

CORRECTION

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# Correction: Genome-wide and expression analysis of B-box gene family in pepper

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**Correction:** *BMC Genomics* 22, 883 (2021)  
<https://doi.org/10.1186/s12864-021-08186-w>

Following publication of the original article [1], it was reported that there was an error in Fig. 8C, which was a duplicate of Fig. 7D.

Furthermore, the description of Fig. 8C in the section “Expression analysis of *CaBBX* genes in response to exogenous hormones” was incorrect.

- The incorrect description was: “*CaBBX13* was also up-regulated under early treatment stage, but down-regulated under later treatment stage (Fig. 8C).”
- The correct description is: “Except for *CaBBX3* and *CaBBX5*, all the other four selected *CaBBX* genes were induced to expressed at a higher level by exogenous SA at different treatment stages (Fig. 8C).”

In the section “Expression analysis of *BBX* genes under abiotic stress in pepper” the text describing Fig. 7D was also inaccurate:

- The incorrect description was: “Additionally, *CaBBX13* showed an up-regulation at 3 and 6 h, but rapidly decreased with a more lower expression level than control at 12 h treatment (Fig. 7D).”
- The correct description is: “Additionally, *CaBBX13* showed an up-regulation at 6 h, but rapidly decreased with a more lower expression level than control at 12 h treatment (Fig. 7D).”

The correct Fig. 8C is given in this Correction article and the original article has been updated.

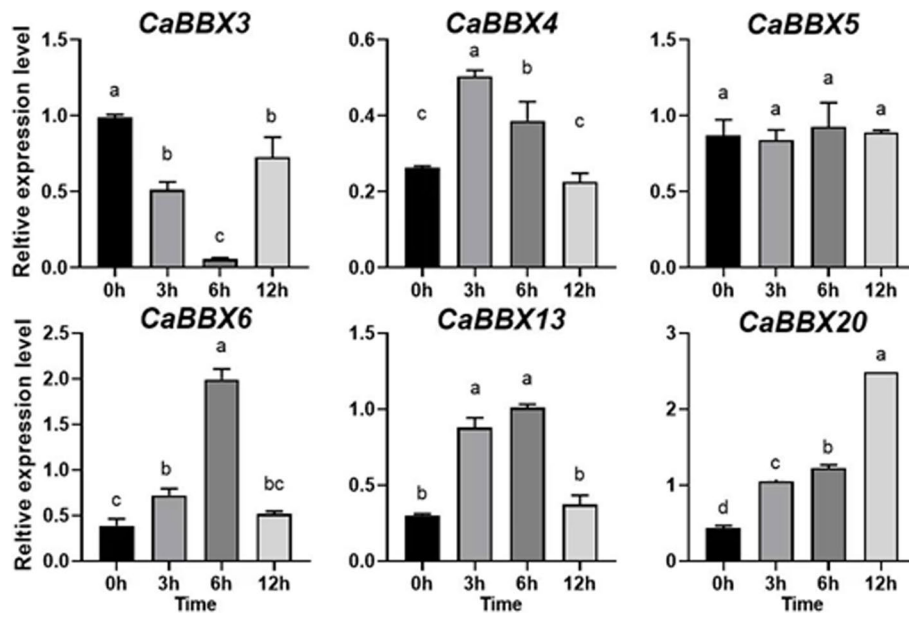
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The original article can be found online at <https://doi.org/10.1186/s12864-021-08186-w>.

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**Fig. 8 C** qRT-PCR transcript analysis of 6 selected *CaBBX* genes under SA treatment. Three independent biological experiments were performed ( $P < 0.05$ )

Published online: 18 December 2023

**Reference**

1. Ma J, Dai Jx, Liu Xw, et al. Genome-wide and expression analysis of B-box gene family in pepper. BMC Genomics. 2021;22:883. <https://doi.org/10.1186/s12864-021-08186-w>.