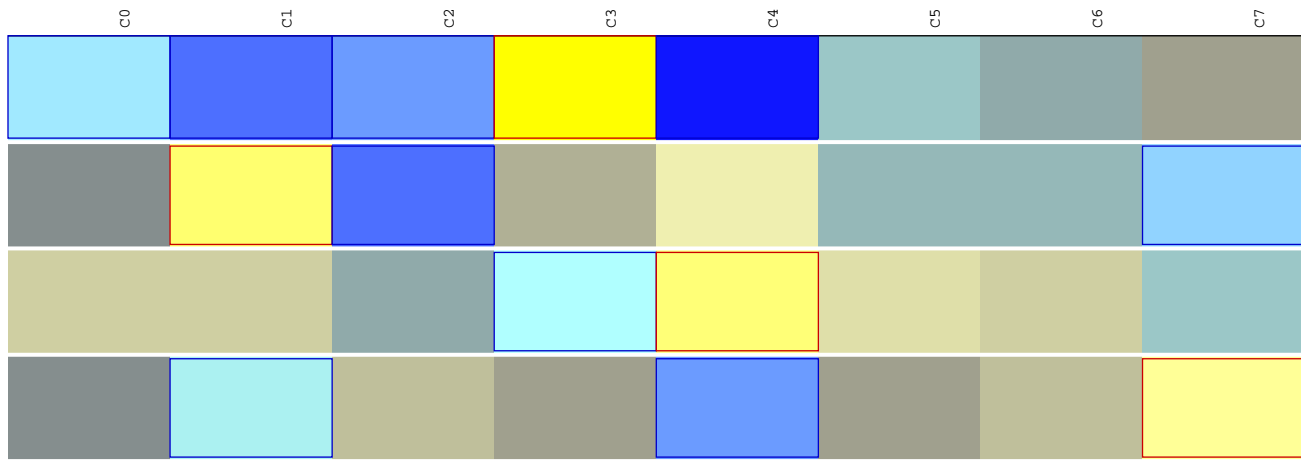
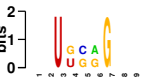
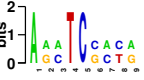
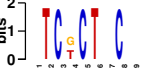



over-representation

representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	Motif name	Protein array
	3'UTR	0.008	39.0	10/10	Y	↔	0.98	UUUCAGA	-	-
	5'	0.004	17.7	10/10	-	↔	0.98	AATCGAT	PIE_1	-
	5'	0.003	11.7	6/10	-	→	0.91	TCTCTTC	EOR_1.2, EOR-1, PIE_1	-
	5'	0.003	11.2	10/10	-	-	0.21	ACATTTC	-	-