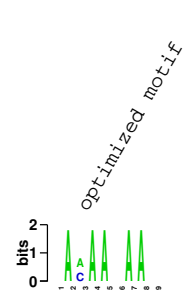
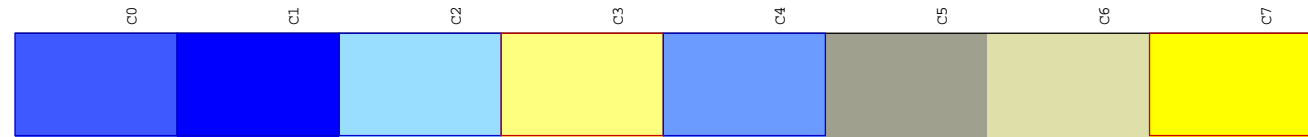


on
-10
10
over-representation



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
3'UTR	0.010	50.6	10/10	Y	→	0.99	AAAAUAA	-

protein array