

OPEN ACCESS

APPROVED BY

Frontiers Editorial Office, Frontiers Media SA, Switzerland

*CORRESPONDENCE

Zhi-Qiang Wu

Wen-Chuang He

Ying-Xue Yang

[†]These authors have contributed equally to this work and share last authorship

RECEIVED 28 September 2023 ACCEPTED 29 September 2023 PUBLISHED 18 October 2023

CITATION

Xiang K-L, Mao W, Peng H-W, Erst AS, Yang Y-X, He W-C and Wu Z-Q (2023) Corrigendum: Organization, phylogenetic marker exploitation, and gene evolution in the plastome of *Thalictrum* (Ranunculaceae). *Front. Plant Sci.* 14:1303516. doi: 10.3389/fpls.2023.1303516

COPYRIGHT

© 2023 Xiang, Mao, Peng, Erst, Yang, He and Wu. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Corrigendum: Organization, phylogenetic marker exploitation, and gene evolution in the plastome of *Thalictrum* (Ranunculaceae)

Kun-Li Xiang^{1,2}, Wei Mao³, Huan-Wen Peng^{2,4}, Andrey S. Erst^{5,6}, Ying-Xue Yang^{1*†}, Wen-Chuang He^{1*†} and Zhi-Qiang Wu^{1,7*†}

¹Shenzhen Branch, Guangdong Laboratory of Lingnan Modern Agriculture, Genome Analysis Laboratory of the Ministry of Agriculture and Rural Affairs, Agricultural Genomics Institute at Shenzhen, Chinese Academy of Agricultural Sciences, Shenzhen, China, ²State Key Laboratory of Systematic and Evolutionary Botany, Institute of Botany, Chinese Academy of Sciences, Beijing, China, ³College of Ecology and Environment, Hainan University, Haikou, China, ⁴College of Life Sciences, University of Chinese Academy of Sciences, Beijing, China, ⁵Central Siberian Botanical Garden, Russian Academy of Sciences, Novosibirsk, Russia, ⁶Laboratory Herbarium (TK), Tomsk State University, Tomsk, Russia, ⁷Kunpeng Institute of Modern Agriculture at Foshan, Chinese Academy of Agricultural Sciences, Foshan, China

KEYWORDS

Thalictrum, plastid genome, genome structure, molecular markers, phylogeny

A Corrigendum on

Organization, phylogenetic marker exploitation, and gene evolution in the plastome of Thalictrum (Ranunculaceae)

by Xiang K-L, Mao W, Peng H-W, Erst AS, Yang Y-X, He W-C and Wu Z-Q (2022) Front. Plant Sci. 13:897843. doi: 10.3389/fpls.2022.897843.

Error in Author List

In the published article, there was an error in the **author list**. Authors Ying-Xue Yang, Wen-Chuang He and Zhi-Qiang Wu did not have an indication of equal contribution. The corrected author list appears below:

Kun-Li Xiang^{1,2}, Wei Mao³, Huan-Wen Peng^{2,4}, Andrey S. Erst^{5,6}, Ying-Xue Yang^{1*†}, Wen-Chuang He^{1*†} and Zhi-Qiang Wu^{1,7*†}

¹Shenzhen Branch, Guangdong Laboratory of Lingnan Modern Agriculture, Genome Analysis Laboratory of the Ministry of Agriculture and Rural Affairs, Agricultural Genomics Institute at Shenzhen, Chinese Academy of Agricultural Sciences, Shenzhen, China, ²State Key Laboratory of Systematic and Evolutionary Botany, Institute of Botany, Chinese Academy of Sciences, Beijing, China, ³College of Ecology and Environment, Hainan University, Haikou, China, ⁴College of Life Sciences, University of Chinese Academy of Sciences, Beijing, China, ⁵Central Siberian Botanical Garden, Russian Academy of Sciences, Novosibirsk, Russia, ⁶Laboratory Herbarium (TK), Tomsk State University, Tomsk, Russia, ⁷Kunpeng Institute of Modern Agriculture at Foshan, Chinese Academy of Agricultural Sciences, Foshan, China

Xiang et al. 10.3389/fpls.2023.1303516

*Correspondence: Zhi-Qiang Wu, wuzhiqiang@caas.cn; Wen-Chuang He, hewenchuang@caas.cn; Ying-Xue Yang, yyxue32@163.com

[†]These authors have contributed equally to this work and share last authorship.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.