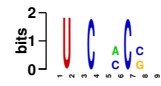
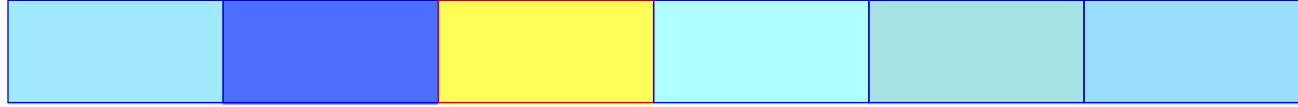


on
-10
over-representation
10



Optimized motif

location

3'UTR

MI (bits)

0.005

z-score

26.8

robustness

10/10

position bias

Y

orientation bias

→

conservation bias

0.99

seed

UUCCCCC

motif name

-

protein array

-