

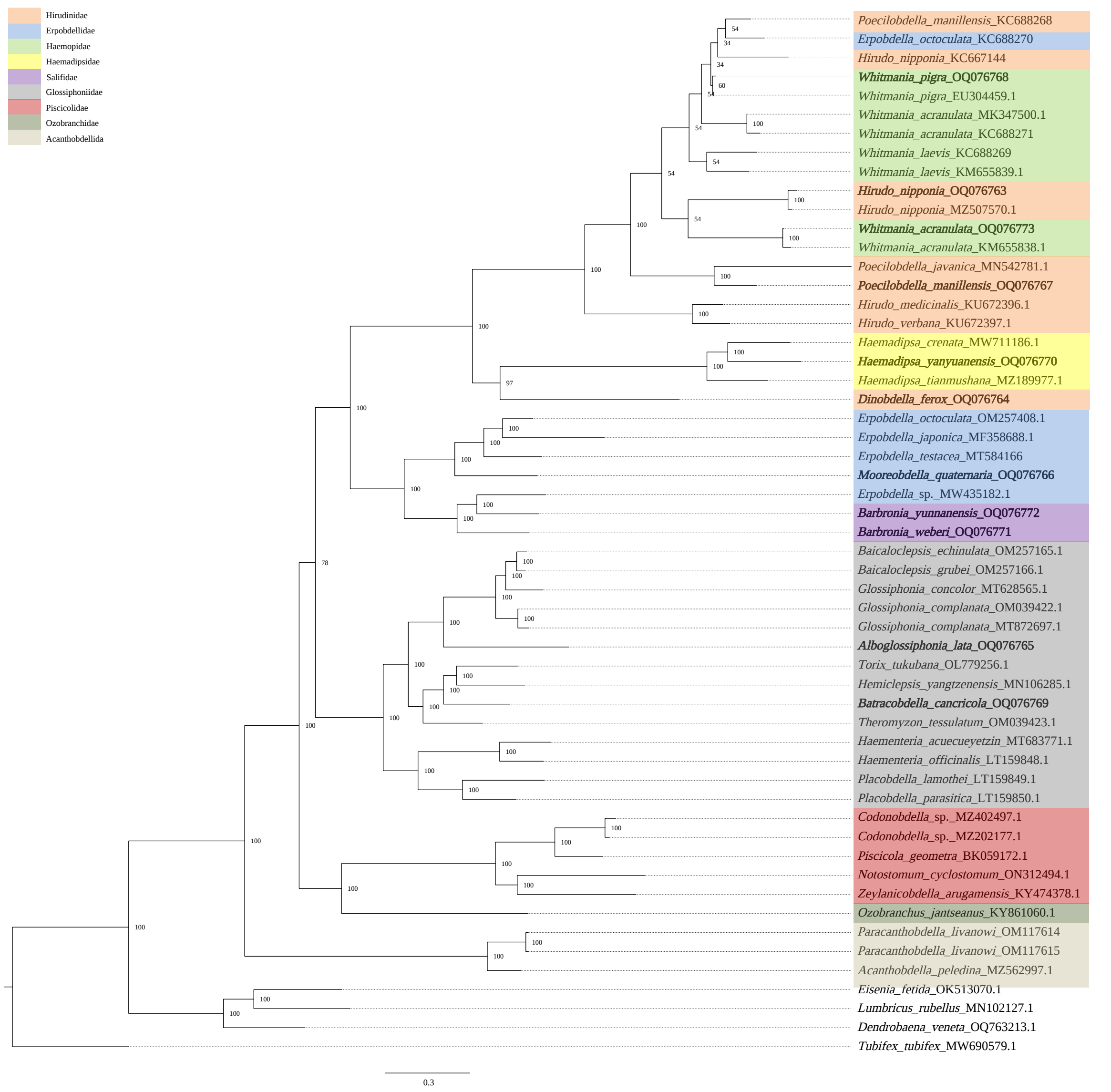


Fig. S12 Phylogenetic relationships inferred from all codon positions of PCGs (PCG123). Numbers are posterior probabilities calculated according to Bayesian inference method (BI tree).