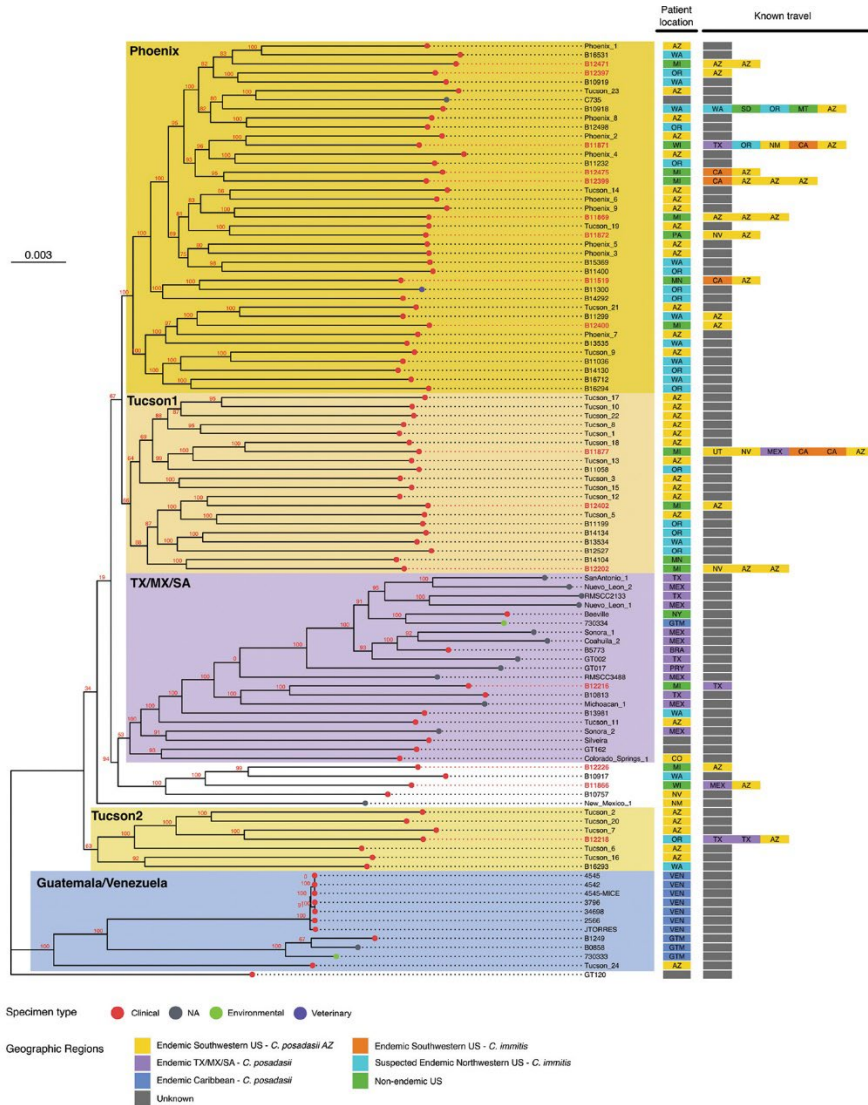
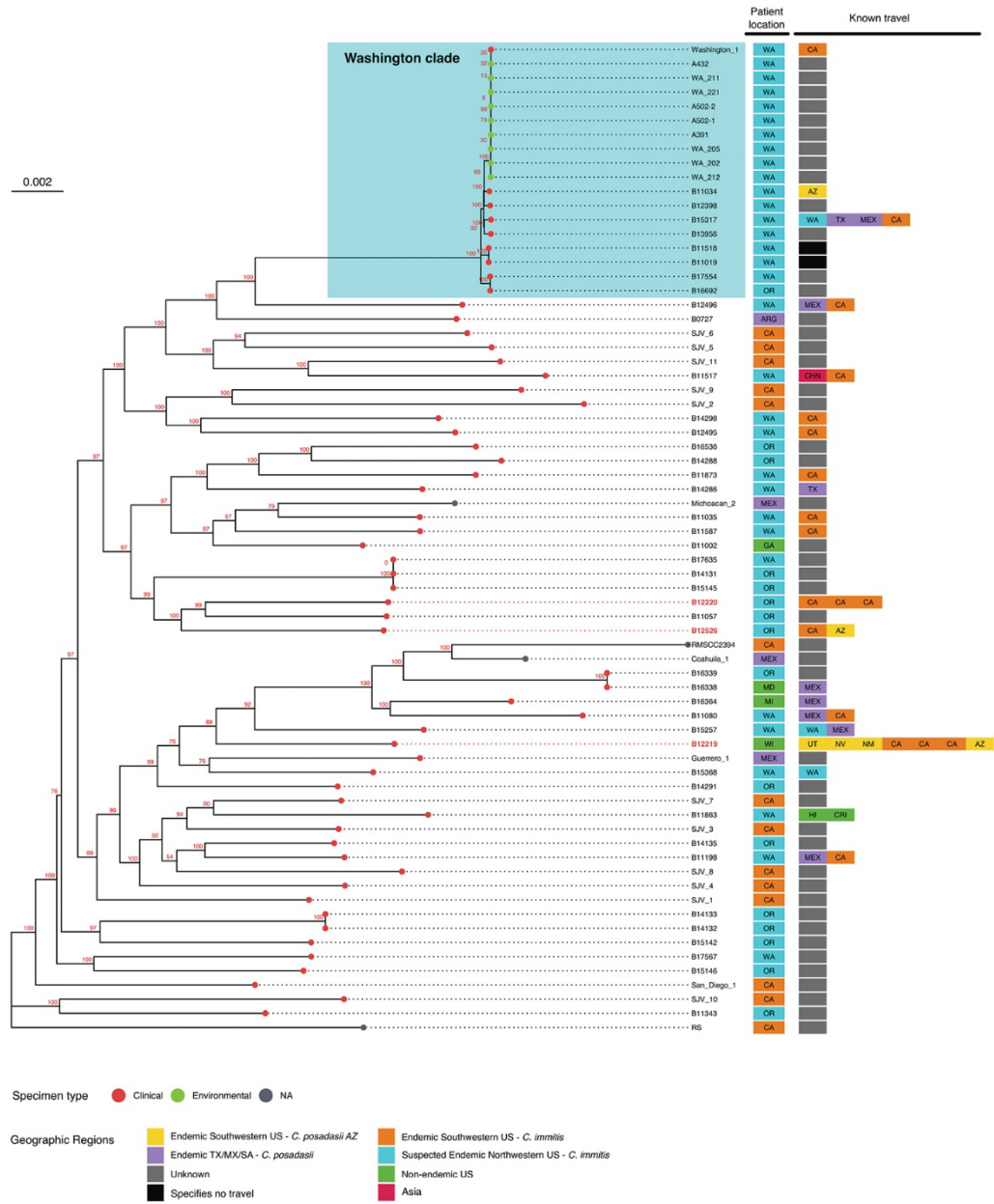


Genomic Epidemiology Linking Nonendemic Coccidioidomycosis to Travel

Appendix 2



Appendix 2 Figure 1. Maximum likelihood phylogenetic tree for *Coccidioidomycosis posadasii*. Each recognized phylogeographic clade is highlighted with a colored box labeled in its top right corner. Each leaf is labeled with locations of isolation and all known patient travel, with colors for the geographic regions. The samples sequenced for this study are shown with red bolded tip labels. Bootstrap values for each node are shown in red.



Appendix 2 Figure 2. Maximum likelihood phylogenetic tree for *Coccidioimycosis immitis*. Each recognized phylogeographic clade is highlighted with a colored box labeled in its top right corner. Each leaf is labeled with locations of isolation and all known patient travel, with colors for the geographic regions. The samples sequenced for this study are shown with bolded tip labels colored by the specimen type. Bootstrap values for each node are shown in red.