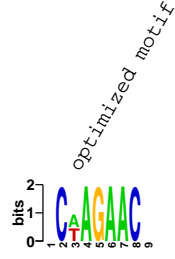


1
0



c0

c1 nucleobase, nucleoside, nucleotide and nucleic acid metab



optimized motif

location

5'

MI (bits)

0.009

z-score

15.6

robustness

6/10

position bias

-

orientation bias

-

conservation index

-

seed

CAAGAAC

motif name

-