	optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
		3'UTR	0.010	49.7	10/10	Y	$\leftrightarrow$	0.99	UUGAAAA	–	–
		3'UTR	0.004	17.5	10/10	–	$\rightarrow$	0.99	UUCCCCC	–	–