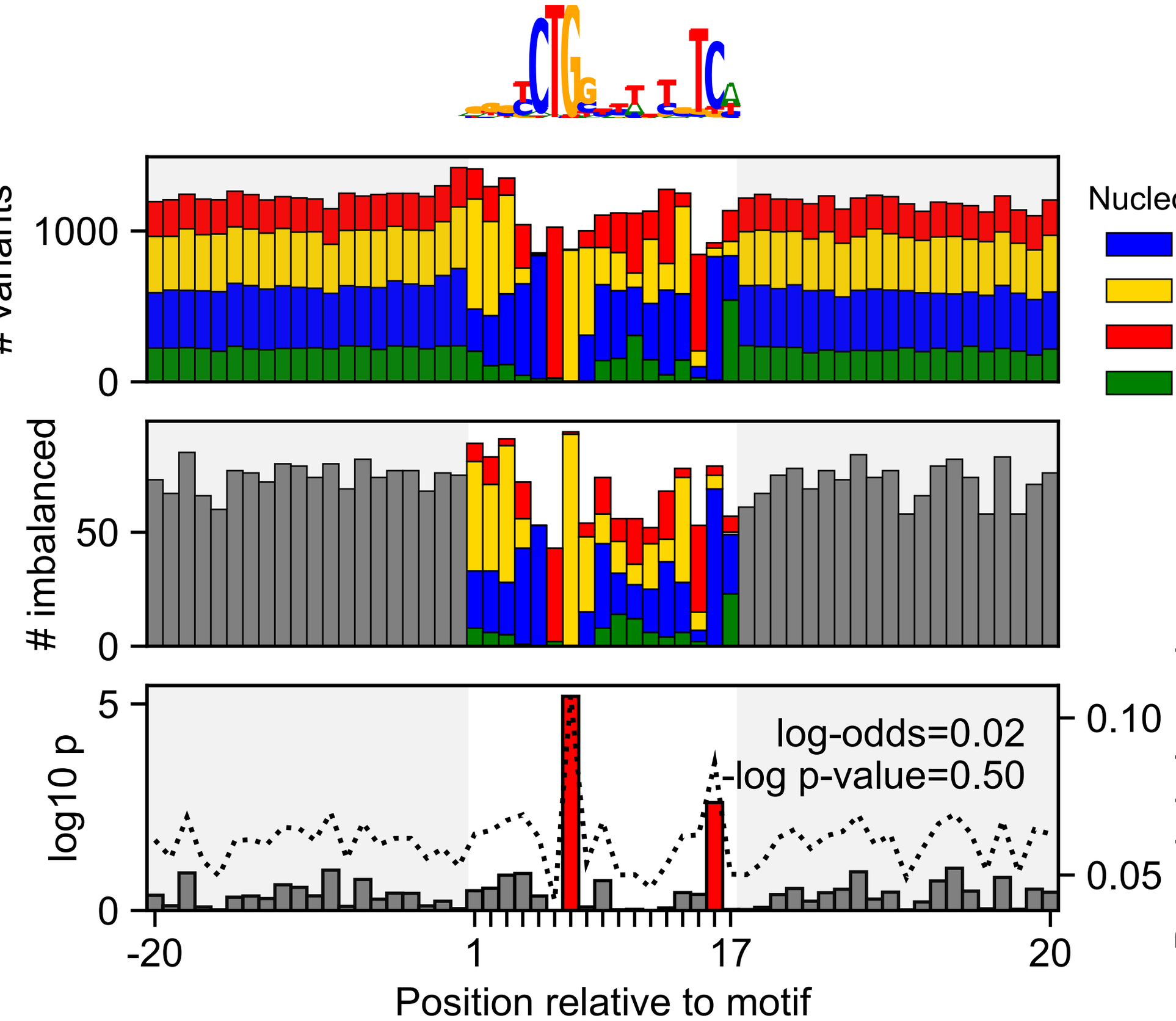
Nucle
variants
1000
0
imbalanced
50
0
log10 p
5
0
0.10
0.05
log-odds=0.02
-log p-value=0.50
-20
1
17
20
Position relative to motif