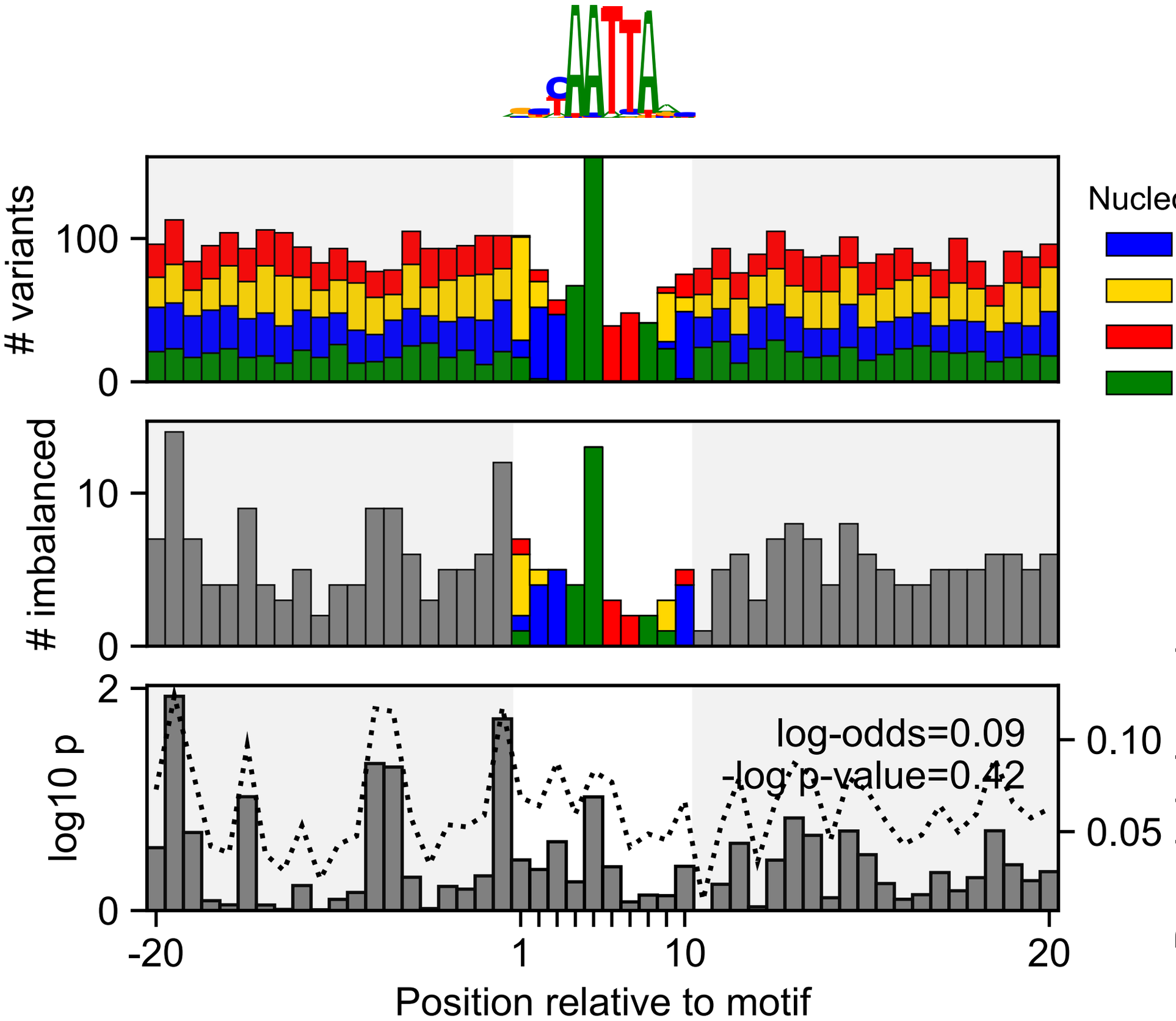
Nucle
# variants
100
0
# imbalanced
10
0
log10 p
2
0
log-odds=0.09
-log p-value=0.42
0.10
0.05
-20
1
10
20
Position relative to motif