

Supplementary material

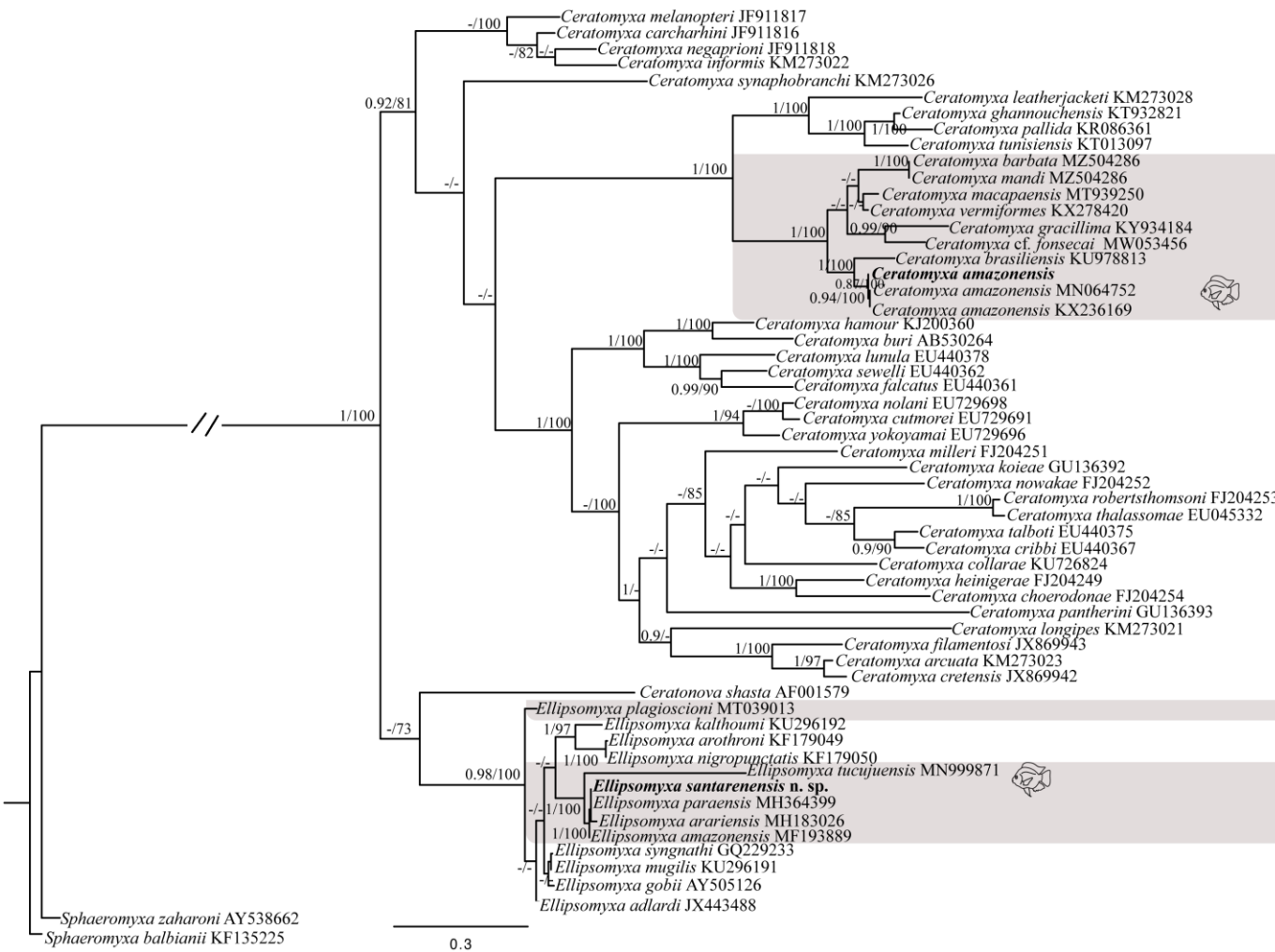


Figure S1. Maximum Likelihood (ML) consensus phylogenetic tree obtained from the SSU-rDNA sequences from *Ellipsomyxa santarenensis* n. sp., parasites of *Satanoperca jurupari*. GenBank accession numbers are presented in front of the name of each species.