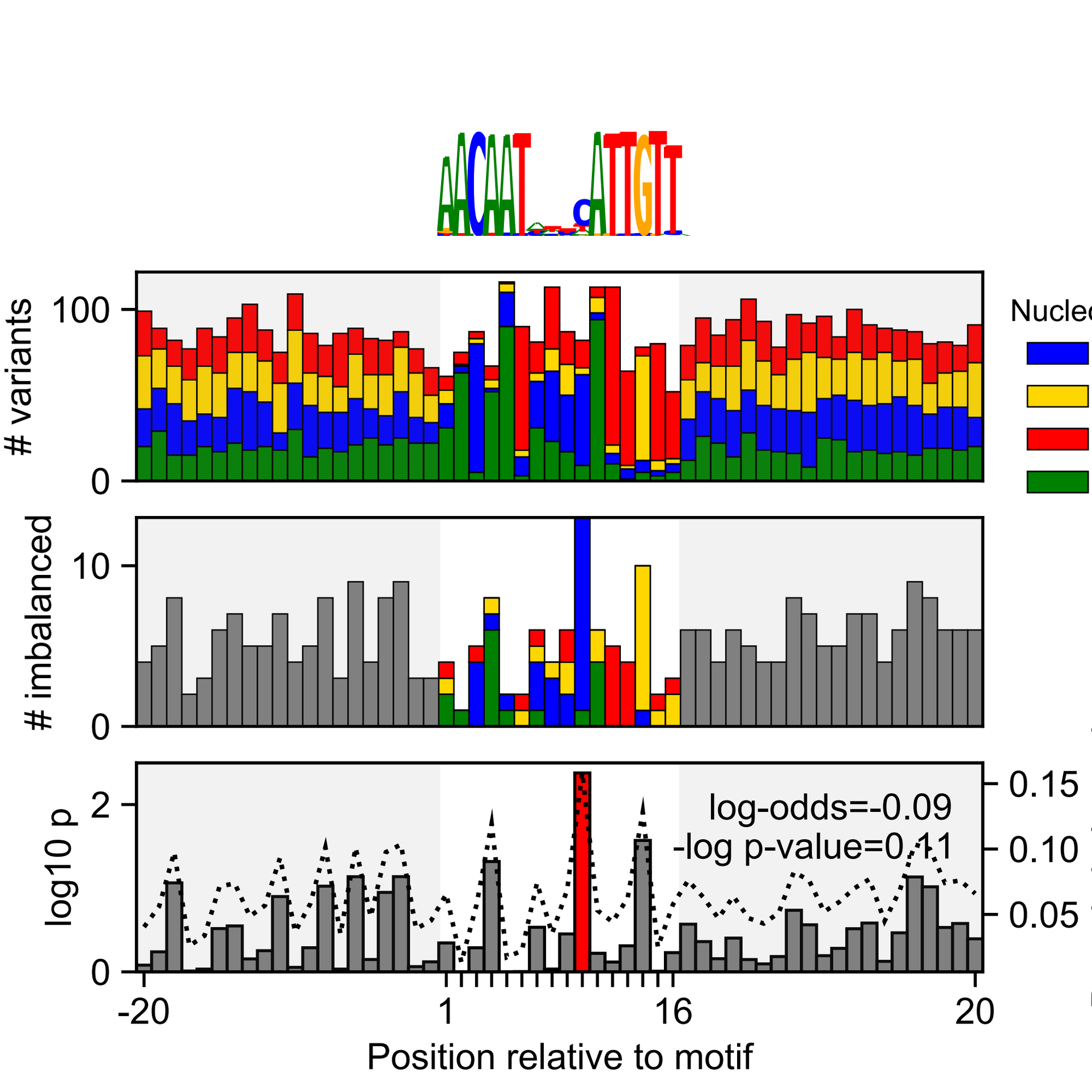
AACAAT_gATTGTT
# variants
100
0
Nucle
# imbalanced
10
0
log10 p
2
0
log-odds=-0.09
-log p-value=0.11
0.15
0.10
0.05
-20
1
16
20
Position relative to motif