



S6 Fig. Phylogenetic tree of NAC TFs from *B. distachyon*, Arabidopsis and rice. The deduced full-length amino acid sequences of 101 NAC TFs in *B. distachyon*, 99 in Arabidopsis and 132 in rice were used to construct an unrooted phylogenetic tree using MEGA5 by the neighbor-joining method with 1000 bootstrap iterations. Each NAC group was indicated in a specific color. Members of *B. distachyon*, Arabidopsis and rice were denoted by red circle, green triangle and blue diamond, respectively.