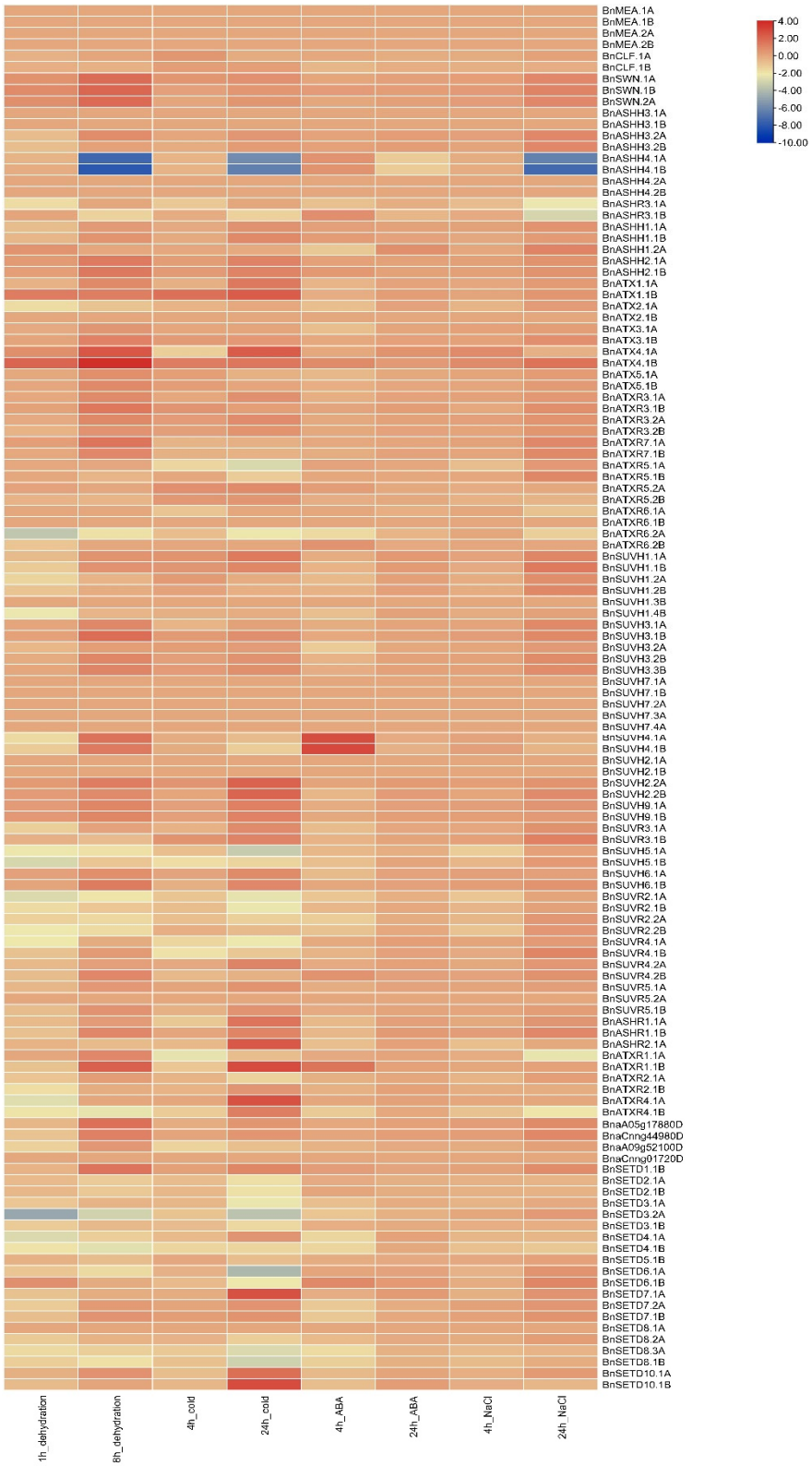


Figure S1. Heat map of in silico expression (Log2 fold change) of *BnSDG*s under dehydration, cold, ABA and salinity stress. The h reperesents the sample collection at specific hours after the treatments.

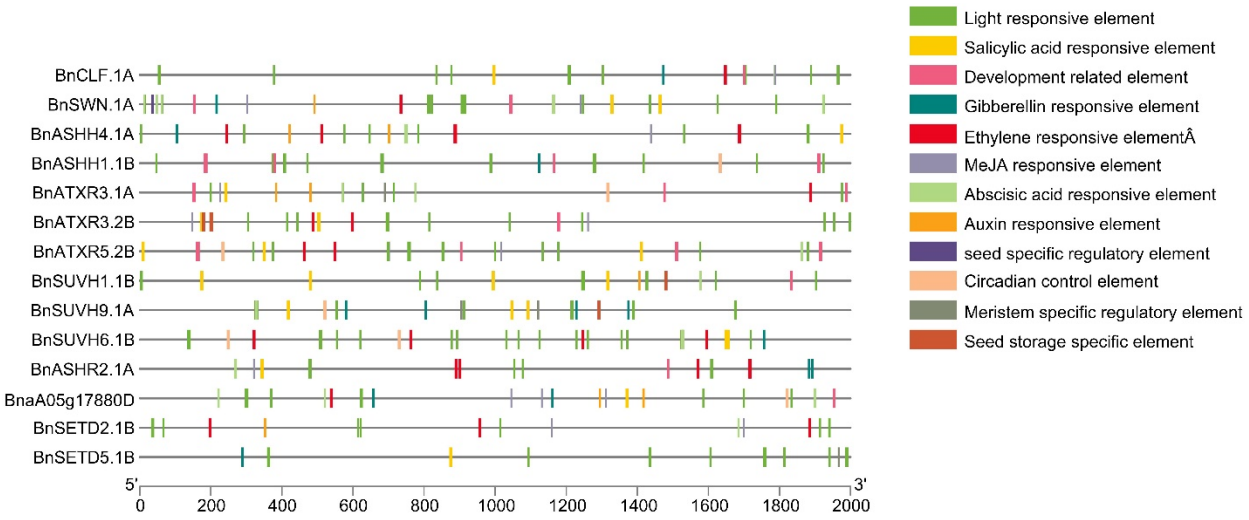


Figure S2. The potential cis-regulatory elements in the 2kb upstream coding region of 15 candidate *BnSDGs*.