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Supplementary material, Fig. S8

Figure S8. Maximum likelihood phylogeny of terrestrial isopod 18*S* sequences aligned with the L-INS-I algorithm. Sequences are coloured by family. Families represented by a single sequence are coloured black. *Nagurus cristatus* sequences (Trachelipodidae) are at the top of the phylogeny in a clade containing Scyphacidae, Porcellionidae, and *Levantoniscus*. Numbers above nodes show ultrafast bootstrap support. Nodes with < 70% ultrafast bootstrap support are collapsed.