

Supplementary Fig. S1 The comparison of bGMYC results between COI and COI haplotype data sets. The tree on the left is the BEAST tree inferred from the entire COI data set, with ABGD results on the corresponding clades. GAB, GA and Gb within the parentheses stands for PHSS suggested by, respectively both ABGD and bGMYC, or only by ABGD or bGMYC. The tree on the right is the BEAST tree estimated from the COI haplotypes data set. Individuals and their haplotypes are connected by lines between two trees. Two grayscale histograms represent the uncertainty of species delimitations in bGMYC analysis.

