Supplementary material

Phylogeography and species delimitation of Cherax destructor (Decapoda: Parastacidae) using genome-wide SNPs

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\textbf{Fig. S1.} Maximum-likelihood tree for 381 Cherax individuals (see following pages), based on an analysis of 9140 concatenated sequence fragments. Each operational taxonomic unit (OTU) code is based on the site number and locality as per Table 1 and Fig. 1.