

doi: 10.1093/gigascience/giaa040 Expression of concern

EXPRESSION OF CONCERN

Expression of concern: Dissection of soybean populations according to selection signatures based on whole-genome sequences

Jae-Yoon Kim ଢ1,2,†, Seongmun Jeong1,†, Kyoung Hyoun Kim1,2, Won-Jun Lim1,2, Ho-Yeon Lee1,2, Namhee Jeong3, Jung-Kyung Moon3 and Namshin Kim ଢ1,2,*

¹Genome Editing Research Center, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Gwahak-ro 125, Yuseong-gu, Daejeon 34141, Republic of Korea; ²Department of Bioinformatics, KRIBB School of Bioscience, University of Science and Technology (UST), Gajeong-ro 217, Yuseong-gu, Daejeon 34141, Republic of Korea and ³National Institute of Crop Science, Rural Development Administration, Nongsaengmyeong-ro 370, Deokjin-gu, Jeon-Ju 54874, Republic of Korea

*Correspondence address. Namshin Kim, Genome Editing Research Center, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Gwahak-ro 125, Yuseong-gu, Daejeon 34141, Republic of Korea. Tel/Fax: +82-42-879-8162. E-mail: deepreds@kribb.re.kr http://orcid.org/0000-0001-6361-274X these authors contributed equally to this research.

In 2019, GigaScience published the article "Dissection of soybean populations according to selection signatures based on whole-genome sequences," by Jae-Yoon Kim and colleagues: https://doi.org/10.1093/gigascience/giz151. In January 2020, the editors received credible information about attribution problems with the source of the data in the article. In an

attempt to address the concerns raised, the authors flagged some additional, potential issues with the presentation and content of their article. We alert our readers that concerns have been raised about this article, and that a full institutional investigation is in progress.