

SCIENTIFIC REPORTS

OPEN

Author Correction: The bHLH transcription factor CgbHLH001 is a potential interaction partner of CDPK in halophyte *Chenopodium glaucum*

Juan Wang^{1,2}, Gang Cheng¹, Cui Wang¹, Zhuanzhuan He¹, Xinxin Lan¹, Shiyue Zhang¹ & Haiyan Lan¹

Correction to: *Scientific Reports* <https://doi.org/10.1038/s41598-017-06706-x>, published online 16 August 2017

This Article contains errors in the Results section, under the subheading ‘Analysis of interaction between CDPK and bHLH in *C. glaucum*’,

“The three top ranking models of the CgCDPK-CgbHLH001 interaction complex were presented in Fig. 5c”
should read:

“The three top ranking models of the CgCDPK-CgbHLH001 interaction complex were presented in Fig. 6c”

In addition,

“Prediction of phosphorylation sites of CgbHLH001 revealed 39 potential amino acid residues (Fig. 6d), in which the motif of ⁶¹-GKRLKS-⁶⁶ was a putative action site for CgCDPK.”

should read:

“Prediction of phosphorylation sites of CgbHLH001 revealed 39 potential amino acid residues (Fig. 6d), in which the motif of ⁹¹-GKRLKS-⁹⁶ was a putative action site for CgCDPK.”



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article’s Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article’s Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2018

¹Xinjiang Key Laboratory of Biological Resources and Genetic Engineering, College of Life Science and Technology, Xinjiang University, Urumqi, 830046, China. ²Institute of Economic Crops, Xinjiang Academy of Agricultural Sciences, Urumqi, 830091, China. Juan Wang and Gang Cheng contributed equally. Correspondence and requests for materials should be addressed to H.L. (email: lanhaiyan@xju.edu.cn)