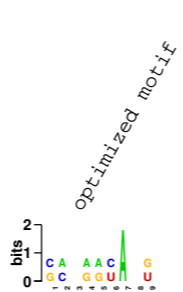
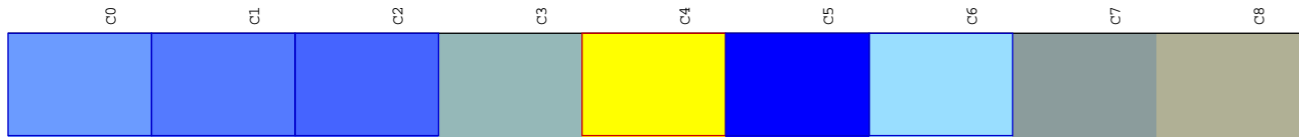


on
-10
10 over-representation



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.012	55.6	10/10	Y	↔	0.94	AAAAUAA	-	-