



OPEN ACCESS

APPROVED BY
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

*CORRESPONDENCE
Ming Lei
✉ leiming@gxyzyzw.com

[†]These authors have contributed equally to this work

RECEIVED 16 January 2026
REVISED 13 January 2026
ACCEPTED 16 January 2026
PUBLISHED 03 February 2026

CITATION
Wan L, Huang Q, Li C, Yu H, Tan G, Wei S, El-Sappah AH, Sooranna S, Zhang K, Pan L, Zhang Z and Lei M (2026) Correction: Integrated metabolome and transcriptome analysis identifies candidate genes involved in triterpenoid saponin biosynthesis in leaves of *Centella asiatica* (L.) Urban. *Front. Plant Sci.* 17:1784161. doi: 10.3389/fpls.2026.1784161

COPYRIGHT
© 2026 Wan, Huang, Li, Yu, Tan, Wei, El-Sappah, Sooranna, Zhang, Pan, Zhang and Lei. 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: Integrated metabolome and transcriptome analysis identifies candidate genes involved in triterpenoid saponin biosynthesis in leaves of *Centella asiatica* (L.) Urban

Lingyun Wan^{1,2,3†}, Qiulan Huang^{4†}, Cui Li^{2,3}, Haixia Yu^{1,2,3}, Guiyu Tan^{1,2,3}, Shugen Wei^{1,2,3}, Ahmed H. El-Sappah^{4,5}, Suren Sooranna^{3,6}, Kun Zhang^{1,2,3}, Limei Pan^{1,2,3}, Zhanjiang Zhang^{1,2,3} and Ming Lei^{2,3*}

¹Guangxi Key Laboratory for High-Quality Formation and Utilization of Dao-Di Herbs, Guangxi Botanical Garden of Medicinal Plants, Nanning, China, ²National Center for Traditional Chinese Medicine (TCM) Inheritance and Innovation, Guangxi Botanical Garden of Medicinal Plants, Nanning, China, ³National Engineering Research Center for the Development of Southwestern Endangered Medicinal Materials, Guangxi Botanical Garden of Medicinal Plants, Nanning, China, ⁴Faculty of Agriculture, Forestry and Food Engineering, Yibin University, Yibin, China, ⁵Genetics Department, Faculty of Agriculture, Zagazig University, Zagazig, Egypt, ⁶Department of Metabolism, Digestion and Reproduction, Imperial College London, London, United Kingdom

KEYWORDS

Centella asiatica, transcriptome, metabolome, triterpenoid saponin, candidate gene

A Correction on

Integrated metabolome and transcriptome analysis identifies candidate genes involved in triterpenoid saponin biosynthesis in leaves of *Centella asiatica* (L.) Urban

By Wan L, Huang Q, Li C, Yu H, Tan G, Wei S, El-Sappah AH, Sooranna S, Zhang K, Pan L, Zhang Z and Lei M (2024) *Front. Plant Sci.* 14:1295186. doi: 10.3389/fpls.2023.1295186

The funder “the Guangxi Science and Technology Base and Talent Project”, “Guike AD22080016” to “Ming Lei” was erroneously omitted.

The correct **Funding** statement is “The author(s) declare financial support was received for the research, authorship, and/or publication of this article. This research was funded by the Guangxi Science and Technology Base and Talent Project (Guike AD22080016), the Natural Science Foundation of Guangxi Province (2023GXNSFAA026330 and 2020GXNSFAA159151), Guangxi Science and Technology Base and Special Talents (Guike AD22035026), Scientific Research Funding Project of Guangxi Botanical Garden of Medicinal Plants (GYJ202008), Innovative Team for Traditional Chinese Medicinal Materials Quality of Guangxi (GZKJ2305) and the Key Laboratory Construction Program of Guangxi Health commission (ZJC2020003).”

The original version of this 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.