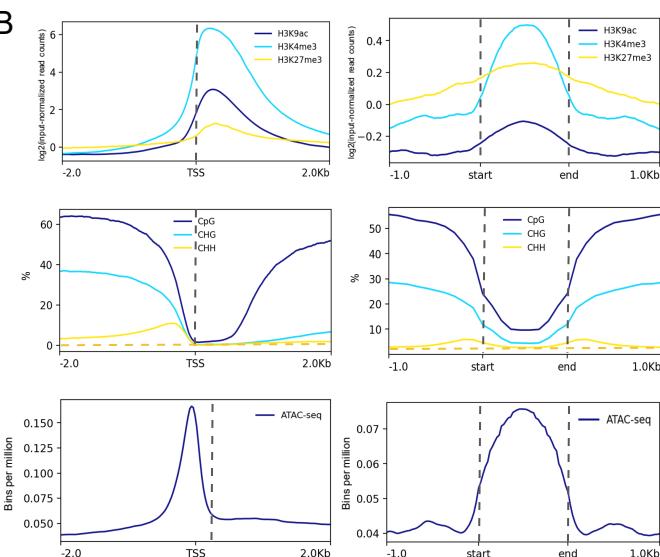
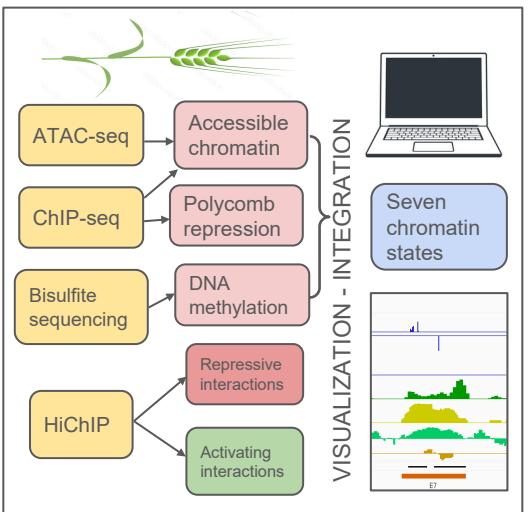


Figure 1

A



C

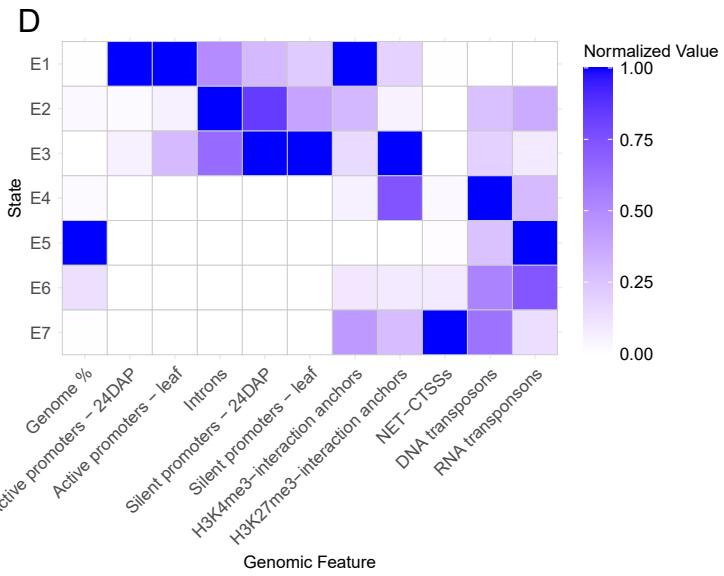
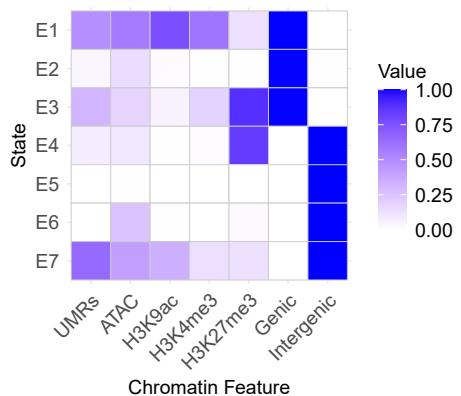
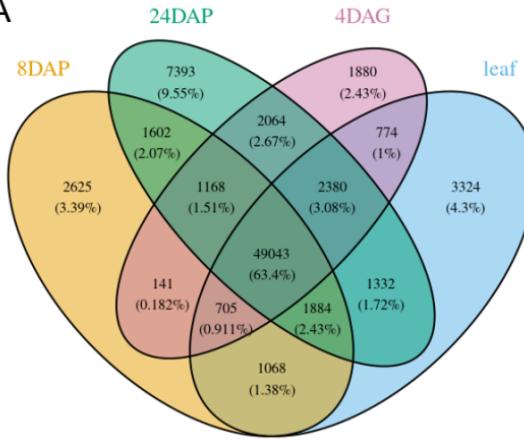
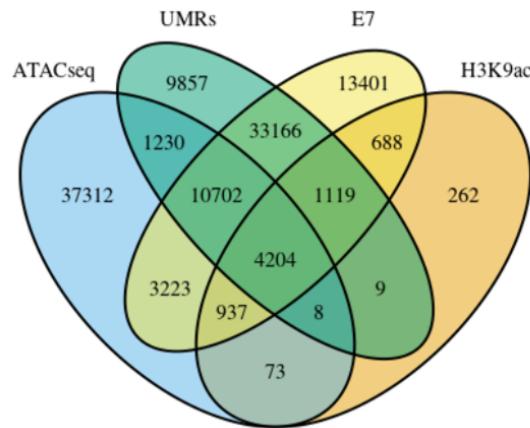


Figure 2

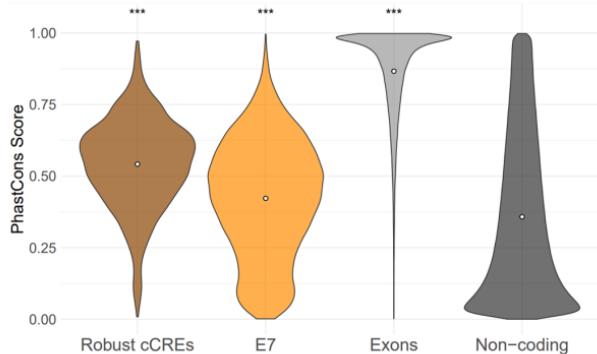
A



B



C



D

24DAP robust CRE motif	e-value	Transcription factor
	2.3e-35	ERF family protein
	2.3e-35	Trihelix, GT-family protein; WRKY family protein
	7.2e-31	HD-ZIP family protein; M-type_MADS family protein
	7.2e-31	Basic leucine zipper factors (bZIP)
	8.9e-07	B3 domain-containing protein LFL1-like

Figure 3

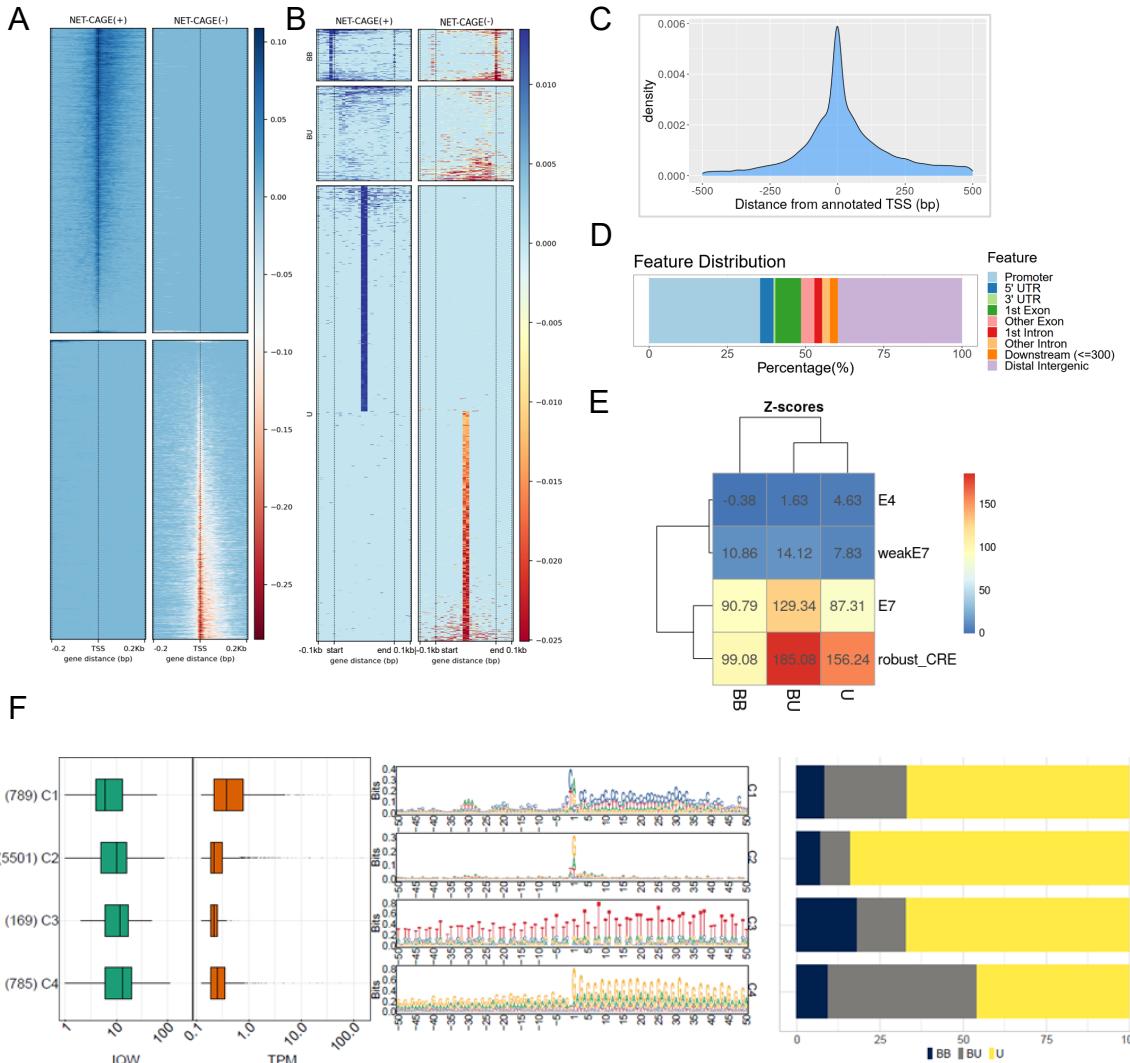
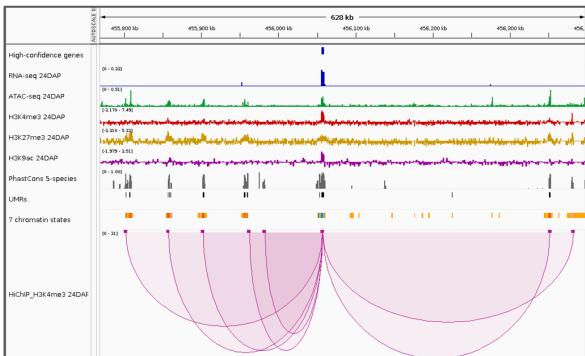
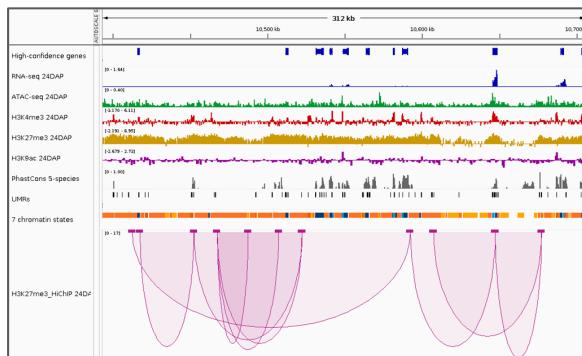


Figure 4

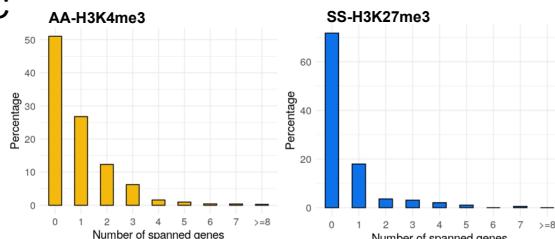
A



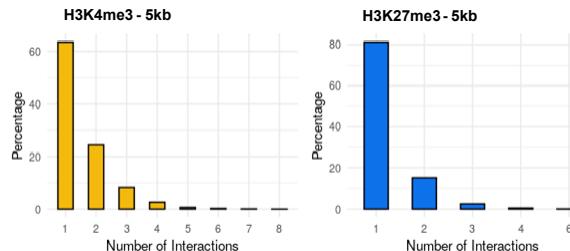
B



C



D



E

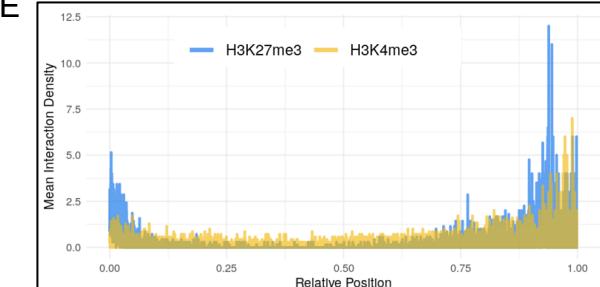
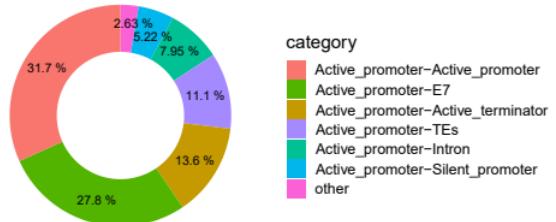


Figure 5

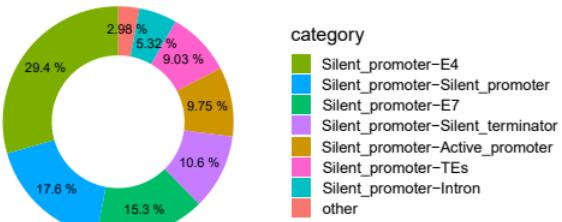
A

Interaction Classes



B

Interaction Classes



C

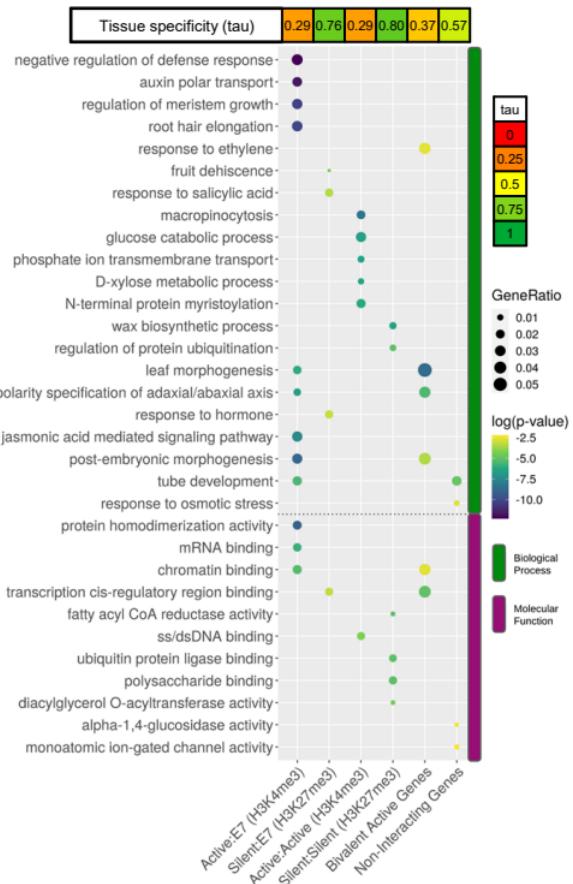
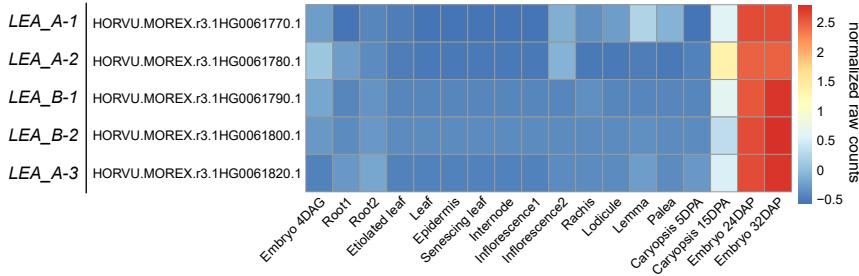
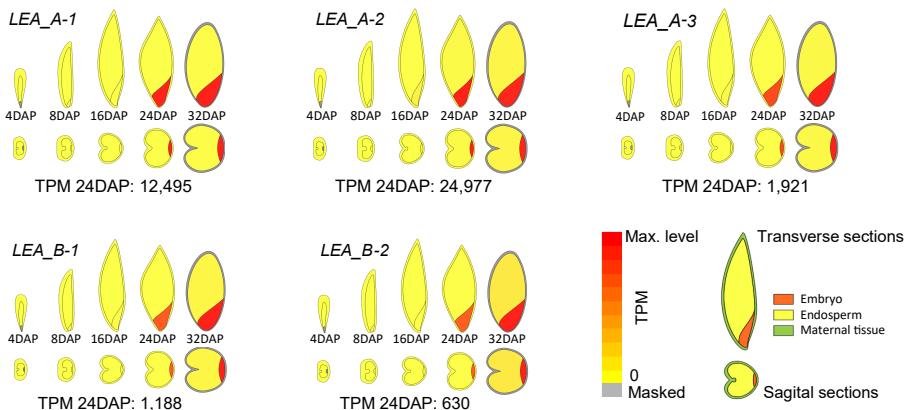


Figure 6

A



B



C

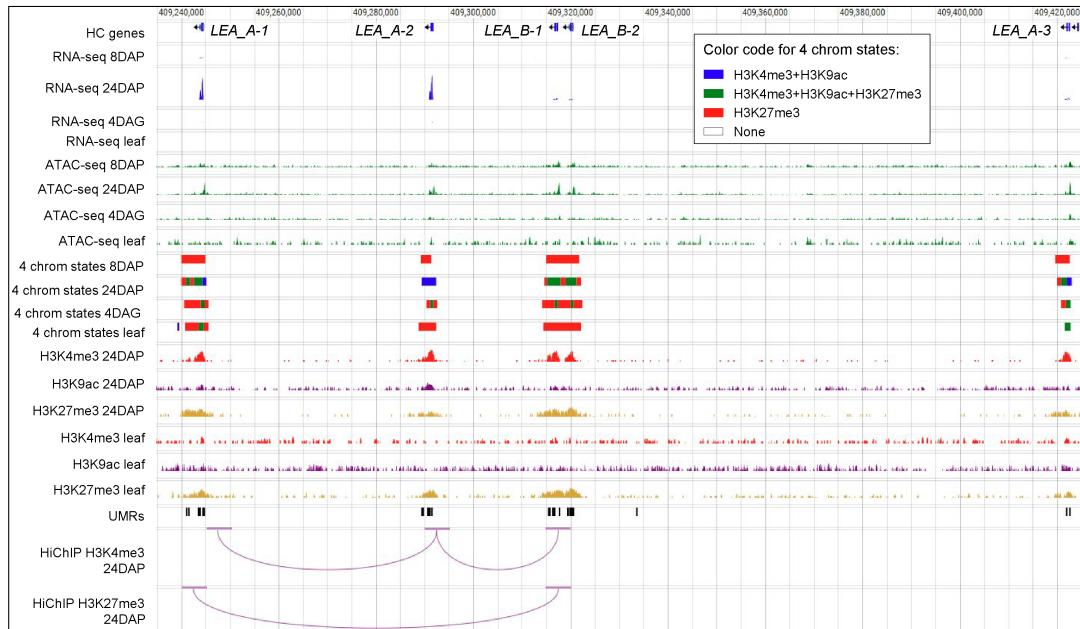
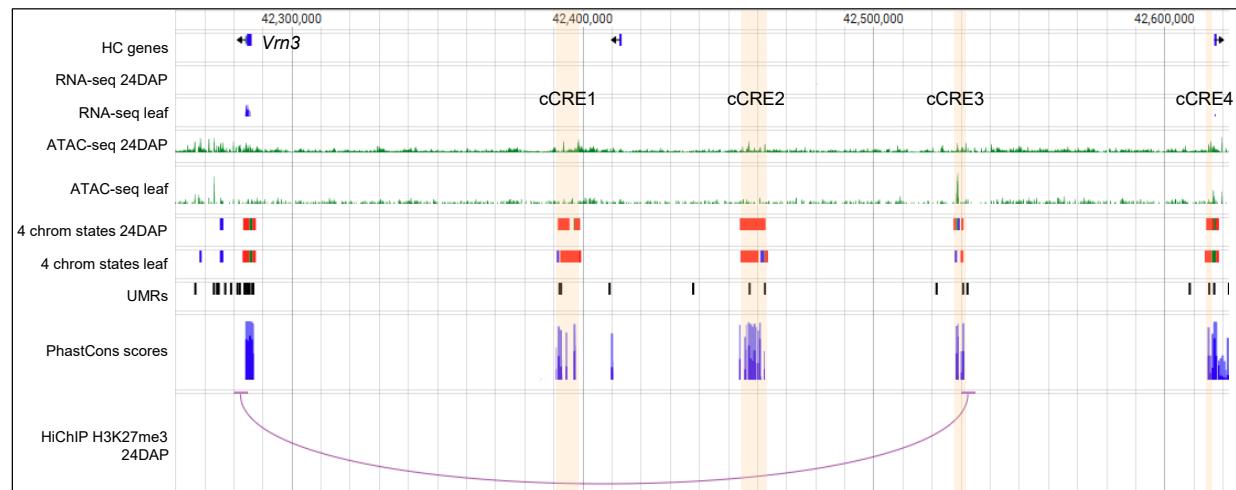
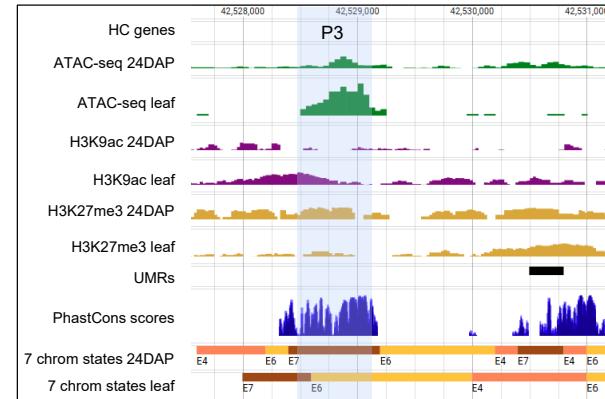


Figure 7

A



B



C

