



**QUEEN'S
UNIVERSITY
BELFAST**

Author Correction: Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways

Howard, D. M., Adams, M. J., Shirali, M., Clarke, T. K., Marioni, R. E., Davies, G., Coleman, J. R. I., Alloza, C., Shen, X., Barbu, M. C., Wigmore, E. M., Gibson, J., Agee, M., Alipanahi, B., Auton, A., Bell, R. K., Bryc, K., Elson, S. L., Fontanillas, P., ... McIntosh, A. M. (2021). Author Correction: Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways. *Nature Communications*, 12, Article 2012. <https://doi.org/10.1038/s41467-021-22411-w>

Published in:

Nature Communications

Document Version:

Publisher's PDF, also known as Version of record

Queen's University Belfast - Research Portal:

[Link to publication record in Queen's University Belfast Research Portal](#)

Publisher rights

Copyright 2022 the authors.

This is an open access article published under a Creative Commons Attribution License (<https://creativecommons.org/licenses/by/4.0/>), which permits unrestricted use, distribution and reproduction in any medium, provided the author and source are cited.

General rights

Copyright for the publications made accessible via the Queen's University Belfast Research Portal is retained by the author(s) and / or other copyright owners and it is a condition of accessing these publications that users recognise and abide by the legal requirements associated with these rights.

Take down policy

The Research Portal is Queen's institutional repository that provides access to Queen's research output. Every effort has been made to ensure that content in the Research Portal does not infringe any person's rights, or applicable UK laws. If you discover content in the Research Portal that you believe breaches copyright or violates any law, please contact openaccess@qub.ac.uk.

Open Access










This research has been made openly available by Queen's academics and its Open Research team. We would love to hear how access to this research benefits you. – Share your feedback with us: <http://go.qub.ac.uk/oa-feedback>



<https://doi.org/10.1038/s41467-021-22411-w>

OPEN

Author Correction: Genome-wide association study of depression phenotypes in UK Biobank identifies variants in excitatory synaptic pathways

David M. Howard , Mark J. Adams , Masoud Shirali , Toni-Kim Clarke, Riccardo E. Marioni, Gail Davies, Jonathan R. I. Coleman , Clara Alloza, Xueyi Shen , Miruna C. Barbu, Eleanor M. Wigmore, Jude Gibson & 23andMe Research Team*, Saskia P. Hagenaars, Cathryn M. Lewis , Joey Ward, Daniel J. Smith , Patrick F. Sullivan, Chris S. Haley , Gerome Breen, Ian J. Deary & Andrew M. McIntosh 

Correction to: *Nature Communications* <https://doi.org/10.1038/s41467-018-03819-3>, published online 16 April 2018.

The original version of this Article contained an error in the ‘Methods’ subsection entitled ‘Broad depression phenotype’, which incorrectly read “‘Have you ever seen a psychiatrist for nerves, anxiety, tension or depression’? (UK Biobank field 2010)’. The same error was made three times in the subsections ‘Probable MDD phenotype’ and ‘ICD-coded MDD phenotype’, which read ‘(UK Biobank fields: 2090 and 2010)’. The correct version states ‘2100’ in place of ‘2010’ in each case. This has been corrected in both the PDF and HTML versions of the Article.

Published online: 25 March 2021



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article’s Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article’s Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2021

*A list of authors and their affiliations appears online.