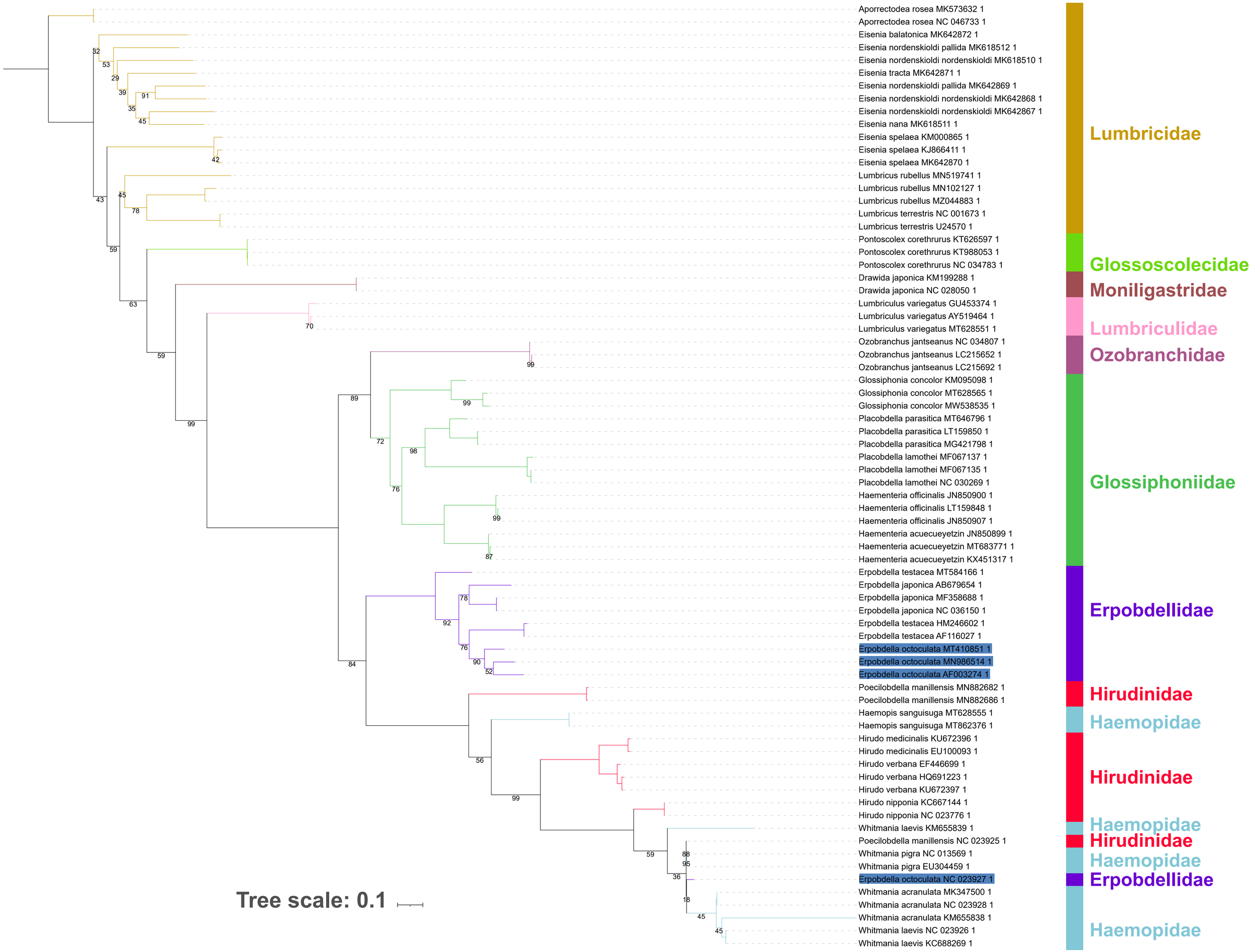Additional file 13. Phylogeny of all Clitellata species from the 174 mitogenomes dataset, minus the Megascolecidae. The phylogram was inferred using the cox1 gene and RAXML. The figure shows (from left to right): a tree with bootstrap values < 100% shown at nodes, species names and taxonomy (family).