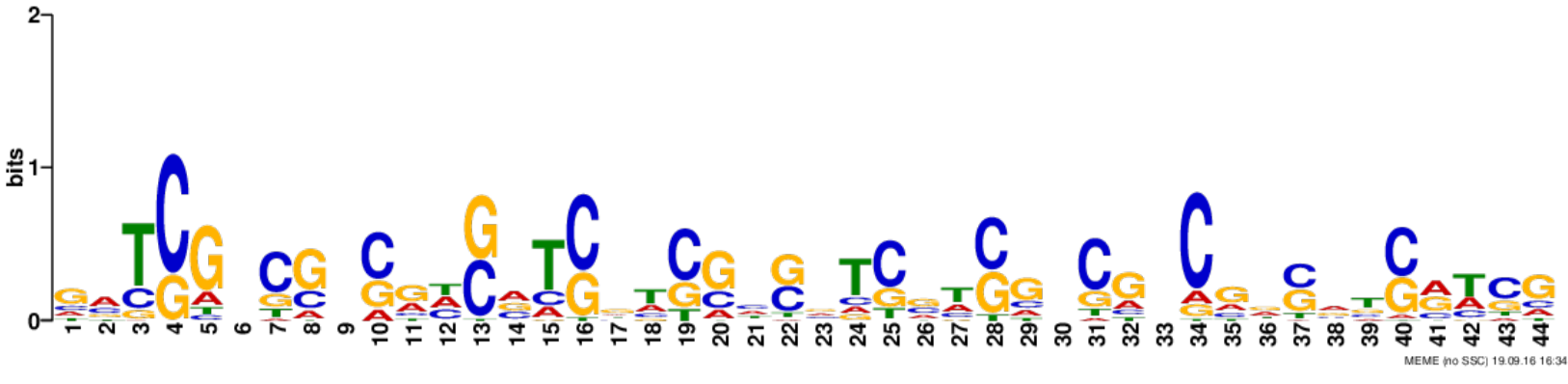
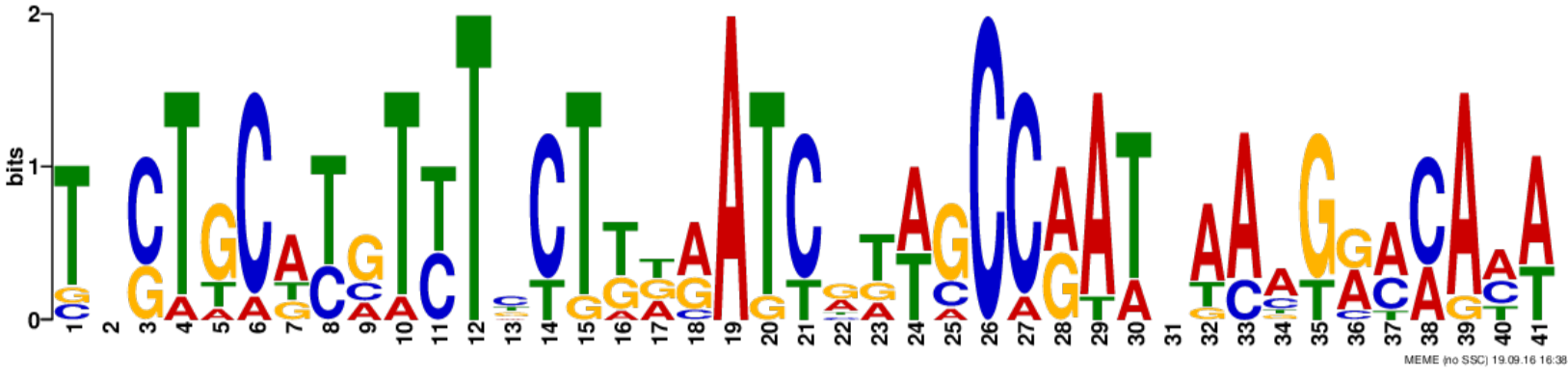
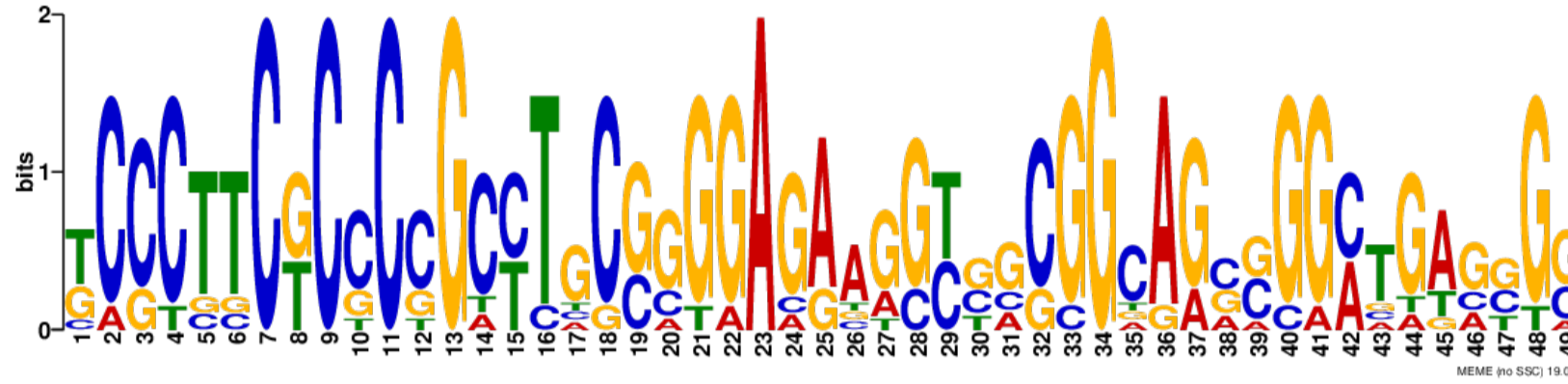
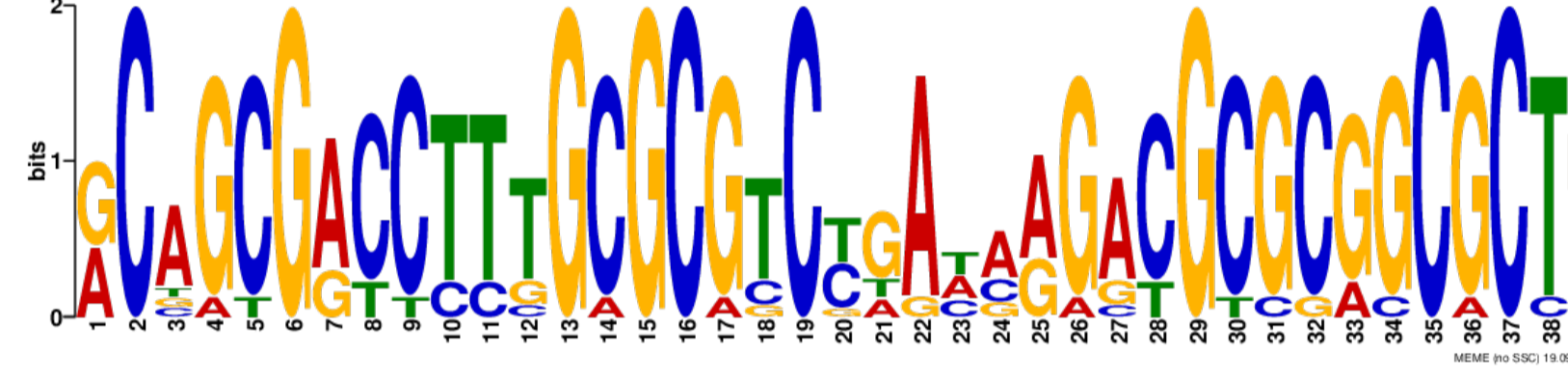
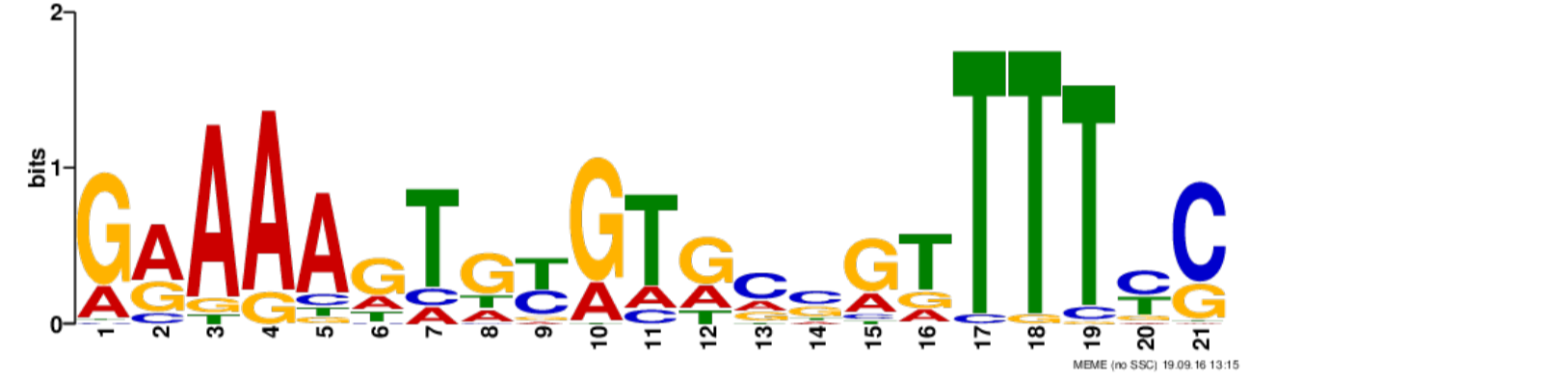
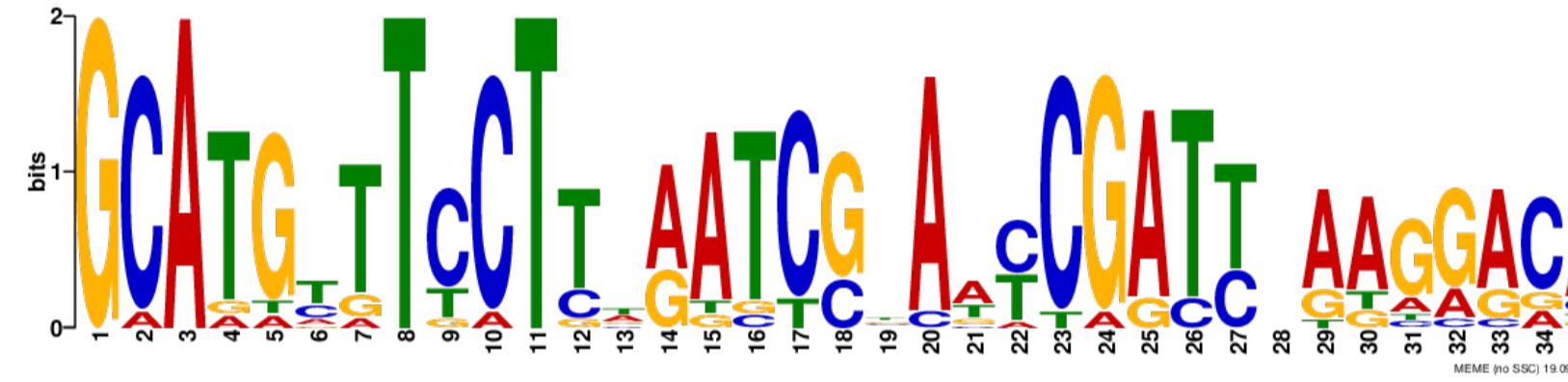
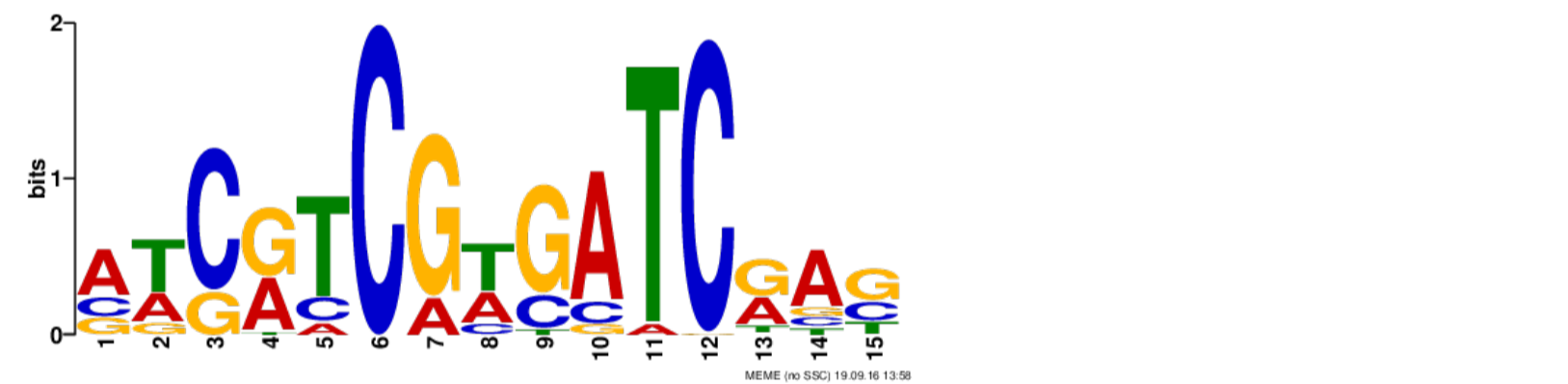
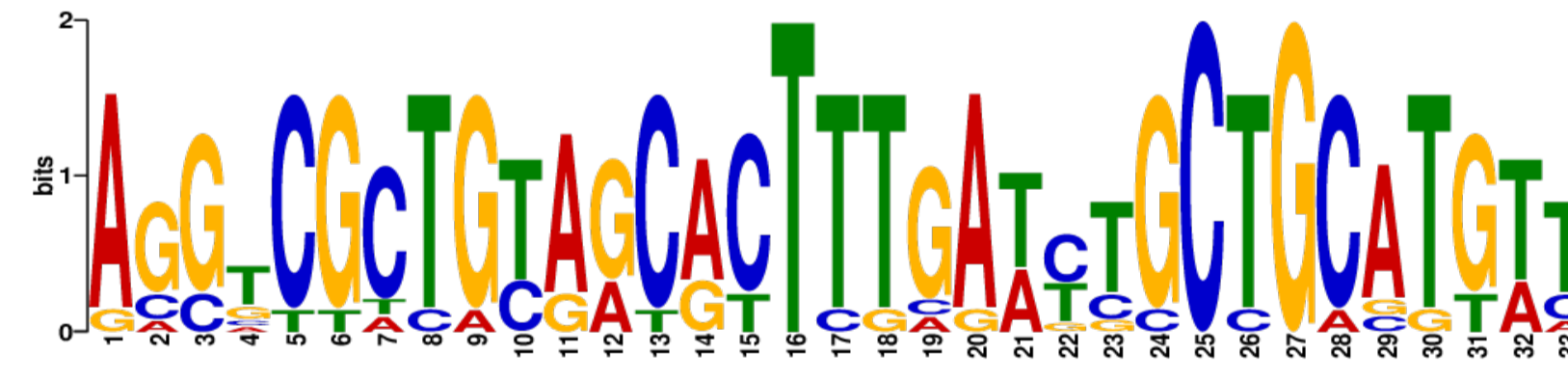
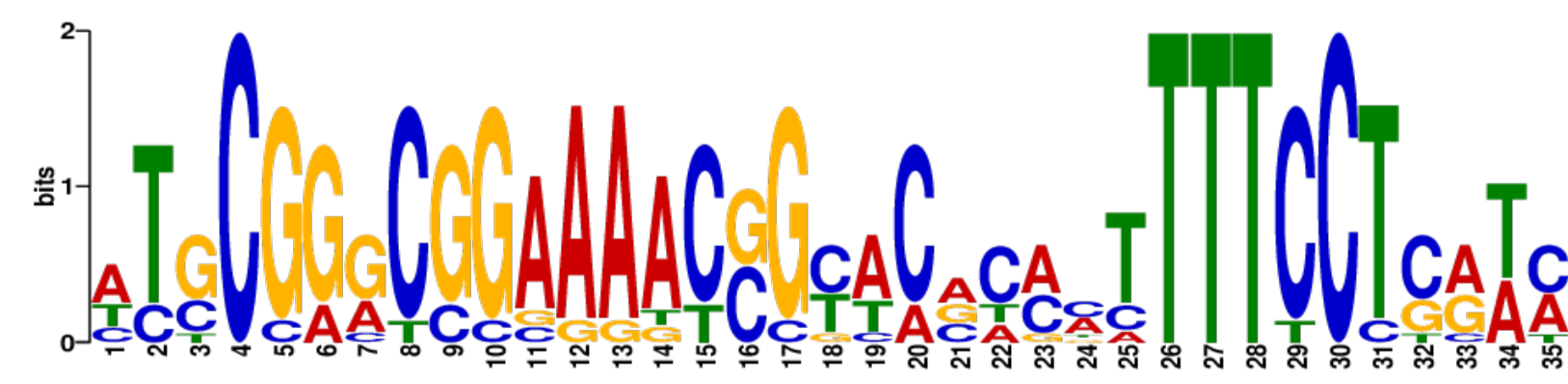
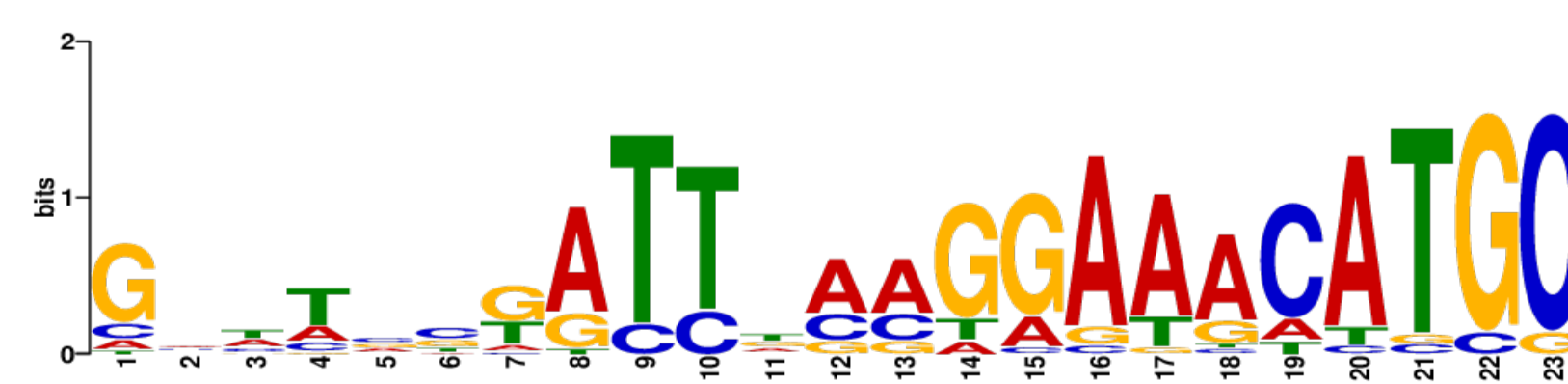
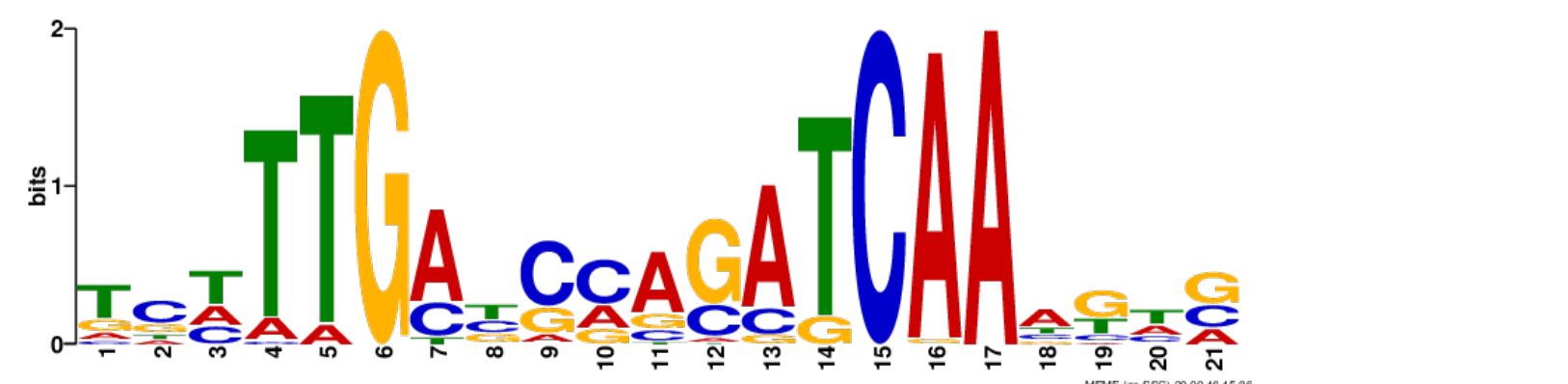
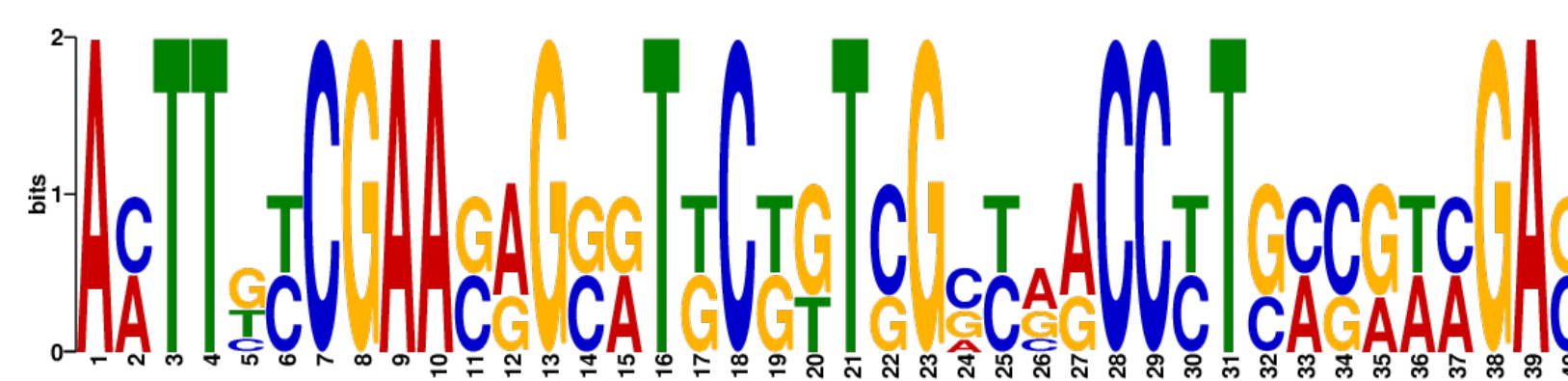


cluster ^a	motif	e-value	sites ^b	genes in cluster (%)
B1		4.6e-41	163	44.4
		5.4e-16	9	2.4
B2		6.8e-20	9	2.9
B3		1e-51	11	1.5
		9.6e-35	51	7.1
		2.2e-43	14	2.0
		1.2e-16	84	11.7
B4B		1.1e-35	10	2.7
		3.8e-39	10	2.7
		9.5e-16	21	5.7
A		3.3e-80	47	31.1
		5.3e-17	6	4.0