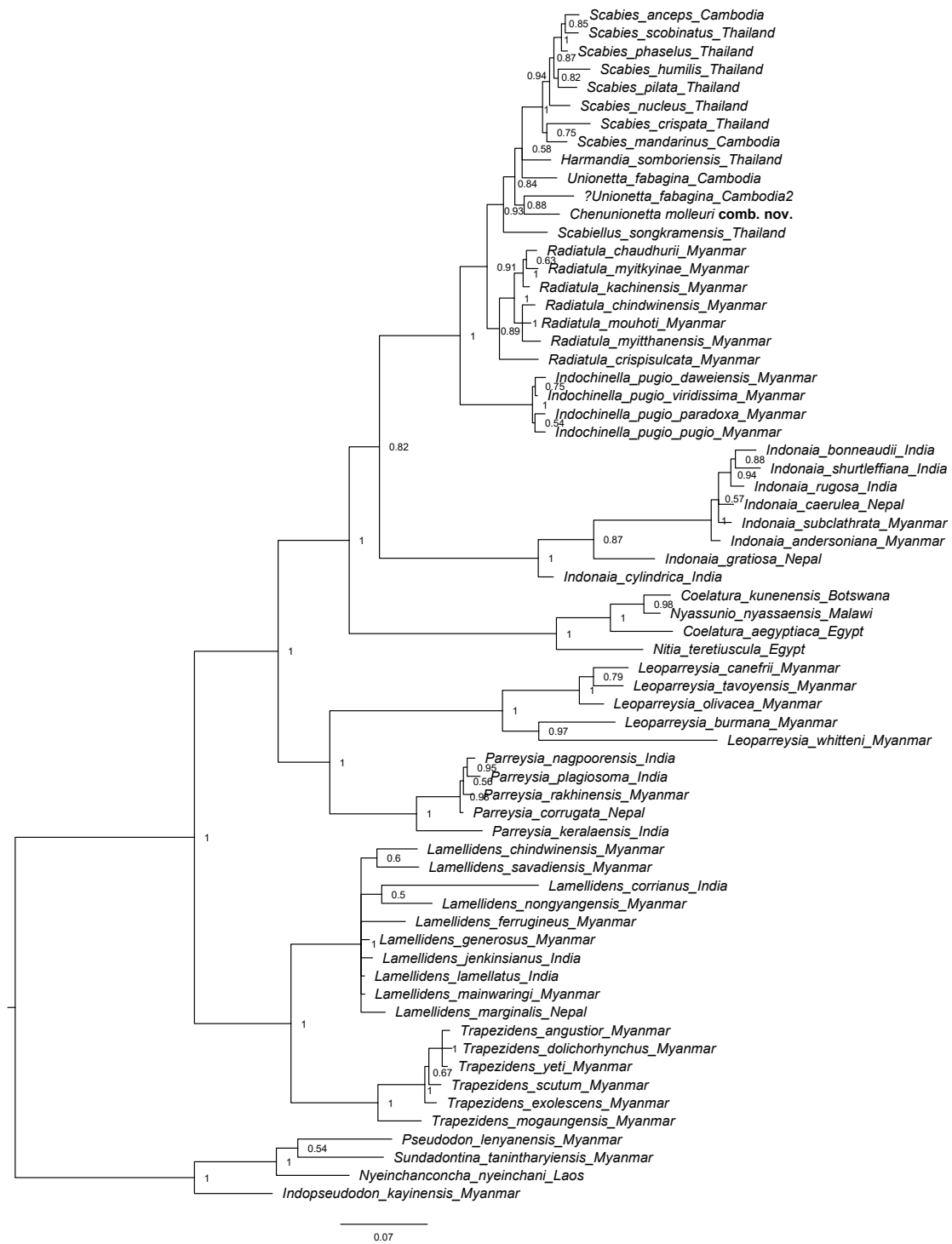




Supplementary Figure S3. Bayesian inference phylogeny inferred from the concatenated three gene dataset (COI, 16S, and 28S). Posterior probabilities are indicated at nodes.