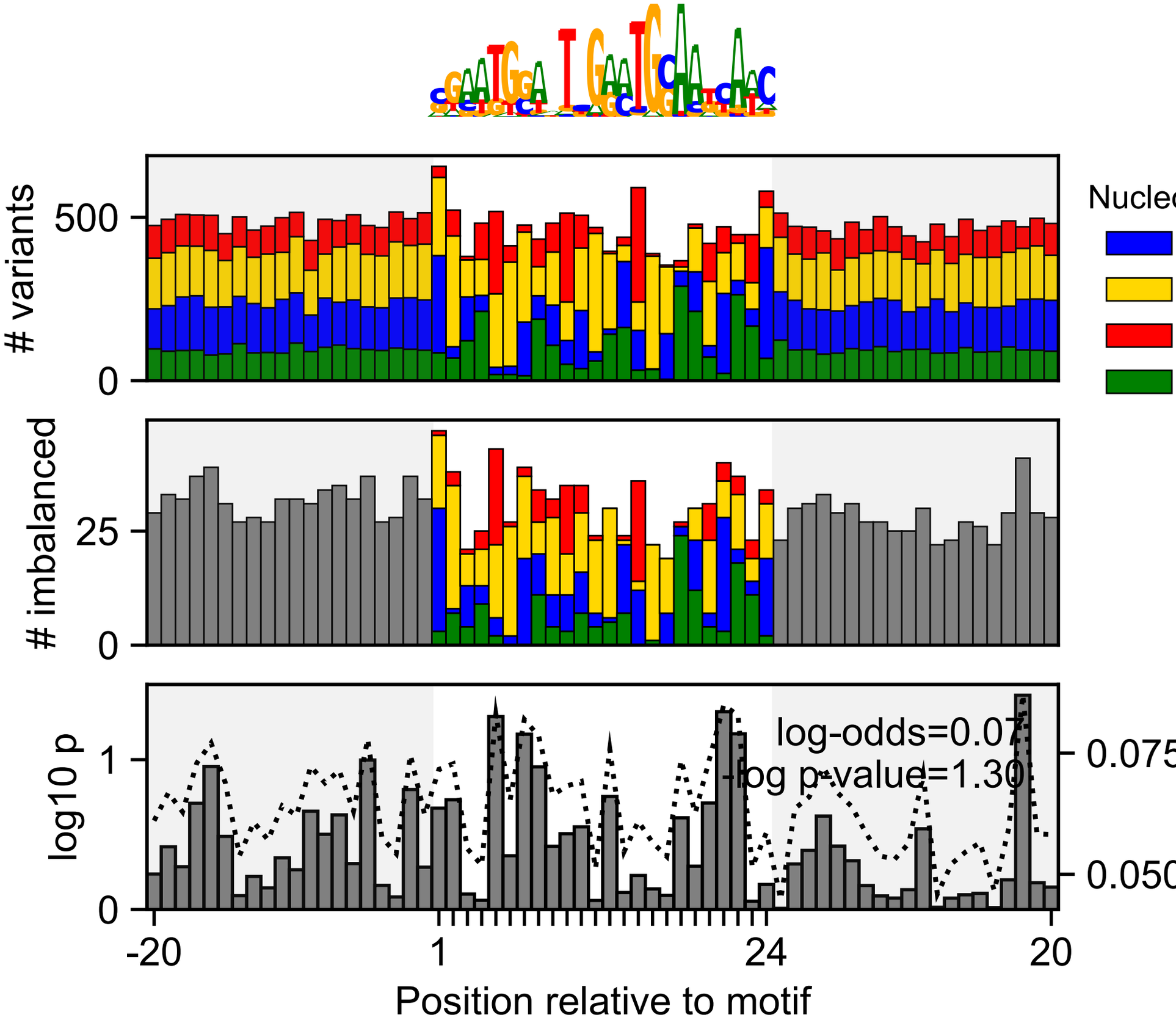
Nucle
# variants
500
0
# imbalanced
25
0
log10 p
1
0
log-odds=0.07
-log p-value=1.30
0.075
0.050
-20
1
24
20
Position relative to motif