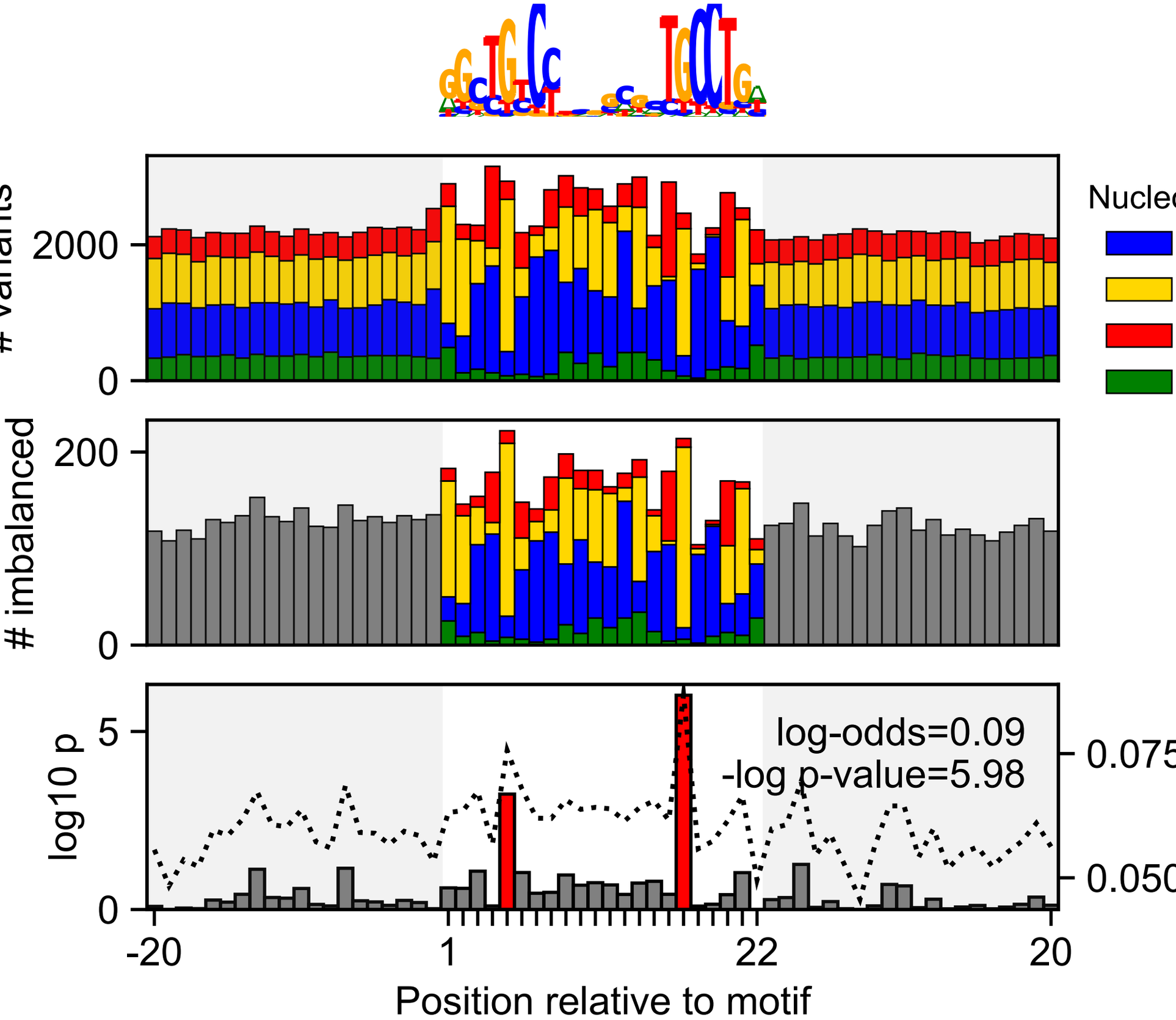
Nucle
variants
2000
0
imbalanced
200
0
log10 p
5
0
log-odds=0.09
-log p-value=5.98
0.075
0.050
-20
1
22
20
Position relative to motif