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Corrigendum

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The first mitogenome of the genus Amphalius (Siphonaptera: Ceratophyllidae) and its phylogenetic implications – CORRIGENDUM

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The author regrets the inclusion of the below errors in the above article. These errors concern the numbering of several figures and supplementary figures.

The correct list is as follows:

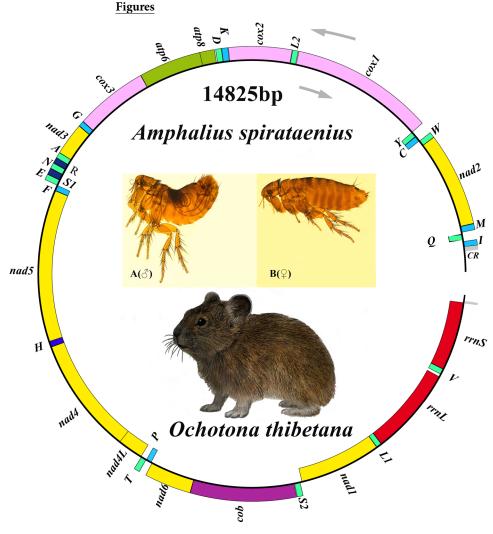


Figure 1. Organization of the Amphalius spirataenius mitogenome. tRNA genes were shown with the single-letter abbreviations of their corresponding amino acids. (note: The morphological figure of Ochotona thibetana from the volume 7 of The Mammals of The World (Wilson et al., 2017))

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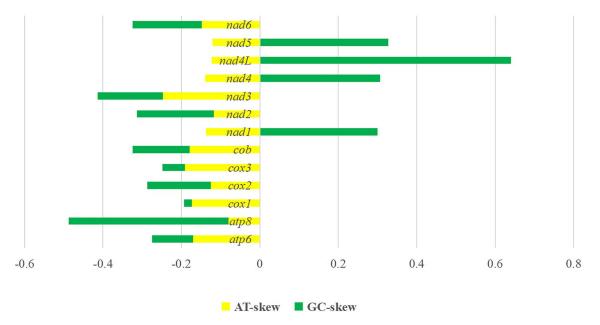


Figure 2. Skewness of 13 protein-coding genes of Amphalius spirataenius

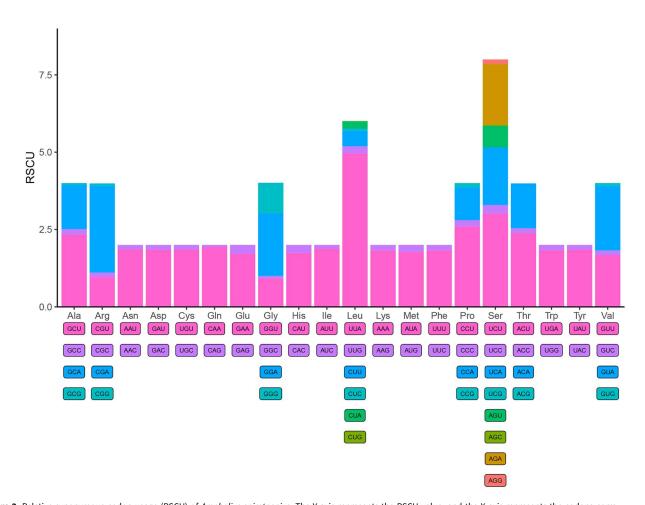


Figure 3. Relative synonymous codon usage (RSCU) of Amphalius spirataenius. The Y-axis represents the RSCU value, and the X-axis represents the codons corresponding to each amino acid

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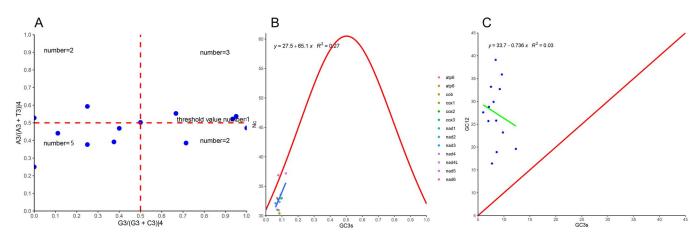


Figure 4. Analysis of 13 protein-coding genes of Amphalius spirataenius. A PR2; B ENC-plot; C Neutral curve

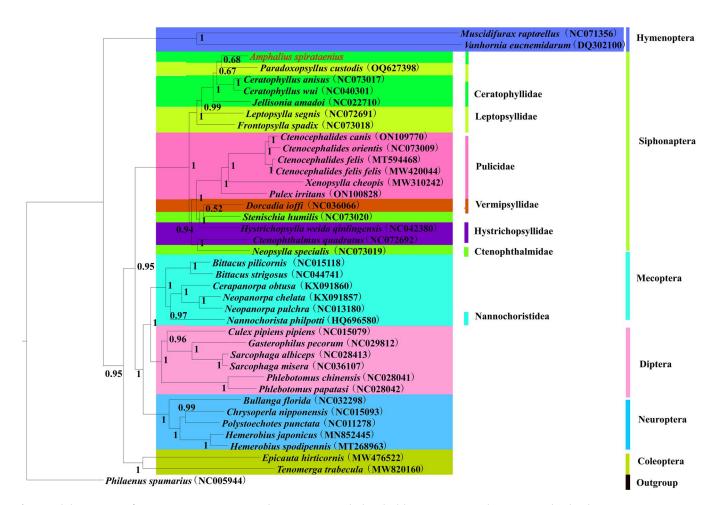


Figure 5. Phylogenetic tree of 40 insect species was constructed using Bayesian methods with *Philaenus spumarius* as the outgroup and node values as posterior probability values (PP). *Amphalius spirataenius* was labelled in red

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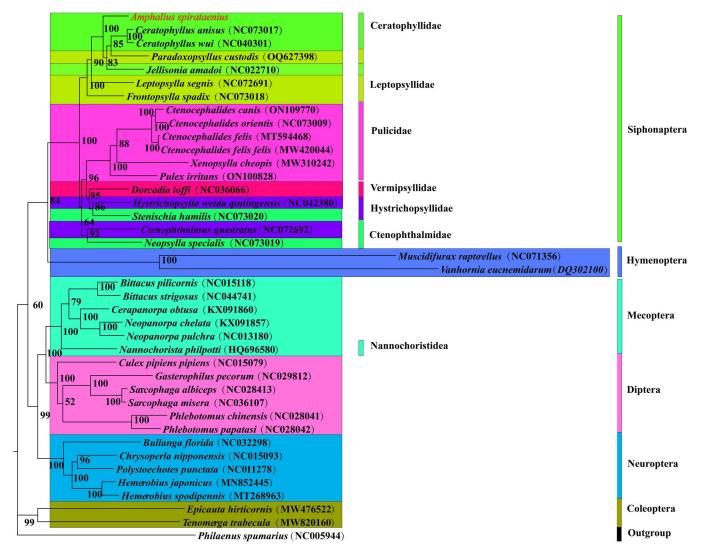


Figure 6. Phylogenetic tree of 40 insect species was constructed by Maximum likelihood method with *Philaenus spumarius* as an outgroup and node values as bootstrap values (BS). *Amphalius spirataenius* was labelled in red

Supplementary Figures

The author apologises for these errors and wishes to correct them through this notice.

Reference

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