



Publisher Correction: Multi-ancestry genome-wide association analyses of refractive error augment genetic discovery and polygenic prediction

Correction to: *Nature Genetics*

<https://doi.org/10.1038/s41588-026-02576-0>,
published online 20 April 2026.

<https://doi.org/10.1038/s41588-026-02643-6>

Published online: 1 June 2026

Fei-Fei Cheng, Xiaoxi Liu, Hao Mi, Lizhong Wang¹, Ruilei Ma¹, Yazhou Guo¹, Julia Sidorenko¹, Chen Jiang, Tania Islam, Akira Meguro¹, Keiko Hikino¹, Yuki Ishikawa¹, Senwei Tang, Teng Li, Ruoyan Chen, Likun Wang¹, Reedik Mägi¹, Andres Metspalu¹, Estonian Biobank Research Team, 23andMe Research Team*, Masaki Takeuchi¹, Nobuhisa Mizuki¹, Hélène Choquet¹, Zi-Bing Jin, Gang Chen¹, Kun Zhou¹, Chikashi Terao¹, Jian Zeng¹ & Jian Yang¹

In the version of the article initially published, the 23andMe Research Team was missing from the online author list. The HTML version of the article has now been amended.

Open Access This article is licensed under a Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License, which permits any non-commercial use, sharing, 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 licence, and indicate if you modified the licensed material. You do not have permission under this licence to share adapted material derived from this article or parts of it. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence 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 licence, visit <http://creativecommons.org/licenses/by-nc-nd/4.0/>.

© The Author(s) 2026