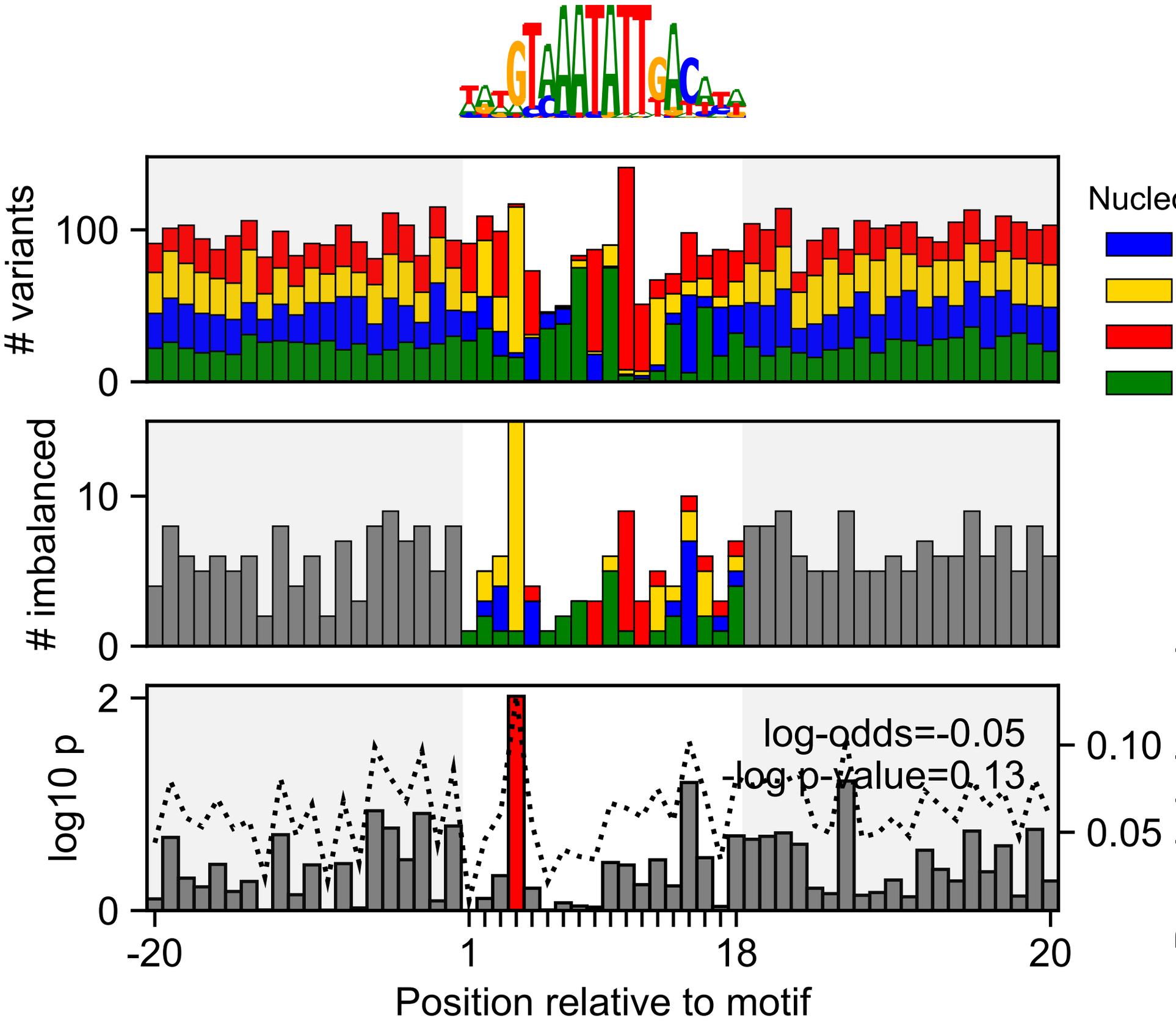
Nucle
variants
100
0
imbalanced
10
0
log10 p
2
0
log-odds=-0.05
-log p-value=0.13
0.10
0.05
-20
1
18
20
Position relative to motif