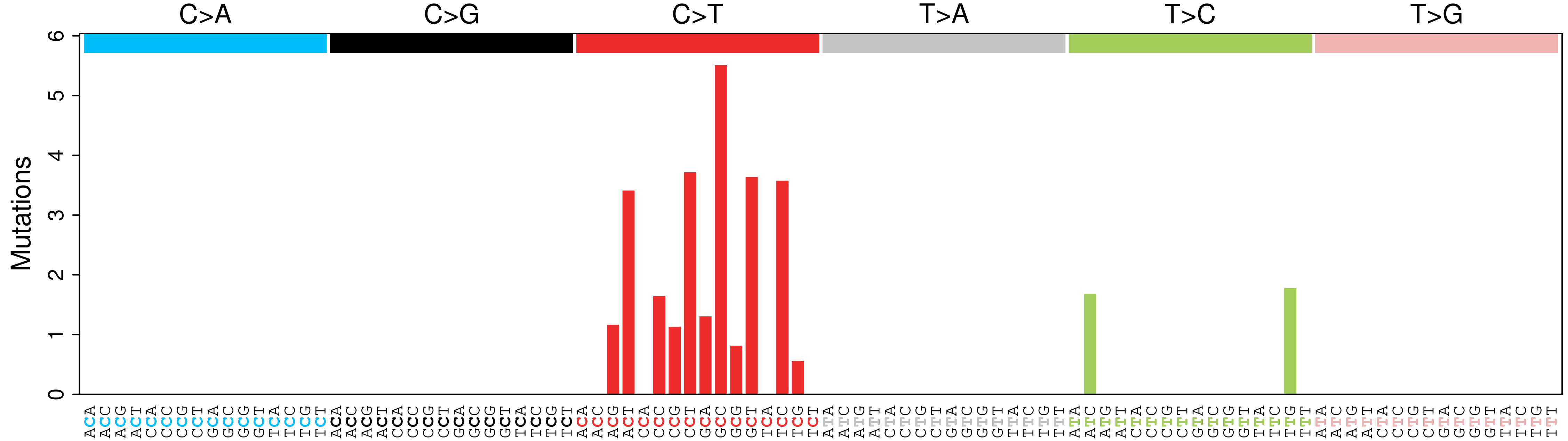
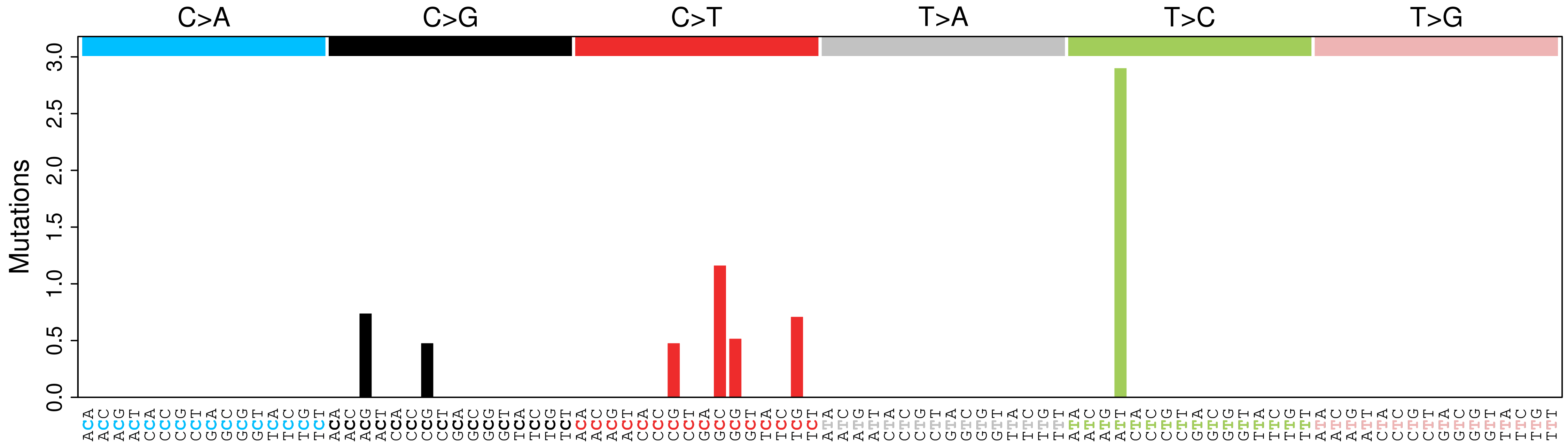


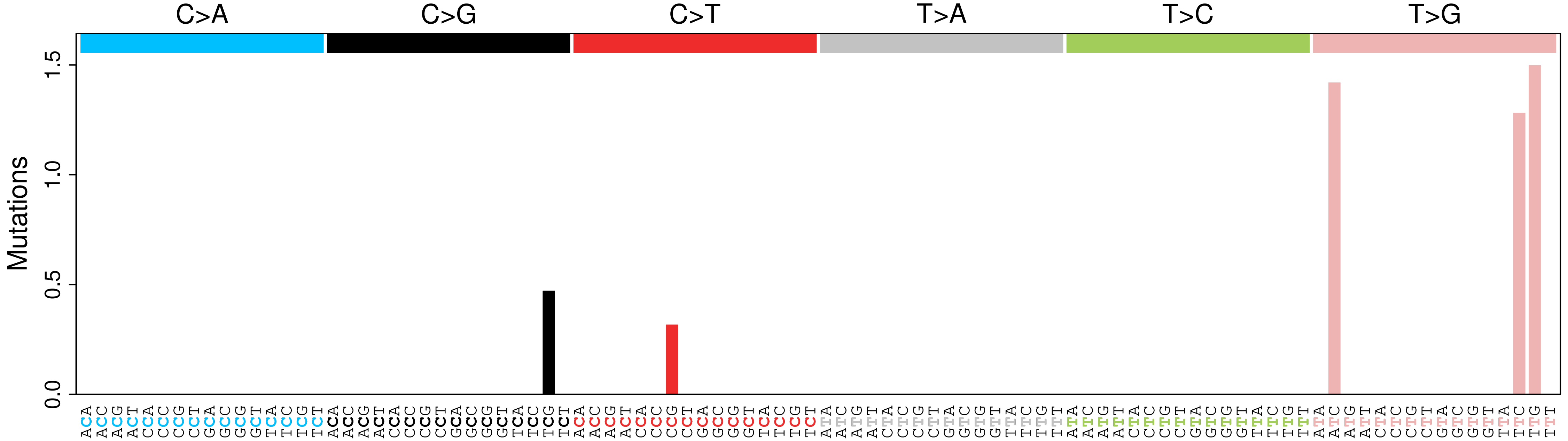
CATD0393a (30 mutations)



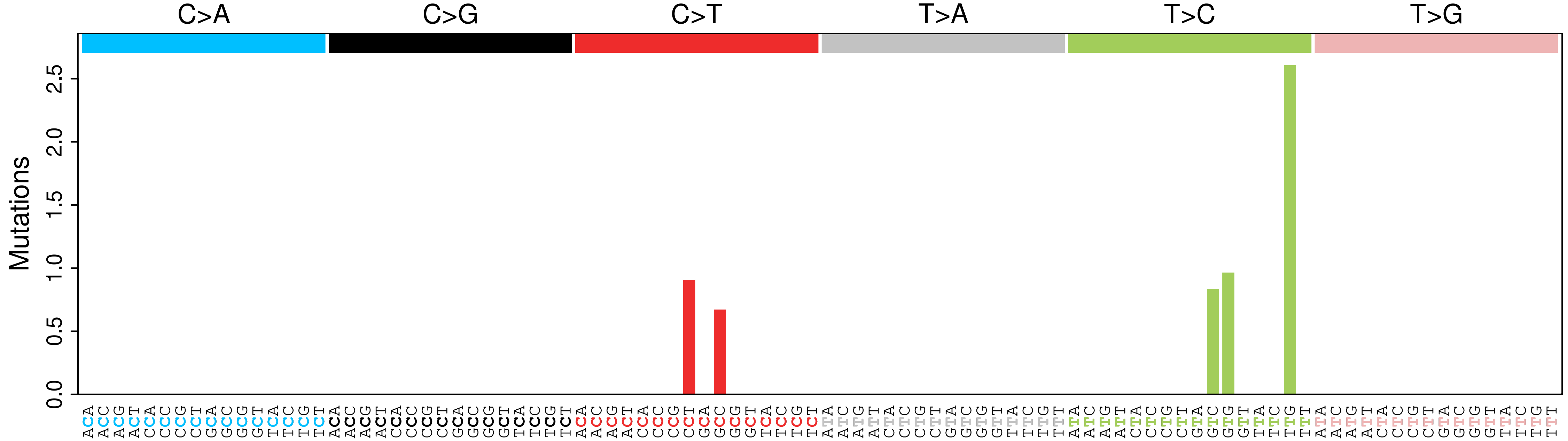
CATD0657a (7 mutations)



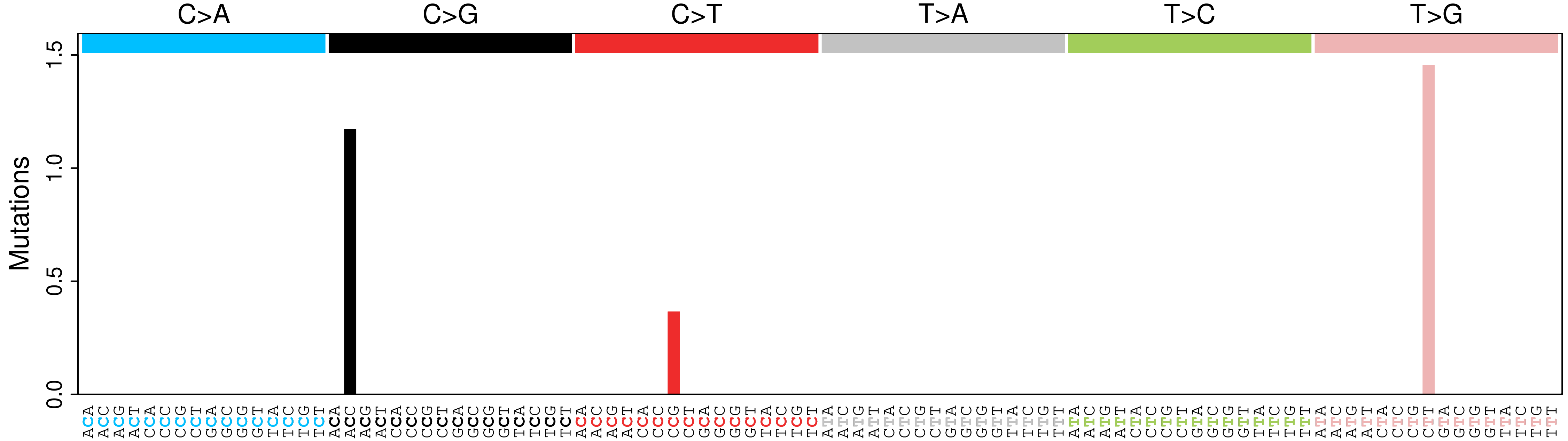
CATD0388a (5 mutations)



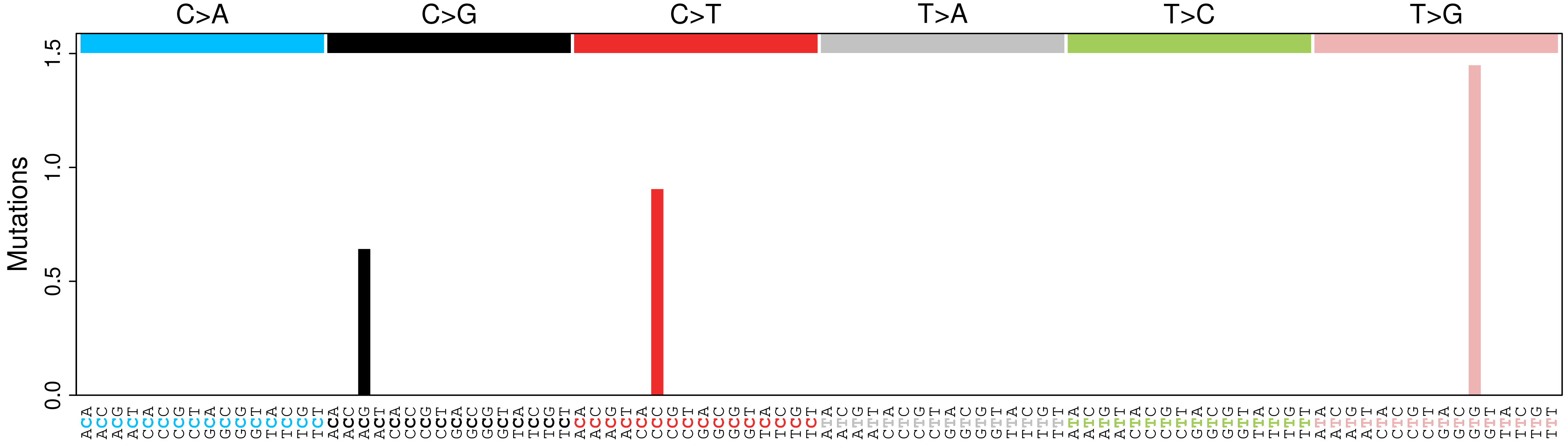
CATD0395a (6 mutations)



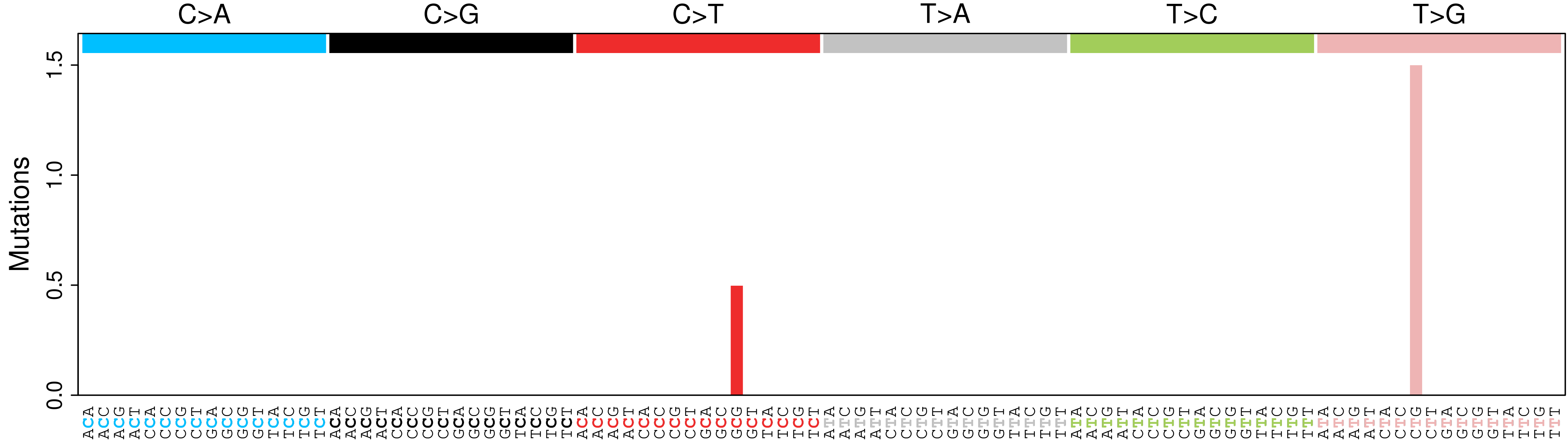
CATD0434a (3 mutations)



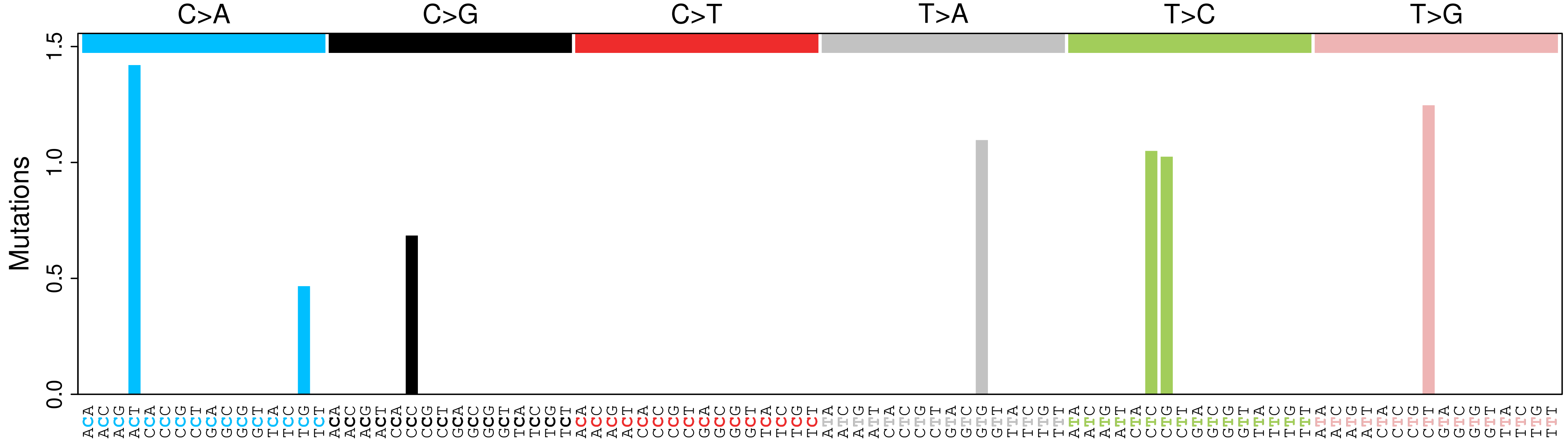
CATD0666a (3 mutations)



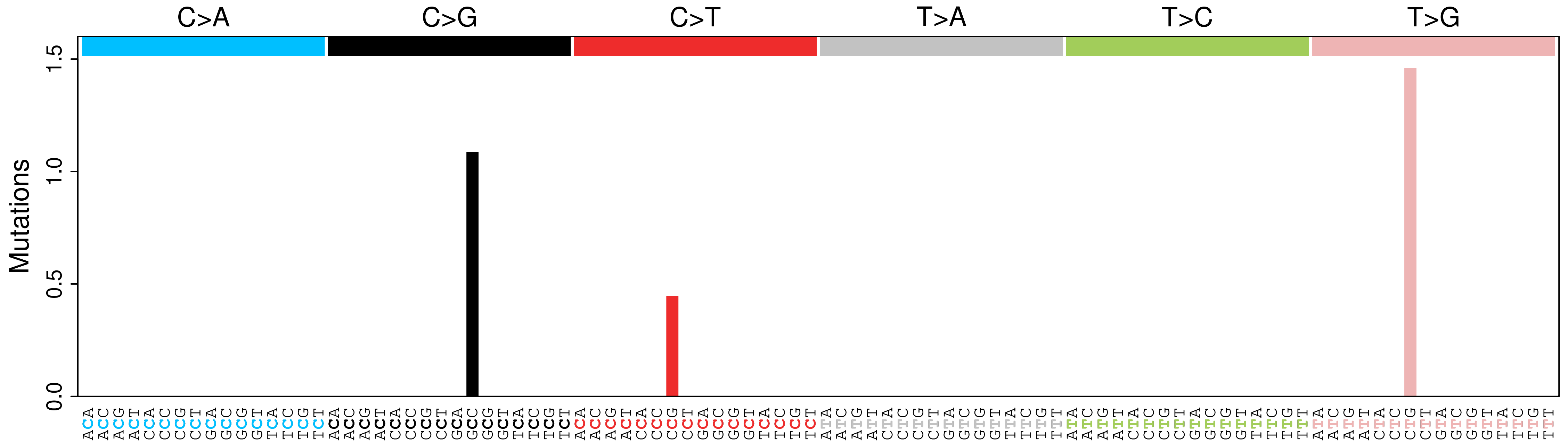
CATD0391a (2 mutations)



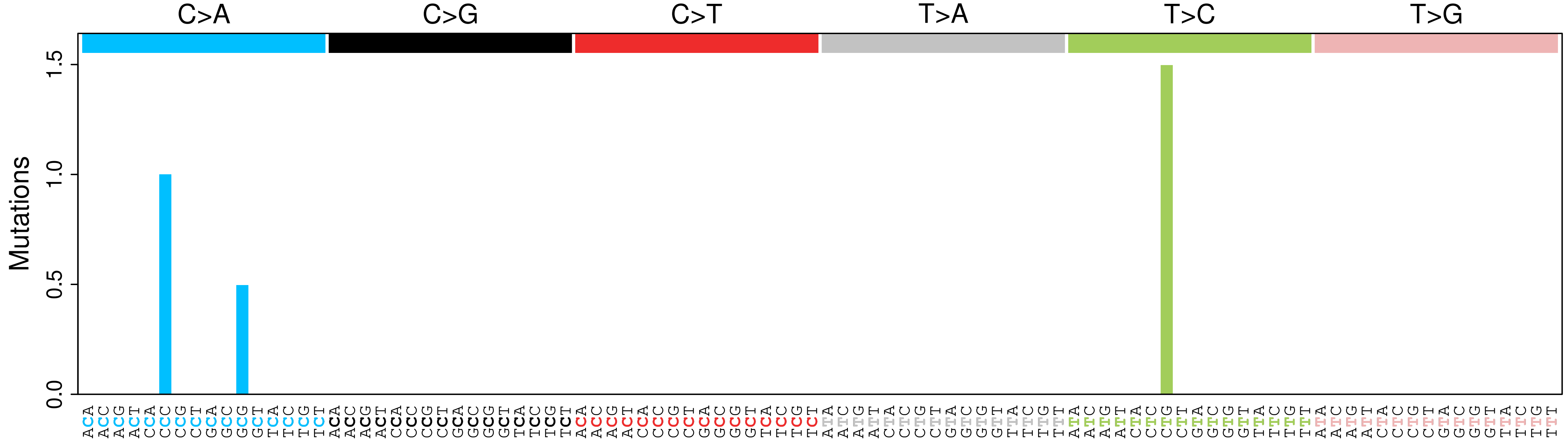
CATD0381a (7 mutations)



CATD0390a (3 mutations)



CATD0423a (3 mutations)



CATD0398a (3 mutations)

$C > A$

C > G

$$C > T$$

T > A

$$T > C$$

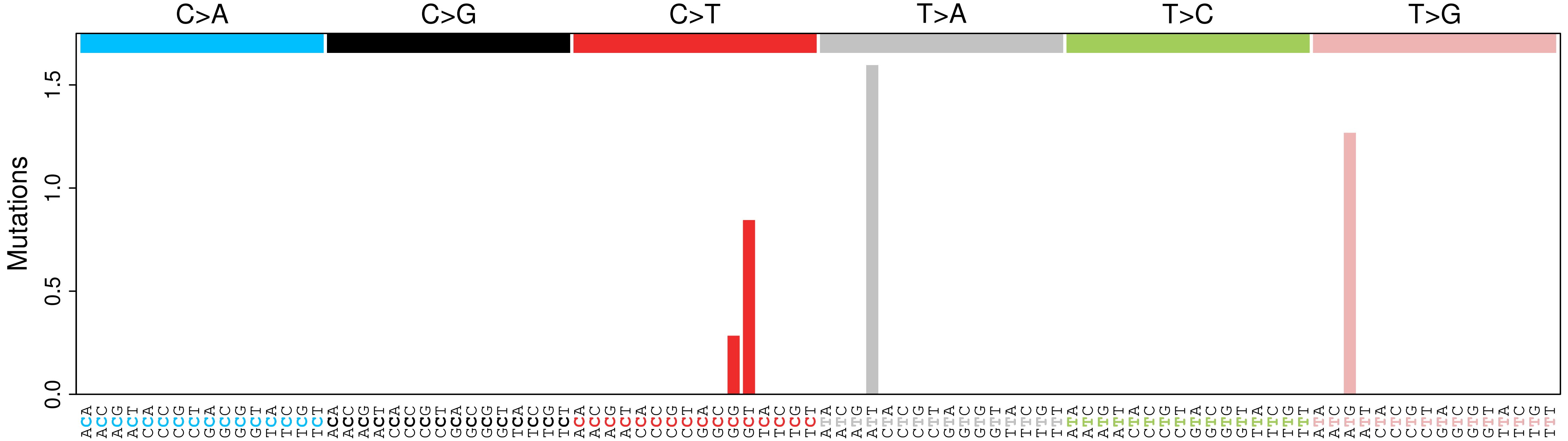
T > G

Mutations

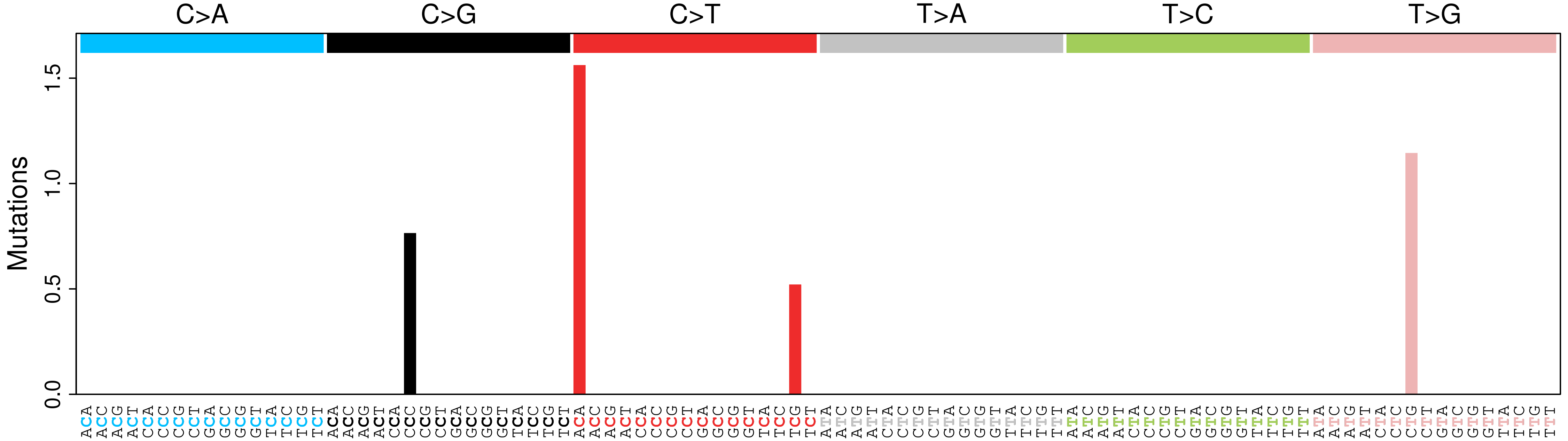
0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

[illegible]

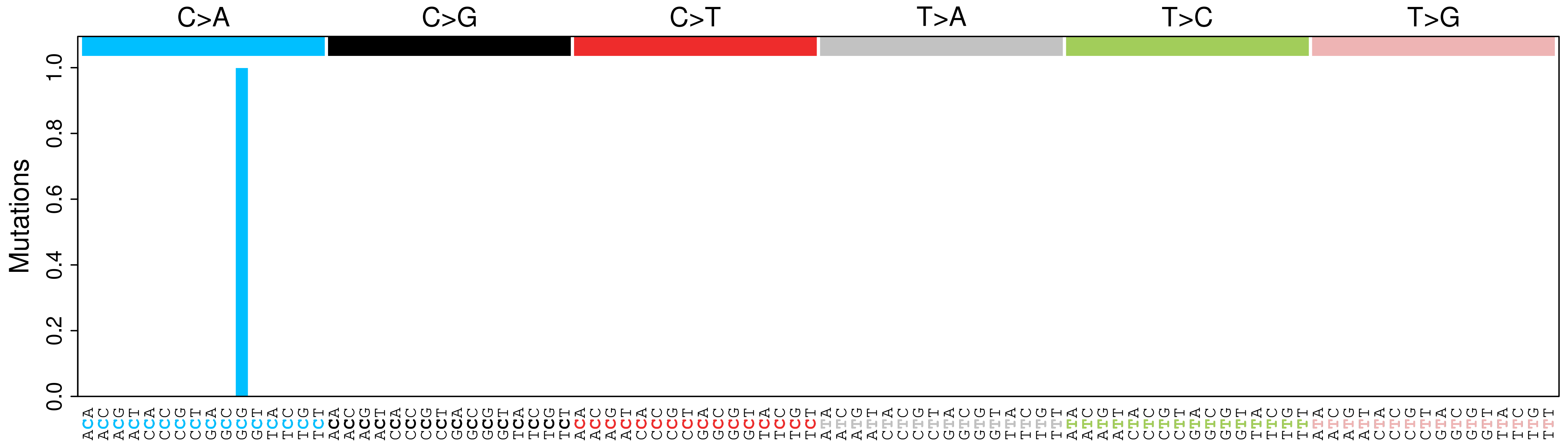
CATD0428a (4 mutations)



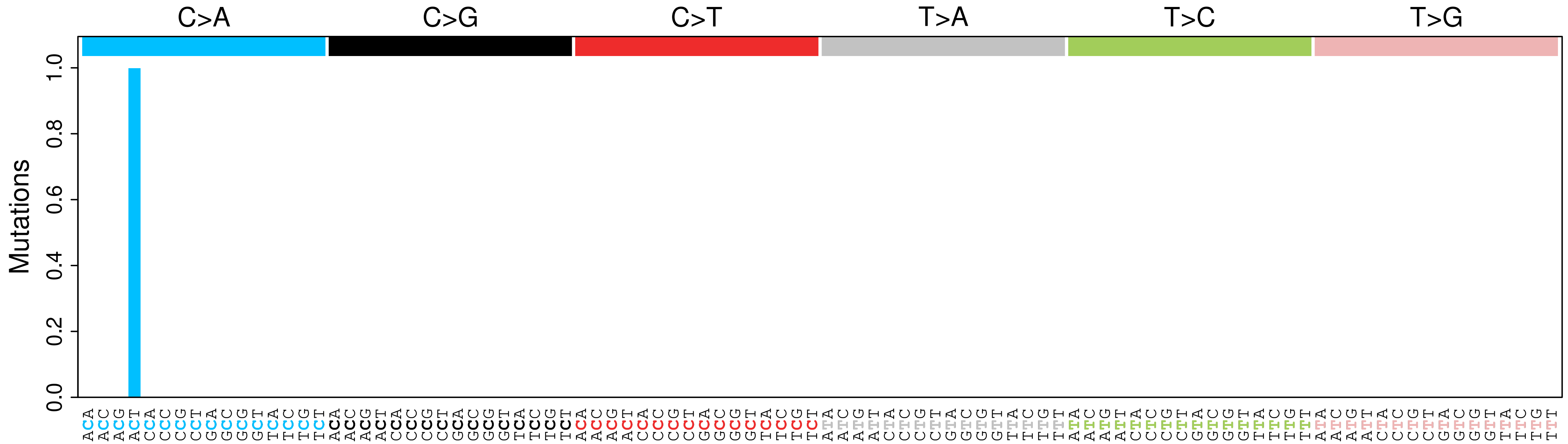
CATD0378a (4 mutations)



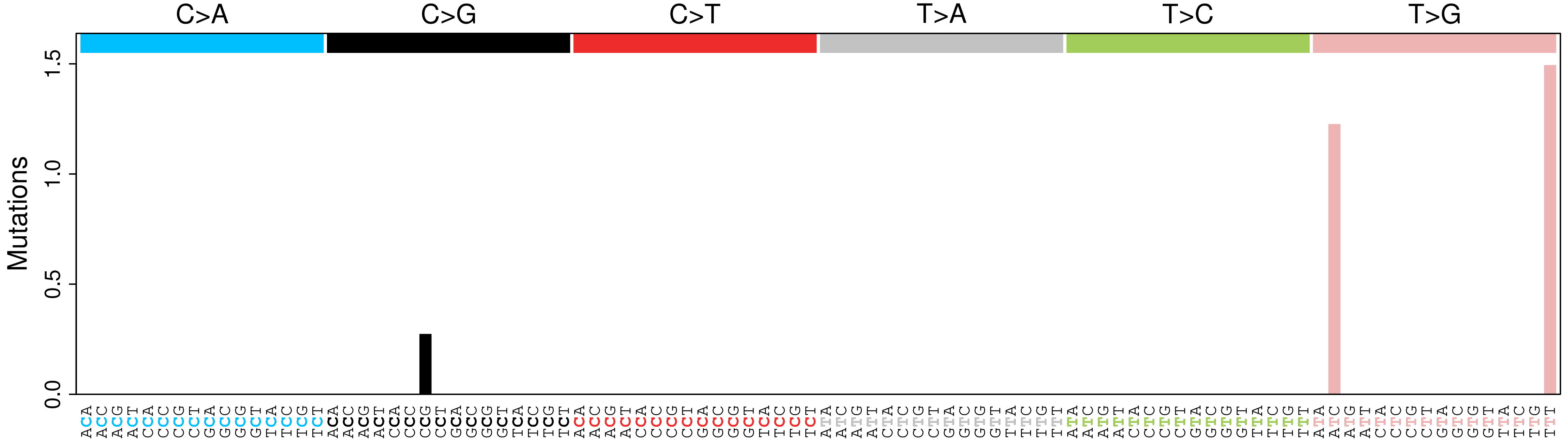
CATD0380a (1 mutations)



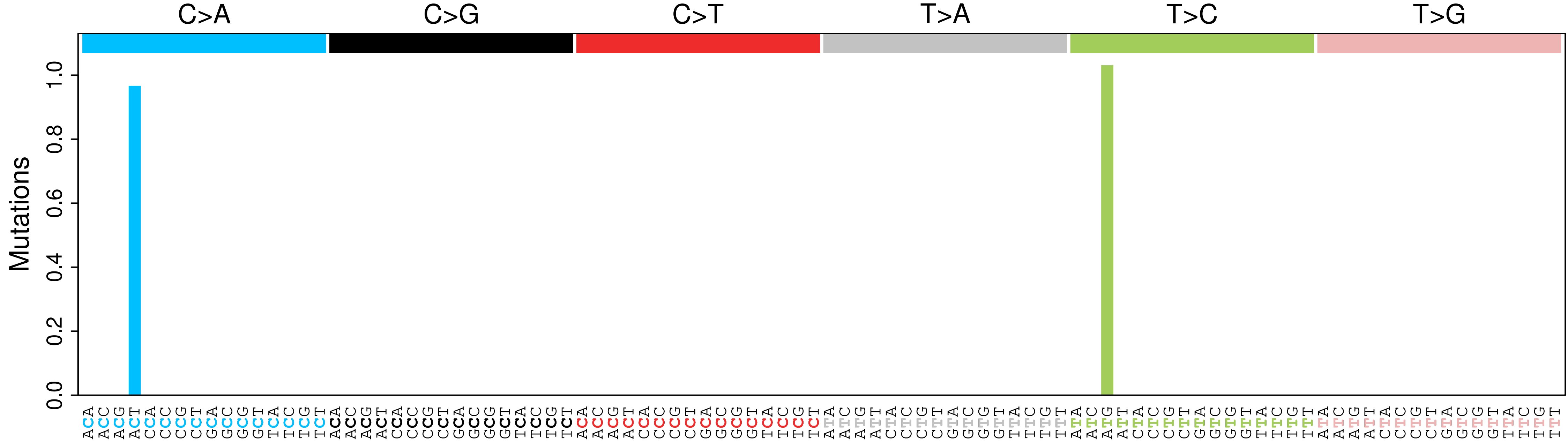
CATD0662a (1 mutations)



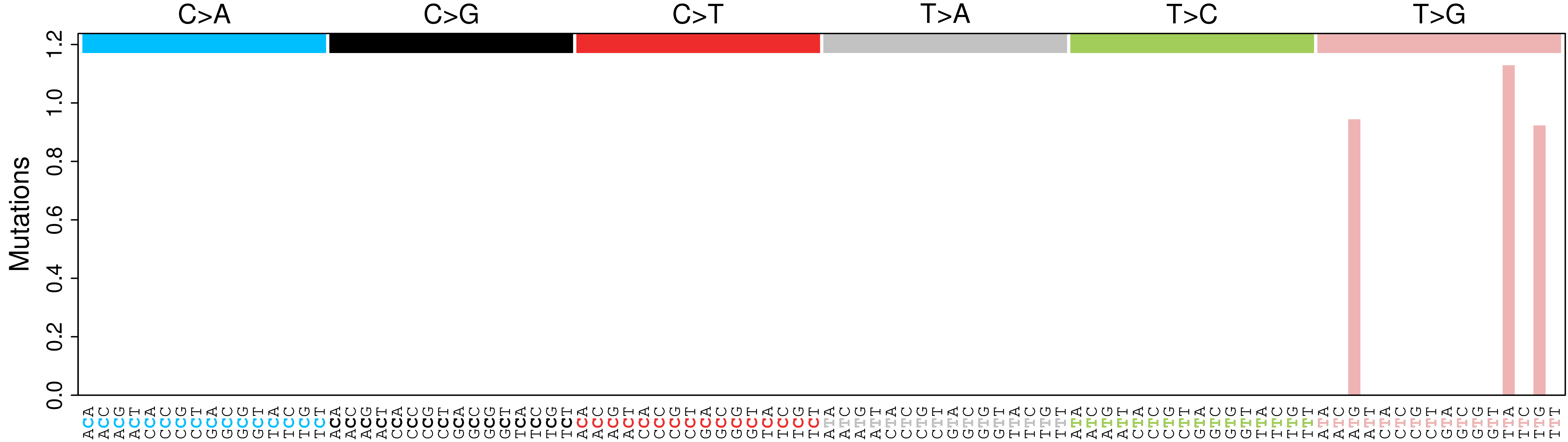
CATD0399a (3 mutations)



CATD0659a (2 mutations)



CATD0389a (3 mutations)



CATD0658a (2 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

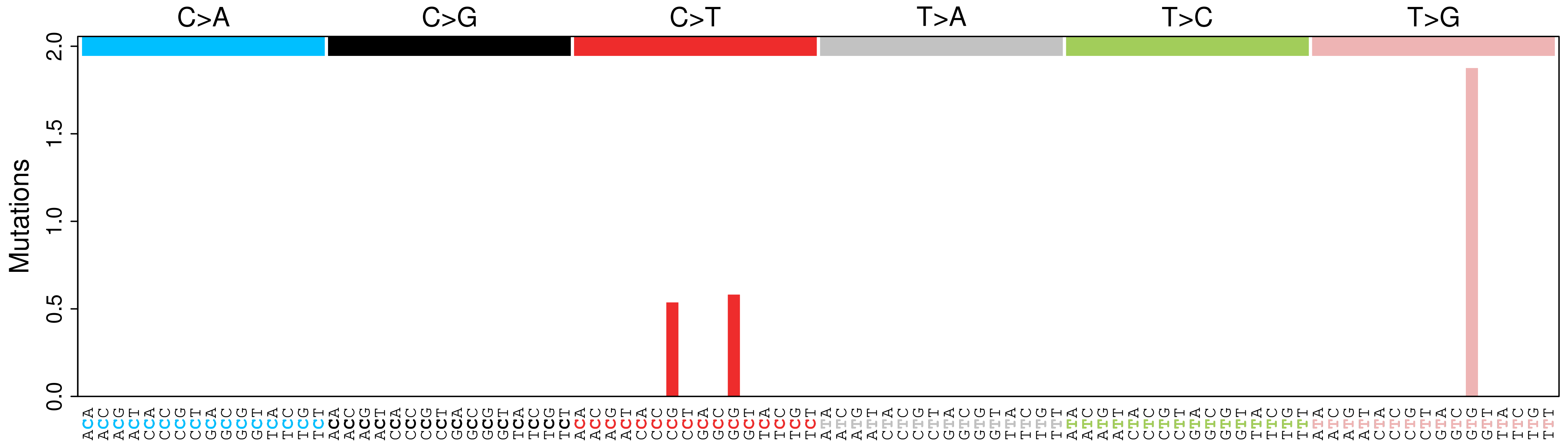
T > G

Mutations

0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

[illegible]

CATD0665a (3 mutations)



CATD0377a (3 mutations)

C > A

C > G

$C > T$

T > A

$T > C$

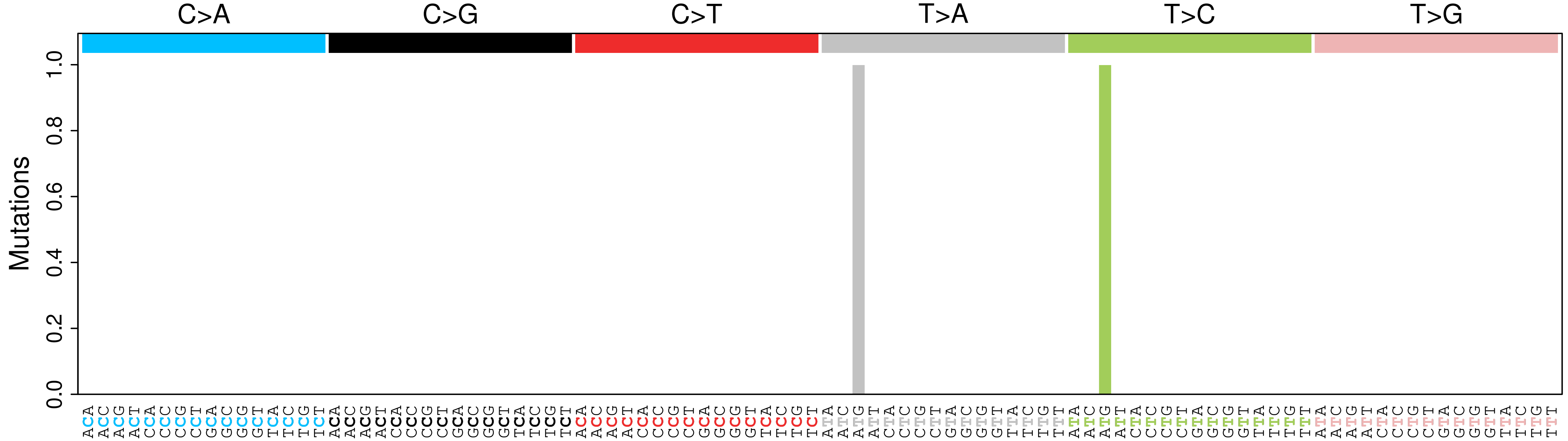
T>G

Mutations

0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

[illegible]

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)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

T > G

Mutations

α	$\langle n_s \rangle$
0.0	0.00
0.1	0.75
0.15	0.85
0.2	0.75
0.3	0.55
0.35	0.40
0.4	0.45
0.5	0.50
0.6	0.55
0.8	0.58
1.0	0.60
1.2	0.62

[illegible]

CATD0424a (1 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

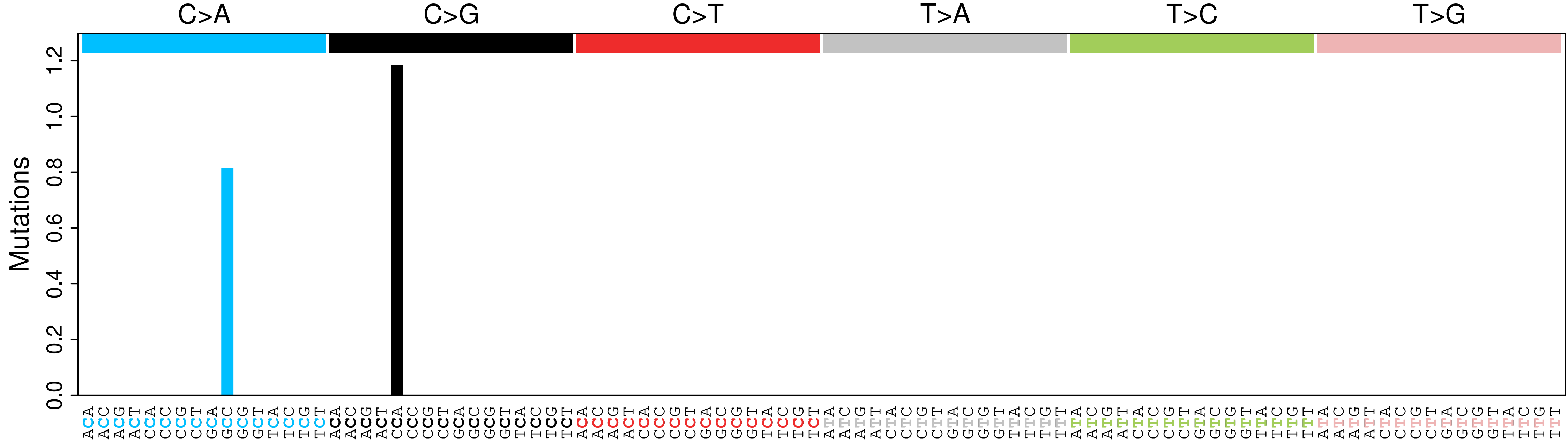
T > G

Mutations

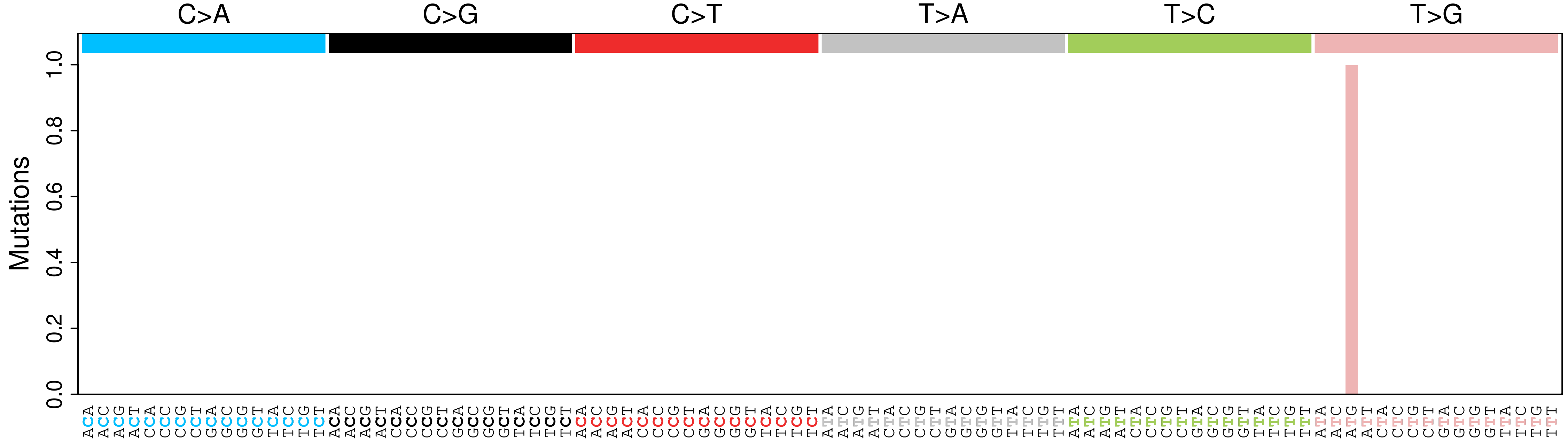
0.0 0.2 0.4 0.6 0.8 1.0

[illegible]

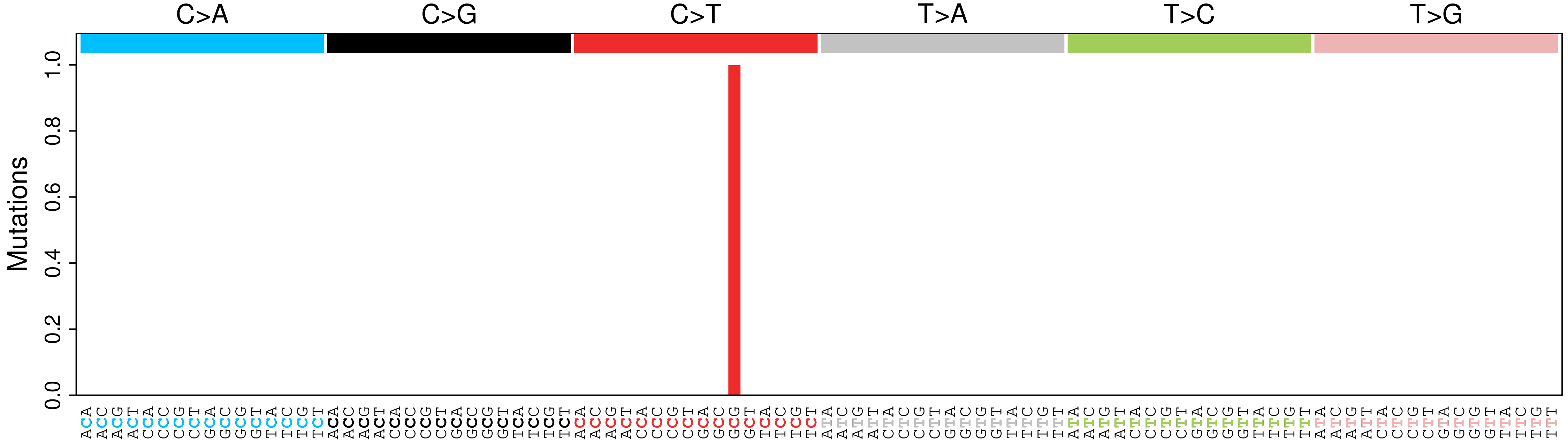
CATD0426a (2 mutations)



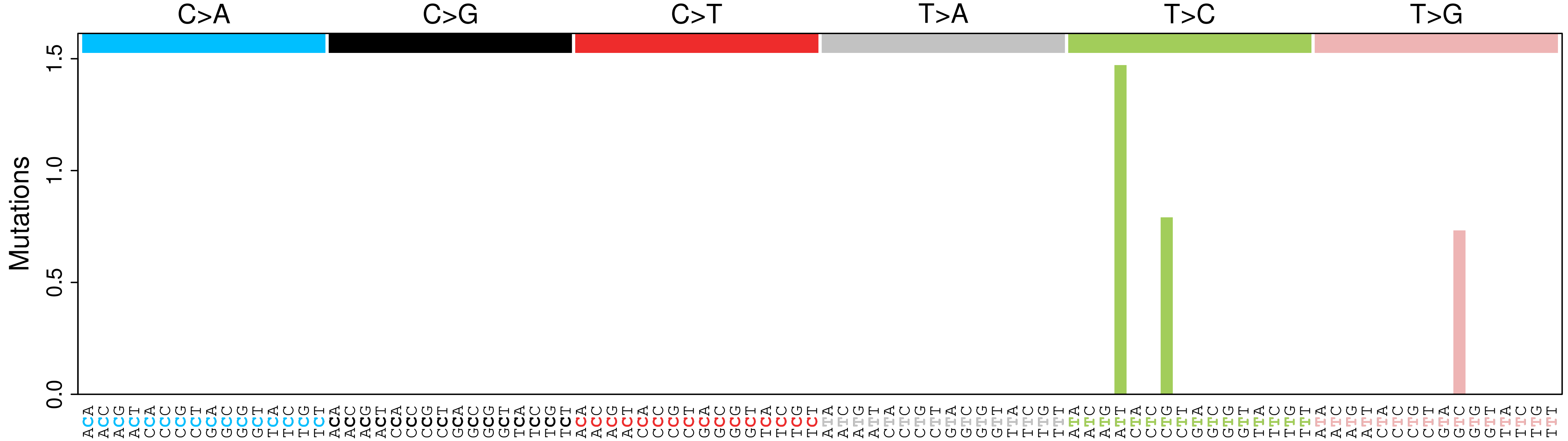
CATD0383a (1 mutations)



CATD0385a (1 mutations)



CATD0382a (3 mutations)



CATD0660a (1 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

T > G

Mutations

1.0

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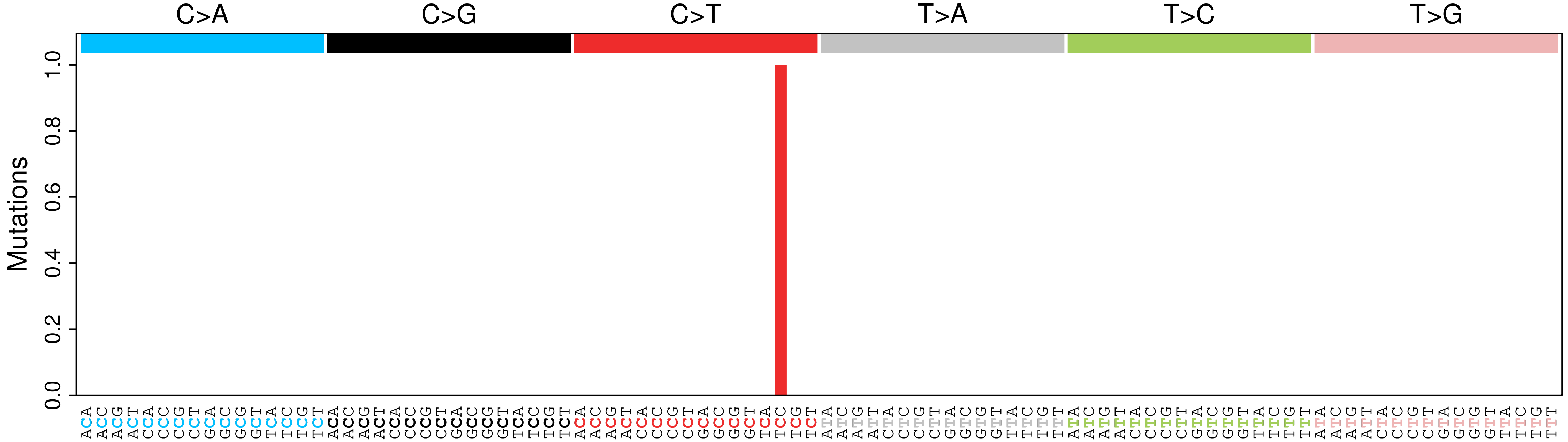
1

2

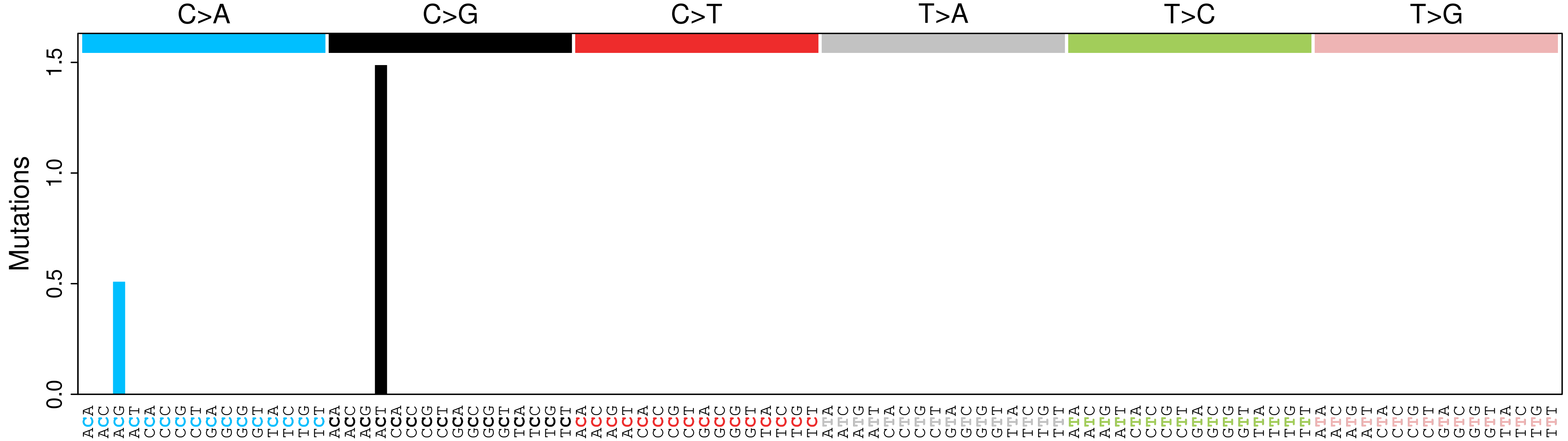
0.

[illegible]

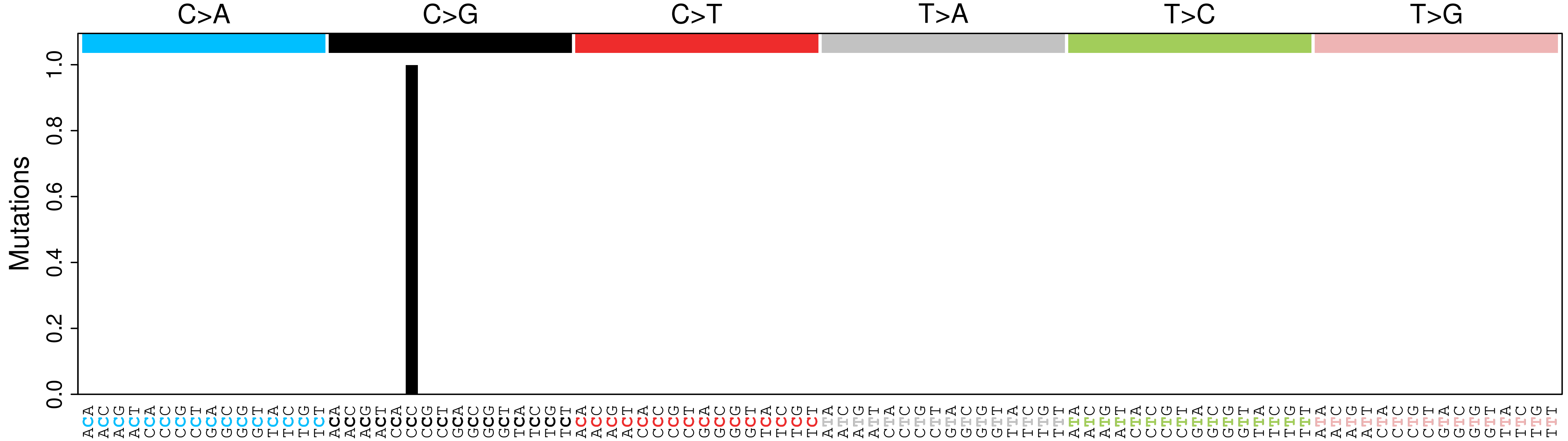
CATD0437a (1 mutations)



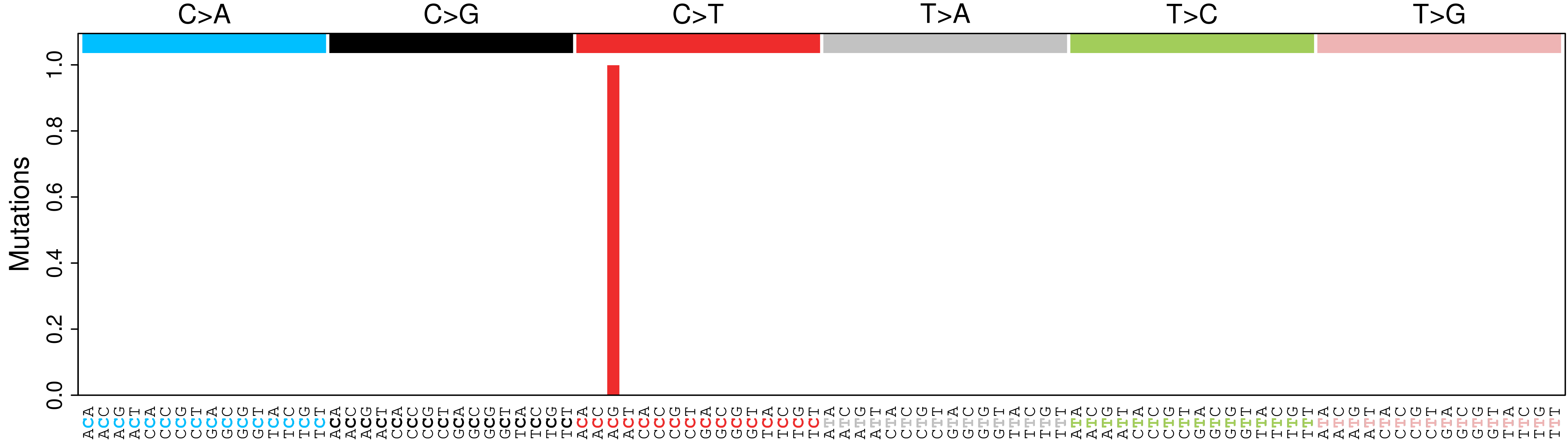
CATD0439a (2 mutations)



CATD0663a (1 mutations)



CATD0379a (1 mutations)



CATD0397a (1 mutations)

