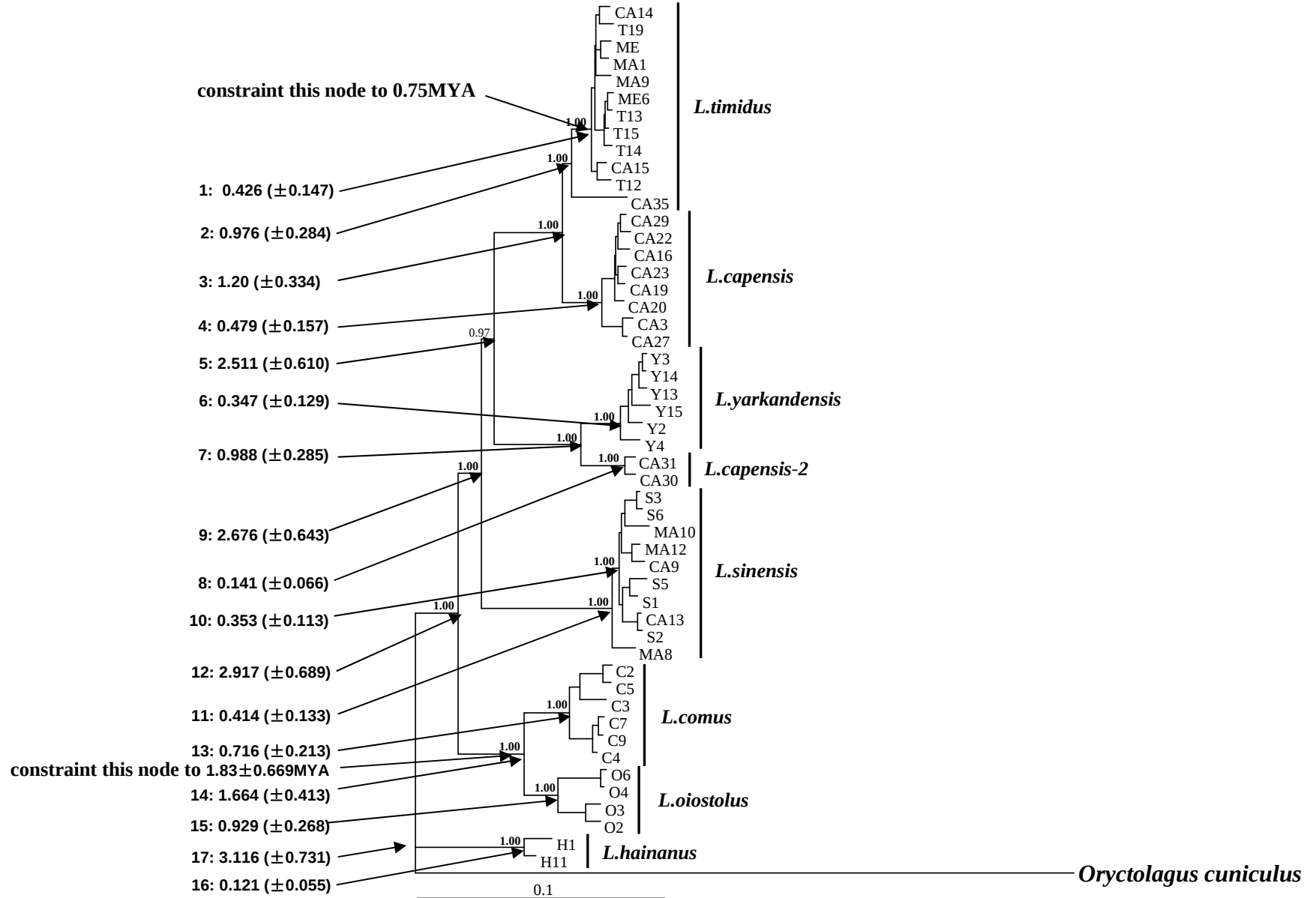




Additional file 3 Bayesian tree used for divergence time estimates and the divergence time for each major clades. Phylogenetic analyses are rooted with *Oryctolagus cuniculus*. The best substitute model GTR+I+G was used for Bayesian inference. The numbers above the branches are the Bayesian posterior probabilities (PP).