ERRATUM

BMC Genomics





Erratum to: Chromosome doubling to overcome the chrysanthemum cross barrier based on insight from transcriptomic and proteomic analyses

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Erratum

Unfortunately, the original version of this article [1] contained an error. The Figure legends for Figs. 3, 4, 5 and 6 are incorrect. The correct version of Figs. 3, 4, 5 and 6 can be found below.

The correct match is as follows:

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Reference

China

 Zhang F, et al. Chromosome doubling to overcome the chrysanthemum cross barrier based on insight from transcriptomic and proteomic analyses. BMC Genomics. 2016;17:585. doi:10.1186/s12864-016-2939-0.

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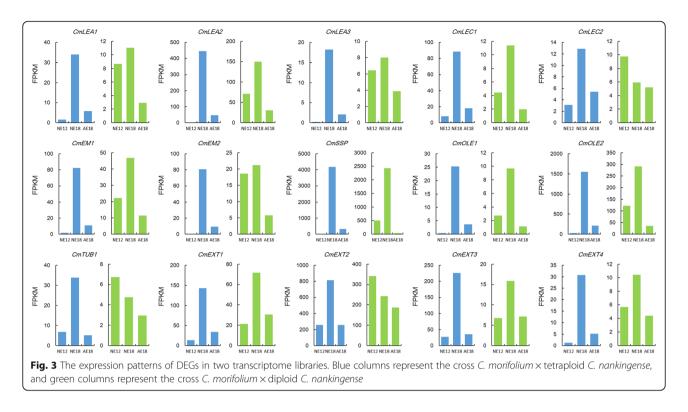
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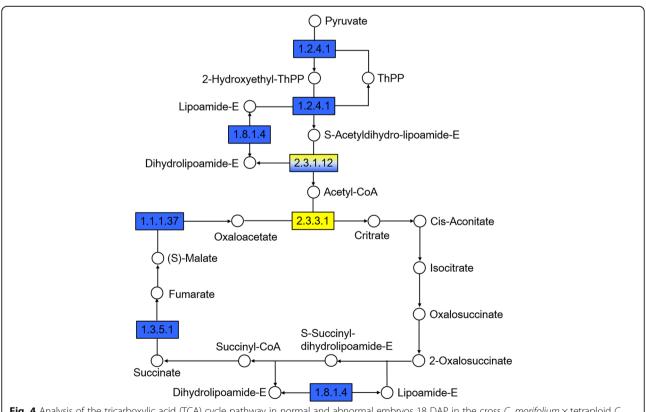


Fig. 4 Analysis of the tricarboxylic acid (TCA) cycle pathway in normal and abnormal embryos 18 DAP in the cross *C. morifolium* × tetraploid *C. nankingense*. The map displays selected steps from the KEGG pathway ko00020 'Citrate cycle (TCA cycle)'. Yellow indicates higher relative levels and blue indicates lower levels in AE18. Enzymes are given as EC numbers: 1.2.4.1, pyruvate dehydrogenase; 1.8.1.4, dihydrolipoamide dehydrogenase; 2.3.1.12, dihydrolipoamide acetyltransferase; 2.3.3.1, citrate synthase; 1.3.5.1, succinate dehydrogenase; and 1.1.1.37, malate dehydrogenase

