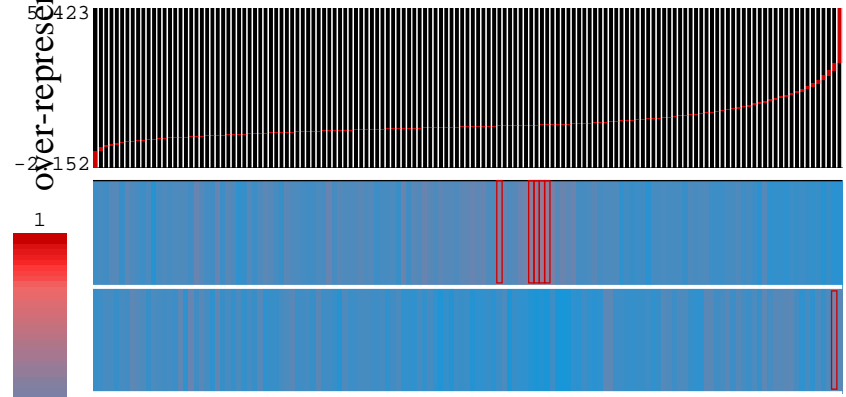


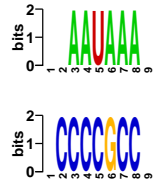
over-representation

1

0



protein serine/threonine phosphatase complex, p<0.05



5'

0.017

11.4

10/10

-

-

1.00

CCCCGCC

Sp1



3'UTR

0.018

12.3

10/10

Y

→

1.00

CAAUAAA

-

Optimized motif

location

MI (bits)

z-score

robustness

position bias

orientation bias

conservation index

seed

motif name