RETRACTION

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

Jae-Yoon Kim [1,2,†], Seongmun Jeong[1,†], Kyoung Hyoun Kim[1,2], Won-Jun Lim[1,2], Ho-Yeon Lee[1,2], Namhee Jeong[3], Jung-Kyung Moon[3] and Namshin Kim [1,2,*]

[1]Genome Editing Research Center, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Gwahak-ro 125, Yuseong-gu, Daejeon 34141, Republic of Korea; [2]Department of Bioinformatics, KRIBB School of Bioscience, University of Science and Technology (UST), Gajeong-ro 217, Yuseong-gu, Daejeon 34141, Republic of Korea and [3]National 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 December 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 2020, the editors received credible information about attribution problems with the source of the data in the article and subsequently alerted the readers with a statement of concern: https://doi.org/10.1093/gigascience/giaa040. A full institutional investigation has now been completed, which has concluded that research misconduct was carried out. As a result, this article is being retracted by the editors and authors.