

CORRECTION Open Access

Correction: Transcriptomic analysis of *Stropharia rugosoannulata* reveals carbohydrate metabolism and cold resistance mechanisms under low-temperature stress

Haibo Hao^{1†}, Jinjing Zhang^{2†}, Shengdong Wu¹, Jing Bai², Xinyi Zhuo², Jiaxin Zhang², Benke Kuai^{1*} and Hui Chen^{2*}

Correction to: AMB Express (2022) 12(1): 1-5 https://doi.org/10.1186/s13568-022-01400-2

Following publication of the original article (Hao et al. 2022), the authors reported an error in the author group. The typesetter has inadvertently missed to include the dagger symbol for first author. Now this has been corrected with this erratum.

The original article (Hao et al. 2022) has been updated.

Reference

Hao H, Zhang J, Wu S, Bai J, Zhuo X, Zhang J, Kuai B, Chen H (2022) Transcriptomic analysis of Stropharia rugosoannulata reveals carbohydrate metabolism and cold resistance mechanisms under low-temperature stress. AMB Express 12(1):1–5

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Published online: 13 June 2022

The original article can be found online at https://doi.org/10.1186/s13568-022-01400-2.

[†]Haibo Hao and Jinjing Zhang contributed equally to this work

National Research Center for Edible Fungi Biotechnology and Engineering, Key Laboratory of Applied Mycological Resources and Utilization, Ministry of Agriculture, Shanghai Key Laboratory of Agricultural Genetics and Breeding, Institute of Edible Fungi, Shanghai Academy of Agricultural Sciences, 309 Room, No. 1000, Jinqi Road, FengXian District, Shanghai 201403, China



© The Author(s) 2022. **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 licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence 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 licence, visit http://creativecommons.org/licenses/by/4.0/.

^{*}Correspondence: bkkuai@fudan.edu.cn; Huichen js@aliyun.com

¹ State Key Laboratory of Genetic Engineering and Fudan Center for Genetic Diversity and Designing Agriculture, Institute of Plant Biology, School of Life Sciences, Fudan University, 413 Room, No. 2005, Songhu Road, YangPu District, Shanghai 200438, China