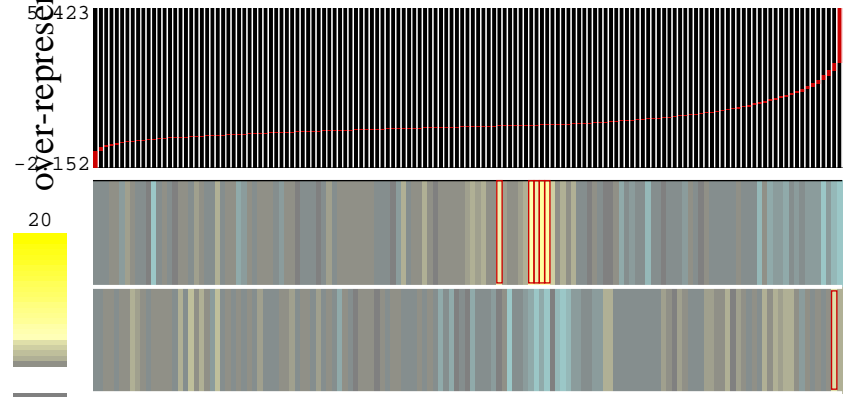
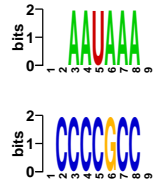


over-representation

20



protein serine/threonine phosphatase complex, p<0.



Optimized motif

3'UTR

location

0.018

MI (bits)

12.3

z-score

10/10

robustness

Y

position bias

→

orientation bias

1.00

conservation index

seed

CAAUAAA

-

motif name



5'

0.017

11.4

10/10

-

-

1.00

CCCCGCC

Sp1