

PUBLISHER CORRECTION

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Publisher Correction: The haplotype-resolved genome assembly of an ancient citrus variety provides insights into the domestication history and fruit trait formation of loose-skin mandarins

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The original article can be found online at <https://doi.org/10.1186/s13059-025-03535-4>.

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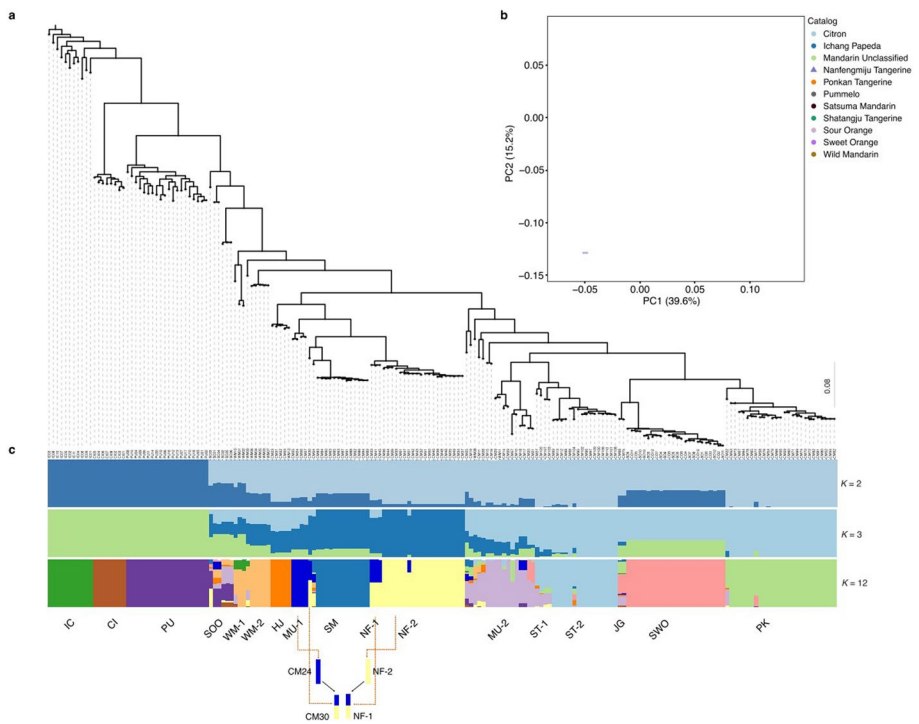
<https://doi.org/10.1186/s13059-025-03535-4>

Following publication of the original article [1], the authors identified a typesetting error in the PDF version of the article. Figure 2b was erroneously missing content.

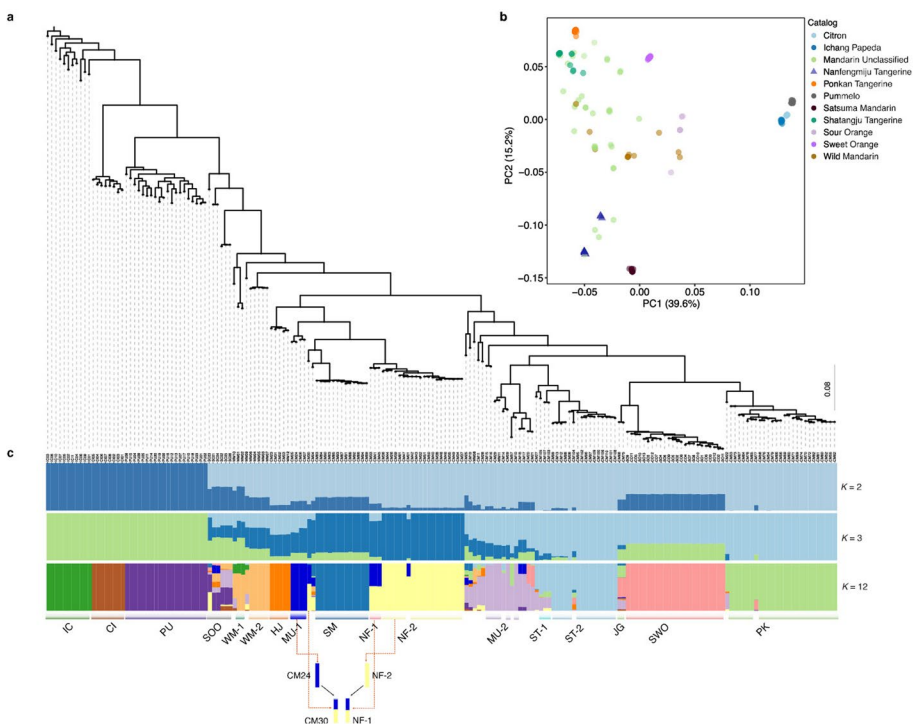


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The incorrect Fig. 2 is as follows:



The correct Fig. 2 is as follows:



The original article [1] has been corrected.

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Reference

1. Yin M, Song X, He C, et al. The haplotype-resolved genome assembly of an ancient citrus variety provides insights into the domestication history and fruit trait formation of loose-skin mandarins. *Genome Biol.* 2025;26:61. <https://doi.org/10.1186/s13059-025-03535-4>.