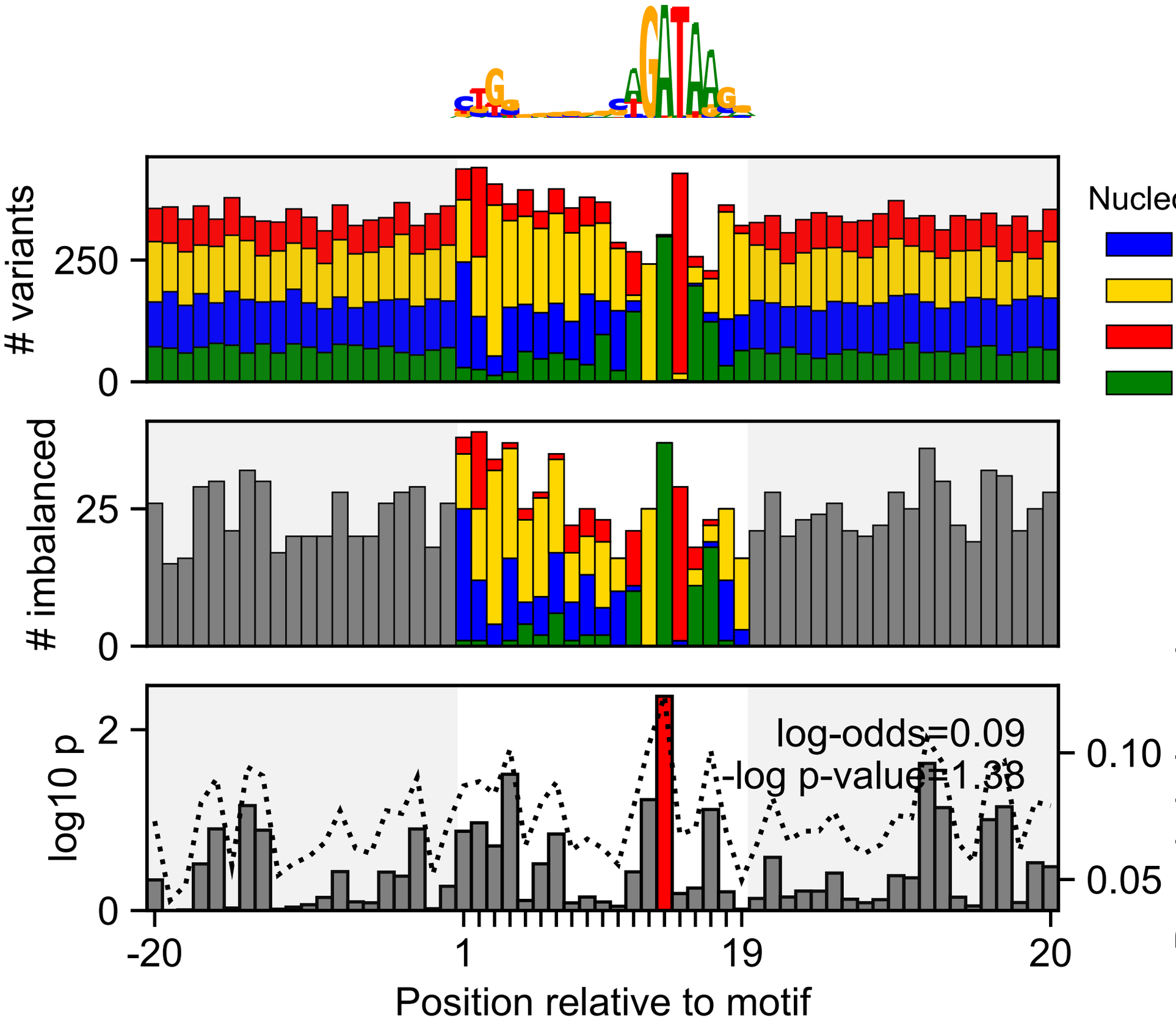
Nucle
variants
250
0
imbalanced
25
0
log10 p
2
0
log-odds=0.09
-log p-value=1.38
0.10
0.05
-20
1
19
20
Position relative to motif