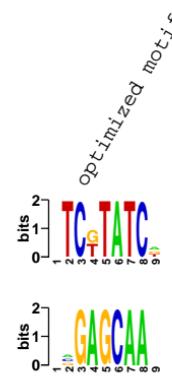
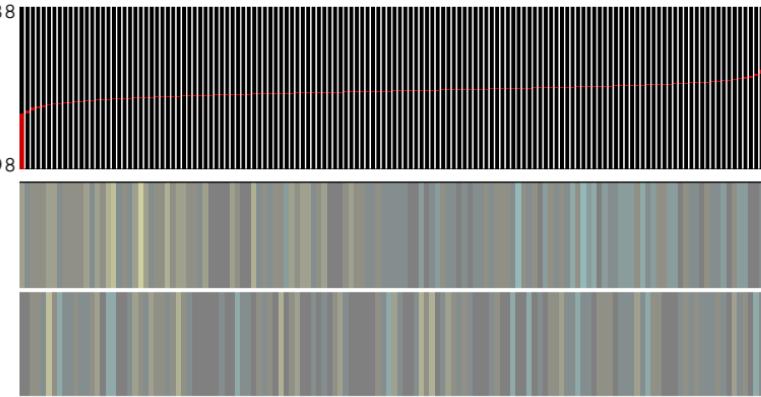


Over-representation

structural molecule activity, p<0.07



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