



Fig. S6. Transcriptomic data of *BnaCRKs*. **(A)** *BnaCRKs* expression profile in response to heat, drought, cold, freezing, salinity and flood. **(B)** *BnaCRKs* expression profile in response to *S. s* (*Sclerotinia sclerotiorum*) and clubroot. The expression data was normalized by Log2 method and the values are listed in Table S9.1. The color bar shows the expression level ranging from lower to higher.