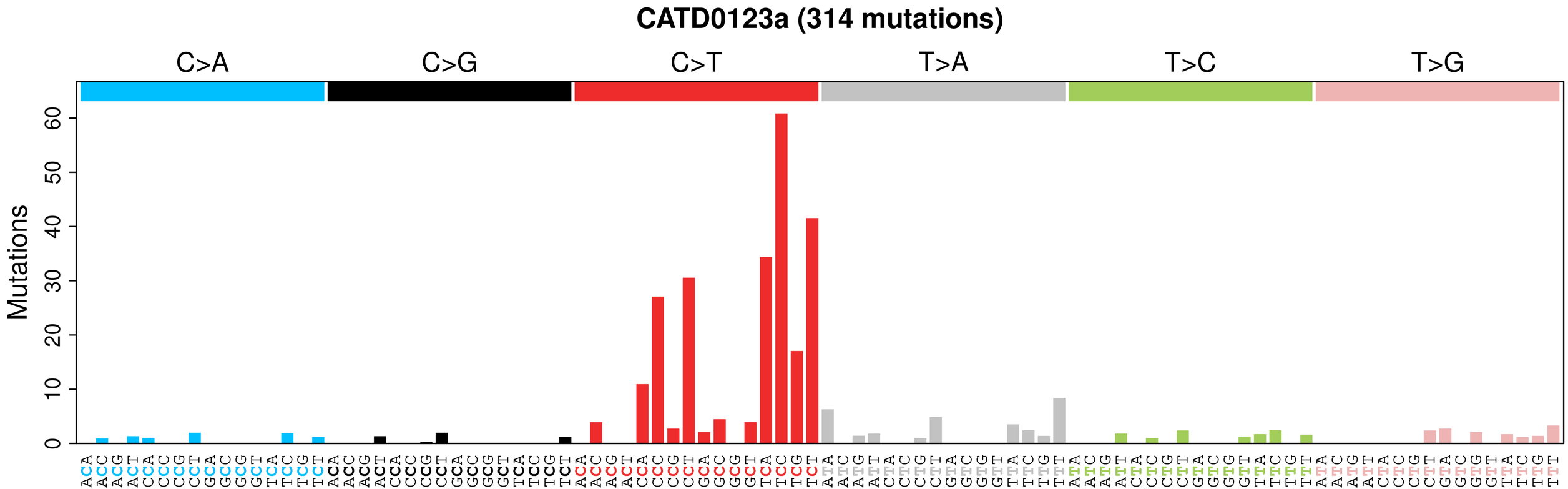
CATD0123a (314 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
10
20
30
40
50
60