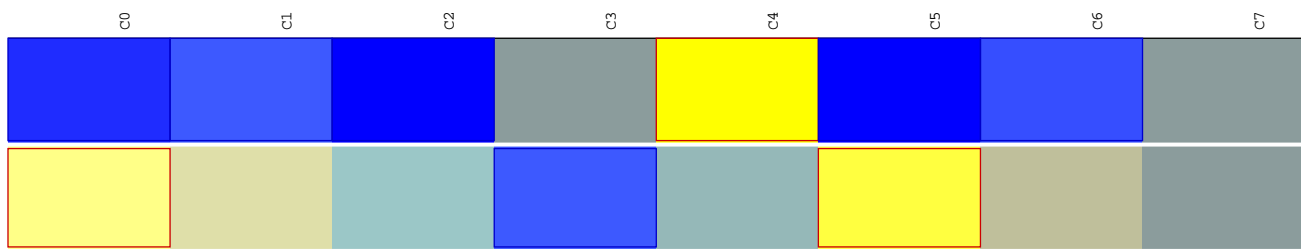
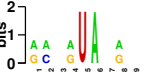


on
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name	Protein array
	3'UTR	0.018	87.3	10/10	Y	↔	0.99	AAAUAAA	-	-
	3'UTR	0.005	21.5	10/10	Y	↔	1.00	UCCCCU	-	-