



Fig. S5 Phylogenetic reconstruction by maximum likelihood based on combined protein-coding subunits concatenated of the species sequenced in this study and two other taxa with data available from GenBank. a The combination of ND2 and ND4 subunits. b The combination of COI, ND2, ND4 and ND5 subunits. The values of bootstrapping support are shown to the left of the branch point.