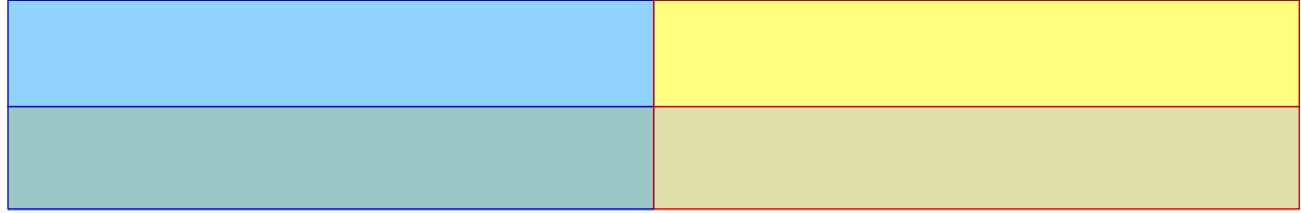




on
over-representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.005	48.4	10/10	Y	→	0.98	-	-	-
	3'UTR	0.002	16.7	6/10	Y	→	0.21	-	-	-