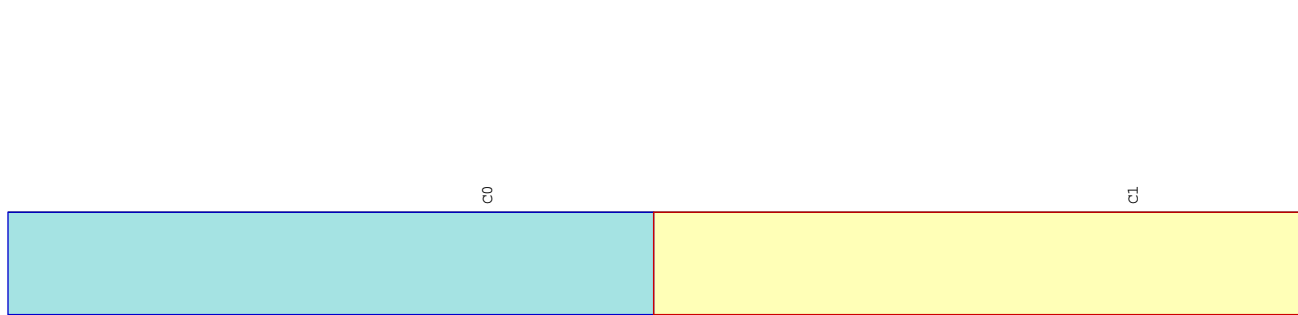


over-representation  
on

20  
-20



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name	protein array
3'UTR	0.001	23.8	9/10	Y	-	0.98		-	-