

OPEN ACCESS

APPROVED BY

Frontiers Editorial Office, Frontiers Media SA, Switzerland

*CORRESPONDENCE
Hua Chen,

☑ 936483206@qq.com
Di Liao,
☑ liaodi1118@csu.edu.cn

RECEIVED 23 March 2024 ACCEPTED 25 March 2024 PUBLISHED 10 April 2024

CITATION

Yang Z, He M, Zhang Q, Li S, Chen H and Liao D (2024), Corrigendum: Exploring the bidirectional relationship and shared genes between depression and stroke via NHANES and bioinformatic analysis. Front. Genet. 15:1405696. doi: 10.3389/fgene.2024.1405696

COPYRIGHT

© 2024 Yang, He, Zhang, Li, Chen and Liao. 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.

Corrigendum: Exploring the bi-directional relationship and shared genes between depression and stroke via NHANES and bioinformatic analysis

Zhanghuan Yang¹, Maokun He², Qian Zhang^{3,4}, Shifu Li^{3,4}, Hua Chen⁵* and Di Liao⁶*

¹Department of Oncology, Xiangya Cancer Center, Xiangya Hospital, Central South University, Changsha, Hunan, China, ²Hainan Medical University, Haikou, Hainan, China, ³Department of Neurosurgery, Xiangya Hospital, Central South University, Changsha, Hunan, China, ⁴National Clinical Research Center for Geriatric Disorders, Central South University, Changsha, Hunan, China, ⁵Department of Neurosurgery, The First people's Hospital of Changde, Changde, China, ⁶Department of Neurology, Xiangya Hospital, Central South University, Changsha, Hunan, China

KEYWORDS

ischemic stroke, major depressive disorder, bioinformatics, shared genes, immune infiltration, NHANES

A Corrigendum on

Exploring the bi-directional relationship and shared genes between depression and stroke via NHANES and bioinformatic analysis

by Yang Z, He M, Zhang Q, Li S, Chen H and Liao D (2023). Front. Genet. 14:1004457. doi: 10.3389/fgene.2023.1004457

In the published article, there was an error in the **Funding** statement. The correct **Funding** statement appears below.

Funding

This work was supported by the National Natural Science Foundation of China (82101947), the China Postdoctoral Science Foundation (2021TQ0371 and 2021M703636), and the Fundamental Research Funds for the Central Universities of Central South University (2020zzts269).

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.