

## Corrigendum: Multi-Omics and miRNA Interaction Joint Analysis Highlight New Insights Into Anthocyanin Biosynthesis in Peanuts (*Arachis hypogaea* L.)

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## Keywords: peanut, anthocyanin, testa, multi-omics joint analysis, miRNA interaction, qRT-PCR

## A Corrigendum on

## Multi-Omics and miRNA Interaction Joint Analysis Highlight New Insights Into Anthocyanin Biosynthesis in Peanuts (*Arachis hypogaea* L.)

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In the original article, there were two mistakes in **Figure 7** and **Supplementary Figure 1** as published. The author mistakenly wrote "AhmiRNA398" as "AhmiRNA50" in **Figure 7** and the paternal parent "Zizhenzhu" had been wrongly written as "Z18-40" in **Supplementary Figure 1**, which caused a discrepancy with the original text and affected readers' understanding. The corrected **Figure 7** and **Supplementary Figure 1** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

Citation:

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preferred pathways, and light blue arrows indicate non-preferred pathways yet. The dashed rectangle indicates miRNAs-DEGs interaction. The green labels beside the arrows mean DEGs, the miRNAs corresponding to DEGs are marked in a gray ellipse, and both of them are connected by light gray arrows, and the regulation is indicated below the miRNAs. "o" with pink and purple means 30DAF of G110 and Z18-40, respectively. "D" with pink and purple that there is a positive, negative and none regulations plotween DEGs and miRNAs in the comparison on the left. The red cross indicates that

there are no regulation expression in the metabolites or genes.

Cinnamic acid

Ah4CL

Phenylalanine

AhPAL

