

CATD0393a (30 mutations)

$C > A$

C > G

$C > T$

T>A

 $T > C$

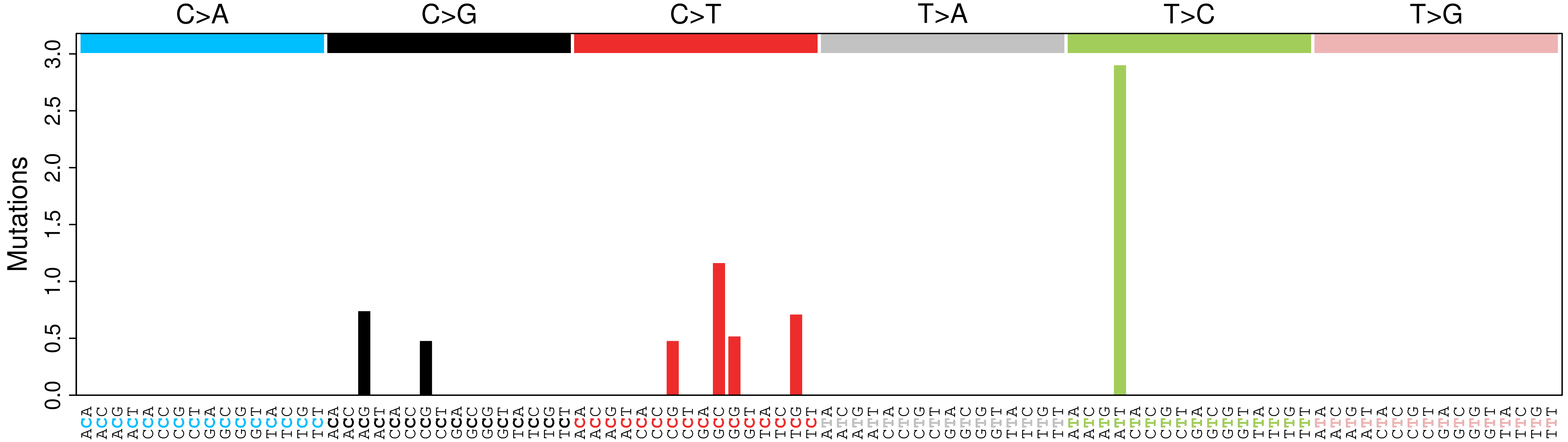
T>G

Mutations

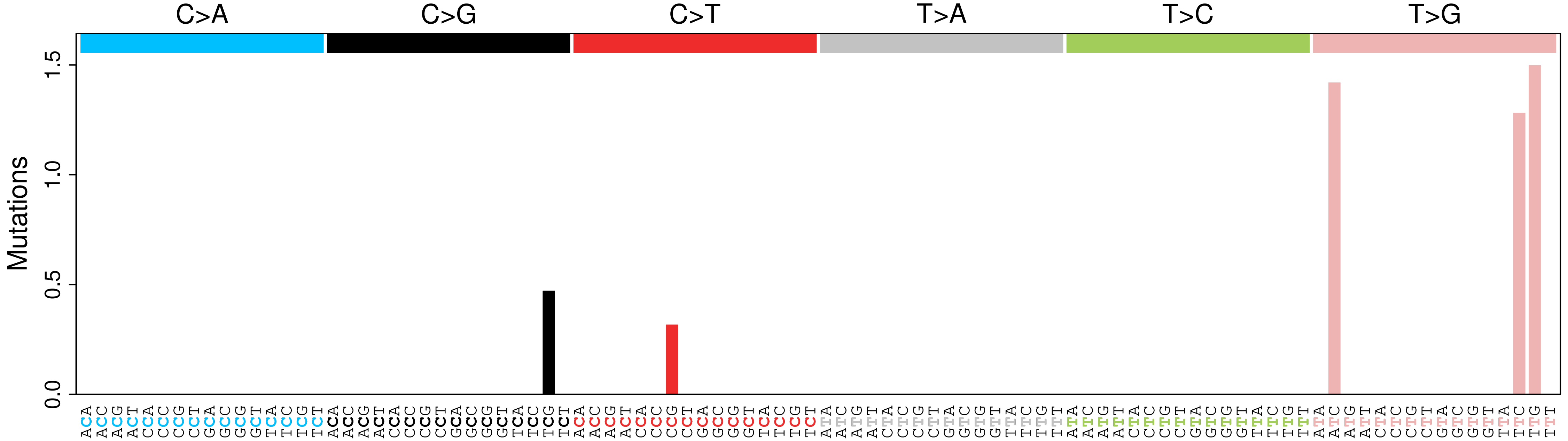
6
5
4
3
2
1
0

[illegible]

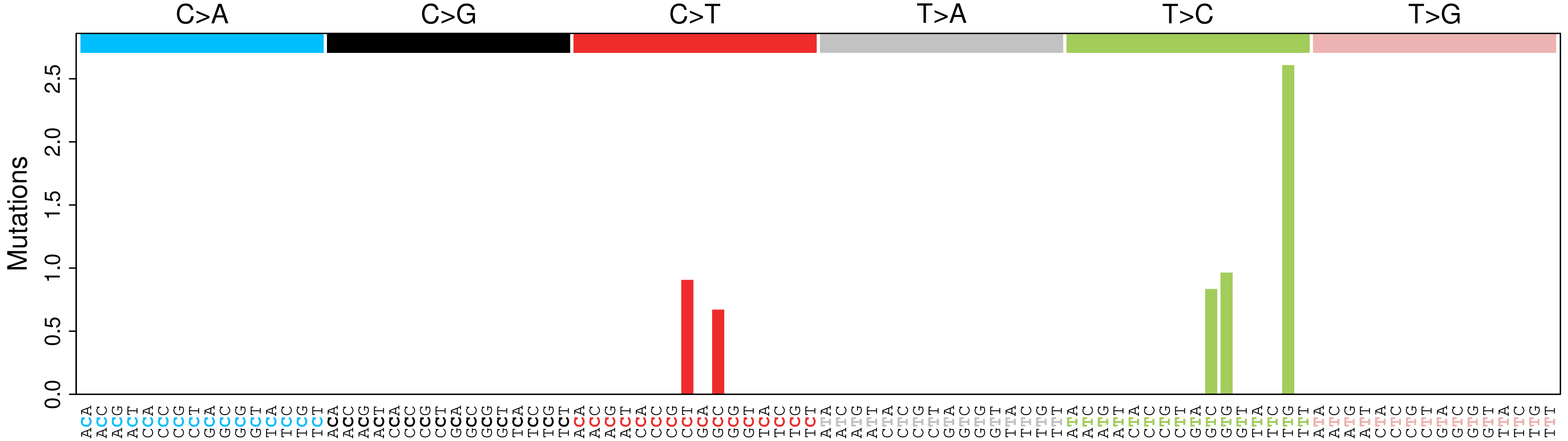
CATD0657a (7 mutations)



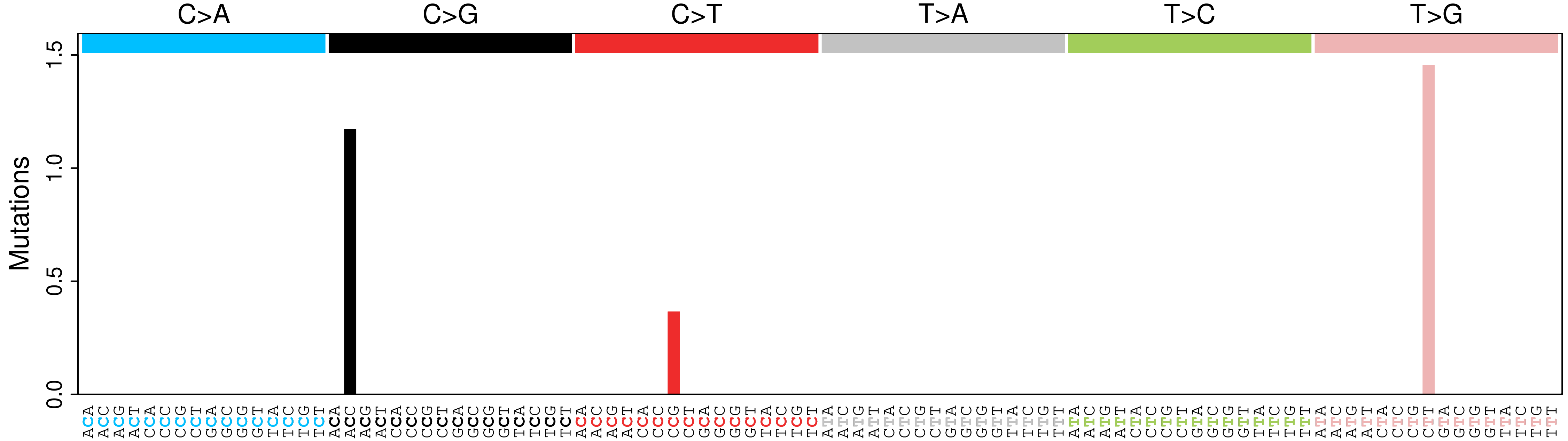
CATD0388a (5 mutations)



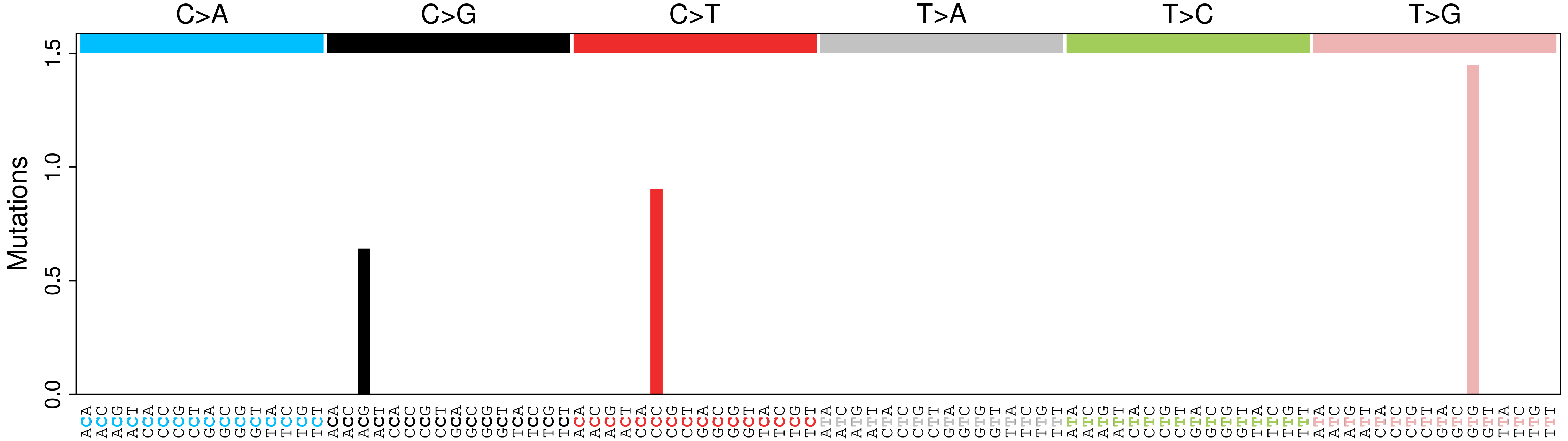
CATD0395a (6 mutations)



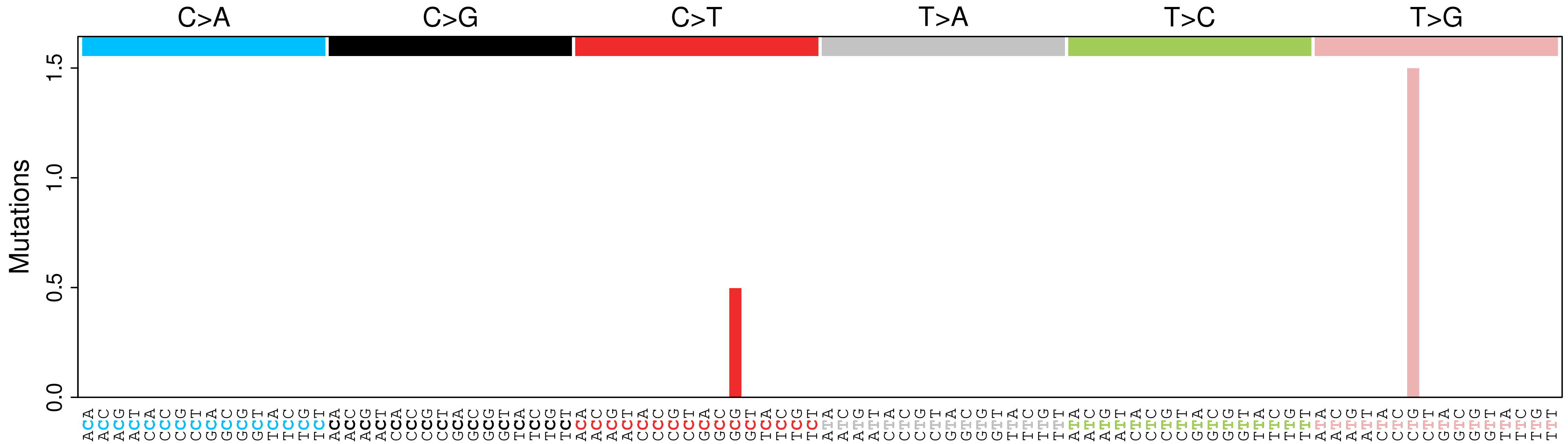
CATD0434a (3 mutations)



CATD0666a (3 mutations)



CATD0391a (2 mutations)



CATD0387a (2 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

T > G

Mutations

1.5

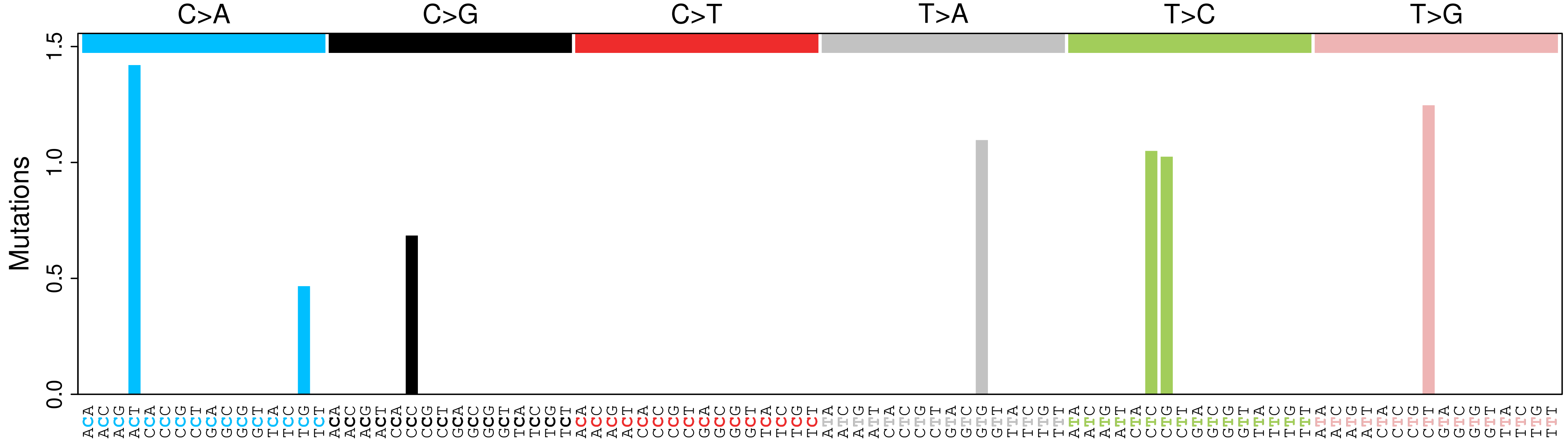
1.

0.

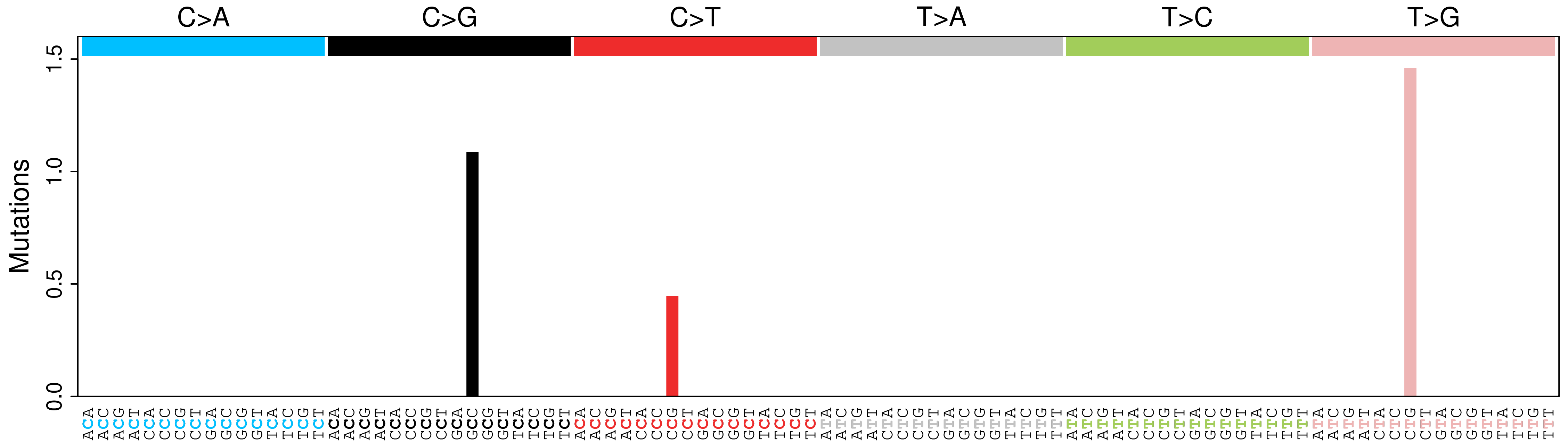
0.0

[illegible]

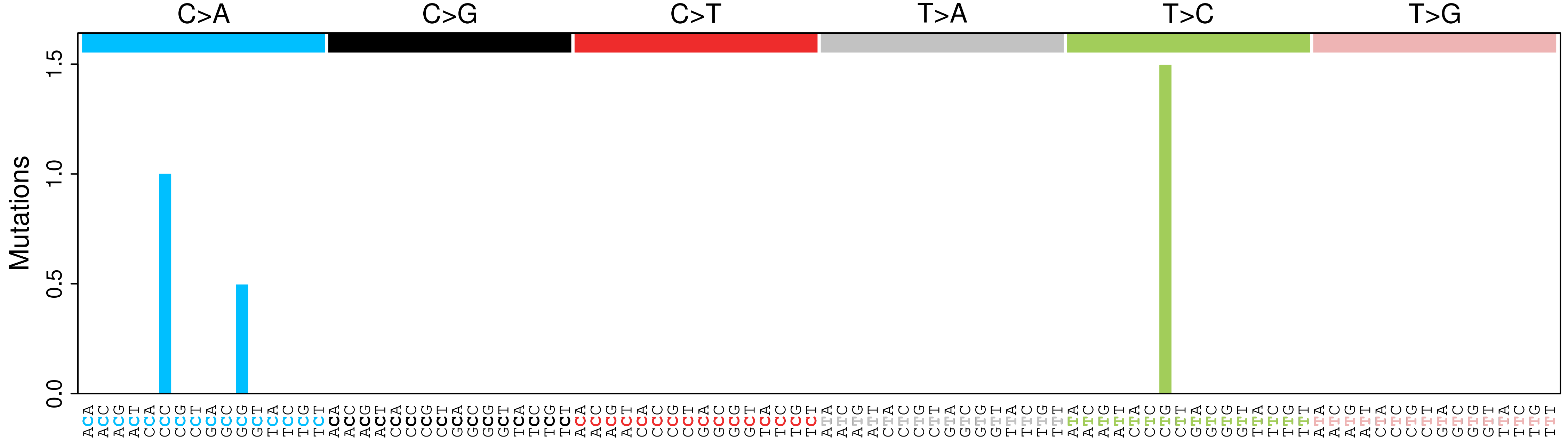
CATD0381a (7 mutations)



CATD0390a (3 mutations)



CATD0423a (3 mutations)



CATD0398a (3 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

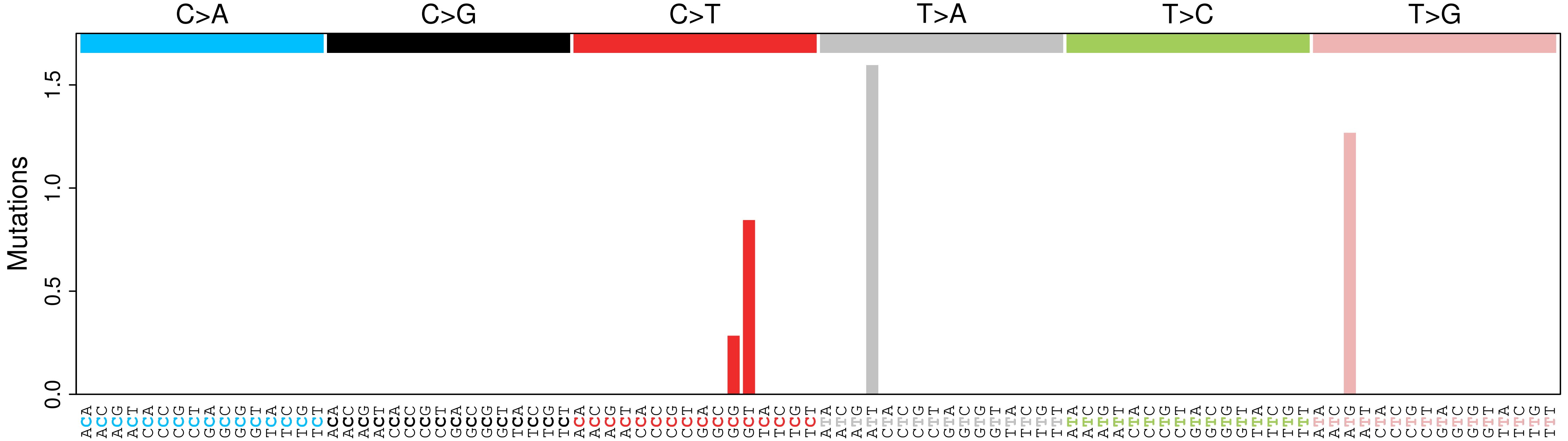
T > G

Mutations

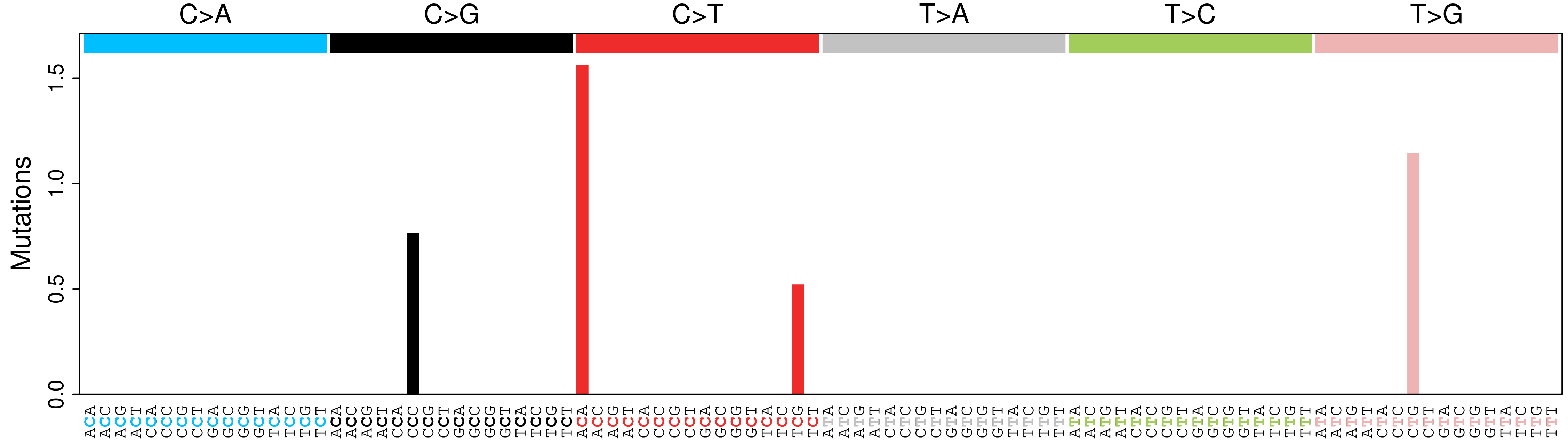
0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

[illegible]

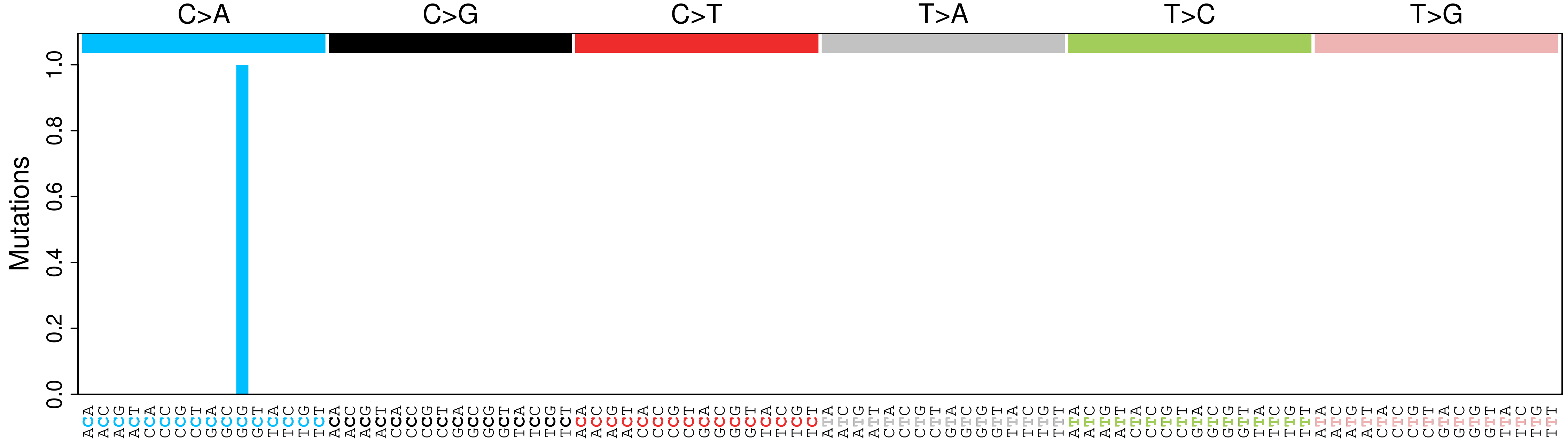
CATD0428a (4 mutations)



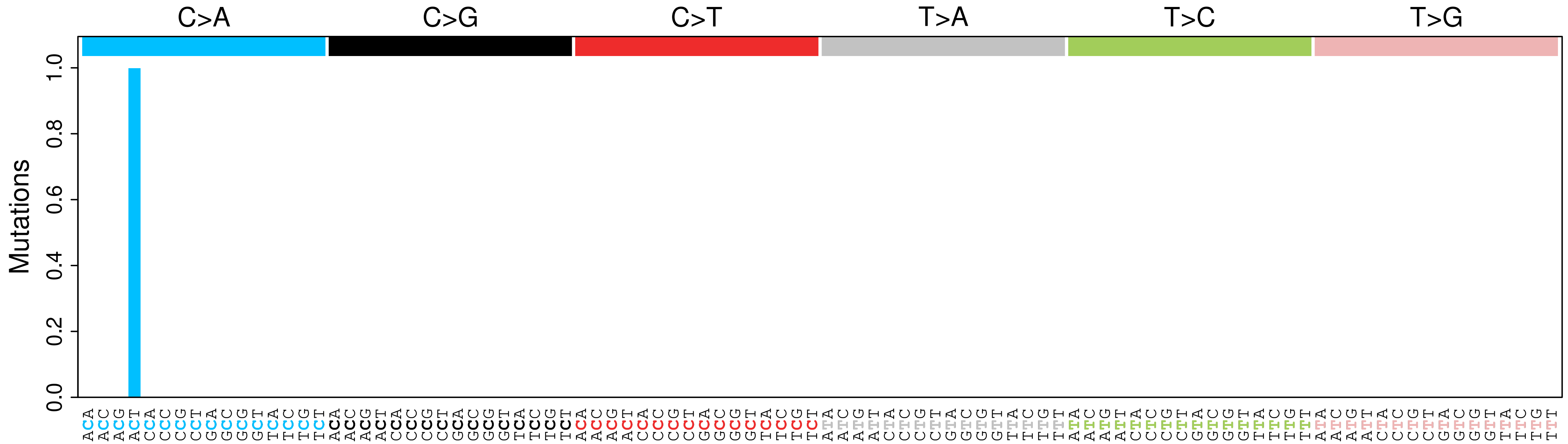
CATD0378a (4 mutations)



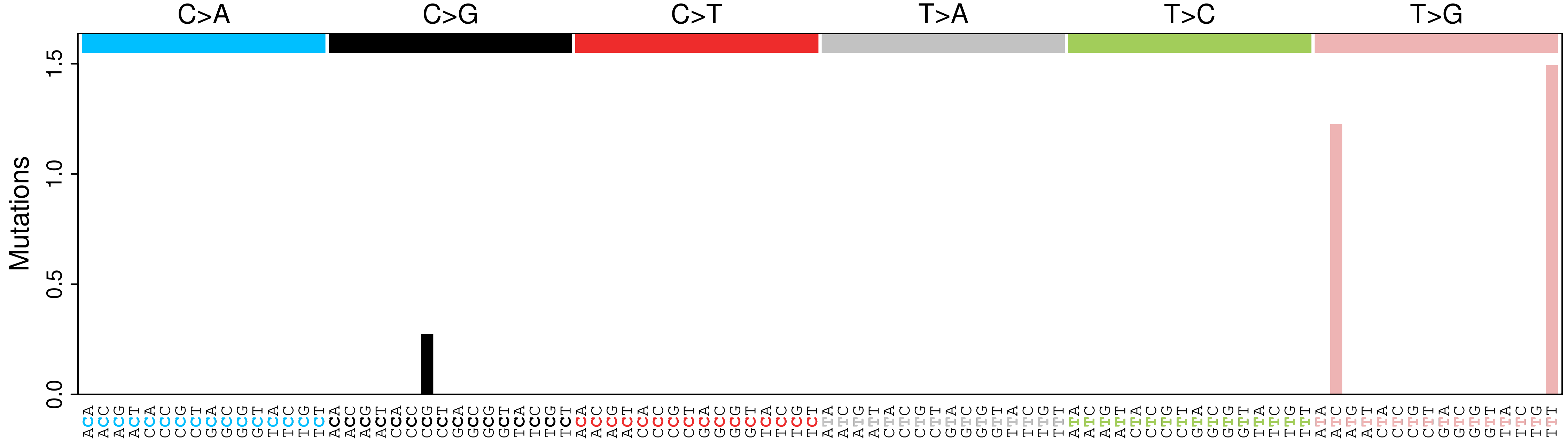
CATD0380a (1 mutations)



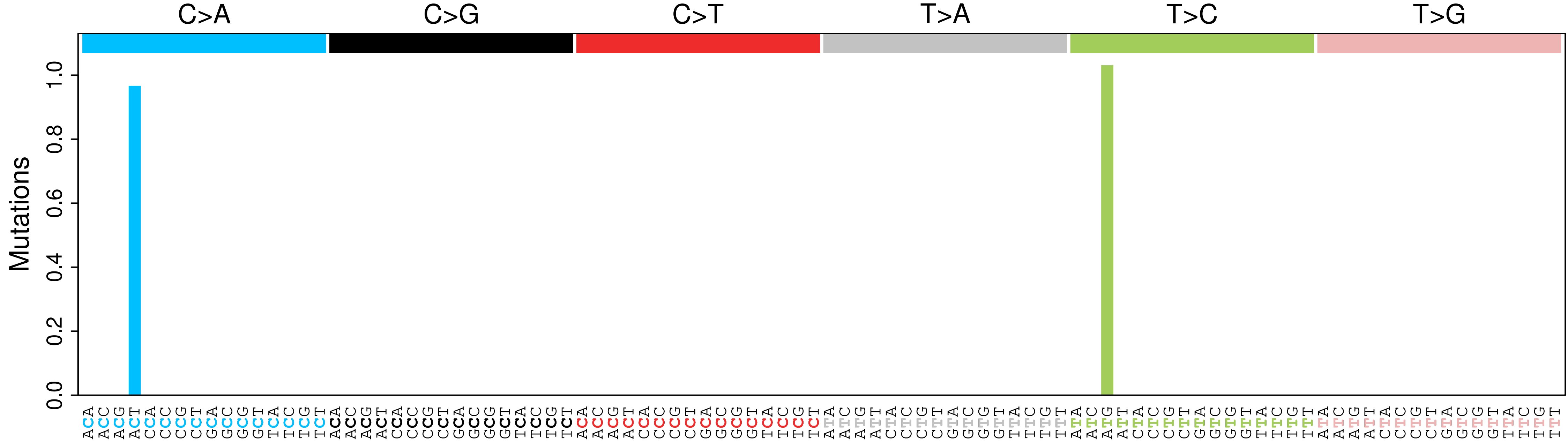
CATD0662a (1 mutations)



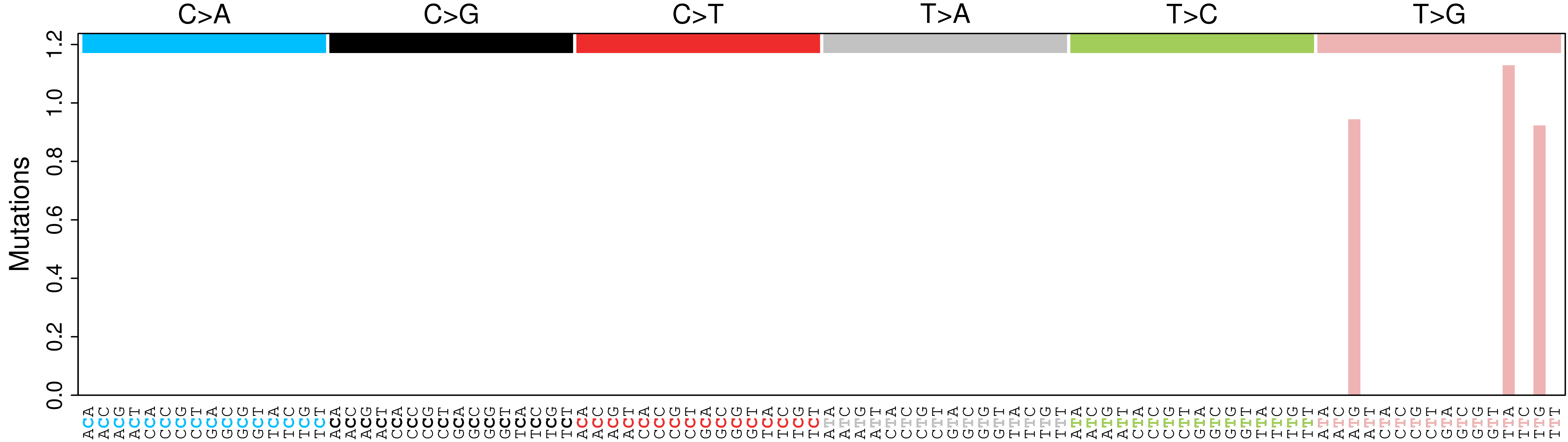
CATD0399a (3 mutations)



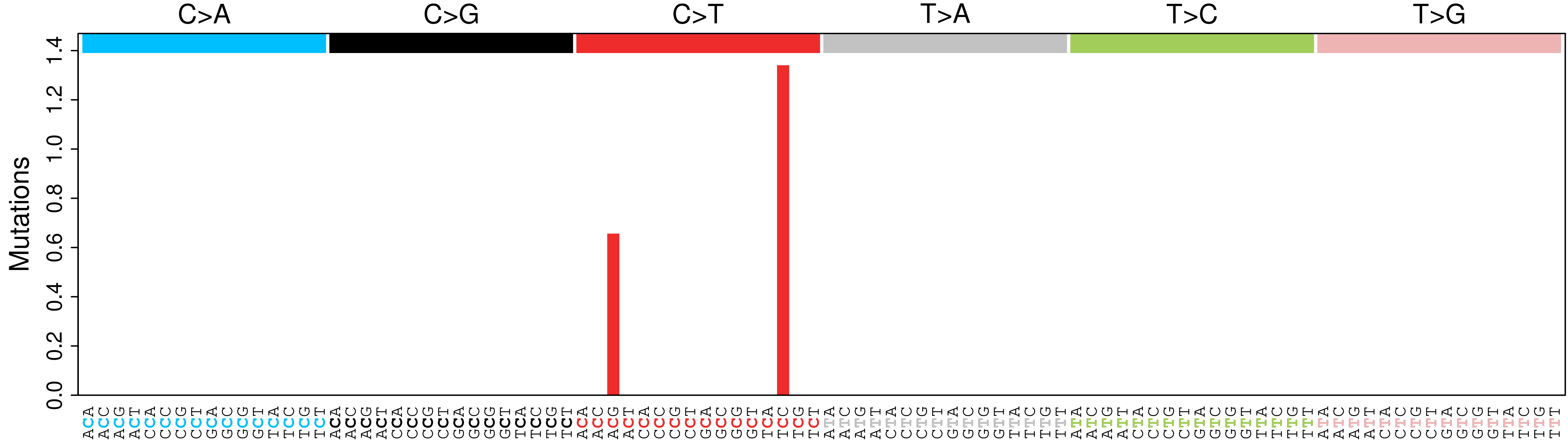
CATD0659a (2 mutations)



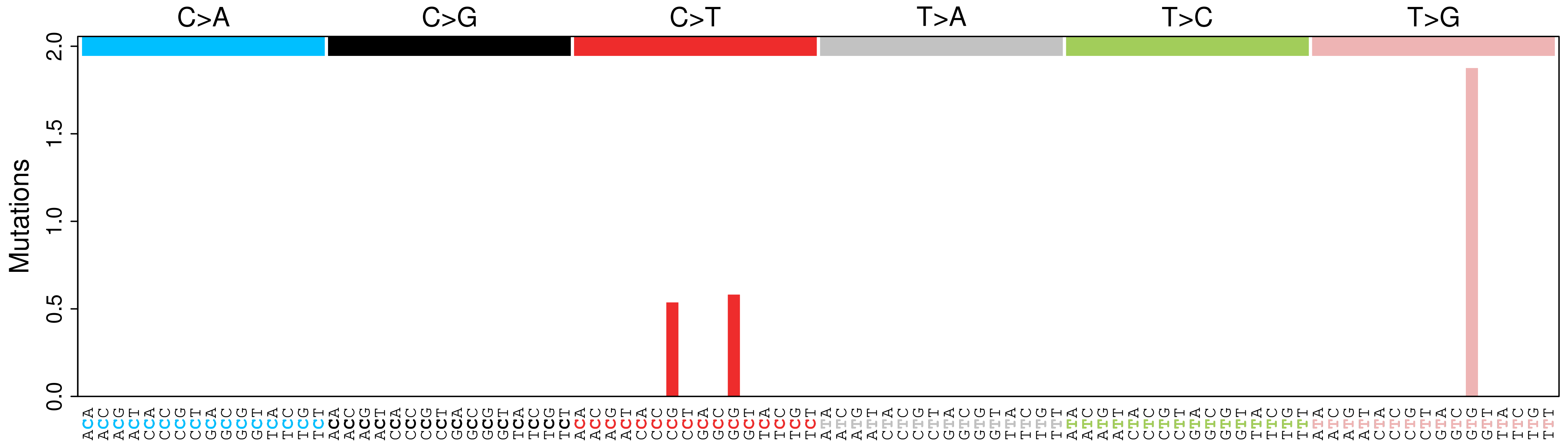
CATD0389a (3 mutations)



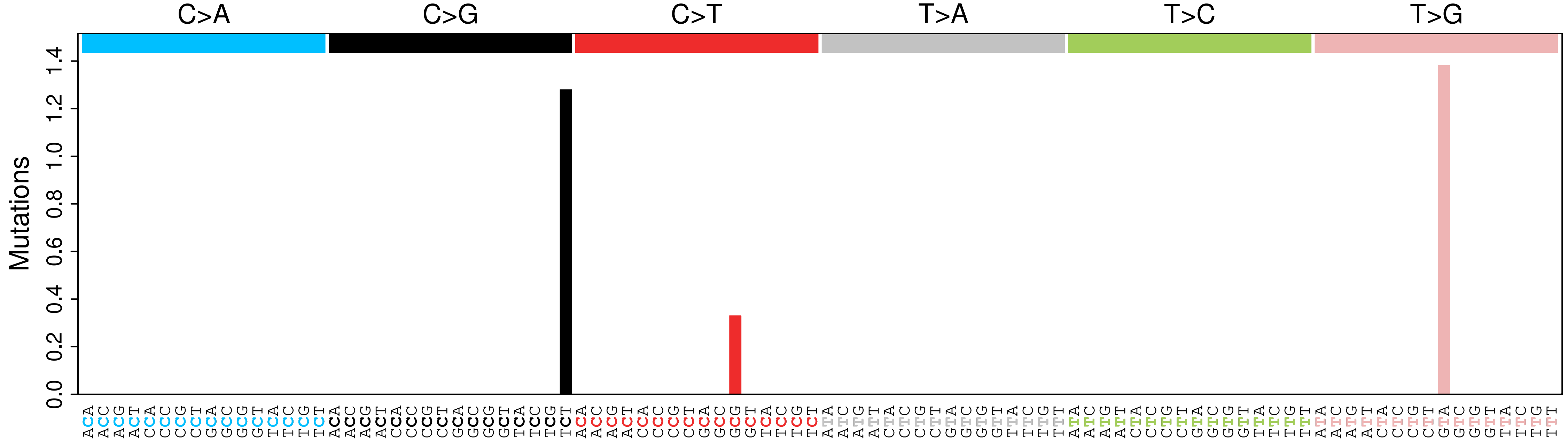
CATD0658a (2 mutations)



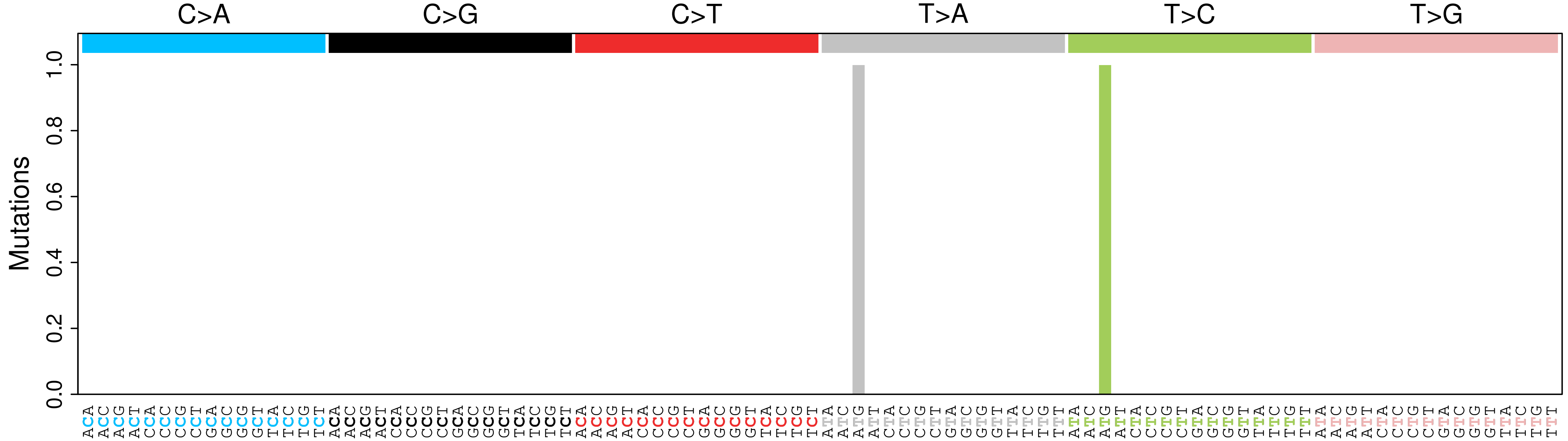
CATD0665a (3 mutations)



CATD0377a (3 mutations)



CATD0422a (2 mutations)



CATD0392a (2 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

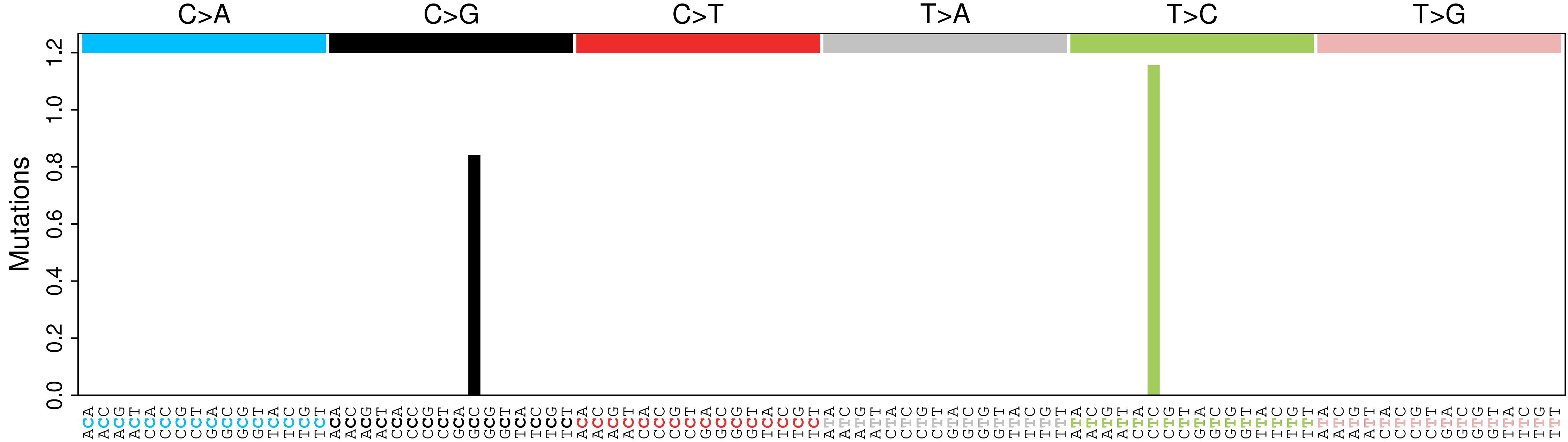
T > G

Mutations

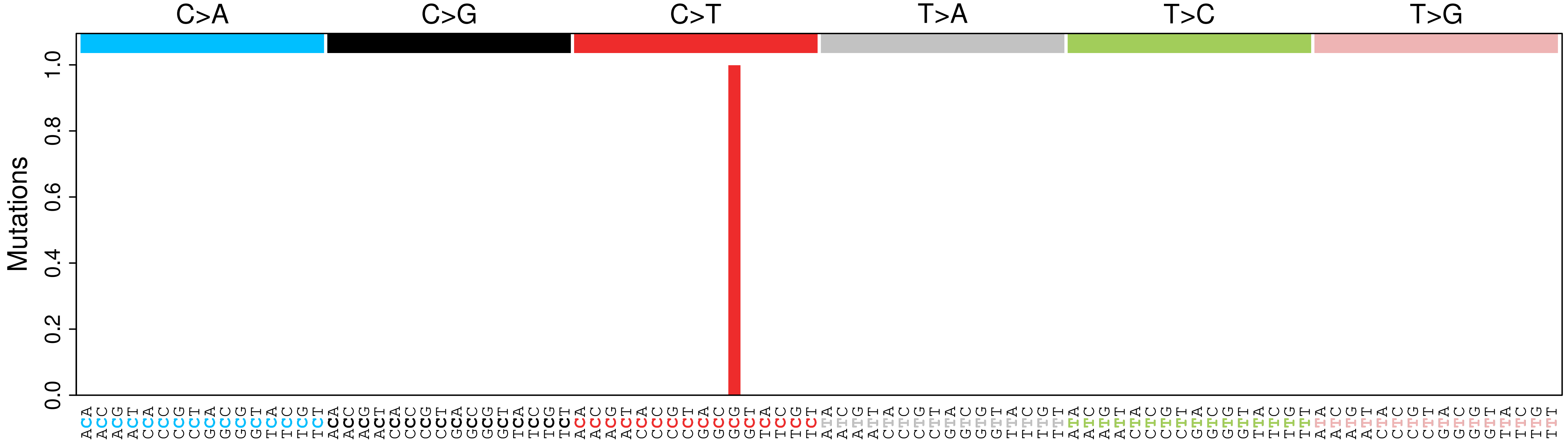
0.0 0.2 0.4 0.6 0.8 1.0 1.2

[illegible]

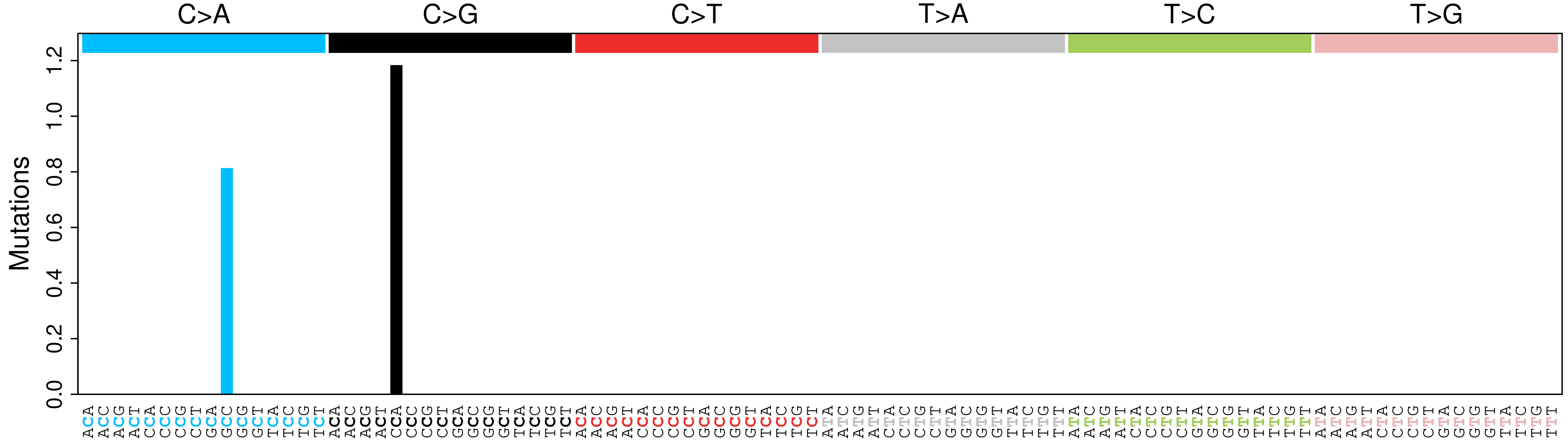
CATD0661a (2 mutations)



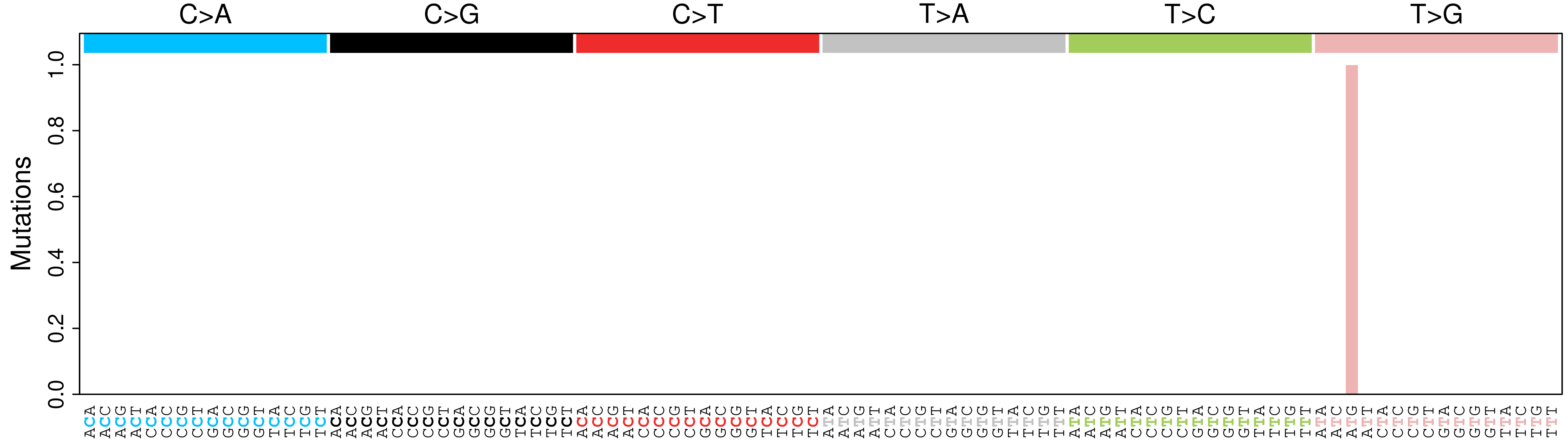
CATD0424a (1 mutations)



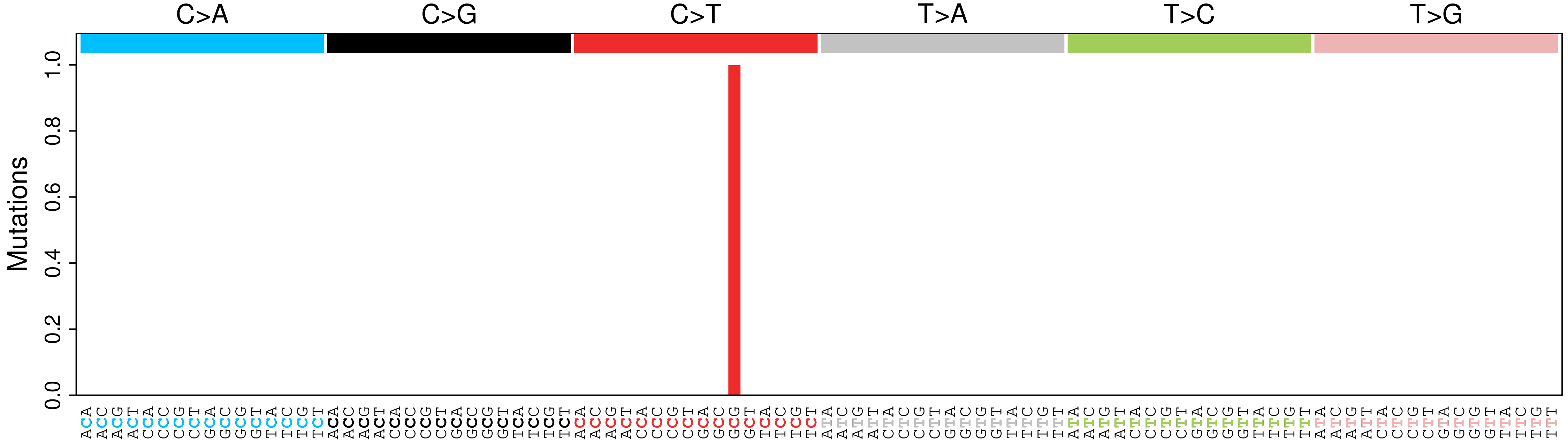
CATD0426a (2 mutations)



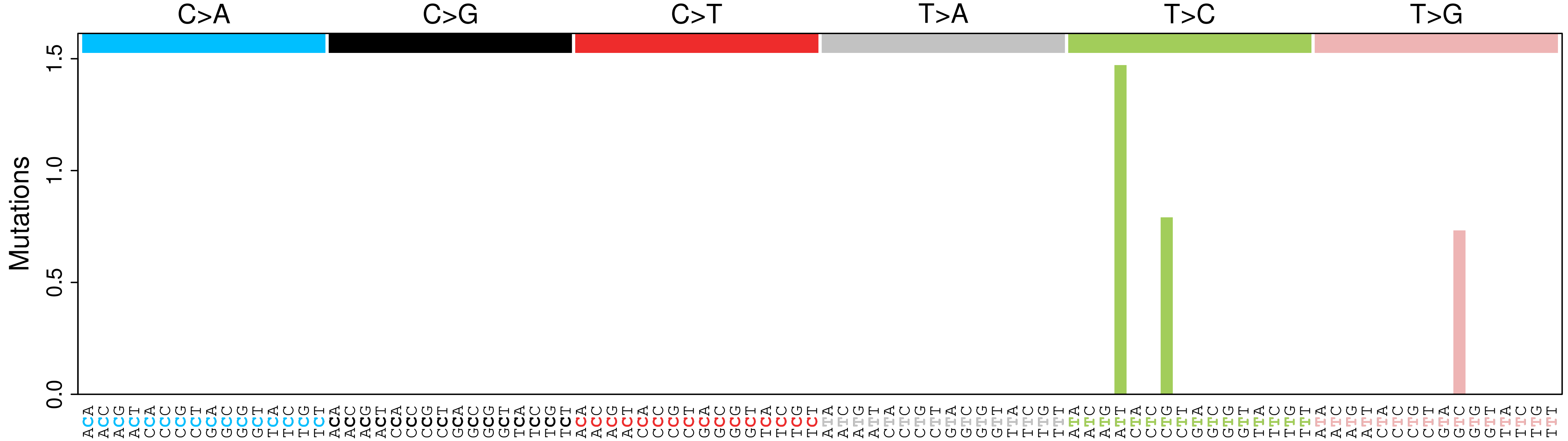
CATD0383a (1 mutations)



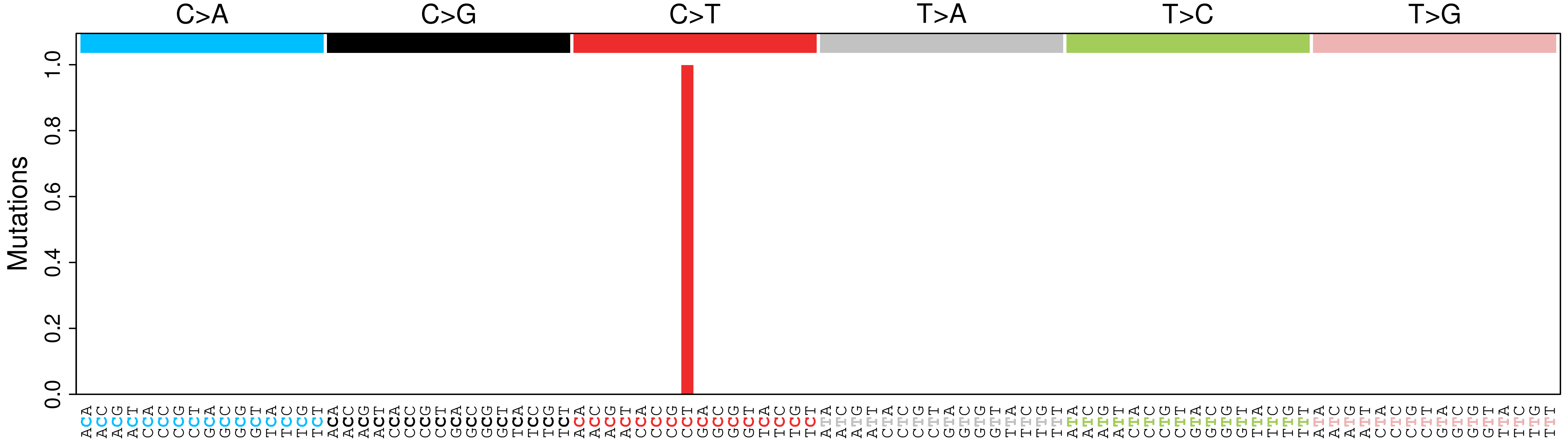
CATD0385a (1 mutations)



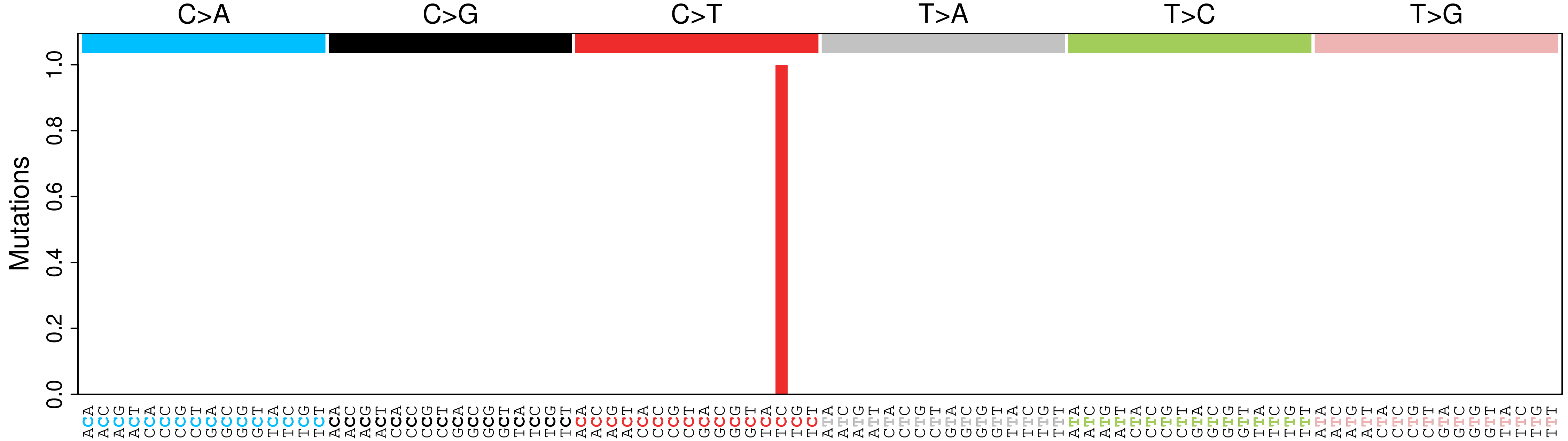
CATD0382a (3 mutations)



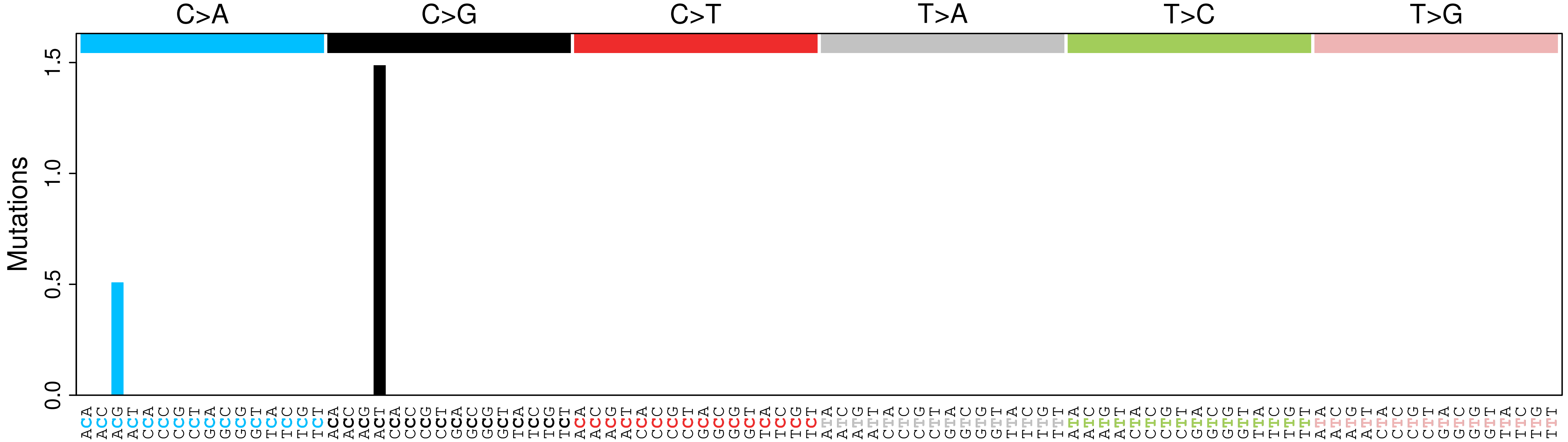
CATD0660a (1 mutations)



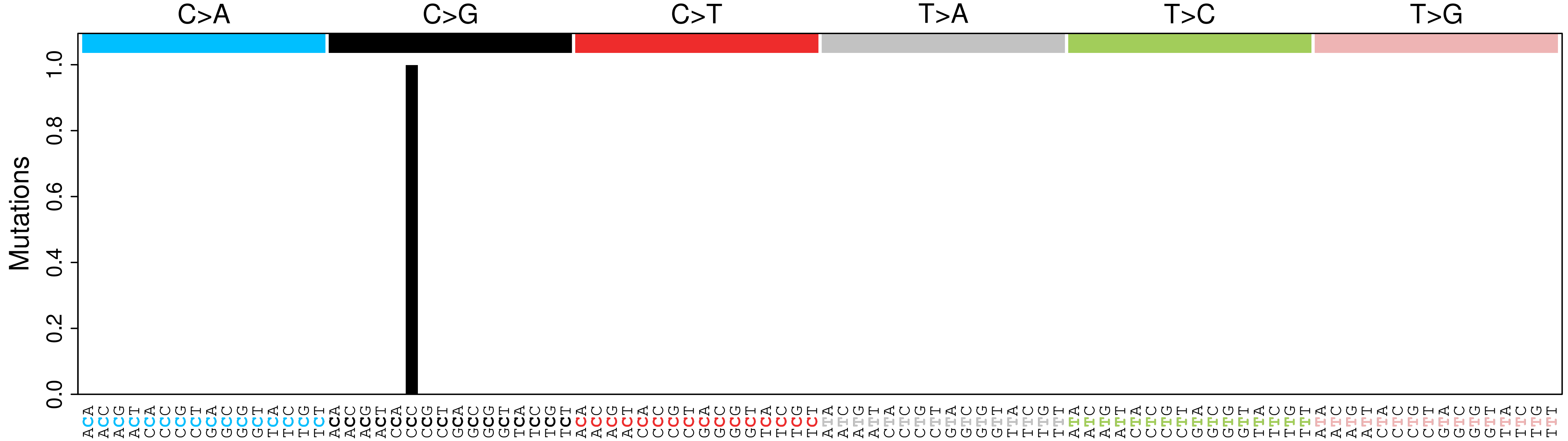
CATD0437a (1 mutations)



CATD0439a (2 mutations)



CATD0663a (1 mutations)



CATD0379a (1 mutations)

$C > A$

C > G

C>T

T>A

$$T > C$$

T > G

Mutations

1.0

3.0

0

U

[illegible]

CATD0397a (1 mutations)

