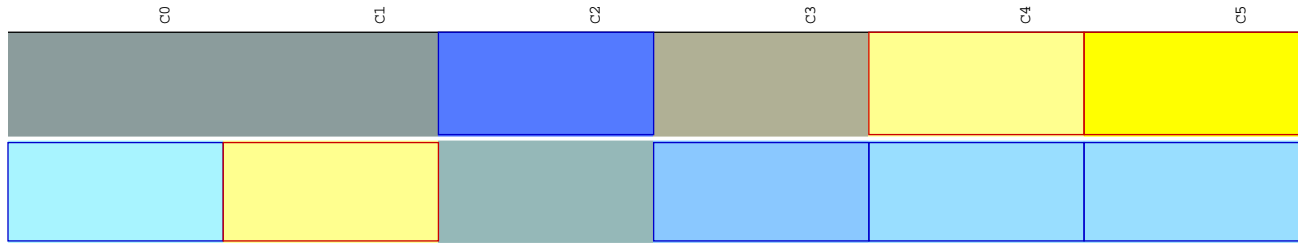


over-representation  
on



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.005	29.3	10/10	Y	↔	1.00	AGAAAAA	-	-
	3'UTR	0.004	20.7	10/10	Y	→	0.99	UUUCCCC	-	-