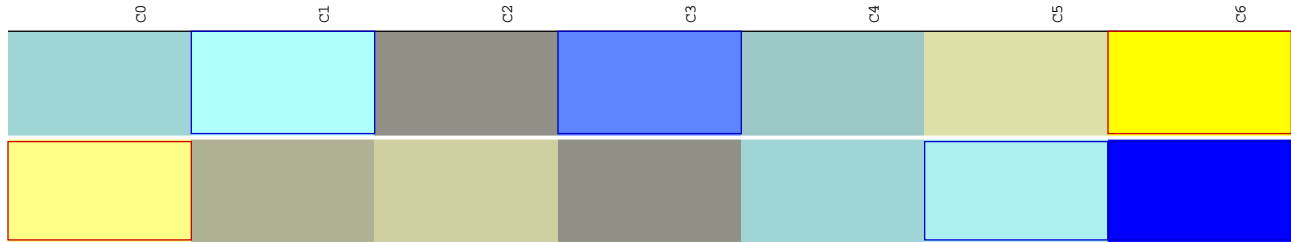

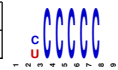


on
over-representation

10
-10



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.006	33.0	10/10	Y	→	0.94	UUUCAGA	-	-
	3'UTR	0.004	21.3	10/10	-	→	1.00	UUUUUUU	-	-