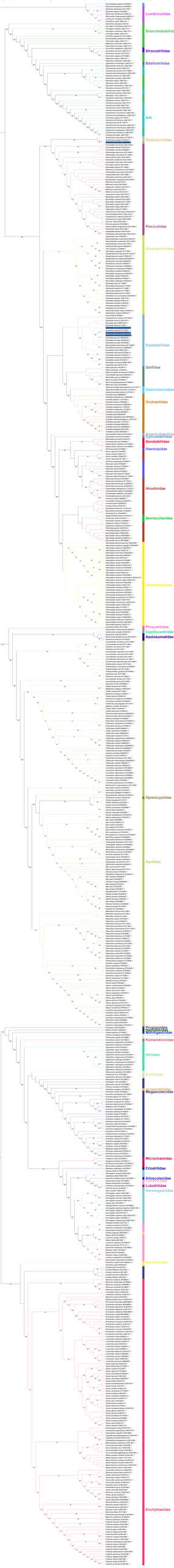Additional file 14. Phylogeny of all Clitellata species in the GenBank. The phylogeny was inferred using the 18S gene and the Maximum Likelihood methodology implemented in IQ-TREE. The figure shows (from left to right): the tree, species and taxonomy (family).