



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.006	30.7	10/10	Y	→	0.98	AAAUAAA	-	-
3'UTR	0.004	18.2	10/10	-	→	0.99	UUUCCCC	-	-