



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.004	22.4	10/10	Y	←	1.00	AATCGAT	CEH_28, CEH-38, CEH-28 ...	-
	5'	0.003	13.7	7/10	-	←	0.79	ACGGCGG	-	-
	5'	0.002	11.0	7/10	-	-	0.30	TACCAAA	PIE_1, TBX_40	-