



### OPEN ACCESS

#### Edited by:

Yasir Muhammad, King Abdulaziz University, Saudi Arabia

#### Reviewed by:

J. Christopher Fenno, University of Michigan, United States

#### \*Correspondence:

Sascha Knauf sknauf@dpz.eu

#### <sup>†</sup>Present address:

Sascha Knauf, Division of Microbiology and Animal Hygiene, University of Göttingen, Göttingen, Germany

#### Specialty section:

This article was submitted to Evolutionary and Genomic Microbiology, a section of the journal Frontiers in Microbiology

Received: 18 September 2019 Accepted: 24 October 2019 Published: 08 November 2019

#### Citation:

Hallmaier-Wacker LK, Lüert S, Gronow S, Spröer C, Overmann J, Buller N, Vaughan-Higgins RJ and Knauf S (2019) Corrigendum: A Metataxonomic Tool to Investigate the Diversity of Treponema. Front. Microbiol. 10:2581. doi: 10.3389/fmicb.2019.02581

# **Corrigendum: A Metataxonomic Tool** to Investigate the Diversity of *Treponema*

## Luisa K. Hallmaier-Wacker<sup>1,2</sup>, Simone Lüert<sup>1,2</sup>, Sabine Gronow<sup>3</sup>, Cathrin Spröer<sup>3</sup>, Jörg Overmann<sup>3,4</sup>, Nicky Buller<sup>5</sup>, Rebecca J. Vaughan-Higgins<sup>6</sup> and Sascha Knauf<sup>1\*†</sup>

<sup>1</sup> Neglected Tropical Diseases Work Group, Infection Biology Unit, German Primate Center, Leibniz Institute for Primate Research, Göttingen, Germany, <sup>2</sup> Primate Genetics Laboratory, German Primate Center, Leibniz Institute for Primate Research, Göttingen, Germany, <sup>3</sup> Leibniz Institute DSMZ – German Collection of Microorganisms and Cell Cultures, Braunschweig, Germany, <sup>4</sup> Department of Microbiology, Braunschweig University of Technology, Braunschweig, Germany, <sup>5</sup> Animal Pathology – Bacteriology Laboratory, Department of Primary Industries and Regional Development, South Perth, WA, Australia, <sup>6</sup> Department of Conservation Medicine, College of Veterinary Medicine, School of Veterinary and Life Sciences, Murdoch University, Murdoch, WA, Australia

#### Keywords: metagenomics, metataxonomics, one health, spirochete, 16S rRNA, Treponema, marsupial, Potorous

#### A Corrigendum on

#### A Metataxonomic Tool to Investigate the Diversity of Treponema

by Hallmaier-Wacker, L. K., Lüert, S., Gronow, S., Spröer, C., Overmann, J., Buller, N., et al. (2019). Front. Microbiol. 10:2094. doi: 10.3389/fmicb.2019.02094

In the original article, there was a mistake in **Figure 1** as published. The figure contains an error in the number of cycles for the "add indices and adapters" step. The step should read 8 cycles instead of 20 cycles. The corrected **Figure 1** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Copyright © 2019 Hallmaier-Wacker, Lüert, Gronow, Spröer, Overmann, Buller, Vaughan-Higgins and Knauf. This is an openaccess article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

