

## ERRATUM

Reassignment of the land tortoise haemogregarine *Haemogregarina fitzsimonsi* Dias 1953 (Adeleorina: Haemogregarinidae) to the genus *Hepatozoon* Miller 1908 (Adeleorina: Hepatozoidae) based on parasite morphology, life cycle and phylogenetic analysis of 18S rDNA sequence fragments – ERRATUM

COURTNEY A. COOK, SCOTT P. LAWTON, ANGELA J. DAVIES *and* NICO J. SMIT

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Key words: terrestrial chelonian, tick-vector, herpetological haematozoan taxonomy, reptile apicomplexan phylogeny, *Amblyomma*, erratum.

In the above published article by Cook *et al.* Fig. 4 has been reproduced to a low resolution making it difficult to read. A higher quality illustration is published on the next page.

## REFERENCES

- Cook, C. A., Lawton, S. P., Davies, A. J. and Smit, N. J. Reassignment of the land tortoise haemogregarine *Haemogregarina fitzsimonsi* Dias 1953 (Adeleorina: Haemogregarinidae) to the genus *Hepatozoon* Miller 1908 (Adeleorina: Hepatozoidae) based on parasite morphology, life cycle and phylogenetic analysis of 18S rDNA sequence fragments, *Parasitology*, First published online 13 June 2014, 1–10, doi: <http://dx.doi.org/10.1017/S003118201400081X>.
- Dias, J. A. T. S. (1953). Subsídios para o estudo dos hematozoários dos répteis de Moçambique. *Boletim da Sociedade de Estudos Moçambique* **23**, 41–73.

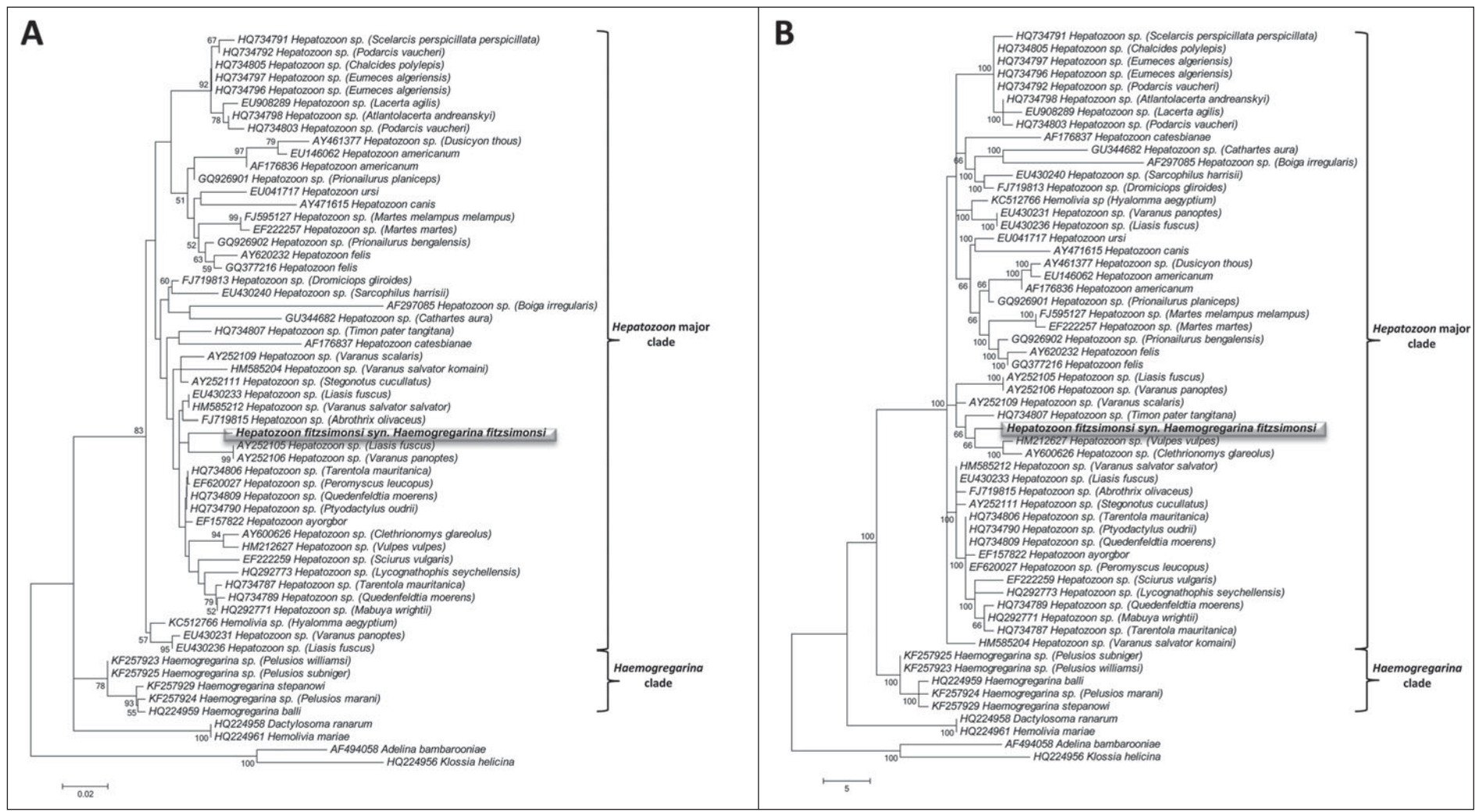


Fig. 4. Phylogenetic analysis, implemented in MEGA5, for *Haemogregarina fitzsimonsi* Dias, 1953 (highlighted) against species of *Hepatozoon*, *Haemogregarina* and *Hemolivia*, with *Adelina bambarooniae* and *Klossia helicina* as an outgroup. Nodal support is provided by bootstrap values with only those >50 shown. (A) Maximum likelihood analysis under the conditions of the Tamura 3-parameter model. The tree with the highest log likelihood (– 1984.703) is shown. (B) Maximum parsimony analysis inferred from 11 648 most parsimonious trees.