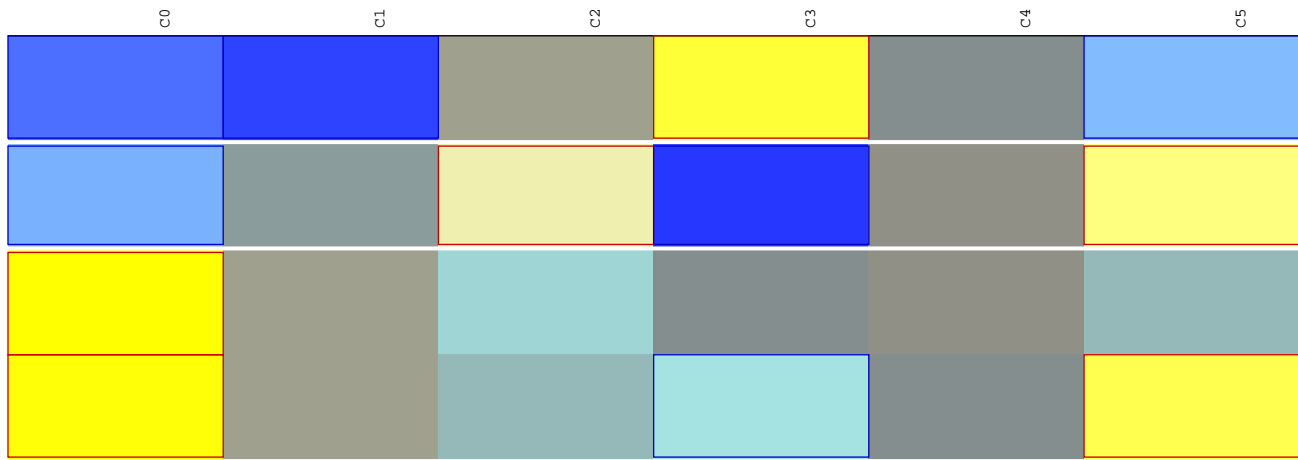
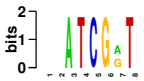

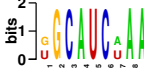
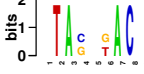


over-representation

representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.005	32.7	10/10	-	↔	0.99	AATCGAT	CEH_28, CEH-28, CEH_48 ...	-
	3'UTR	0.004	22.7	10/10	-	→	0.97	AUUUGAA	-	-
	3'UTR	0.004	20.8	10/10	Y	→	0.06	GCAUCUA	-	-
	5'	0.004	21.5	10/10	-	↔	0.88	TACATAC	-	-