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Correction to: New insights on key genes involved in drought stress response of barley: gene networks reconstruction, hub, and promoter analysis



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Correction to: J Genet Eng Biotechnol 19, 2 (2021) https://doi.org/10.1186/s43141-020-00104-z

Following publication of the original article [1], the authors reported an error in the PDF version for Figs. 2, 3 and 4 which weren't the latest version even though the HTML version is. The revised Figures are indicated hereafter.

The changes requested are implemented in this correction and the original article [1] has been corrected.

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Published online: 28 May 2021

Reference

 Javadi SM, Shobbar ZS, Ebrahimi A et al (2021) New insights on key genes involved in drought stress response of barley: gene networks reconstruction, hub, and promoter analysis. J Genet Eng Biotechnol 19:2. https://doi.org/10.1186/s43141-020-00104-z

The original article can be found online at https://doi.org/10.1186/s43141-020-00104-z.

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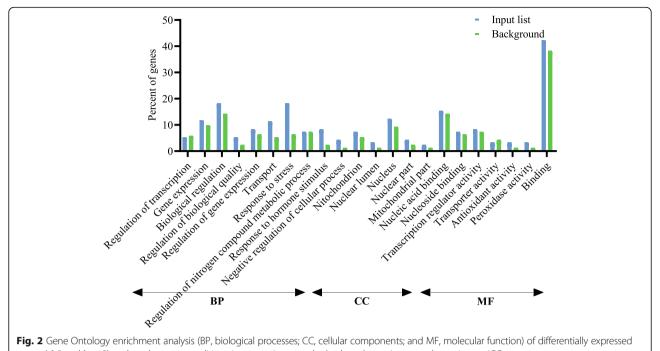


Fig. 2 Gene Ontology enrichment analysis (BP, biological processes; CC, cellular components; and MF, molecular function) of differentially expressed genes (\leq 5 and \geq - 5) at drought stress conditions in vegetative stage barley based on microarray data using agriGO

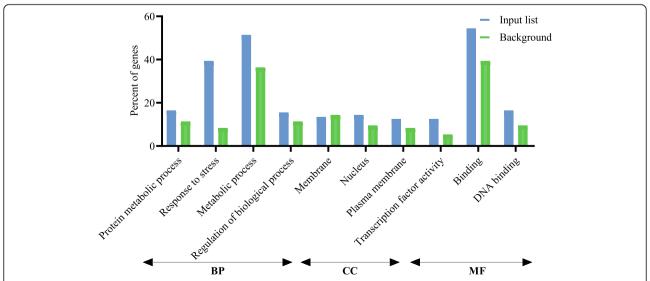


Fig. 3 Gene Ontology enrichment analysis (BP, biological processes; CC, cellular components; and MF, molecular function) of differentially expressed genes (\leq 5 and \geq -5) at drought stress conditions in reproductive stage barley based on microarray data using agriGO

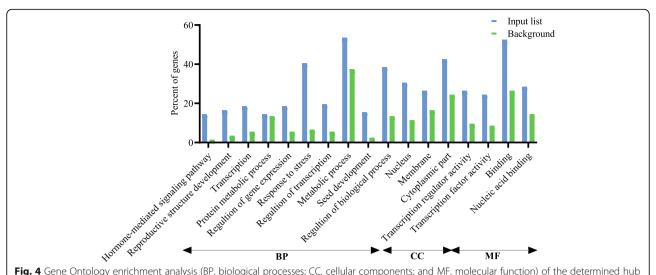


Fig. 4 Gene Ontology enrichment analysis (BP, biological processes; CC, cellular components; and MF, molecular function) of the determined hub genes at drought stress conditions in vegetative and reproductive stage barley based on microarray data using agriGO