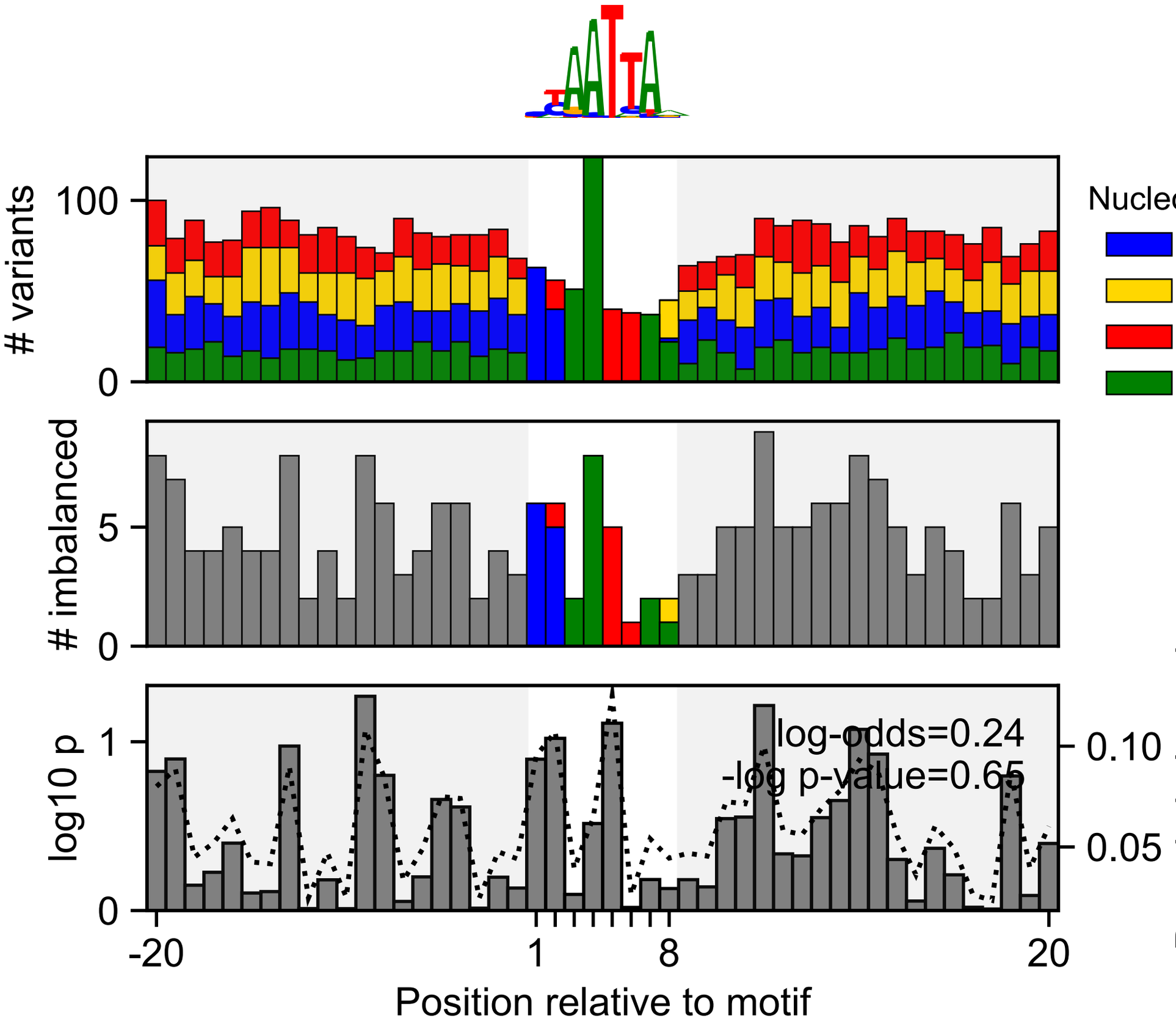
Nucle
variants
100
0
imbalanced
5
0
log10 p
1
0
log-odds=0.24
-log p-value=0.65
0.10
0.05
-20
1
8
20
Position relative to motif