

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

Ju Pu , Xiaoxia Lin and Wenge Dong

## Corrigendum

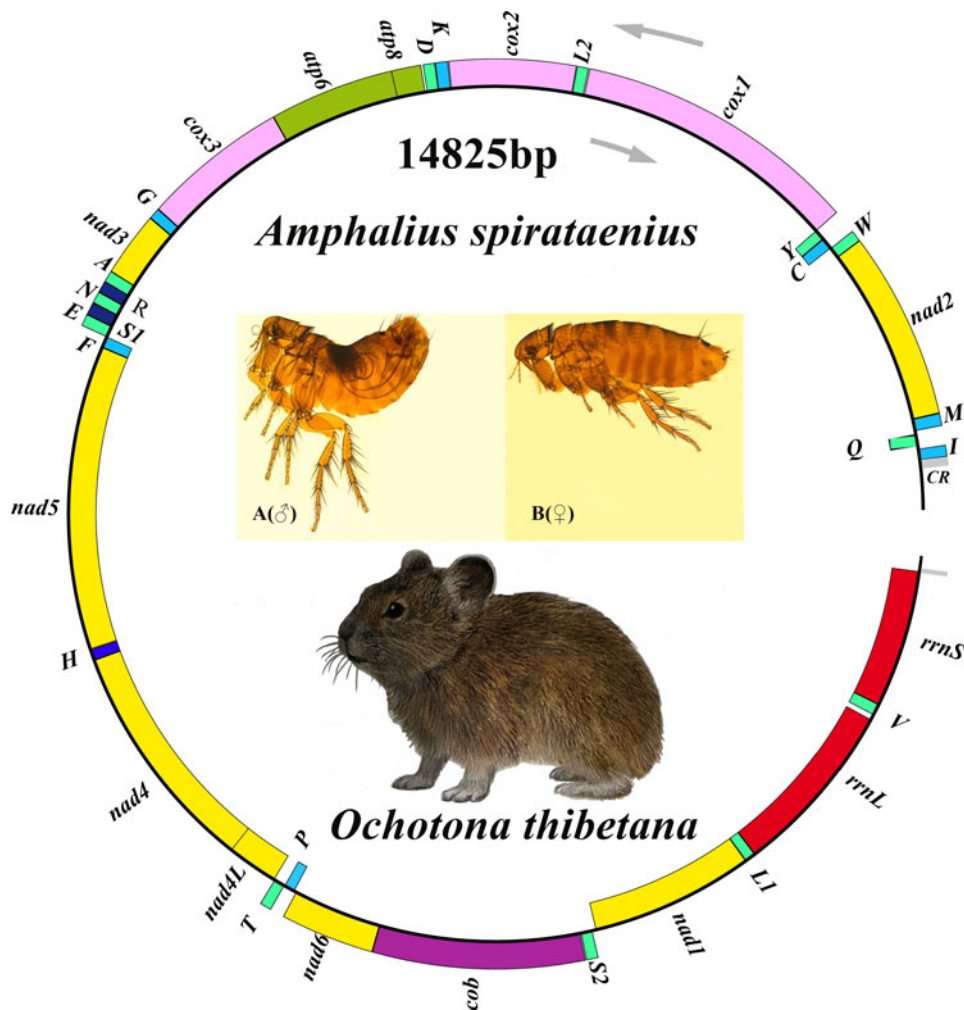
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The author regrets the inclusion of the below errors in the above article. These errors concern the numbering of several figures and supplementary figures.

The correct list is as follows:

### Figures



**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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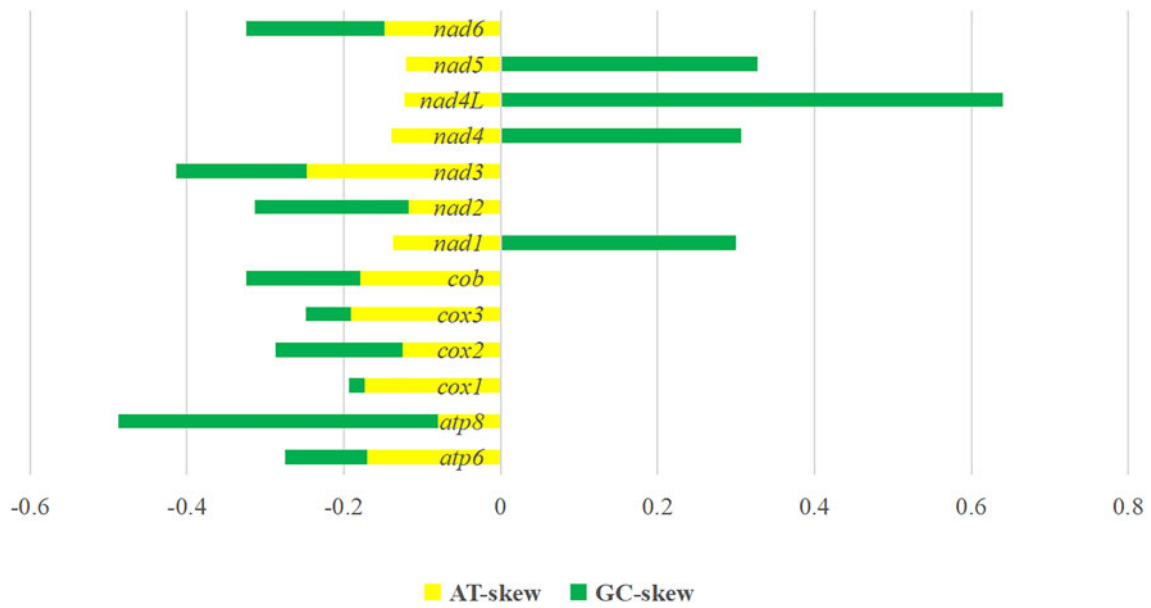


Figure 2. Skewness of 13 protein-coding genes of *Amphilius spirataenius*

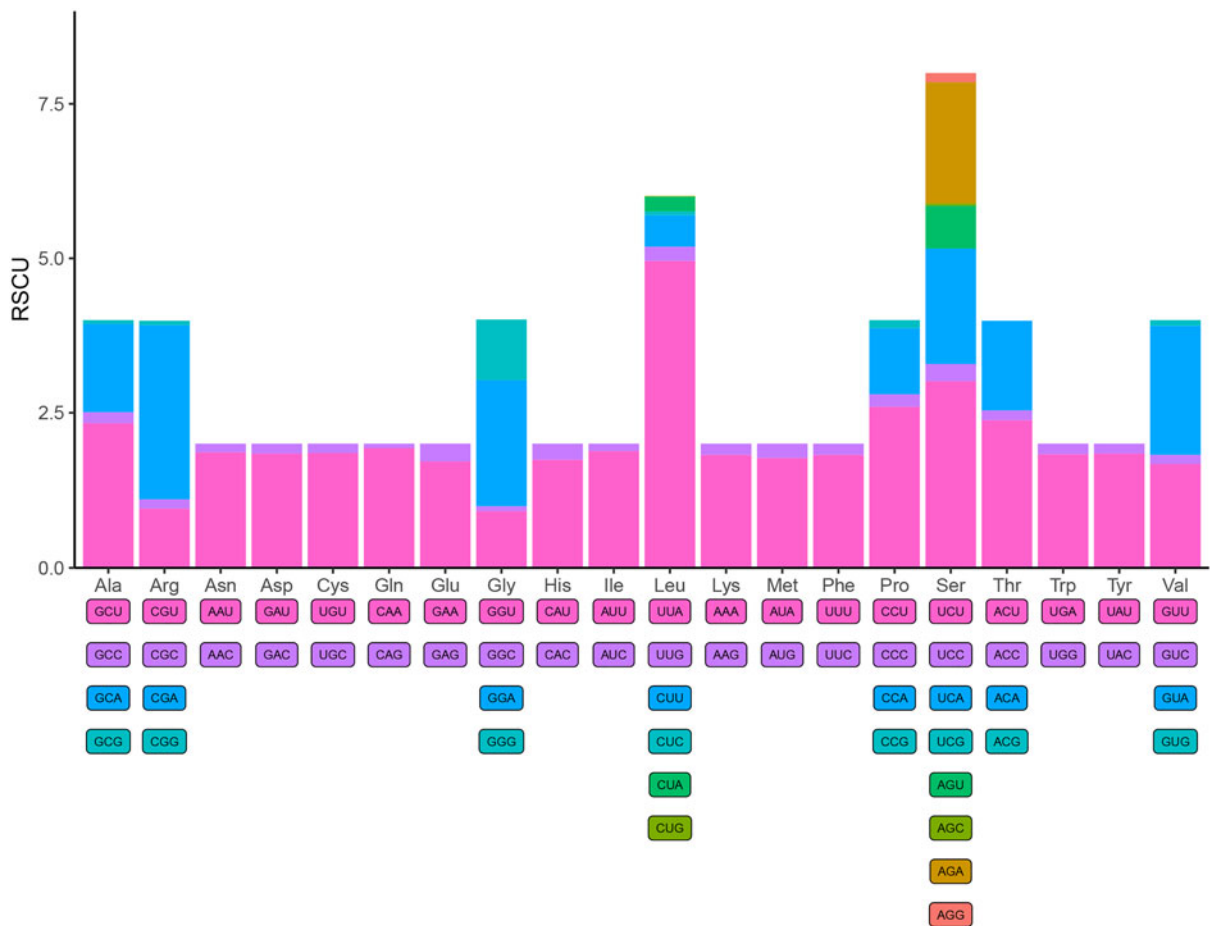


Figure 3. Relative synonymous codon usage (RSCU) of *Amphilius spirataenius*. The Y-axis represents the RSCU value, and the X-axis represents the codons corresponding to each amino acid

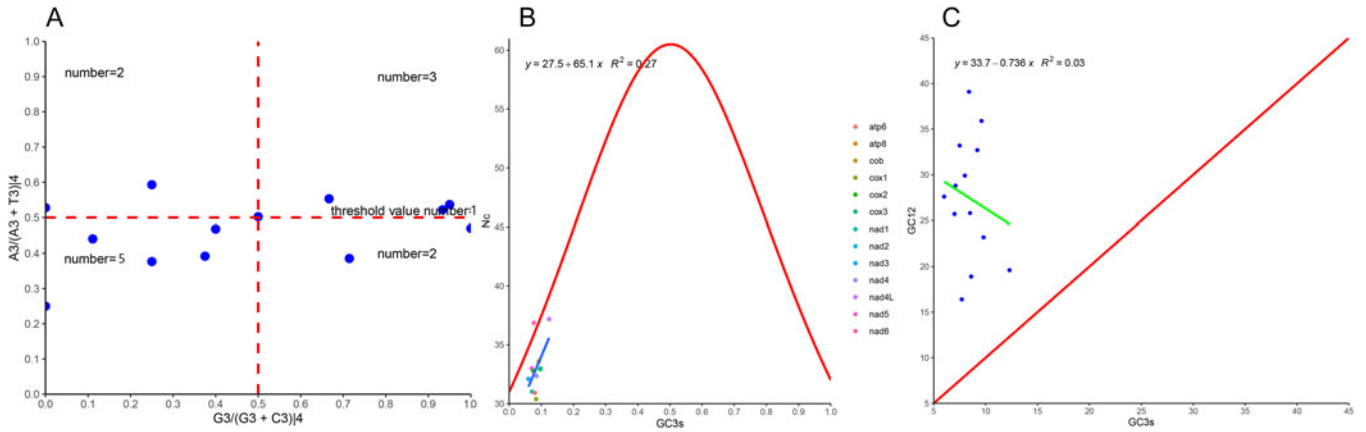


Figure 4. Analysis of 13 protein-coding genes of *Amphialius spirataenius*. A PR2; B ENC-plot; C Neutral curve

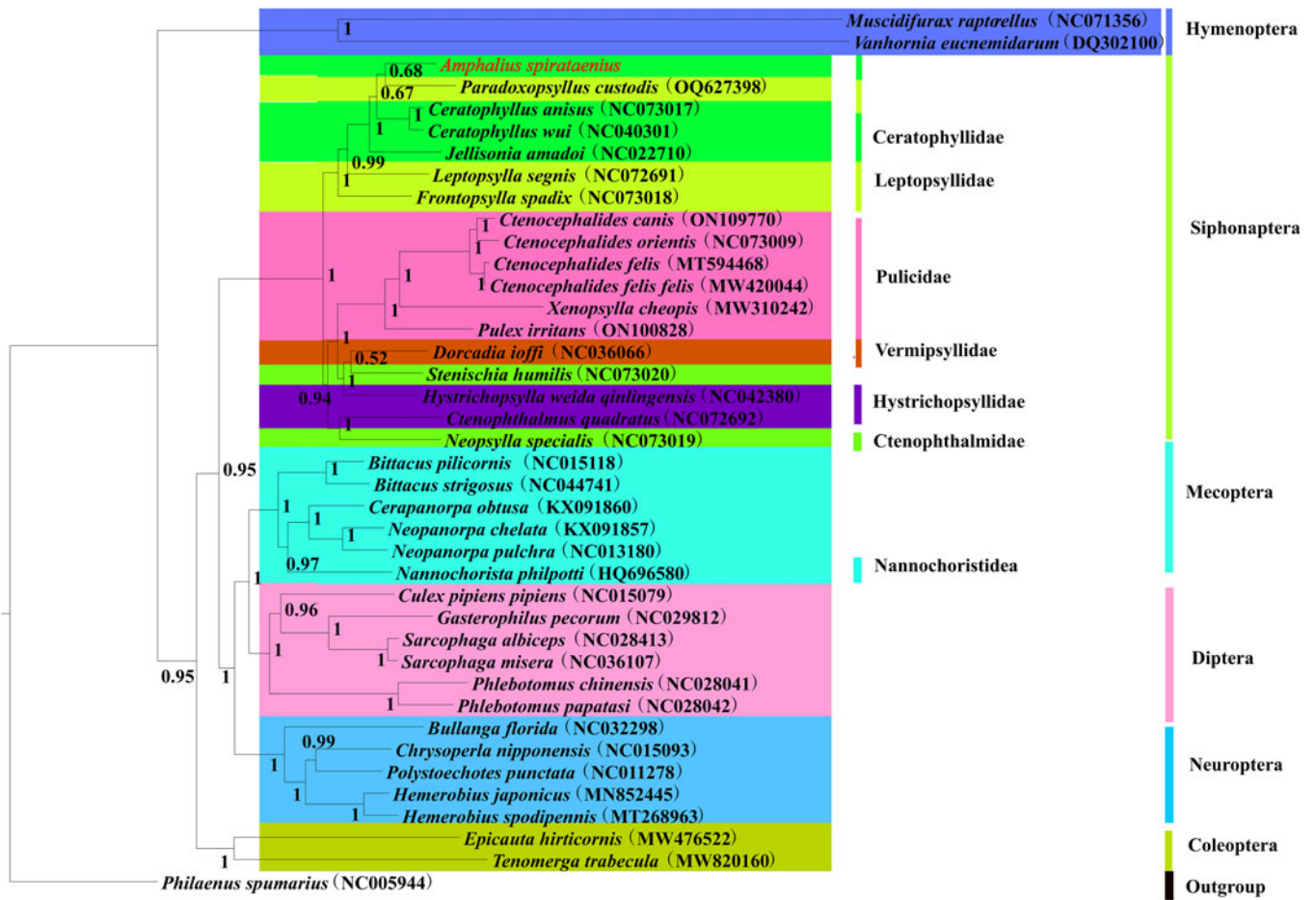
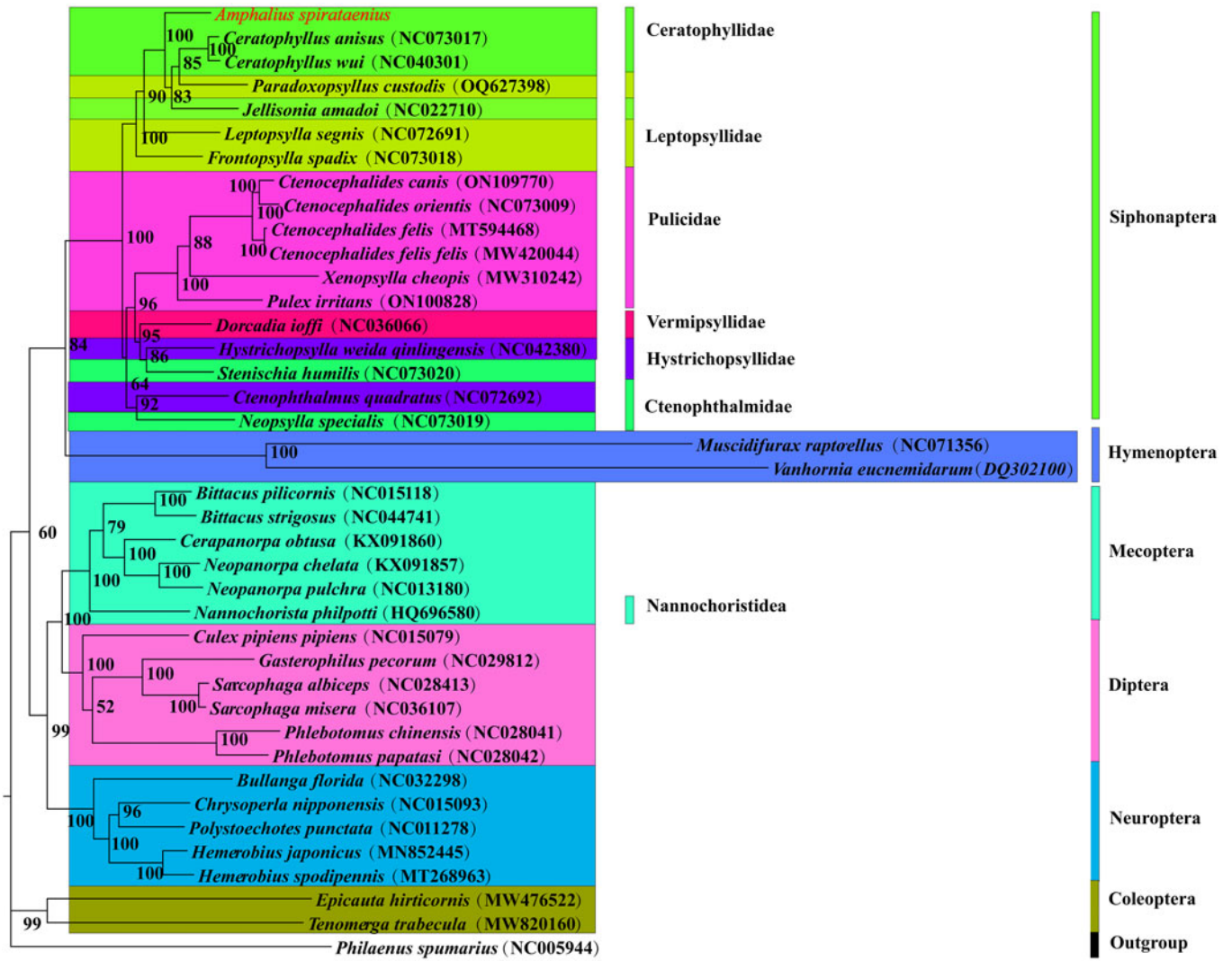


Figure 5. Phylogenetic tree of 40 insect species was constructed using Bayesian methods with *Philaenus spumarius* as the outgroup and node values as posterior probability values (PP). *Amphialius spirataenius* was labelled in red



**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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