

CATD231a (1 mutations)

$C > A$

C > G

$C > T$

T > A

$$T > C$$

$T > G$

Mutations

1.0

3.0

 \circ

0

U

[illegible]

CATD0716a (2 mutations)

C > A

C > G

$C > T$

T > A

$$T > C$$

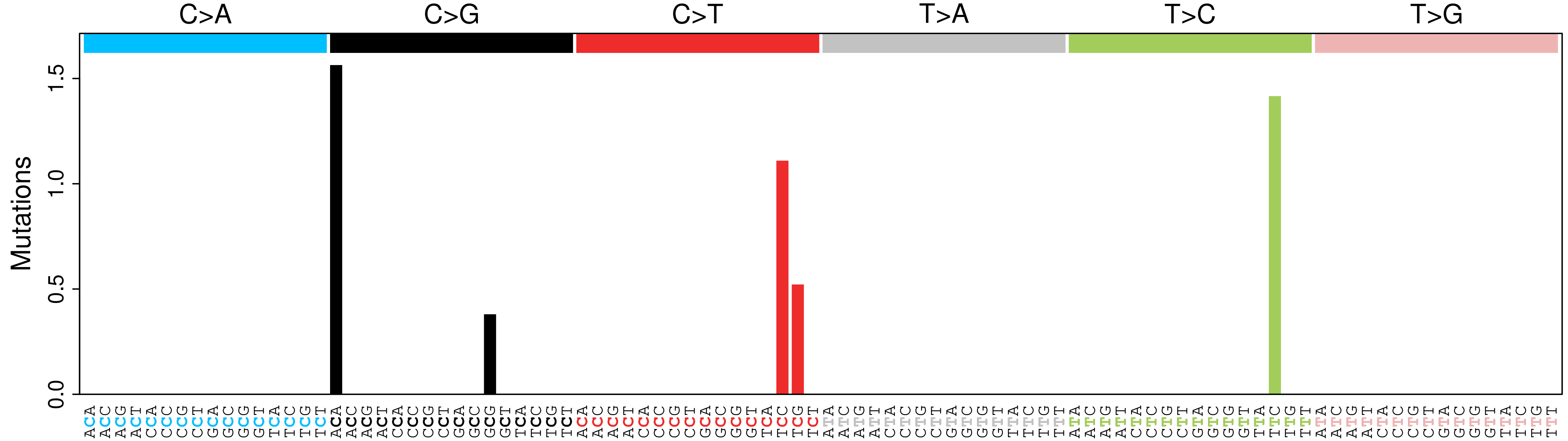
T > G

Mutations

0.0 0.2 0.4 0.6 0.8 1.0

[illegible]

CATD217a (5 mutations)



CATD227a (2 mutations)

$C > A$

$C > G$

$C > T$

$T > A$

 $T > C$

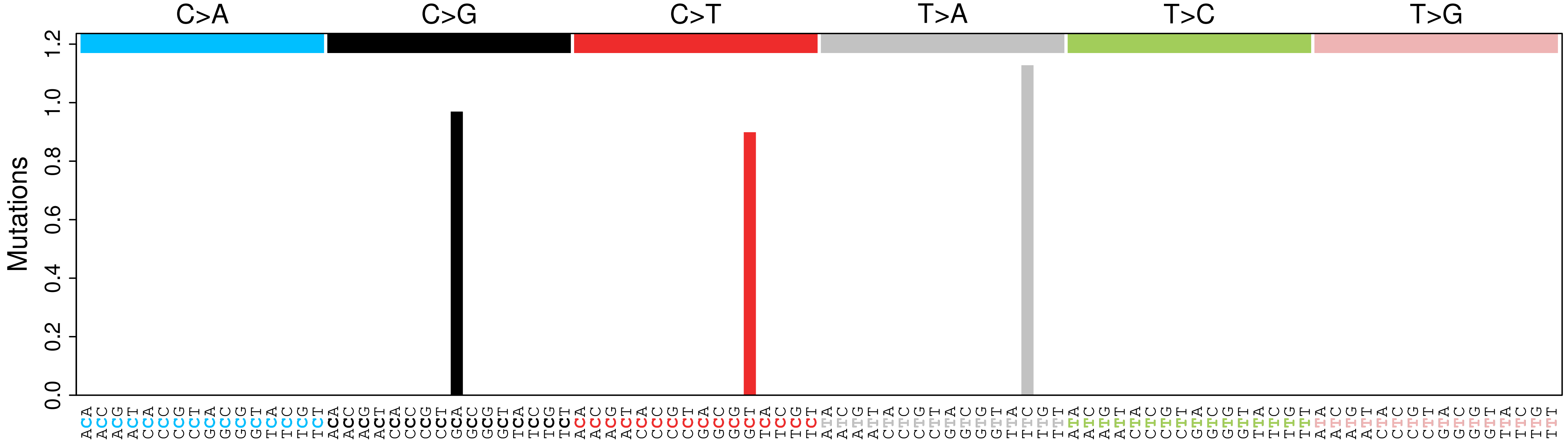
T>G

Mutations

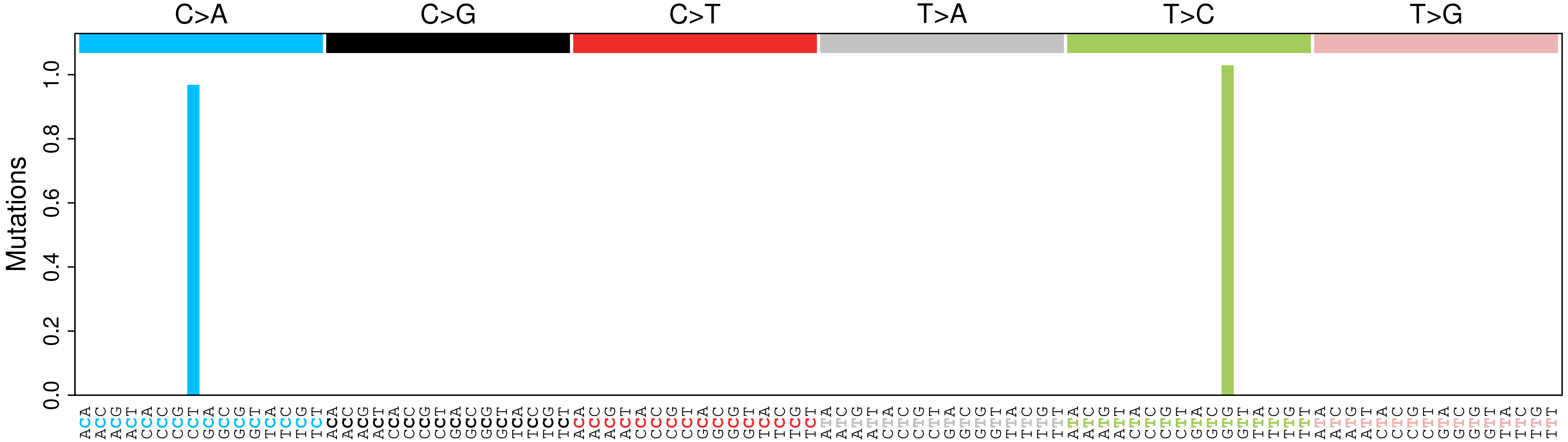
0.0 0.2 0.4 0.6 0.8 1.0 1.2

[illegible]

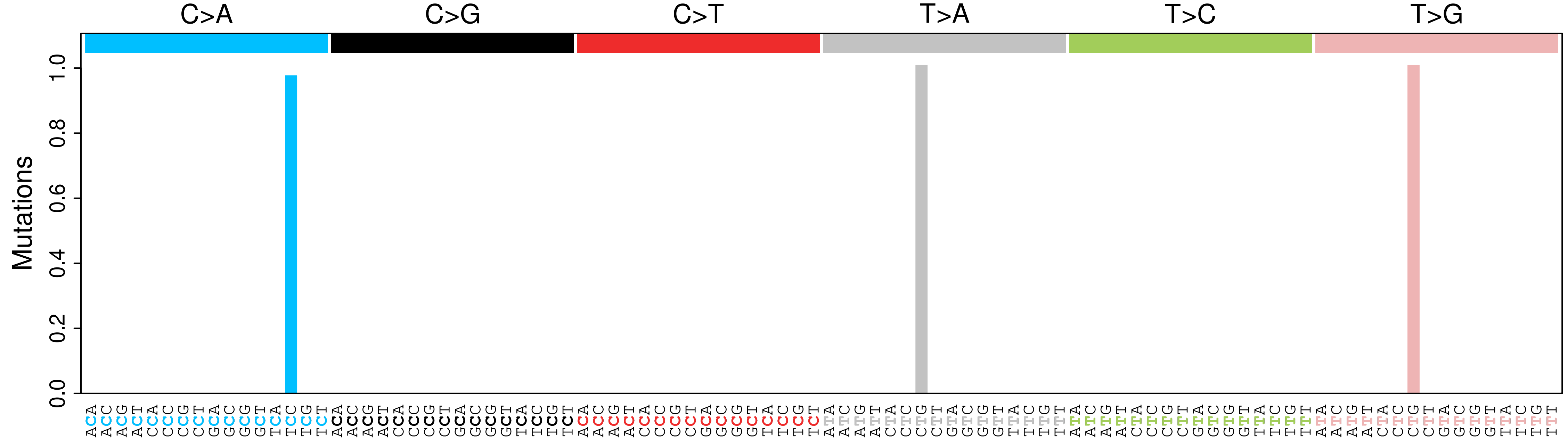
CATD0710a (3 mutations)



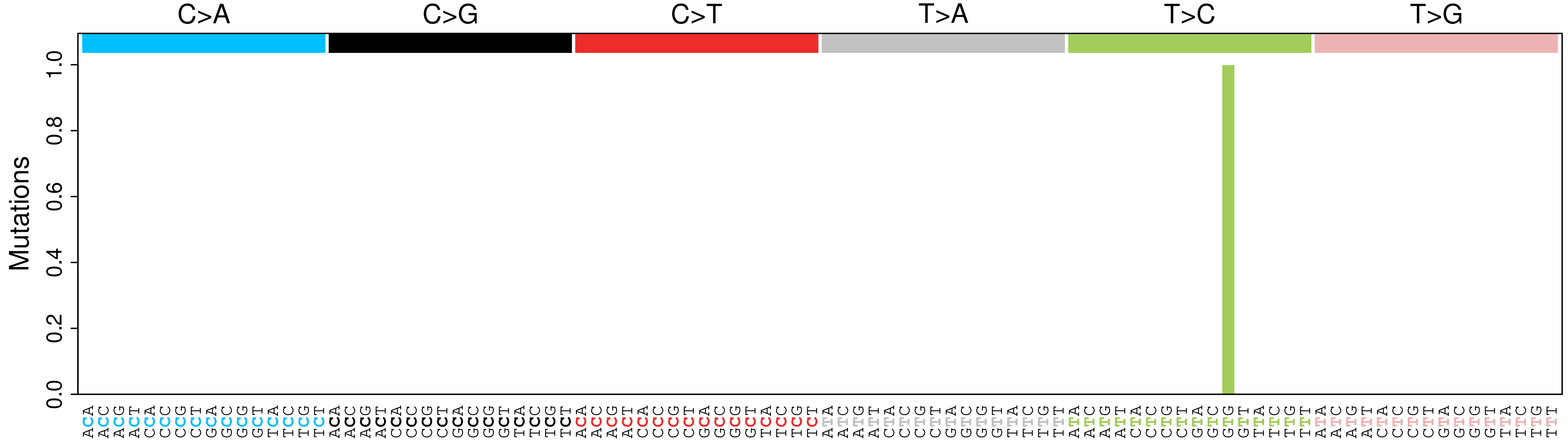
CATD0717a (2 mutations)



CATD242a (3 mutations)



CATD0709a (1 mutations)



CATD239a (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

[illegible]

CATD216a (1 mutations)

$C > A$

$C > G$

$C > T$

T>A

 $T > C$

T>G

Mutations

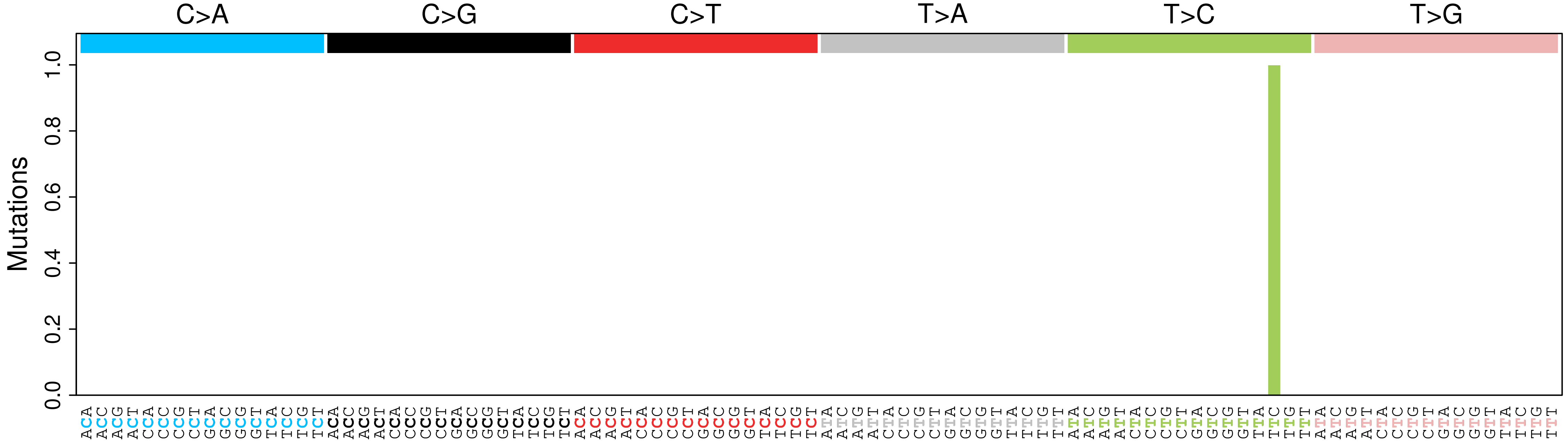
1.0

0

2

0.

CATD232a (1 mutations)



CATD229a (3 mutations)

$C > A$

C > G

$$C > T$$

T > A

$$T > C$$

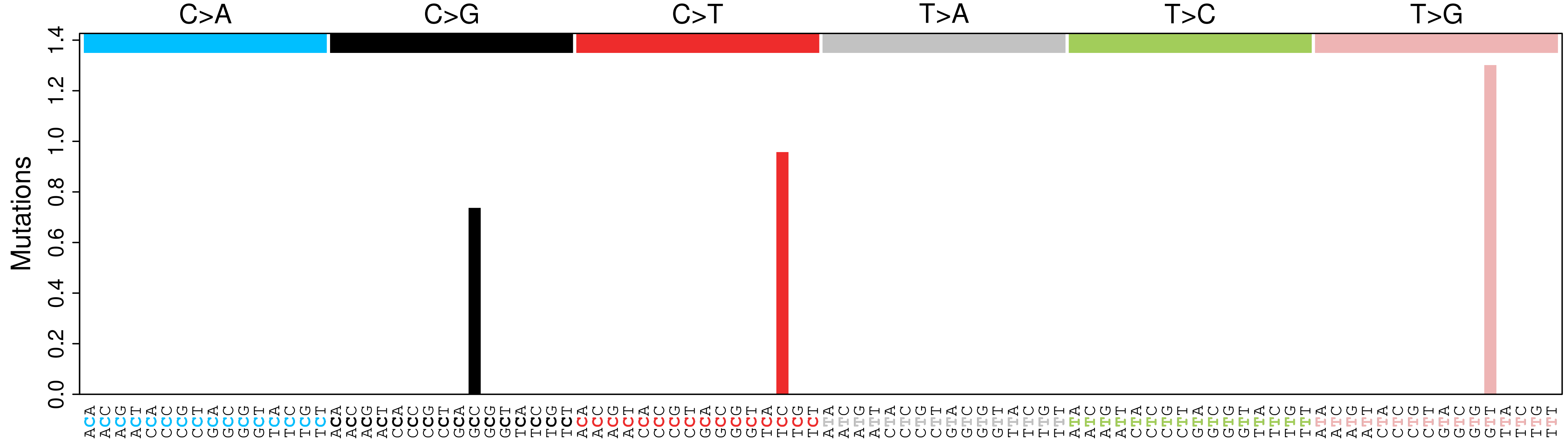
T>G

Mutations

0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

[illegible]

CATD0712a (3 mutations)



CATD226a (1 mutations)

$C > A$

$C > G$

$C > T$

$T > A$

 $T > C$

T>G

Mutations

1.0

 \dot{O}

1

2

0.

[illegible]

CATD0715a (1 mutations)

$C > A$

$C > G$

$C > T$

T > A

$T > C$

T > G

Mutations

1.0

 \dot{O}

1

2

0.

[illegible]

CATD224a (2 mutations)

$C > A$

C > G

$$C > T$$

T>A

$T > C$

T>G

Mutations

1.5

1.0

0.5

0.

[illegible]

CATD0561e (1 mutations)

C > A

C > G

$C > T$

T > A

$$T > C$$

$T > G$

Mutations

1.0

 \dot{O}

1

2

0.

[illegible]