

Trees in R

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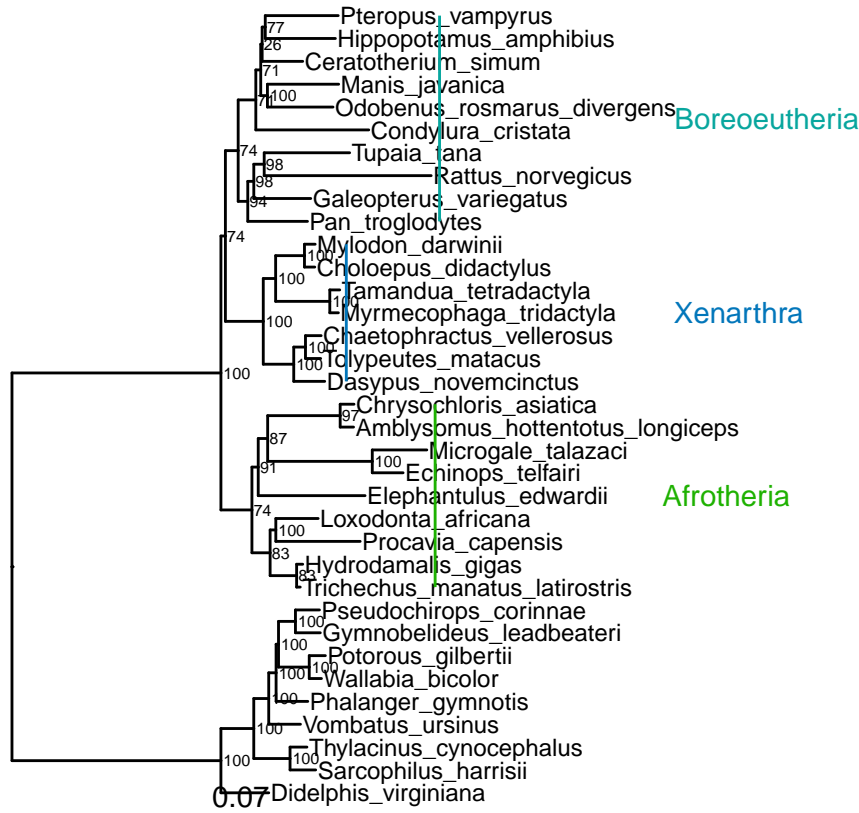


Figure 2. Maximum Likelihood Phylogenetic Analysis performed in RAxML for 30 placental mammal and 10 marsupial outgroup taxa. Orthologous loci were obtained from raw genomic reads using the SISRS pipeline. Support values represent 100 bootstrap replicates. Placental mammal clades are indicated by color-coded vertical lines. These preliminary results appear to support the Afrotheria hypothesis for the branching pattern at the root of placental mammals.