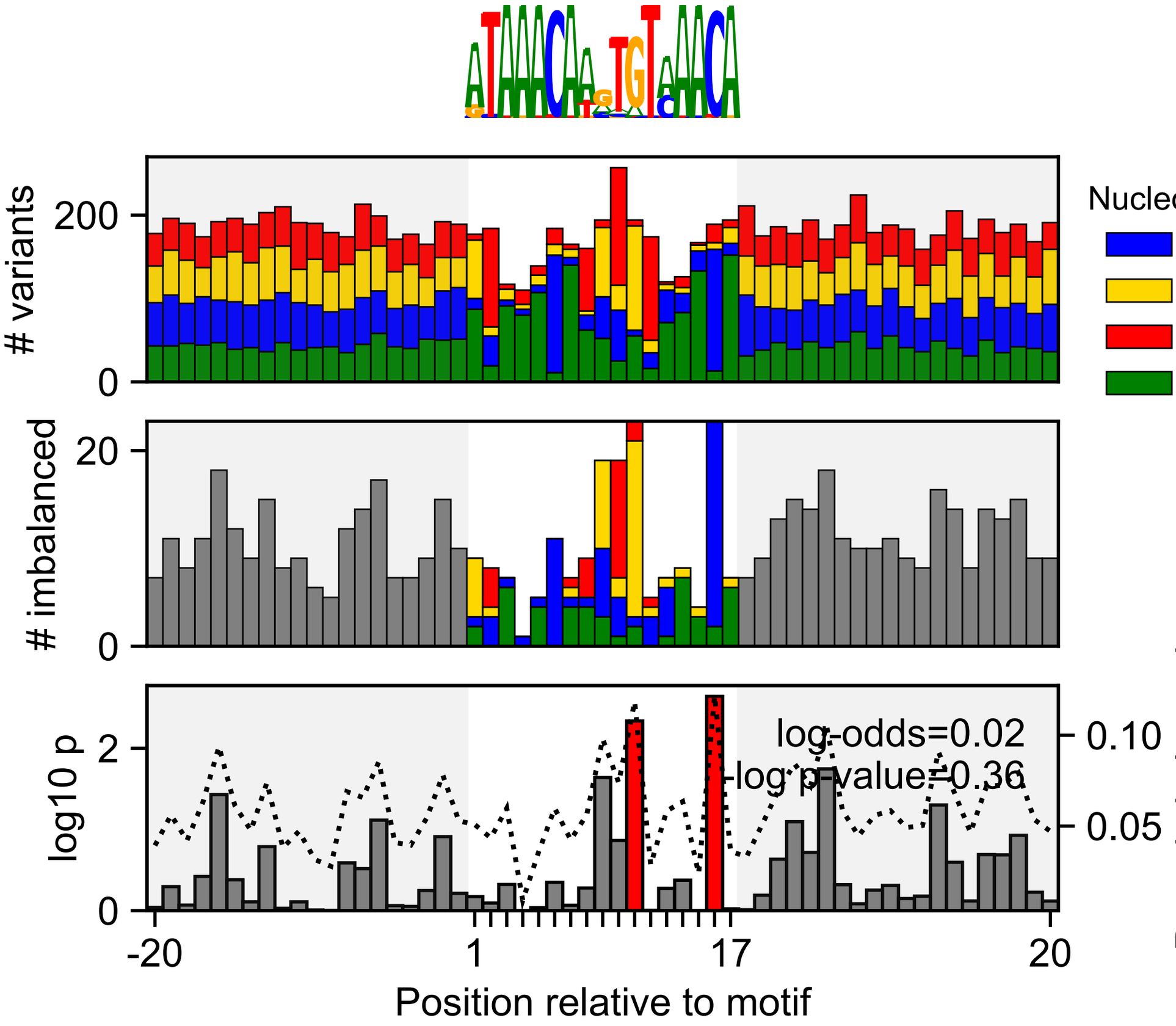
Nucle
variants
200
0
imbalanced
20
0
log10 p
2
0
log-odds=0.02
log p-value=-0.36
0.10
0.05
-20
1
17
20
Position relative to motif