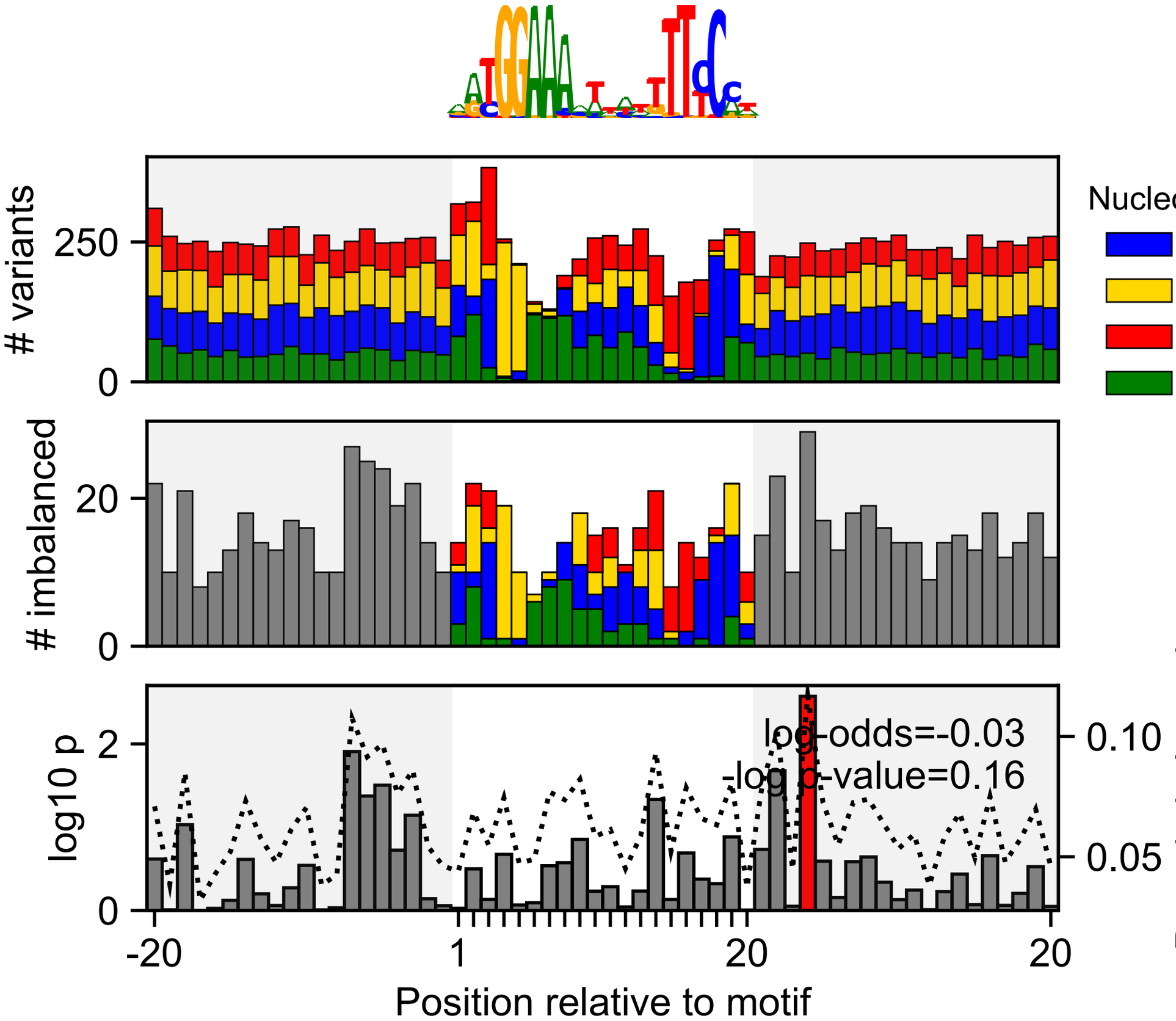
Nucle
variants
250
0
imbalanced
20
0
log10 p
2
0
log-odds=-0.03
-log p-value=0.16
0.10
0.05
-20
1
20
20
Position relative to motif