



# **OPEN ACCESS**

## Approved by:

Frontiers in Editorial Office, Frontiers Media SA, Switzerland

## \*Correspondence:

Susan R. McCouch srm4@cornell.edu Georgia C. Eizenga georgia.eizenga@usda.gov

# †ORCID:

Liakat Ali orcid.org/0000-0002-1180-5846 Ehsan Shakiba orcid.org/0000-0001-7264-6351 Diane R. Wang orcid.org/0000-0002-2290-3257

<sup>‡</sup>These authors share first authorship

# §Present address:

Diane R. Wana. Department of Agronomy, Purdue University, West Lafayette, IN, United States I iakat Ali Department of Plant Sciences, University of Missouri-Columbia Delta Center, Portageville, MO, United States HyunJung Kim, LG Chemical, Ltd., Seoul, South Korea Ju-Won Kana. Department of Southern Area Crop Science, National Institute of Crop Science, RDA, Miryang, South Korea

# Specialty section:

This article was submitted to Plant Breeding, a section of the journal Frontiers in Plant Science

Received: 10 December 2020 Accepted: 11 December 2020 Published: 02 March 2021

# Citation:

Singh N, Wang DR, Ali L, Kim H,
Akther KM, Harrington SE, Kang J-W,
Shakiba E, Shi Y, DeClerck G,
Meadows B, Govindaraj V, Ahn S-N,
Eizenga GC and McCouch SR (2021)
Corrigendum: A Coordinated Suite of
Wild-Introgression Lines in Indica and
Japonica Elite Backgrounds.
Front. Plant Sci. 11:640122.
doi: 10.3389/fpls.2020.640122

# Corrigendum: A Coordinated Suite of Wild-Introgression Lines in *Indica* and *Japonica* Elite Backgrounds

Namrata Singh<sup>1‡</sup>, Diane R. Wang<sup>1†‡§</sup>, Liakat Ali<sup>2†§</sup>, HyunJung Kim<sup>1§</sup>, Kazi M. Akther<sup>1</sup>, Sandra E. Harrington<sup>1</sup>, Ju-Won Kang<sup>3§</sup>, Ehsan Shakiba<sup>2†</sup>, Yuxin Shi<sup>1</sup>, Genevieve DeClerck<sup>1</sup>, Byron Meadows<sup>1</sup>, Vishnu Govindaraj<sup>1</sup>, Sang-Nag Ahn<sup>3</sup>, Georgia C. Eizenga<sup>4\*</sup> and Susan R. McCouch<sup>1\*</sup>

<sup>1</sup> Plant Breeding and Genetics Section, School of Integrative Plant Science, Cornell University, Ithaca, NY, United States, <sup>2</sup> Rice Research and Extension Center, University of Arkansas, Stuttgart, AR, United States, <sup>3</sup> Department of Agronomy, Chungnam National University, Daejeon, South Korea, <sup>4</sup> USDA-ARS Dale Bumpers National Rice Research Center, Stuttgart, AR, United States

Keywords: Oryza sativa, crop wild relatives, Oryza rufipogon Species Complex, chromosome segment substitution line, pre-breeding resources

# A Corrigendum on

A Coordinated Suite of Wild-Introgression Lines in *Indica* and *Japonica* Elite Backgrounds by Singh, N., Wang, D. R., Ali, L., Kim, H., Akther, K. M., Harrington, S. E., et al. (2020). Front. Plant Sci. 11:564824. doi: 10.3389/fpls.2020.564824

In the original article, there was a mistake in **Data Availability Statement** as published. A correction has been made to **Data Availability Statement**. The corrected **Statement** appears below.

The genotypic datasets presented in this study can be found in Supplementary Data 2—Genotype information (C7AIR) for CSSL populations. Sequencing data for grain and hull color genes from the wild and cultivated parents and selected CSSLs is available in GenBank with the following Accession IDs: *Bh4* (MW310629 - MW310643), *Rd* (MW310644 - MW310658), *Rc* (MW310659 - MW310674), *Ph* (MW310675 - MW310688).

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Copyright © 2021 Singh, Wang, Ali, Kim, Akther, Harrington, Kang, Shakiba, Shi, DeClerck, Meadows, Govindaraj, Ahn, Eizenga and McCouch. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

1