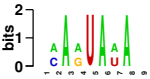

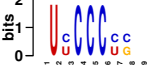


optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.008	43.6	10/10	Y	→	0.99	AAAUAAA	-	-
	3'UTR	0.004	25.4	10/10	-	→	1.00	UAAUAAA	-	-
	3'UTR	0.004	21.8	10/10	-	↔	1.00	UUUUUUU	-	-