optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5′	0.004	21.8	10/10	-	↔	1.00	AATCGAT	PIE_1	-
	5′	0.003	14.5	8/10	-	→	0.99	TGCGCTC	LIN_15B, EFL_1, PIE_1	-
	5′	0.003	15.1	10/10	Y	↔	0.96	TACAAAC	NHR_28_NHR_53_NHR_70, IRX_1	-