

CORRECTION

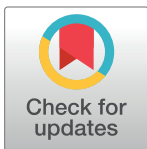
Correction: Leveraging correlations between variants in polygenic risk scores to detect heterogeneity in GWAS cohorts

Jie Yuan, Hengrui Xing, Alexandre Louis Lamy, The Schizophrenia Working Group of the Psychiatric Genomics Consortium, Todd Lencz, Itsik Pe'er

The second author's name is spelled incorrectly. The correct name is: Hengrui Xing.

Reference

1. Yuan J, Xing H, Lamy AL, The Schizophrenia Working Group of the Psychiatric Genomics Consortium, Lencz T, Pe'er I (2020) Leveraging correlations between variants in polygenic risk scores to detect heterogeneity in GWAS cohorts. PLoS Genet 16(9): e1009015. <https://doi.org/10.1371/journal.pgen.1009015> PMID: [32956347](https://pubmed.ncbi.nlm.nih.gov/32956347/)



OPEN ACCESS

Citation: Yuan J, Xing H, Lamy AL, The Schizophrenia Working Group of the Psychiatric Genomics Consortium, Lencz T, Pe'er I (2020) Correction: Leveraging correlations between variants in polygenic risk scores to detect heterogeneity in GWAS cohorts. PLoS Genet 16(10): e1009158. <https://doi.org/10.1371/journal.pgen.1009158>

Published: October 23, 2020

Copyright: © 2020 Yuan et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.