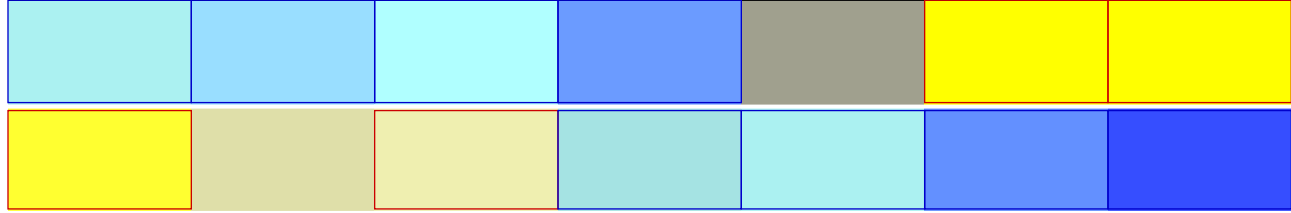



on
over-representation

10
-10



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.014	75.5	10/10	Y	↔	0.98	AAAUAAA	-	-
	3'UTR	0.006	31.1	10/10	Y	→	1.00	UUUUUUU	-	-