

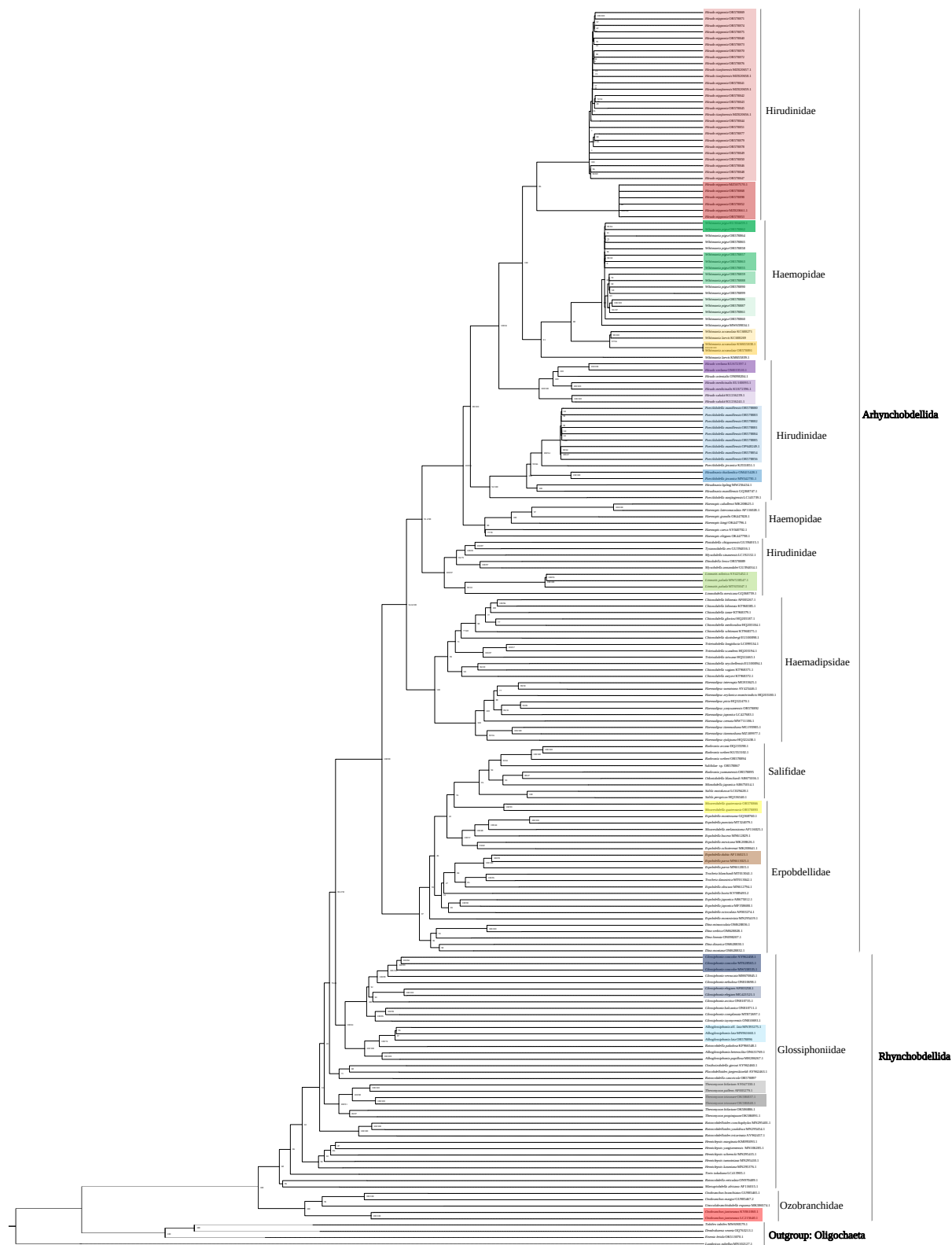


Fig. 6 Maximum likelihood phylogenetic tree based on 190 Hirudinea *COXI* sequences. Numbers are bootstrap values. Species represented by the same color block are delimited as a single MOTU (molecular operational taxonomic unit) by species delimitation analysis, and each unmarked species corresponds to an individual MOTU.