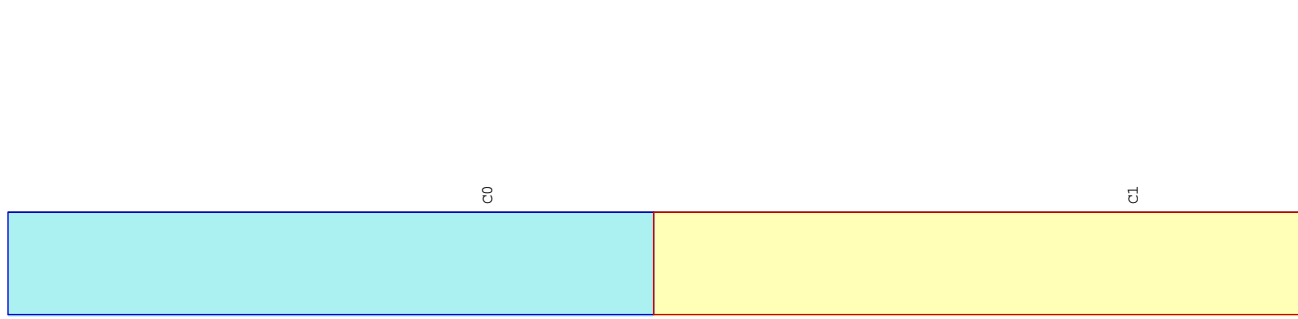


on

over-representation

20

-20



location	MI (bits)	z-score	robustness	position bias	orientation bias	conservation index	seed	motif name
3'UTR	0.005	25.7	10/10	-	→	0.98	-	-

protein array