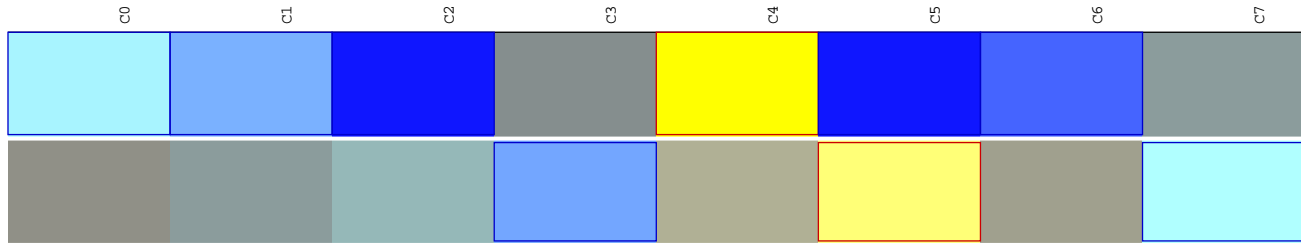


on  
over-representation

10  
-10



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.011	55.8	10/10	Y	↔	0.99	AAAAUAA	-	-
	3'UTR	0.003	12.6	10/10	-	→	0.95	UCCCCC	-	-