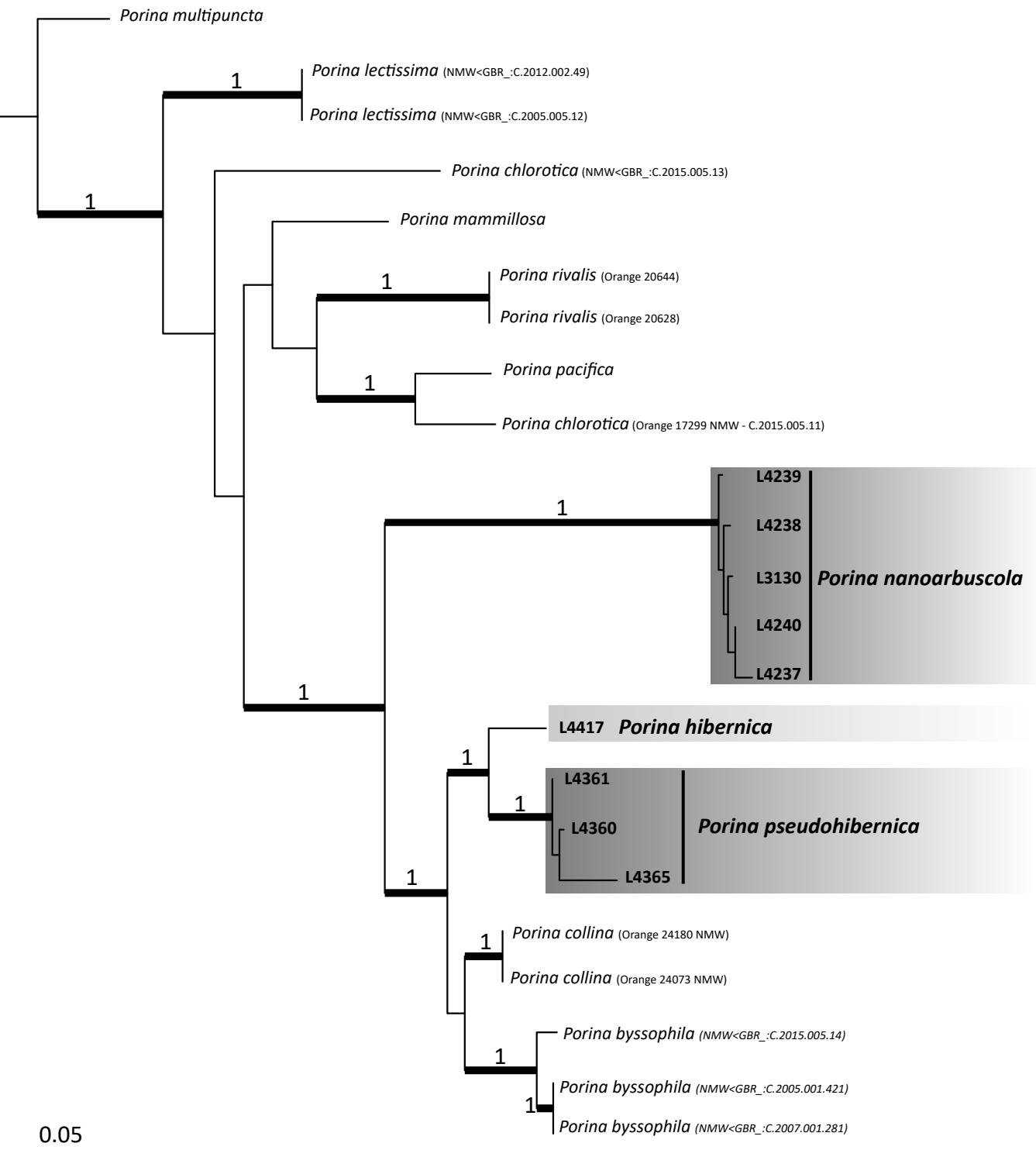


*Porina austroatlantica*

Outgroup



Supplementary Material Fig. S1. Phylogenetic hypothesis based on the combined ITS and mtSSU loci; 50% majority rule consensus tree obtained by Bayesian analysis. ML bootstrap values >70% are in boldface branches; Bayesian PP values > 0.8 are reported above branches. The newly sequenced samples are highlighted in bold.