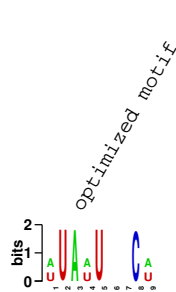
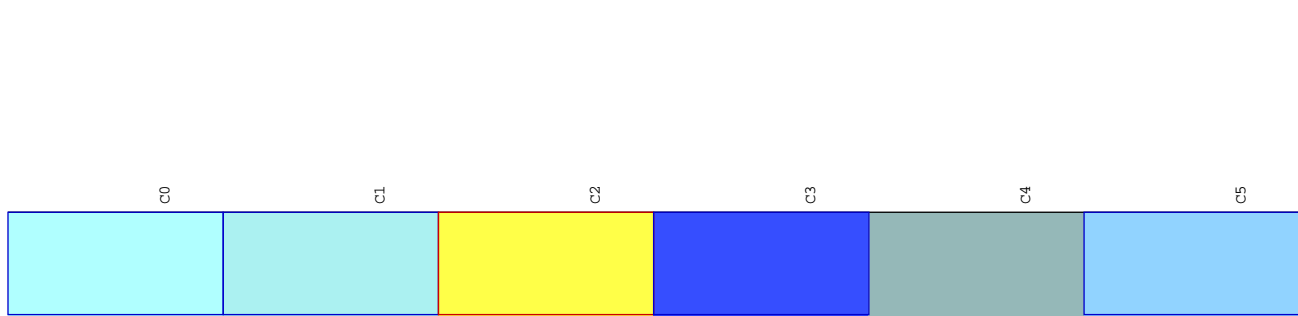


on
-10
10 over-representation



location
3'UTR

MI (bits)
0.005

z-score
27.8

robustness
10/10

position bias
Y

orientation bias
→

conservation index
0.87

seed
UAUUUUC

motif name
-

protein array
-