CORRECTION

Correction: Gene Expression Profiling of Development and Anthocyanin Accumulation in Kiwifruit (*Actinidia chinensis*) Based on Transcriptome Sequencing

Wenbin Li, Yifei Liu, Shaohua Zeng, Gong Xiao, Gan Wang, Ying Wang, Ming Peng, Hongwen Huang

The Excel file is absent from the published [S5 File](#). Please view the complete [S5 File](#) here.

S5 File. Supporting Tables. Table A. The expression profiles of all expressed genes in developing fruit of *A. chinensis* cv. 'Hongyang'. Table B. List of novel transcripts, alternative splicing and genes extended identified through analysis of the 'Hongyang' transcriptome. Table C. Verification of RNA-seq results by qPCR in *A. chinensis* cv. 'Hongyang'. Table D. Differentially expressed genes between two neighboring stages of fruit development in *A. chinensis* cv. 'Hongyang'. Table E. List of genes involved in ABA, CK, AUX and GA3 biosynthesis and signal transduction in developing fruit of *A. chinensis* cv. 'Hongyang'. Table F. List of genes involved in biosynthesis and metabolism of sugar and starch in developing fruit of *A. chinensis* cv. 'Hongyang'. Table G. List of genes involved in L-ascorbic acid biosynthesis and metabolism in developing fruit of *A. chinensis* cv. 'Hongyang'. Table H. List of genes involved in flavonoid biosynthesis and regulation in developing fruit of *A. chinensis* cv. 'Hongyang'. (XLS)

Reference

1. Li W, Liu Y, Zeng S, Xiao G, Wang G, Wang Y, et al. (2015) Gene Expression Profiling of Development and Anthocyanin Accumulation in Kiwifruit (*Actinidia chinensis*) Based on Transcriptome Sequencing. PLoS ONE 10(8): e0136439. doi:[10.1371/journal.pone.0136439](https://doi.org/10.1371/journal.pone.0136439) PMID: [26301713](https://pubmed.ncbi.nlm.nih.gov/26301713/)

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