



Corrigendum: Reprogramming of Cell Fate During Root Regeneration by Transcriptional and Epigenetic Networks

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A Corrigendum on

Reprogramming of Cell Fate During Root Regeneration by Transcriptional and Epigenetic Networks

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Jing T, Ardiansyah R, Xu Q, Xing Q and Müller-Xing R (2021) Corrigendum: Reprogramming of Cell Fate During Root Regeneration by Transcriptional and Epigenetic Networks. Front. Plant Sci. 12:698412. doi: 10.3389/fpls.2021.698412 In the original article, **Tables 1** and **2** contained several errors as published. The errors are indicated below and the article has been updated.

In Table 1, of the 24 genes named, the following eight were spelled incorrectly:

- "ABCB19 (MDR1)" was presented incorrectly as "ABCB19 (MDRlj)"
- "ASA1" was presented incorrectly as "ASM"
- "COII" was presented incorrectly as "con"
- "GA1 (CPS1), GA5 (GA20OX1)" was presented incorrectly as "GA1 (1), GA5 (GA200X1)"
- "IAA14 (SLR)" was presented incorrectly as "IAA14 SLR (IAA14)"
- "TAA1 (WEI8), TAR2" was presented incorrectly as "TAA1 (WEIS), TAR2"
- "WOX5,7" was presented incorrectly as " $W0 \times 5,7$ "

- "WOX11,12" was presented incorrectly as " $W0 \times 11,12$ "

Several classifications of expression and/or mutant phenotypes were incorrect in Table 1.

In **Table 1**, the citation to reference "Du and Scheres (2018)" should have been to "**Du and Scheres** (2017)"; the reference in question was omitted from the original article and appears in the Reference section below.

Table 1 had several minor typographical errors and omitted the explanations of the following abbreviations from the footnote: AR, adventitious root; ARP, adventitious root primordia; LRP, lateral root primordia; OE, overexpression; QC, quiescent center; RAM, root apical meristem.

Finally, the order of the genes in Table 1 was incorrect, with "AUX1, LAX3" appearing out of sequence.

Table 2 had several minor typographical errors and omitted the explanations of the following abbreviations from the footnote: AR, adventitious root; LRP, lateral root primordia; RAM, root apical meristem; QC, quiescent center.

The authors apologize for these errors and state that they do not change the scientific conclusions of the review article in any way. The original article has been updated.

REFERENCES

Du, Y., and Scheres, B. (2017). PLETHORA transcription factors orchestrate de novo organ patterning during Arabidopsis lateral root outgrowth. Proc. Natl. Acad. Sci. U.S.A. 114, 11709–11714. doi: 10.1073/pnas.171441 0114 Copyright © 2021 Jing, Ardiansyah, Xu, Xing and Müller-Xing. This is an openaccess 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.