



# Corrigendum: Forest Soil Fungal Community Elevational Distribution Pattern and Their Ecological Assembly Processes

Yuyu Sheng<sup>1</sup>, Wei Cong<sup>1</sup>, Linsen Yang<sup>2</sup>, Qiang Liu<sup>2</sup> and Yuguang Zhang<sup>1\*</sup>

<sup>1</sup> Research Institute of Forest Ecology, Environment and Protection, Chinese Academy of Forestry, and the Key Laboratory of Biological Conservation of National Forestry and Grassland Administration, Beijing, China, <sup>2</sup> Shennongjia National Park Administration, and Hubei Provincial Key Laboratory on Conservation Biology of the Shennongjia Golden Monkey, Shennongjia, China

## OPEN ACCESS

**Edited and reviewed by:**  
Manuel Delgado Baquerizo,  
University of Alicante, Spain

**\*Correspondence:**  
Yuguang Zhang  
yugzhang@sina.com.cn

**Specialty section:**  
This article was submitted to  
Terrestrial Microbiology,  
a section of the journal  
Frontiers in Microbiology

**Received:** 24 October 2019  
**Accepted:** 18 November 2019  
**Published:** 05 December 2019

**Citation:**  
Sheng Y, Cong W, Yang L, Liu Q and  
Zhang Y (2019) Corrigendum: Forest  
Soil Fungal Community Elevational  
Distribution Pattern and Their  
Ecological Assembly Processes.  
*Front. Microbiol.* 10:2802.  
doi: 10.3389/fmicb.2019.02802

**Keywords:** soil fungal diversity, community assembly processes, elevational gradient, environmental factors, Illumina MiSeq sequencing

## A Corrigendum on

### Forest Soil Fungal Community Elevational Distribution Pattern and Their Ecological Assembly Processes

by Sheng, Y., Cong, W., Yang, L., Liu, Q., and Zhang, Y. (2019). *Front. Microbiol.* 10:2226. doi: 10.3389/fmicb.2019.02226

In the original article, there was a mistake in the legend for **Figure 2** as published. The sentence “Relationships between (a)  $\alpha$ -diversity of plant and soil fungi and (b) soil fungal  $\alpha$ -diversity and elevational gradient” in the legend should be deleted. The correct legend appears below.

**Figure 2 |** Function predictions of soil fungal communities using FUNGuild method. The y axis represents the relative OTUs abundance and different letters above the columns mean significant difference among samples ( $P < 0.05$ ).

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 Sheng, Cong, Yang, Liu and Zhang. This is an open-access 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.