

Number of consensus motif Hits (2kb upstream)		HSF	HMB	E2F	HMB/WRKY	2β 3α DBD	HMB(STaR)	HMB	bZIP	HSF
		TAGAA	TAGAATTCCAGAA	TTCCCCGC	CCACTTG	GCCGACACA	GTAATGATT	ATCATTA	CACGTG	CGCC
	Complete genome	43609	1	1730	8812	347	652	13390	13997	43477
Tolerant	Up regulated	2628	0	97	556	18	35	793	1037	2641
	Down regulated	2624	0	92	630	22	27	853	904	2603
Susceptible	Up regulated	3672	0	148	744	24	39	1072	1273	3666
	Down regulated	2333	0	88	573	22	29	790	882	2347