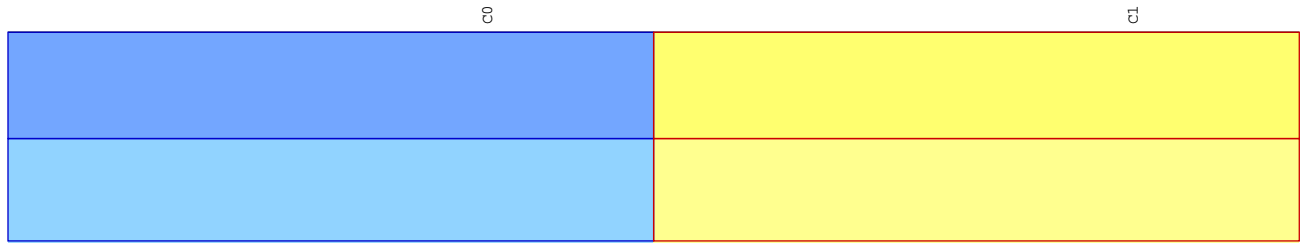




over-representation
on



optimized motif	location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
	3'UTR	0.007	62.0	10/10	-	-	0.21		-	-
	3'UTR	0.005	44.6	10/10	-	→	0.98		-	-