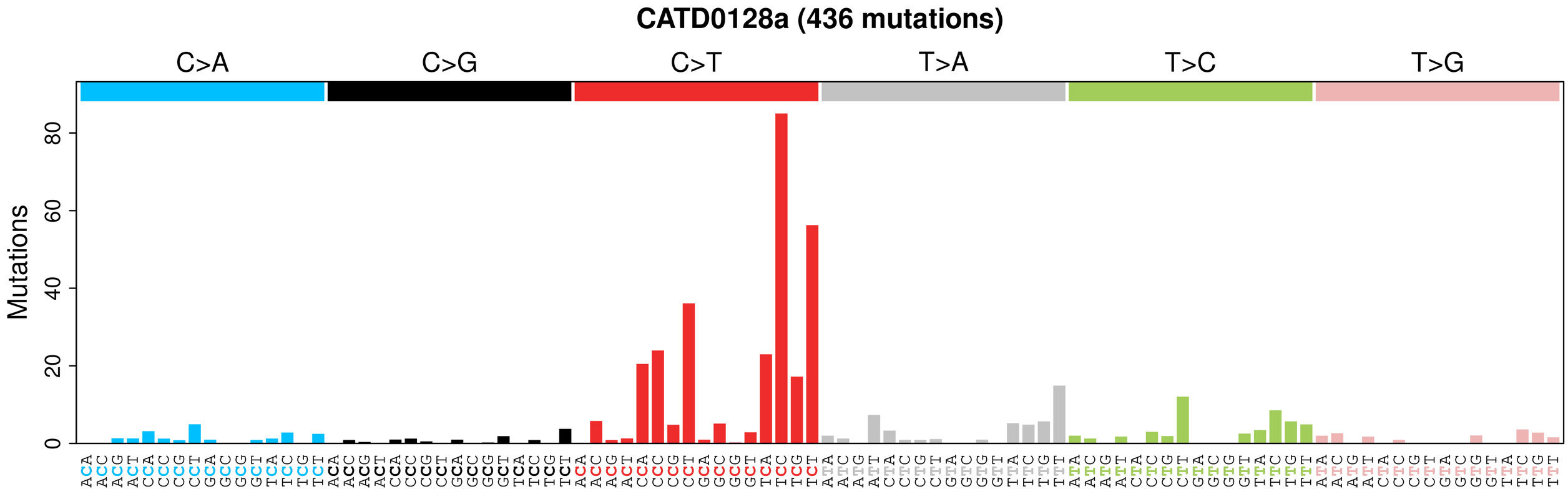
CATD0128a (436 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
20
40
60
80