

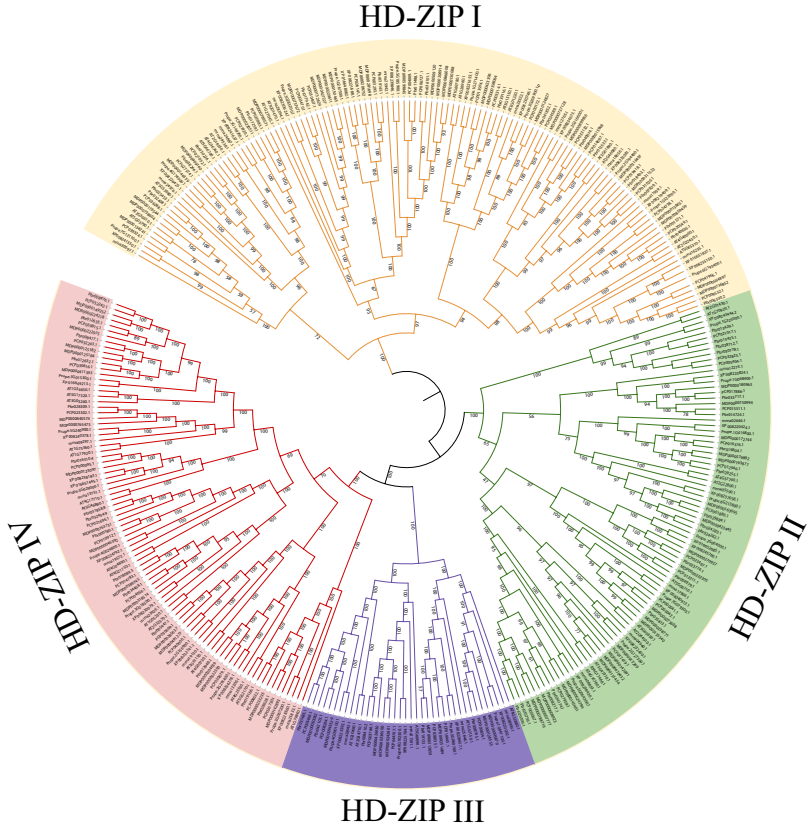


Fig. S2 Phylogenetic analysis of HD-ZIPs from 6 Rosaceae species and Arabidopsis. The tree was constructed with IQ-TREE v.1.6 software using the neighbor-joining method with 1000 bootstrap replicates. The PbHBs were clustered into 4 distinct clades, marked by curves of different colors.