optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3’UTR	0.024	119.5	10/10	Y	↔	0.99	AAAAUAA	-	-
	3’UTR	0.007	34.4	10/10	-	↔	0.99	UUUCCCC	-	-