






optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.016	1490.3	10/10	Y	-	0.32	CCCCCC	-	-
	3'UTR	0.013	1243.0	10/10	-	-	0.84	UCAAUUG	-	-
	3'UTR	0.001	48.5	10/10	-	-	0.19	ACGCUAG	-	-
	3'UTR	0.015	1416.4	10/10	Y	-	0.97	GAAGAAG	-	-
	3'UTR	0.006	500.3	10/10	Y	-	0.23	CGGCGGC	-	-