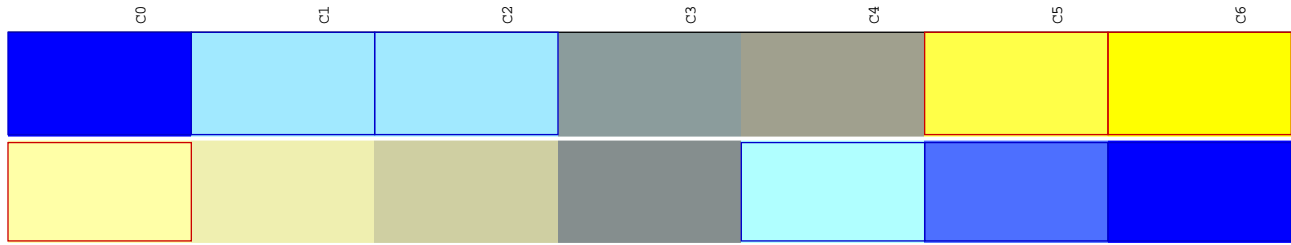
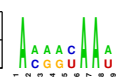
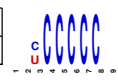


on
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.017	88.7	10/10	Y	↔	0.98	AAAAUAA	-	-
	3'UTR	0.005	26.9	10/10	-	→	0.99	UUC C C C C	-	-