Supplementary Material

Combining mitochondrial DNA and morphological data to delineate four new millipede species and provisional assignment to the genus *Apeuthes* Hoffman & Keeton (Diplopoda: Spirobolida: Pachybolidae: Trigoniulinae)

Piyatida Pimvichai^{A,*}, Somsak Panha^B and Thierry Backeljau^{C,D}

^ADepartment of Biology, Faculty of Science, Mahasarakham University, Maha Sarakham 44150, Thailand.

^BAnimal Systematics Research Unit, Department of Biology, Faculty of Science, Chulalongkorn University, Bangkok 10330, Thailand.

^CRoyal Belgian Institute of Natural Sciences, Vautierstraat 29, BE-1000 Brussels, Belgium.

^DEvolutionary Ecology Group, University of Antwerp, Universiteitsplein 1, BE-2610 Antwerp, Belgium

^{*}Correspondence to: Email: piyatida.p@msu.ac.th

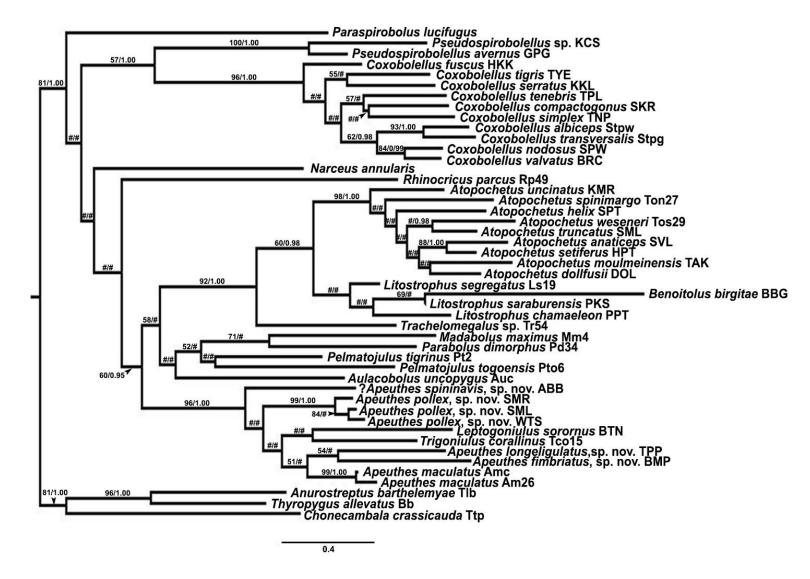


Fig. S1. Phylogenetic relationships of the subfamily Trigoniulinae based on maximum likelihood analysis (ML) and Bayesian Inference (BI) of 660 bp of cytochrome *c* oxidase I (*COI*). Numbers at nodes indicate branch support based on bootstrapping (ML) and posterior probabilities (BI). Scale bar: 0.4 substitutions per site. The number sign (#) indicates branches with <50% ML bootstrap support and <0.95 Bayesian posterior probability.

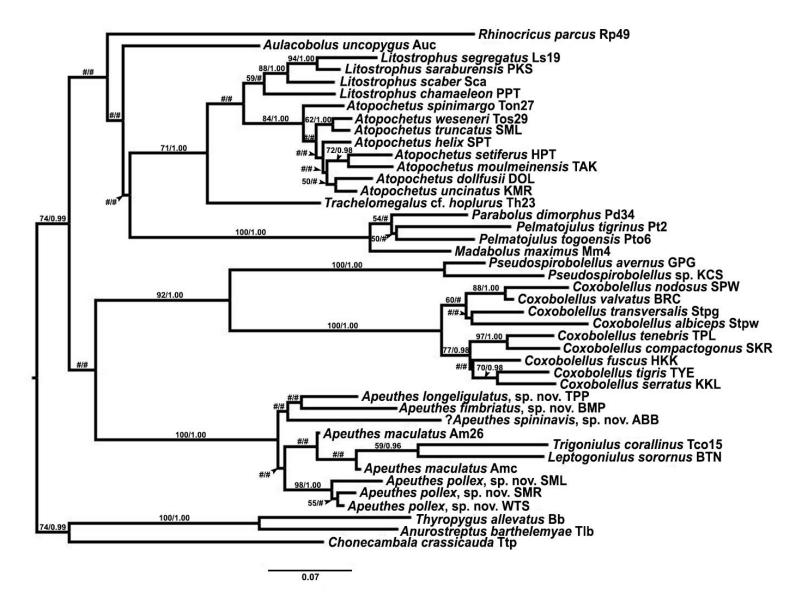


Fig. S2. Phylogenetic relationships of the subfamily Trigoniulinae based on maximum likelihood analysis (ML) and Bayesian Inference (BI) of *16S* ribosomal RNA (458 bp). Numbers at nodes indicate branch support based on bootstrapping (ML) and posterior probabilities (BI). Scale bar: 0.07 substitutions per site. The number sign (#) indicates branches with <50% ML bootstrap support and <0.95 Bayesian posterior probability.