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Author Correction: Landscape of multi-nucleotide variants in 125,748 human exomes and 15,708 genomes

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Correction to: *Nature Communications* <https://doi.org/10.1038/s41467-019-12438-5>, published online 27 May 2020.

The original version of this Article omitted from the Genome Aggregation Database consortium the member Marquis P. Vawter, from the Department of Psychiatry & Human Behavior, University of California Irvine, Irvine, CA, USA. Additionally, the following was added to the Author Contributions: 'All authors listed under The Genome Aggregation Database Consortium contributed to the generation of the primary data incorporated into the gnomAD resource'.

This has been corrected in both the PDF and HTML versions of the Article.

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