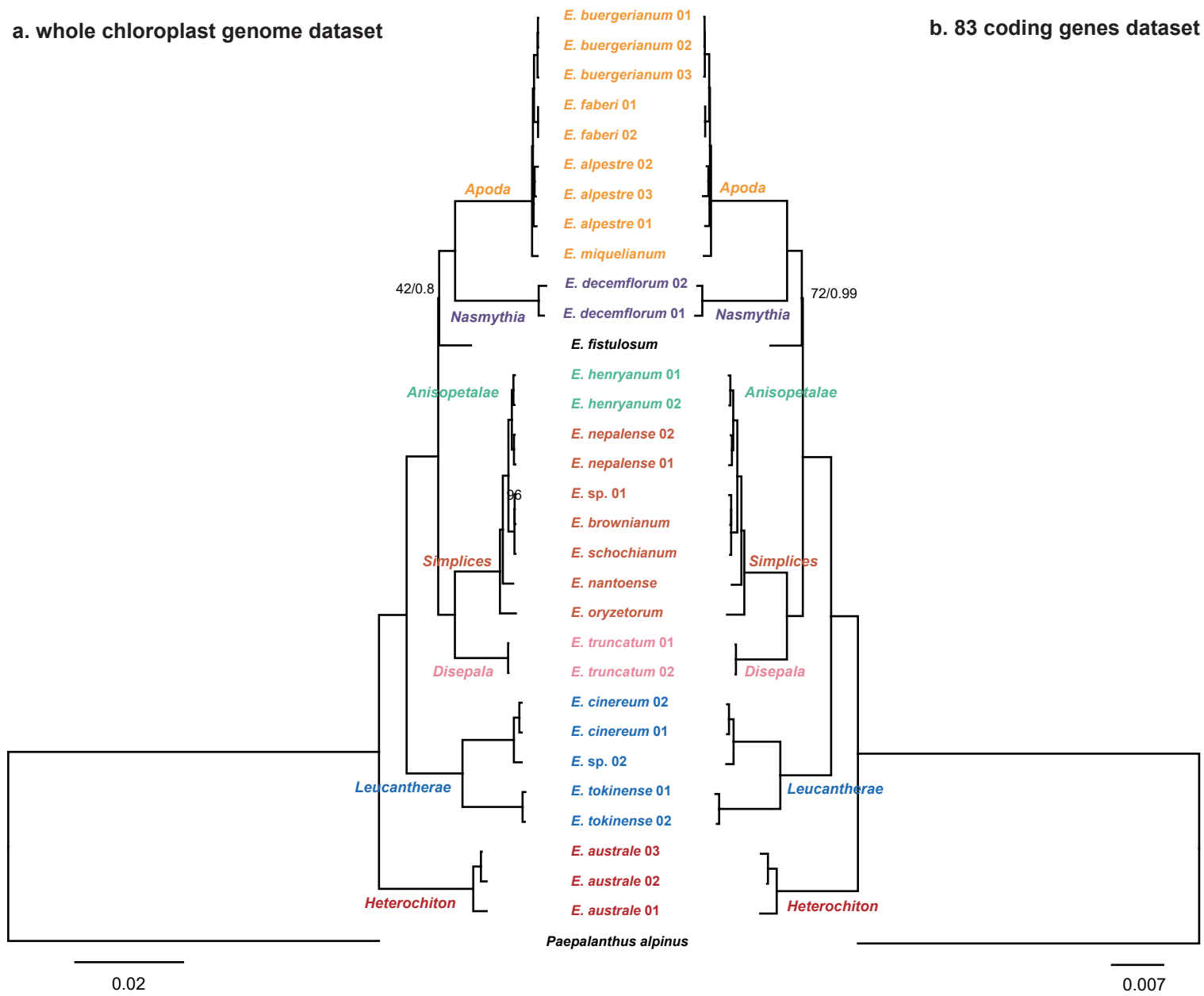


a. whole chloroplast genome dataset

b. 83 coding genes dataset



**Figure S3.** Phylogenetic trees of *Eriocaulon*. a. The whole chloroplast genome dataset. b. The 83-genes dataset. The number above the lines indicates the ML bootstrap values (BS) and BI posterior probability (PP). BS=100 and PP=1.0 are not shown.