

Published online: 10 March 2020

OPEN Author Correction: Evolutionary and expression analysis of CAMTA gene family in Nicotiana tabacum yielded insights into their origin, expansion and stress responses

Kaleem U. Kakar, Zarqa Nawaz, Zhouqi Cui, Peijian Cao 🕟, Jingjing Jin, Qingyao Shu 🗈 & **Xueliang Ren**

Correction to: Scientific Reports https://doi.org/10.1038/s41598-018-28148-9, published online 09 July 2018

In the original version of this Article, Kaleem U. Kakar was incorrectly affiliated with 'Faculty of Life Sciences, University of Balochistan, Quetta, 87300, Pakistan'. The correct affiliations are listed below.

Molecular Genetics Key Laboratory of China Tobacco, Guizhou Academy of Tobacco Science, Guiyang, 550081, China

State Key Laboratory of Rice Biology, Institution of Crop Science, Zhejiang University, Hangzhou, 310058, China

Department of Microbiology, Faculty of Life Sciences & Informatics, Balochistan University of Information Technology, Engineering, and Management Sciences, Quetta, 87300, Pakistan

This error has now been corrected in the PDF and HTML versions of the Article.

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) 2020