Supplementary figures to the paper: Computational identification of functional introns: high positional conservation of introns that harbor RNA genes

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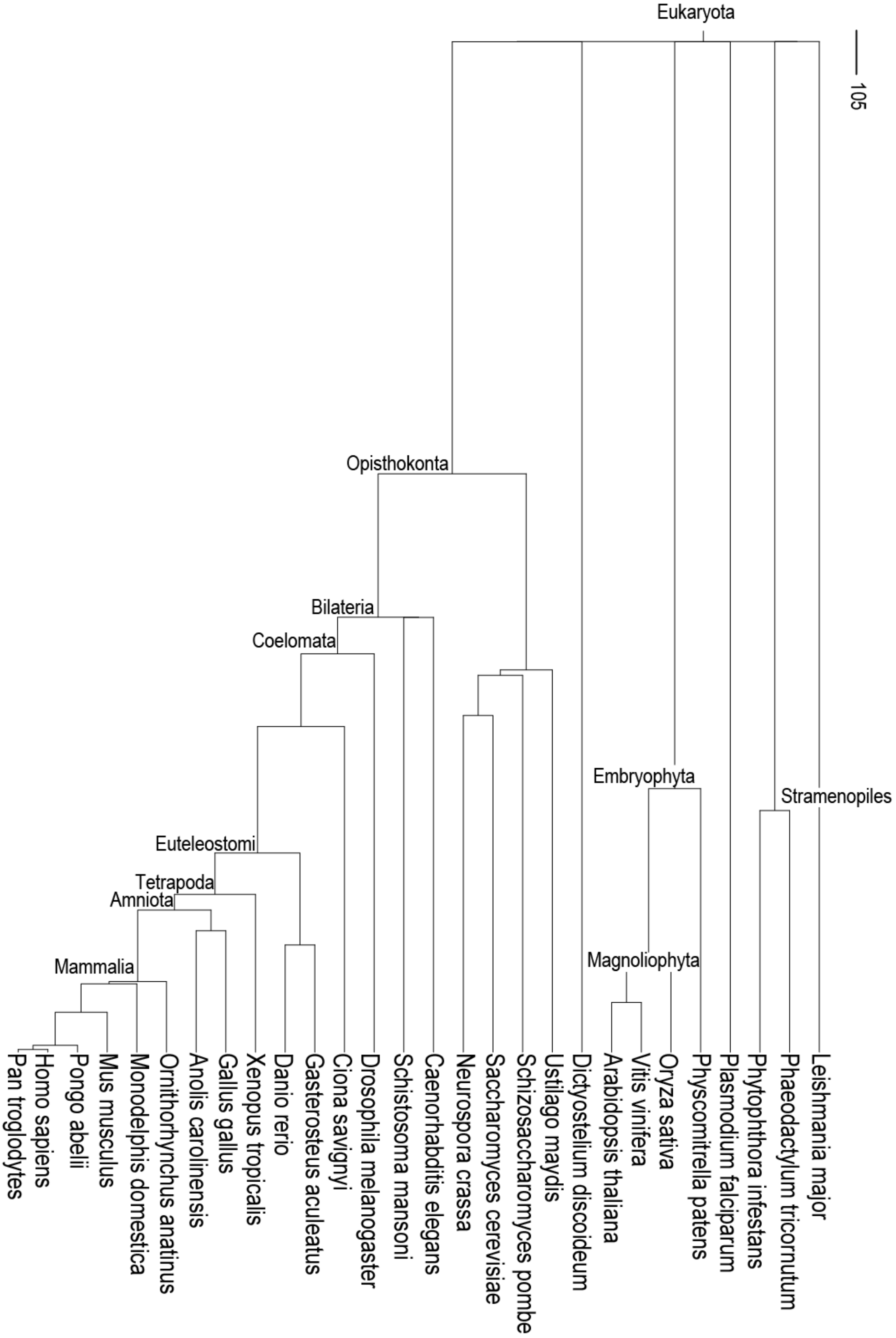

Figure S1. The phylogenetic tree of the final set of 28 species. Scale in million years.

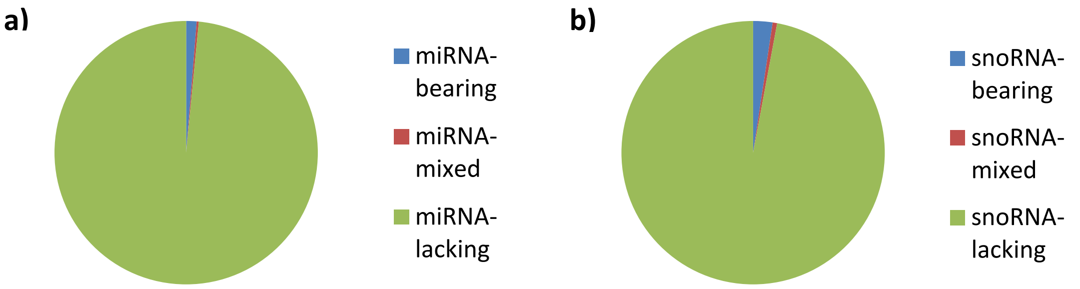

Figure S2. Division of all unique patterns. **a)** There are 53 miRNA-bearing, 11 miRNA-mixed and 4,099 miRNA-lacking unique patterns. **b)** There are 99 snoRNA-bearing, 23 snoRNA-mixed and 4,041 snoRNA-lacking unique patterns.