


Author Correction: Association analysis in over 329,000 individuals identifies 116 independent variants influencing neuroticism


Michelle Luciano , Saskia P. Hagenaars, Gail Davies, W. David Hill, Toni-Kim Clarke, Masoud Shirali , Sarah E. Harris, Riccardo E. Marioni, David C. Liewald , Chloe Fawns-Ritchie, Mark J. Adams, David M. Howard , Cathryn M. Lewis , Catharine R. Gale, Andrew M. McIntosh  and Ian J. Deary

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-017-0013-8>, published online 18 December 2017.

In the version of this article initially published, in Table 2, the descriptions of pathways and definitions in the first and last columns did not correctly correspond to the values in the other columns. The error has been corrected in the HTML and PDF versions of the article.

Published online: 29 January 2019
<https://doi.org/10.1038/s41588-019-0357-3>

Publisher Correction: An evolutionary framework for measuring epigenomic information and estimating cell-type-specific fitness consequences

Brad Gulko  and Adam Siepel 

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-018-0300-z>, published online 17 December 2018.

In the version of this article initially published, in the Methods section ‘Statistics and data analysis’, subsection ‘Measuring entropy with *INSIGHT*’, the equation for the maximized log likelihood incorrectly duplicated the equation for entropy; the equation read $H(X) = -C \sum_x P(x|\hat{\theta}) \log P(x|\hat{\theta})$ but should have read $\mathcal{L}(\hat{\theta}, X) = \max_{\theta} \log P(X|\theta) = \max_{\theta} \sum_i \log P(X_i|\theta)$. The error has been corrected in the HTML and PDF versions of the article.

Published online: 20 February 2019
<https://doi.org/10.1038/s41588-019-0369-z>