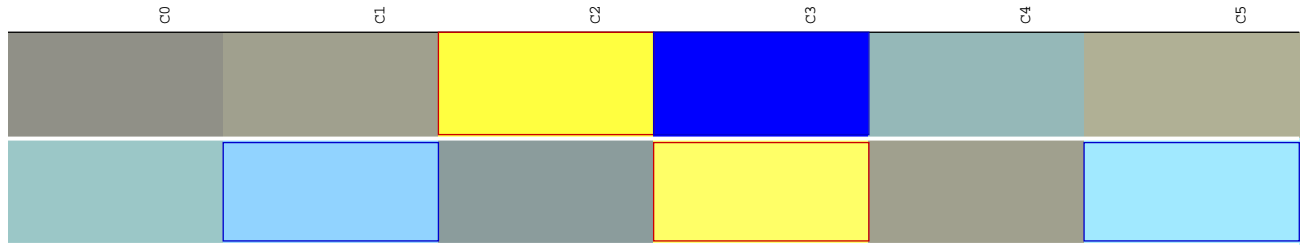
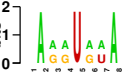
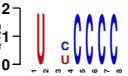


over-representation
on



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	Protein array
	3'UTR	0.006	37.3	10/10	Y	→	0.99	AAAUAAA	-	-
	3'UTR	0.003	15.0	10/10	Y	→	1.00	UUUGCCC	-	-