CORRECTION

Correction: Exploring the Distribution of Genetic Markers of Pharmacogenomics Relevance in Brazilian and Mexican Populations

The *PLOS ONE* Staff

There is an error in the second sentence in the third paragraph of the “Genetic differentiation and haplotype diversity for DMET Plus markers” section of the Results. The correct sentence is: Although, the array captures many of the functional variants described for these genes, unfortunately *UGT1A1**28 was not questioned in our study.

The [S5 Table](#) file does not open properly. Please view the correct [S5 Table](#) below.

Supporting Information

S5 Table. Supplementary dataset. Minor Allele Frequencies (MAF) for 1,647 genetic markers. The markers are those included in the DMET Plus array and for all populations: Europeans (Hapmap CEU), Africans (Hapmap YRI and LWK), Native Americans (Zapotecas), Brazilians and Mexican mestizos (Guanajuato (GUA); Guerrero (GUE); Sonora (SON); Veracruz (VER) and Yucatan (YUC)).
(XLSX)

Reference

1. Bonifaz-Peña V, Contreras AV, Struchiner CJ, Roela RA, Furuya-Mazzotti TK, Chammas R, et al. (2014) Exploring the Distribution of Genetic Markers of Pharmacogenomics Relevance in Brazilian and Mexican Populations. PLoS ONE 9(11): e112640. doi: [10.1371/journal.pone.0112640](https://doi.org/10.1371/journal.pone.0112640) PMID: [25419701](https://pubmed.ncbi.nlm.nih.gov/25419701/)

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Citation: The *PLOS ONE* Staff (2015) Correction: Exploring the Distribution of Genetic Markers of Pharmacogenomics Relevance in Brazilian and Mexican Populations. PLoS ONE 10(3): e0122161. doi:10.1371/journal.pone.0122161

Published: March 23, 2015

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