



Figure S1. Heatmap of the accumulation of 10 superclasses of metabolites in HW, SW, and TZ

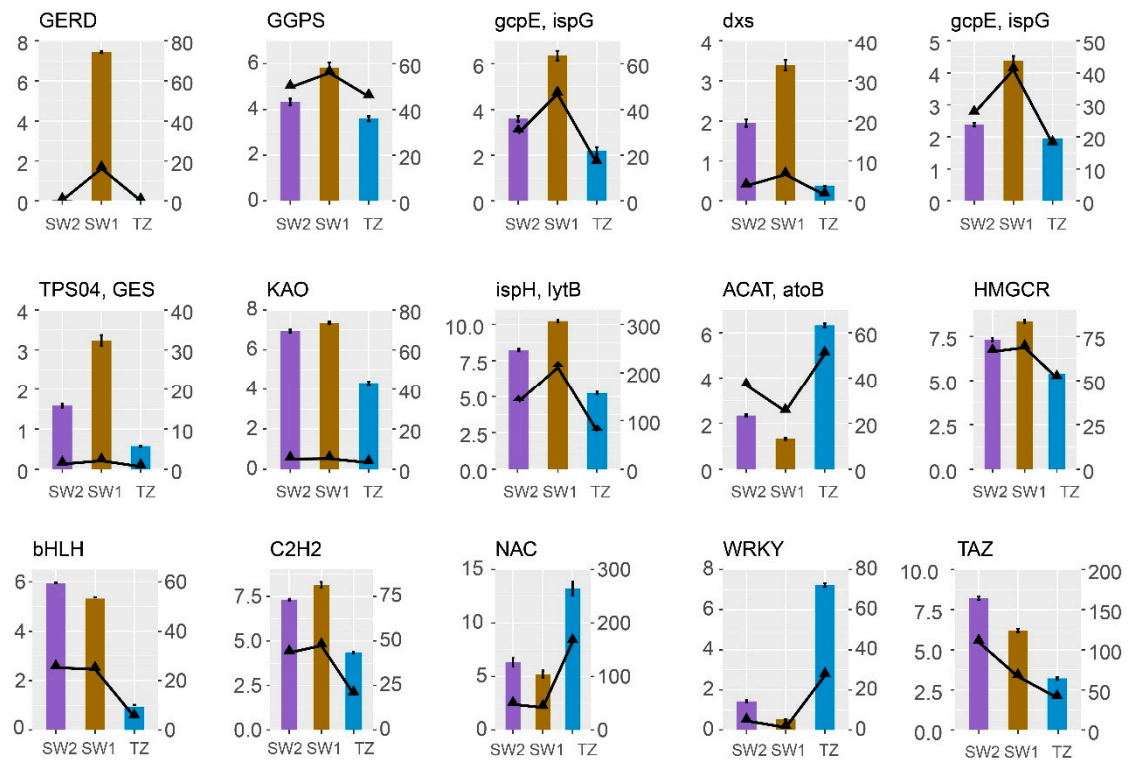


Figure S2. The RT-qPCR analysis of the relative level (left) and fragment per kilobase per million reads (FPKM) (right) of 10 enzyme genes and 5 transcription factors in SW1, SW2, and TZ.

The column showed the relative expression level, the line showed the FPKM value.