Correction

Correction: The Role of Fusion in Ant Chromosome Evolution: Insights from Cytogenetic Analysis Using a Molecular Phylogenetic Approach in the Genus *Mycetophylax*

The PLOS ONE Staff

There is an error in Table 1. During the production process, part of the table caption got merged with the information under the table. Please see the corrected Table 1 here.

Table 1. Cytogenetic data of the *Mycetophylax* species. Summary of chromosome number, chromosome morphology and banding patterns of observed karyotypes of all the specimens analyzed here.

	Chromosome number (n)	Chromosome morphology			C-banding positive blocks			Fluorochrome staining	
		M	SM	A	C	PC	SA	AT+ bands	GC+ bands
<i>M. simplex</i>	18	10	8	no	yes	yes	yes	yes	yes
<i>M. conformis</i>	15	11	4	no	yes	yes	yes	yes	yes
<i>M. morschi</i>	15	9	5	1	yes	no	no	no	no
<i>M. morschi</i>	13	9	3	1	yes	no	no	no	no

M: metacentric; SM: submetacentric; A: acrocentric.
C: centromeric; PC: pericentromeric; SA: Short arm.
doi:10.1371/journal.pone.0087473.t001

Reference

- Cardoso DC, das Graças Pompolo S, Cristiano MP, Tavares MG (2014) The Role of Fusion in Ant Chromosome Evolution: Insights from Cytogenetic Analysis Using a Molecular Phylogenetic Approach in the Genus *Mycetophylax*. PLoS ONE 9(1): e87473. doi:10.1371/journal.pone.0087473

Citation: The PLOS ONE Staff (2014) Correction: The Role of Fusion in Ant Chromosome Evolution: Insights from Cytogenetic Analysis Using a Molecular Phylogenetic Approach in the Genus *Mycetophylax*. PLoS ONE 9(4): e95408. doi:10.1371/journal.pone.0095408

Published: April 10, 2014

Copyright: © 2014 The PLOS ONE Staff. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.