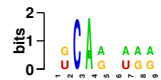


on
over-representation

10
-10



Optimized motif

3'UTR

location

0.008

MI (bits)

42.5

z-score

10/10

robustness

Y

position bias

→

orientation bias

0.94

conservation index

UCAAAAA

seed

-

motif name

-

protein array