

















































	3-AF3 binding site		CBFHV		MBS
	AAGAA motif		CGCG-box		MNF1
	ACE		CGTCA/TGACG-motif		MRE
	AC-II		Chs-CMA1a		O2-site
	ABRE		circadian		P-box
	ACTTTA motif		E-box		Py-rich region
	AE-box		ERE		RAA-motif
	AG-box		Eri box3		RY-element
	ARE		GAG-motif		Skn-1
	AS-2-box		GA-motif		Sp1
	ATCT-motif		GARE-motif		TCA-element
	AT1-motif		GATA-box		TC-rich repeats
	AT-rich element		G-box		TCT-motif
	AU-RR core		GCN4-motif		TGA-element
	Box 1		GT1-box		T/G box
	Box 4		HSE		W1-box
	CAT-box		L-box		WUN