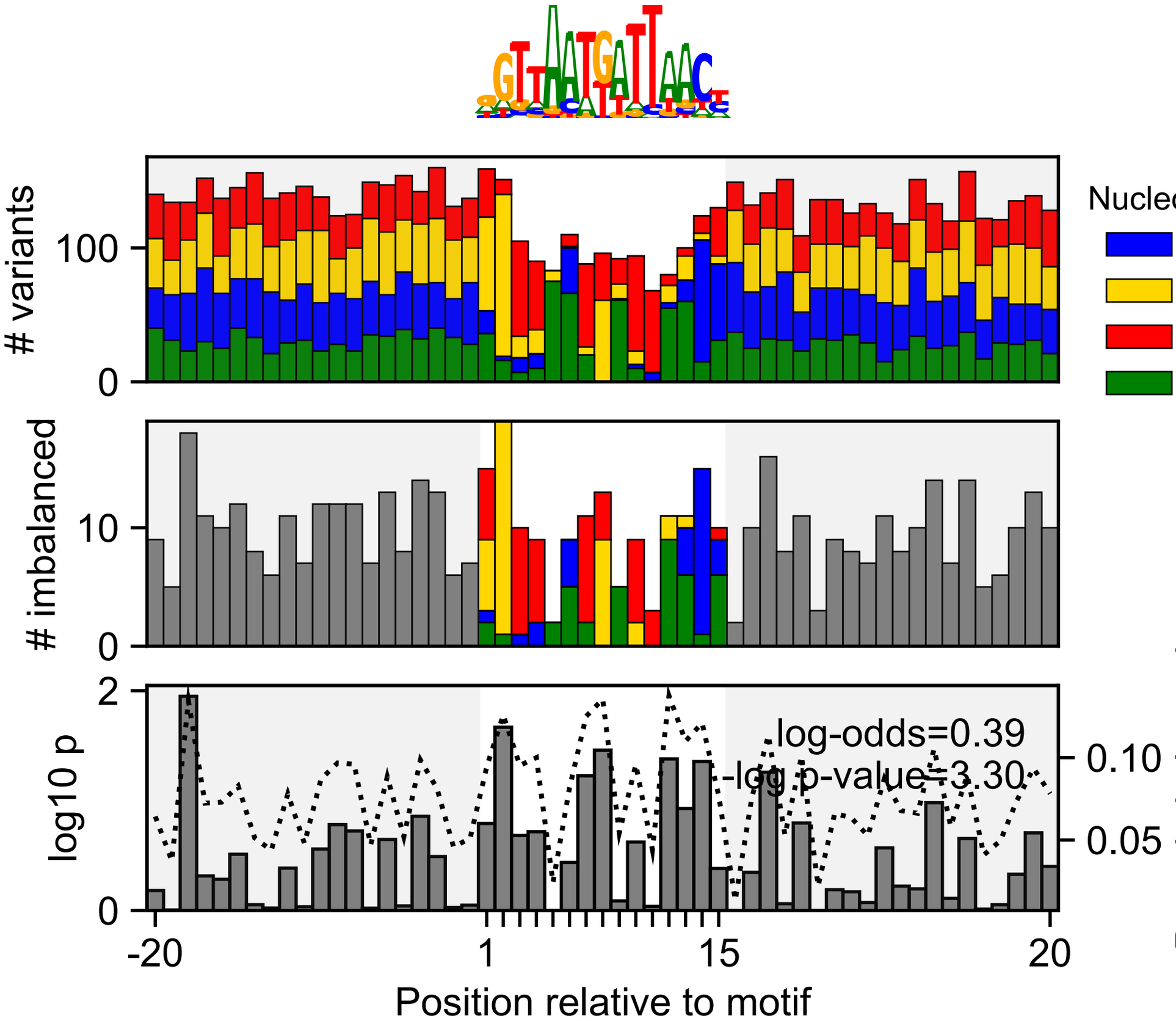
GTTAATGATTAAC
variants
100
0
Nucle
imbalanced
10
0
log10 p
2
0
log-odds=0.39
-log p-value=3.30
0.10
0.05
-20
1
15
20
Position relative to motif