

Fig. S9

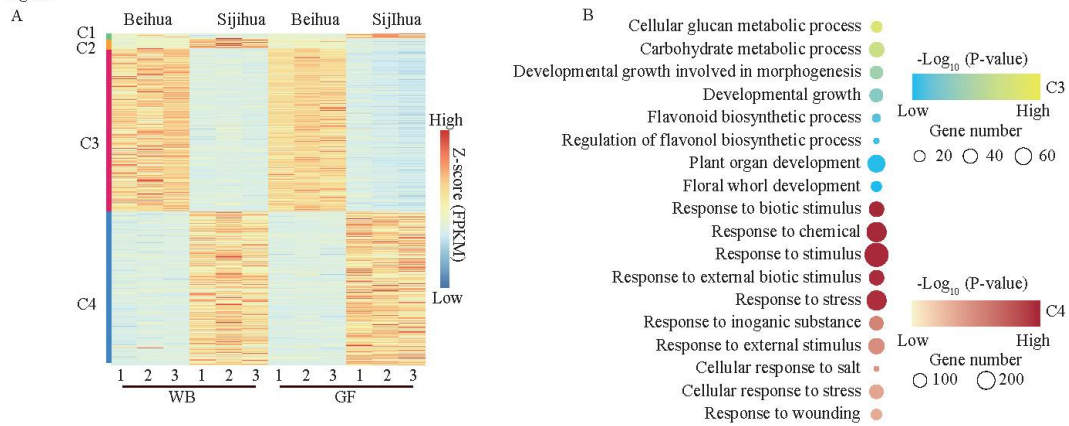


Figure S9. The pattern of differentially expressed genes related to SNP-associated DMCs. A, Heatmap showing the expression levels of the DEGs during the WB and GF stage in Beihua and Sijihua (There are four clusters of expression: C1: the expression levels of Beihua>Sijihua in WB stage and Beihua<Sijihua in GF stage, C2: the expression levels of Beihua<Sijihua in WB stage and Beihua>Sijihua in GF stage, C3: the expression levels of Beihua in WB stage and GF stage is higher than Sijihua, C4: the expression levels of Beihua in WB stage and GF stage is lower than Sijihua). **B,** GO enrichment analysis of C3 and C4 gene clusters.