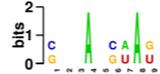


bits over-representation



-10

on



Optimized motif

location

3'UTR

MI (bits)

0.007

z-score

38.7

robustness

10/10

position bias

Y

orientation bias

→

conservation index

0.97

seed

AAAACAA

motif name

-

protein array

-