

**Figure S4.** Analysis of genes related to Auxin pathway. (a) A heatmap is presented to illustrate the transcriptional changes of genes in the flavonoid pathway between WT and transgenic calli (red represents upregulated and blue represents downregulated). (b) qPCR analysis of key structural genes. Data are means  $\pm$  SD (n = 3). \*\*P < 0.01.