

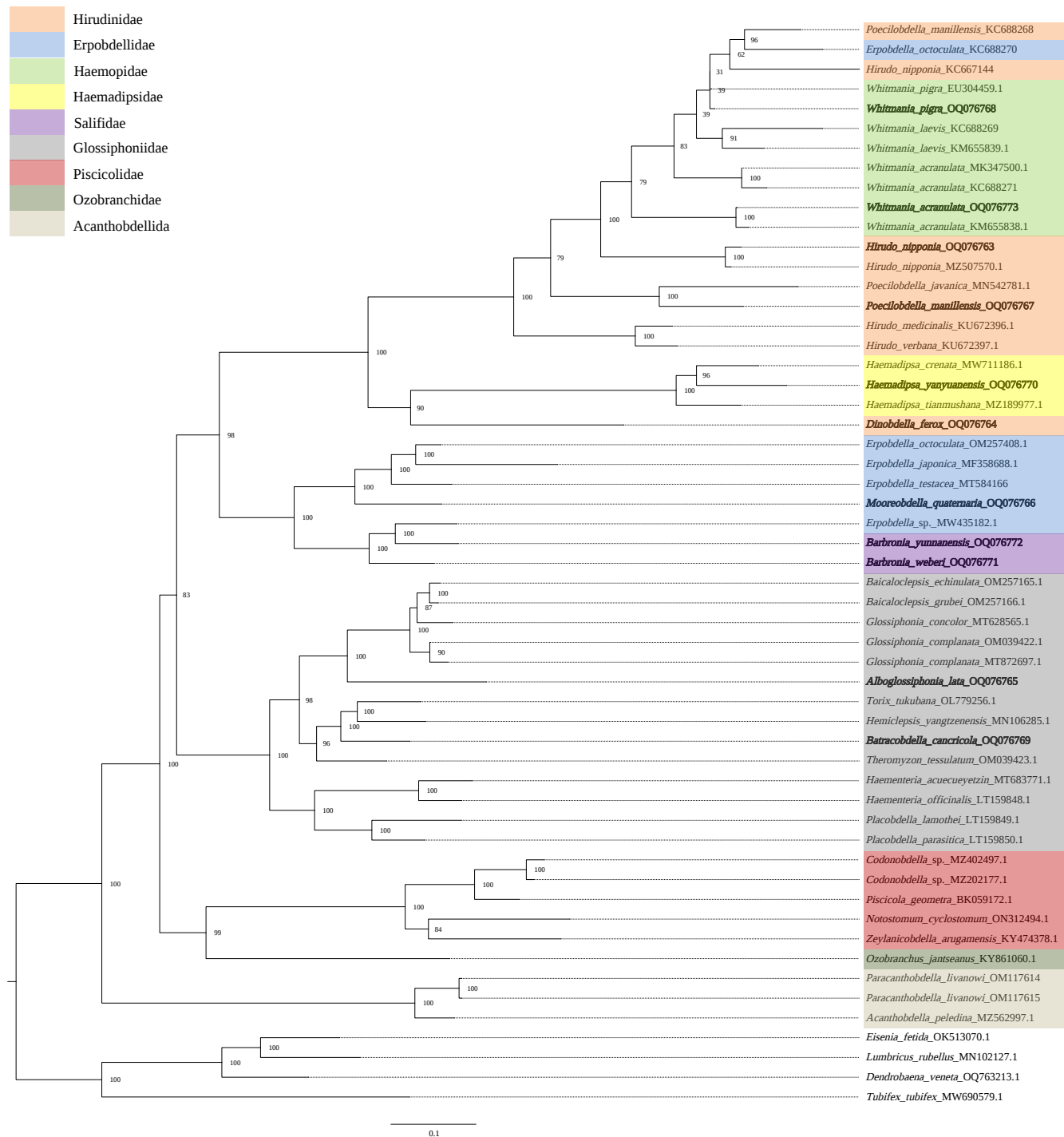


Fig. S8 Phylogenetic relationships inferred from first and second codon positions of PCGs (PCG12). Numbers are bootstrap values calculated according to maximum likelihood method (ML tree).