
Correction

Correction to: MetaSquare: an integrated metadatabase of 16S rRNA gene amplicon for microbiome taxonomic classification

Bioinformatics (2022) <https://doi.org/10.1093/bioinformatics/btac184>

In the originally published version of this manuscript, the funding information was incorrect but it has been updated as:

This project was funded by the grants MOST 108-2321-B-037 -001 and 110-2314-B-001 -006 from the Ministry of Science and Technology, Taiwan, and AS-GCS-109-07 from Academia Sinica, Taiwan, to financially support this research and publication.

This error has been corrected (online).

Reference

Chun-Chieh, L. *et al.* (2022) MetaSquare: an integrated metadatabase of 16S rRNA gene amplicon for microbiome taxonomic classification, *Bioinformatics*. <https://doi.org/10.1093/bioinformatics/btac184>.