



## Correction to: Genome-wide characterization of the cellulose synthase gene superfamily in *Pyrus bretschneideri* and reveal its potential role in stone cell formation

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The above article was published online with the same Figures 7 and 8. The correct figure 8 is presented here. The Original Article has been corrected.

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The online version of the original article can be found at <https://doi.org/10.1007/s10142-020-00747-8>

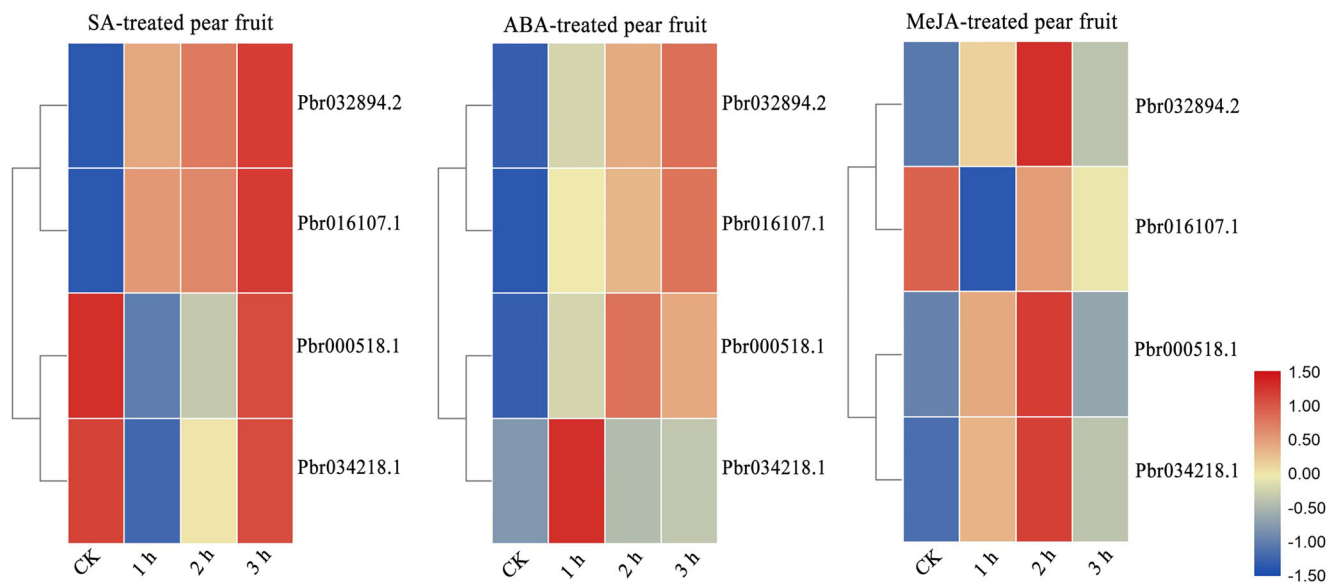
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**Fig. 8** Expression patterns of *PbCes/Csl* genes in response to different hormone treatments. The heatmap was generated by TBtools software using the *PbCes/Csl* genes expression data with hormonal treatments.

Log2-based-fold changes were used to create a heatmap. As shown in the bar at the lower right corner, the gene transcription level is expressed in different colors on the map. Data represent the mean of 3 replicates