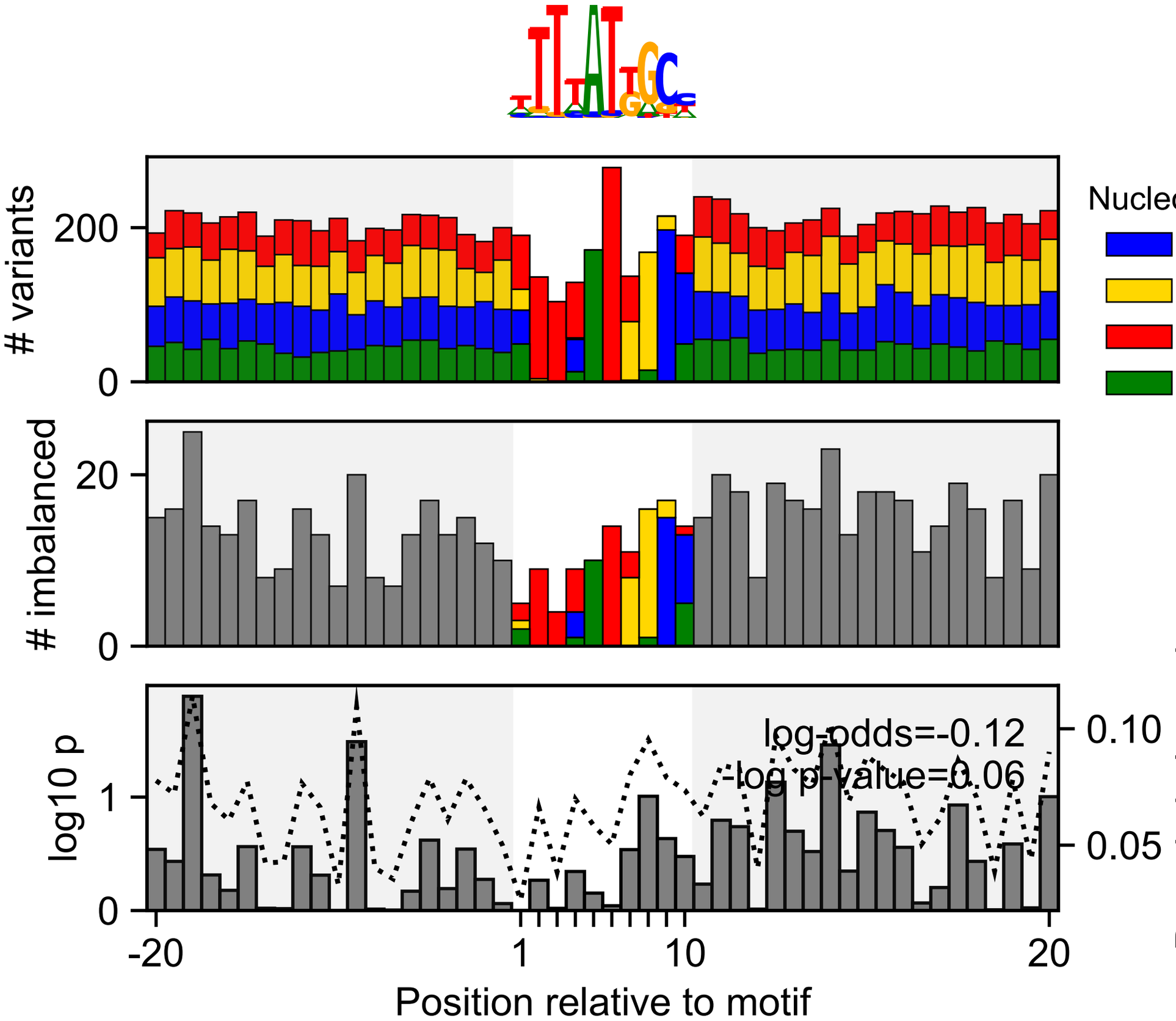
Nucle
variants
200
0
imbalanced
20
0
log10 p
1
0
log-odds=-0.12
-log p-value=0.06
0.10
0.05
-20
1
10
20
Position relative to motif