

Author Correction: A genome-wide association study with 1,126,563 individuals identifies new risk loci for Alzheimer's disease

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Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-021-00921-z>, published online 7 September 2021.

In the version of this Article initially published, the author list and affiliations were incomplete. Karl Heilbron and Adam Auton, both with 23andMe, Inc., have been added to the author list. Further, the location for the 23andMe Research Team, now reading “Sunnyvale, CA” initially appeared as Mountain View, CA. The changes have been made to the online version of the paper.





*A list of authors and their affiliations appears online.

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Author Correction: A high-resolution HLA reference panel capturing global population diversity enables multi-ancestry fine-mapping in HIV host response

Yang Luo , Masahiro Kanai , Wanson Choi, Xinyi Li, Saori Sakaue , Kenichi Yamamoto, Kotaro Ogawa, Maria Gutierrez-Arcelus , Peter K. Gregersen , Philip E. Stuart , James T. Elder , Lukas Forer , Sebastian Schönherr, Christian Fuchsberger , Albert V. Smith , Jacques Fellay , Mary Carrington , David W. Haas, Xiuqing Guo, Nicholette D. Palmer , Yii-Der Ida Chen, Jerome I. Rotter , Kent D. Taylor , Stephen S. Rich , Adolfo Correa , James G. Wilson, Sekar Kathiresan, Michael H. Cho , Andres Metspalu , Tonu Esko , Yukinori Okada , Buhm Han , NHLBI Trans-Omics for Precision Medicine (TOPMed) Consortium*, Paul J. McLaren  and Soumya Raychaudhuri 

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In the version of this Analysis initially published, author Buhm Han was missing an affiliation. In addition to the Interdisciplinary Program in Bioengineering, Seoul National University, Seoul, South Korea, his affiliations should also have included the Department of Biomedical Sciences, Seoul National University College of Medicine, Seoul, South Korea.

The change has been made to the online version of the article.

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