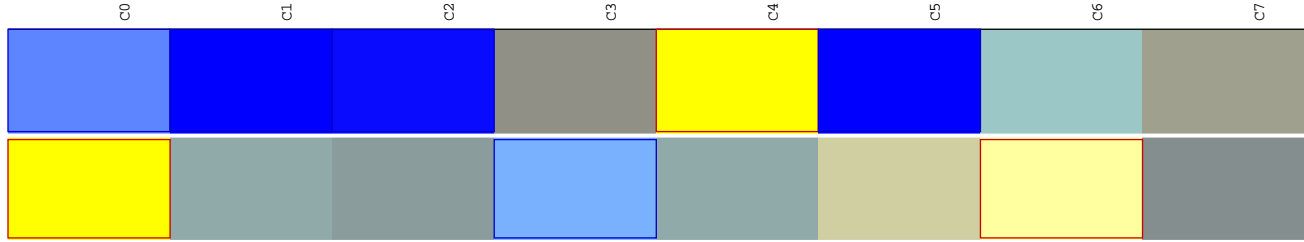
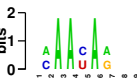
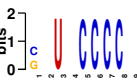


on
over-representation

10
-10



Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name	Protein array
	3'UTR	0.013	64.2	10/10	Y	↔	0.99	AAAUAAA	-	-
	3'UTR	0.004	19.6	10/10	-	→	0.96	UUUUUUU	-	-