

over-representation

under-representation



		C0		C1		optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
							3'UTR	0.006	52.7	10/10	-	→	0.75	AGAUUAG	-	-
							3'UTR	0.005	47.0	10/10	-	→	0.54	UAAGGU	-	-
							3'UTR	0.005	45.5	10/10	-	→	0.81	CUAACA	-	-
							3'UTR	0.006	49.7	10/10	-	→	0.20	CCAUCG	-	-
							3'UTR	0.005	48.3	10/10	Y	→	0.58	GGUCAA	-	-
							3'UTR	0.004	35.1	10/10	Y	→	0.78	GUCGAU	-	-
							3'UTR	0.003	28.8	10/10	-	→	0.74	CAUGUA	-	-
							3'UTR	0.003	22.6	8/10	-	→	0.58	GUGAGUA	-	-
							3'UTR	0.002	21.9	8/10	-	→	0.35	CGACCA	-	-