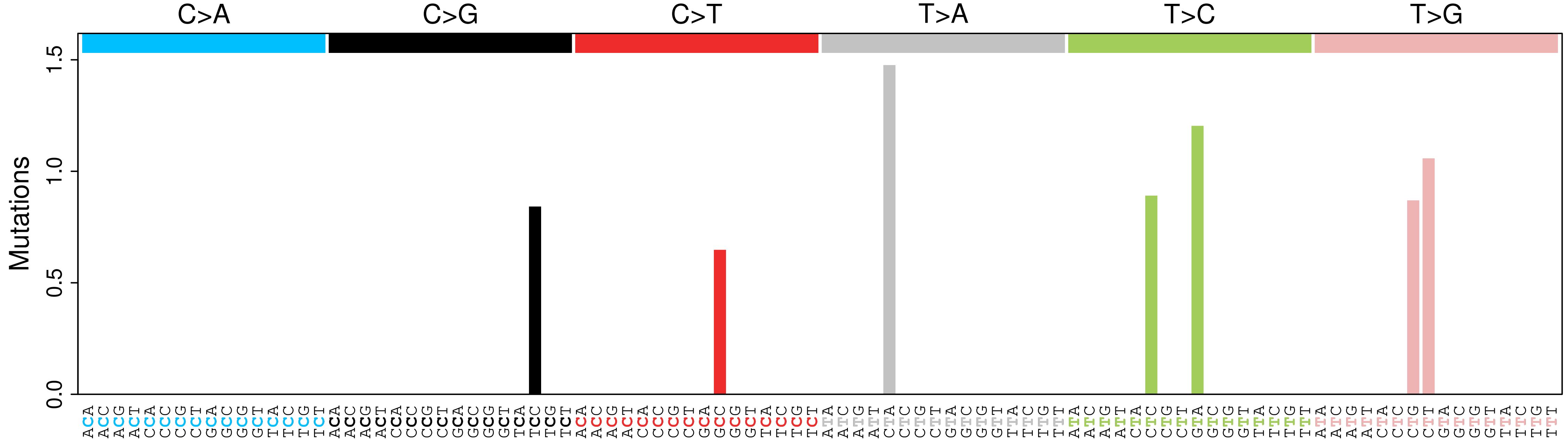
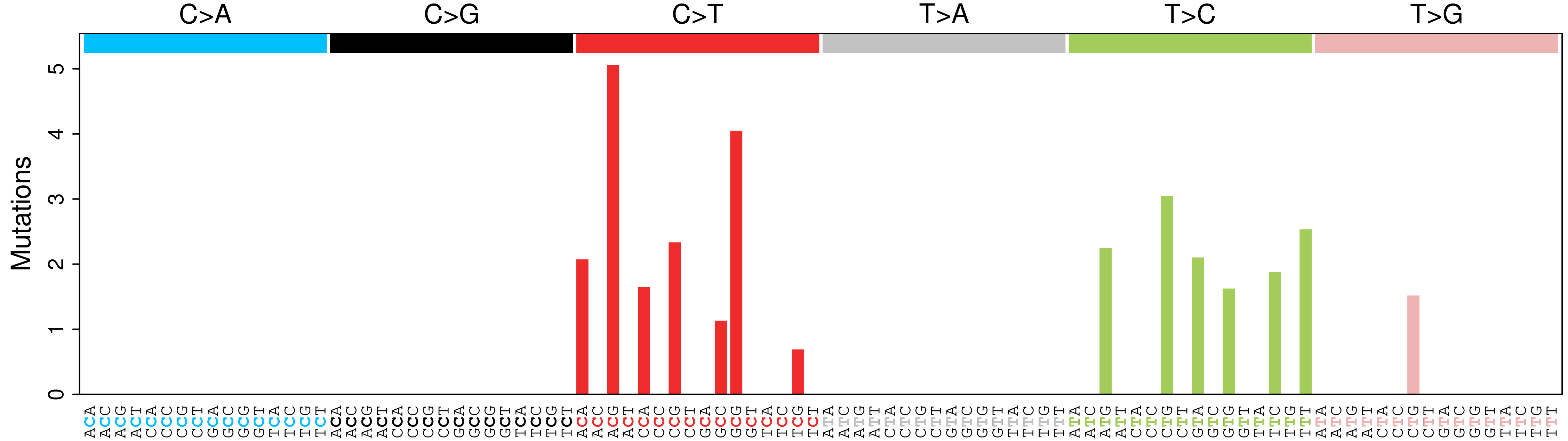


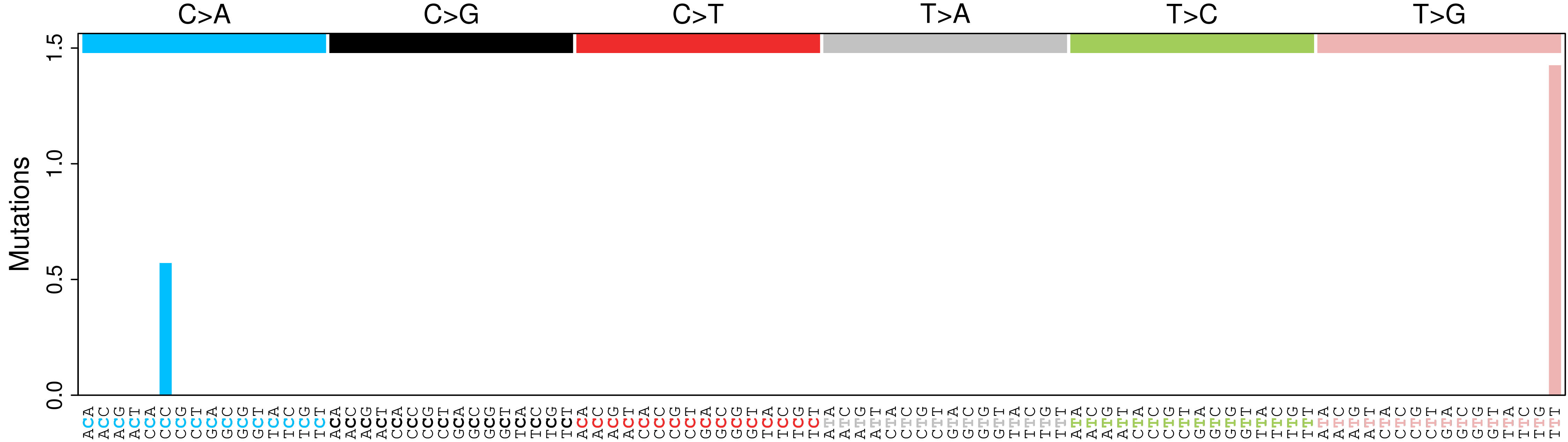
CATD0507a (7 mutations)



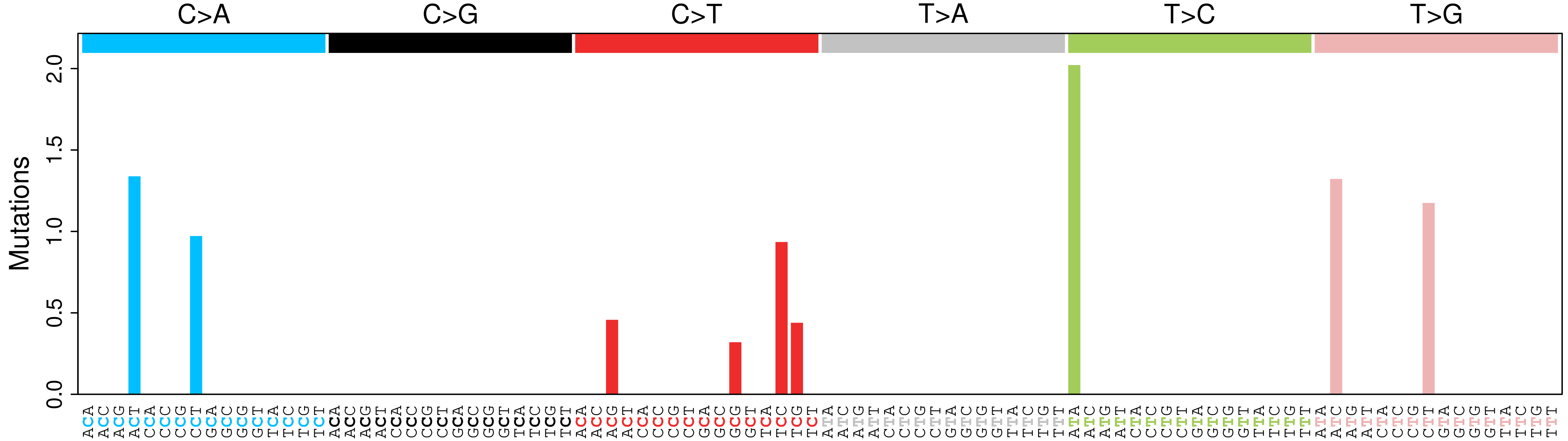
CATD0500a (32 mutations)



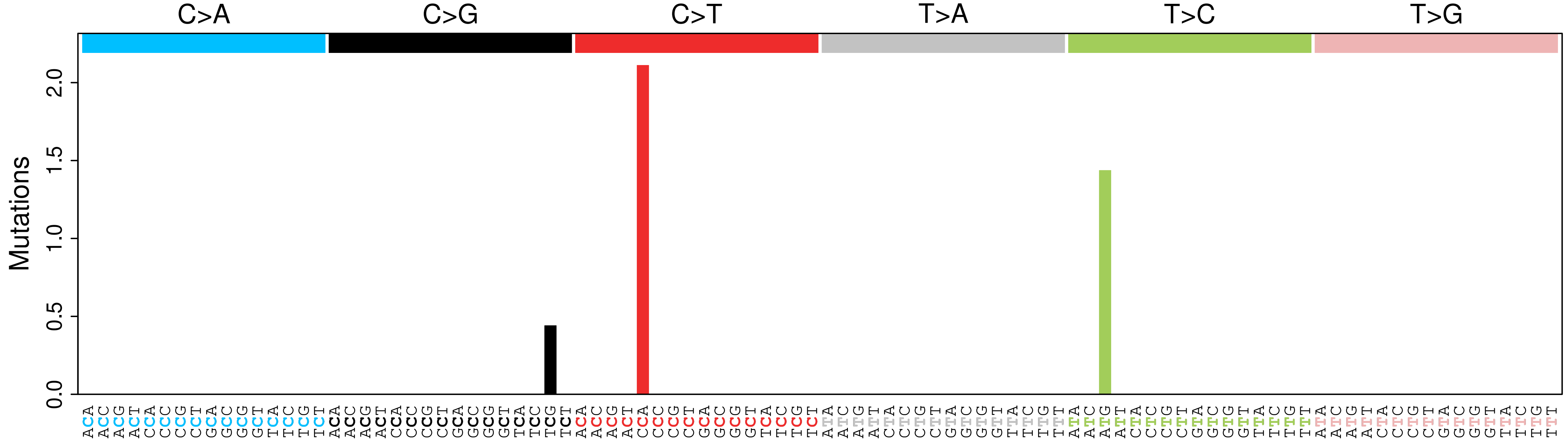
CATD0467a (2 mutations)



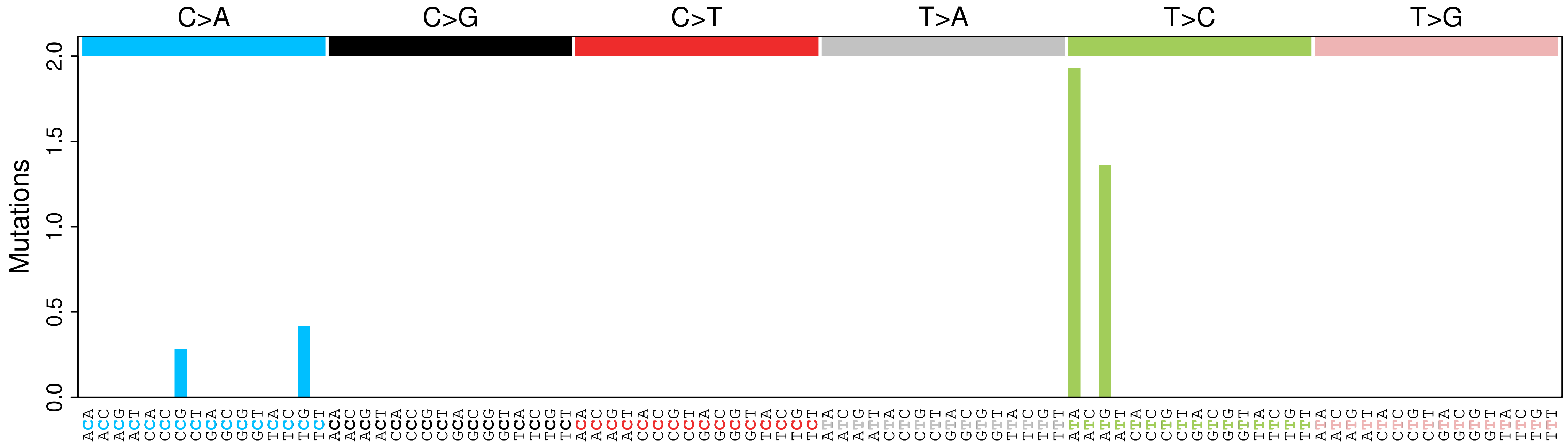
CATD0698a (9 mutations)



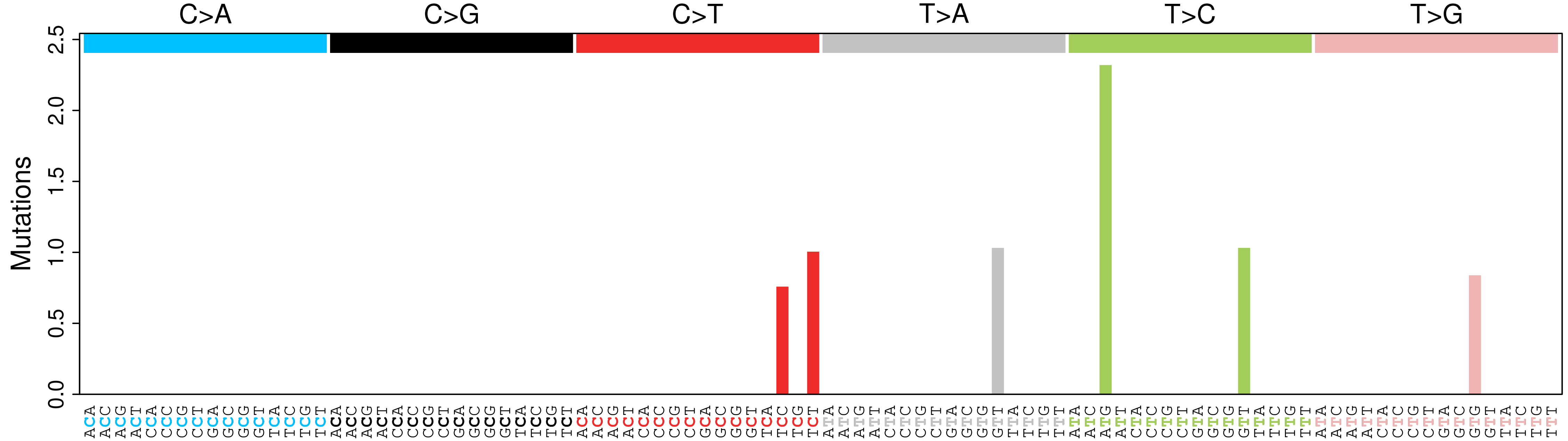
CATD0480a (4 mutations)



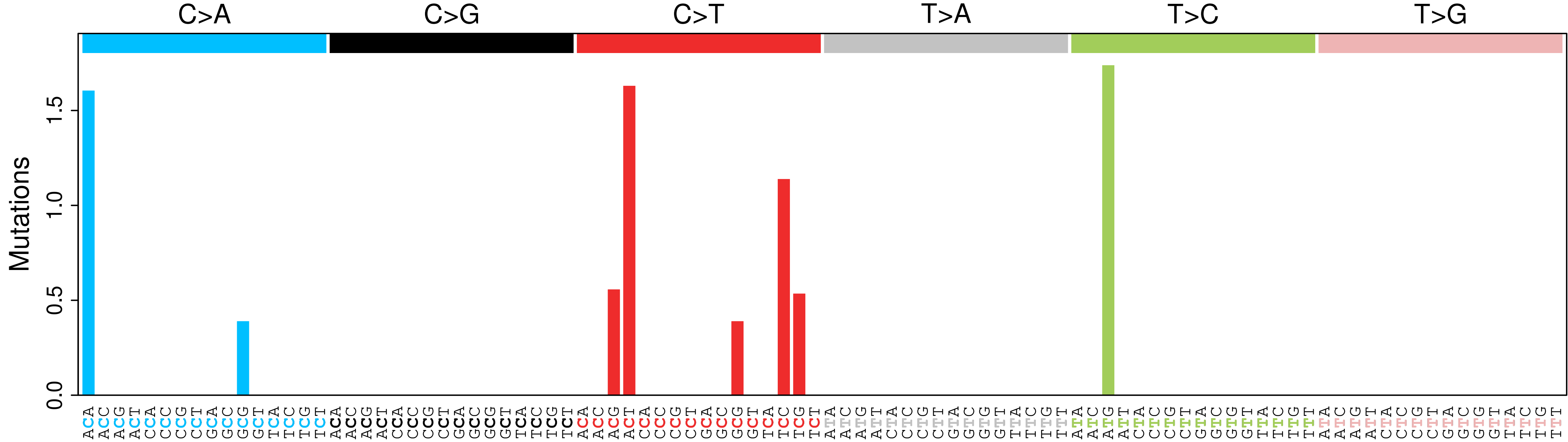
CATD0491a (4 mutations)



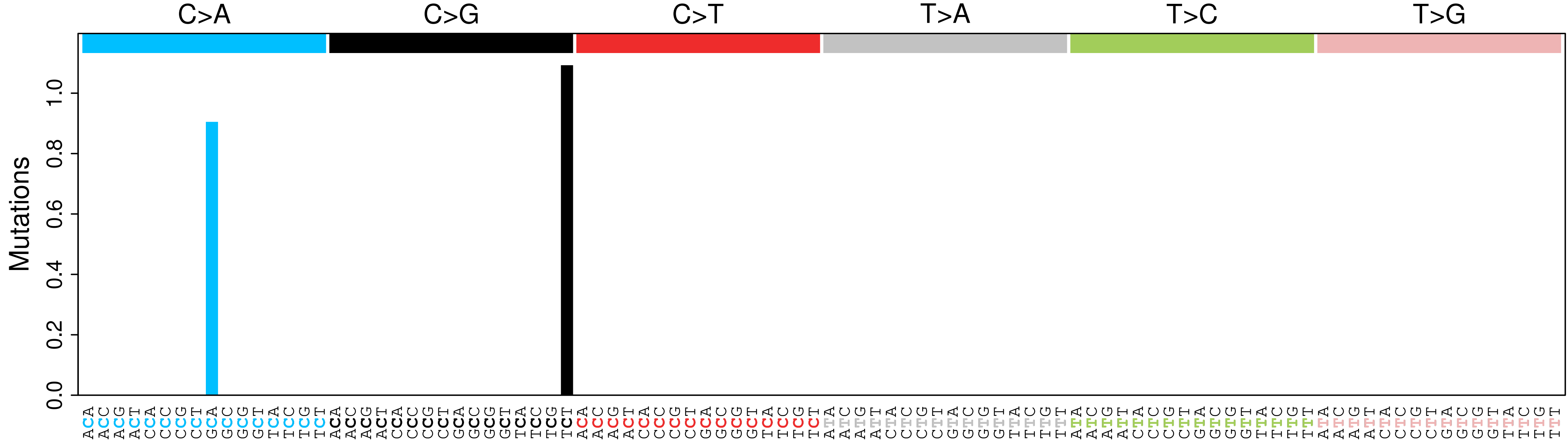
CATD0689a (7 mutations)



CATD0701a (8 mutations)



CATD0495a (2 mutations)



CATD0702a (7 mutations)

C>A

C>G

C>T

T>A

T>C

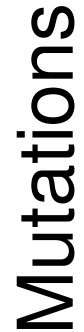
T>G

Mutations

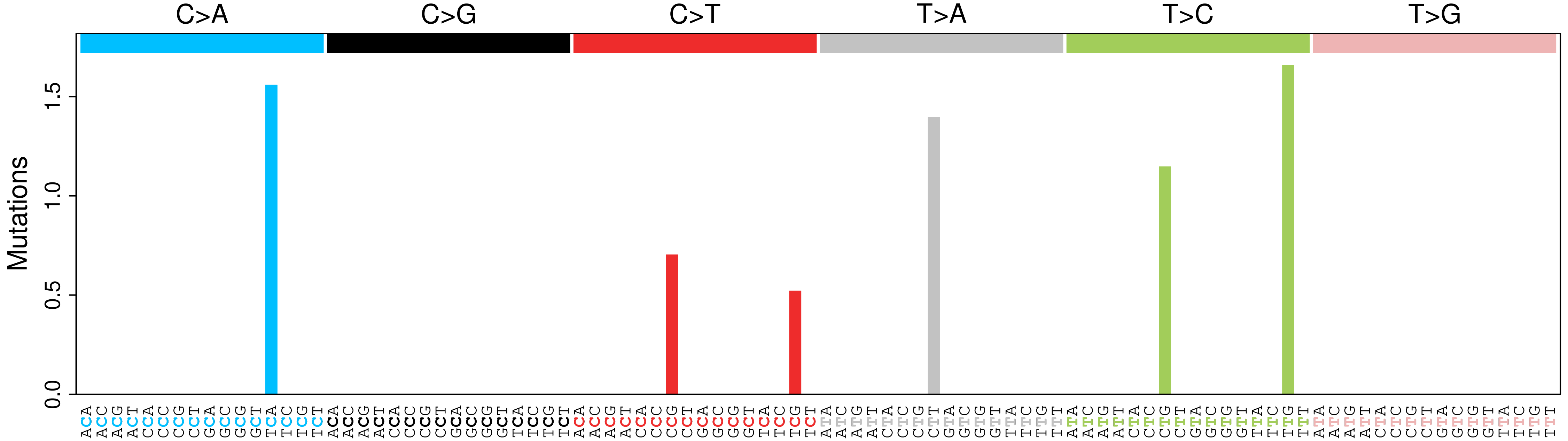
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CATD0505a (8 mutations)



CATD0482a (7 mutations)



CATD0694a (4 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

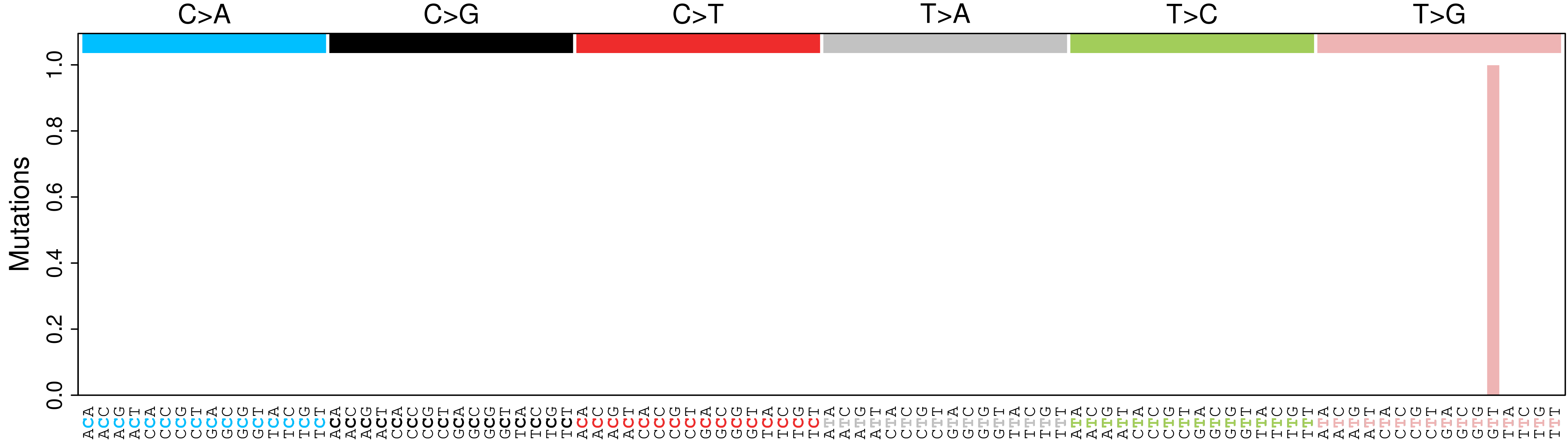
T > G

Mutations

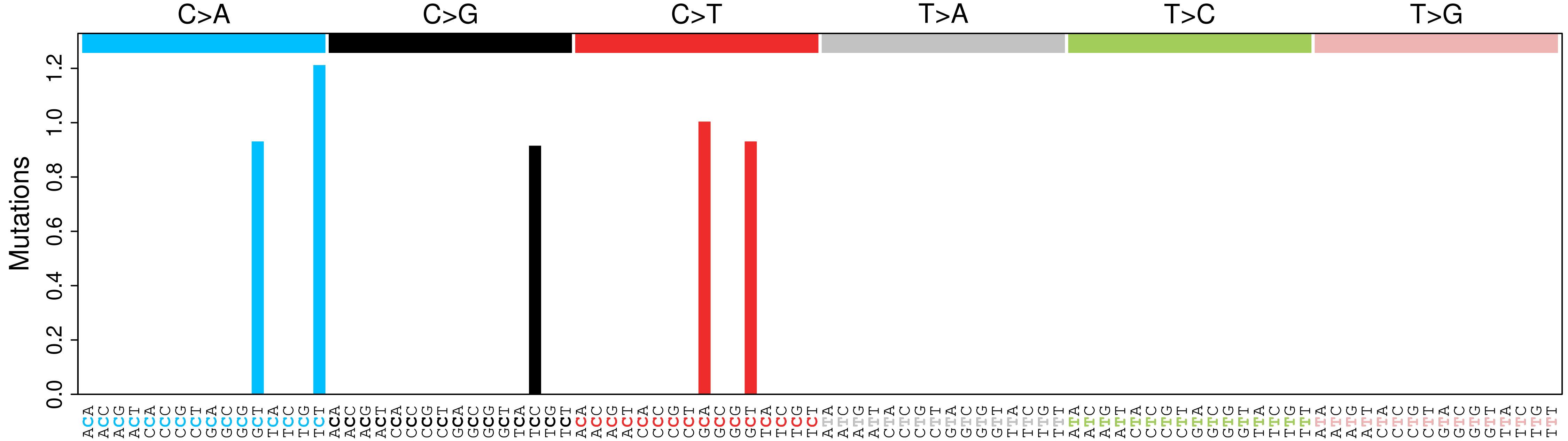
0.0 0.5 1.0 1.5

[illegible]

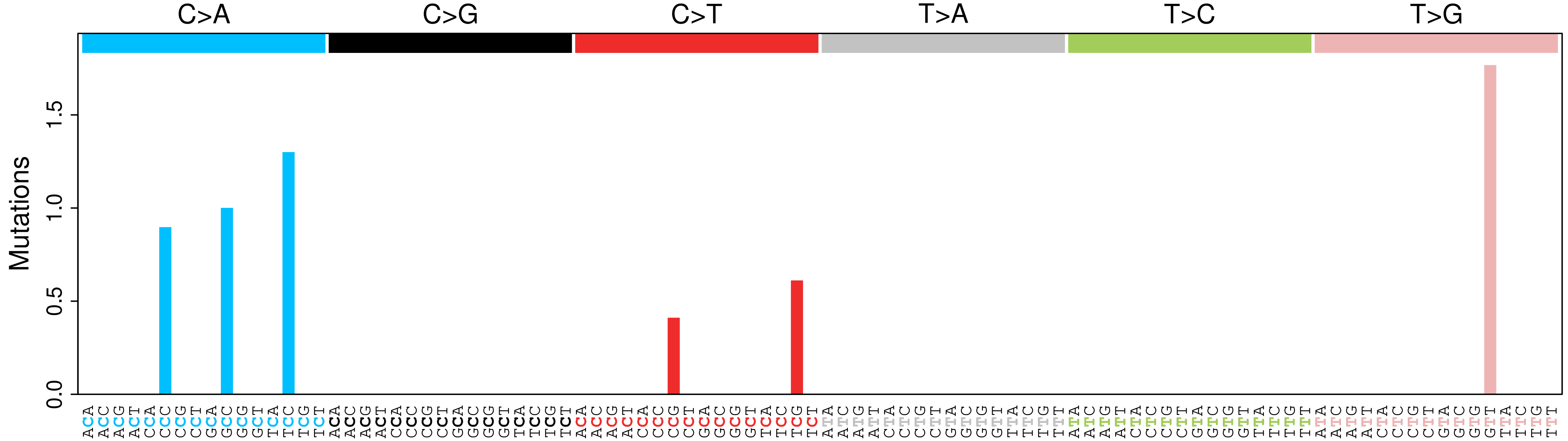
CATD0499a (1 mutations)



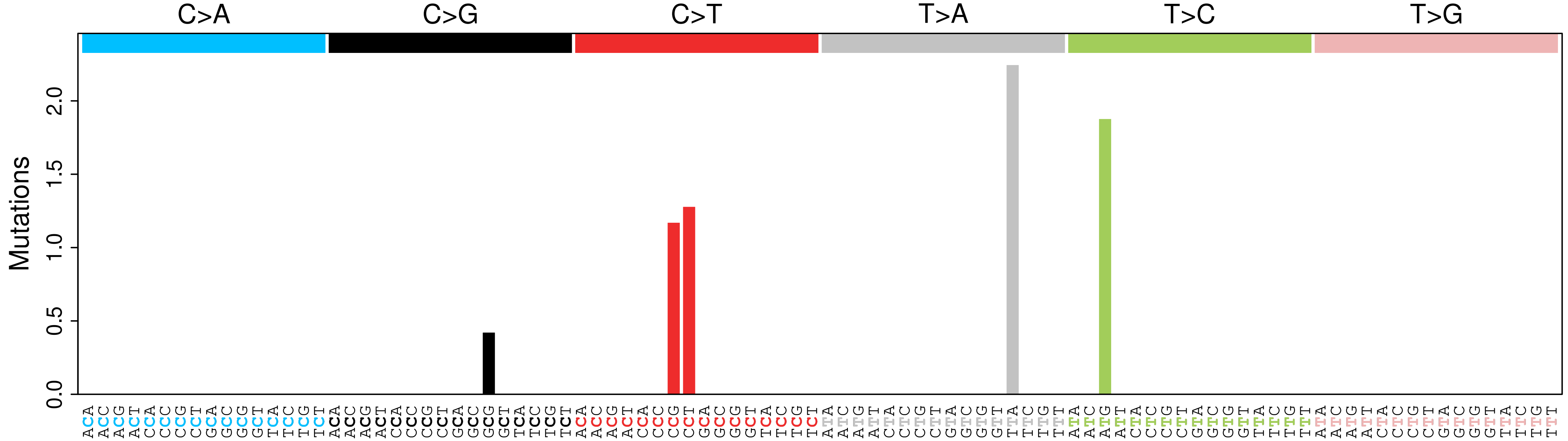
CATD0504a (5 mutations)



CATD0700a (6 mutations)



CATD0469a (7 mutations)



CATD0468a (3 mutations)

$C > A$

C > G

C>T

T > A

$$T > C$$

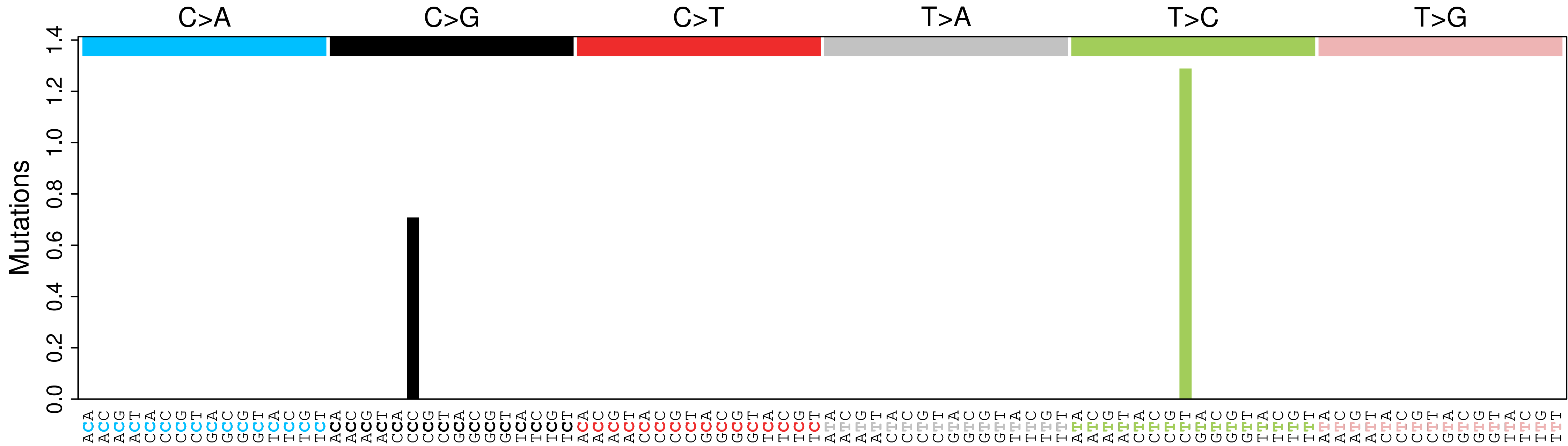
T>G

Mutations

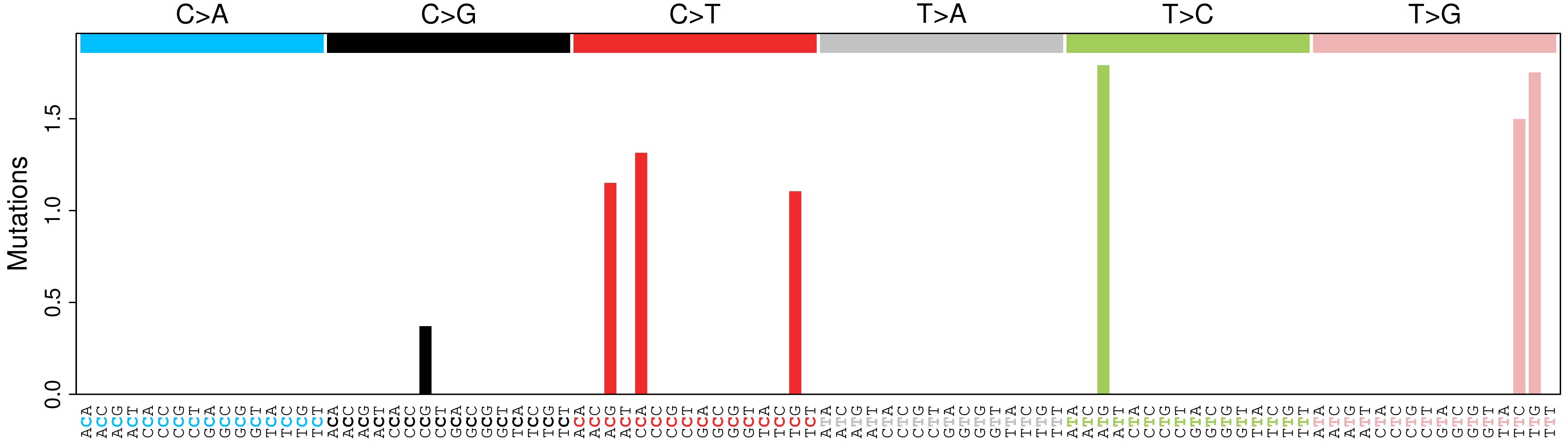
0.0 0.2 0.4 0.6 0.8 1.0 1.2

[illegible]

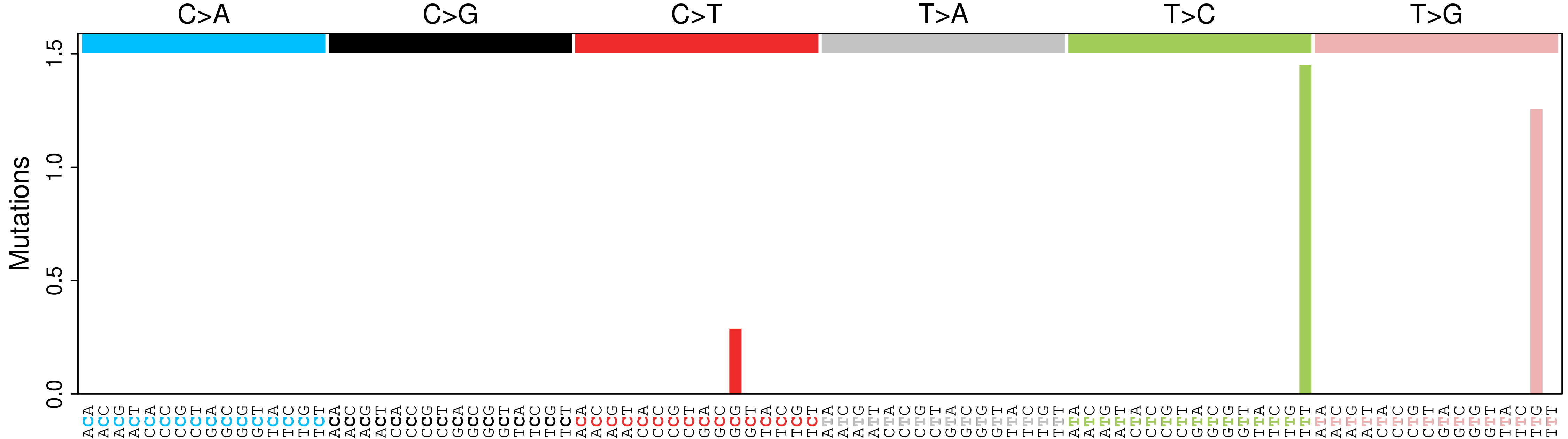
CATD0691a (2 mutations)



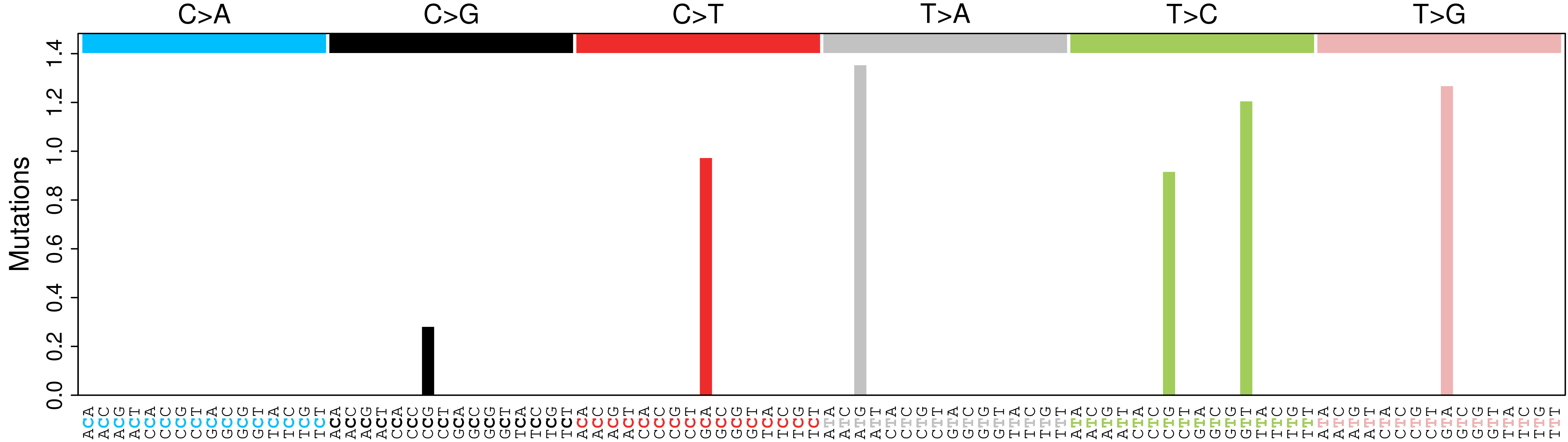
CATD0479a (9 mutations)



CATD0688a (3 mutations)



CATD0494a (6 mutations)



CATD0742a (2 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

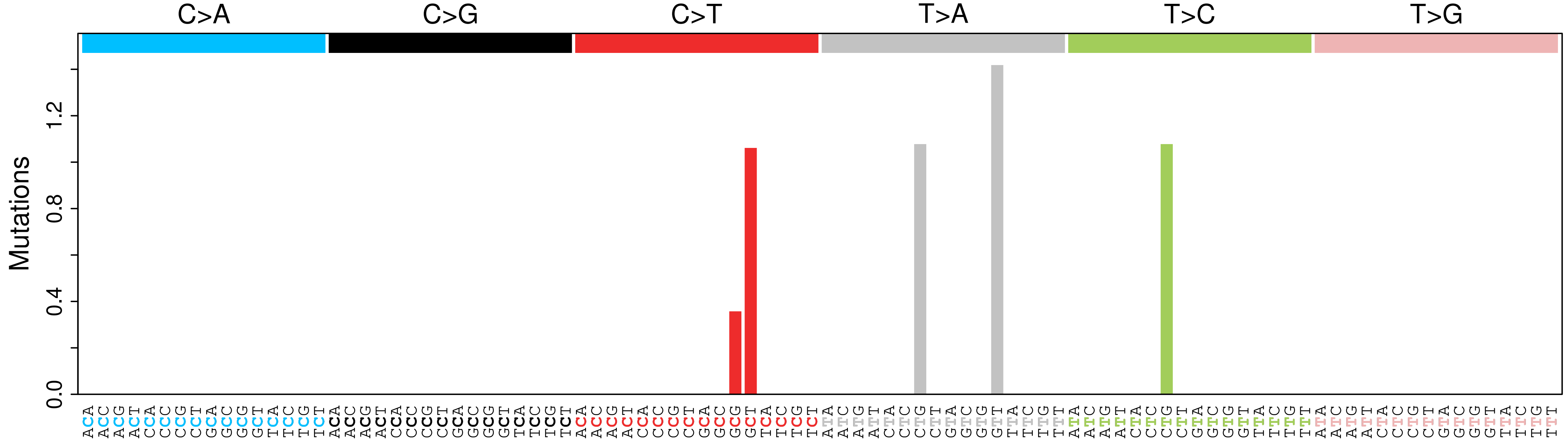
T > G

Mutations

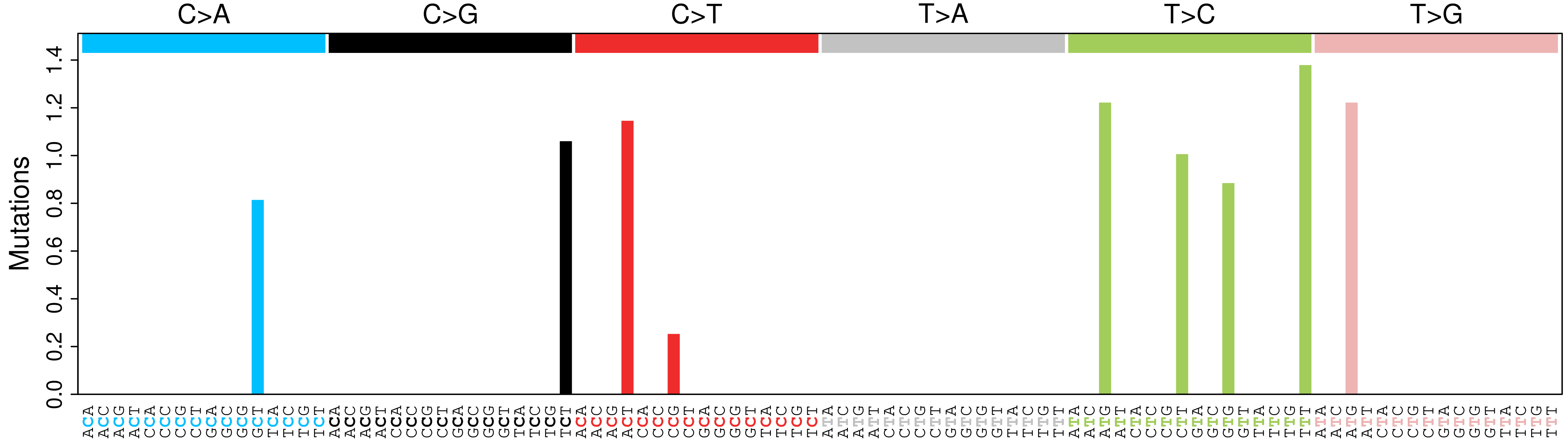
0.0 0.2 0.4 0.6 0.8 1.0 1.2

[illegible]

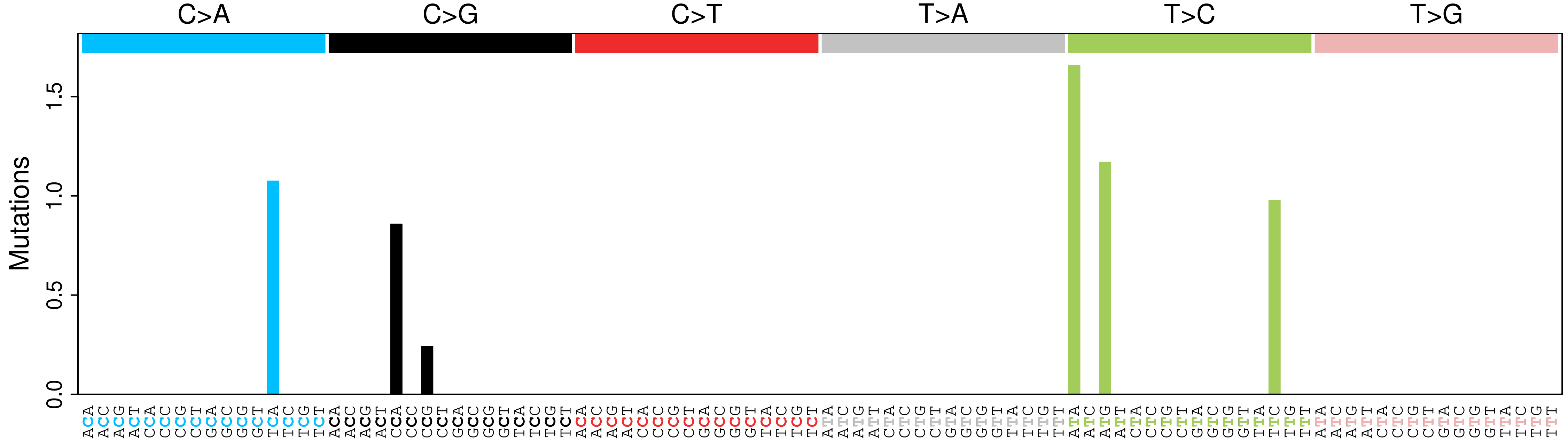
CATD0746a (5 mutations)



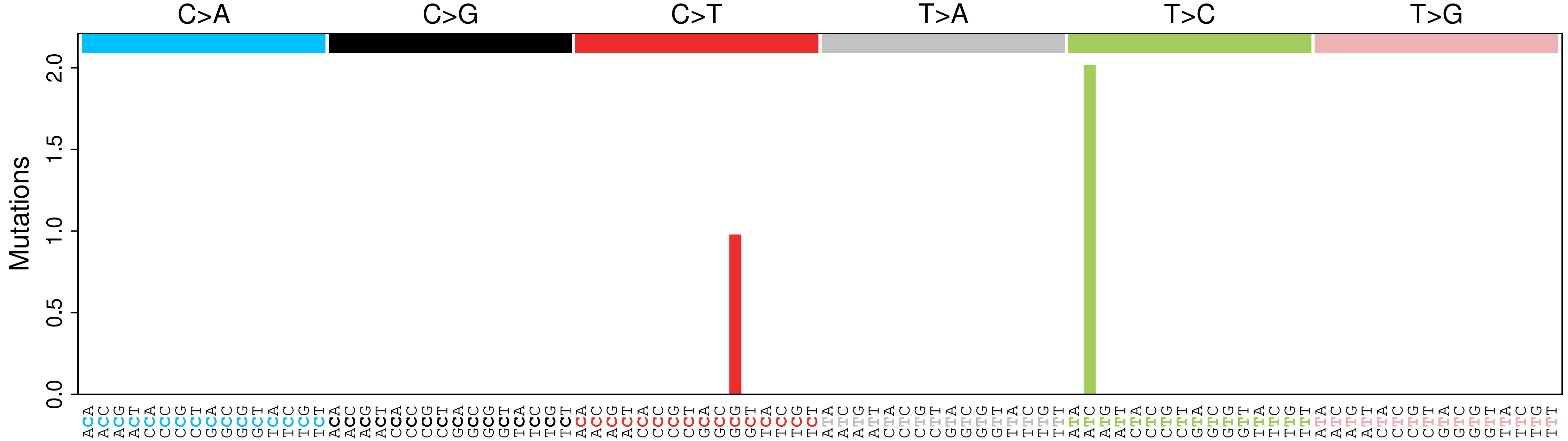
CATD0745a (9 mutations)



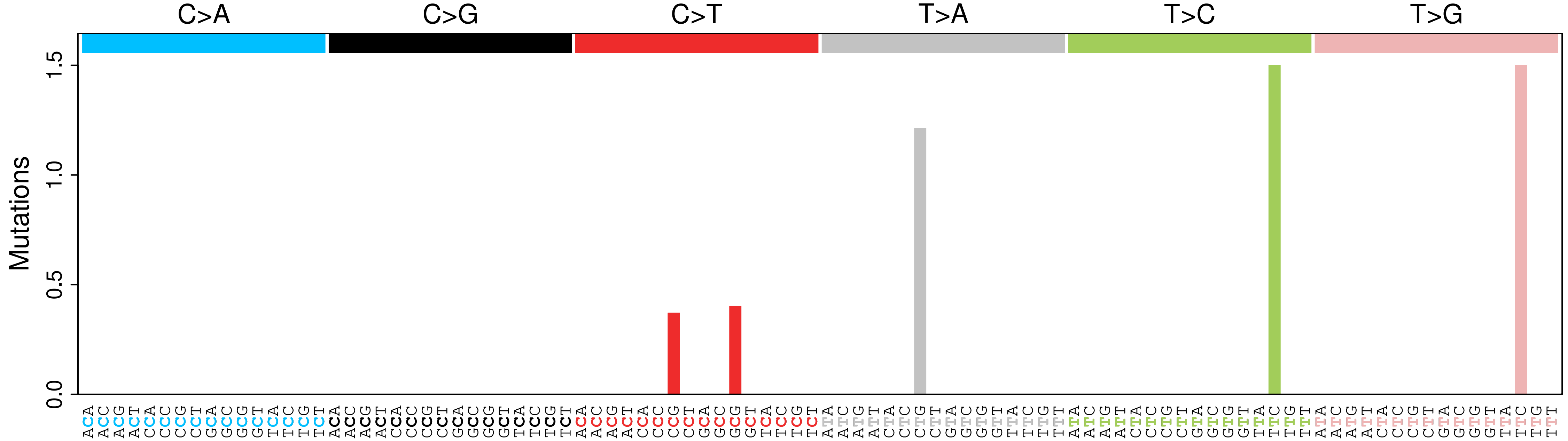
CATD0475a (6 mutations)



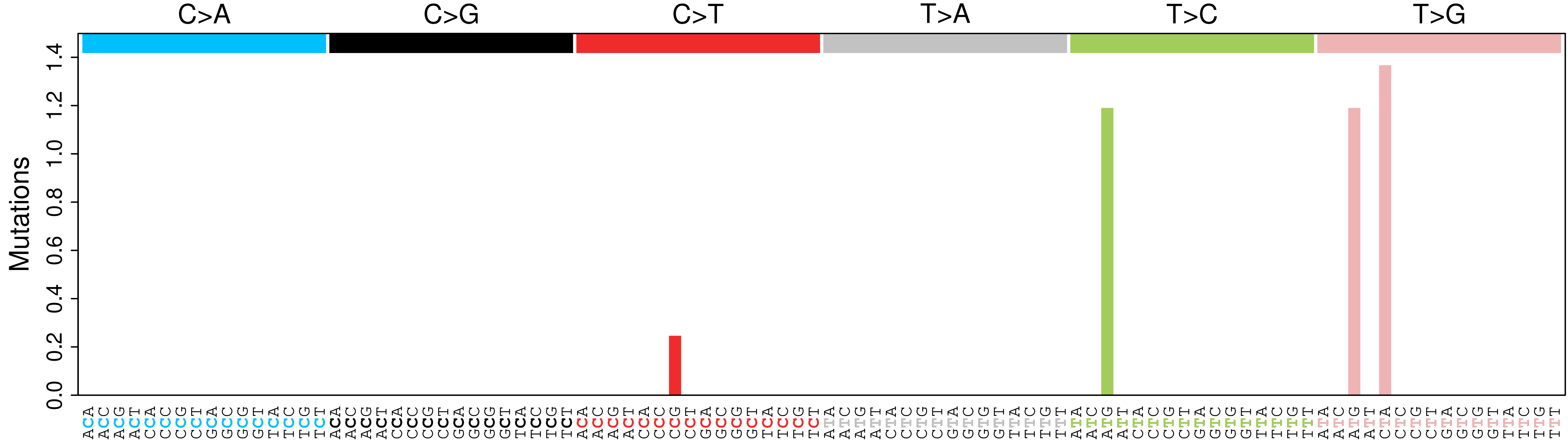
CATD0503a (3 mutations)



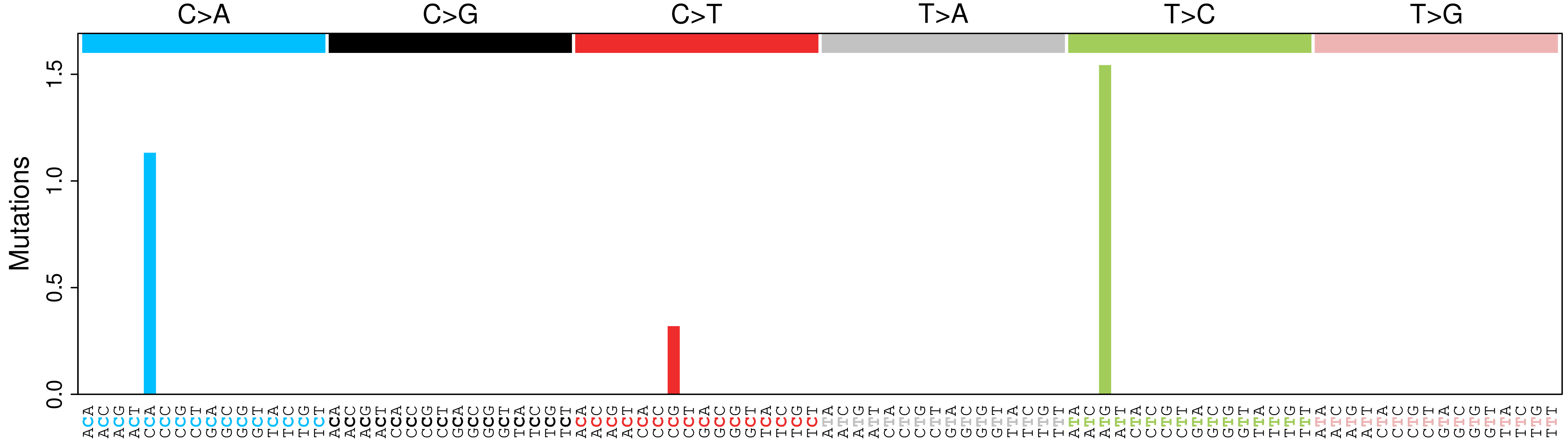
CATD0743a (5 mutations)



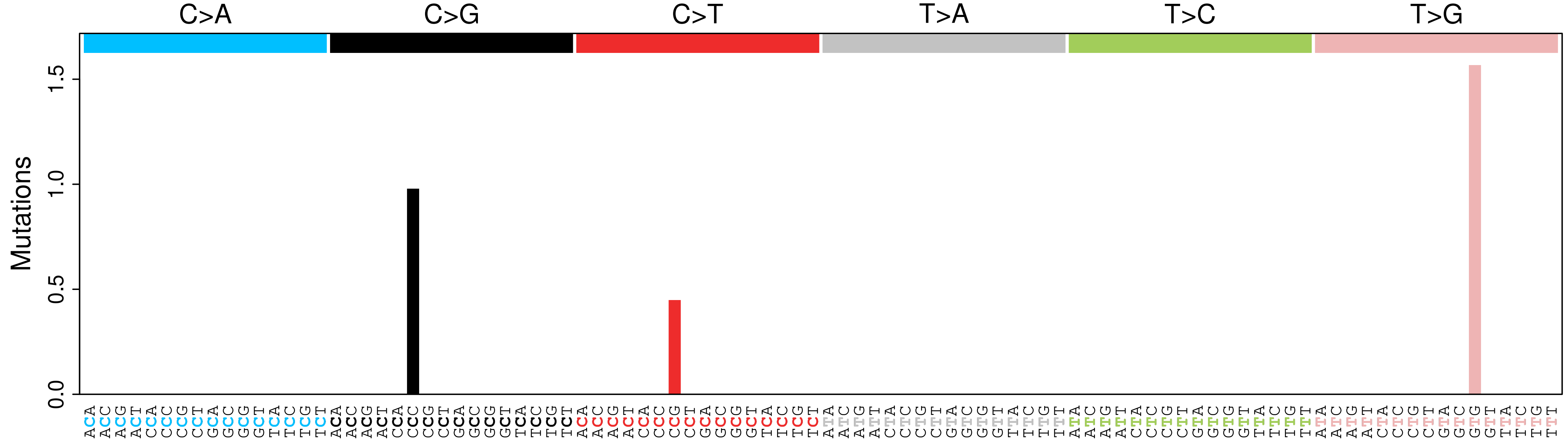
CATD0478a (4 mutations)



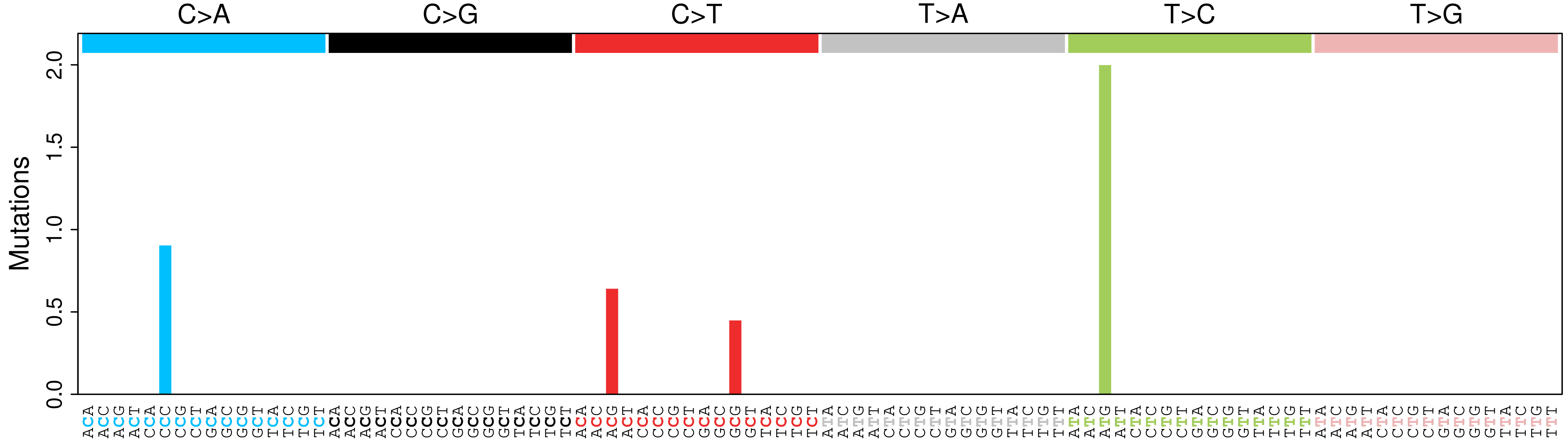
CATD0488a (3 mutations)



CATD0697a (3 mutations)



CATD0485a (4 mutations)



CATD0486a (4 mutations)

C>A

C>G

C>T

T>A

T>C

T>G

Mutations

1.2

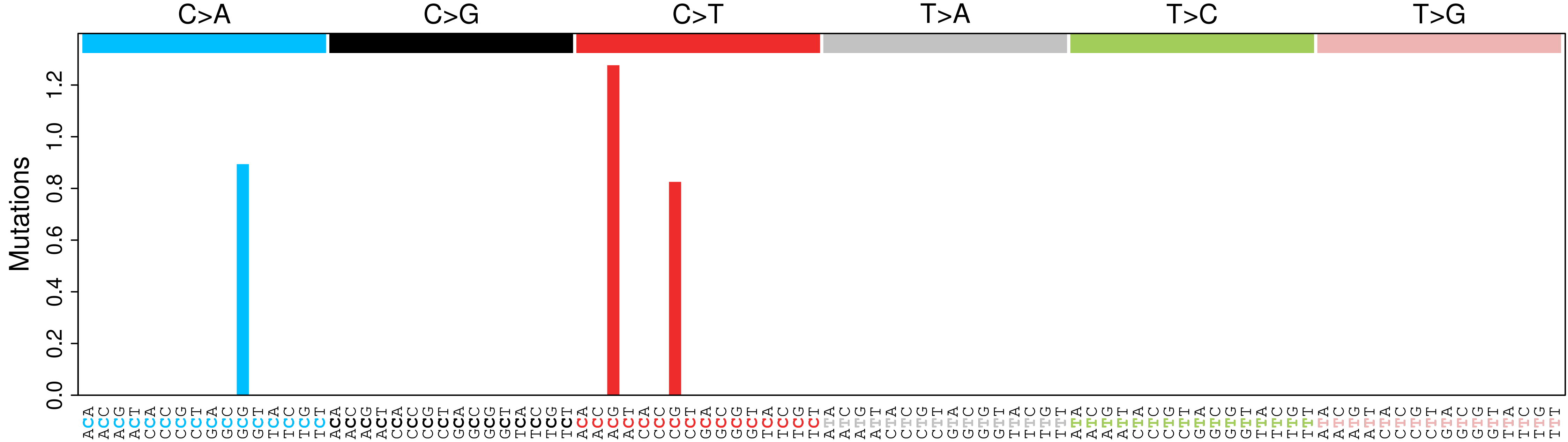
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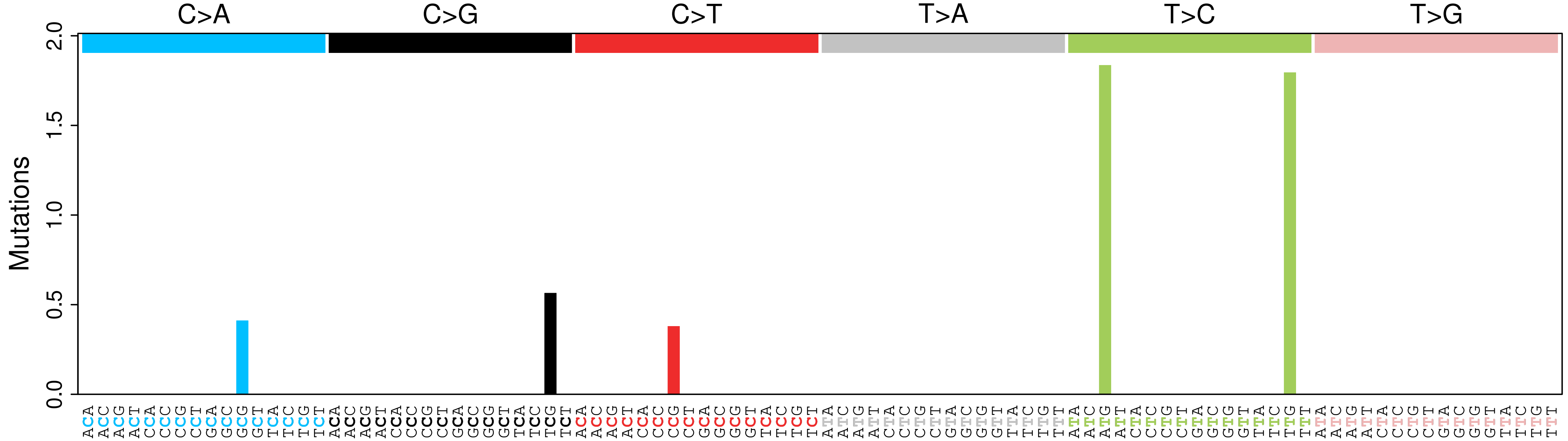
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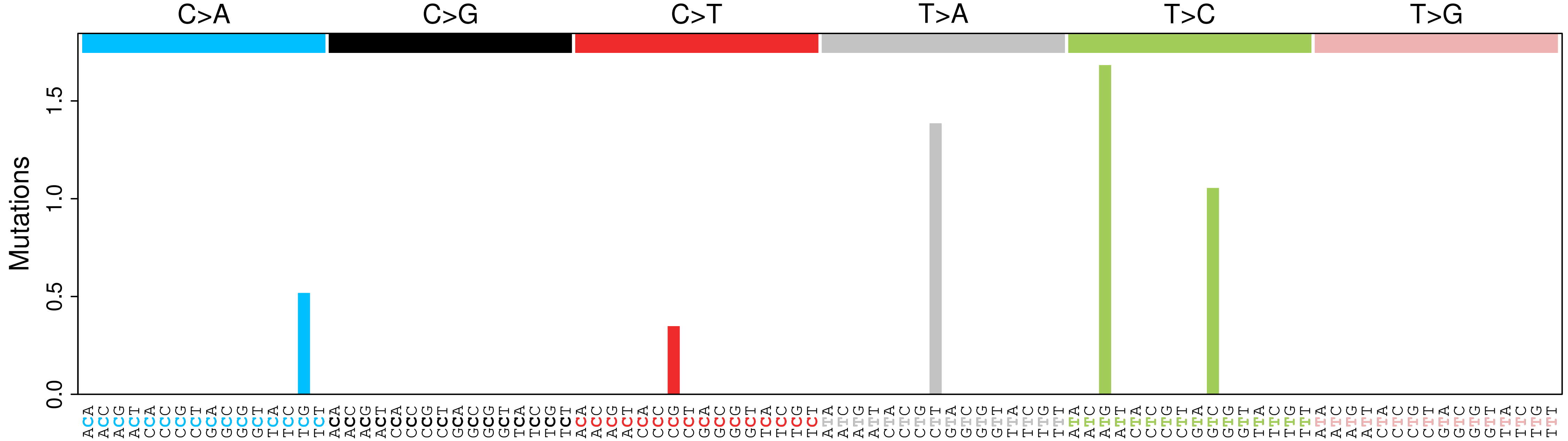
CATD0487a (3 mutations)



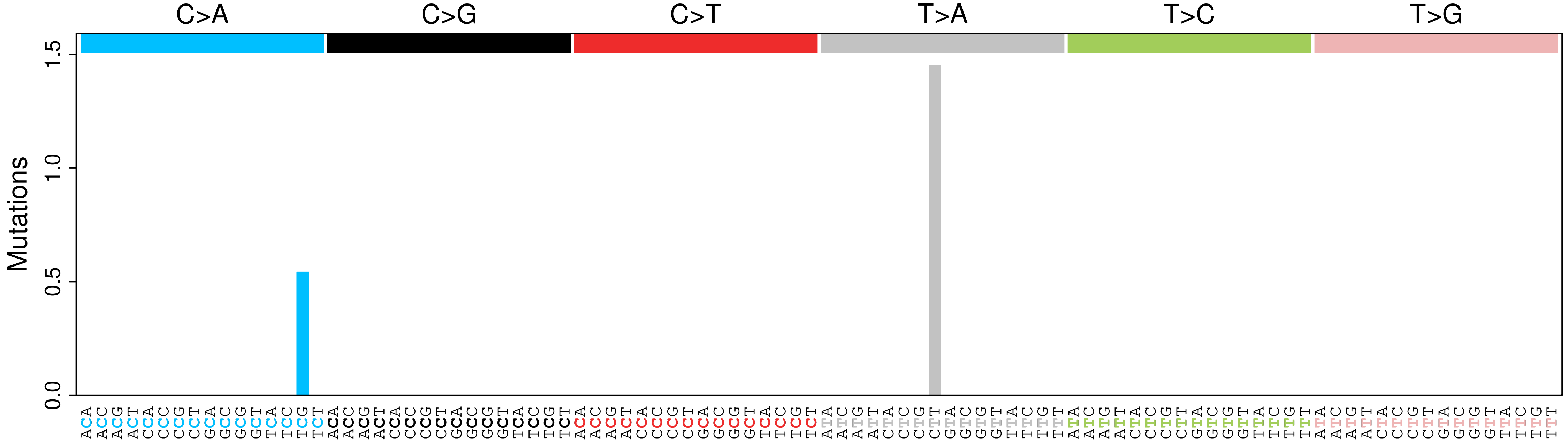
CATD0744a (5 mutations)



CATD0471a (5 mutations)



CATD0741a (2 mutations)



CATD0739a (3 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

T > G

Mutations

1.5

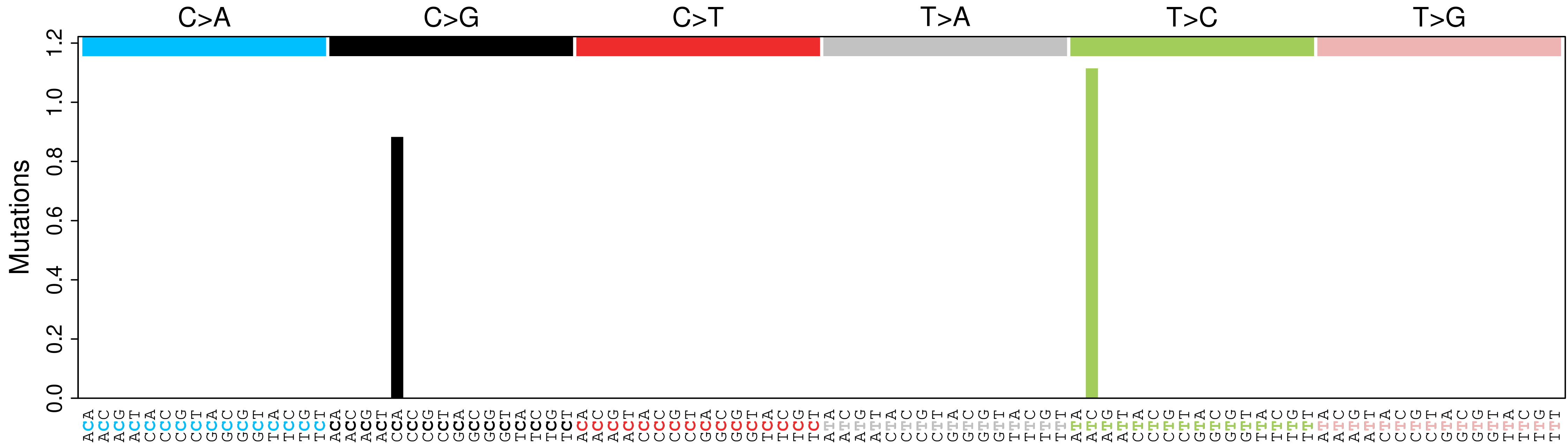
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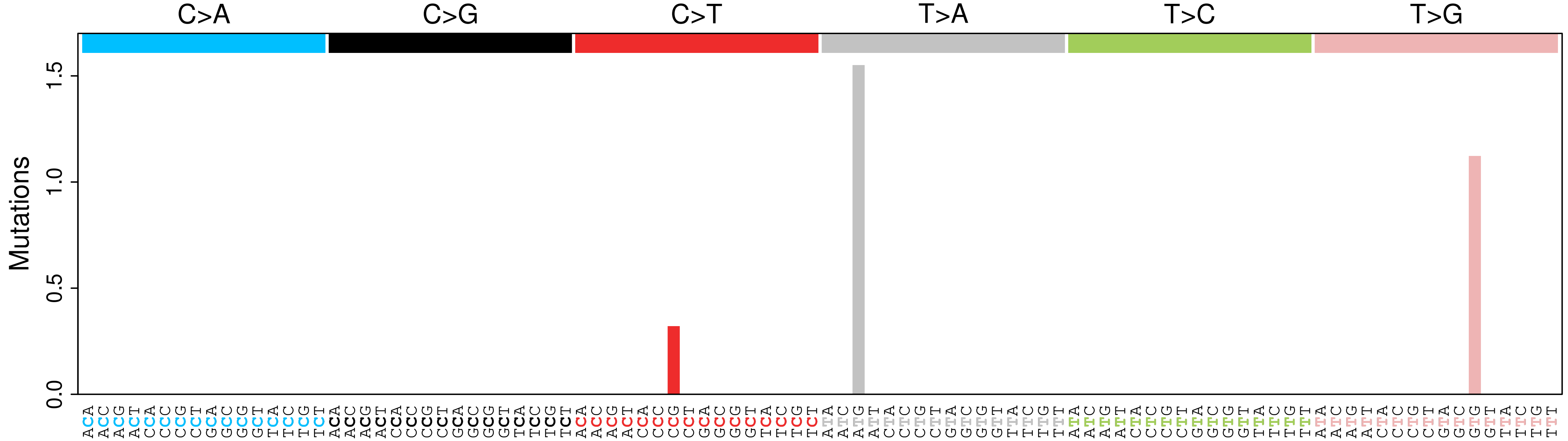
0

[illegible]

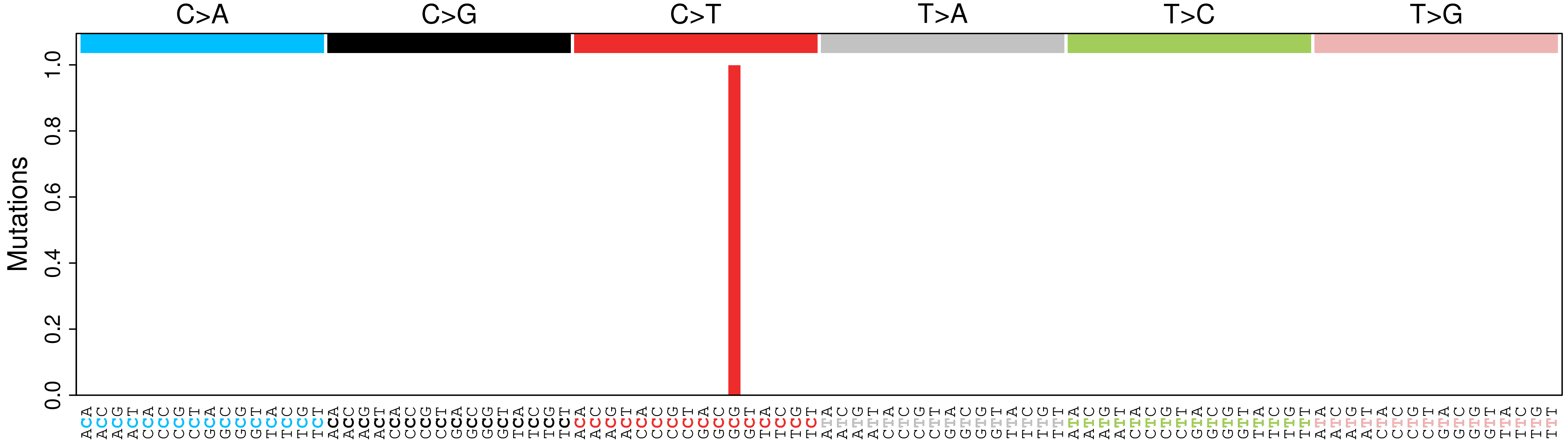
CATD0738a (2 mutations)



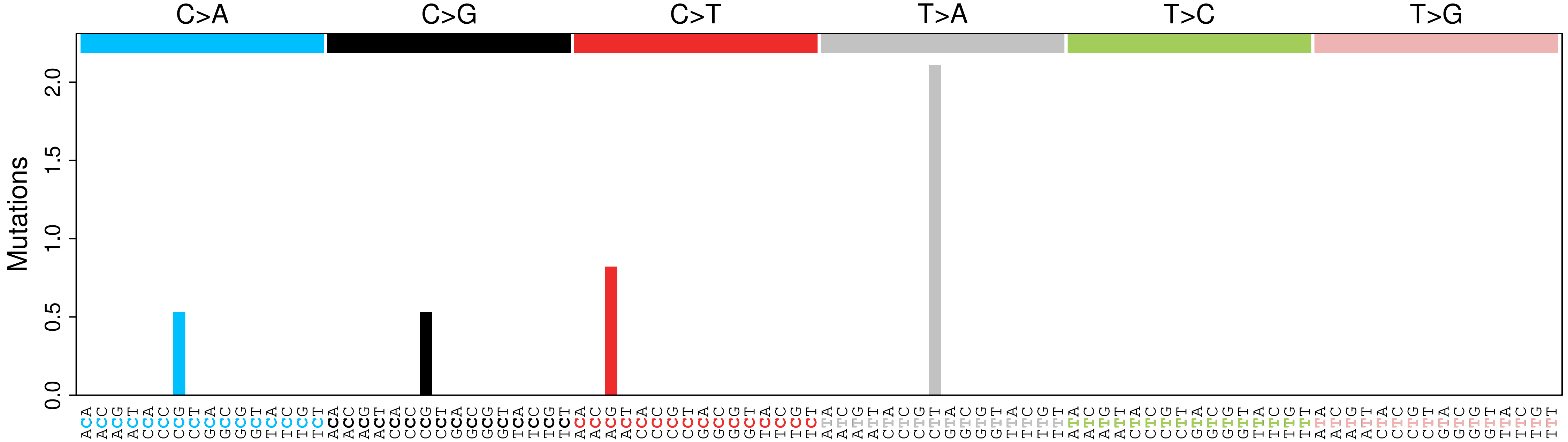
CATD0692a (3 mutations)



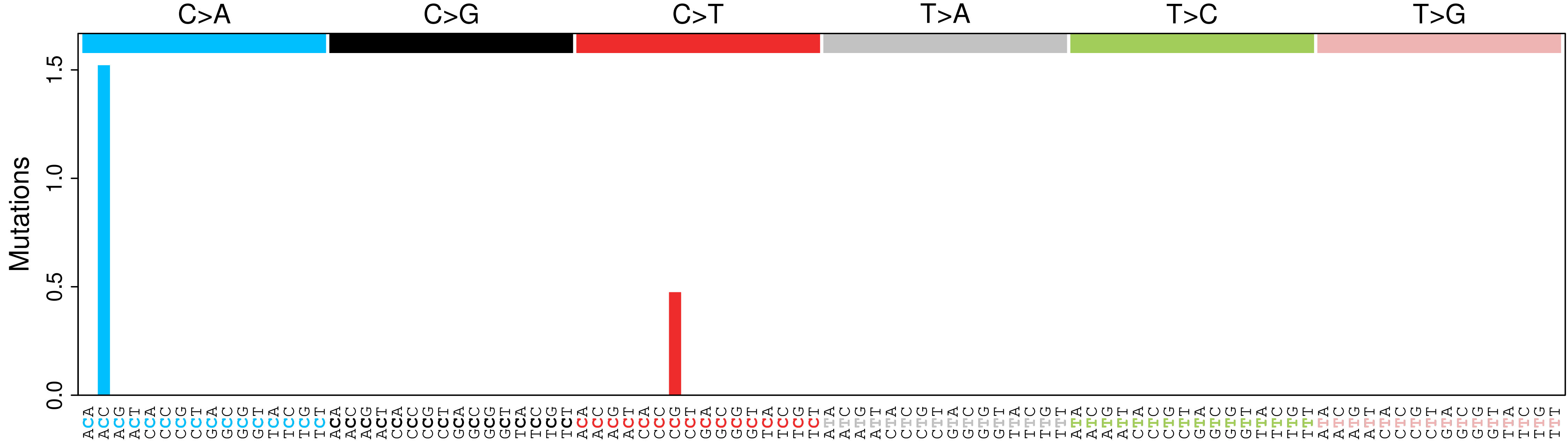
CATD0497a (1 mutations)



CATD0498a (4 mutations)



CATD0699a (2 mutations)



CATD0695a (1 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

T > G

Mutations

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0.

[illegible]