

Figure S1: Gene structure and conserved motif analysis of 180 *HbbHLH* genes.

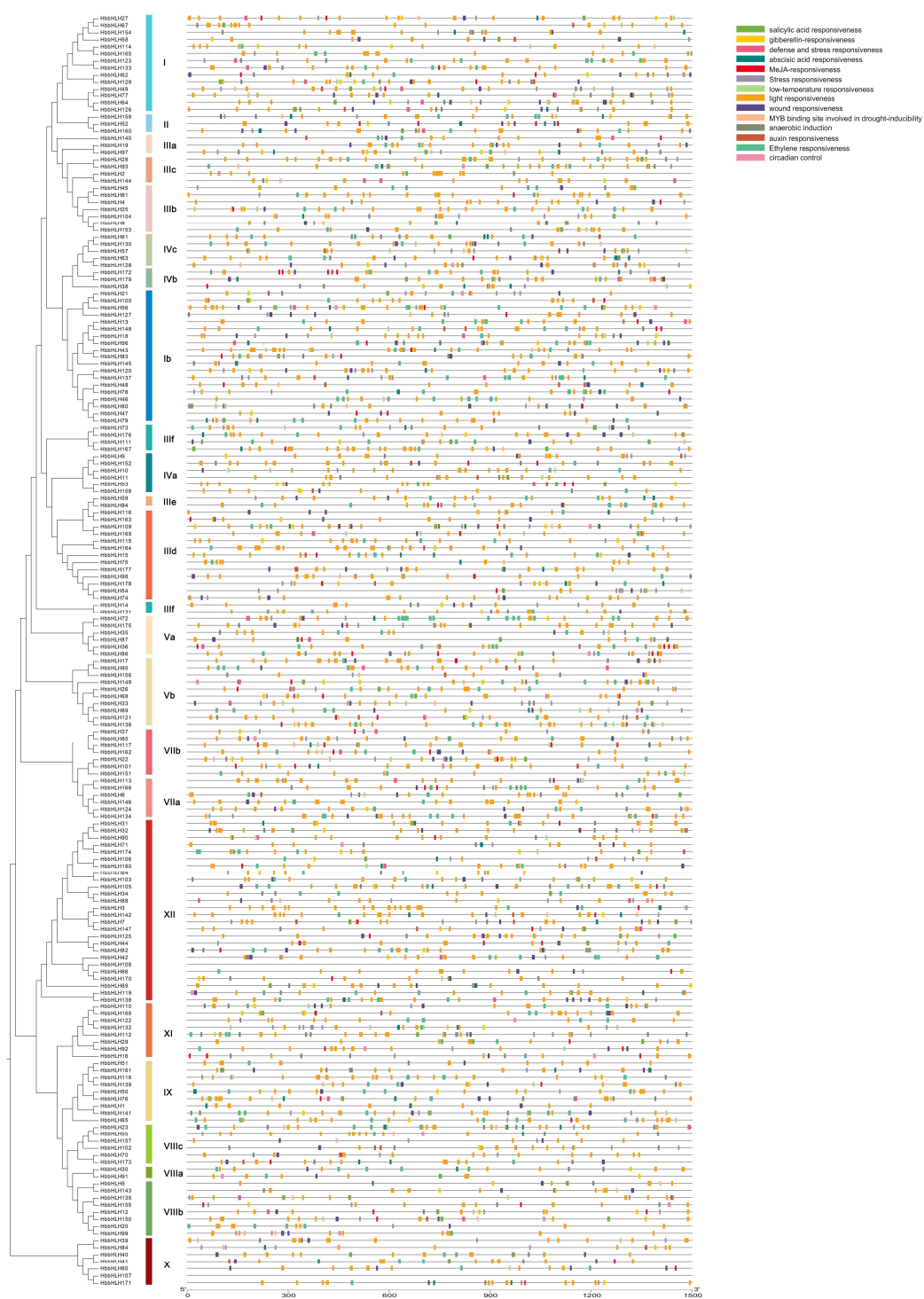


Figure S2: Analysis of cis-regulatory elements in the promoter regions of 180 *HbbHLH* genes.