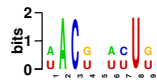
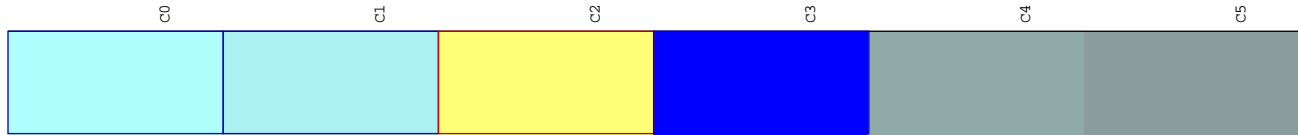
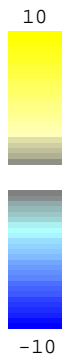


on
-10
over-representation



optimized motif

location

MI (bits)

z-score

robustness

position bias

orientation bias

conservation index

seed

motif name

3'UTR

0.005

26.6

10/10

-

↔

0.66

ACUUUUU

-

protein array

-