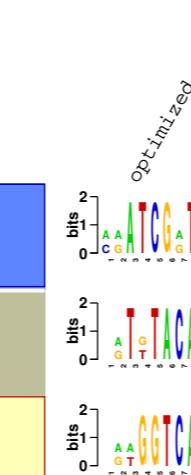


over-representation  
ratio

c0 c1 c2 c3 c4 c5 c6

10

-10



location

5'

0.003

17.5

10/10

-

↔

0.99

AATCGAT

PIE\_1, CEH\_28, CEH-28 ...

MI (bits)

0.003

12.8

10/10

-

↑

0.91

ATGTACA

DAF-16, PHA\_4.3, FKH\_8

z-score

10.3

8/10

-

↑

0.99

AAGGTCA

NHR\_91, SEX\_1, NHR\_84 ...

robustness

position bias

orientation bias

conservation index

seed

motif name

protein array