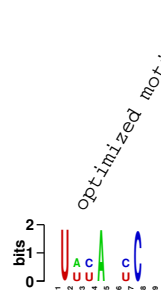
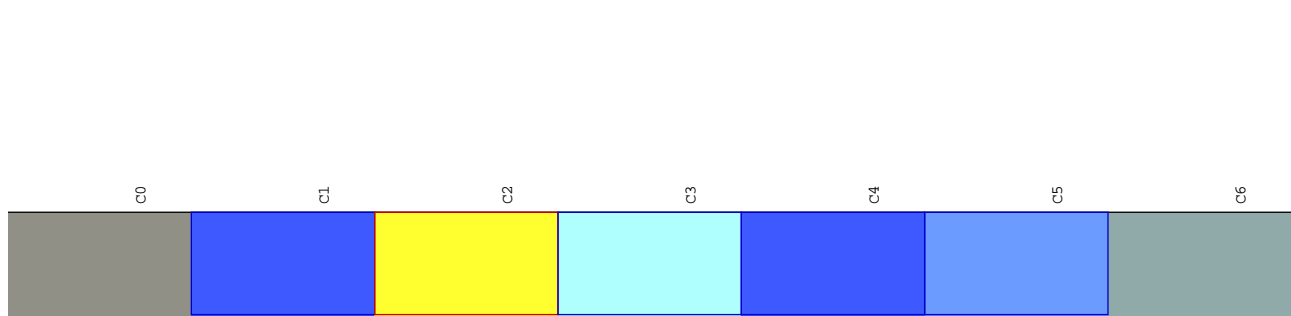


on
-10
10 over-representation



Optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
UUUUUU	3'UTR	0.007	33.9	10/10	Y	↔	0.98	UUUAUUC	-	-