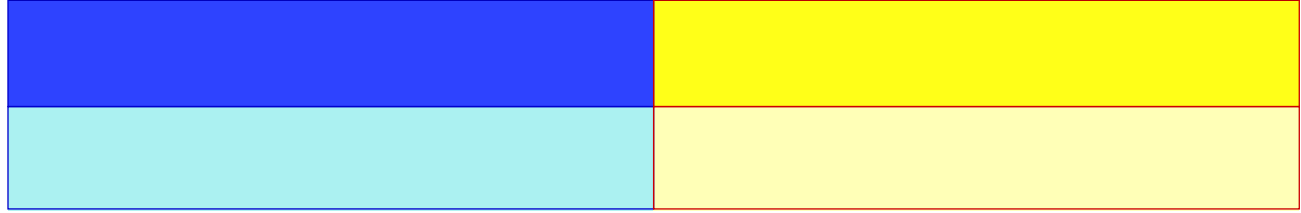
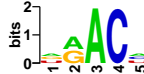



on
over-representation



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.001	99.0	10/10	Y	-	0.21		-	-
	3'UTR	0.000	24.4	9/10	Y	-	0.98		-	-