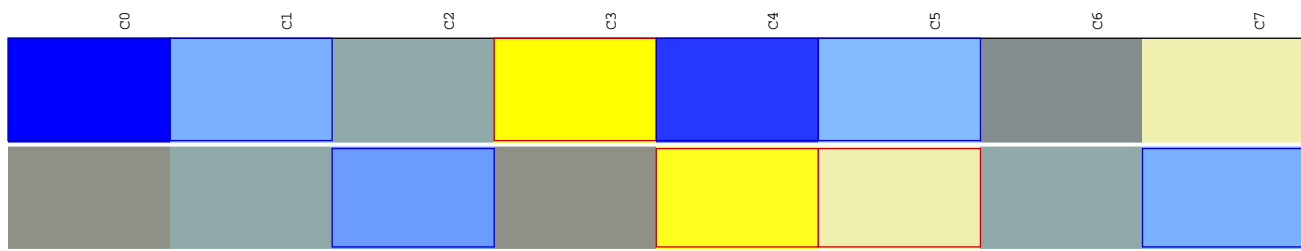
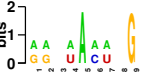
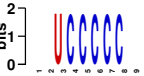


on  
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name	Protein array
	3'UTR	0.008	41.2	10/10	Y	↔	0.97	AAAAAUU	-	-
	3'UTR	0.004	20.6	10/10	-	→	0.99	AUCCCCC	-	-