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Correction

Correction: Liu et al. The Genome-Wide Identification of Stable Internal Reference Genes Related to Delayed Spoilage for Accurate qRT-PCR Normalization in Ethephon-Treated *Pueraria thomsonii* Benth. *Horticulturae* 2023, *9*, 1014

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Missing Citation

In the original publication [1], **References [46,47]** were not cited. The citation has now been inserted in ***2.8. Gene Expression Stability Analysis of Candidate IRGs*** and should read:

In addition, RefFinder (https://blooge.cn/RefFinder/?type=reference#tabs-1, accessed on 20 October 2020) was employed to check the output of three programs (GeNorm, NormFinder, and BestKeeper) [45–47], as well an additional stability ranking method (ΔCt method) to rank all the genes in the *P. thomsonii* samples.

In this article, we utilized the software tool "RefFinder" by Xie et al to perform a thorough analysis of the results obtained from four different analysis software (delta Ct method, Bestkeeper, NormFinder, and GeNorm). In order to acknowledge the original publications mentioned on the RefFinder, we have cited the original references in the relevant sections of the article. The citation order of the references in the corrected manuscript have been adjusted accordingly following the numerical order after the change. The authors apologize for any inconvenience caused and state that the scientific conclusions are unaffected. This correction was approved by the Academic Editor. The original publication has also been updated.

Reference

1. Liu, Y.; Li, Y.; He, B.; Yu, B.; Liu, B.; Ma, C.; Chen, J.; Zhang, T.; Zhang, H.; Liu, Y.; et al. The Genome-Wide Identification of Stable Internal Reference Genes Related to Delayed Spoilage for Accurate qRT-PCR Normalization in Ethephon-Treated *Pueraria thomsonii* Benth. *Horticulturae* **2023**, *9*, 1014. [CrossRef]

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Citation: Liu, Y.; Li, Y.; He, B.; Yu, B.; Liu, B.; Ma, C.; Chen, J.; Zhang, T.; Zhang, H.; Liu, Y.; et al. Correction: Liu et al. The Genome-Wide Identification of Stable Internal Reference Genes Related to Delayed Spoilage for Accurate qRT-PCR Normalization in Ethephon-Treated *Pueraria thomsonii* Benth. *Horticulturae* 2023, *9*, 1014. *Horticulturae* **2023**, *9*, 1250. https://doi.org/10.3390/horticulturae9121250

Received: 18 October 2023
Accepted: 19 October 2023
Published: 21 November 2023

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