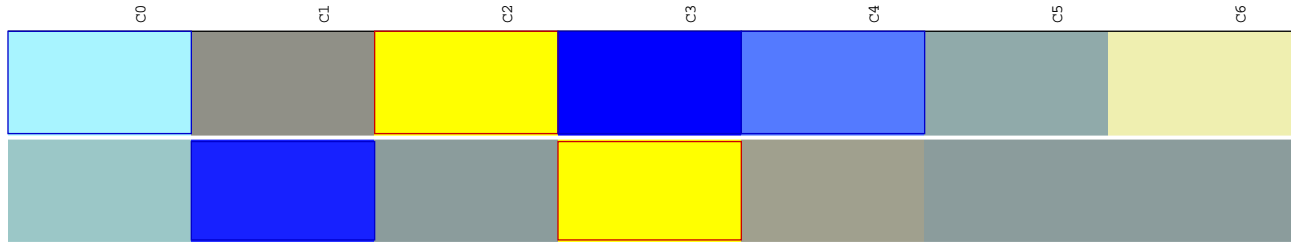
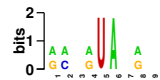
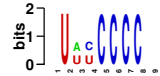


on
over-representation

10
-10



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.010	51.8	10/10	Y	→	0.99	AAAUAAA	-	-
	3'UTR	0.004	23.3	10/10	Y	→	0.99	UUUCCCC	-	-