iScience



Correction

Citrus genomic resources unravel putative genetic determinants of Huanglongbing pathogenicity

Yuxia Gao, Jin Xu, Zhilong Li, Yunzeng Zhang, Nadia Riera, Zhiwei Xiong, Zhigang Ouyang, Xinjun Liu, Zhanjun Lu, Danelle Seymour, Balian Zhong,* and Nian Wang*

(iScience 26, 106024; February 17, 2023)

In the originally published version of this article, the first column of the supplementary table Data S2 contained some errors caused by sorting. This has now been corrected, and the discussion and conclusions of this paper remain unchanged.

In addition, the assembled genome sequences of new sequenced genome for *C. sinensis* "Newhall" were deposited in Zenodo (https://zenodo.org/record/8176803) and Citrus Genome Database (https://www.citrusgenomedb.org/) instead of the NCBI database due to technical issues. The data and code availability section has been updated to reflect this change. The authors apologize for any confusion caused.

*Correspondence: zhongbalian@gnnu.edu.cn (B.Z.), nianwang@ufl.edu (N.W.) https://doi.org/10.1016/j.isci.2023.108255

