Supplementary Figure S4. Nucleotide sequence alignment of 5' upstream regions of BraGy genes. Nucleotide sequence alignment of 1.5 kb sequence upstream to putative translation start site (ATG) of BraaGy1 (Bra007741), BraaGy2 (Bra001894), BraaGy3 (Bra023782), BraaGy4 (Bra20117), BraaGy5 (Bra006568), and Arabidopsis Gy genes (AaAGG1, AaAGG2 and AaAGG3) was performed using Clustal W. The residues differing from consensus are marked in dark background.