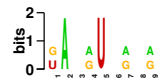
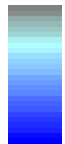


on
-10
over-representation

10



Optimized motif

location

MI (bits)

z-score

robustness

position bias

orientation bias

conservation index

seed

motif name

protein array

3'UTR

0.008

40.3

10/10

Y

↔

0.98

AAAUAAA

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