

Fig. S11

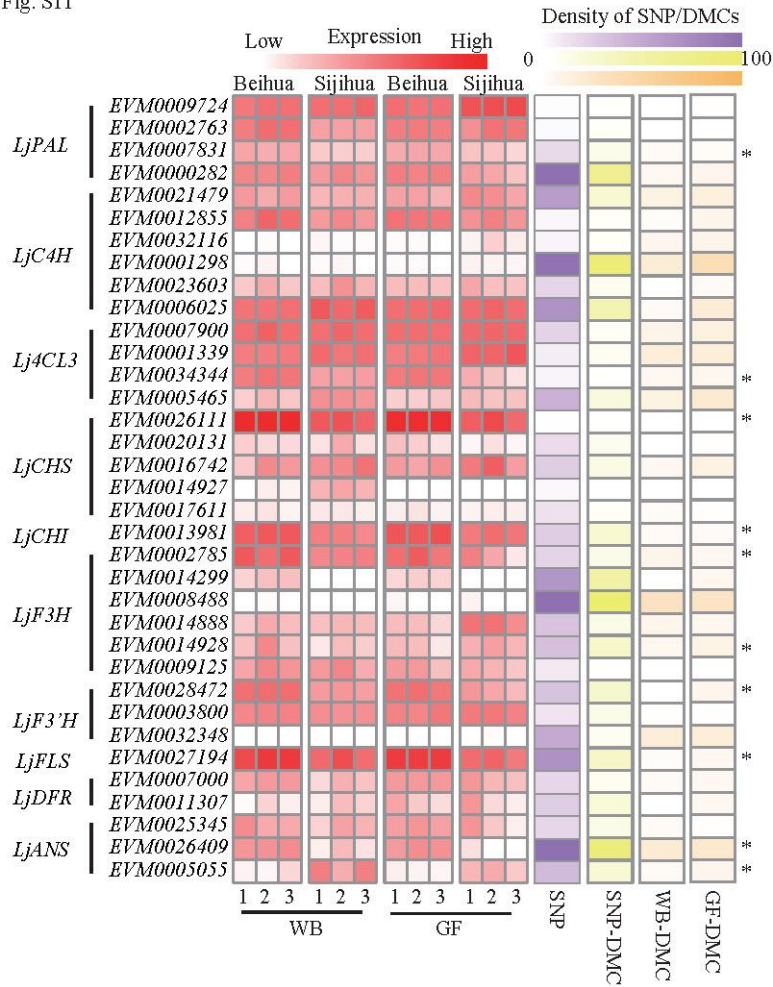


Figure S11. Heatmaps showing the expression levels of 35 key genes, the number of SNPs in the gene body of the genes, the number of SNP-associated DMCs, and the number of non-SNP-associated DMCs in Beihua and Sijihua during the WB and GF stages (* represents genes that showed significant differences in both stages).