

Supplementary File

Supplementary Figure 1. Phylogeny of *Candida auris* benchmark dataset including cases from Clade I. The internal solid circle nodes indicate separations with a bootstrap value of at least 90%, and the number above the branch indicate branch length. Each leaf node represents a unique isolate; the shape of the node refers to the outbreak number code and the node color refers to the facility.

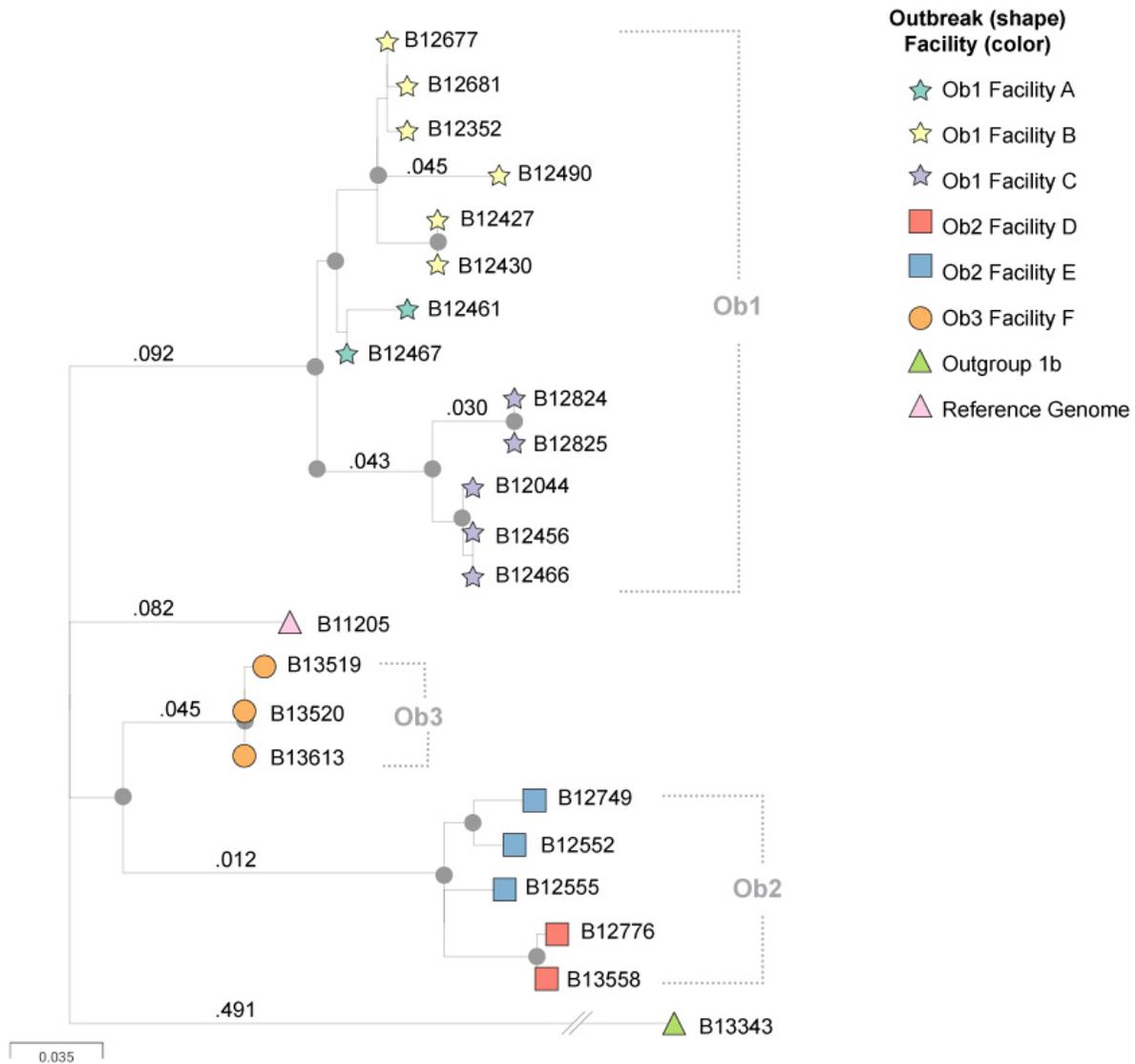
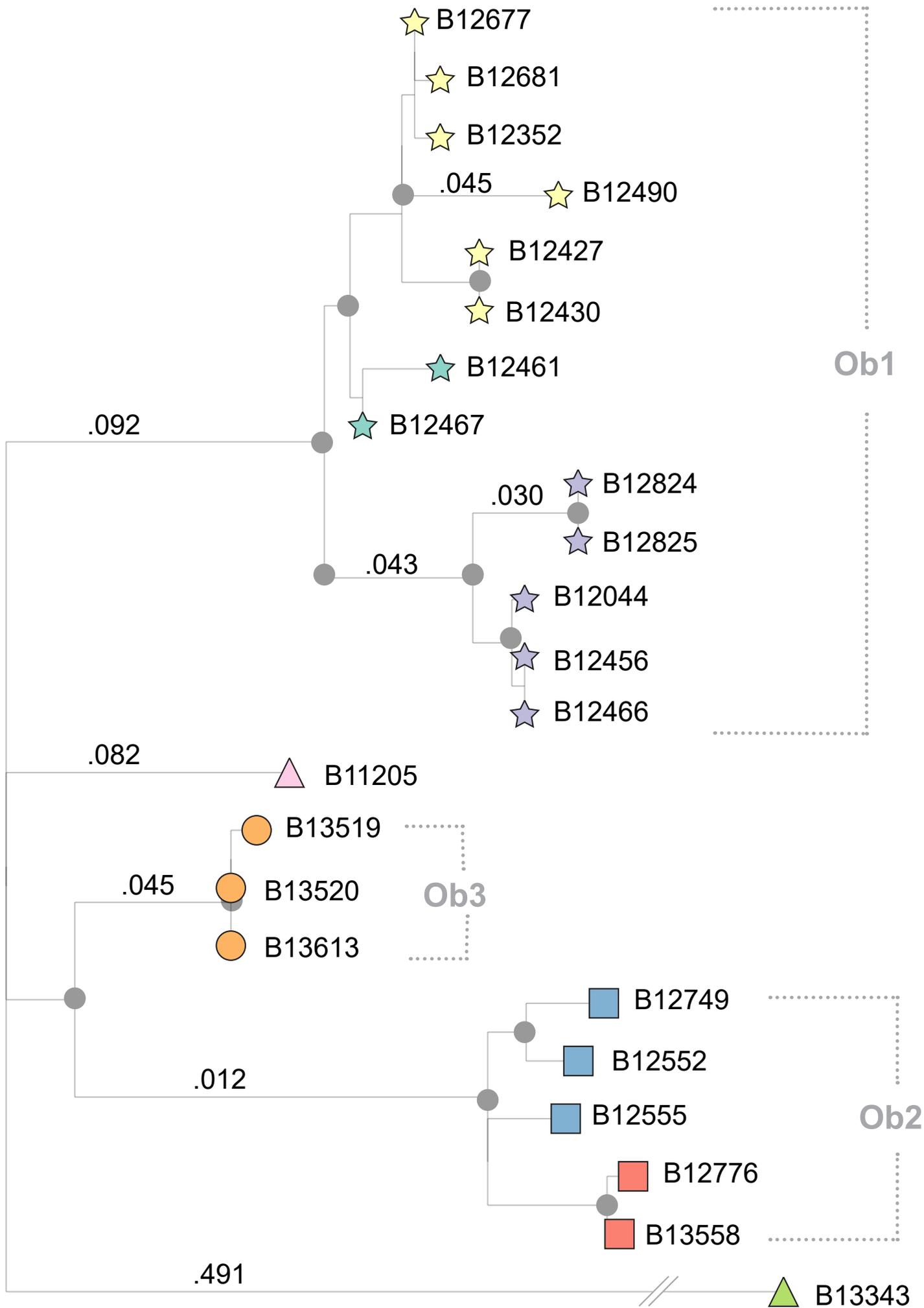


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Outbreak (shape)
Facility (color)

- Ob1 Facility A
- Ob1 Facility B
- Ob1 Facility C
- Ob2 Facility D
- Ob2 Facility E
- Ob3 Facility F
- Outgroup 1b
- Reference Genome



0.035