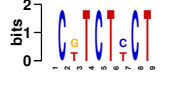
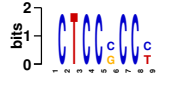
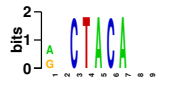
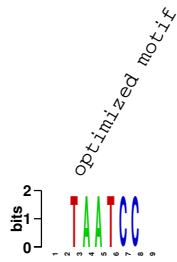
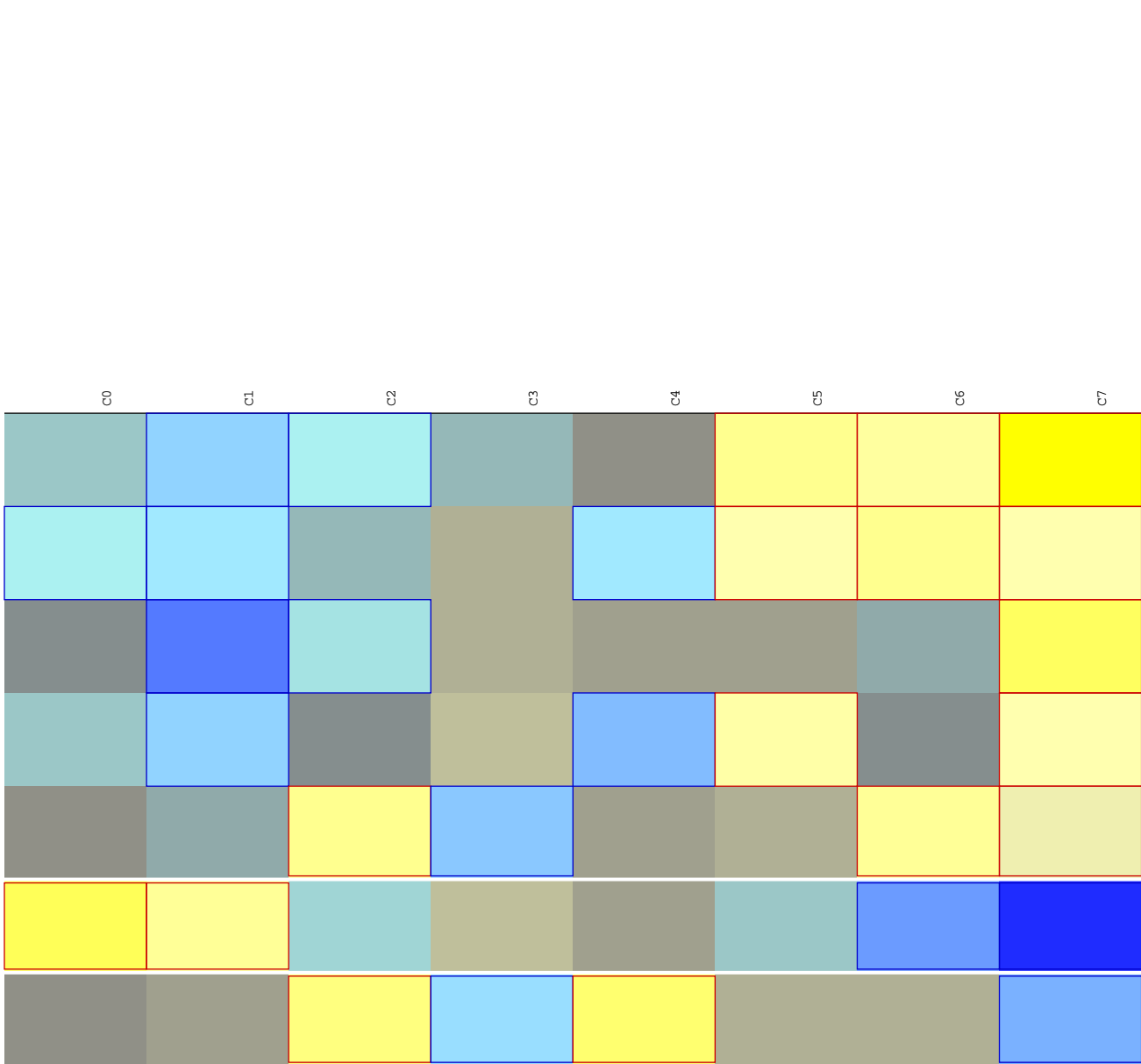


over-representation

under-representation



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	Motif name	Protein array
5'	0.006	28.1	10/10	Y	↔	0.98	ATAATCC	CEH_45, UNC-30 UNC_30, CEH_53 ...	-
5'	0.003	16.3	10/10	-	↔	0.94	ACTACAA	NHR_28_NHR_53_NHR_70, NHR_109_NHR_12_NHR_16_NHR_273_NHR_86, NHR_2...	-
5'	0.003	13.8	9/10	-	←	0.70	AGATTAA	CEH_37, CEH_53, TTX_1 ...	-
5'	0.003	13.6	8/10	-	←	0.31	TATATGA	UNC-86, UNC_86	-
5'	0.003	12.4	7/10	-	↔	1.00	CTCCGCC	PIE_1	-
5'	0.005	26.6	10/10	-	↔	1.00	AATCGAT	CEH_28, CEH-28, CEH-38 ...	-
5'	0.004	17.0	10/10	Y	↔	0.99	CTTCTTC	PIE_1	-