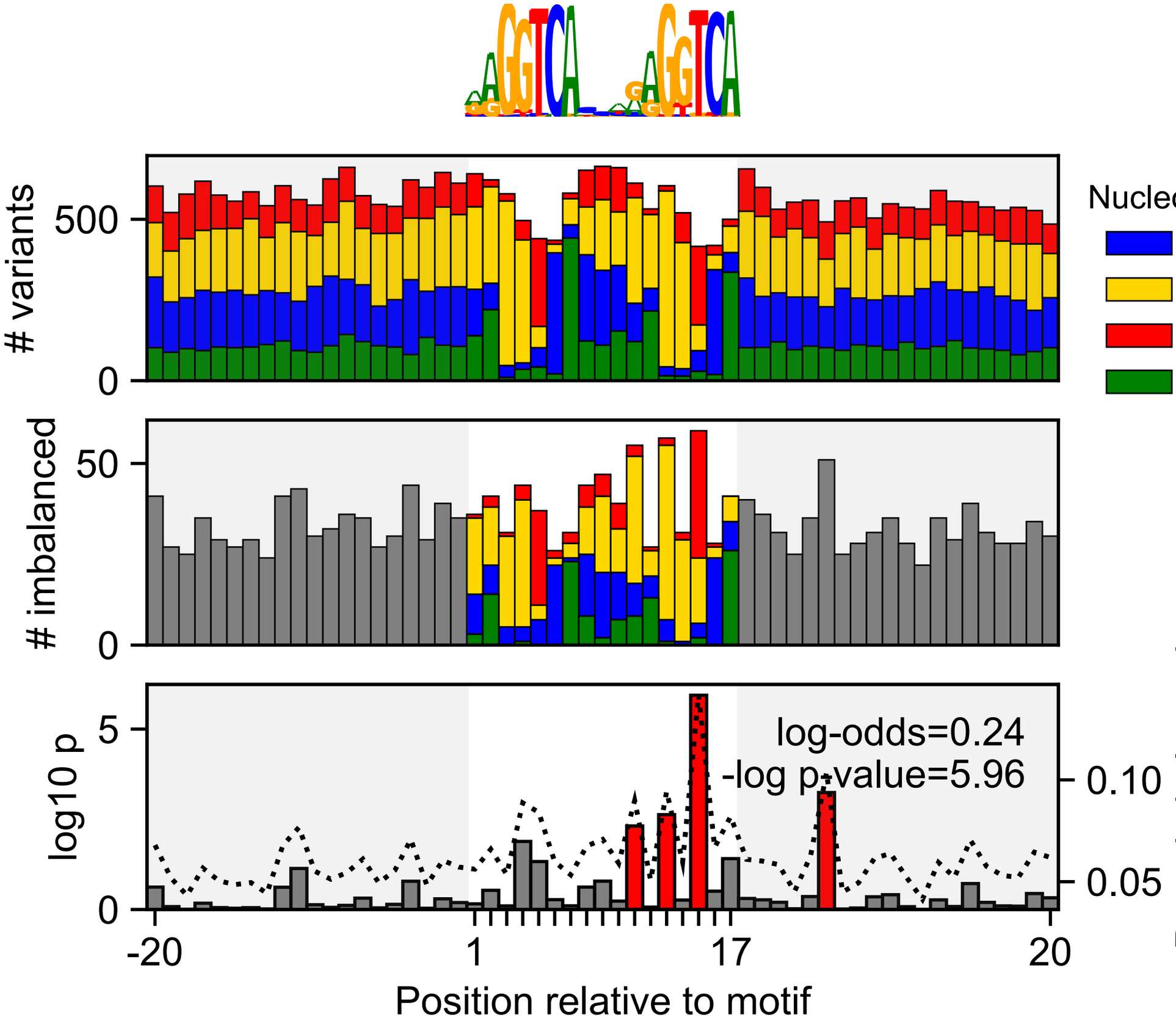
AGGTCA_AGGTCA
variants
500
0
Nucle
imbalanced
50
0
log10 p
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log-odds=0.24
-log p-value=5.96
0.10
0.05
-20
1
17
20
Position relative to motif