

CORRECTION

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Correction: Comprehensive *in silico* analysis of the underutilized crop tef (*Eragrostis tef* (Zucc.) Trotter) genome reveals drought tolerance signatures

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Correction: *BMC Plant Biol* 23, 506 (2023)

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Following publication of the original article [1], author spotted errors in Fig. 8. Revised Fig. 8 was mistakenly not provided during production process. The correct figure is given below:

The corrections do not affect the overall result or conclusion of the article. The original article has been corrected.

The original article can be found online at <https://doi.org/10.1186/s12870-023-04515-1>.

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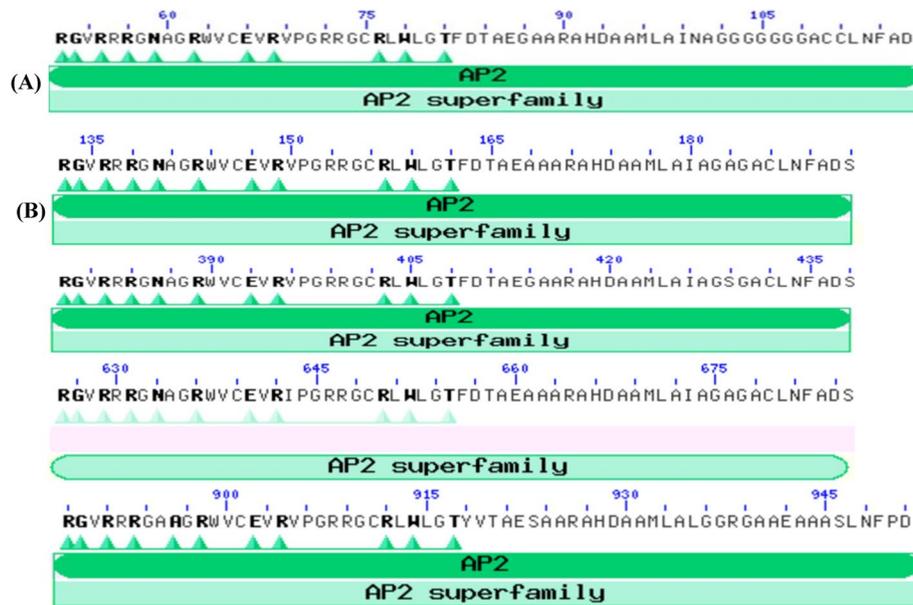


Fig. 8 Conserved domain analysis of selected genes. **A** Rice DREB1A with single AP2 binding site, and **B** Tef DREB1A with four structurally related copes within single transcriptional machinery. The amino acids in bold are predicted significant binding sites

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Reference

1. Bekele-Alemu A, Ligaba-Osena A. Comprehensive *in silico* analysis of the underutilized crop tef (*Eragrostis tef* (Zucc.) Trotter) genome reveals drought tolerance signatures. *BMC Plant Biol.* 2023;23:506. <https://doi.org/10.1186/s12870-023-04515-1>.