

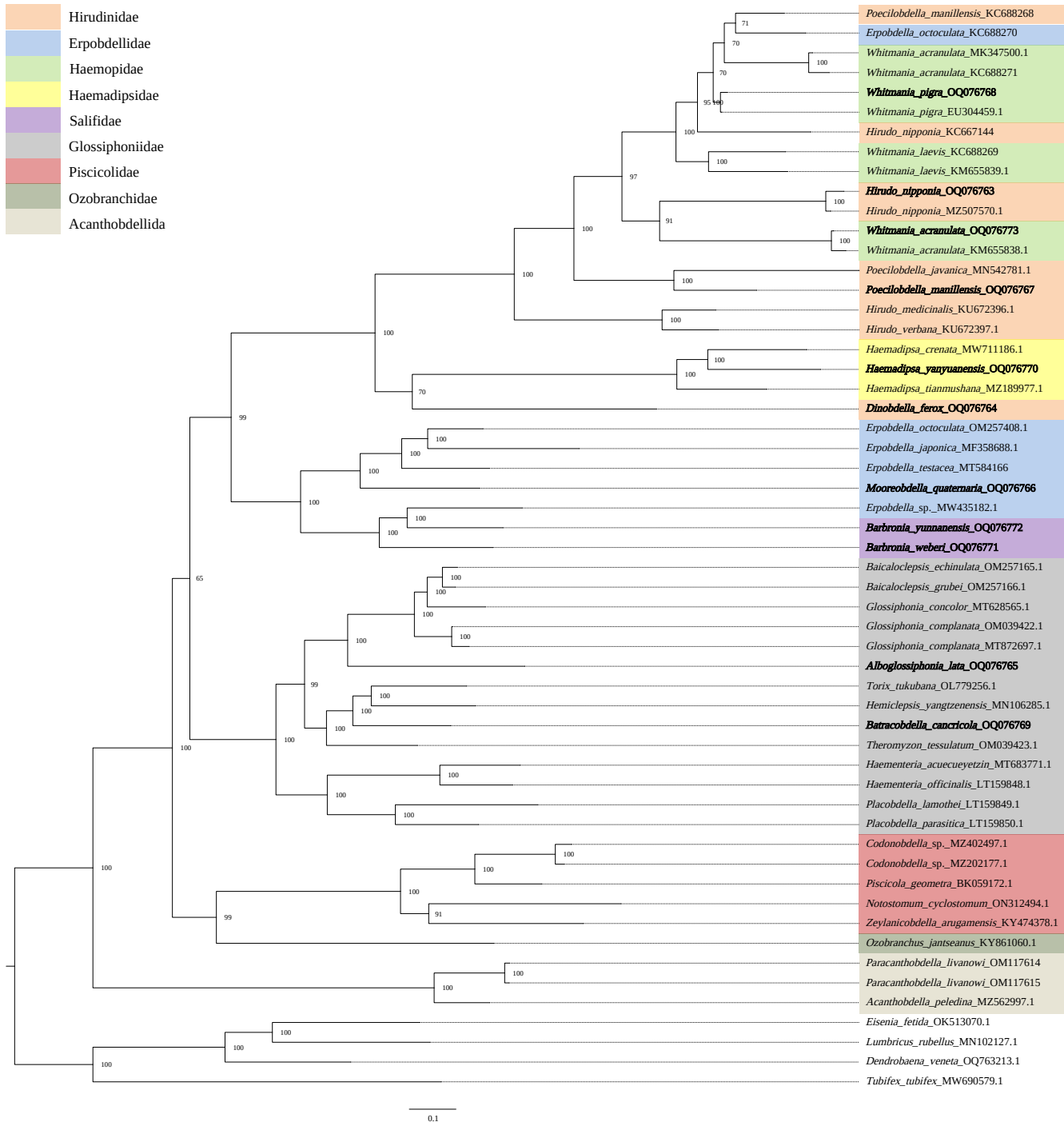


Fig. S10 Phylogenetic relationships inferred from all codon positions of PCGs (PCG123). Numbers are bootstrap values calculated according to maximum likelihood method (ML tree).