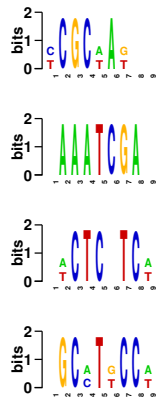
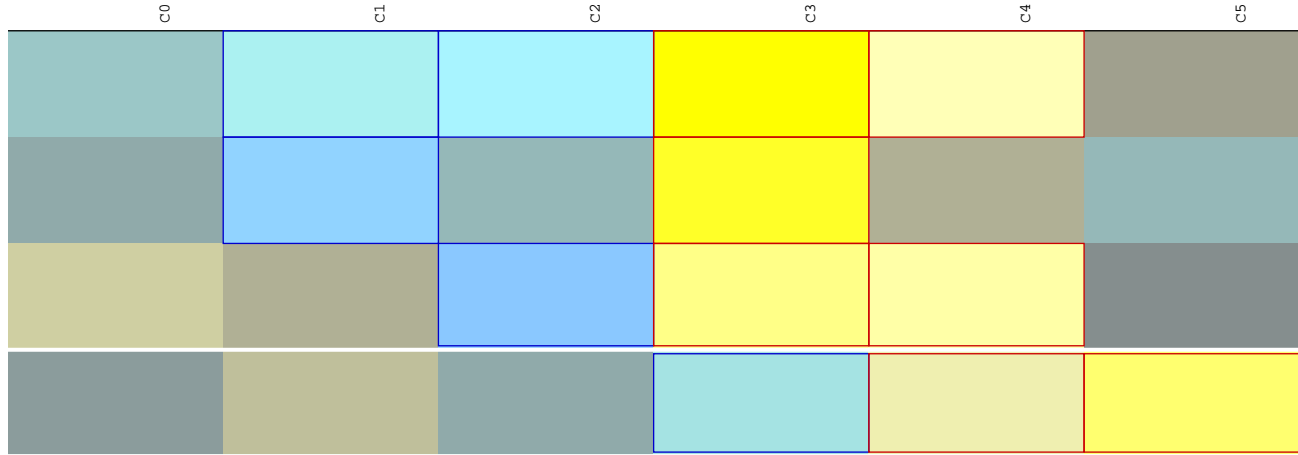


over-representation

representation



Optimized motif

location

MI (bits)

z-score

robustness

position bias

orientation bias

conservation index

seed

motif name

protein array

5'

0.003

19.8

10/10

-

↔

0.87

CGCAATA

-

-

5'

0.003

15.8

10/10

-

↔

0.99

AAATCGA

CEH-38, CEH_28, CEH-28 ...

-

5'

0.002

13.4

9/10

-

→

0.92

TCTCATC

PIE_1

-

5'

0.002

10.2

6/10

-

-

0.45

GCCTGCC

-

-