

Table S1. Matrix of minimum percentage pairwise nucleotide divergence with Kimura 2-parameter (K2P) distances (lower left) and mean values (upper right) based on cytochrome c oxidase subunit I (COI) between 71 speies of Portunoidea from Taiwan (see Table 1).

Table with 71 rows and 71 columns. Rows are grouped into Caruripinae, Lupoeyclinae, Neconectinae, Podophthalminae, Portuninae, and Thalamitinae. Each cell contains a numerical value representing genetic distance or divergence percentage.