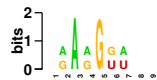
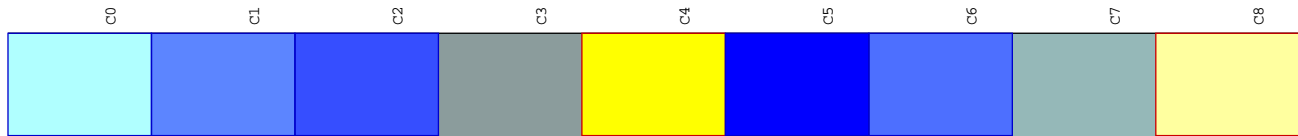


on
-10
10 over-representation



optimized motif

location

MI (bits)

z-score

robustness

position bias

orientation bias

conservation index

seed

motif name

protein array

3'UTR

0.011

51.7

10/10

Y

↔

0.97

AAAGUUU

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