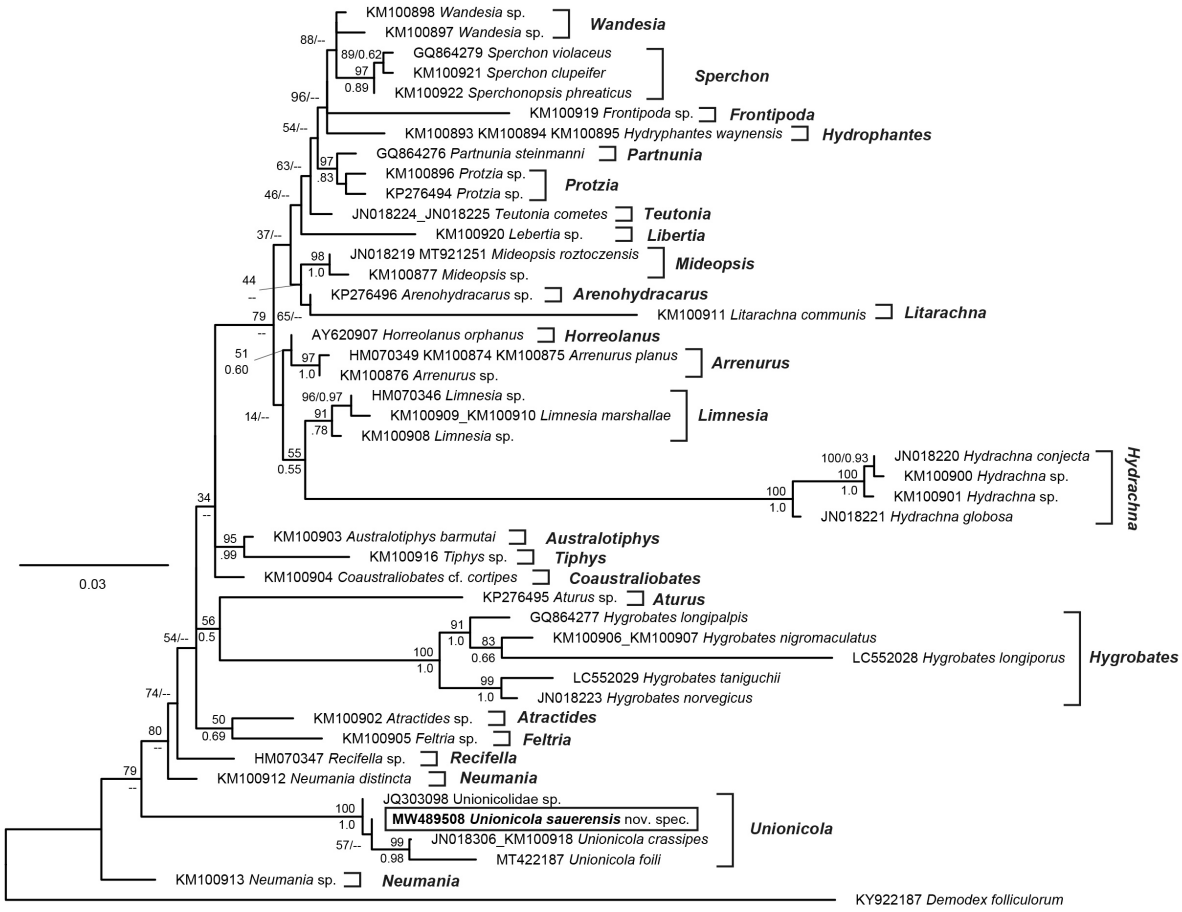
Supporting Information Figure S1



Maximum Likelihood tree based on a 630 bp *18S* section from members of the superfamily Hygrobatoidea. Maximum Likelihood bootstrap support values and Bayesian Inference posterior probabilities are provided for most nodes. The scale bar indicates the number of substitutions according to the model selected.