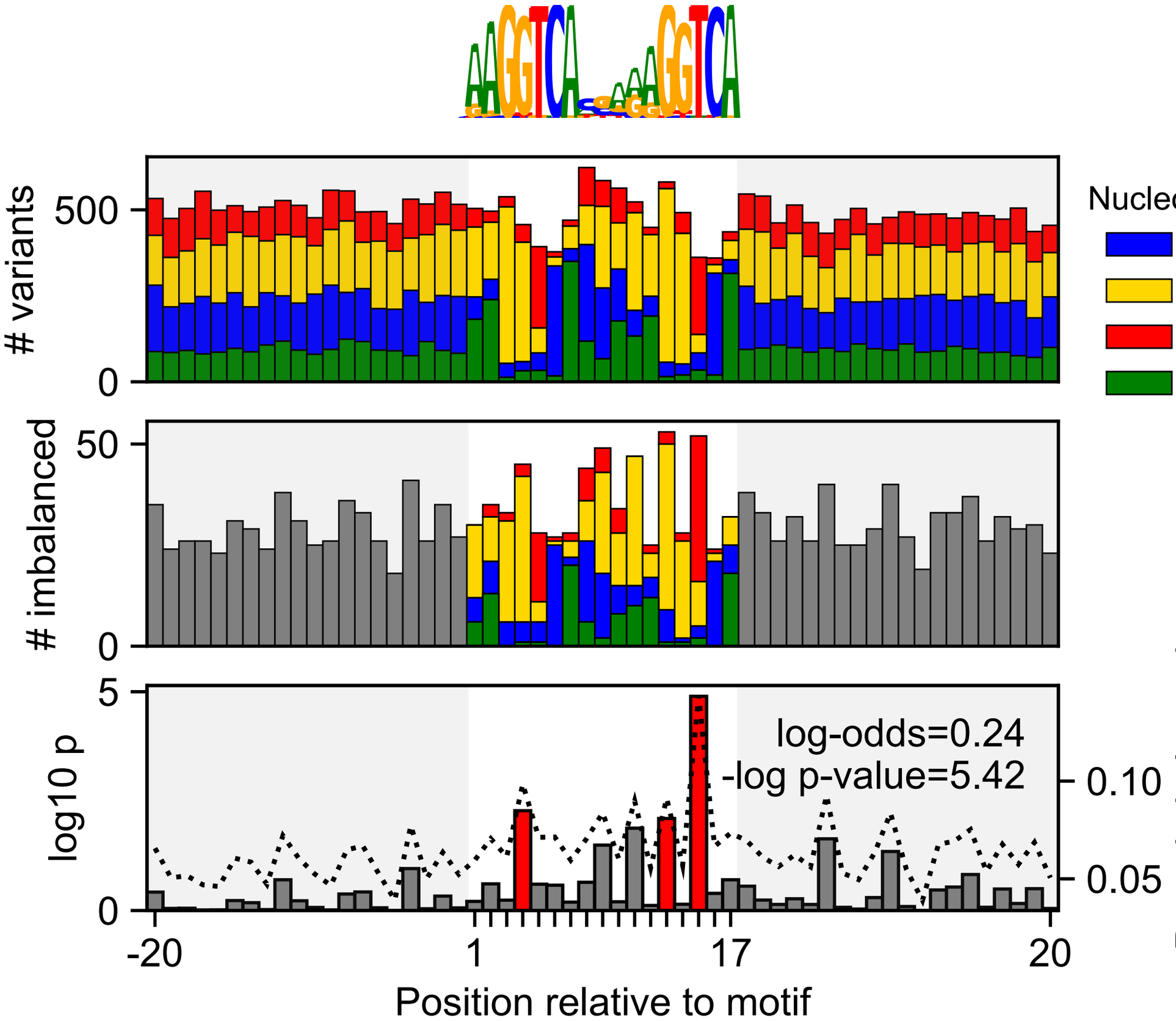
AAGGTCA
variants
500
0
Nucle
imbalanced
50
0
log10 p
5
0
log-odds=0.24
-log p-value=5.42
0.10
0.05
-20
1
17
20
Position relative to motif