



location

*M*T (bits)

*z*-score

robustness

position bias

orientation bias

conservation index

seed

motif name

protein array

bits

bits

3' UTR

3' UTR

0.008

0.004

41.2

20.6

10/10

10/10

Y

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↔

→

0.97

0.99

AAAAAUU

AUCCCCC

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