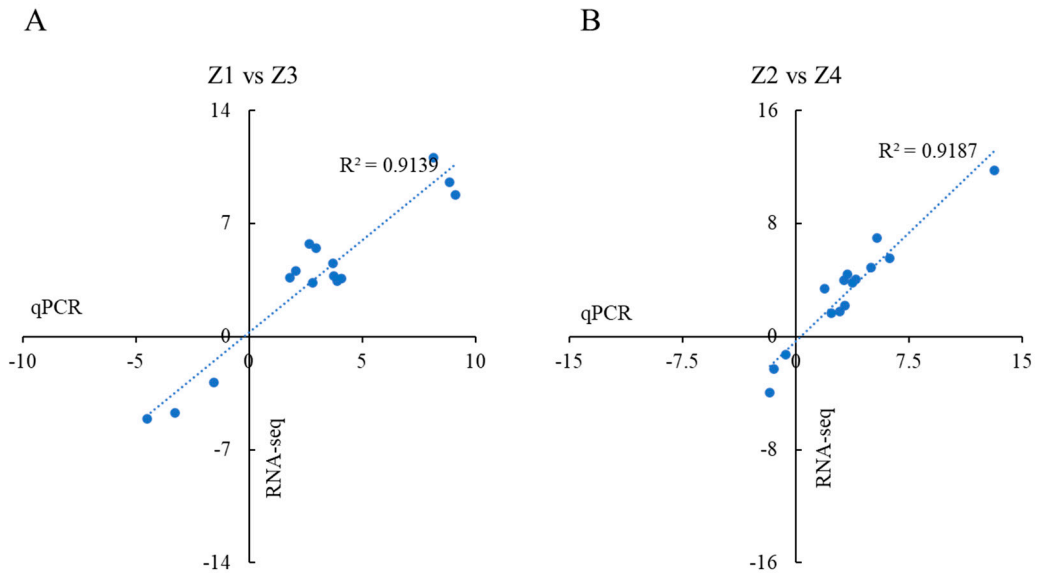

Figure S1. Correlation between qPCR and RNA-seq analyses of 15 DEGs in the pods (A) and the leaves (B) of SGS plants.

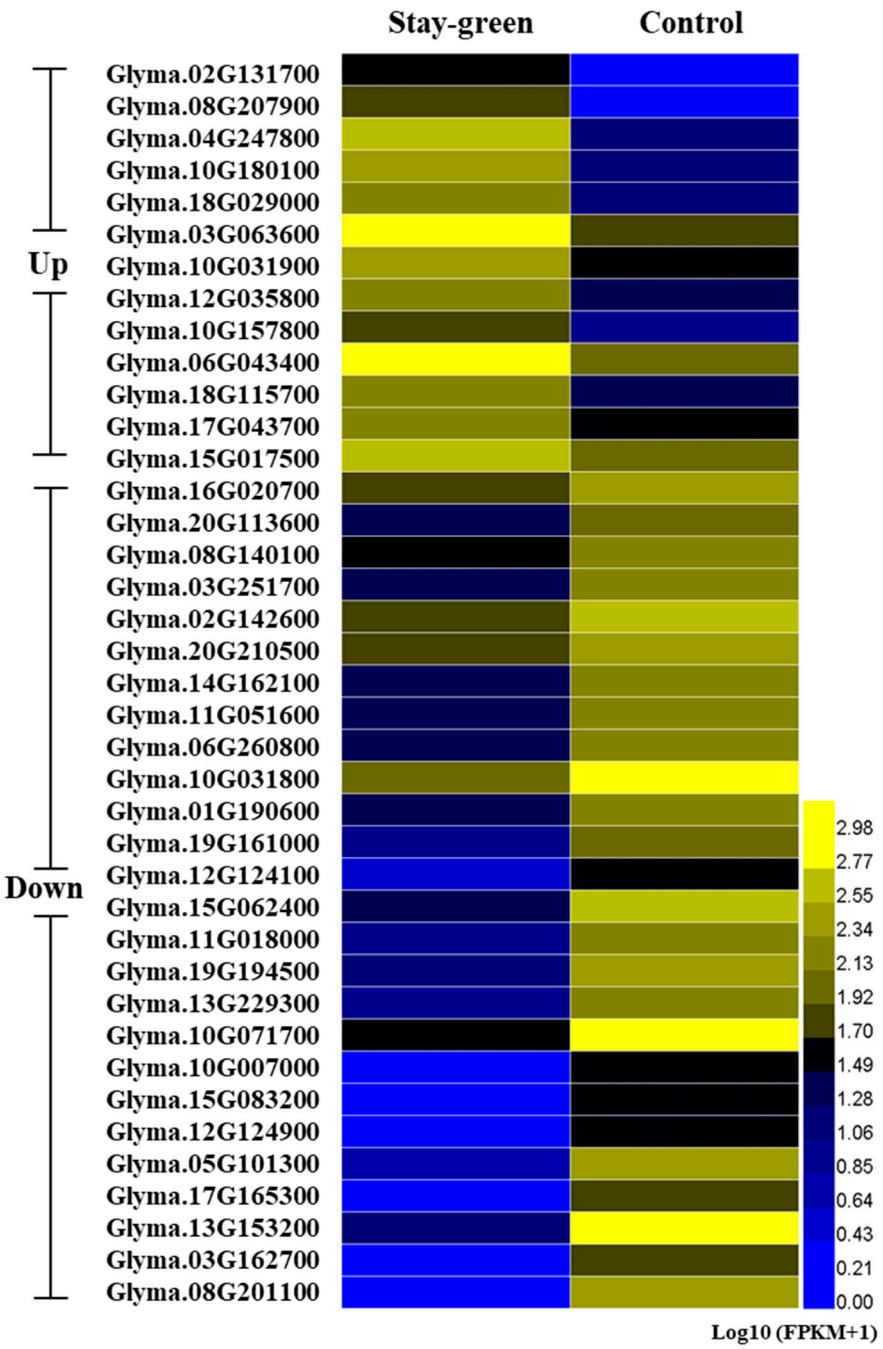

Figure S2. Heat map illustrating the expression of 39 DEGs involved in phytohormone signal transduction pathways.