Corrections & amendments

Author Correction: The mutational constraint spectrum quantified from variation in 141,456 humans

Check for updates

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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. In addition, in the legend to Fig. 1, 'ten' should have been 'seven' in the sentence: "**a**, Uniform manifold approximation and projection (UMAP)[46,47] plot depicting the ancestral diversity of all individuals in gnomAD, using **seven** principal components." The original Article has been corrected online.

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