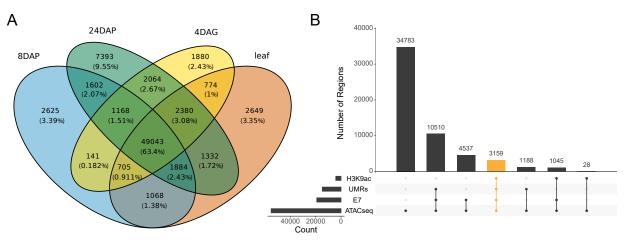
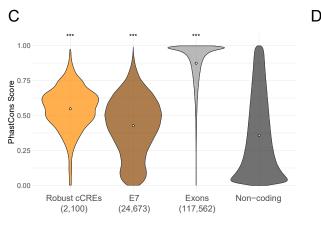
Figure 2





)	24DAP robust CRE motif	e-value	Transcription factor family
	* * * ********************************	2.3e-35	ERF
	a CTTAC	2.3e-35	GT; WRKY
	Section Service	7.2e-31	HD-ZIP; M-type MADS
	* III. TGCA-GTAGTACGTAG	7.2e-31	bZIP
	g of the state of	8.9e-07	B3 domain-containing; LFL1-like