



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.032	110.3	10/10	Y	↔	0.98	AAAAUAA	-	-
	3'UTR	0.014	46.8	10/10	Y	↔	1.00	UUCCCCC	-	-
	3'UTR	0.004	11.2	10/10	Y	→	0.23	GCUGCCU	-	-