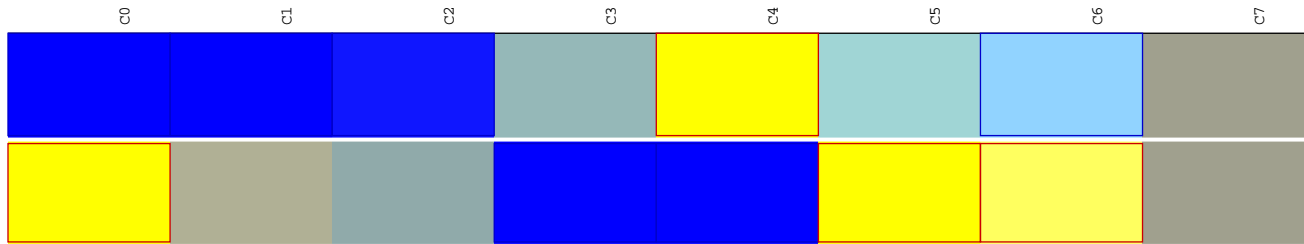


on  
-10  
10  
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name	Protein array
	3'UTR	0.025	128.0	10/10	Y	↔	0.99	AAAUAAA	-	-
	3'UTR	0.014	70.7	10/10	Y	↔	1.00	UUUGUU	-	-