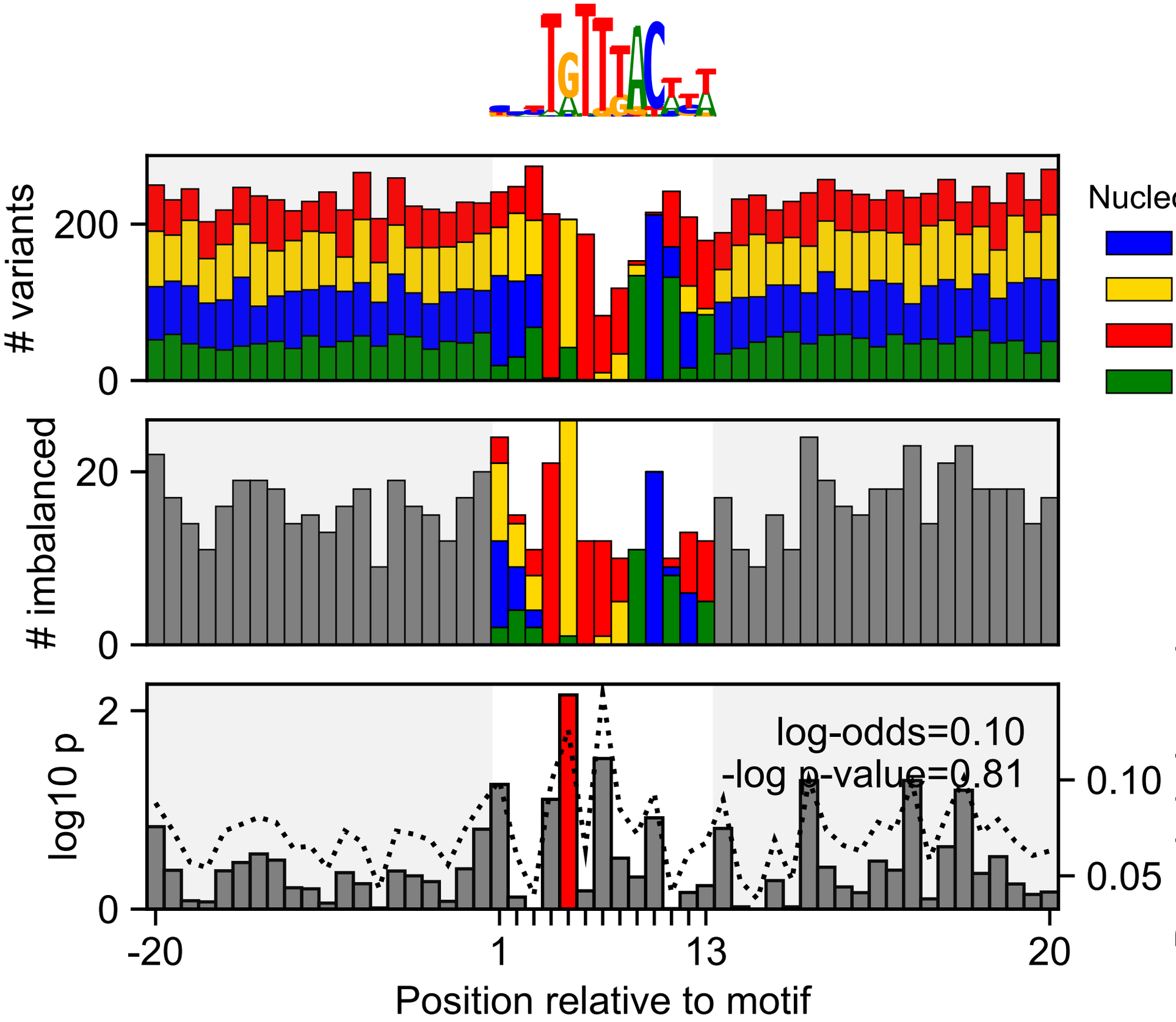
Nucle
variants
200
0
imbalanced
20
0
log10 p
2
0
log-odds=0.10
-log p-value=0.81
0.10
0.05
-20
1
13
20
Position relative to motif