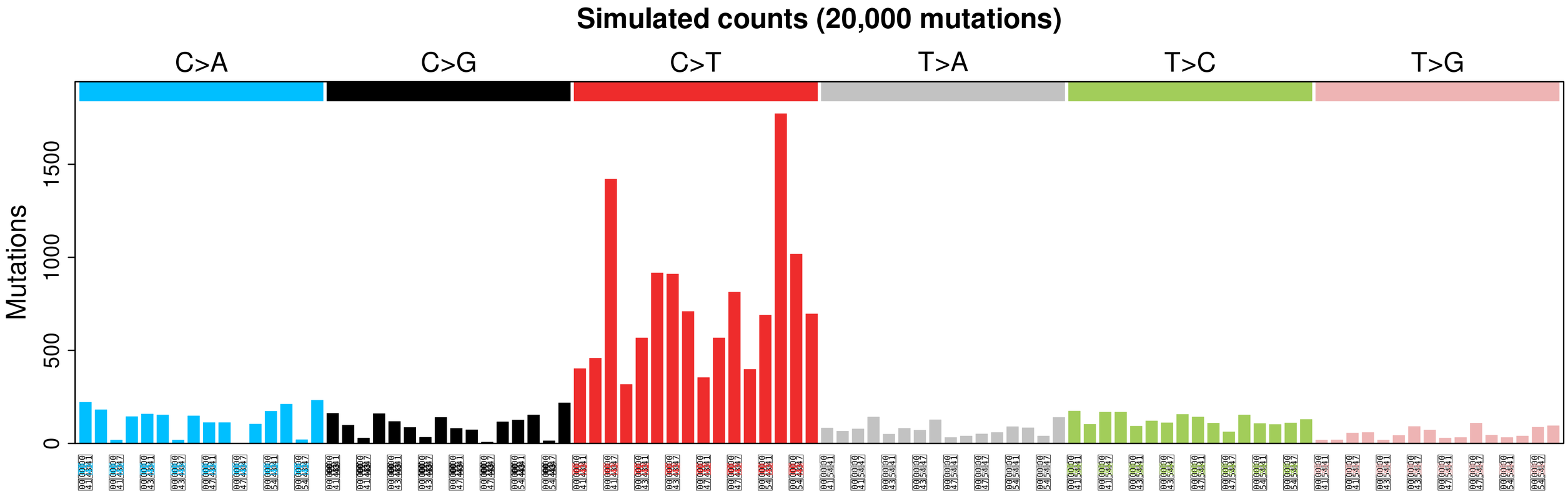
Simulated counts (20,000 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
500
1000
1500