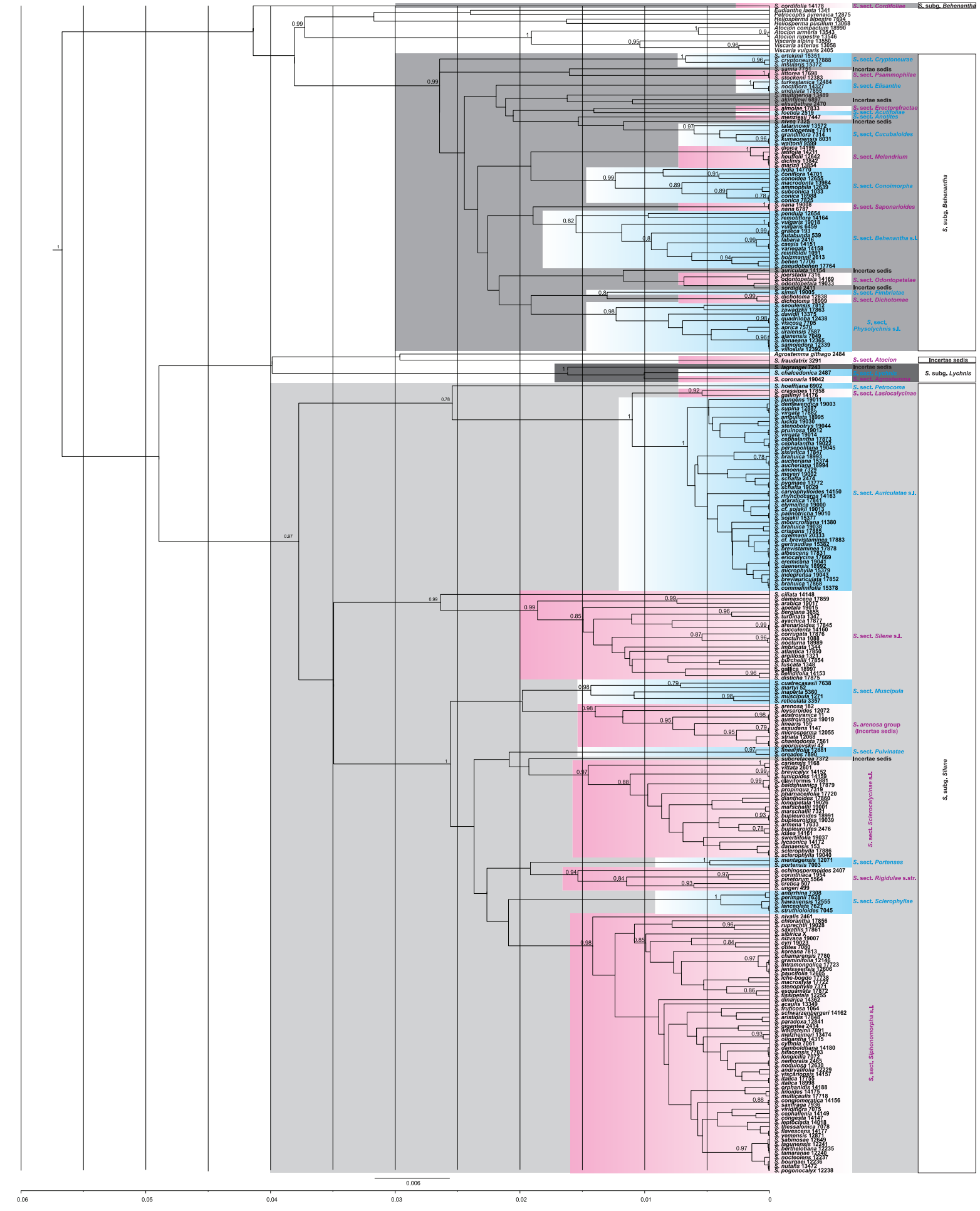

Fig. S1. Species or minimal clusters (SMC) tree from the nrDNA ITS and cpDNA rps16 datasets based on GTR model. Posterior probability values of the analysis based on the HKY model are indicated above branches.