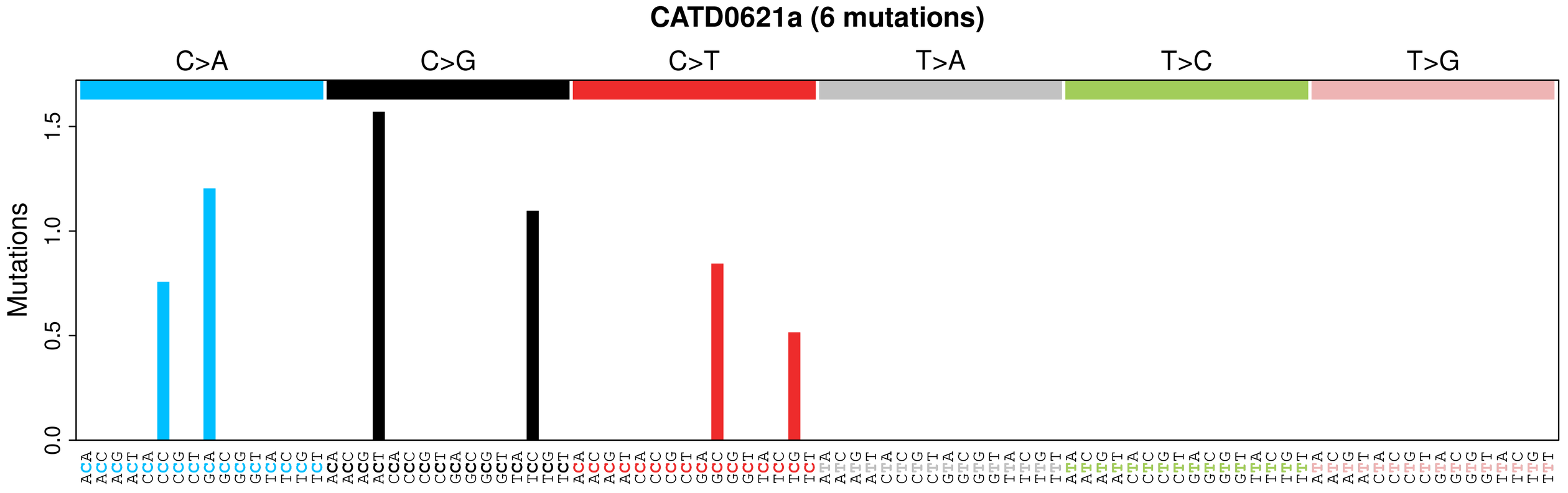
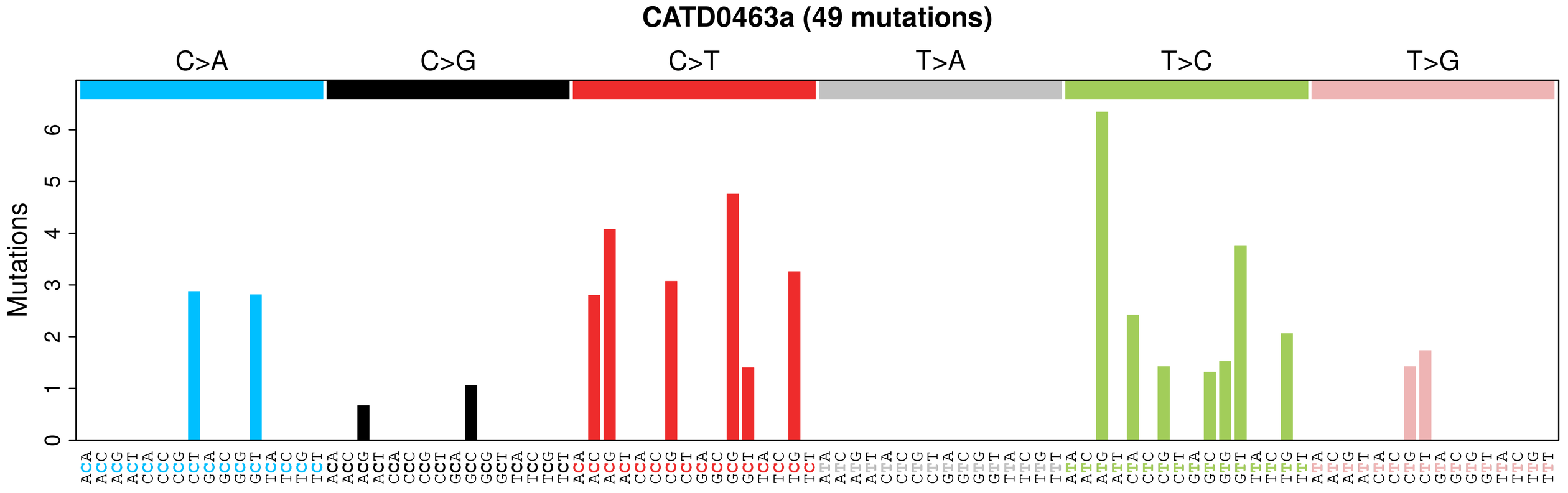
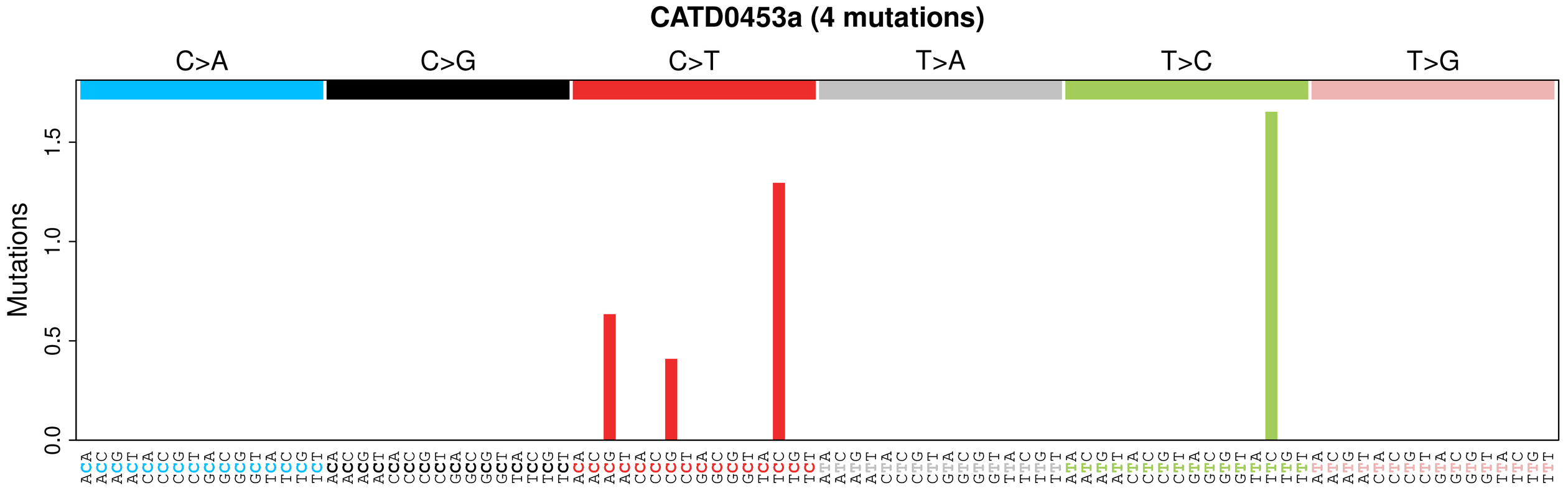
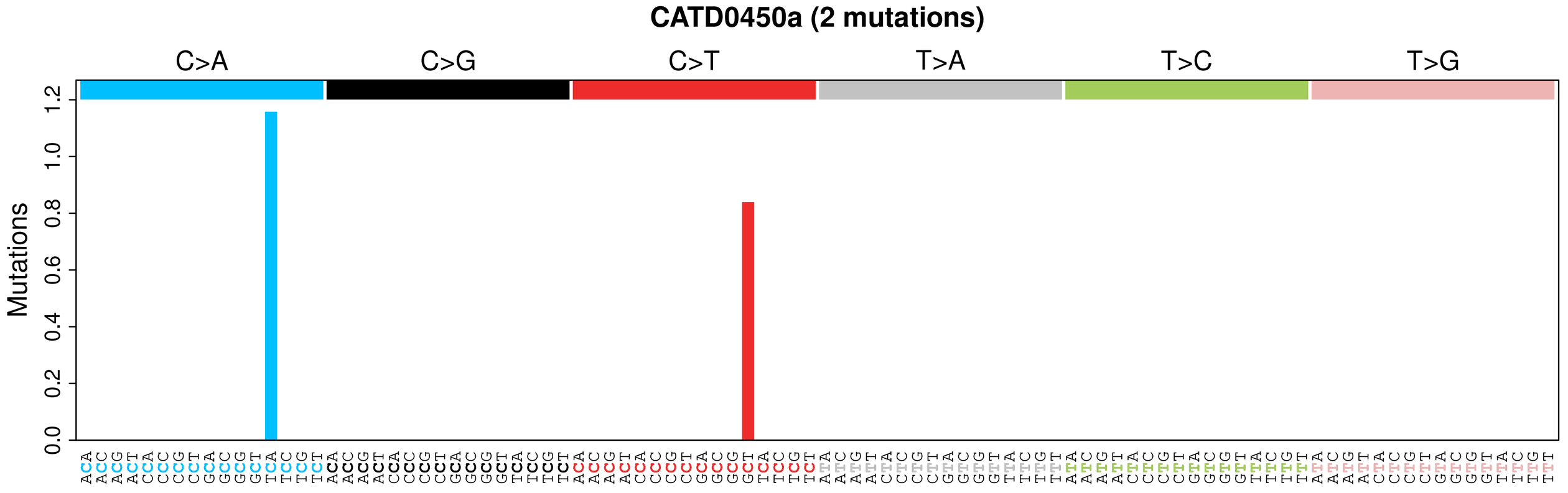
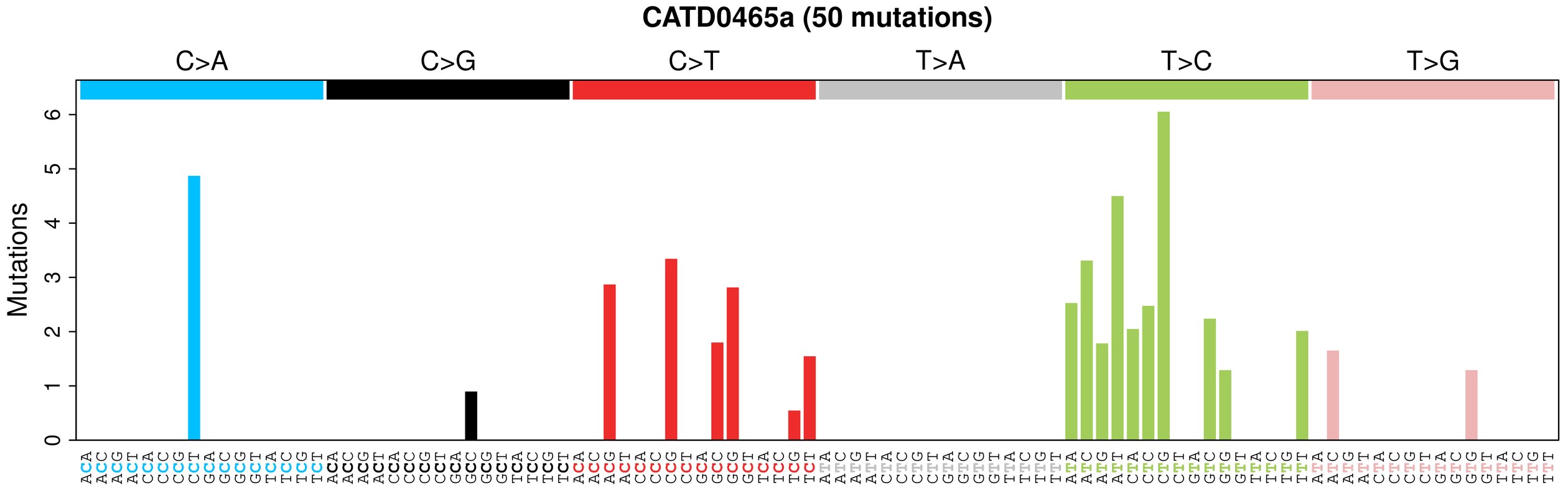
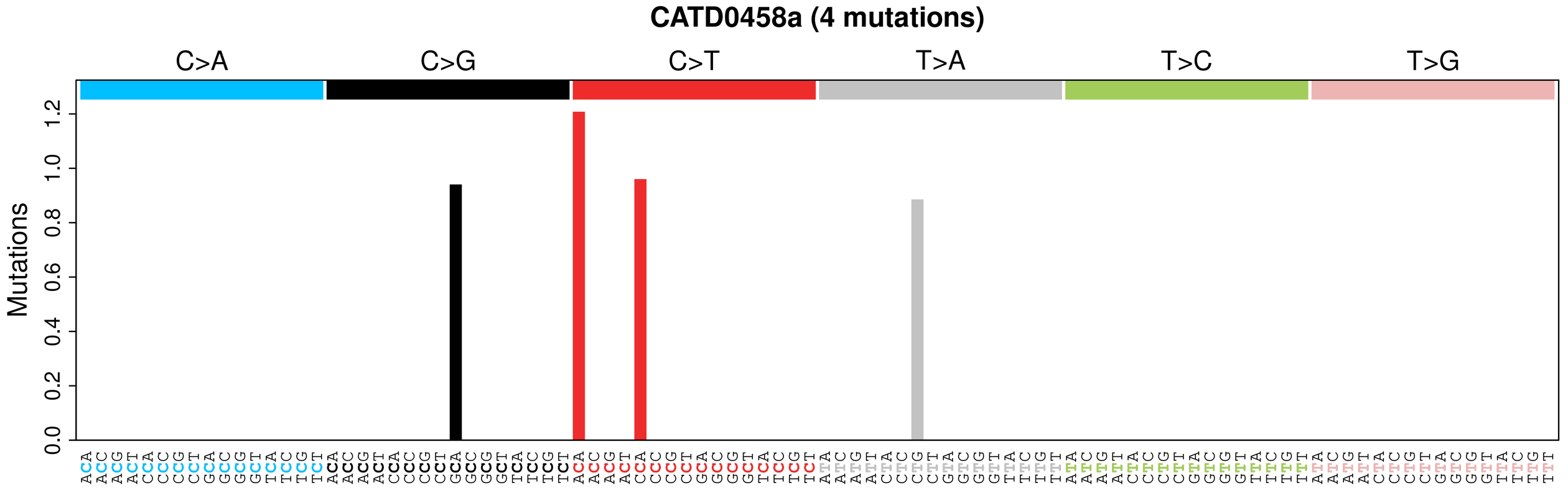

CATD0621a (6 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

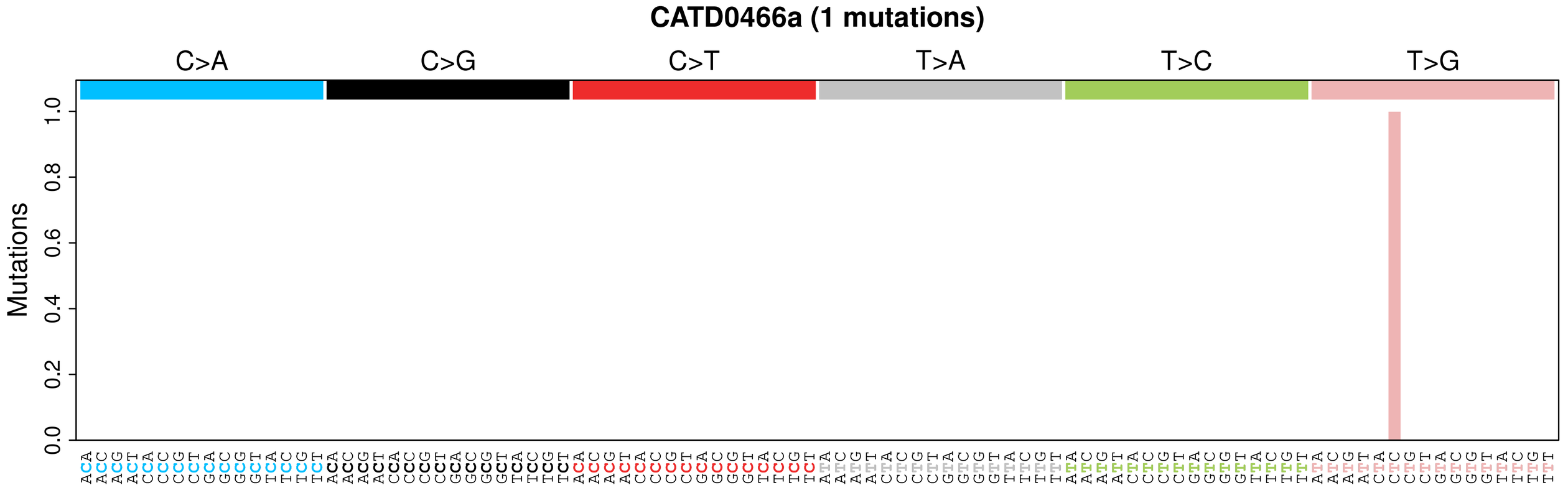

CATD0463a (49 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
1
2
3
4
5
6

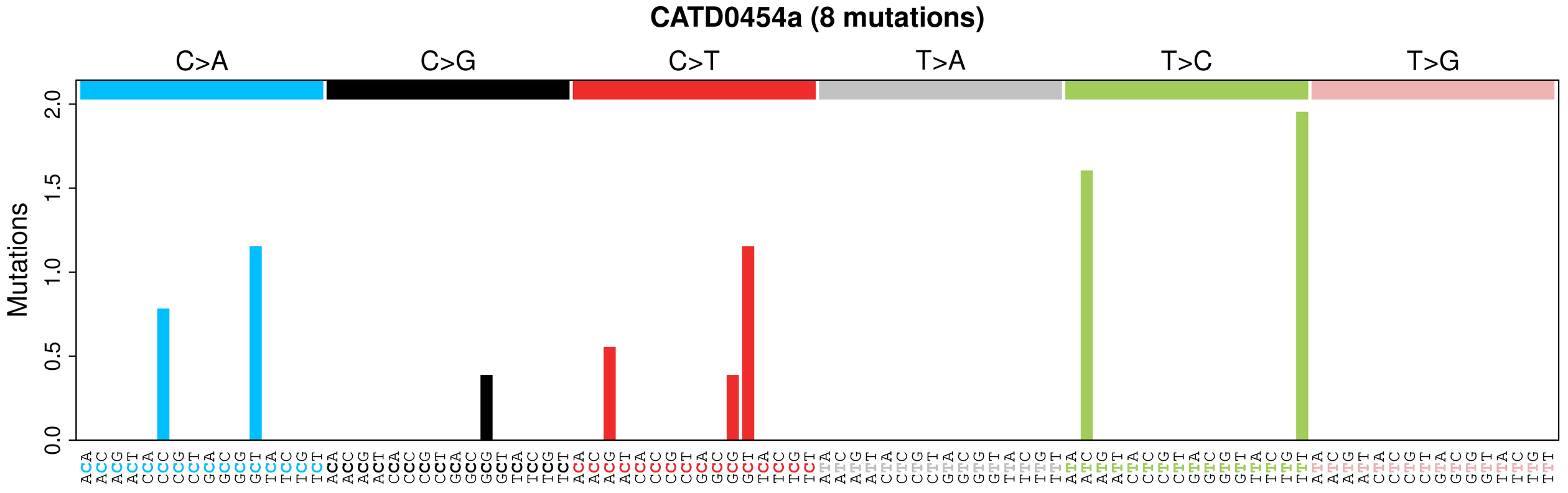

CATD0453a (4 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

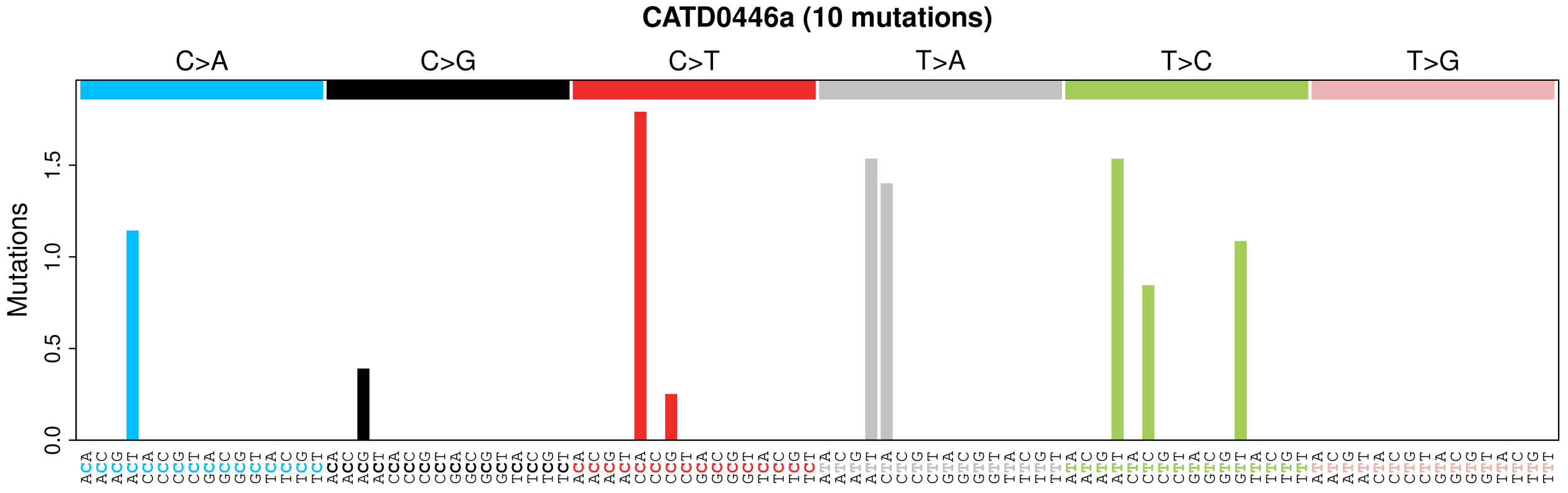

CATD0450a (2 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0
1.2

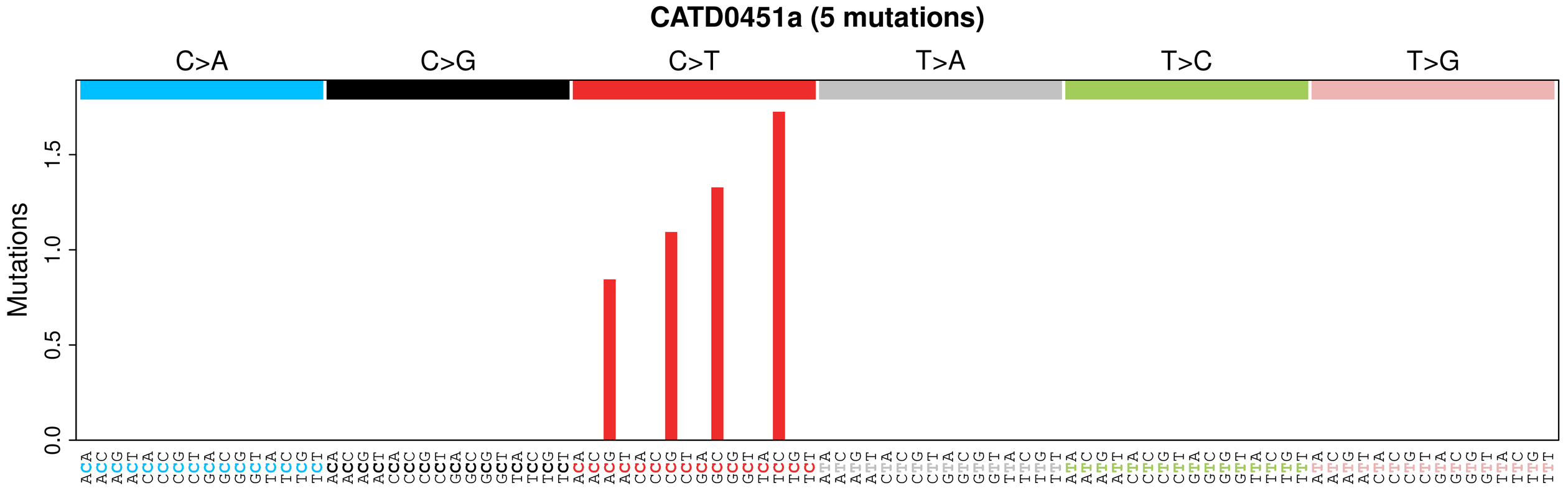

CATD0465a (50 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
1
2
3
4
5
6

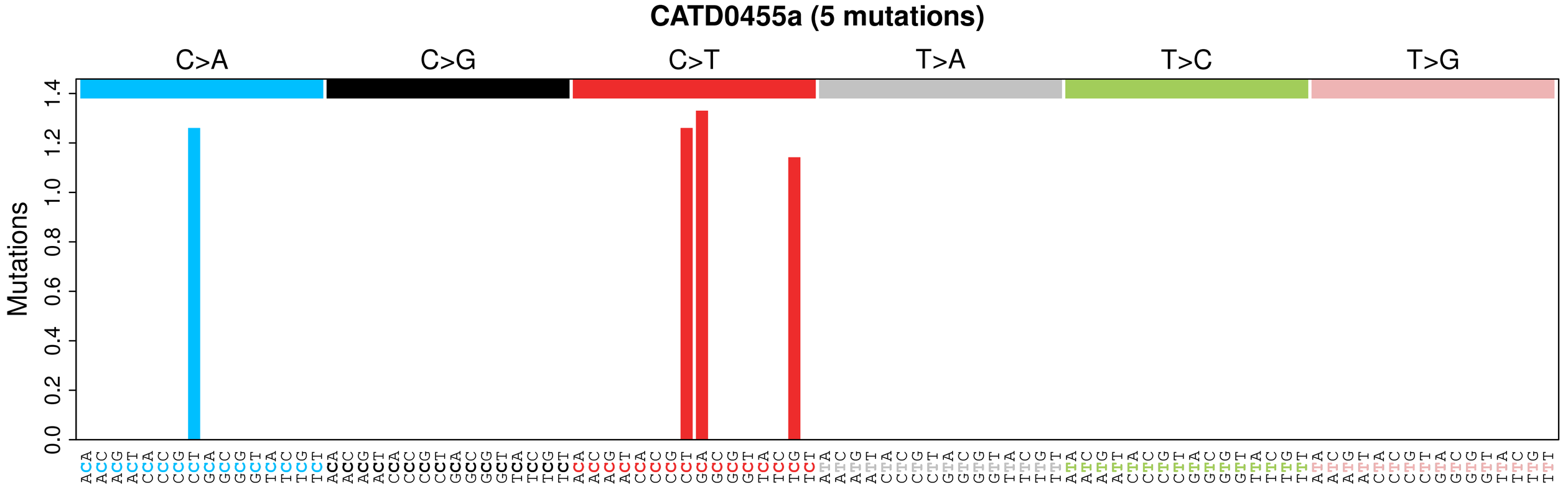

CATD0458a (4 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0
1.2

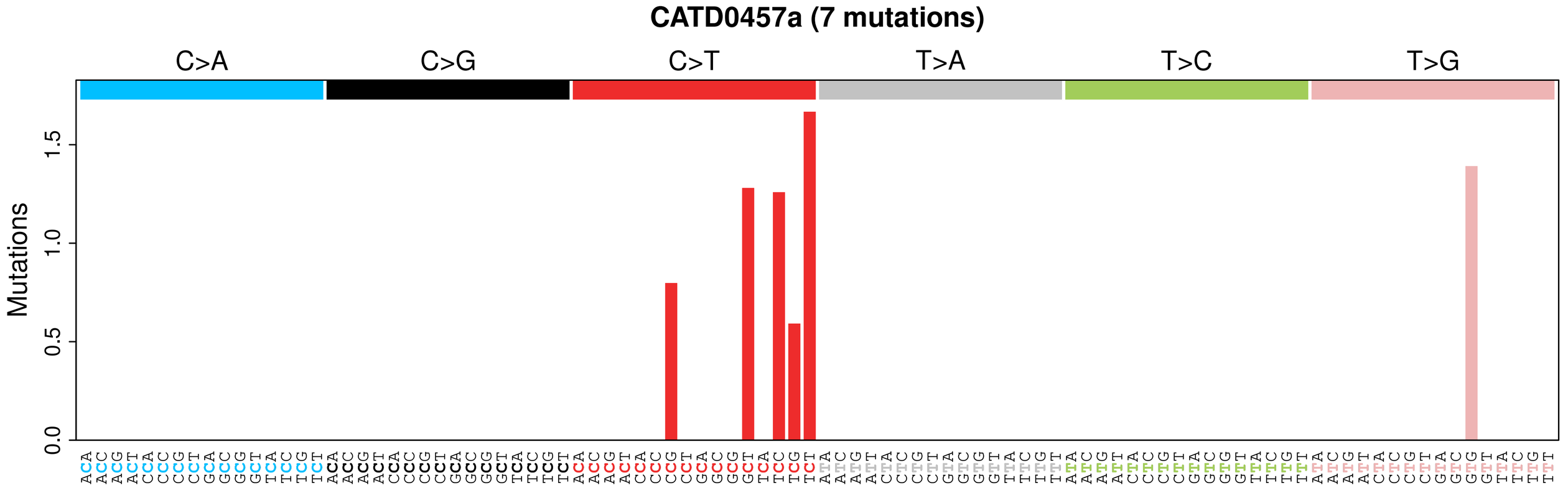

CATD0466a (1 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0

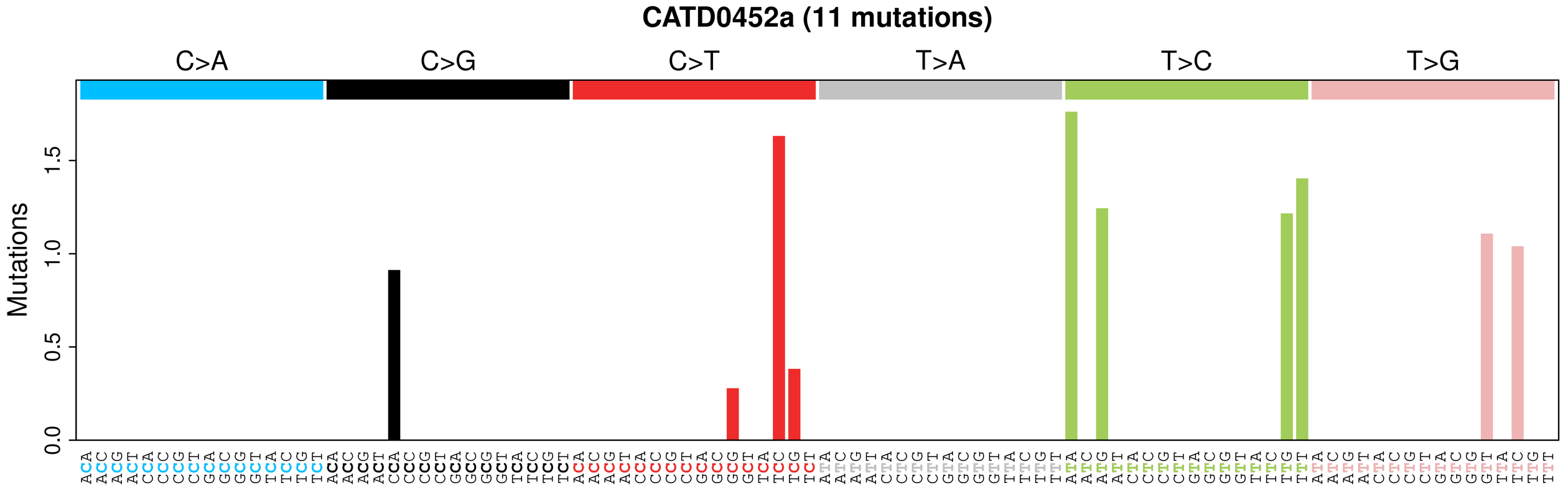

CATD0454a (8 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5
2.0

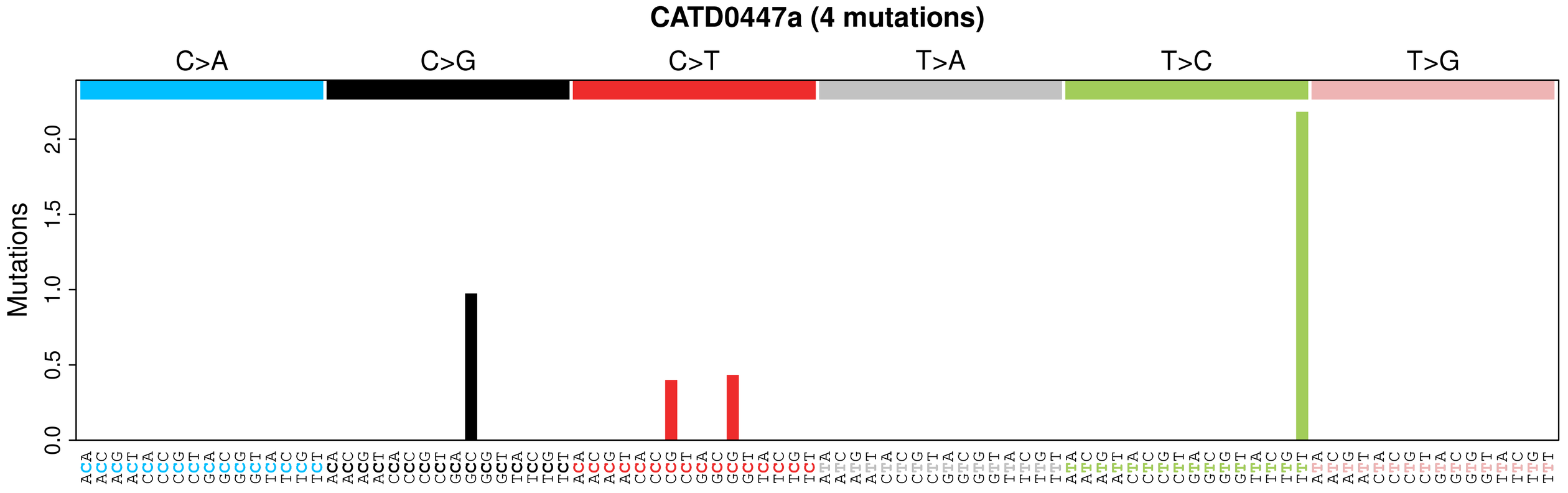

CATD0446a (10 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

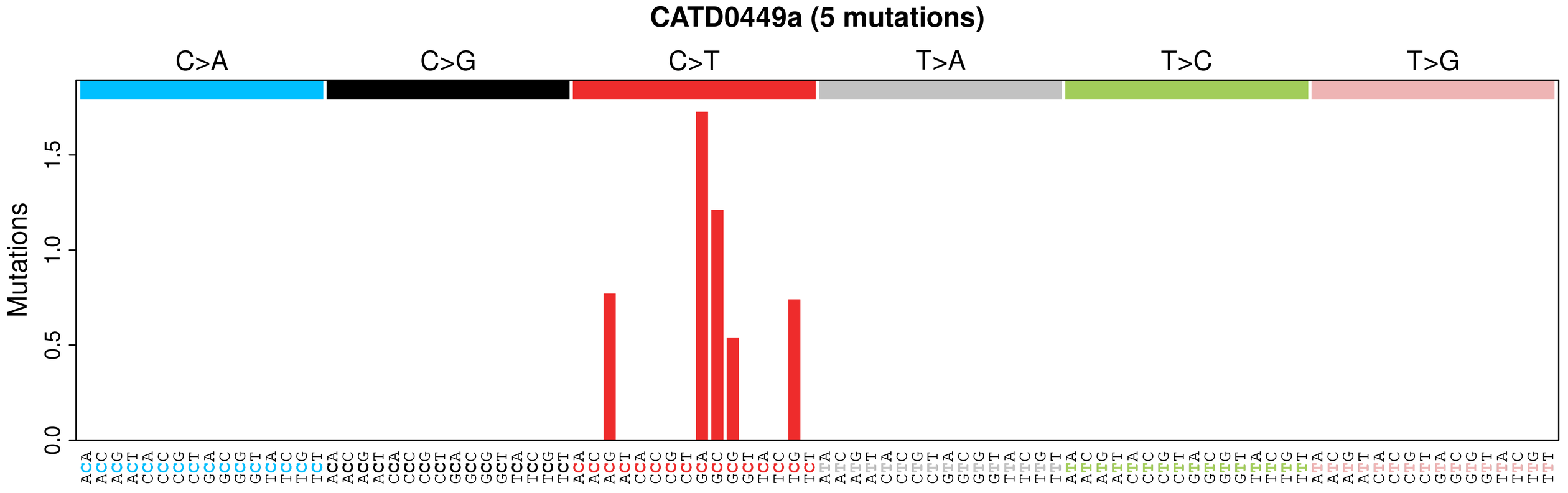

CATD0451a (5 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

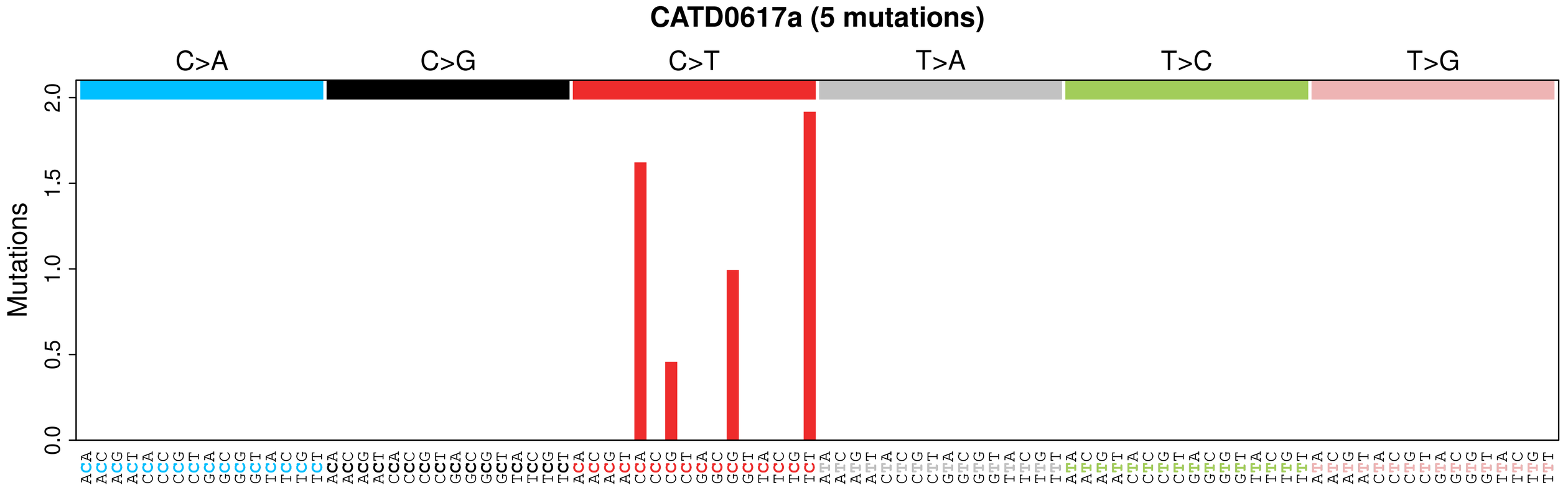

CATD0455a (5 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0
1.2
1.4

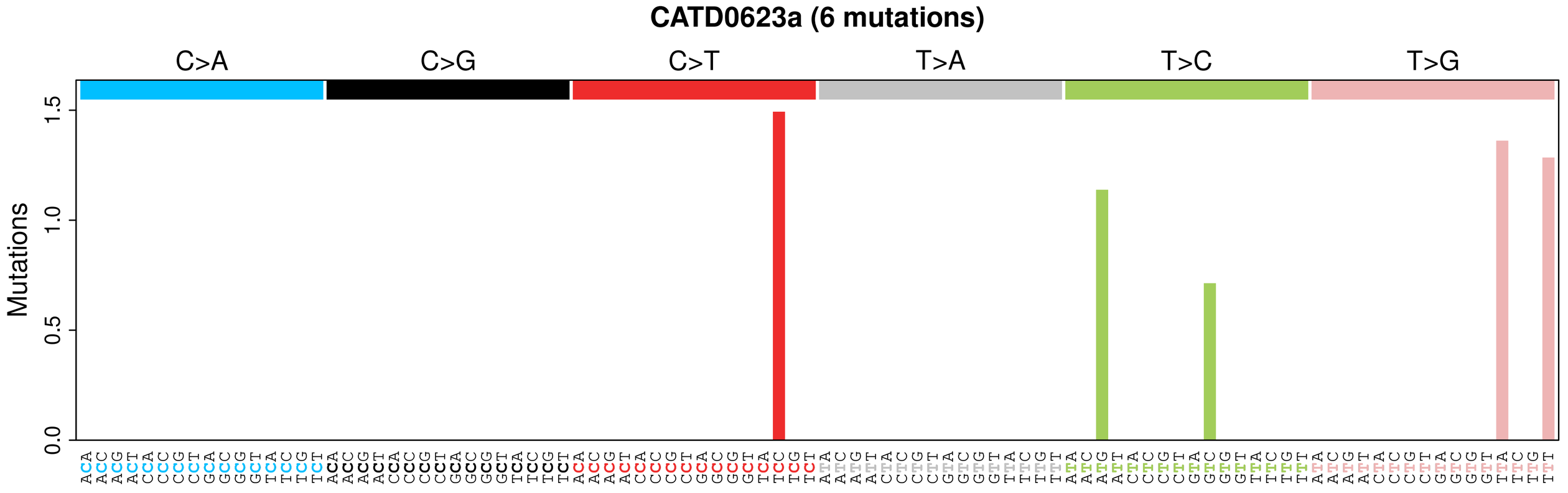

CATD0457a (7 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

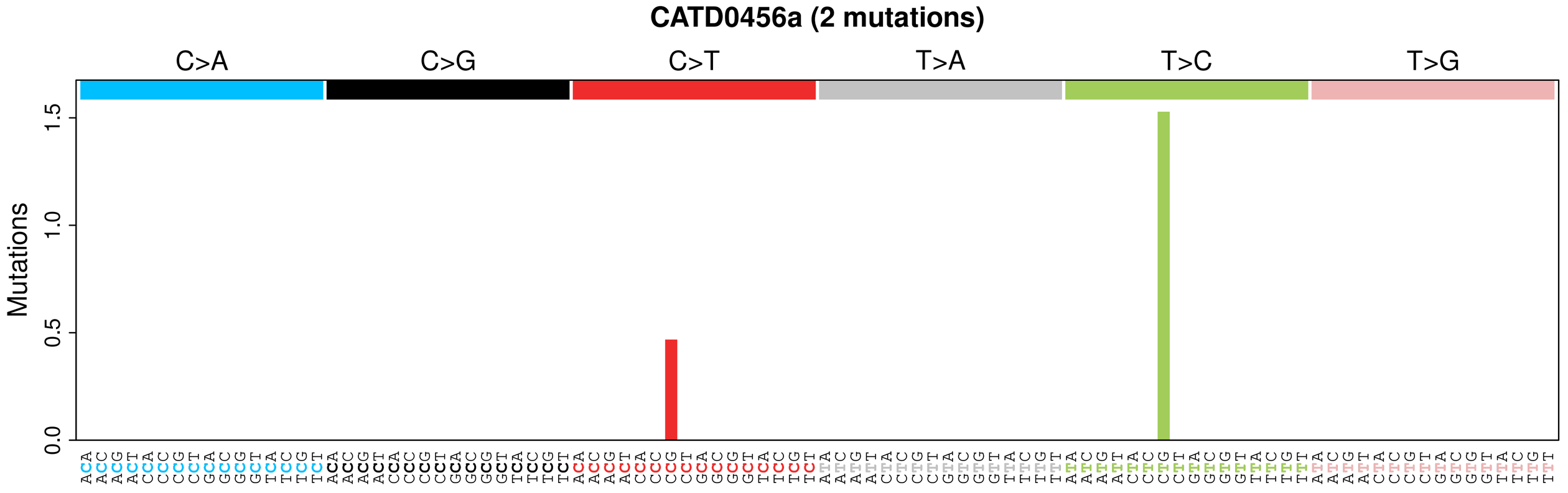

CATD0452a (11 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

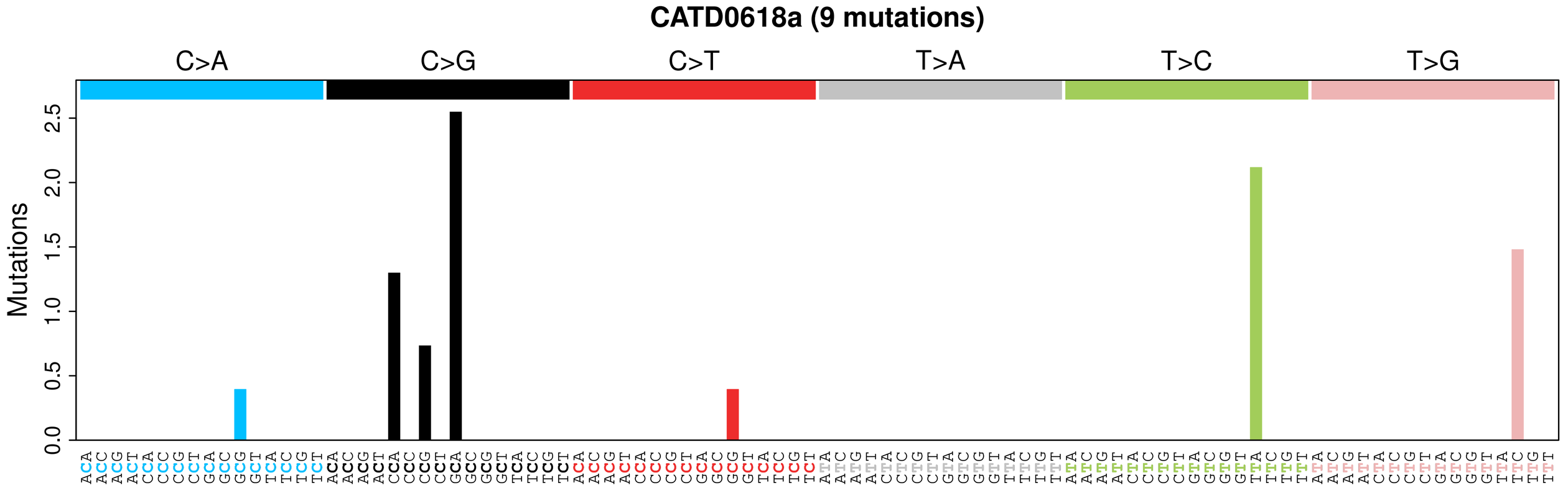

CATD0447a (4 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5
2.0

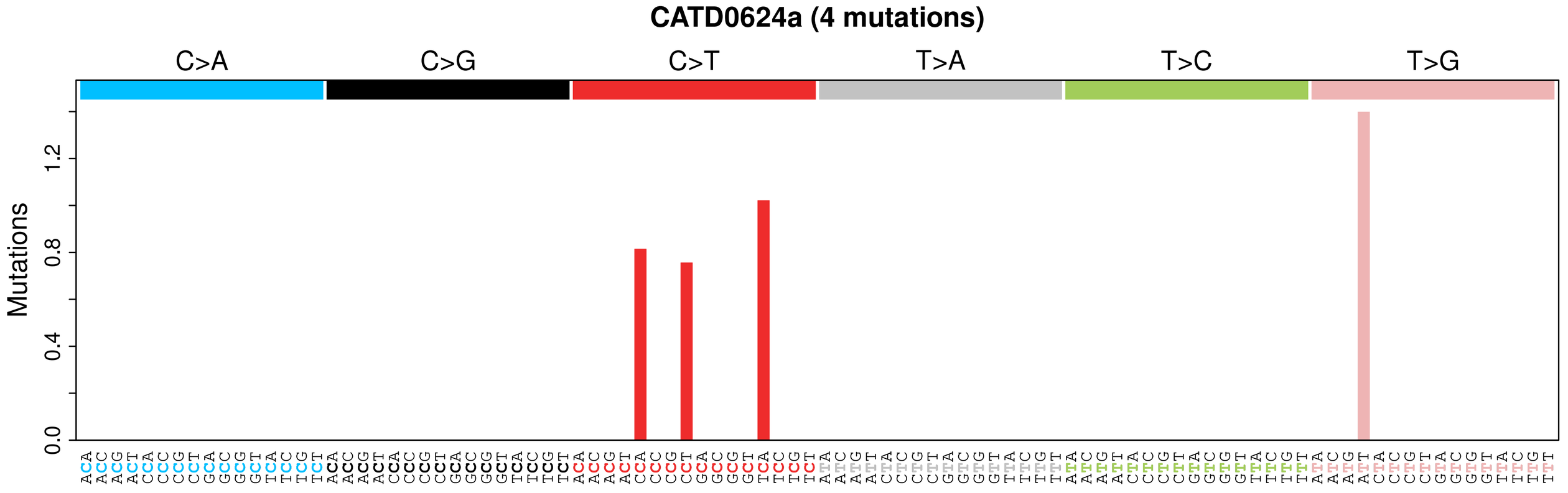

CATD0449a (5 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

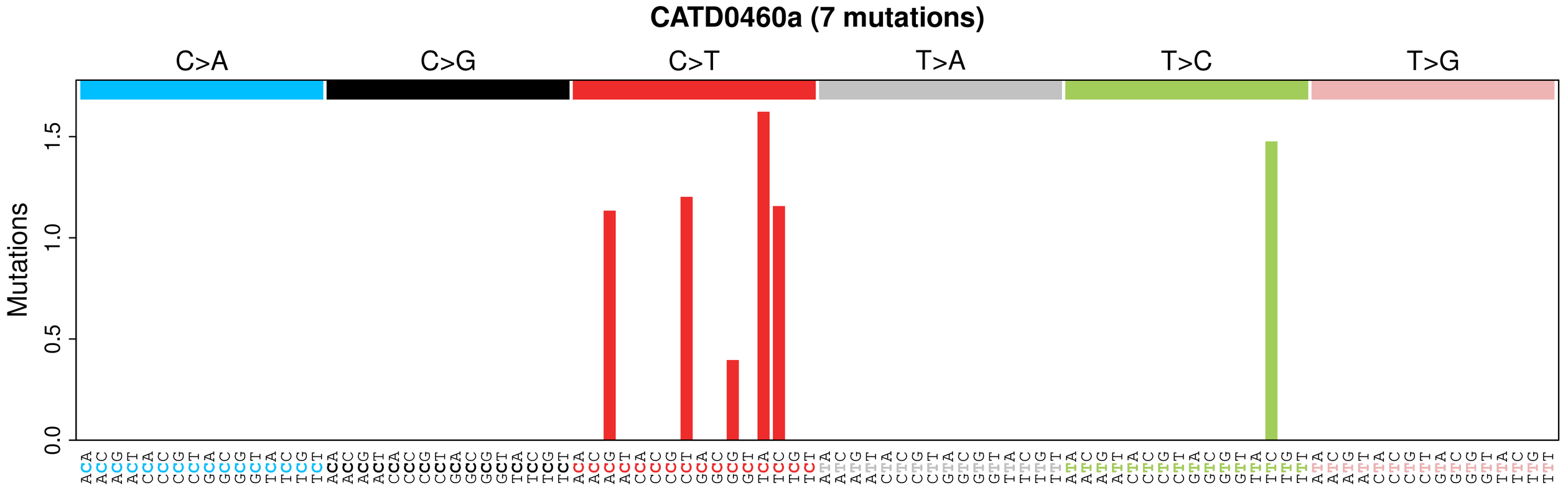

CATD0617a (5 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5
2.0

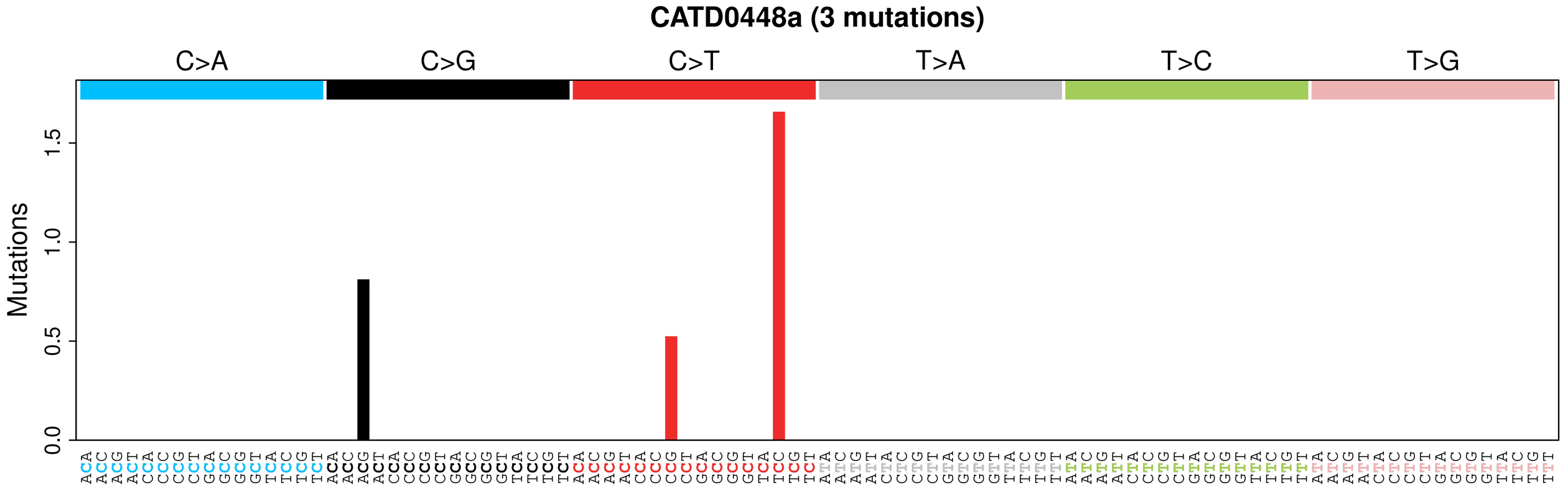

CATD0623a (6 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

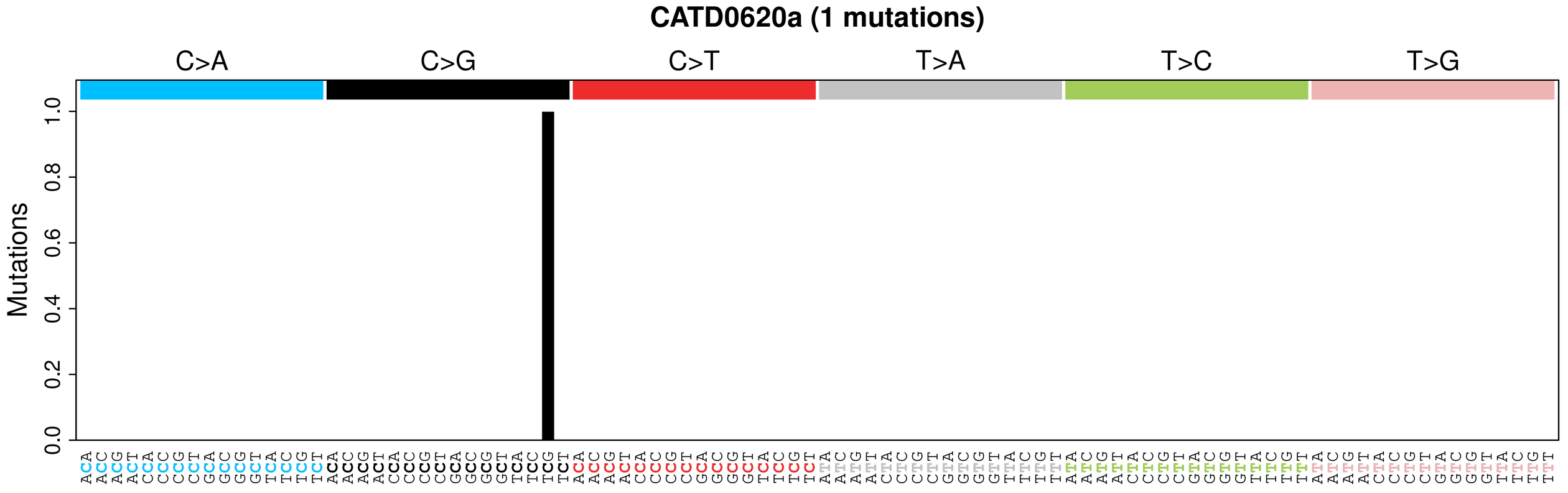

CATD0456a (2 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

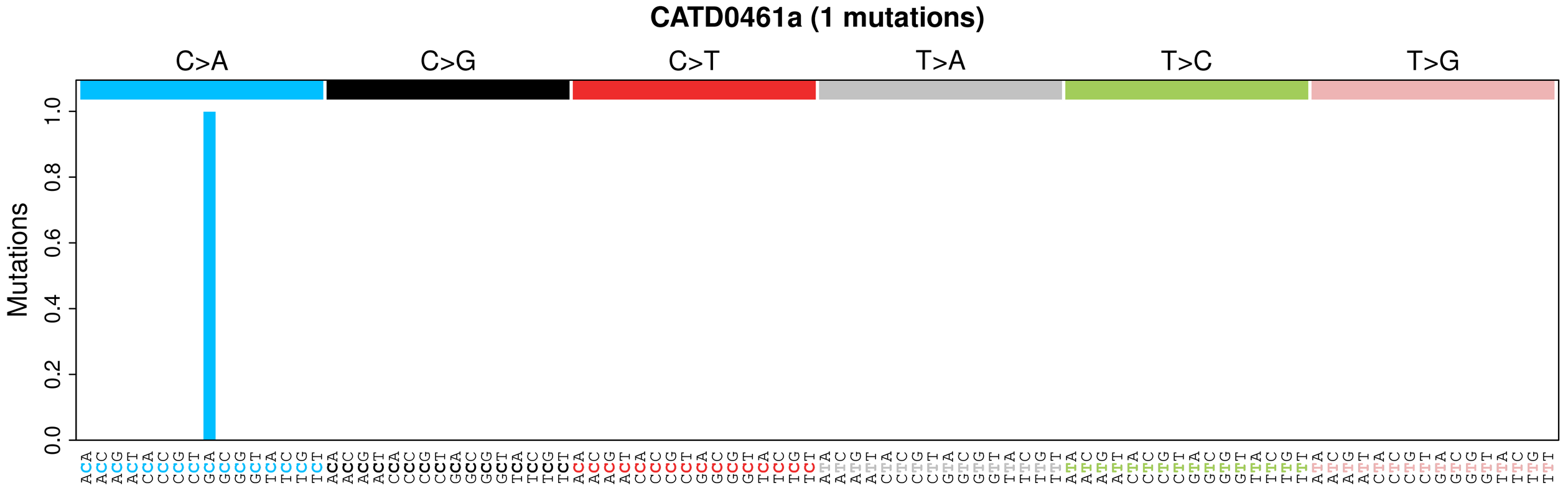

CATD0618a (9 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5
2.0
2.5

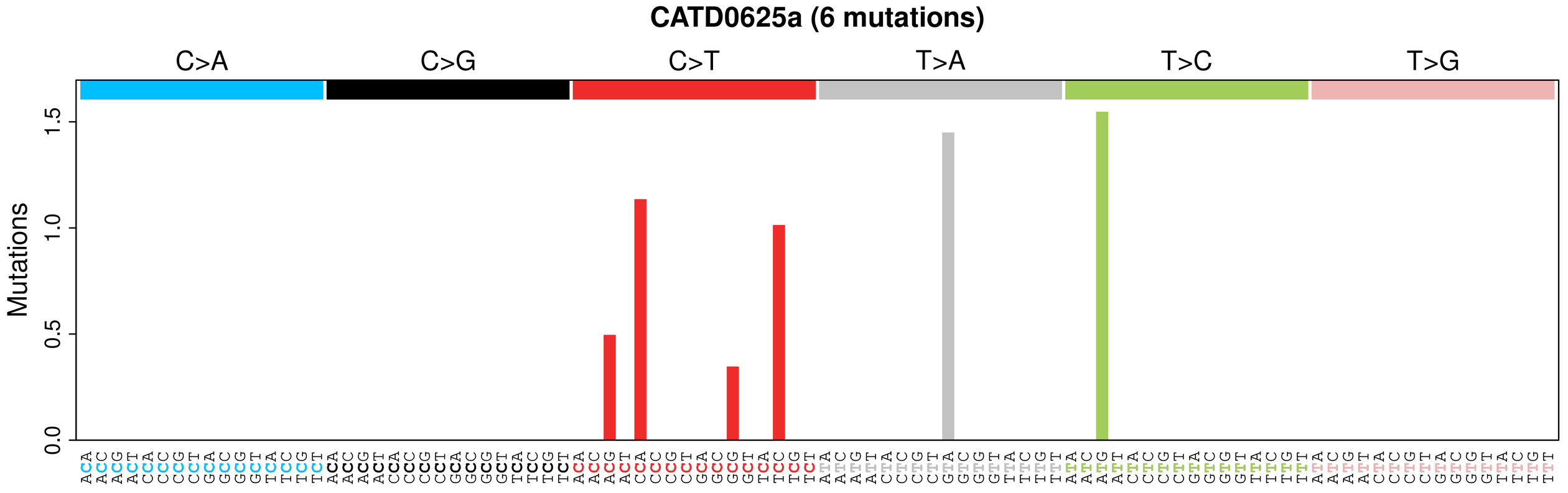

CATD0624a (4 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.4
0.8
1.2

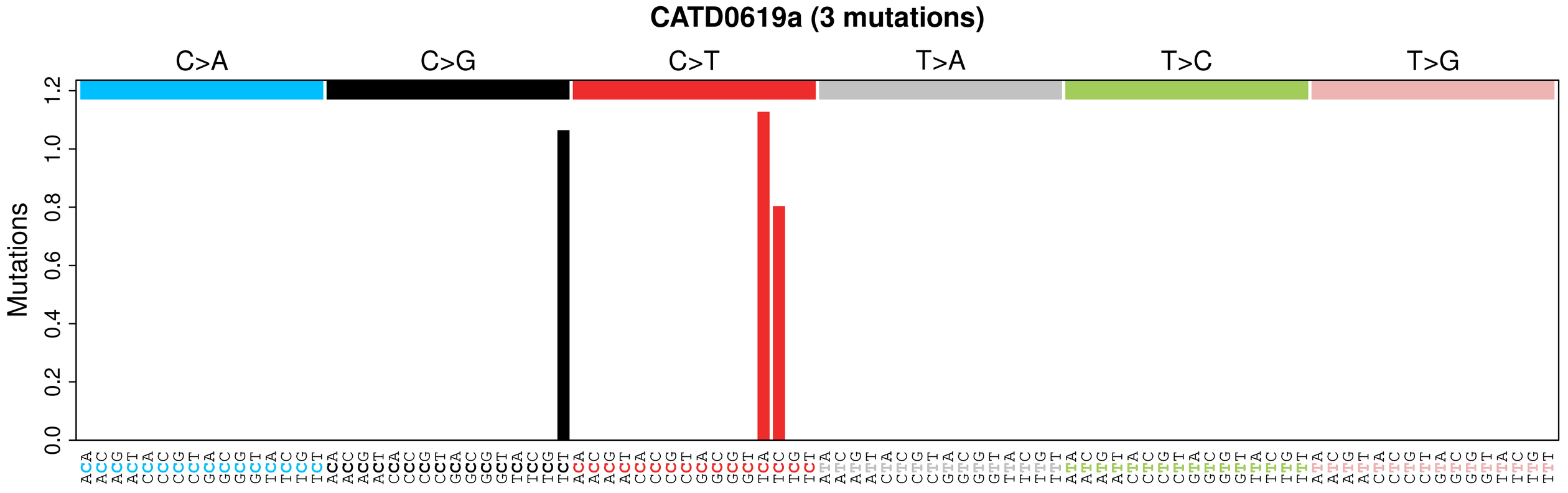

CATD0460a (7 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

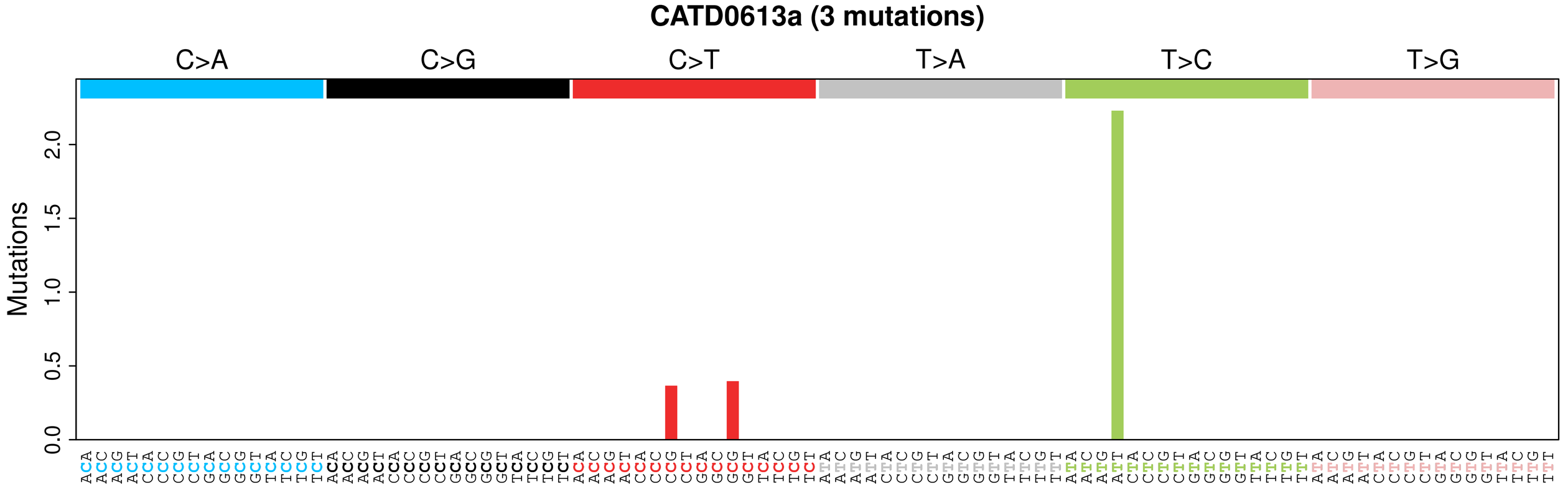

CATD0448a (3 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5

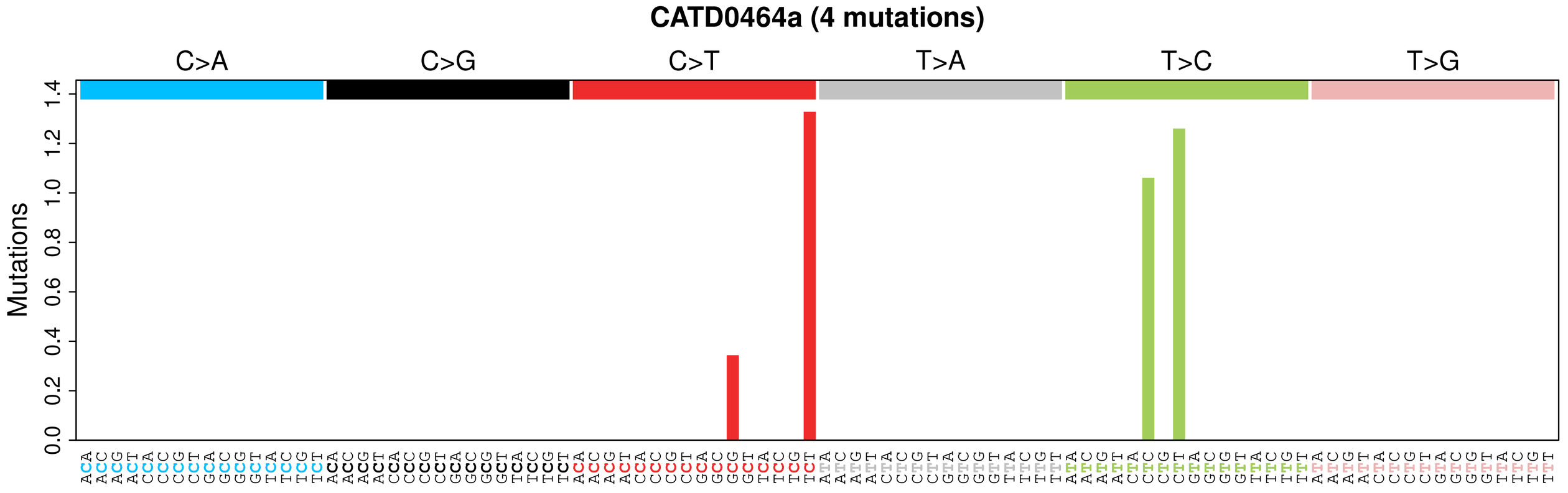

CATD0620a (1 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0


CATD0461a (1 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0


CATD0625a (6 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5


CATD0619a (3 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0
1.2


CATD0613a (3 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.5
1.0
1.5
2.0


CATD0464a (4 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0.0
0.2
0.4
0.6
0.8
1.0
1.2
1.4