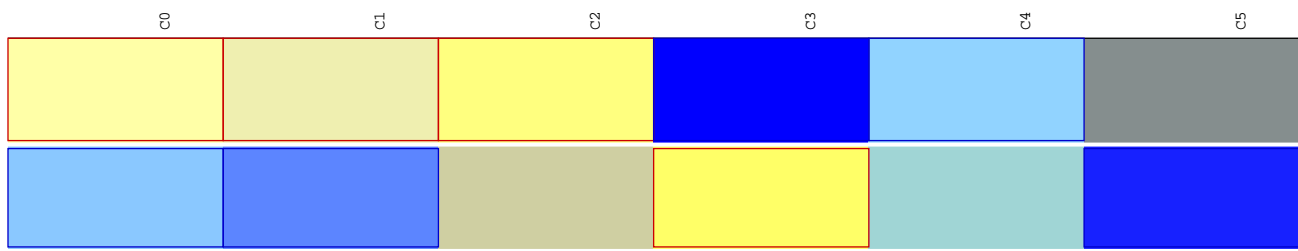
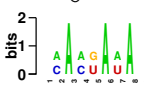
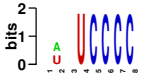


on
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.006	36.9	10/10	Y	→	0.99	AAAUAAA	-	-
	3'UTR	0.006	32.5	10/10	Y	↔	1.00	UUUCCCC	-	-