Erratum

The unique epigenetic features of Pack-MULEs and their impact on chromosomal base composition and expression spectrum

Dongyan Zhao¹,², John P. Hamilton², Brieanne Vaillancourt², Wenli Zhang³,⁴, Georgia C. Eizenga⁵, Yuehua Cui⁶, Jiming Jiang¹,², C. Robin Buell²,* and Ning Jiang¹,⁷,*

¹Department of Horticulture, Michigan State University, East Lansing, MI 48824, USA, ²Department of Plant Biology, Michigan State University, East Lansing, MI 48824, USA, ³Department of Horticulture, University of Wisconsin, Madison, WI 53705, USA, ⁴State Key Laboratory for Crop Genetics and Germplasm Enhancement, Nanjing Agriculture University, Nanjing, Jiangsu 210095, China, ⁵USDA-ARS Dale Bumpers National Rice Research Center, 2890 Highway 130 East, Stuttgart, AR 72160, USA, ⁶Department of Statistics and Probability, Michigan State University, East Lansing, MI 48824, USA and ⁷Program in Ecology, Evolutionary Biology and Behavior, Michigan State University, East Lansing, MI 48824, USA

Nucleic Acids Research, gky025, https://doi.org/10.1093/nar/gky025

The details of the corresponding authors of this paper have been corrected online and in print. Correspondence should be addressed to Robin Buell, Tel: +1 517 353 5597; Email: buell@msu.edu, Ning Jiang. Tel: +1 517 353 0381; Email: jiangn@msu.edu.

*To whom correspondence should be addressed. C. Robin Buell. Tel: +1 517 353 5597; Email: buell@msu.edu, Ning Jiang. Tel: +1 517 353 0381; Email: jiangn@msu.edu

© The Author(s) 2018. Published by Oxford University Press on behalf of Nucleic Acids Research. This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by-nc/4.0/), which permits non-commercial re-use, distribution, and reproduction in any medium, provided the original work is properly cited. For commercial re-use, please contact journals.permissions@oup.com