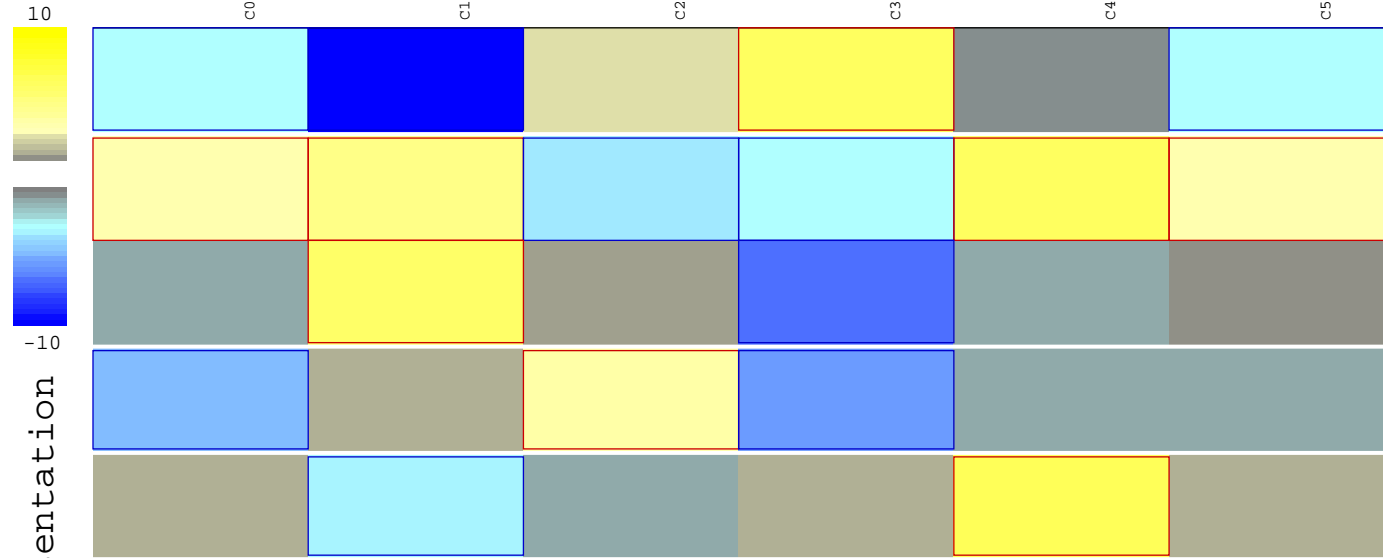
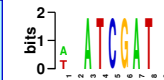
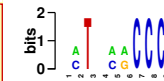
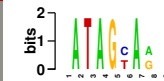
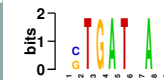
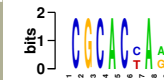


over-representation

representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.005	27.0	10/10	-	↔	1.00	AATCGAT	CEH-28, CEH_28, CEH_38 ...	-
	5'	0.004	23.0	10/10	-	→	0.98	CTCCGCC	KLF_1, KLF_3	-
	5'	0.003	14.9	7/10	-	←	0.61	ATAGTAA	DAF_19	-
	5'	0.003	14.7	9/10	-	←	1.00	CTGATAA	EGL_18_ELT-6_ELT_6, ELT_7, PQM-1 ...	-
	5'	0.002	11.1	6/10	-	→	0.14	CGCACCA	-	-