
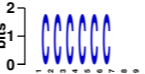
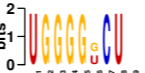


optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.006	27.7	10/10	-	→	0.96	AGUUUUU	-	-
	3'UTR	0.003	14.4	10/10	Y	→	0.90	CCCCCCA	-	-
	3'UTR	0.003	14.6	10/10	-	→	0.06	GGGGGCU	-	-