over-representation

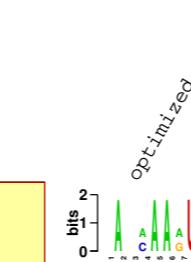10

-10

| | C0 | C1 | C2 | C3 | C4 | C5 | C6 |

optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.006	28.6	10/10	Y	↔	0.97	AAAAAAU	–	–