



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	5'	0.002	13.4	9/10	-	↔	0.98	AACGCGC	LIN_15B, EFL_1, PIE_1	-
	5'	0.002	13.1	8/10	-	-	0.99	AAGCGGA	-	-
	5'	0.002	12.0	6/10	-	→	0.32	AGTACCA	-	-