ERRATUM



Erratum to: Ant and Mite Diversity Drives Toxin Variation in the Little Devil Poison Frog

Jenna R. McGugan¹ · Gary D. Byrd² · Alexandre B. Roland¹ · Stephanie N. Caty¹ · Nisha Kabir³ · Elicio E. Tapia⁴ · Sunia A. Trauger² · Luis A. Coloma^{4,5} · Lauren A. O'Connell¹

Published online: 6 September 2016

© Springer Science+Business Media New York 2016

Erratum to: J Chem Ecol (2016) 42:537–551 DOI: 10.1007/s10886-016-0715-x

There were three minor typos in the original publication, which we have corrected here. The authors regret these errors.

- 1. There was a mistake in the legends of Figure 3 and Online Resource Figure 1. The scale bar is 0.5mm, not 5mm.
- 2. When describing the methods for generating a phylogenetic tree in MEGA 6 using the cytochrome oxidase 1 data for arthropods, we mistakenly wrote the type of clustering method as "Nearest Neighbor Joining", but the correct method was "Neighbor Joining".
- 3. There was a mistake in Online Resource Table 2. The accession number for one of the closest BLAST hits should be KF293453, not KM293453.

The online version of the original article can be found at http://dx.doi.org/10.1007/s10886-016-0715-x

- ☐ Lauren A. O'Connell aloconnell@fas.harvard.edu
- Center for Systems Biology, Harvard University, Cambridge, MA 02138, USA
- ² Small Molecule Mass Spectrometry Facility, Harvard University, Cambridge, MA 02138, USA
- Cambridge Rindge and Latin High School, Cambridge, MA 02138, USA
- ⁴ Centro Jambatu de Investigación y Conservación de Anfibios, Fundación Otonga, Quito, Ecuador
- ⁵ Ikiam, Universidad Regional Amazónica, Muyuna, Tena, Ecuador

