

CORRECTION

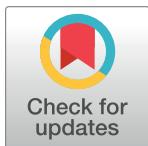
Correction: Enhancing genomic selection by fitting large-effect SNPs as fixed effects and a genotype-by-environment effect using a maize BC₁F_{3:4} population

Dongdong Li, Zhenxiang Xu, Riliang Gu, Pingxi Wang, Demar Lyle, Jialiang Xu, Hongwei Zhang, Guoying Wang

The eighth author's name is spelled incorrectly. The correct name is: Guoying Wang.

Reference

- Li D, Xu Z, Gu R, Wang P, Lyle D, Xu J, et al. (2019) Enhancing genomic selection by fitting large-effect SNPs as fixed effects and a genotype-by-environment effect using a maize BC₁F_{3:4} population. PLoS ONE 14(10): e0223898. <https://doi.org/10.1371/journal.pone.0223898> PMID: 31622400



OPEN ACCESS

Citation: Li D, Xu Z, Gu R, Wang P, Lyle D, Xu J, et al. (2019) Correction: Enhancing genomic selection by fitting large-effect SNPs as fixed effects and a genotype-by-environment effect using a maize BC₁F_{3:4} population. PLoS ONE 14(12): e0226592. <https://doi.org/10.1371/journal.pone.0226592>

Published: December 12, 2019

Copyright: © 2019 Li et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.