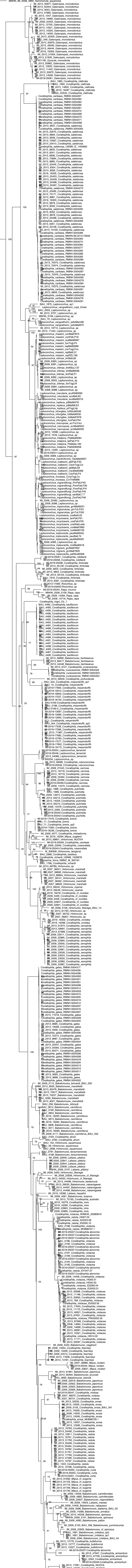
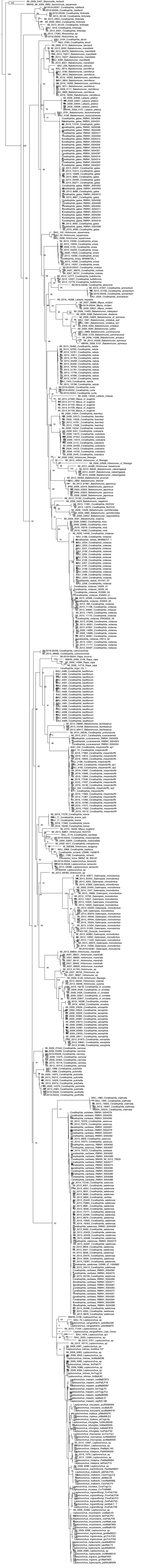


Figure S2. Maximum Likelihood phylogenetic inference on the COI-16S-ITS2 combined dataset. Numbers at nodes indicate ultrafast bootstrap values.









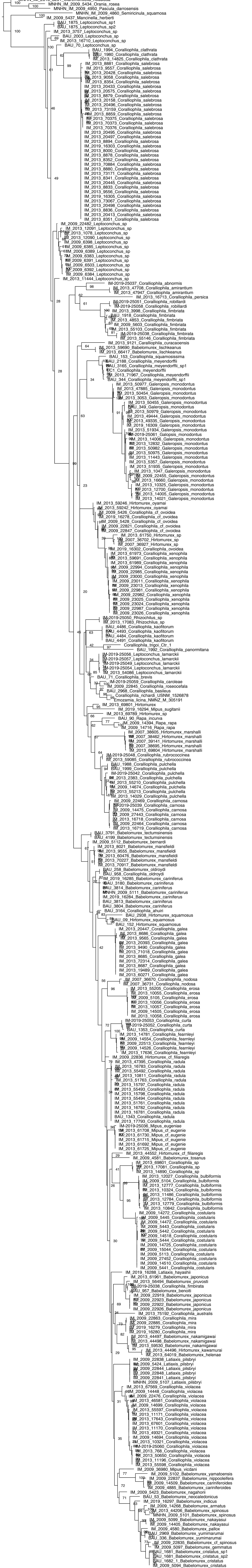
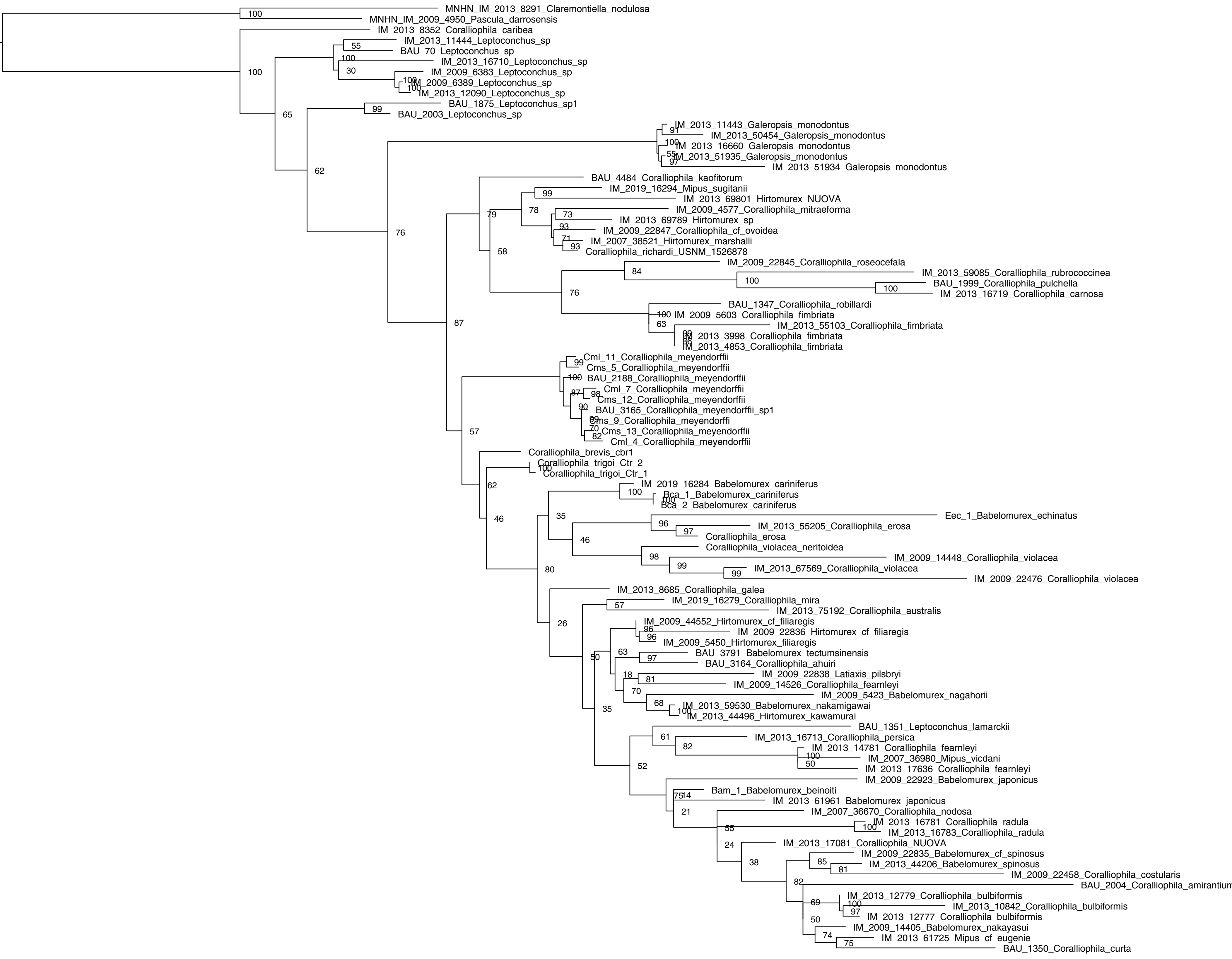


Figure S5. Maximum Likelihood phylogenetic inference on the ITS2 dataset. Numbers at nodes indicate ultrafast bootstrap values.



0.1







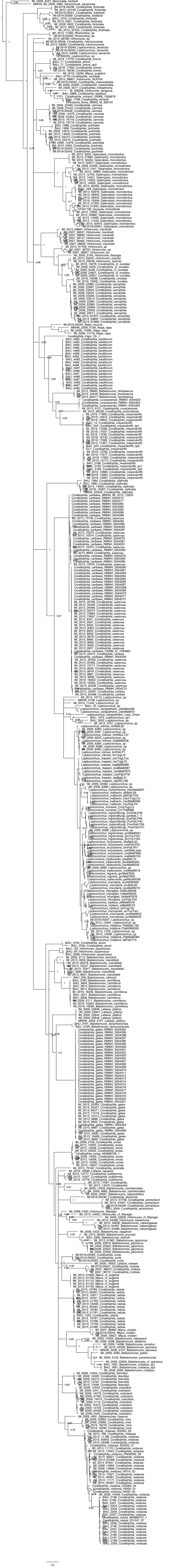
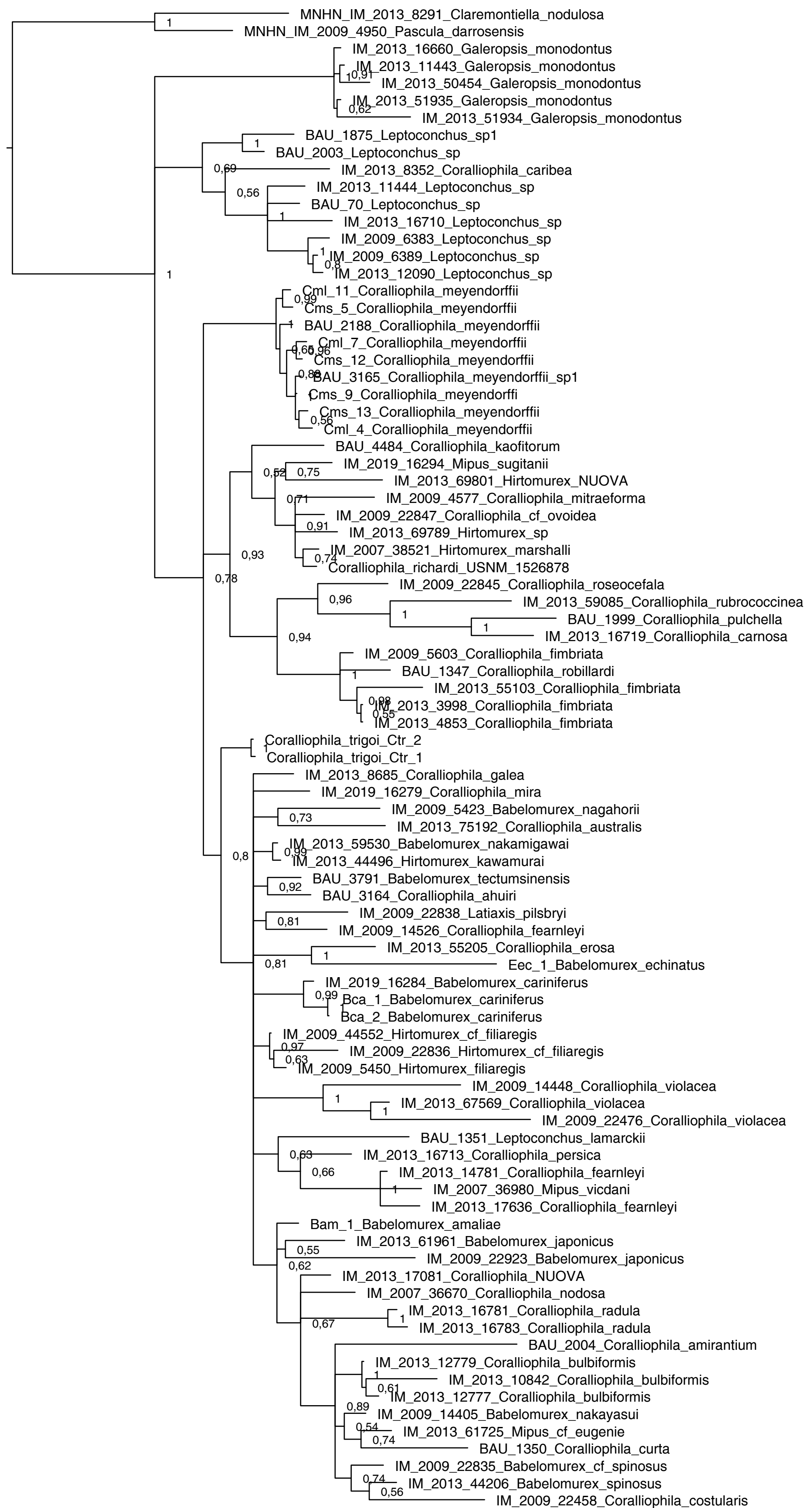






Figure S9. Bayesian phylogenetic inference on the ITS2 dataset. Numbers at nodes indicate posterior probability values.



0.07