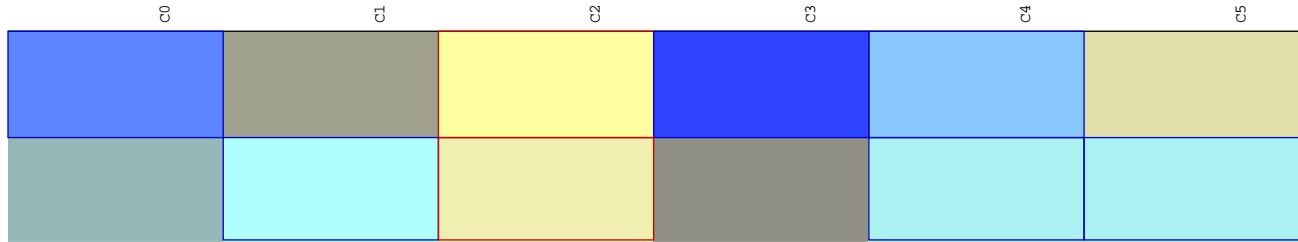
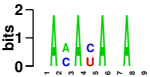
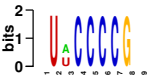


on  
over-representation

10  
-10



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.005	27.2	10/10	Y	↔	0.99	AAAUAAA	-	-
	3'UTR	0.002	10.3	6/10	-	→	0.65	UUCCCCG	-	-