OPEN **Publisher Correction:** Multiple evolutionary lineages for the main vector of *Leishmania guyanensis*, *Lutzomyia umbratilis* (Diptera: Psychodidae), in the Brazilian Amazon

Vera Margarete Scarpassa, Antônio Saulo Cunha-Machado & Ronildo Baiatone Alencar

Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-021-93072-4>, published online 28 July 2021

The original version of this Article contained errors.

In the Abstract,

“The third lineage, first recorded in this study, is in the interfluvium between south of Amazonas River and west of Madeira River, and its involvement in the transmission of this parasite remains to be elucidated.”

now reads:

“The third lineage, first recorded in this study, is in the interfluvium between south of Amazon River and west of Madeira River, and its involvement in the transmission of this parasite remains to be elucidated.”

In addition, a previous rendition of Figure 5 was published. As a result, the colored boxes were incorrectly shown.

The original Figure 5 and accompanying legend appear below.

The original Article has been corrected.

Published online: 19 August 2021

Figure 5. Bayesian Inference (BI) tree (maximum clade of credibility) based on the 89 haplotypes of *COI* and generated in BEAST. Tree inferred under the Time Reversible (GTR) + G + I nucleotide substitution model. The support values, in BPP (Bayesian Posterior Probability), are indicated above of the branches. See Table S1 for identification of the haplotypes. Saint: Saint Georges l' Oyapock, French Guiana. *Lutzomyia anduzei* and *Bichromomyia flaviscutellata* were used as outgroups. The species-genetic lineages delimited by the GMYC, bGMYC and ABGD models are represented by colored boxes. The 9, 8 and 6 indicate the lineage numbers recognized by the GMYC, bGMYC and ABGD models, respectively. Each color represents one lineage. The colors of the lineages delimited by the three models followed the same pattern of the Fig. 2.

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2021