








## Author Correction: Multi-trait analysis of genome-wide association summary statistics using MTAG

Patrick Turley, Raymond K. Walters , Omeed Maghziyan, Aysu Okbay, James J. Lee , Mark Alan Fontana, Tuan Anh Nguyen-Viet, Robbee Wedow , Meghan Zacher, Nicholas A. Furlotte, 23andMe Research Team, Social Science Genetic Association Consortium, Patrik Magnusson , Sven Oskarsson, Magnus Johannesson , Peter M. Visscher , David Laibson, David Cesarini, Benjamin M. Neale and Daniel J. Benjamin 




Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-017-0009-4>, published online 1 January 2018.

In the version of the paper initially published, no competing interests were declared. The ‘Competing interests’ statement should have stated that B.M.N. is on the Scientific Advisory Board of Deep Genomics. The error has been corrected in the HTML and PDF versions of the article.

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<https://doi.org/10.1038/s41588-019-0469-9>

## Author Correction: Linkage disequilibrium-dependent architecture of human complex traits shows action of negative selection

Steven Gazal , Hilary K. Finucane, Nicholas A. Furlotte, Po-Ru Loh, Pier Francesco Palamara, Xuanyao Liu, Armin Schoech, Brendan Bulik-Sullivan, Benjamin M. Neale , Alexander Gusev and Alkes L. Price 









Correction to: *Nature Genetics* <https://doi.org/10.1038/ng.3954>, published online 11 September 2017.

In the version of the paper initially published, information on competing interests for author Benjamin M. Neale was missing. The ‘Competing interests’ statement should have included the sentence ‘B.M.N. is on the Scientific Advisory Board of Deep Genomics’.

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## Author Correction: Atlas of group A streptococcal vaccine candidates compiled using large-scale comparative genomics

Mark R. Davies , Liam McIntyre, Ankur Mutreja, Jake A. Lacey, John A. Lees, Rebecca J. Towers, Sebastián Duchêne, Pierre R. Smeesters, Hannah R. Frost , David J. Price , Matthew T. G. Holden , Sophia David, Philip M. Giffard, Kate A. Worthing, Anna C. Seale, James A. Berkley , Simon R. Harris, Tania Rivera-Hernandez, Olga Berking, Amanda J. Cork, Rosângela S. L. A. Torres, Trevor Lithgow, Richard A. Strugnell , Rene Bergmann, Patric Nitsche-Schmitz, Gusharan S. Chhatwal, Stephen D. Bentley, John D. Fraser, Nicole J. Moreland, Jonathan R. Carapetis, Andrew C. Steer, Julian Parkhill , Allan Saul, Deborah A. Williamson, Bart J. Currie, Steven Y. C. Tong, Gordon Dougan and Mark J. Walker 

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-019-0417-8>, published online 27 May 2019.

In the version of this article initially published, in the subsection ‘GAS vaccine target variation’ in the Results, the text “The analyses revealed that 2,083 isolates carry the J8.0 allele of the M protein” should have read “The analyses revealed that 37% of the 2,083 isolates carry the J8.0 allele of the M protein.” The error has been corrected in the HTML and PDF versions of the article.

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