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## Correction to: Transcriptome and de novo analysis of *Rosa xanthina* f. *spontanea* in response to cold stress

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Correction to: BMC Plant Biol 21, 472 (2021) https://doi.org/10.1186/s12870-021-03246-5

Following publication of the original article [1], due to typesetting problem, Figs. 1 and 2 were published by mistakes. Therefore, Figs. 1 and 2 have been corrected and listed as follows. I am sorry for the inconvenience.

The correction does not have any effect on the results or conclusions of the paper. The original article has been corrected.

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Published online: 30 December 2021

## Reference

 Zhuang D, Ma C, Xue L, et al. Transcriptome and de novo analysis of Rosa xanthina f. spontanea in response to cold stress. BMC Plant Biol. 2021;21:472. https://doi.org/10.1186/s12870-021-03246-5.

The original article can be found online at https://doi.org/10.1186/s12870-021-03246-5.

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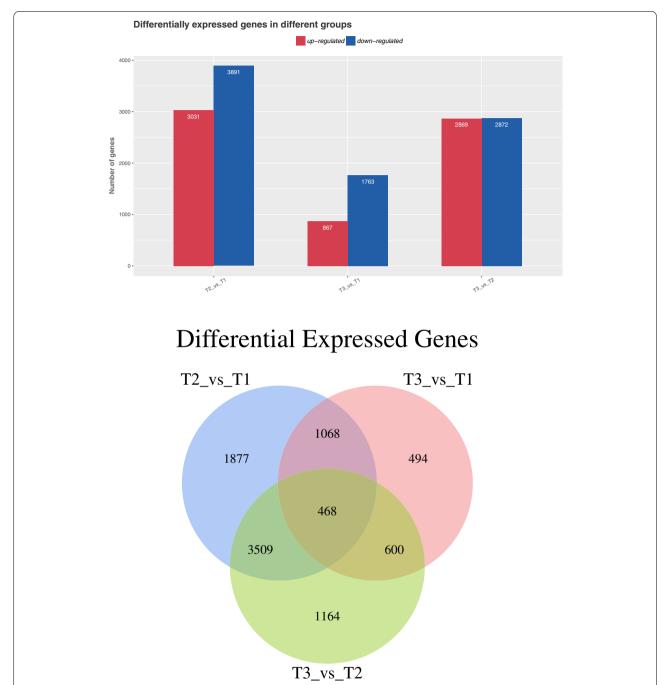


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Zhuang et al. BMC Plant Biology (2021) 21:606 Page 2 of 3



**Fig. 1** The column and Venn diagrams of DEGs assembled under low-temperature stress across three sets of comparisons expressed as  $4^{\circ}$ C vs  $23^{\circ}$ C (control),  $-20^{\circ}$ C vs  $23^{\circ}$ C and  $4^{\circ}$ C vs  $-20^{\circ}$ C, respectively (P < 0.05)

Zhuang et al. BMC Plant Biology (2021) 21:606 Page 3 of 3

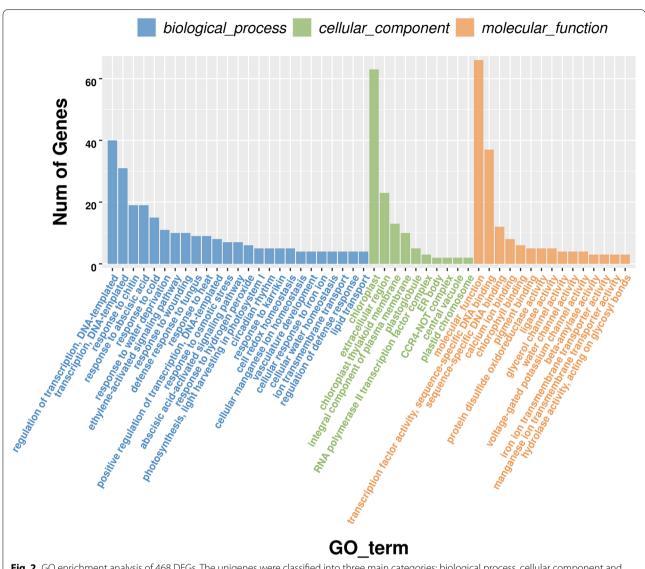


Fig. 2 GO enrichment analysis of 468 DEGs. The unigenes were classified into three main categories: biological process, cellular component and molecular function