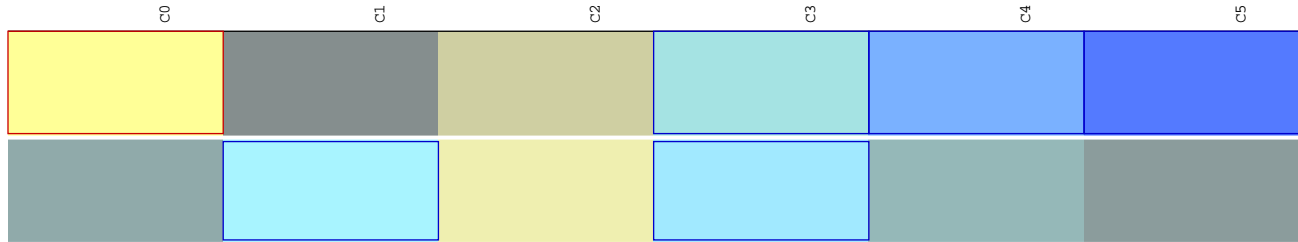
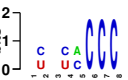



over-representation
on

10
-10



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	Protein array
	3'UTR	0.004	19.2	10/10	Y	→	1.00	UUUCCCC	-	-
	3'UTR	0.002	11.1	8/10	-	→	0.30	AGCUAUG	-	-