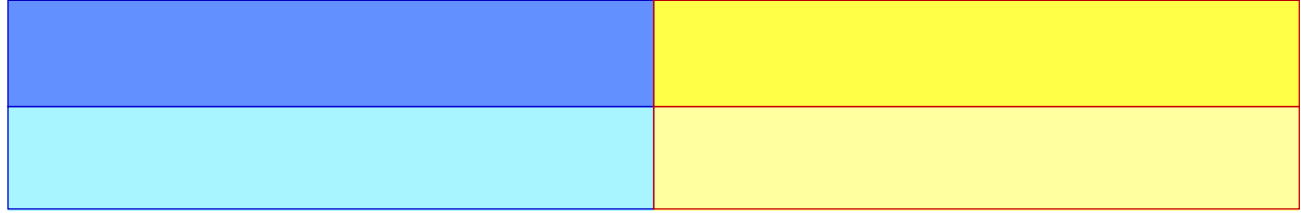

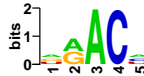


on

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	76.5	10/10	Y	→	0.98		-	-
	3'UTR	0.003	36.4	10/10	Y	→	0.21		-	-