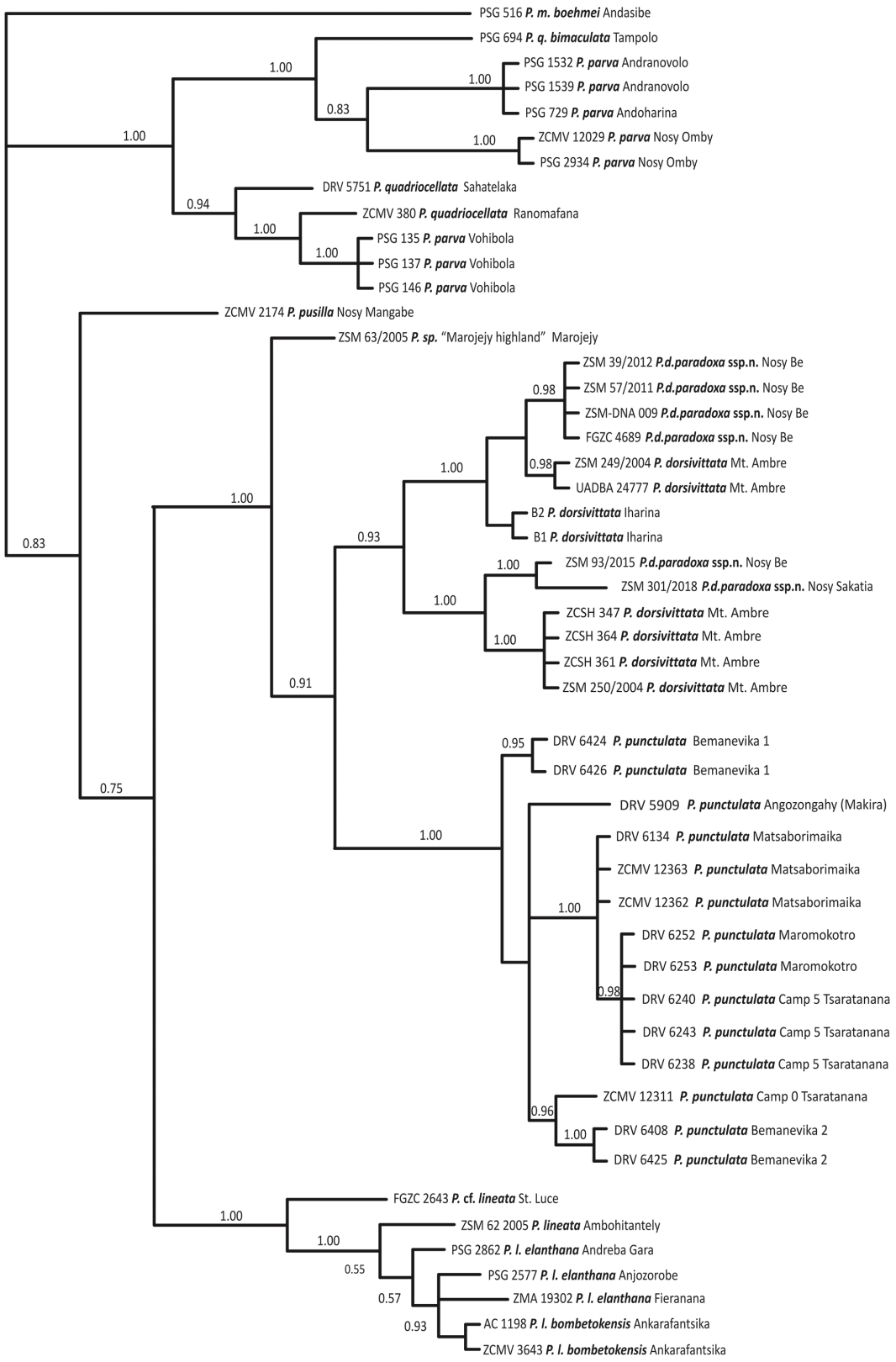

Suplementary Figure S1. Bayesian inference phylogenetic tree of several species in the *Phelsuma lineata* species group, based on 322 bp of the mitochondrial 16S rRNA gene. In the tree, values above branches are Bayesian posterior probabilities. *Phelsuma madagascariensis boehmei* was used as outgroup.