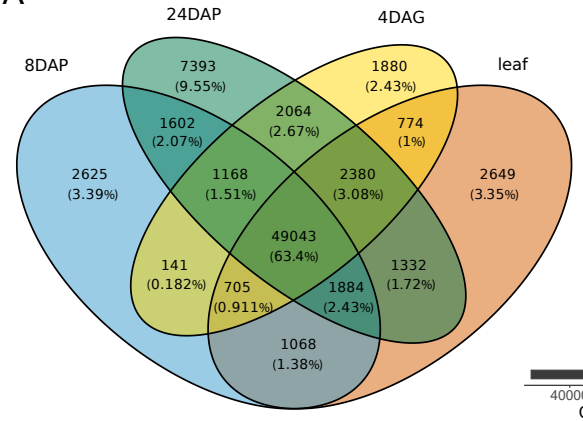
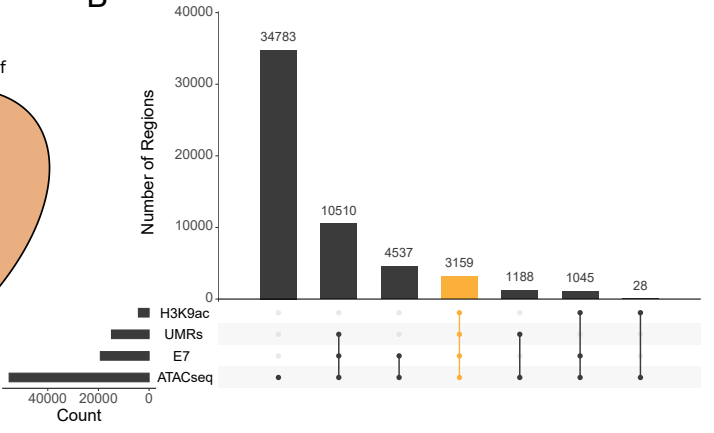
Figure 2

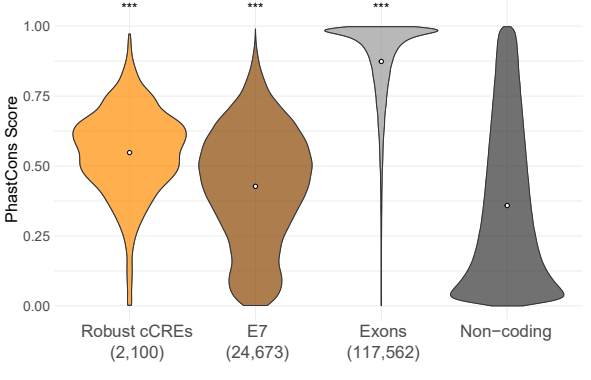
A



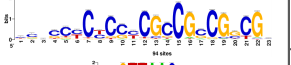
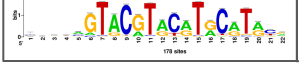
B



C



D

24DAP robust CRE motif	e-value	Transcription factor family
	2.3e-35	ERF
	2.3e-35	GT; WRKY
	7.2e-31	HD-ZIP; M-type MADS
	7.2e-31	bZIP
	8.9e-07	B3 domain-containing; LFL1-like