



## Correction to: Assessing species number and genetic diversity of the Mountainsnails (Oreohelicidae)

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**Correction to: Conservation Genetics**  
<https://doi.org/10.1007/s10592-020-01302-5>

The original version of this article unfortunately contained several mistakes.

Please note the following corrections in the original article.

Under the heading “DNA sequencing, genotyping, and dataset composition”, in third paragraph, the sentence that reads as “The combined 861 sequence dataset contained representatives of 60.9% (52 species) of all currently recognized species and subspecies in Oreohelicidae” should read as “The combined 861 sequence dataset contained representatives of 60.9% (50 species) of all currently recognized species and subspecies in Oreohelicidae”.

Under the heading “Introduction”, in fourth paragraph, the sentence that reads as “The Mountainsnails (family: Oreohelicidae) are a calciphilous family of montane-endemic land snails that includes two genera: Radiocentrum and Oreohelix—purportedly the most diverse genus of land snails in North America (82 currently recognized taxonomic species and subspecies; Pilsbry 1939; Nekola 2014)” should read as “The Mountainsnails (family: Oreohelicidae) are a calciphilous family of montane-endemic land snails that includes two genera: Radiocentrum and Oreohelix—purportedly the most

diverse genus of land snails in North America (77 currently recognized taxonomic species and subspecies; Pilsbry 1939; Nekola 2014)”.

Under the heading “Results”, in third paragraph, the sentence that reads as “Using the criteria that a delimited species is threatened if all the previous taxonomic units that constitute the new delimited species are listed as NatureServe rank G2 or higher, we found 11 threatened species using these two approaches (Supplemental Table 1)” should read as “Using the criteria that a delimited species is threatened if all the previous taxonomic units that constitute the new delimited species are listed as NatureServe rank G2 or higher, we found 13 threatened species using these two approaches (Supplemental Table 1)”.

In the original publication of the article, the Acknowledgements section was published incorrectly. The correct Acknowledgements section is given in this correction.

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