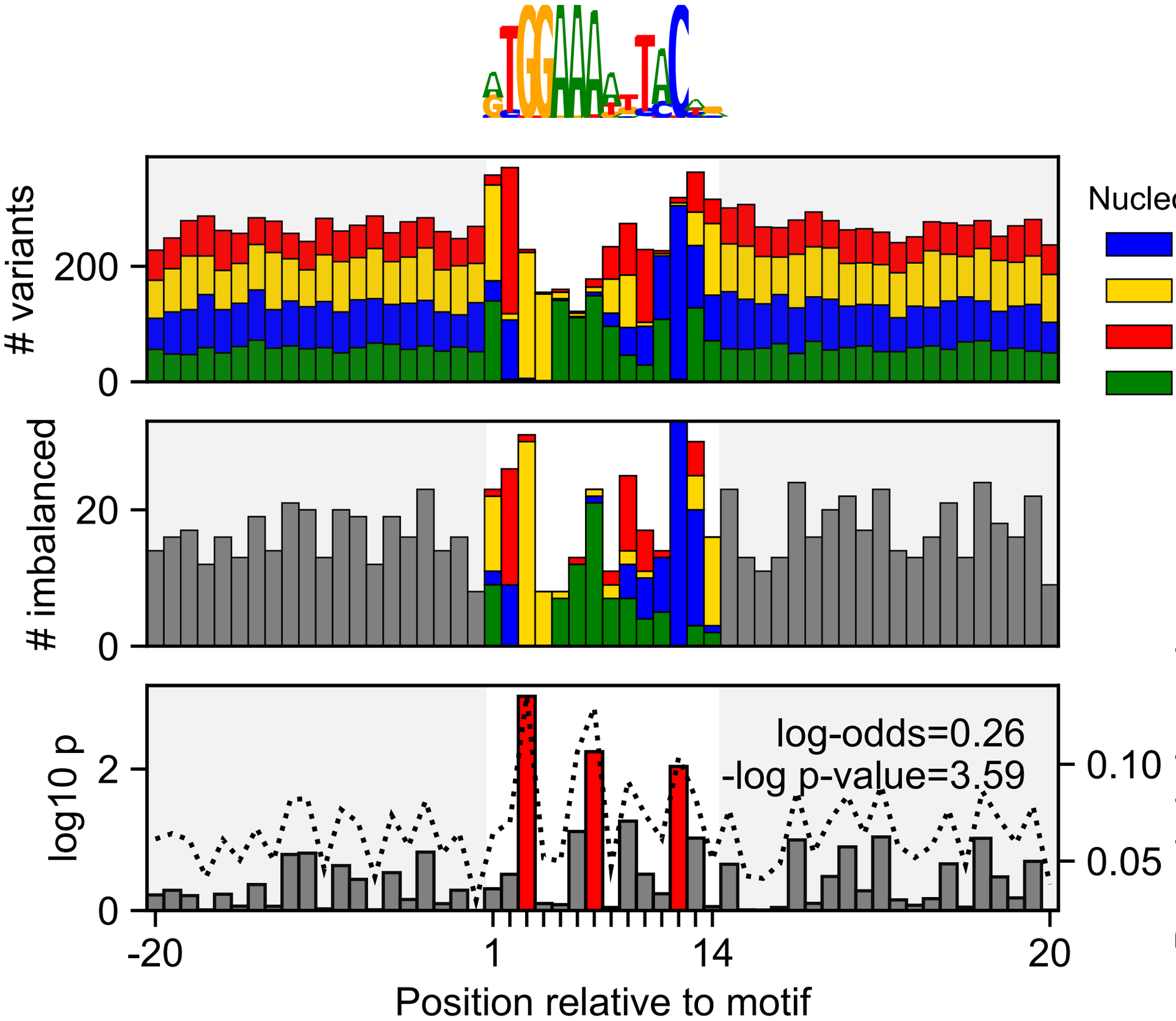
Nucle
variants
200
0
imbalanced
20
0
log10 p
2
0
log-odds=0.26
-log p-value=3.59
0.10
0.05
-20
1
14
20
Position relative to motif