Figure S1. Phylogenetic trees showing the relationships of soybean SPL-Like and Arabidopsis SPL3 and SPL9 proteins constructed using the neighbor-joining method in the CLUSTAL W program. The bootstrap percentage support is indicated at the branches of the tree. The locus IDs of these SPLs are presented in Table S2.

Figure S2. Phylogenetic relationships of soybean FUL-Like and Arabidopsis FUL proteins constructed using the neighbor-joining method in the CLUSTAL W program. The bootstrap percentage support is indicated at the branches of the tree. The locus IDs of these FULs are presented in Table S2.

Figure S3 Schematic diagram of GmFT5a promoter regions.