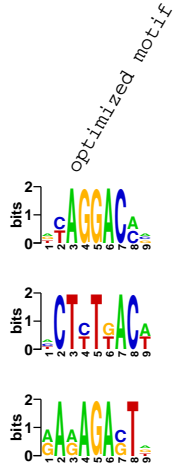




C0 nucleobase, nucleoside, nucleotide and nucleic acid metab

C1 morphogenesis, p<0.01



Optimized motif

location

MI (bits)

z-score

robustness

position bias

orientation bias

conservation index

seed

motif name

5'

0.012

23.0

9/10

-

→

-

CAGGACA

-

5'

0.011

20.5

10/10

-

←

-

CTTTAC

-

5'

0.011

20.8

6/10

-

→

-

AAAGAGT

-