



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.012	55.6	10/10	Y	↔	0.94	AAAAUAA	-	-
	5'	0.005	21.9	10/10	-	↔	1.00	AATCGAT	CEH_28, CEH-28, CEH-38 ...	-
	5'	0.003	13.7	10/10	-	↔	0.95	TCTTCTC	PIE_1	-