

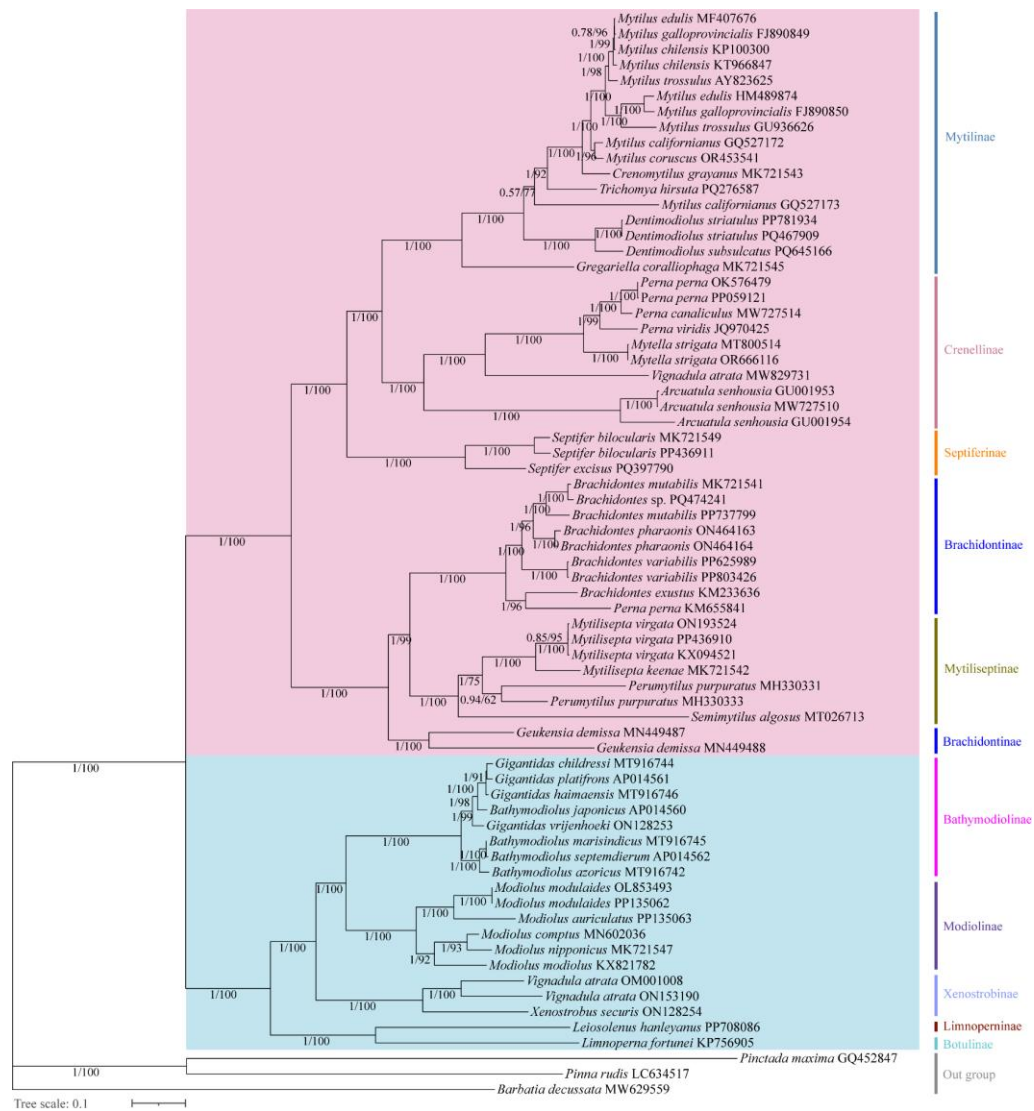


Figure S6. The phylogenetic tree based on mitochondrial amino acid sequences of Mytiloidea. The mitogenomes newly sequenced in this study are marked in red. Posterior probability (PP) / bootstrap (BS) are shown at nodes