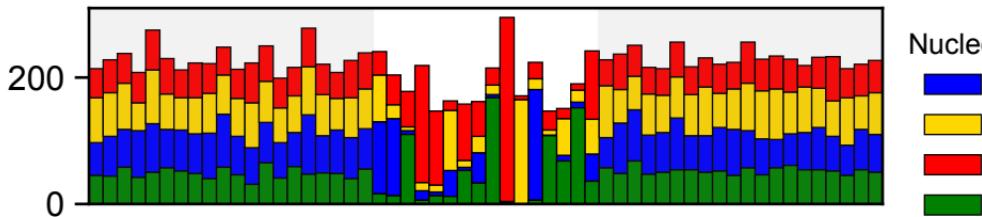


variants



Nucleotide

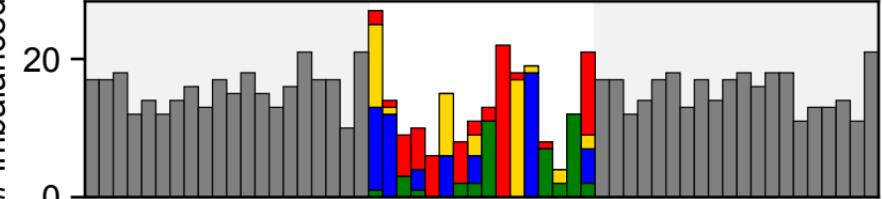
C

G

T

A

imbalanced

 $\log_{10} p$ 

Position relative to motif