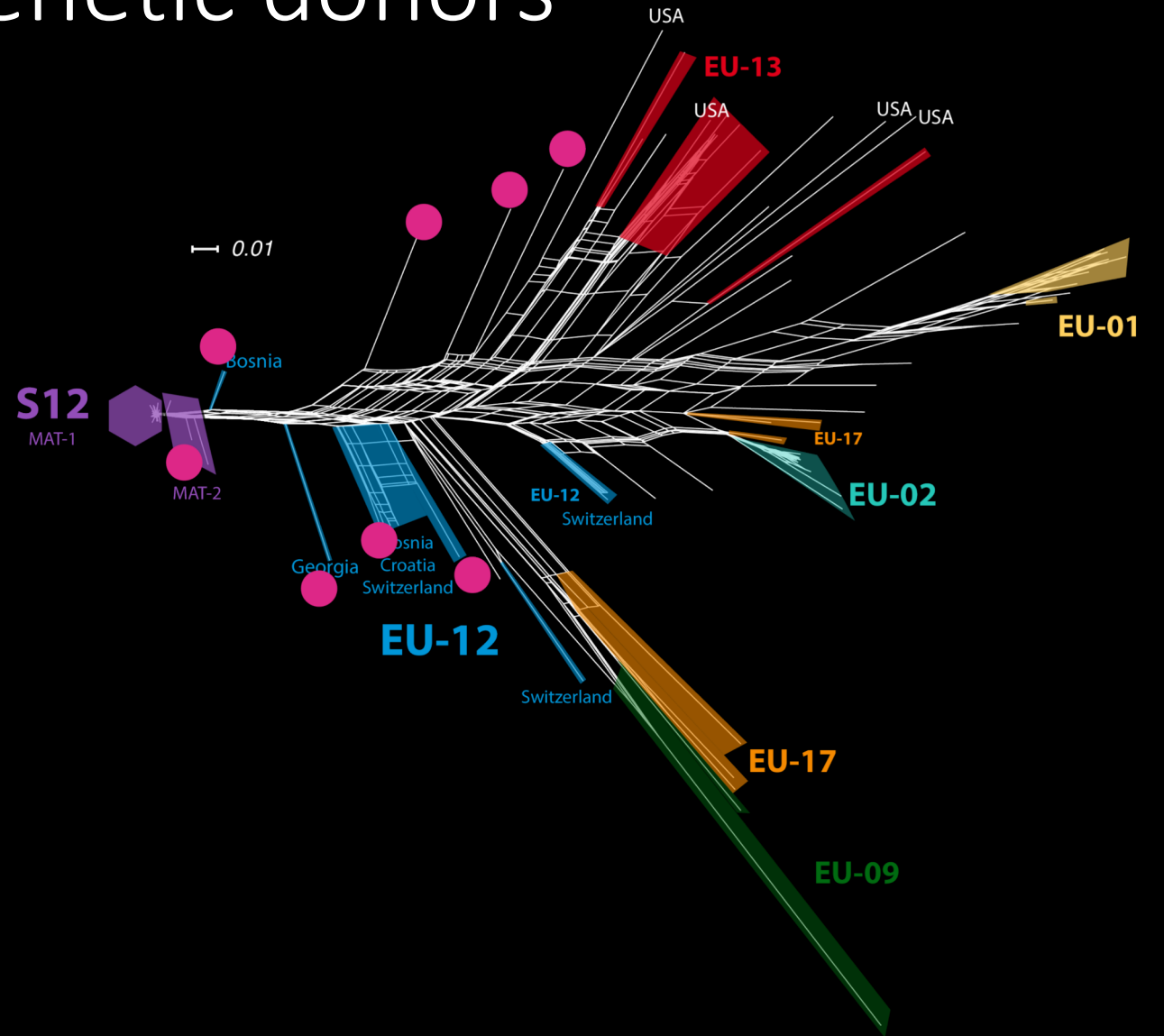
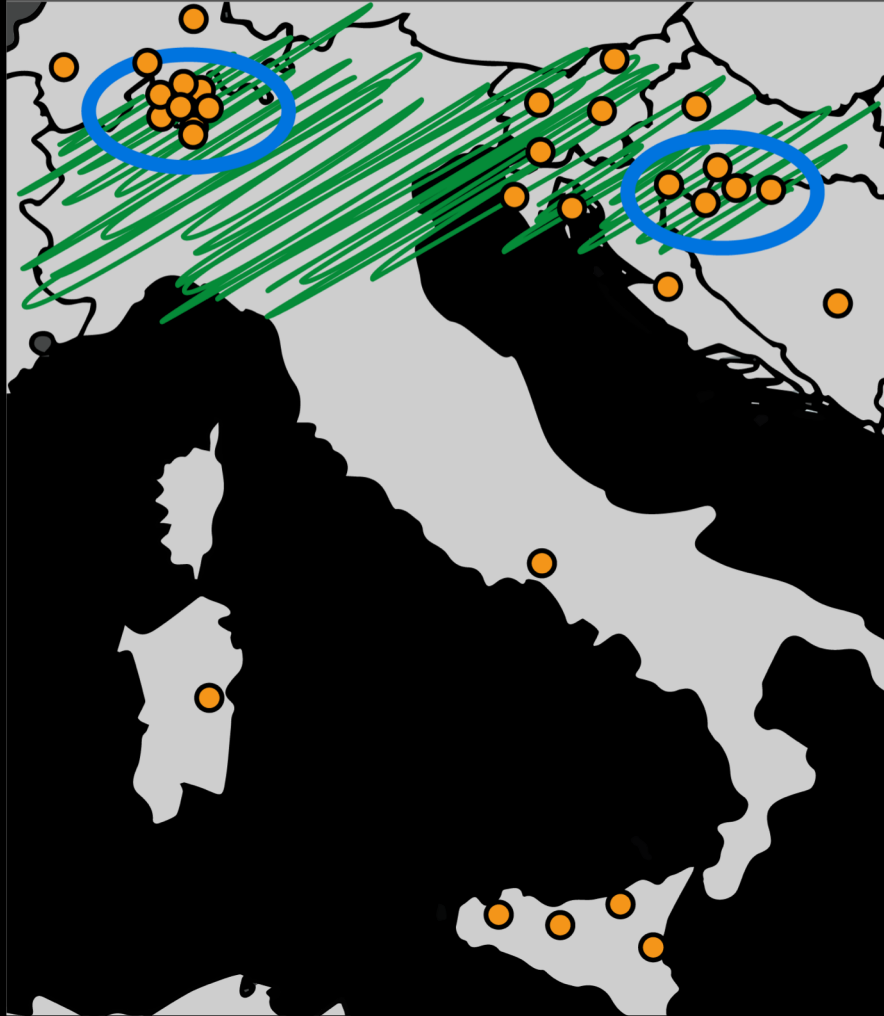
# Origin of S12



# S12 - genetic donors

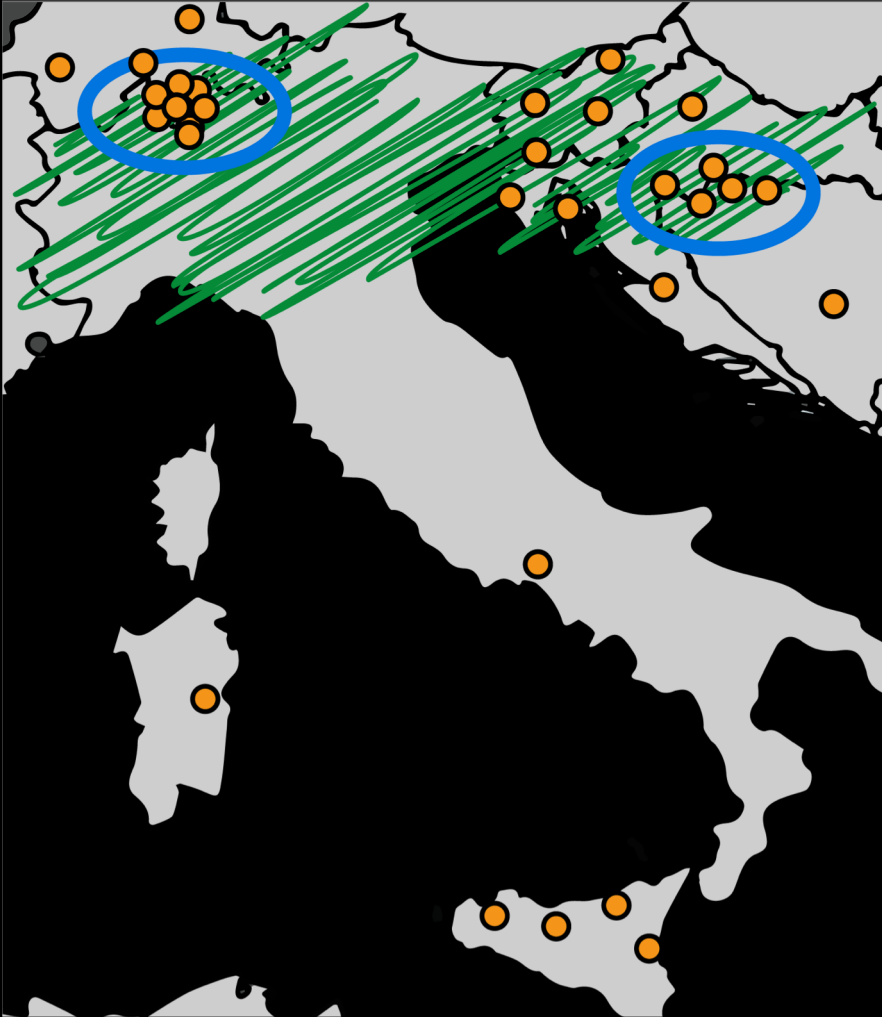


# S12 - genetic donors



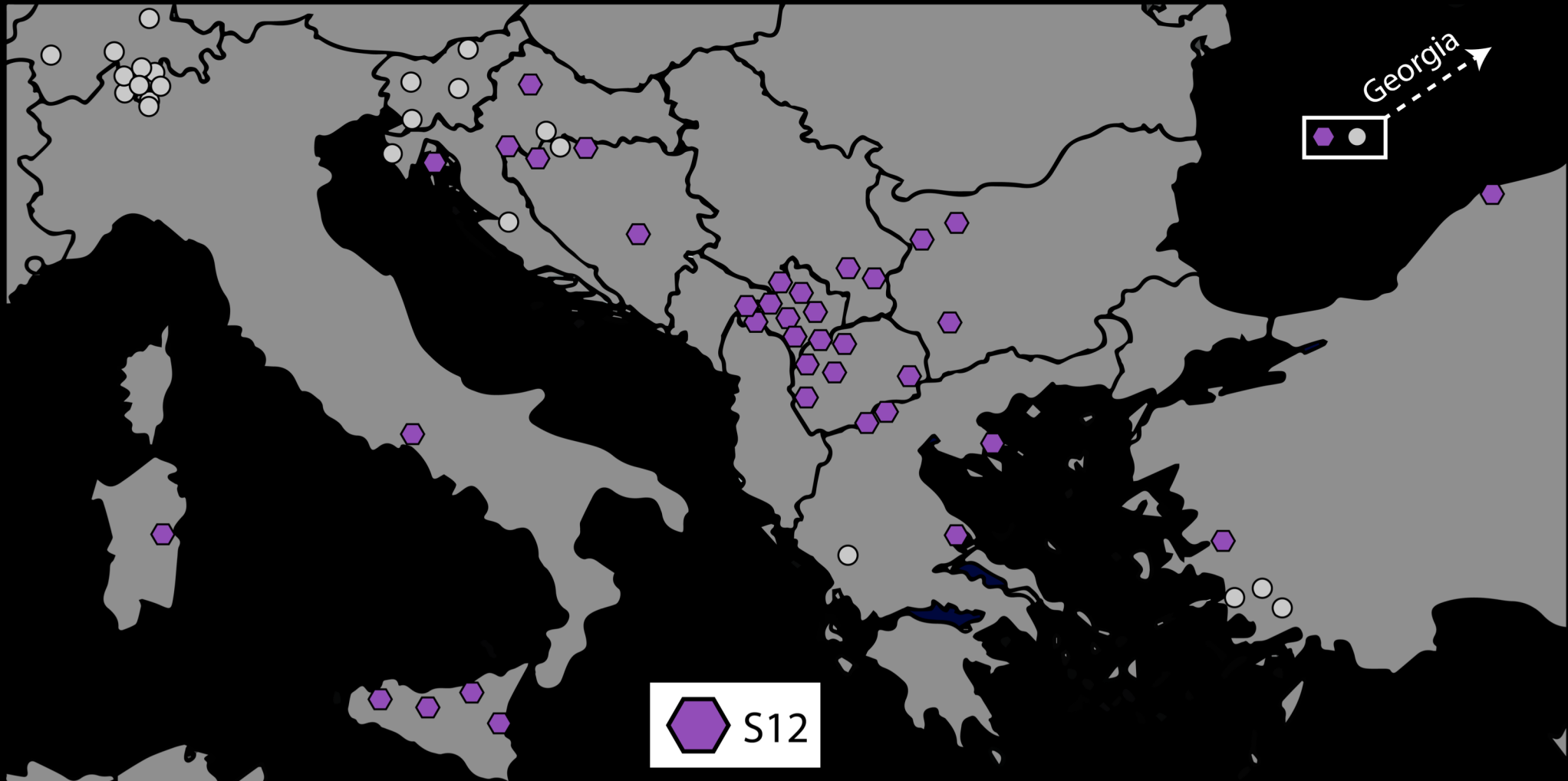


# Origin of S12

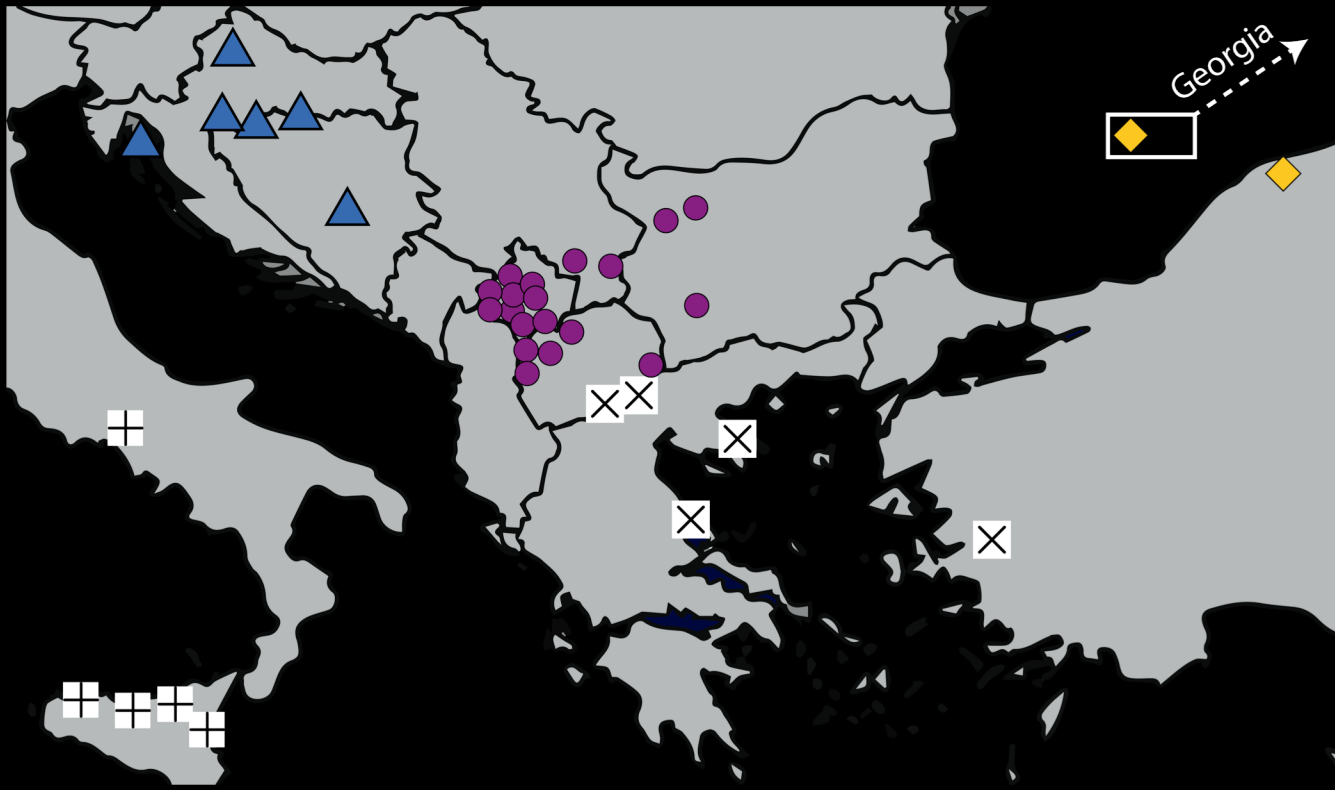


**Likely origin of S12 in northern Italy**

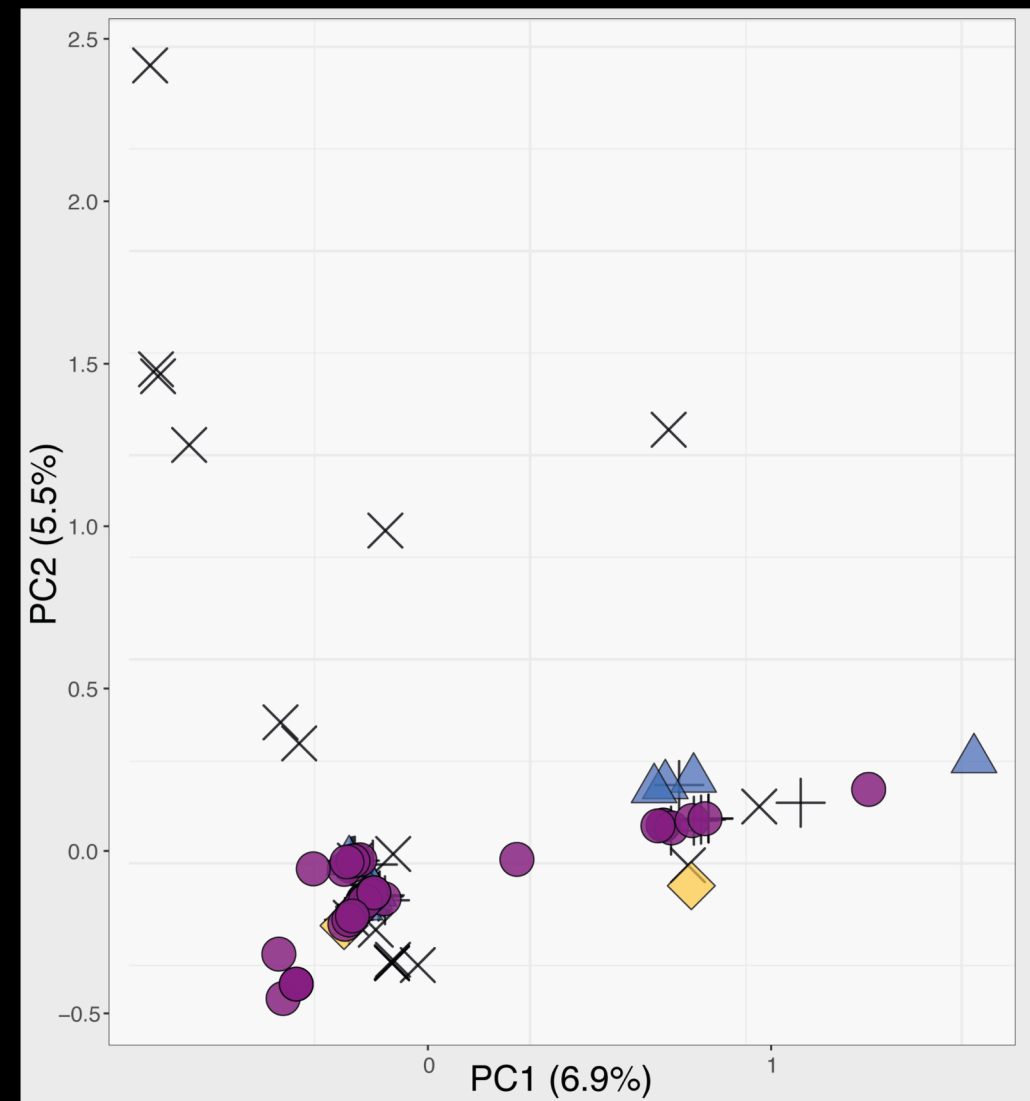
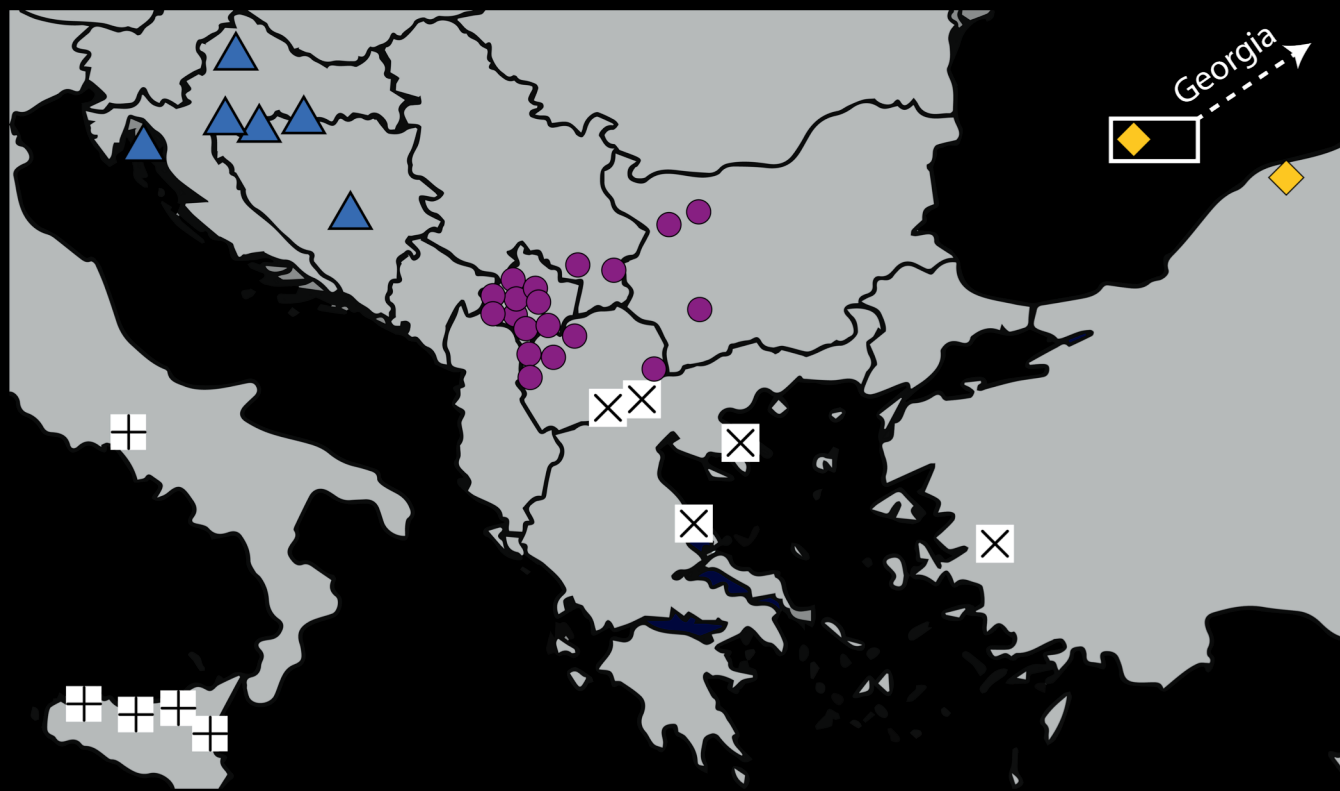
# Diversity in a “clonal lineage”



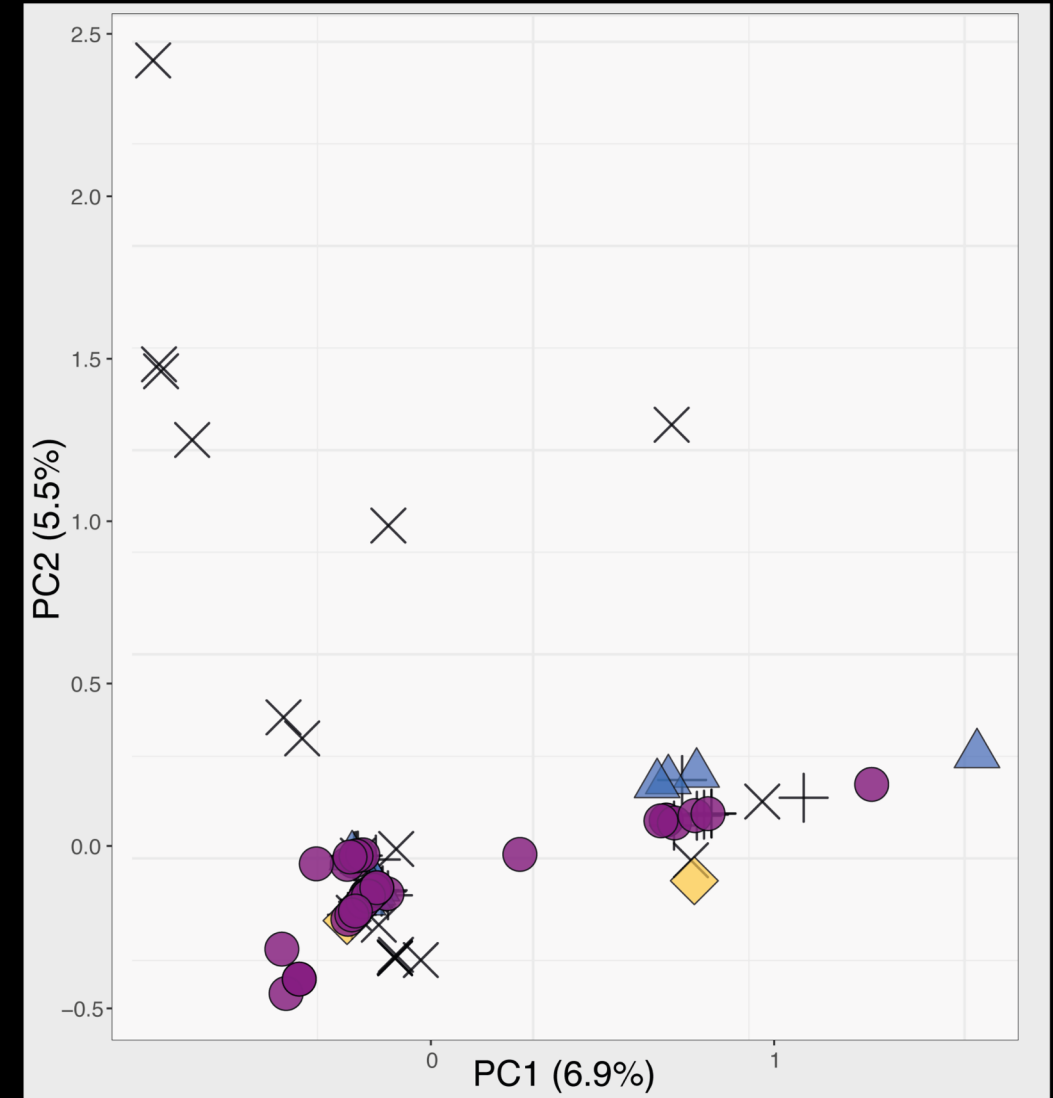
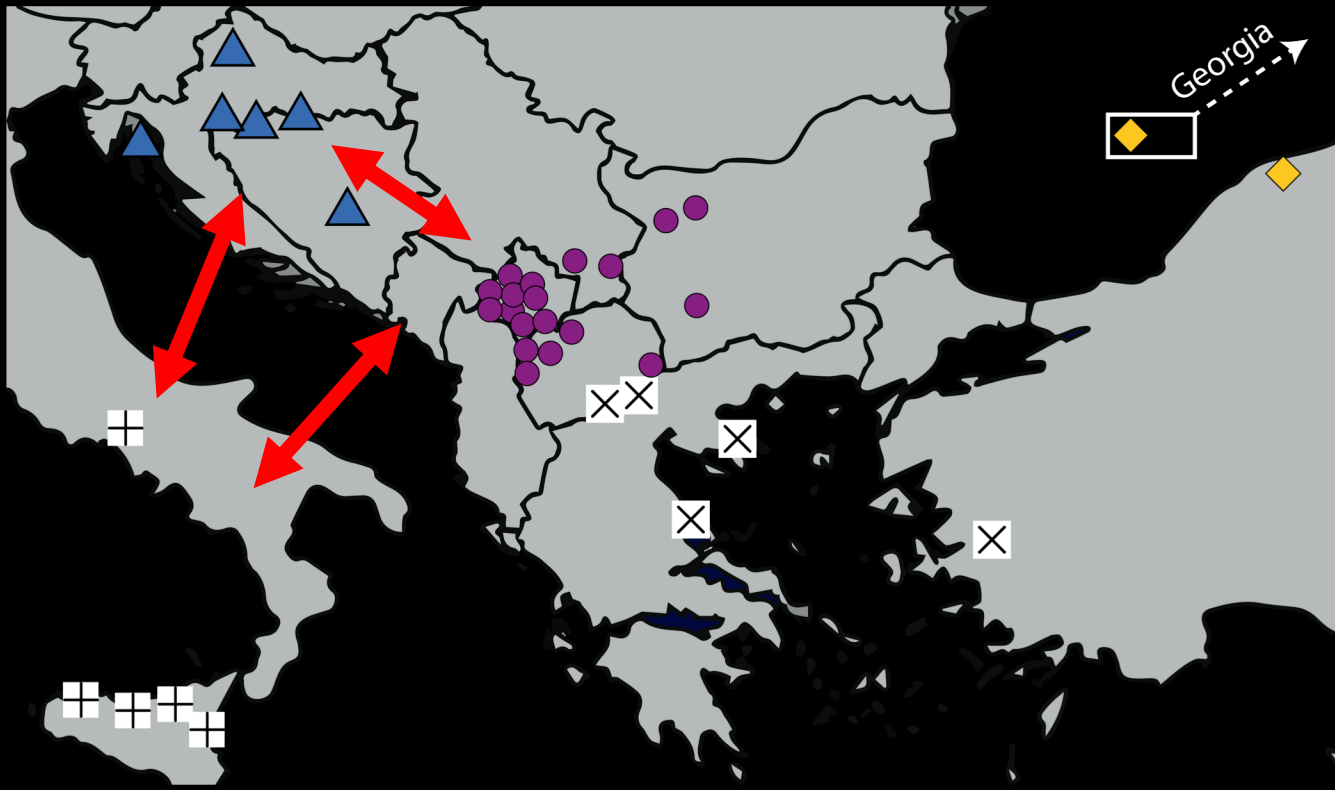
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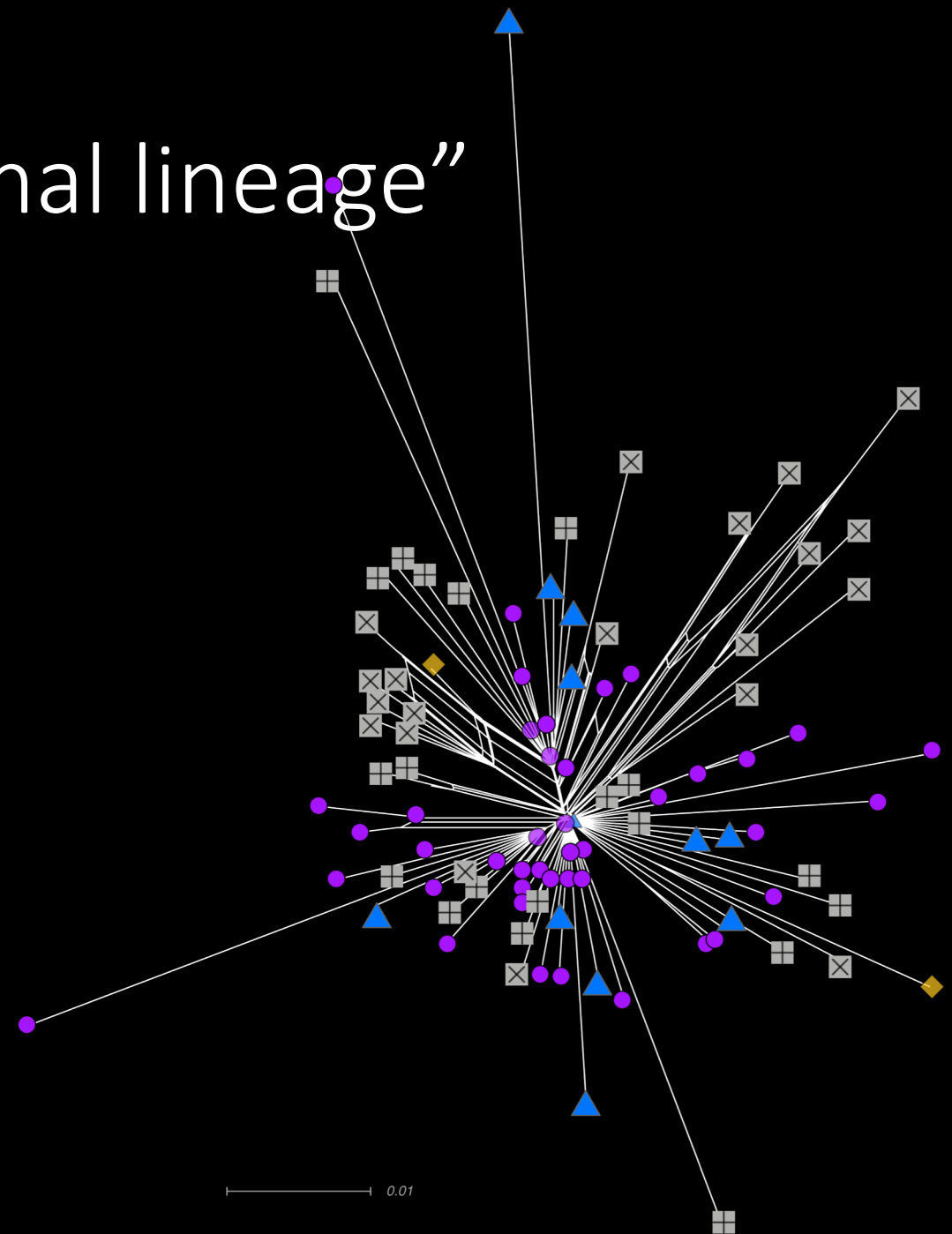


# Genetic drift & human trade



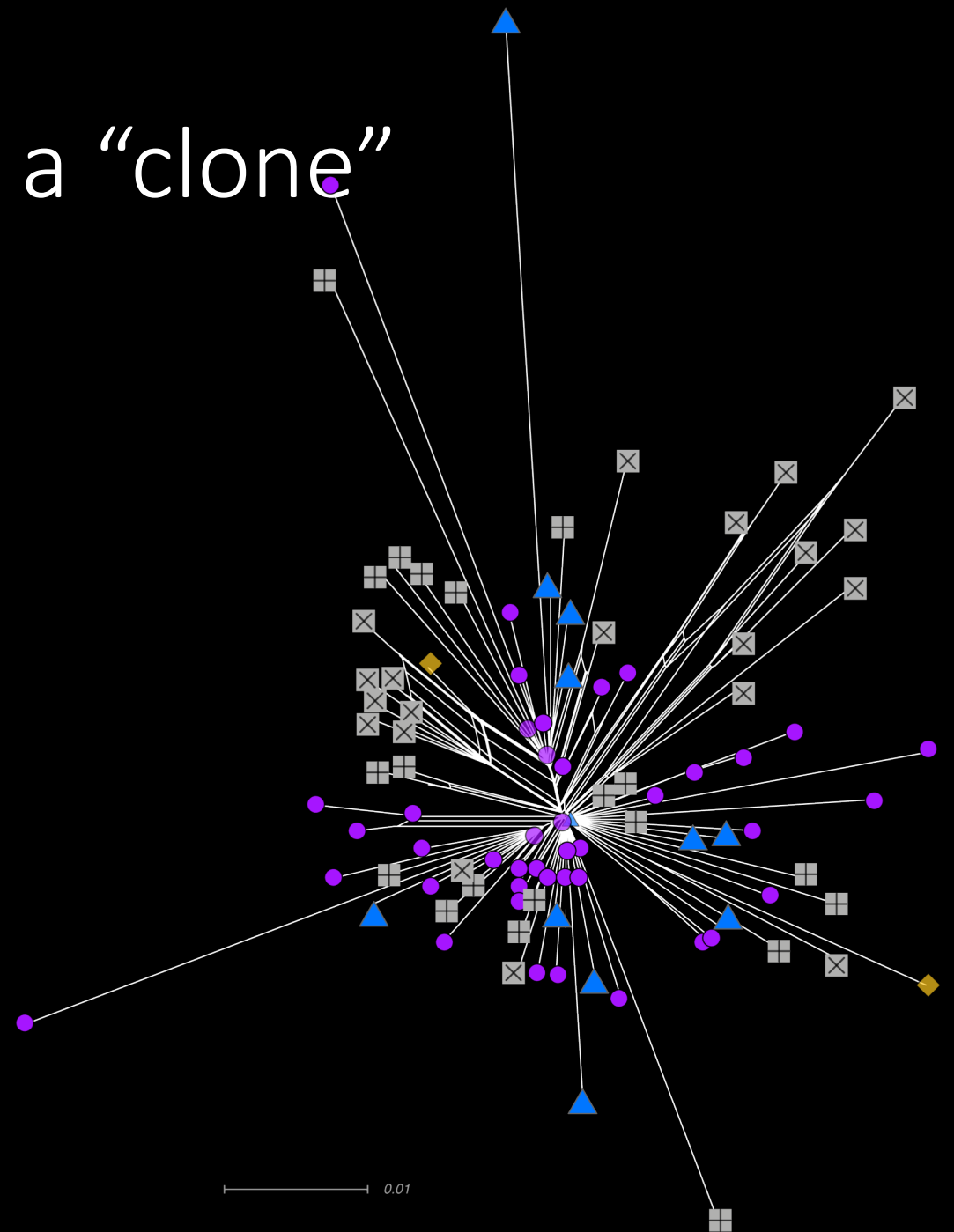
# Diversity in a “clonal lineage”

- Mutation accumulation (~85% singletons!)



# Recombination in a “clone”

- Mutation accumulation (~85% singletons!)
- Recombination (PHI:  $p=0.0035^{**}$ )



# Recombination in a “clone”

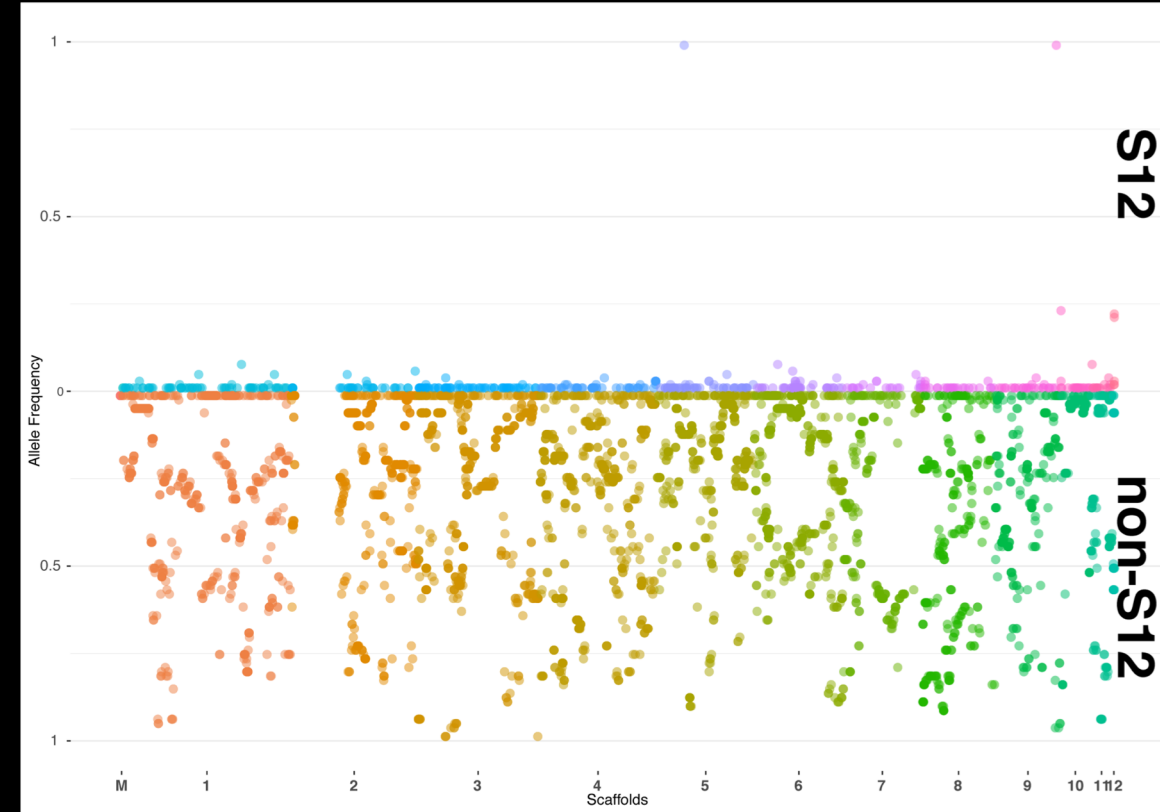
- Mutation accumulation (~85% singletons!)
- Recombination (PHI:  $p=0.0035^{**}$ )





# Purifying selection within lineage

- Mutation accumulation (~85% singletons!)
- Recombination (PHI:  $p=0.0035^{**}$ )
- Few deleterious mutations



**Origin**

European (Italian) bridgehead

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**Diversity**

no geography, mutation accumulation

## **Origin**

European (Italian) bridgehead

## **Diversity**

no geography, mutation accumulation

## **Evolution**

- Purifying selection
- clonality advantageous?
- Recombination possible!

# Thanks to...

Simone Prospero (Swiss Federal Research Institute WSL)

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