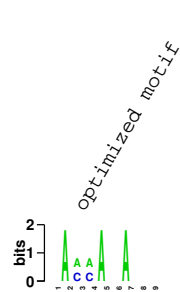


on
-10
10 over-representation



location
3'UTR

MI (bits)
0.013

z-score
65.0

robustness
10/10

position bias
Y

orientation bias
↔

conservation bias
0.98

seed
AAAAUAA

motif name
-

protein array
-