



OPEN ACCESS

APPROVED BY
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

*CORRESPONDENCE
Peng Zhu,
✉ zupeng@cqmu.edu.cn
Jun Liu,
✉ 2022110338@stu.cqmu.edu.cn

†These authors have contributed equally
to this work

RECEIVED 09 March 2026
REVISED 09 March 2026
ACCEPTED 20 March 2026
PUBLISHED 26 March 2026

CITATION
Zhang Y, Qiu C, Li X, Peng Y, Liu J and
Zhu P (2026) Correction: A systematic
exploration of gut microbiota-driven
blood metabolites in sepsis: an integrated
bioinformatics and genetic
association study.
Front. Genet. 17:1826365.
doi: 10.3389/fgene.2026.1826365

COPYRIGHT
© 2026 Zhang, Qiu, Li, Peng, Liu and Zhu.
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.

Correction: A systematic exploration of gut microbiota-driven blood metabolites in sepsis: an integrated bioinformatics and genetic association study

Yanhua Zhang^{1†}, Chenghao Qiu^{1†}, Xingyu Li^{2†}, Yaling Peng³,
Jun Liu^{1*} and Peng Zhu^{1*}

¹Department of Gastrointestinal Surgery, The Second Affiliated Hospital of Chongqing Medical University, Chongqing, China, ²The Second Clinical Medical College of Chongqing Medical University, Chongqing, China, ³Department of Anesthesiology, Western Medicine Department, The Central Hospital of Enshi Tujia and Miao Autonomous Prefecture, Enshi, Hubei, China

KEYWORDS

genome-wide association study, gut microbiota, metabolites, network pharmacology, sepsis

A Correction on

A systematic exploration of gut microbiota-driven blood metabolites in sepsis: an integrated bioinformatics and genetic association study

by Zhang Y, Qiu C, Li X, Peng Y, Liu J and Zhu P (2026). *Front. Genet.* 17:1754817. doi: 10.3389/fgene.2026.1754817

Incorrect funding.

The funder Chongqing Natural Science Foundation, CSTB2024NSCQ-KJFZMSX0101 to Peng Zhu was erroneously omitted.

The original article has been updated.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.