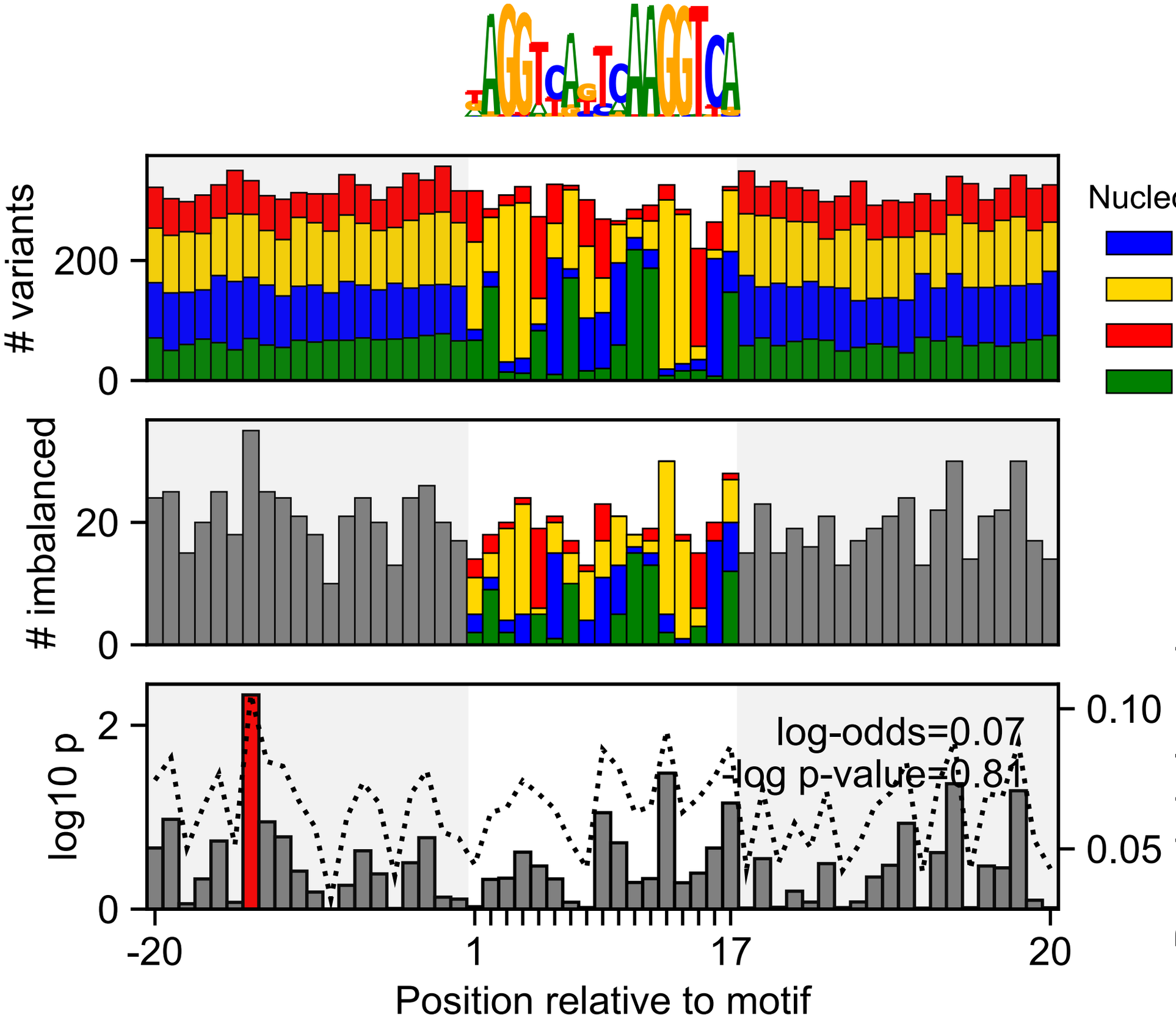
AGGTCATCAAGGTCA
variants
200
0
Nucle
imbalanced
20
0
log10 p
2
0
0.10
0.05
log-odds=0.07
-log p-value=0.81
-20
1
17
20
Position relative to motif