

Published online: 16 November 2018

OPEN Author Correction: Comparing Sanger sequencing and highthroughput metabarcoding for inferring photobiont diversity in lichens

Fiona Paul 1, Jürgen Otte¹, Imke Schmitt^{1,2} & Francesco Dal Grande^{1,3}

Correction to: Scientific Reports https://doi.org/10.1038/s41598-018-26947-8, published online 05 June 2018

In the "Supplementary Table S2" file originally published with this Article, columns E and G contained incorrect accession numbers. These errors have been corrected in the "Supplementary Table S2" file that now accompanies the Article.

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 license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license 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 license, visit http://creativecommons.org/licenses/by/4.0/.

© The Author(s) 2018

¹Senckenberg Biodiversity and Climate Research Centre (SBiK-F), Senckenberganlage 25, 60325, Frankfurt am Main, Germany. 2Institute of Ecology, Evolution and Diversity, Goethe University Frankfurt am Main, Max-von-Laue-Str. 9, 60438, Frankfurt am Main, Germany. ³Departamento de Farmacología, Farmacognosia y Botánica, Universidad Complutense de Madrid, 28040, Madrid, Spain. Correspondence and requests for materials should be addressed to I.S. (email: imke.schmitt@senckenberg.de) or F.D.G. (email: francesco.dalgrande@senckenberg.de)