



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.017	90.1	10/10	Y	↔	0.99	AAAAAUA	-	-
3'UTR	0.006	29.3	10/10	-	→	0.99	UUCCCCC	-	-