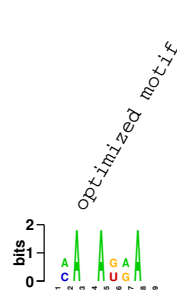


on
-10
10 over-representation



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.007	35.0	10/10	Y	→	0.98	AAAAUAA	-	-