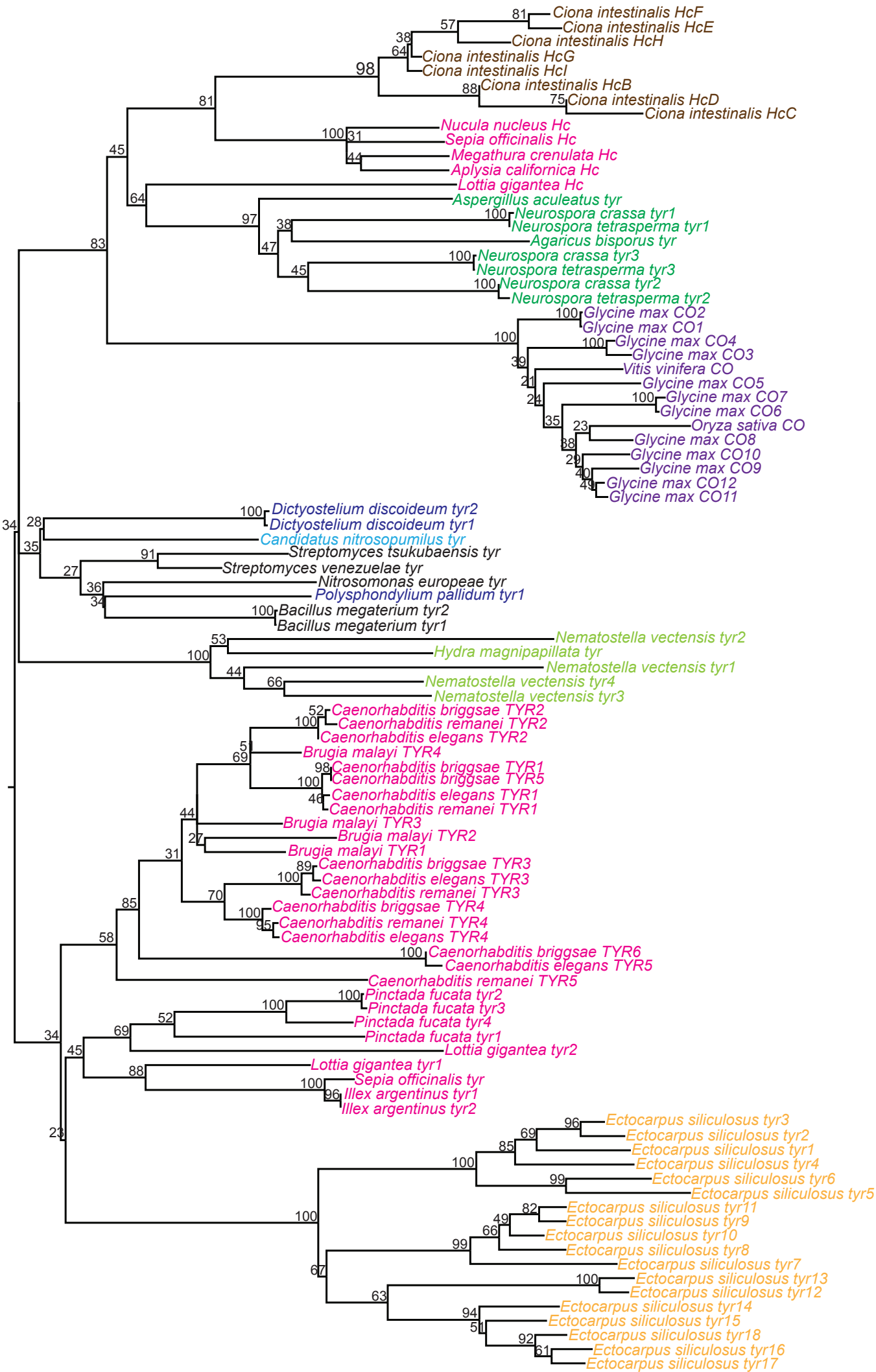


A

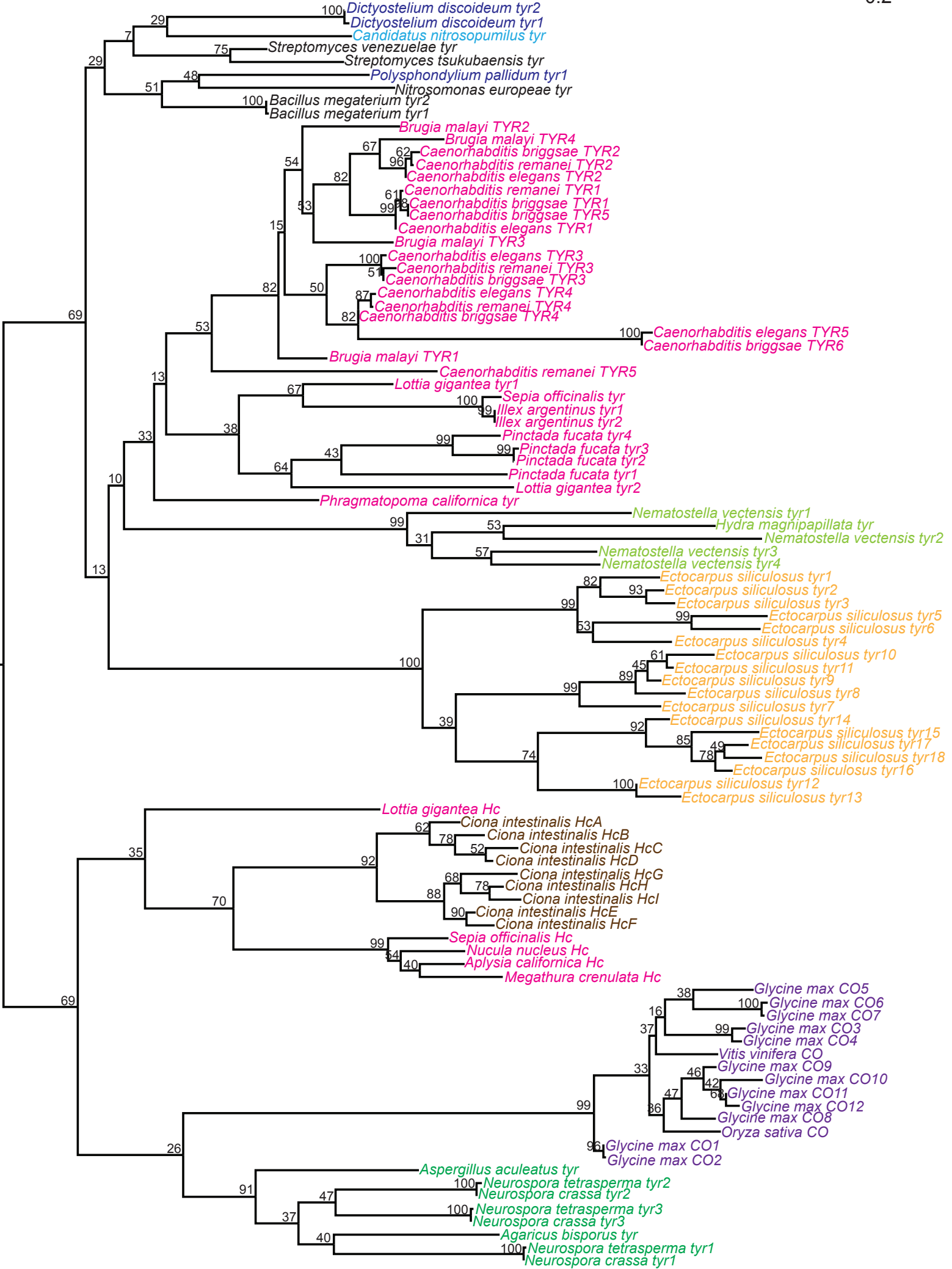
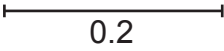
α -subclass phylogenetic tree obtained by Neighbor-Joining method

0.2



B

α-subclass phylogenetic tree obtained by Maximum-Likelihood method



C

 α -subclass phylogenetic tree obtained by Bayesian-Inference method

0.3

