



<div>bits 2 1 0</div> <div>optimized motif</div>	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
<div>bits 2 1 0</div> <div>GAAGGGGUA</div>	3'UTR	0.005	29.3	10/10	Y	↔	1.00	AGAAAAA	-	-
<div>bits 2 1 0</div> <div>TACCGAA</div>	5'	0.002	11.0	7/10	-	-	0.30	TACCAAA	PIE_1, TBX_40	-
<div>bits 2 1 0</div> <div>CAATCGAT</div>	5'	0.004	22.4	10/10	Y	←	1.00	AATCGAT	CEH_28, CEH-38, CEH-28 ...	-
<div>bits 2 1 0</div> <div>UUCGGG</div>	3'UTR	0.004	20.7	10/10	Y	→	0.99	UUUCCCC	-	-
<div>bits 2 1 0</div> <div>ACGGCGGA</div>	5'	0.003	13.7	7/10	-	←	0.79	ACGGCGG	-	-