CORRECTION

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

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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 PMID: 26301713