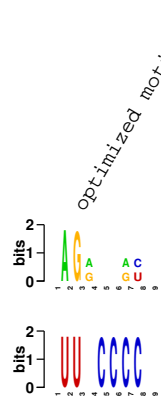
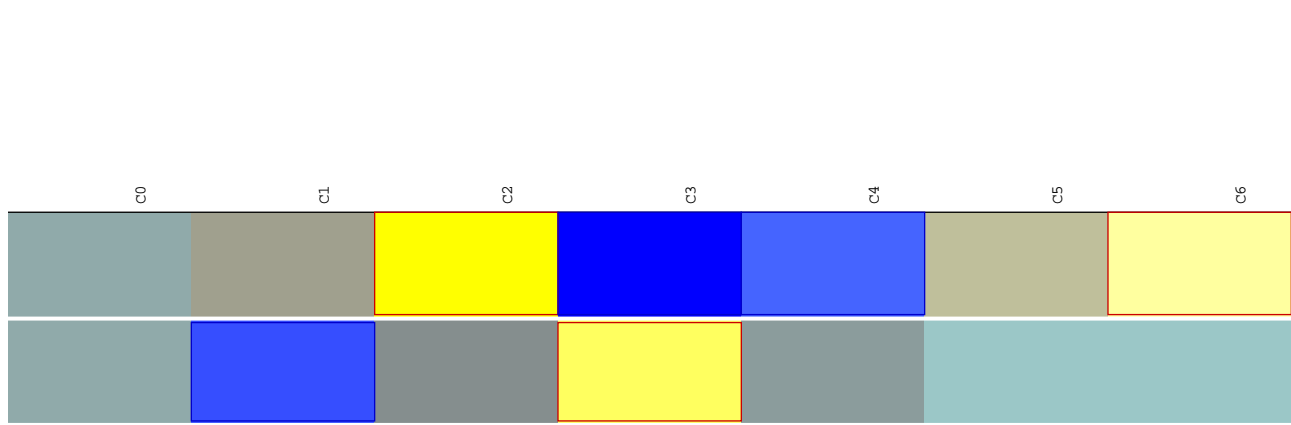


on
over-representation

10
-10



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.012	65.1	10/10	Y	↔	0.99	AGAAAAU	RSP_3	-
3'UTR	0.004	17.6	10/10	-	→	0.99	UUUCCCC	-	-