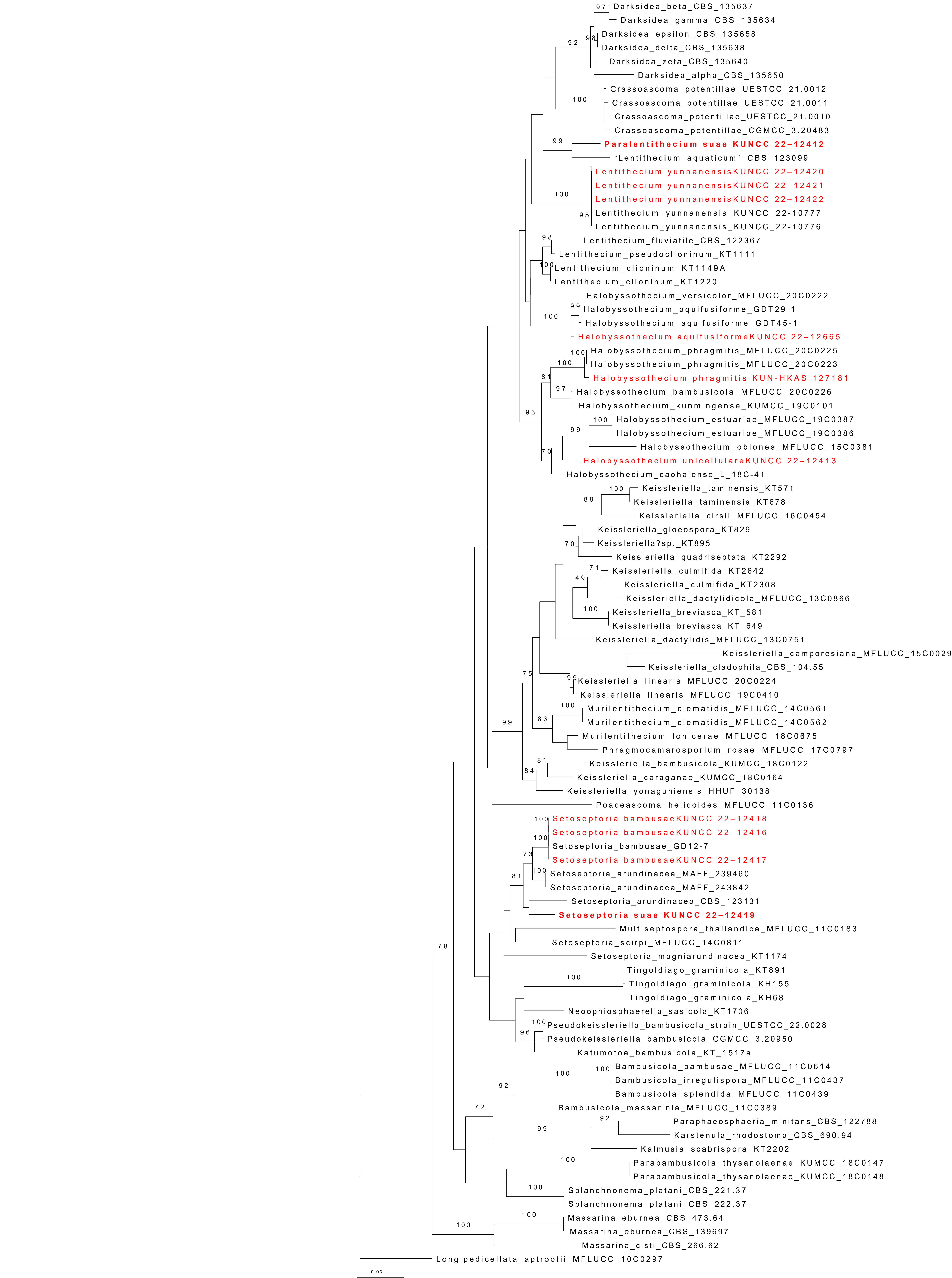




**Figure S4.** ML phylogenetic tree of Lentithecaceae inferred from the TEF sequences (932bp). Support in nodes is indicated above by bootstrap values of more than 70 %. New species are indicated in red bold. New strains are indicated in red.