



cytoplasm organization and biogenesis, p<1e-04
 ribosome, p<1e-07
 cytosolic ribosome (sensu Eukaryota), p<1e-15

optimized motif	location	MI (bits)	z-score	Robustness	Position bias	orientation bias	conservation index	seed	motif name
	5'	0.026	14.2	10/10	-	-	1.00	AAAATTT	RRPE
	3'UTR	0.019	9.0	7/10	-	→	1.00	AUAUAUU	-
	3'UTR	0.018	8.7	8/10	-	→	1.00	UGUAUAA	-