Corrections & amendments

Author Correction: Transcript expressionaware annotation improves rare variant interpretation

https://doi.org/10.1038/s41586-020-03175-7

Correction to: Nature https://doi.org/10.1038/s41586-020-2329-2

Published online 27 May 2020

Open access



Check for updates

Beryl B. Cummings, Konrad J. Karczewski, Jack A. Kosmicki, Eleanor G. Seaby, Nicholas A. Watts, Moriel Singer-Berk, Jonathan M. Mudge, Juha Karjalainen, F. Kyle Satterstrom, Anne H. O'Donnell-Luria, Timothy Poterba, Cotton Seed, Matthew Solomonson, Jessica Alföldi, Genome Aggregation Database Production Team*, Genome Aggregation Database Consortium*, Mark J. Daly & Daniel G. MacArthur

In this Article, author Marquis P. Vawter was missing from the Genome Aggregation Database Consortium list. They are associated with the affiliation: 'Department of Psychiatry & Human Behavior, University of California Irvine, Irvine, CA, USA', and contributed to the generation of the primary data incorporated into the gnomAD resource. The original Article has been corrected online.

*A list of authors and their affiliations appears online.

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/.