

Online Resource Fig. S1 Phylogram from neighbour-joining (NJ) analysis of the *cox1* sequence alignment (42 sequences; 636 nt) for 36 representatives of the subfamily Amphipepleinae (Mollusca: Lymnaeidae). Outgroup: *Galba truncatula*. Bootstrap support values ≥ 70 are provided above branches. The scale-bar indicates the expected number of substitutions per site

