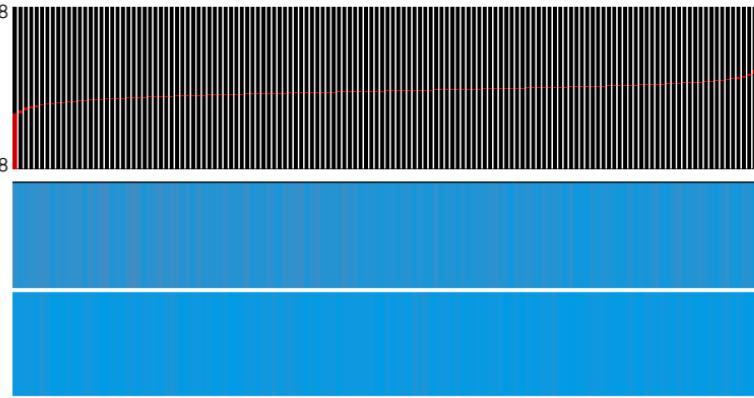


structural molecule activity, $p < 0.01$



motif name	seed	conservation index	orientation bias	position bias	robustness	z -score	location	MI (bits)	Optimized motif
TCTTATC	1.00	1.00	↑	-	9/10	8.8	5'	0.015	
GGAGCAA	0.94	0.94	→	-	6/10	6.2	3' UTR	0.013	