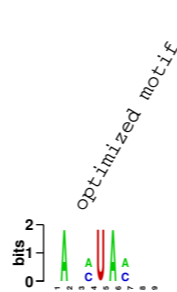


on  
-10  
10 over-representation



location  
3'UTR

MI (bits)  
0.007

z-score  
34.8

robustness  
10/10

position bias  
Y

orientation bias  
↔

conservation bias  
0.97

seed  
AAAUAAA

motif name  
-

protein array  
-