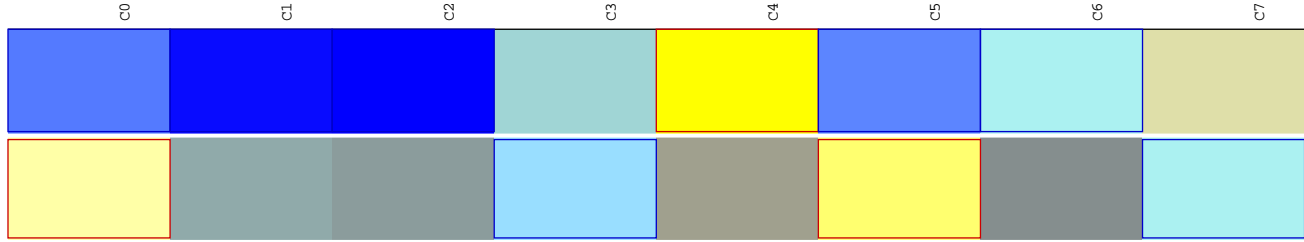
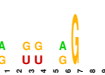
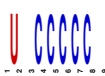


on
over-representation

10
-10



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.012	61.5	10/10	Y	↔	0.99	UUUCAGA	-	-
	3'UTR	0.003	12.7	9/10	-	→	0.98	UUCCCC	-	-