

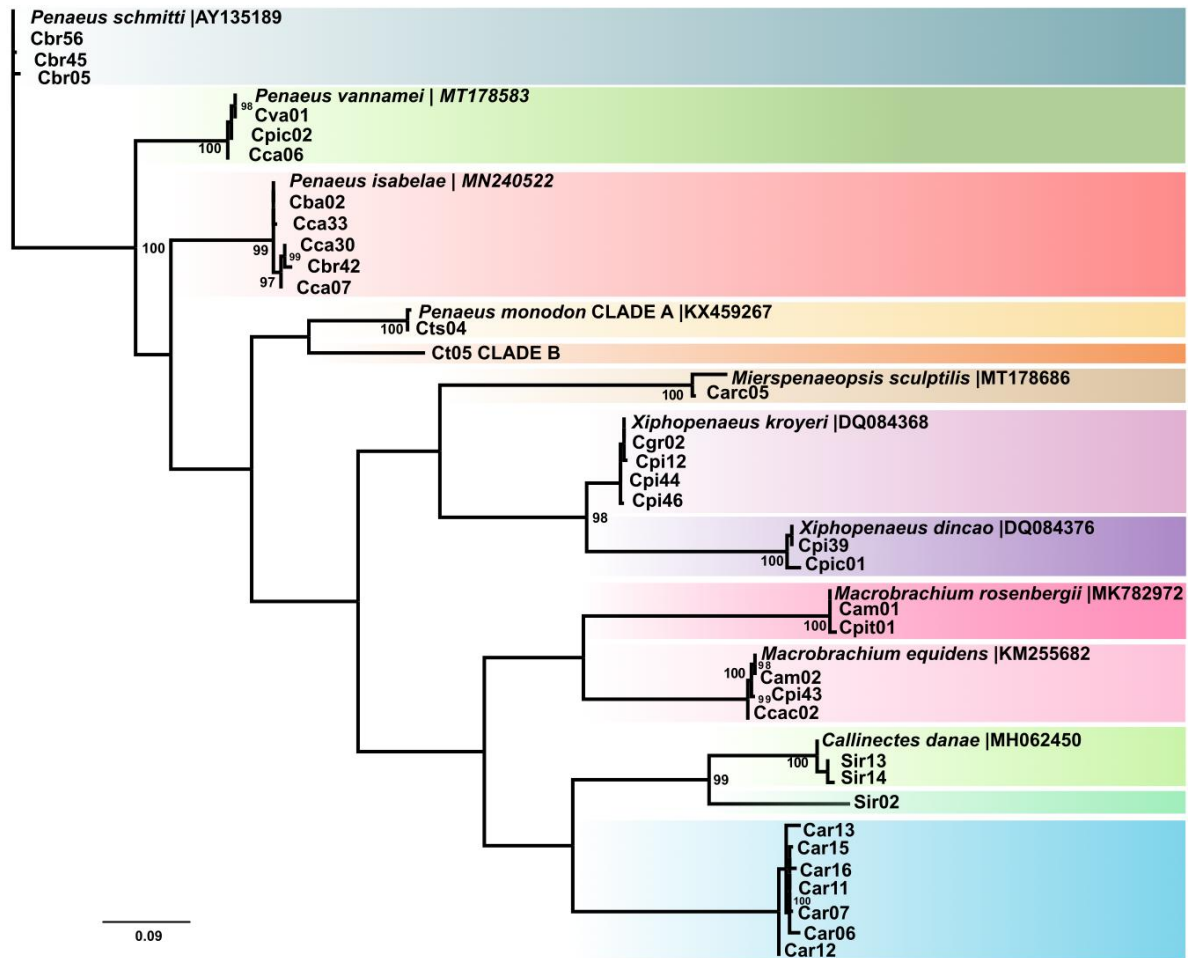


Fig. S1. Maximum likelihood (ML) tree showing the haplotypes of crustaceans based on the second region (region II) of the Cytochrome C Oxidase Subunit I (COI) gene, including the reference sequences from public databases. The statistical support values are shown on internodes.

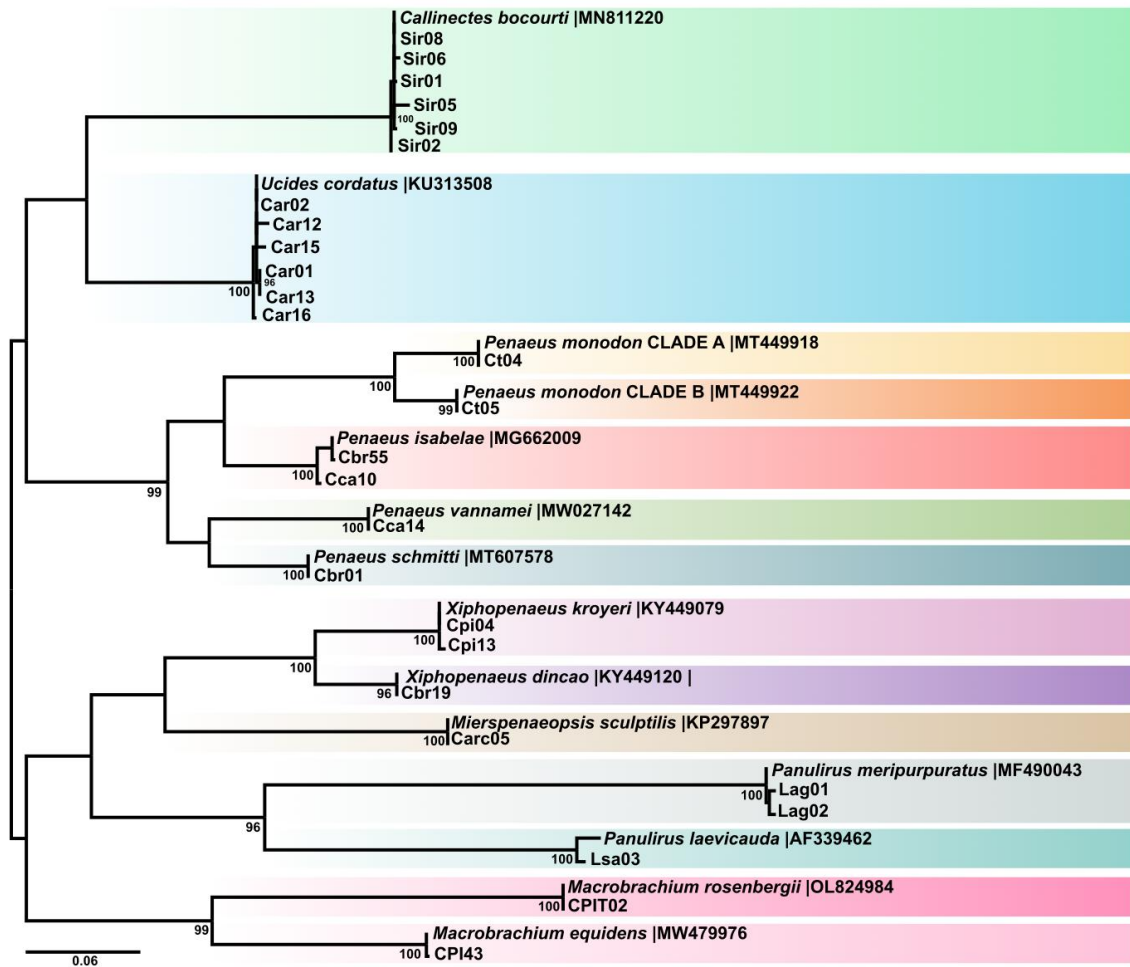


Fig. S2. Maximum likelihood (ML) tree showing the haplotypes of crustaceans based on the first region (barcode region or region I) of the Cytochrome C Oxidase Subunit I (COI) gene, including the reference sequences from public databases. The statistical support values are shown on internodes.