

CORRECTION

[View Article Online](#)
[View Journal](#) | [View Issue](#)

Cite this: *Food Funct.*, 2023, **14**, 5820

Correction: Deep Shotgun metagenomic and 16S rRNA analysis revealed the microbial diversity of lactic acid bacteria in traditional fermented foods of eastern Hainan, China

Yuqing Wu,^a Ao Li,^a Lin Chen,^a Qianxi Chen,^a Jiyang Li,^a Yanqing Xu^{*a} and Dongxue Huo^{*a,b}

DOI: 10.1039/d3fo90047a
rsc.li/food-function

Correction for 'Deep Shotgun metagenomic and 16S rRNA analysis revealed the microbial diversity of lactic acid bacteria in traditional fermented foods of eastern Hainan, China' by Yuqing Wu *et al.*, *Food Funct.*, 2022, **13**, 12938–12952, <https://doi.org/10.1039/D2FO02501A>.

The authors regret that the name of one of the authors (Lin Chen) was shown incorrectly in the original article. The corrected author list is as shown above.

The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

^aSchool of Food Science and Engineering, School of Public Administration, Hainan University, Haikou, China. E-mail: microbiome@vip.163.com, 17784625873@126.com

^bOne Health Institute, Hainan University, Haikou, China

