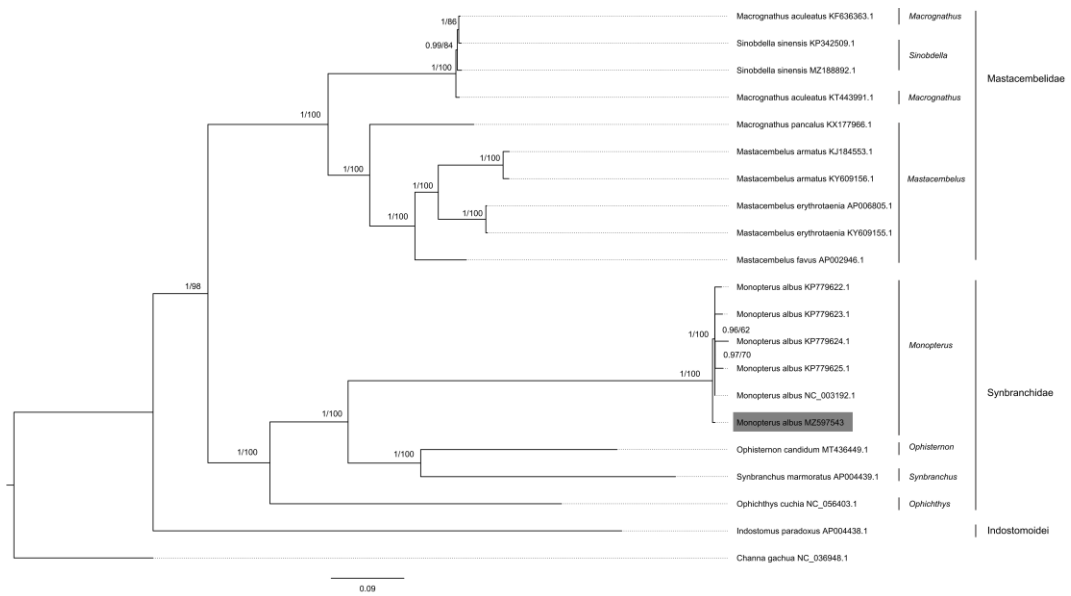

Figure S2: The phylogenetic tree of order Synbranchiformes inferred from the concatenated 13 PCGs, 2 rRNAs, 22 tRNAs and one D-loop of 20 complete mitochondrial sequence obtained using MrBayes v.3.2.6 and IQ-TREE v.1.5.5. *Channa gachua* (NC_036948.1) was used as outgroup.