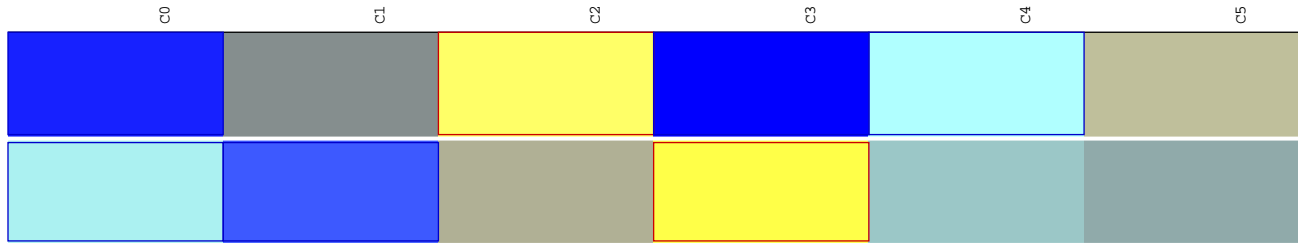
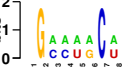
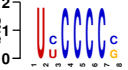


on
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.006	36.0	10/10	Y	→	0.98	GAAAACA	-	-
	3'UTR	0.004	21.4	10/10	-	→	0.97	UCCCCC	-	-