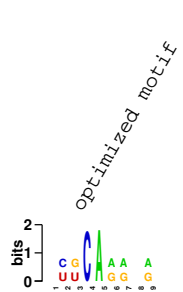
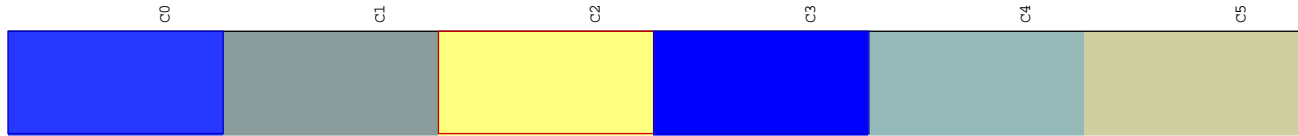


on

over-representation

10

-10



location

3'UTR

MI (bits)

0.006

z-score

32.0

robustness

10/10

position bias

-

orientation bias

↔

conservation bias

0.85

seed

UUCAGAA

motif name

-

protein array

-