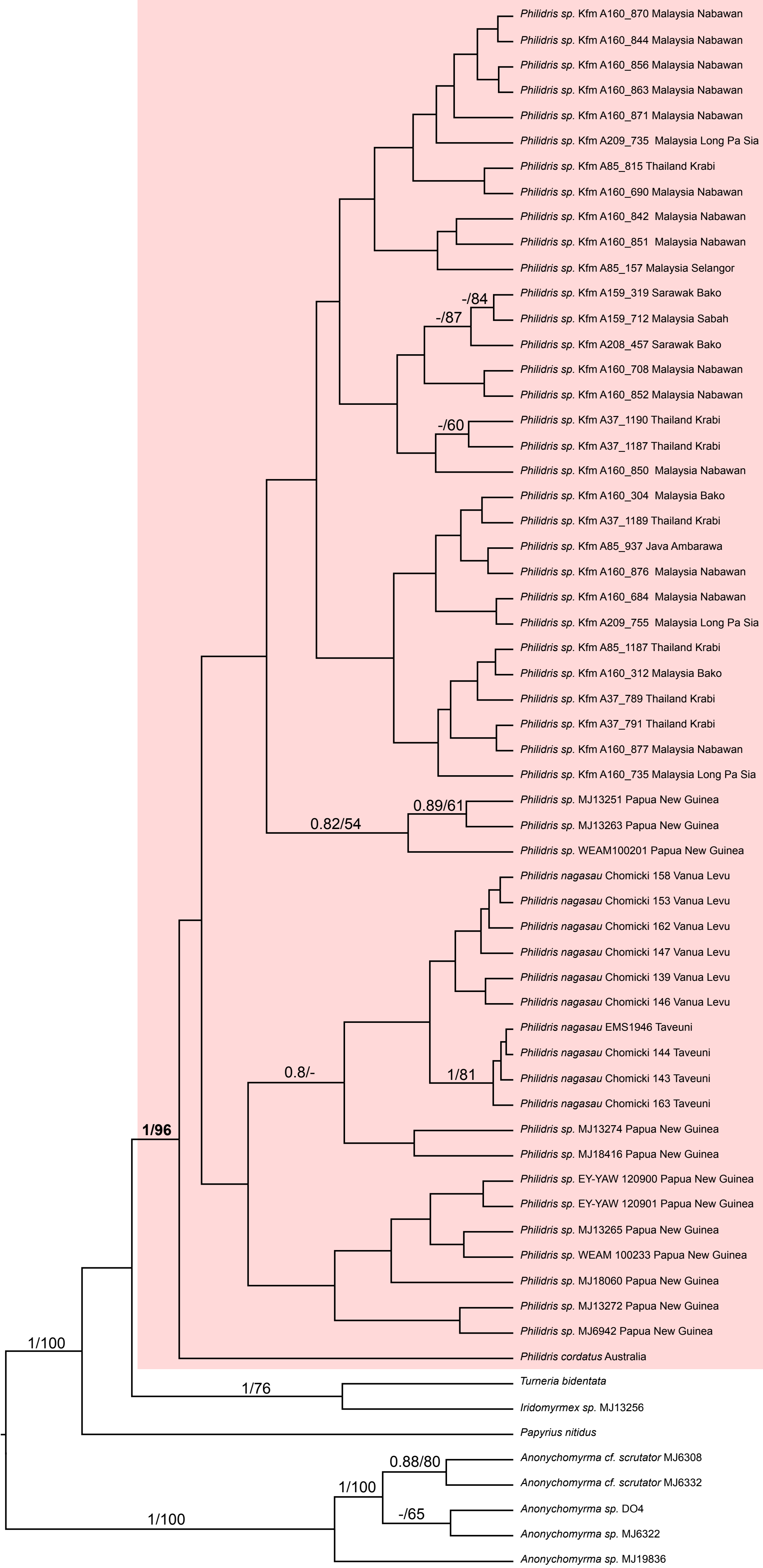

Fig. S1. Phylogeny of *Philidris* ants (shown in red), with support values from likelihood (RAxML) and Bayesian analyses (BEAST) shown when greater than 50% (ML Bootstrap) or 0.8 (Bayesian probabilities).