

Figure. S2 Hierarchical clustering tree of the co-expression modules identified by WGCNA under UV-B radiation (A) and exogenous ABA (B) treatment. Different modules are marked by different colors. Each leaf of the cluster tree represents a gene. The darker the color in the corresponding heat map, the higher the correlation between the gene in the module and the treatment.

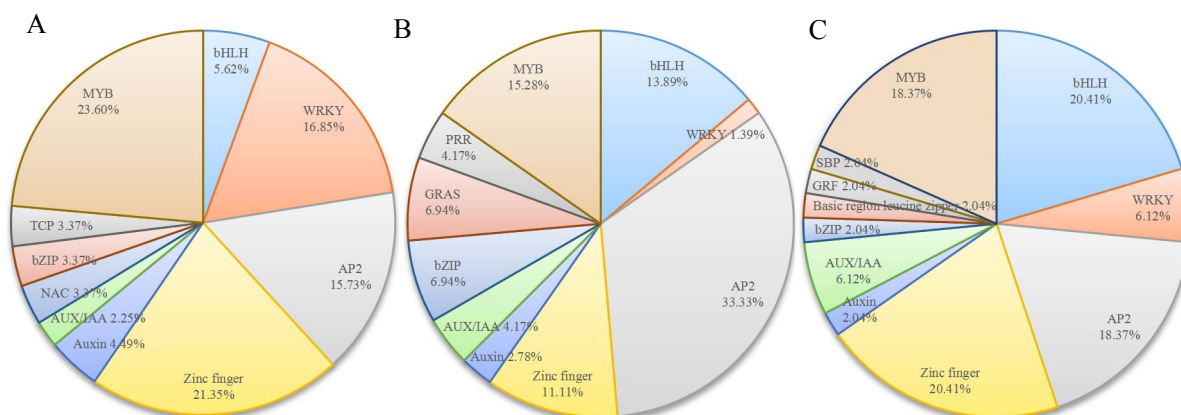


Figure. S3 Transcription factor family clusters under UV-B radiation or exogenous ABA treatment according to WGCNA. **A** The blue module shows transcription factor family clusters under UV-B radiation. **B** The turquoise module shows transcription factor family clusters under UV-B radiation. **C** The blue module shows transcription factor family clusters under exogenous ABA treatment.