

Figure 1

A

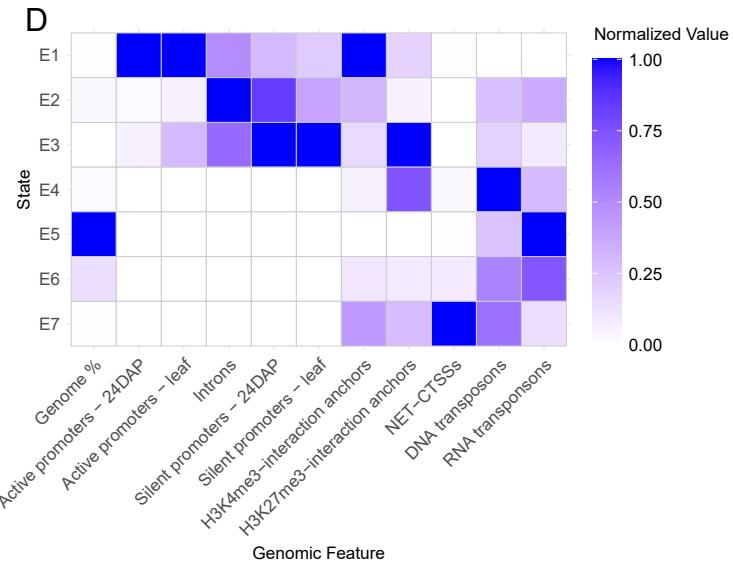
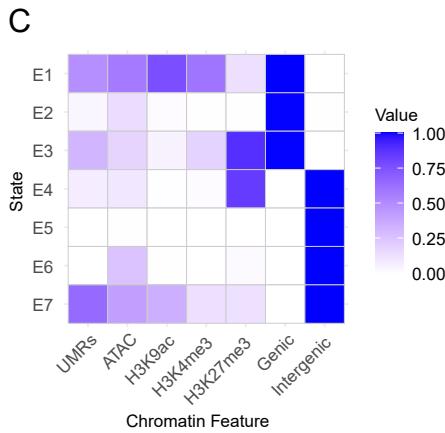
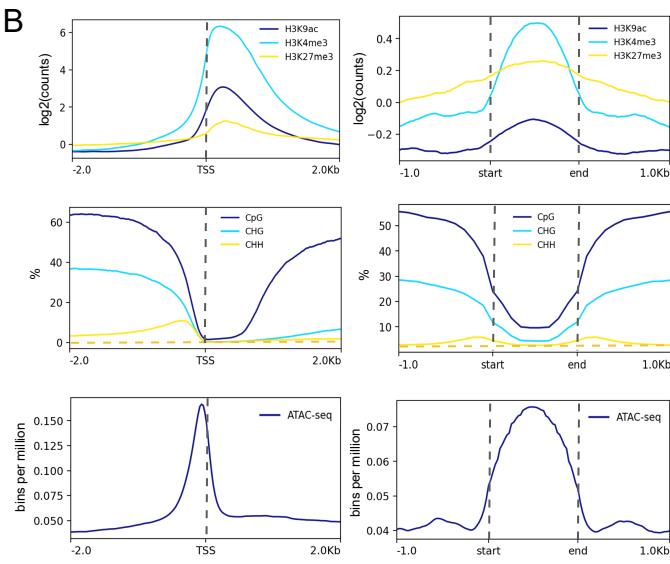
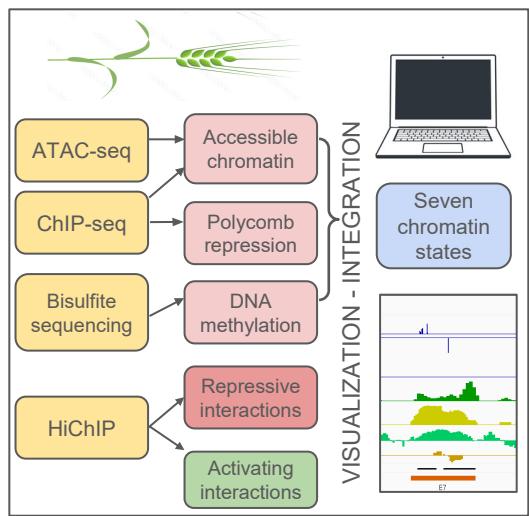
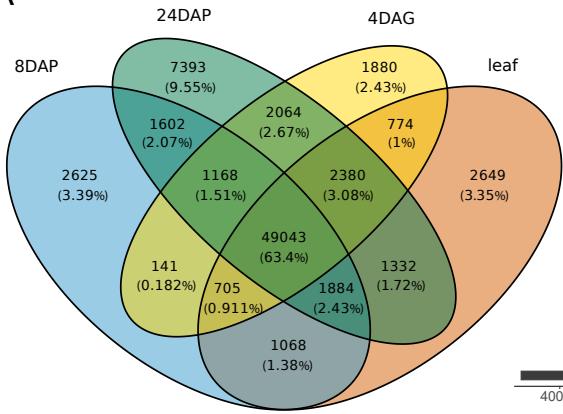
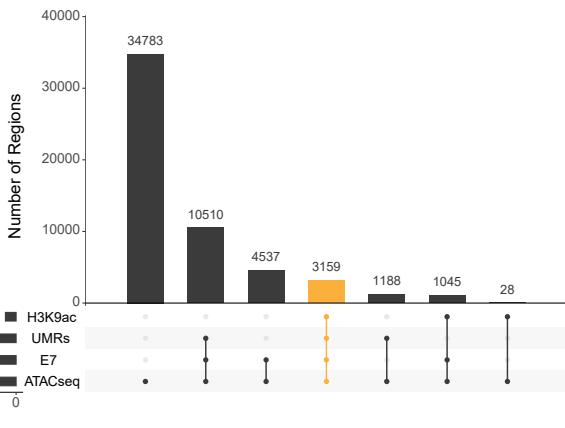


Figure 2

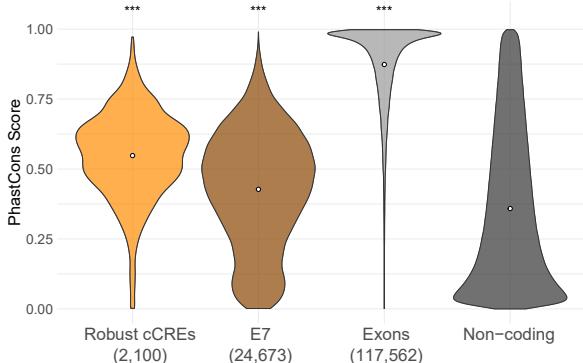
A



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

Figure 3

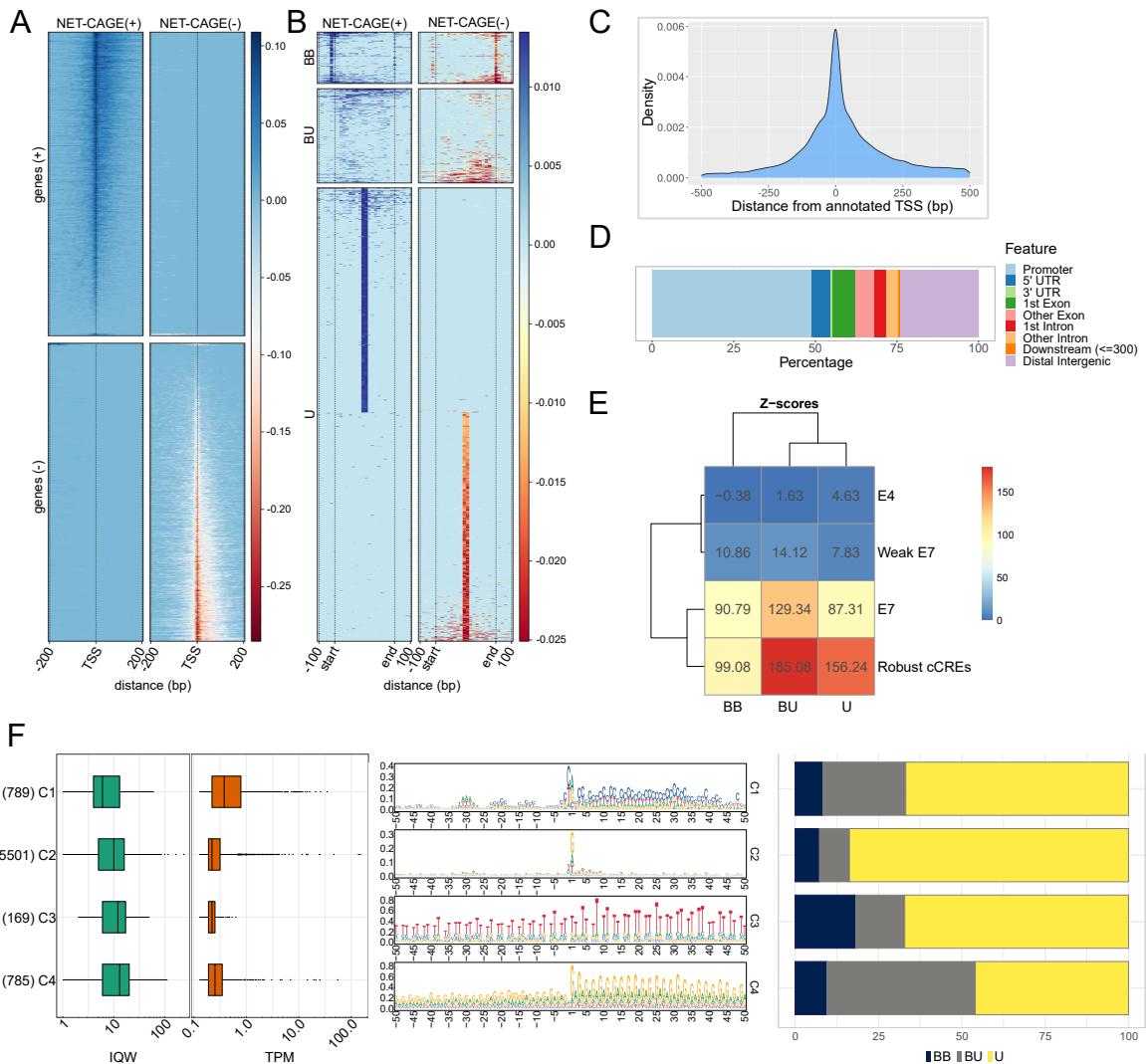
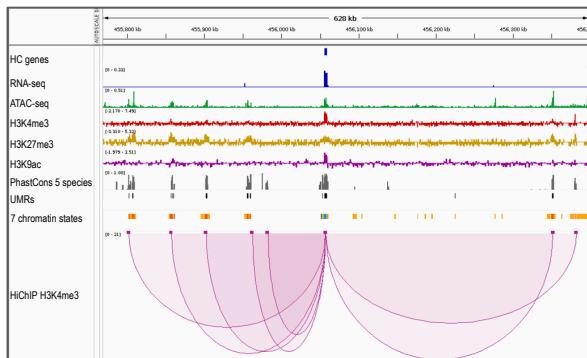
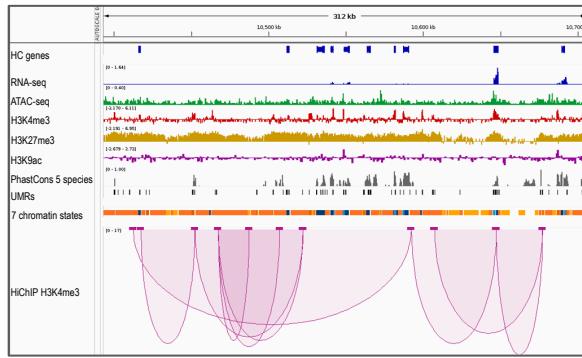


Figure 4

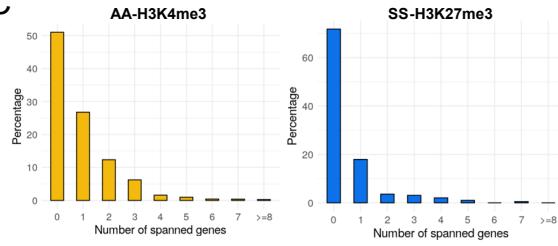
A



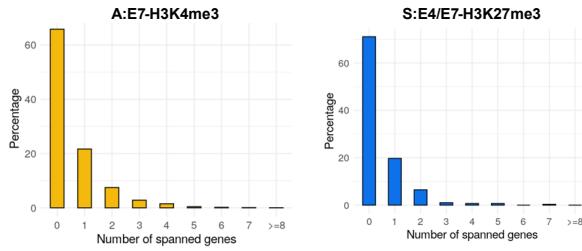
B



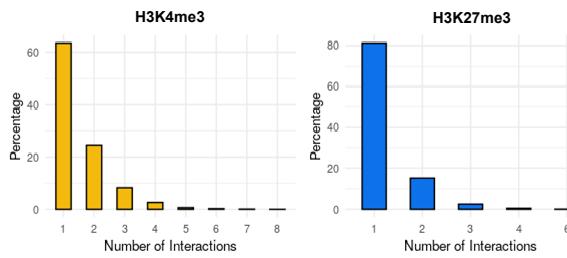
C



A:E7-H3K4me3



D



E

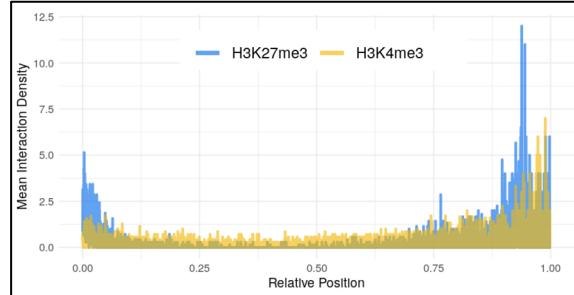
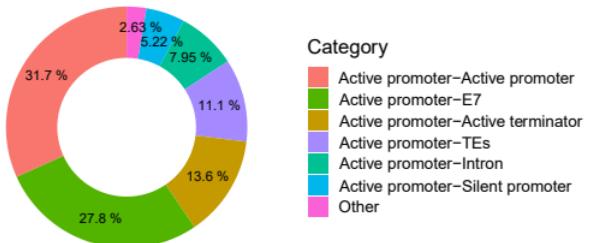
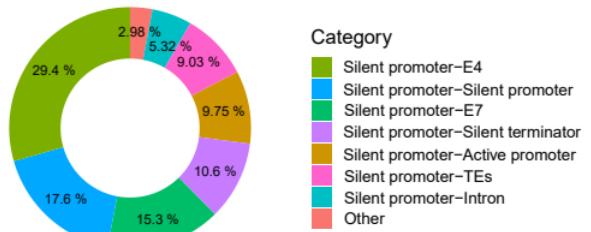


Figure 5

A



B



C

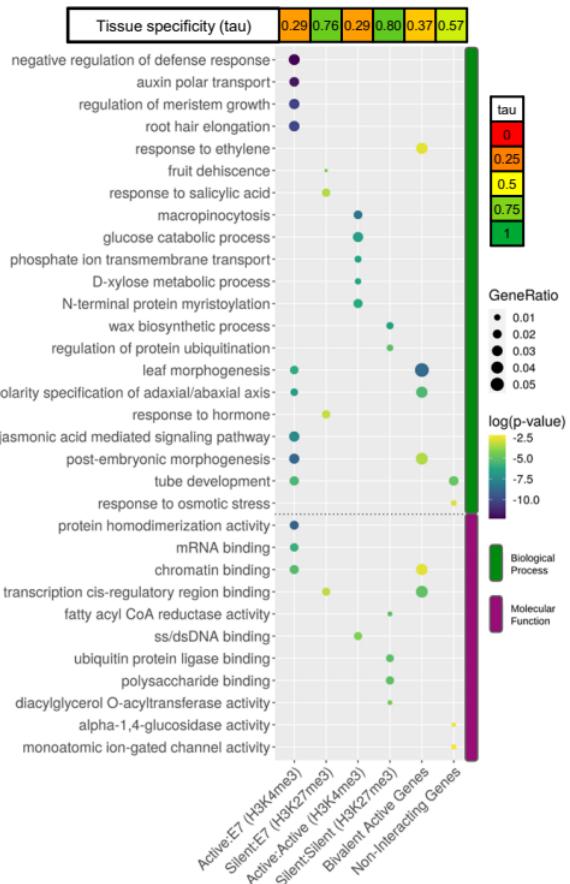
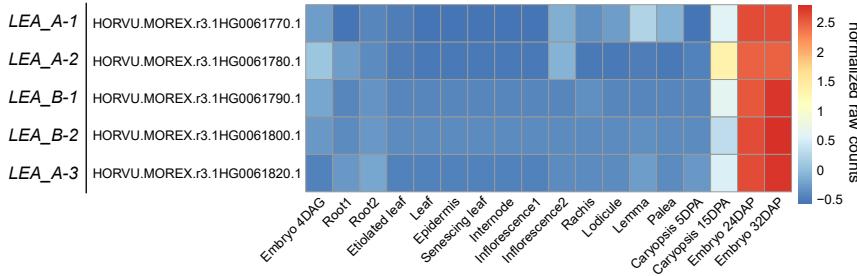
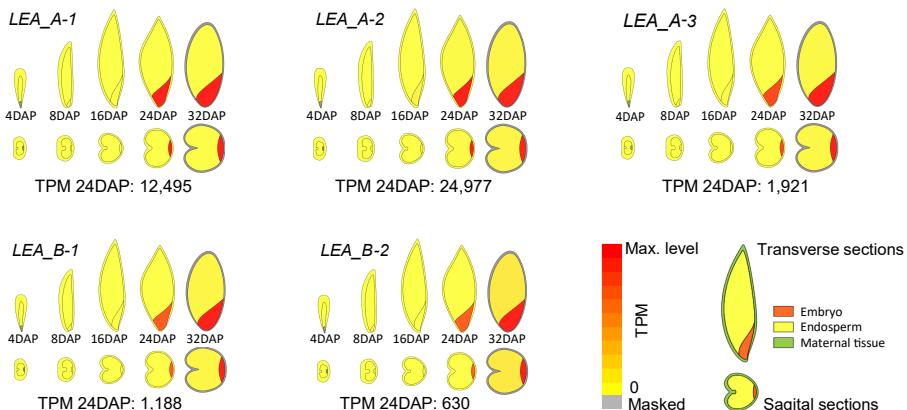


Figure 6

A



B



C

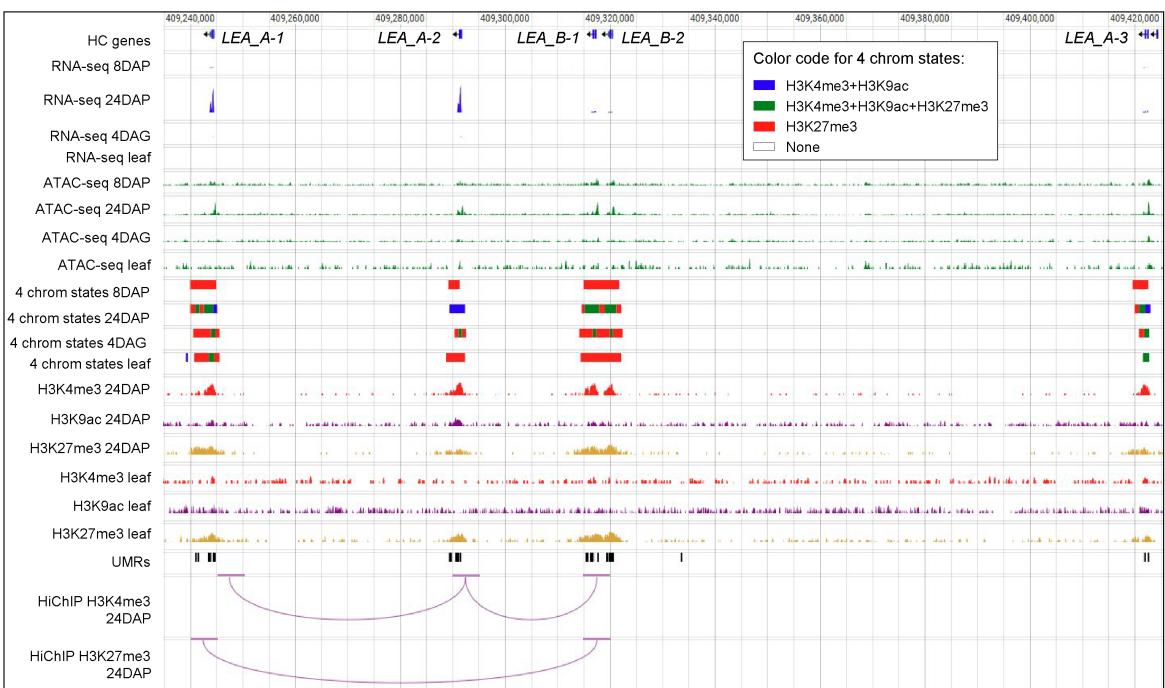
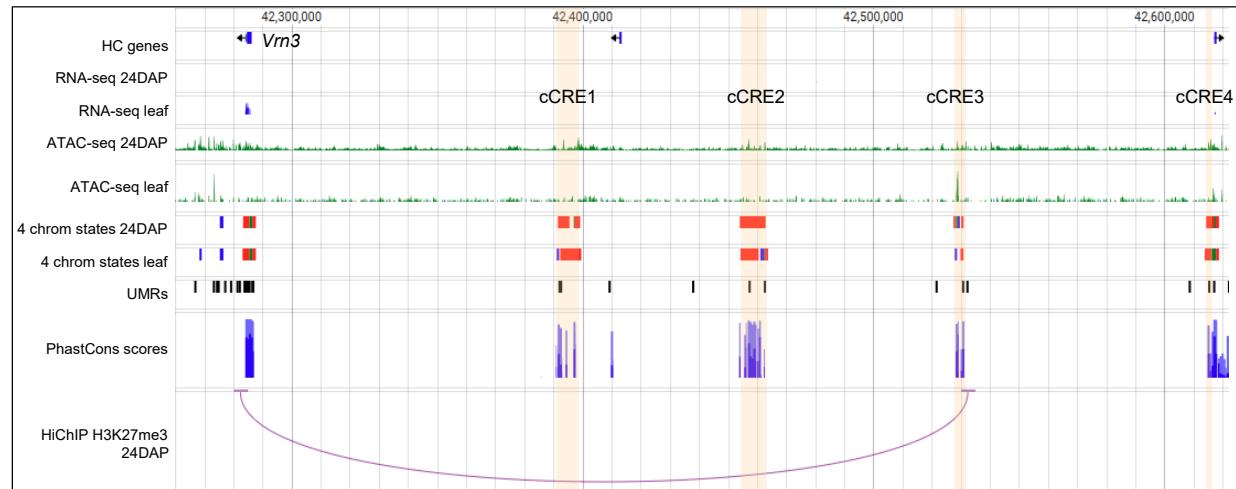
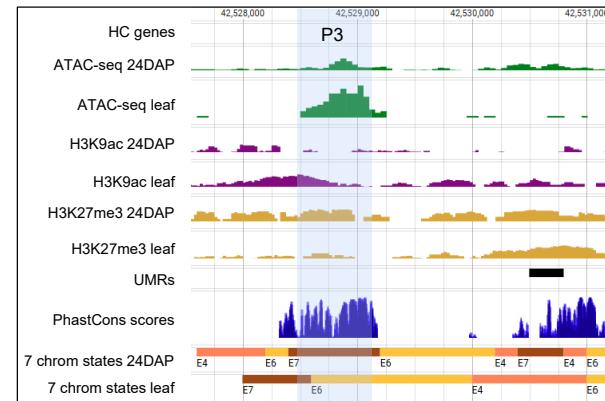


Figure 7

A



B



C

