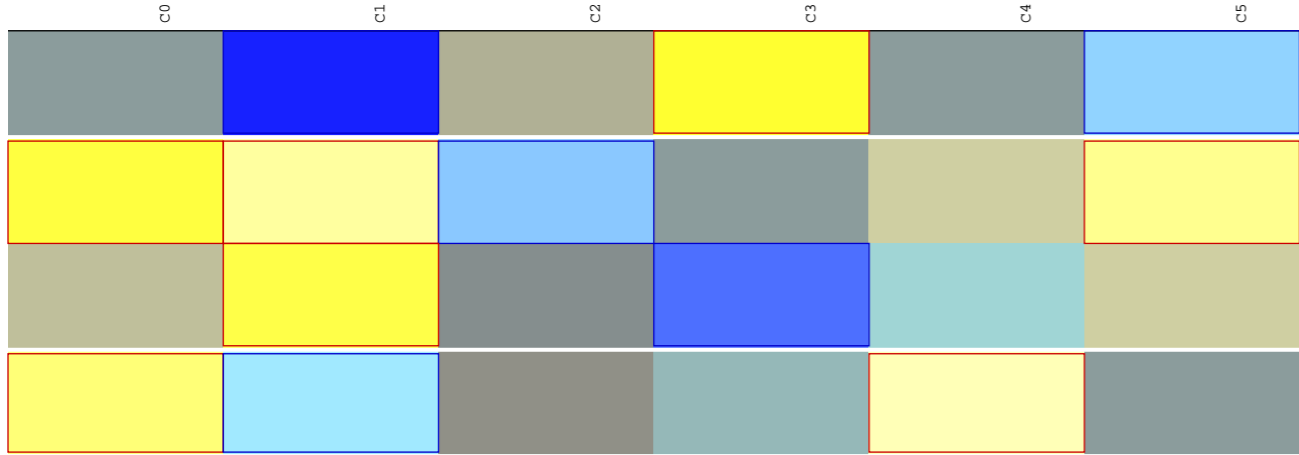
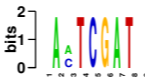
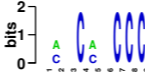
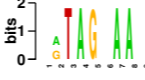



over-representation
representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.004	25.2	10/10	-	↔	1.00	AATCGAT	CEH-28, CEH_28, CEH-38 ...	-
	5'	0.004	21.8	10/10	Y	→	1.00	CTCCGCC	KLF_1	-
	5'	0.003	18.4	10/10	-	↔	0.29	ATAGTAA	DAF_19, PIE_1	-
	5'	0.002	11.5	7/10	-	-	0.51	CAGTGCA	LIN_15B, PIE_1	-