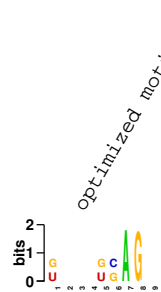
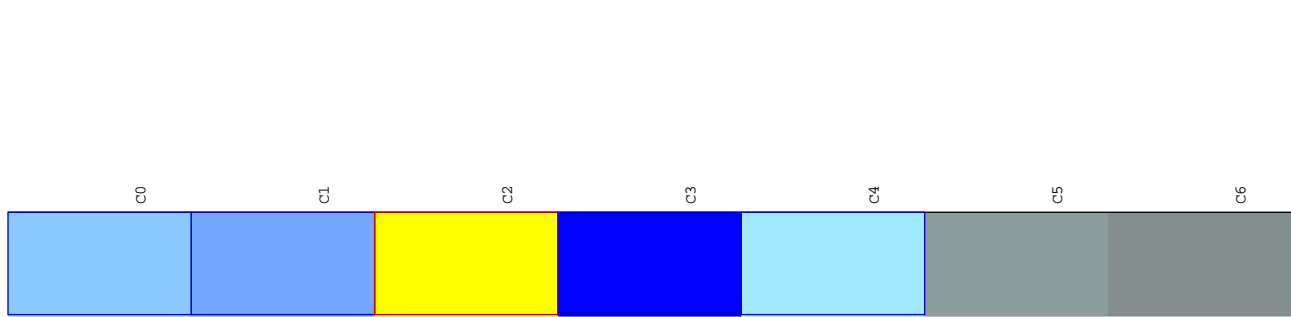


on
over-representation

10
-10



Optimized motif	location	MI (bits)	z-score	robustness	Position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.008	44.7	10/10	Y	↔	0.96	UUUUCAG	-	-	