

CORRECTION

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Correction: The complete mitochondrial genomes of *Paradiplozoon yarkandense* and *Paradiplozoon homoion* confirm that Diplozoidae evolve at an elevated rate

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Correction to: Parasites Vectors 15:149 (2022)
<https://doi.org/10.1186/s13071-022-05275-x>

Following the original publication of this article [1], the authors flagged the following errors: Fig. 3 had been published upside-down; two newly sequenced species (*Paradiplozoon yarkandense* and *Paradiplozoon*

homoion) had been mistakenly omitted from Figs. 4 and 5; and in the section ‘Gene overlaps’, in the sentence “A relatively large putative overlap between *cytb* and *nad4L* was conserved in both species: *P. yarkandense* = 10 bp, *P. homoion* = 16 bp (Table 1)”, *cox1* had been written instead of *cytb*.

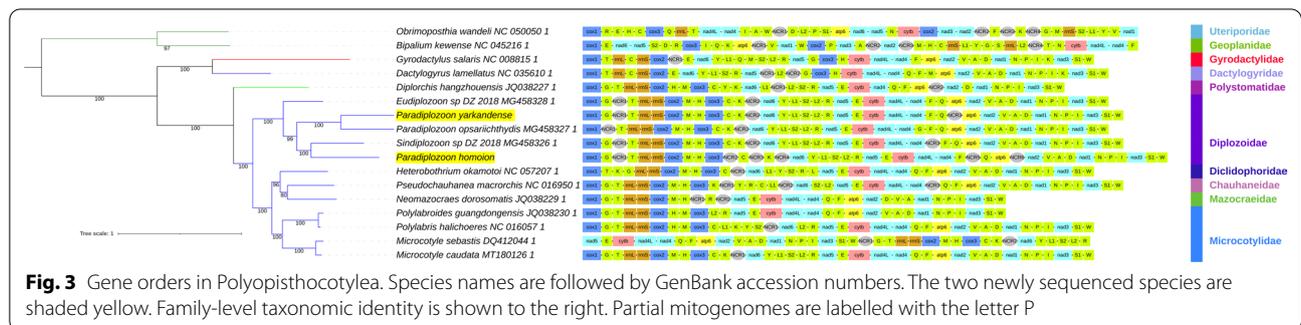


Fig. 3 Gene orders in Polyopisthocytylea. Species names are followed by GenBank accession numbers. The two newly sequenced species are shaded yellow. Family-level taxonomic identity is shown to the right. Partial mitogenomes are labelled with the letter P

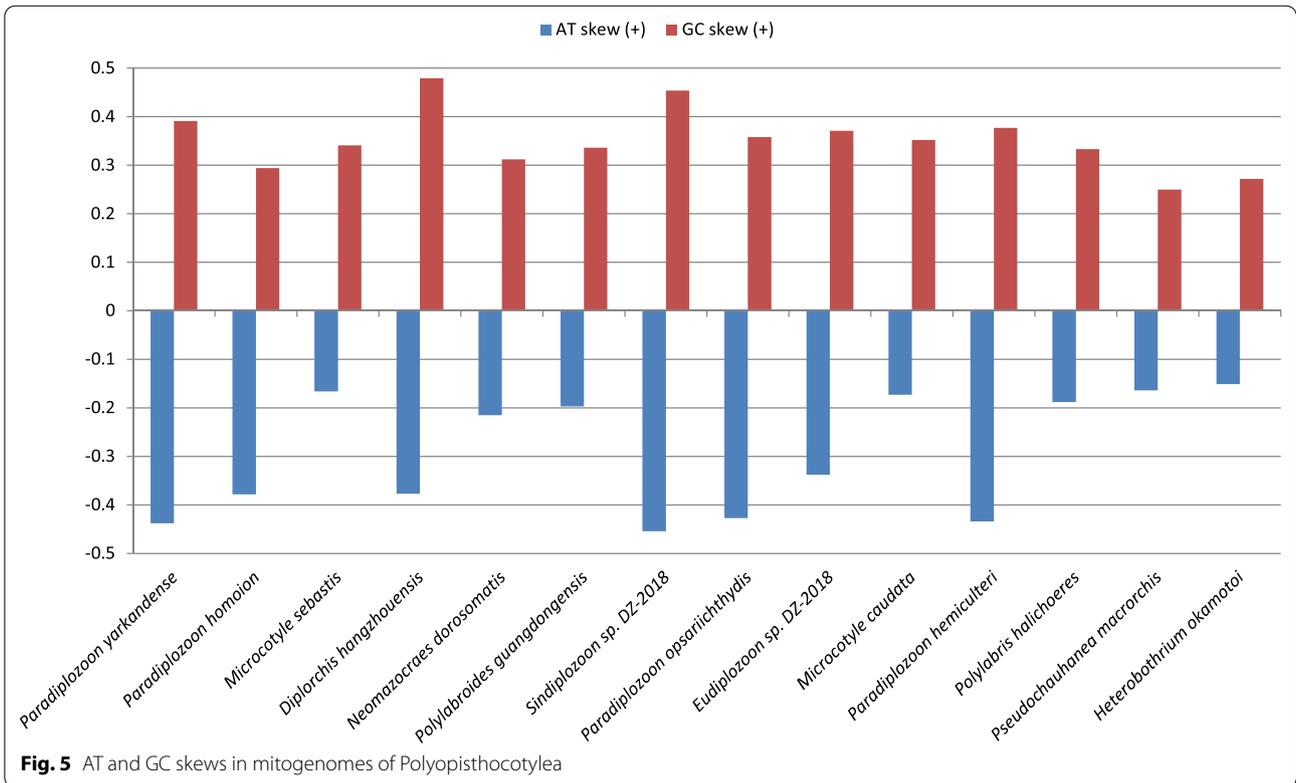
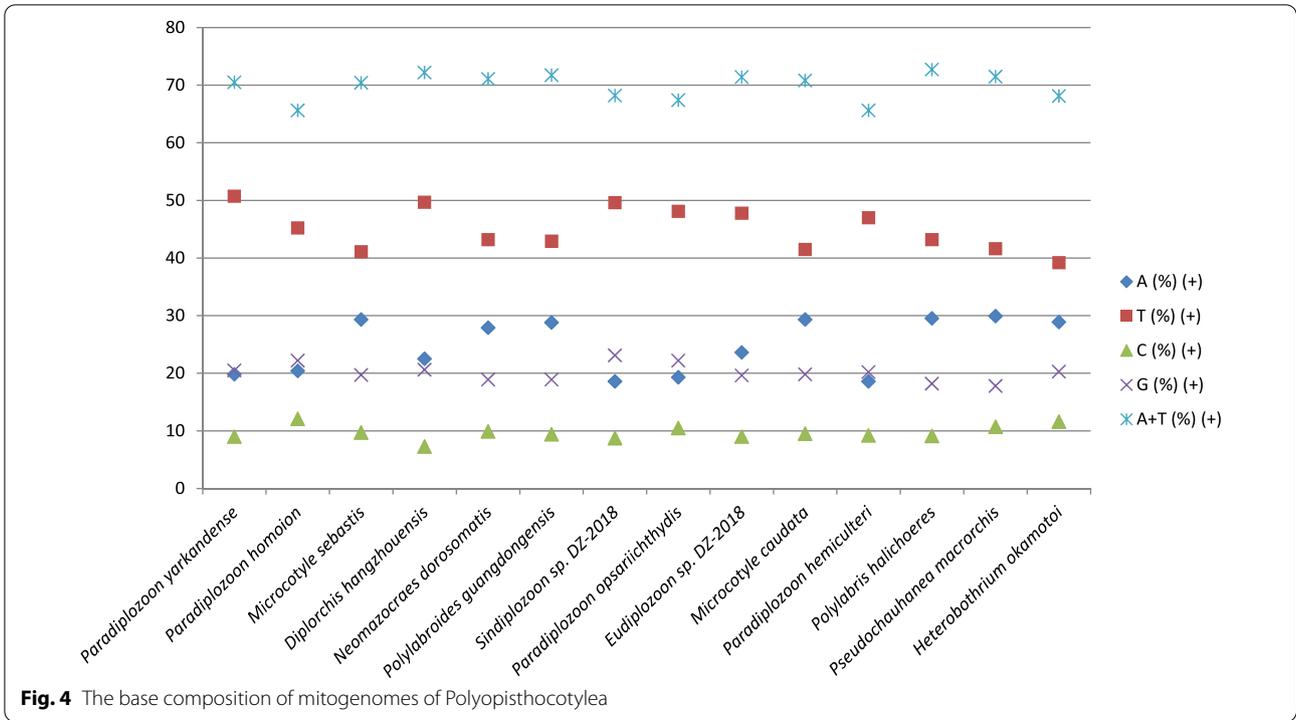
The original article can be found online at <https://doi.org/10.1186/s13071-022-05275-x>.

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The original article has since been corrected and the corrected figures may be found in this erratum for reference.

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Reference

1. Hao C-L, Arken K, Kadir M, Zhang W-R, Rong M-J, Wei N-W, Liu Y-J, Yue C. The complete mitochondrial genomes of *Paradiplozoon yarkandense* and *Paradiplozoon homoion* confirm that Diplozoidae evolve at an elevated rate. *Parasites Vectors*. 2022;15:149. <https://doi.org/10.1186/s13071-022-05275-x>.

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