Erratum to: Transcriptionally and Phylogenetically Analyzing the P Protein Gene of Glycine Decarboxylase for Understanding the Evolution of C₃-C₄ Species in Brassicaceae

Chunyu Zhang¹, In-ae Park², Fangsen Xu¹, Maoteng Li³, Yong Pyo Lim^{2*}, and Jinling Meng¹

¹National Key Laboratory of Crop Genetic Improvement, Huazhong Agricultural University, Wuhan 430070, China

²Department of Horticulture, Chungnam National University, Daejeon 305-764, Korea

³College of Life Science and Technology, Huazhong University of Science and Technology, Wuhan 430074, China

*Corresponding author: yplim@cnu.ac.kr

© Korean Society for Horticultural Science and Springer 2013

Erratum to: Hort. Environ. Biotechnol. 52(4):427-434. 2011.
DOI 10.1007/s13580-011-0215-4

The original version of this article contained an error in the Fig. 2C.
The correct Fig. 2 is reproduced below.

Fig. 2. Deletion analysis of the promoter of *BnGDCP* in C₃ species (*Arabidopsis*). (A) Structures of the deleted promoter/*GUS* chimerical genes used for *Arabidopsis* transformation. *-nP-BnGDCP* represents a series of deleted promoters located at “n” position upstream of AUG initiation codon of *BnGDCP*. (B), (C), and (D) show the histochemical localization of *GUS* activity in a whole leaf, leaf section, and floral tissues of *Arabidopsis* plant transformed with *-0.9P-BnGDCP:GUS* construct, respectively. (E), (F), and (G) show the histochemical localization of *GUS* activity in a whole leaf, leaf section, and floral tissues of *Arabidopsis* plants transformed with *-0.8P-BnGDCP:GUS* construct, respectively.