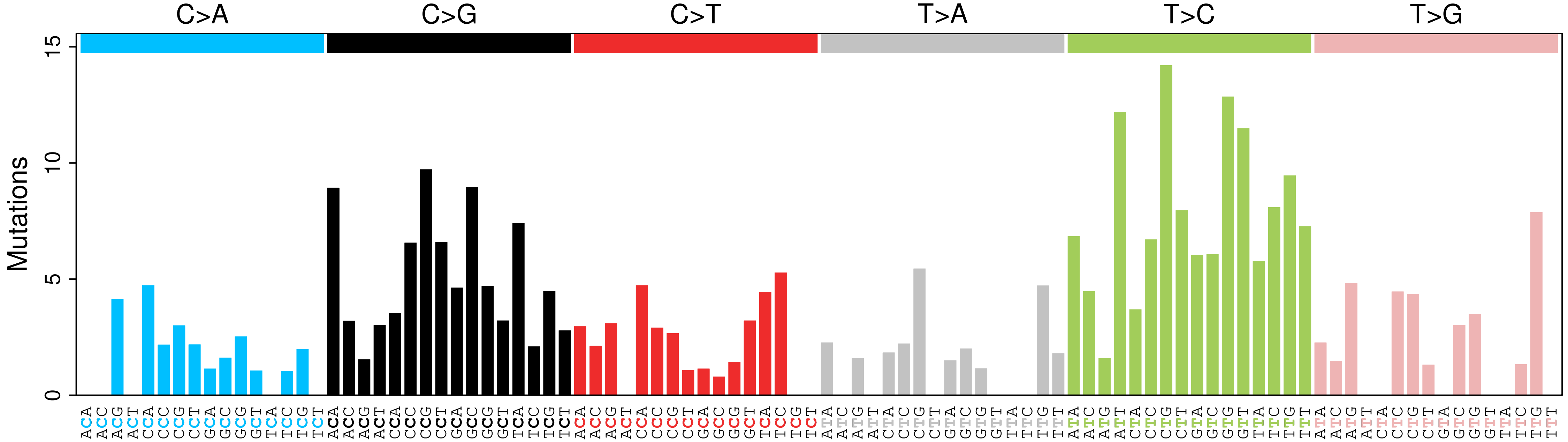
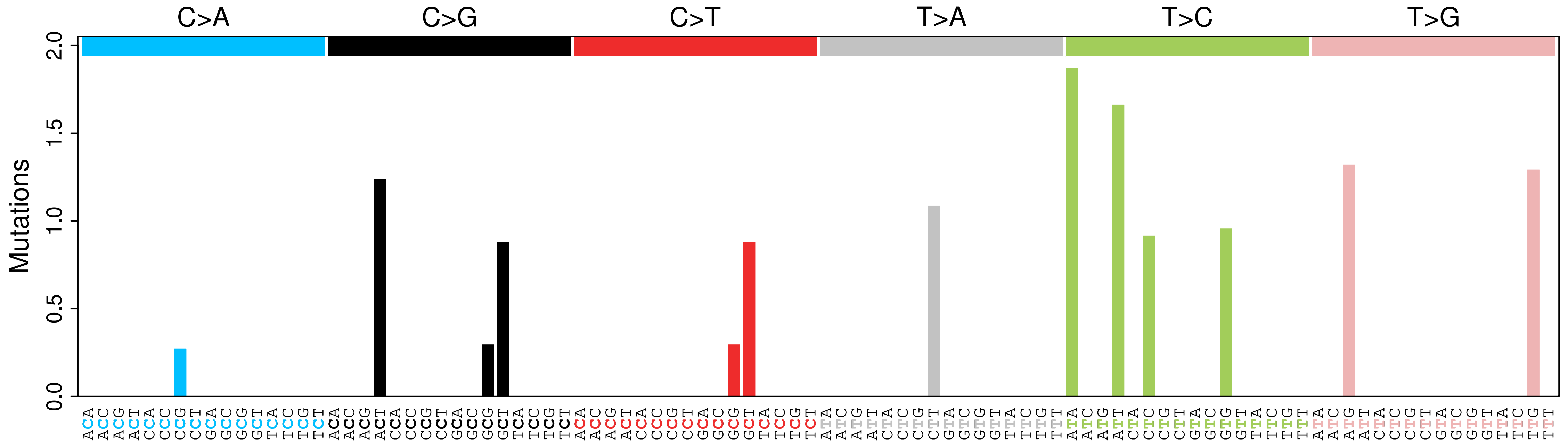


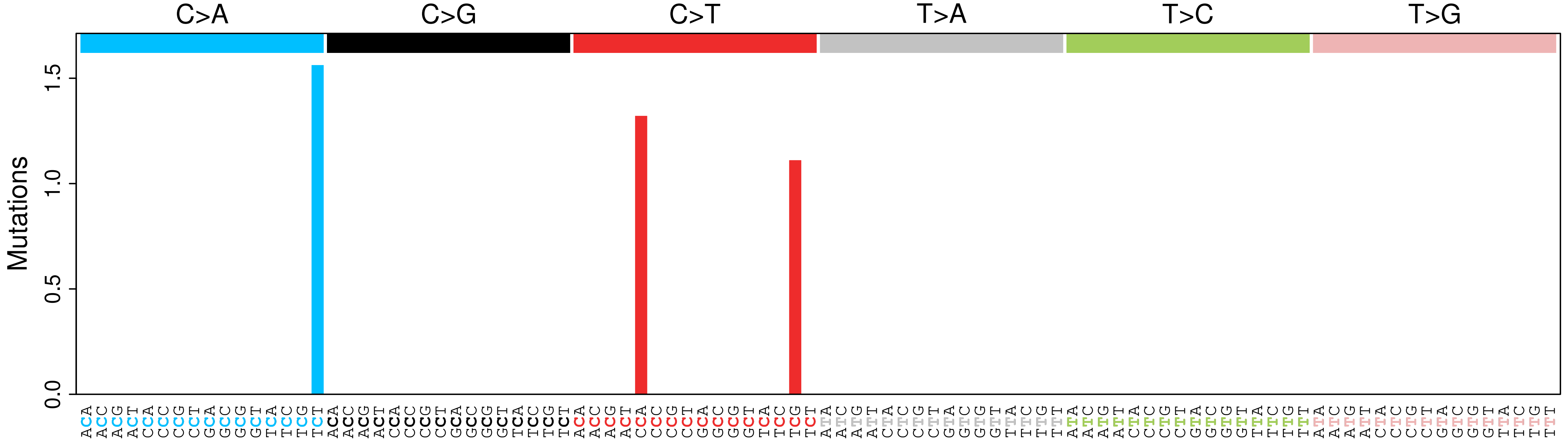
CATD292a (328 mutations)



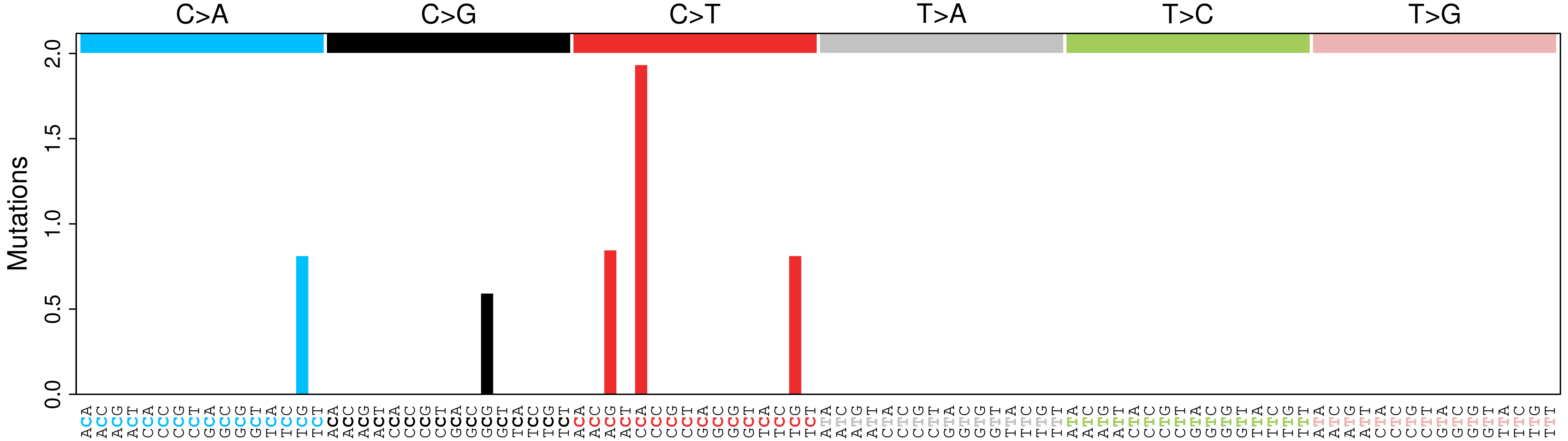
CATD283a (13 mutations)



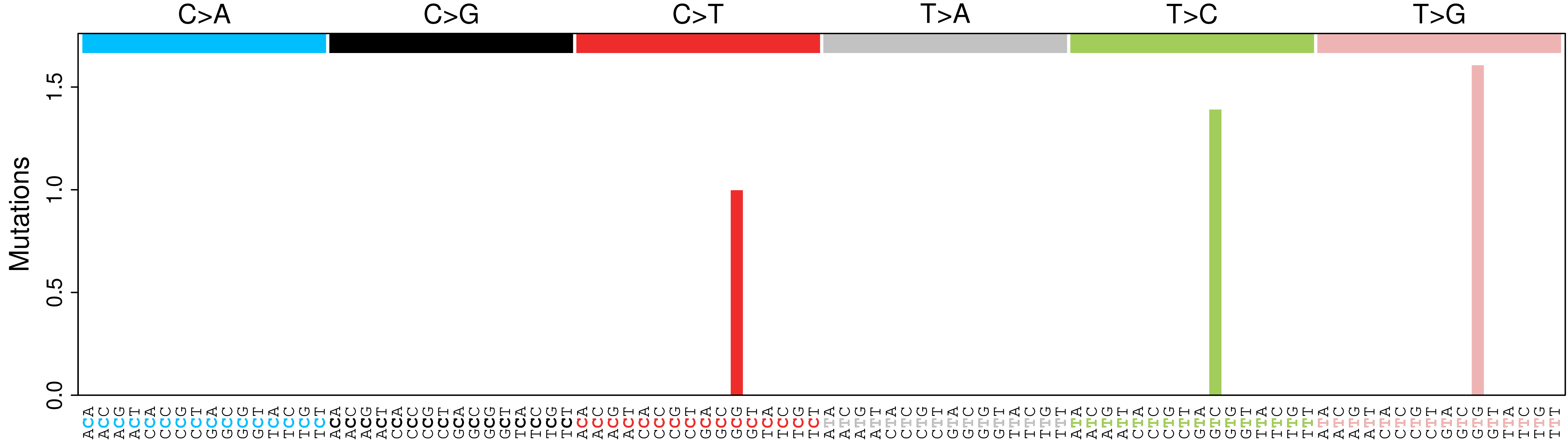
CATD0645a (4 mutations)



CATD0670a (5 mutations)



CATD0650a (4 mutations)



CATD289a (2 mutations)

$C > A$

C > G

$$C > T$$

T>A

$T > C$

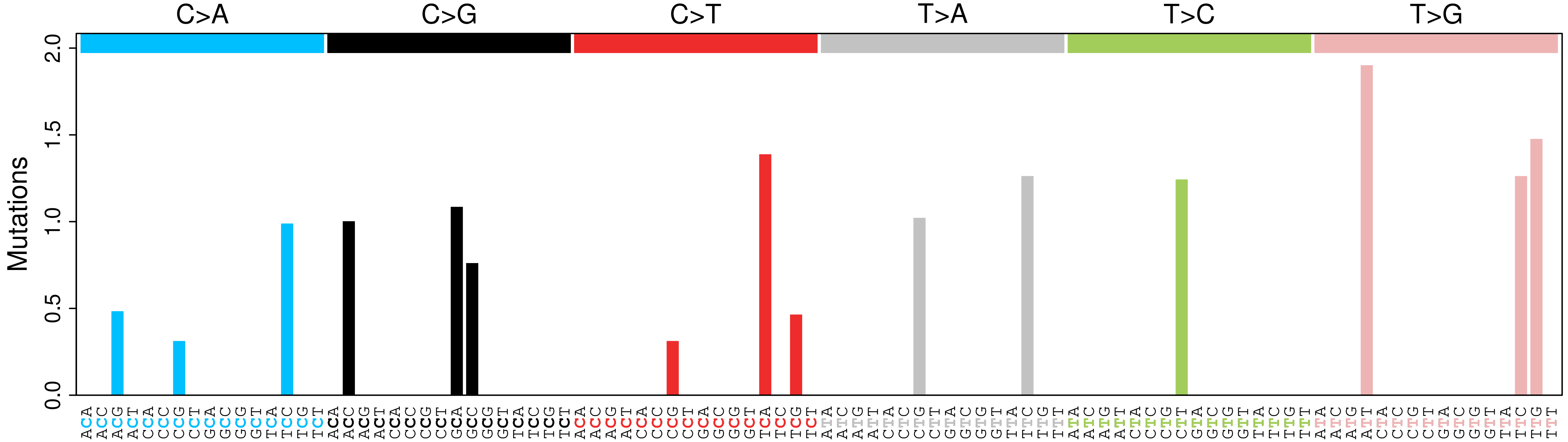
$T > G$

Mutations

0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

[illegible]

CATD0144a (15 mutations)



CATD293a (4 mutations)

$C > A$

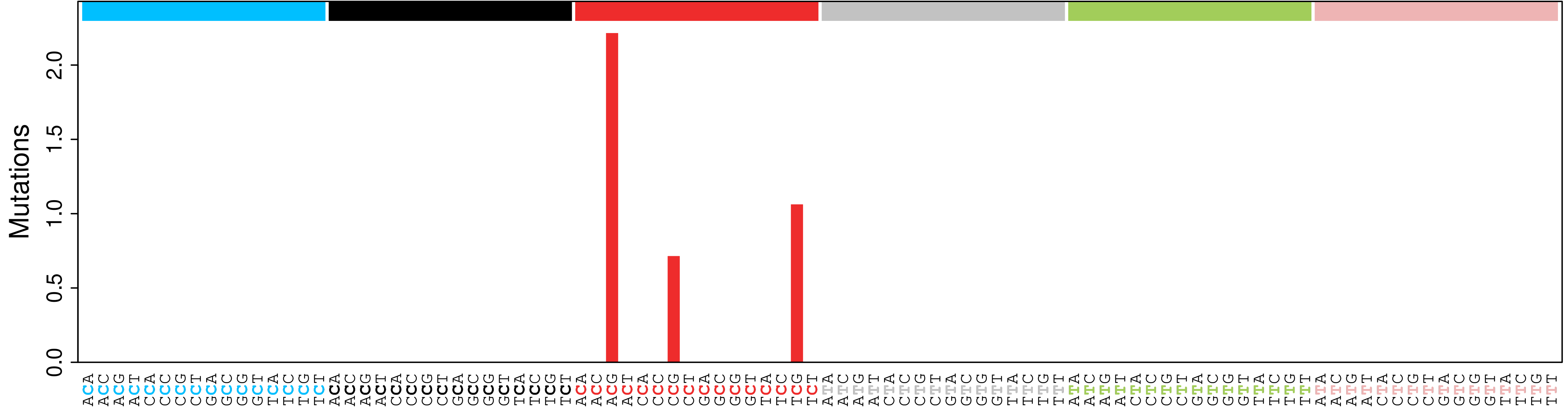
C > G

$$C > T$$

T > A

$$T > C$$

$T > G$



CATD0668a (6 mutations)

C>A

C>G

C>T

T>A

T>C

T>G

Mutations

1.5
1.0
0.5
0.0

ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GGT GTC TCA TCC TCG TCT ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GGT GTC TCA TCC TCG TCT ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TAC TTG TTT ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TAC TTG TTT ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TAC TTG TTT

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

1.5

1.0

0.5

0.0

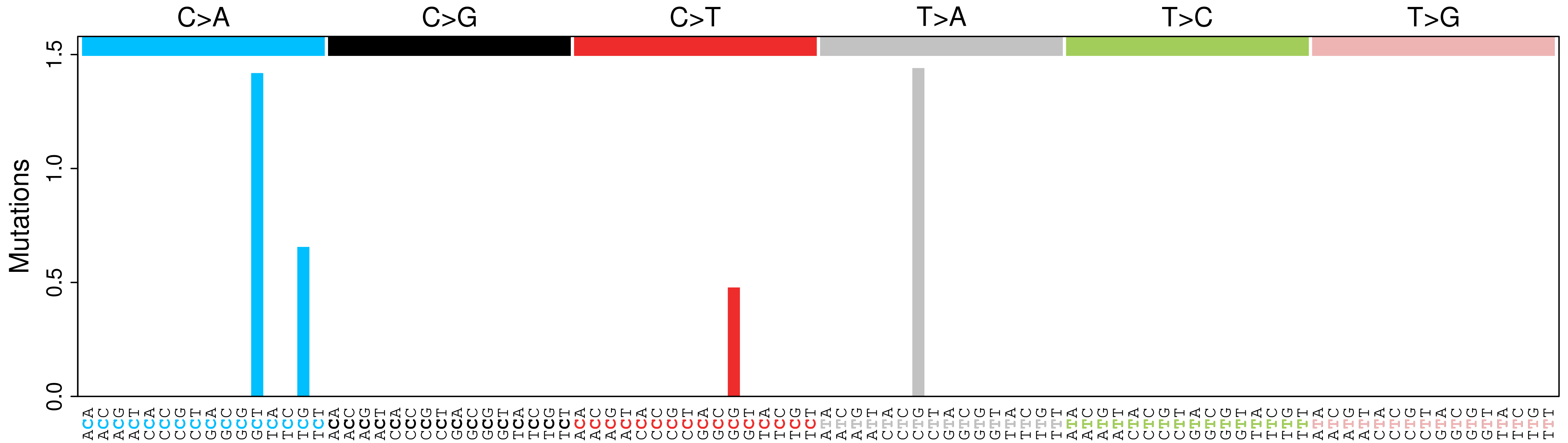
1.5

1.0

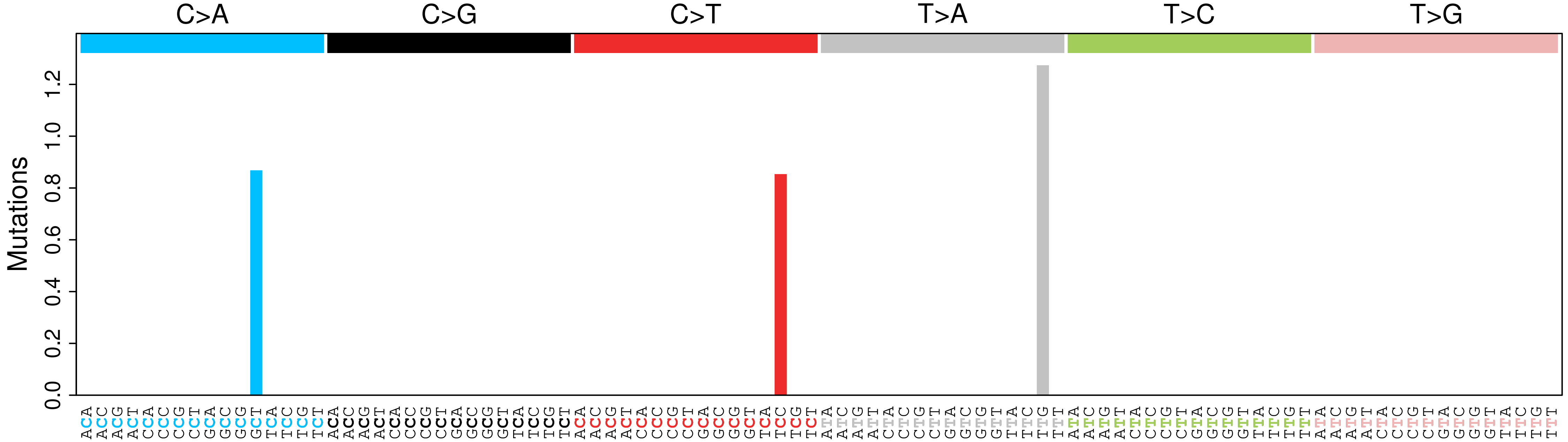
0.5

0.0

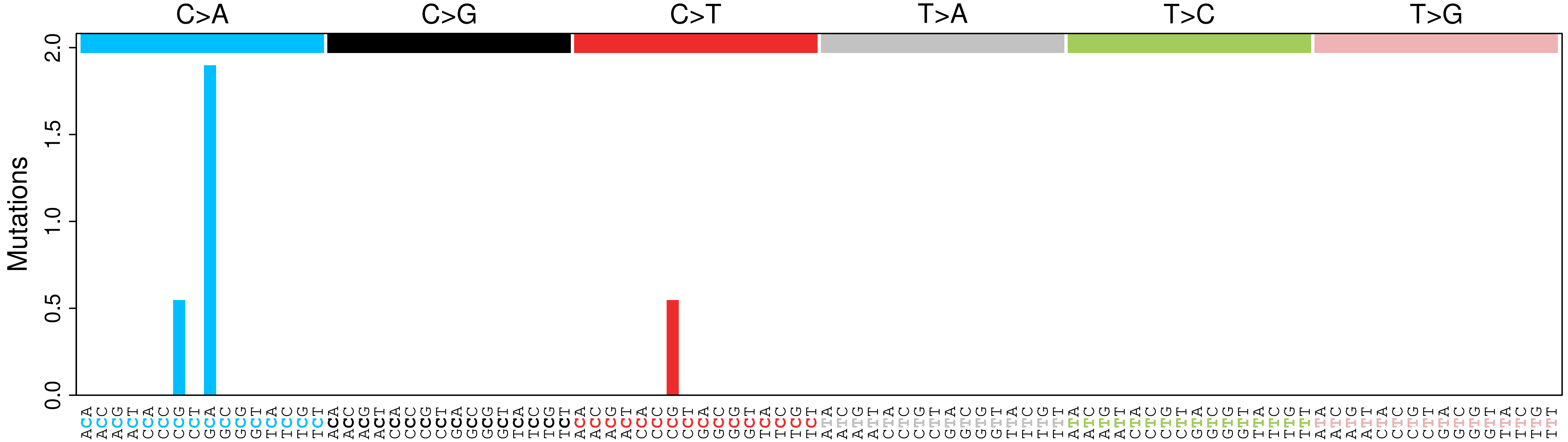
CATD0146a (4 mutations)



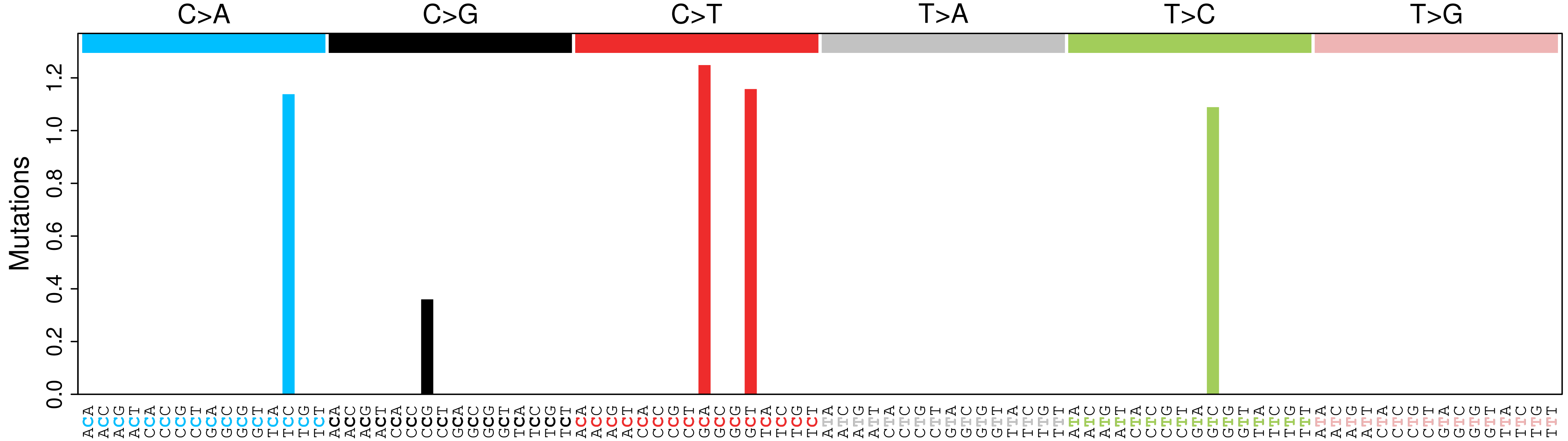
CATD0641a (3 mutations)



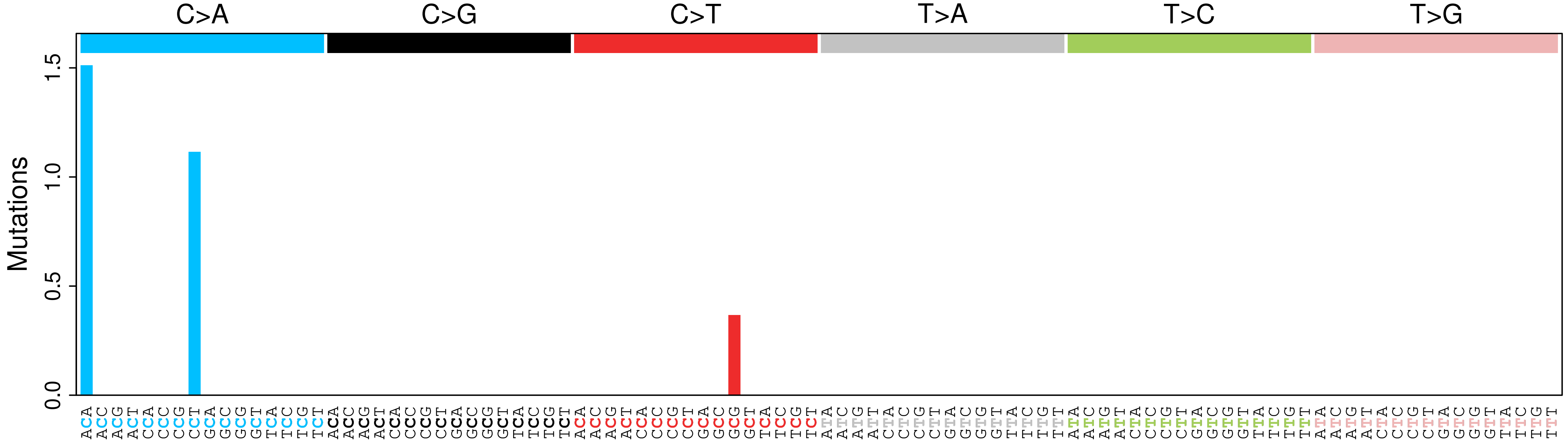
CATD0143a (3 mutations)



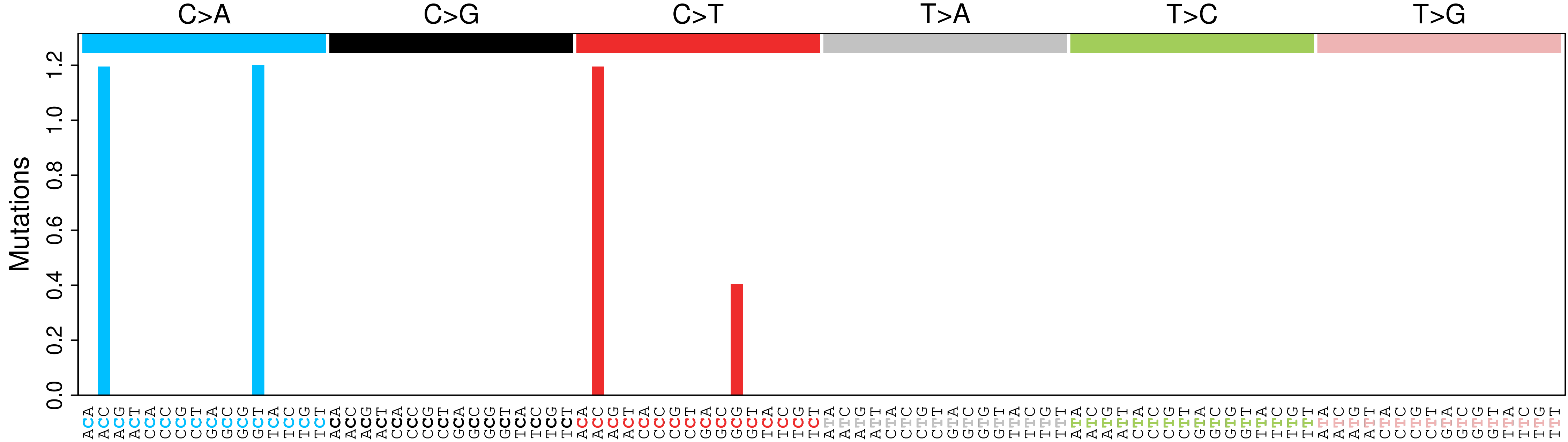
CATD0632a (5 mutations)



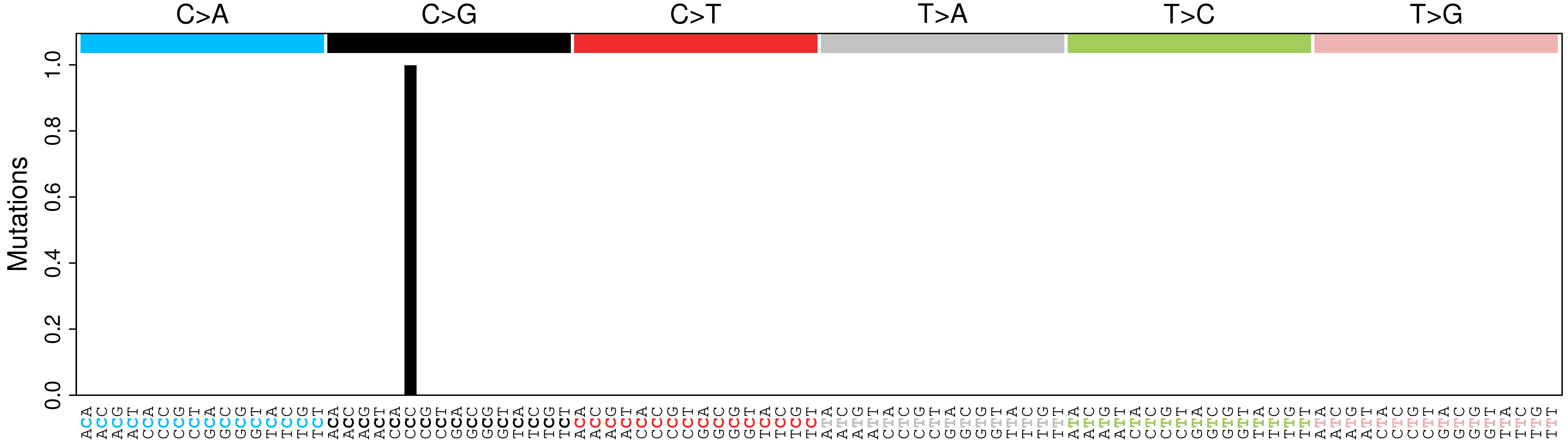
CATD0633a (3 mutations)



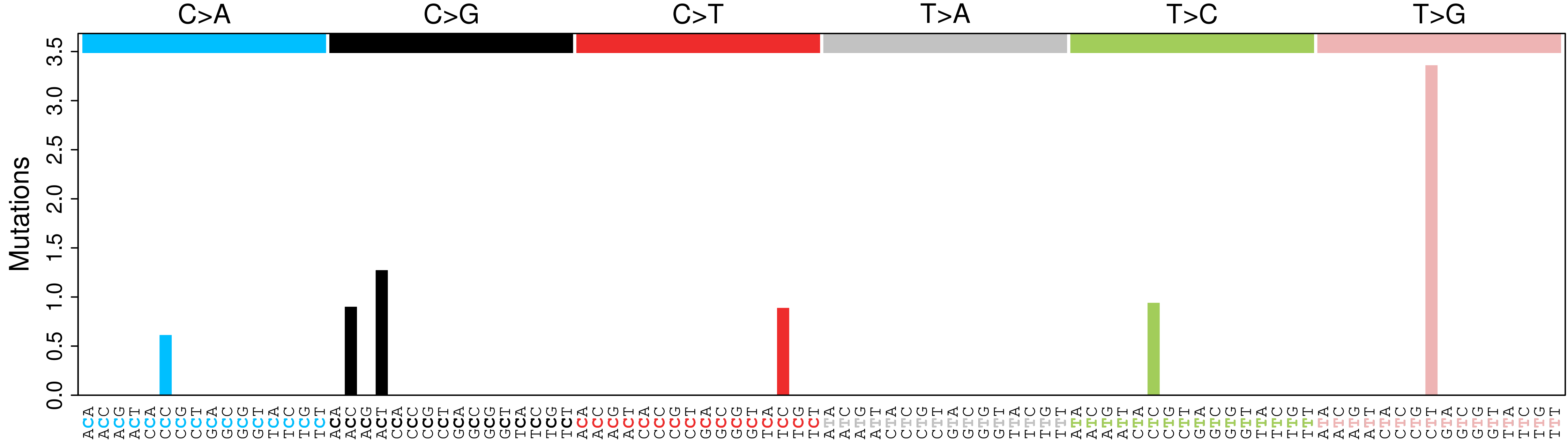
CATD0157a (4 mutations)



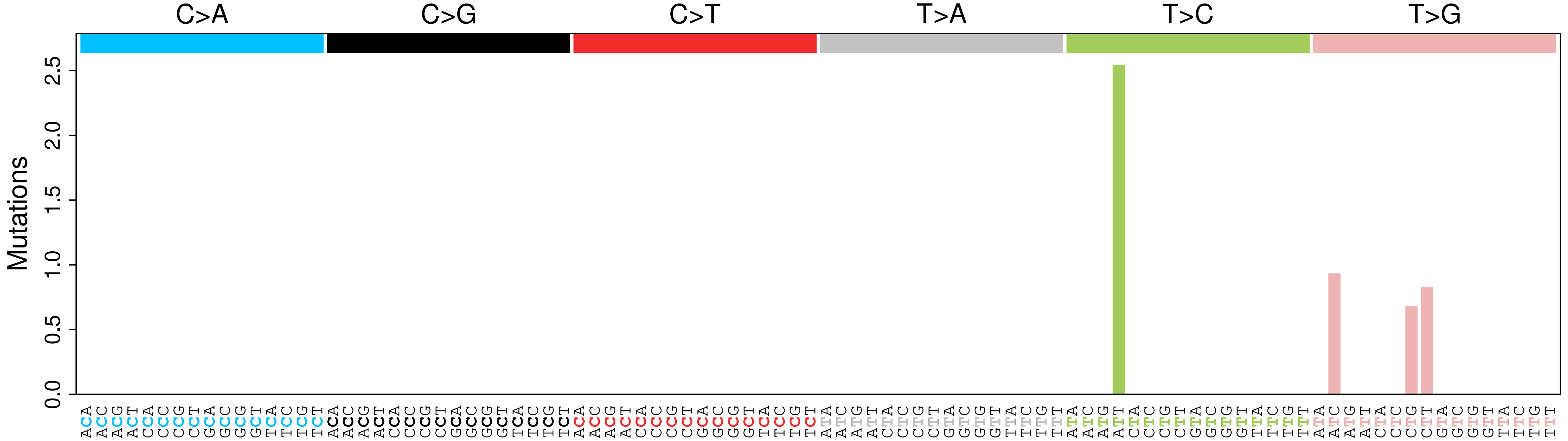
CATD0147a (1 mutations)



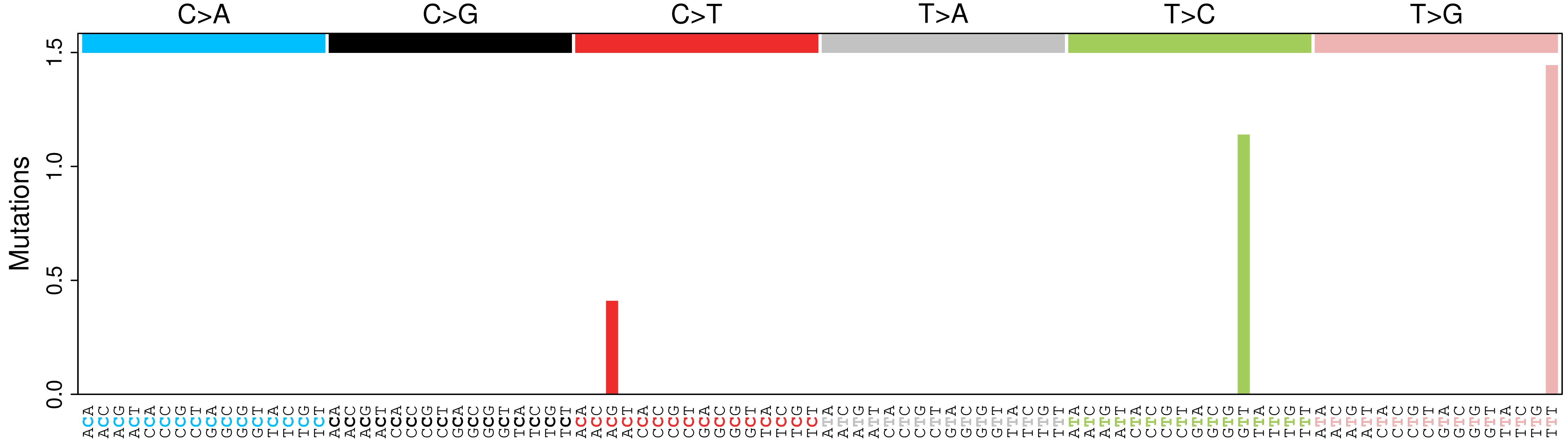
CATD0145a (8 mutations)



CATD0159a (5 mutations)



CATD0148a (3 mutations)



CATD0153a (6 mutations)

$C > A$

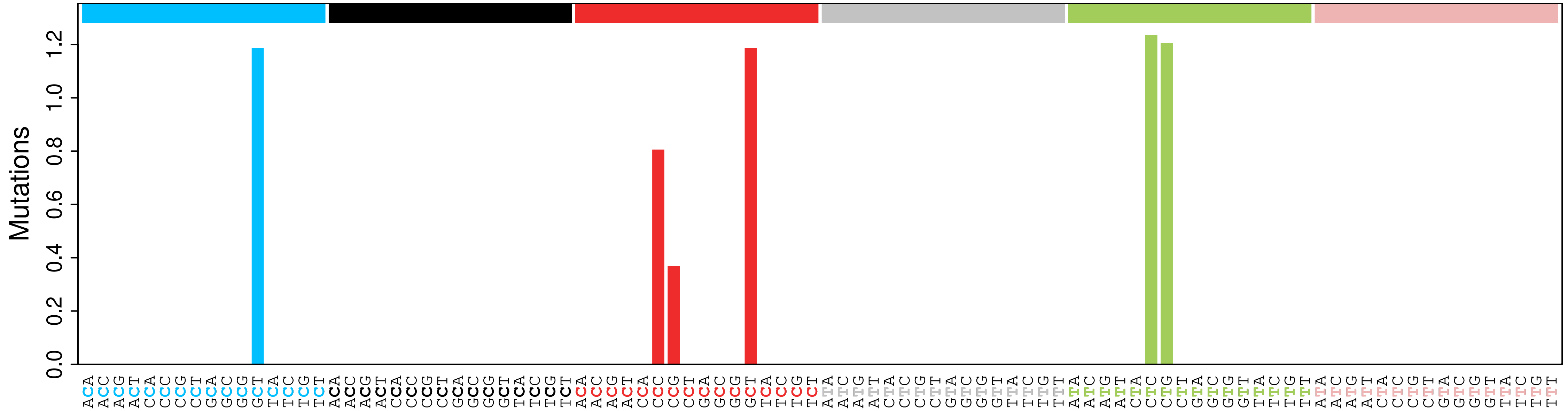
C > G

C>T

T>A

$$T > C$$

T>G



CATD0635a (3 mutations)

$C > A$

C > G

$C > T$

T>A

 $T > C$

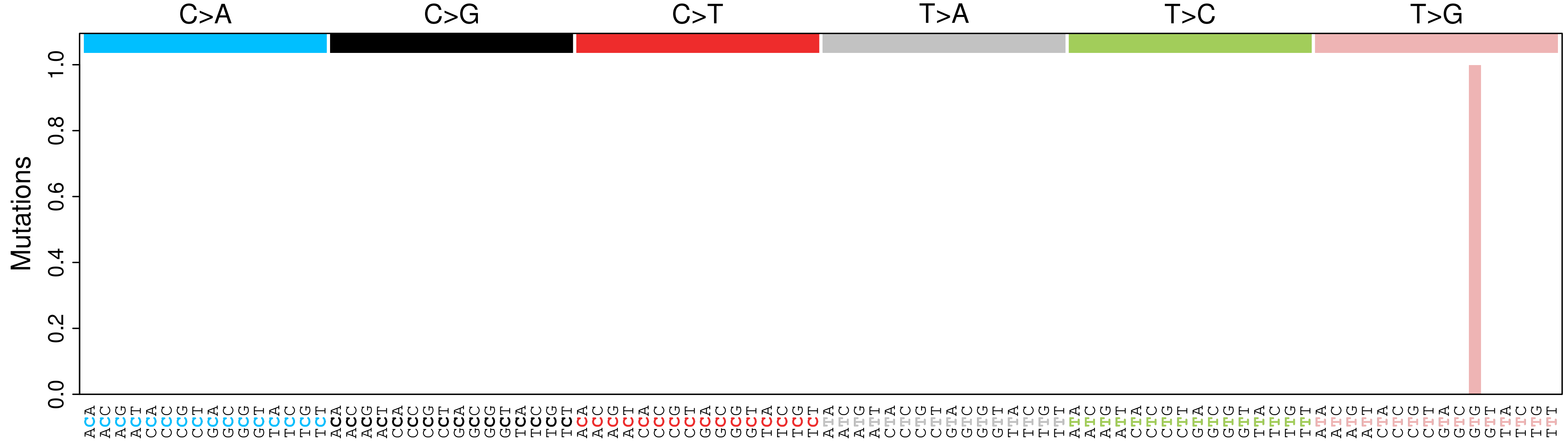
T > G

Mutations

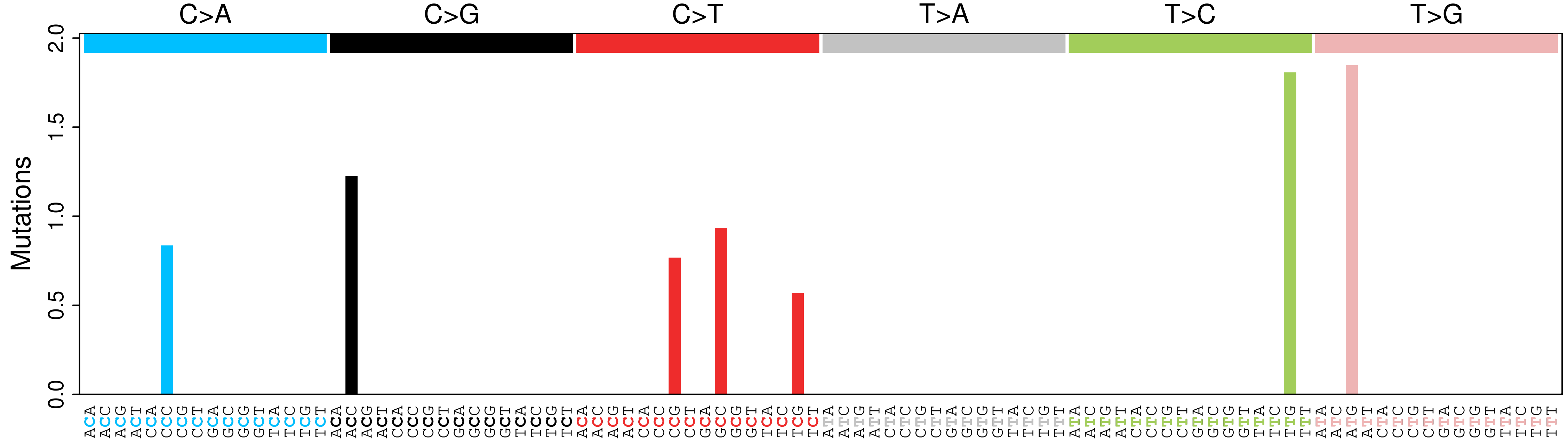
0.0 0.2 0.4 0.6 0.8 1.0 1.2

[illegible]

CATD0646a (1 mutations)



CATD295a (8 mutations)



CATD290a (1 mutations)

$C > A$

$C > G$

$C > T$

T>A

 $T > C$

T>G

Mutations

1.0

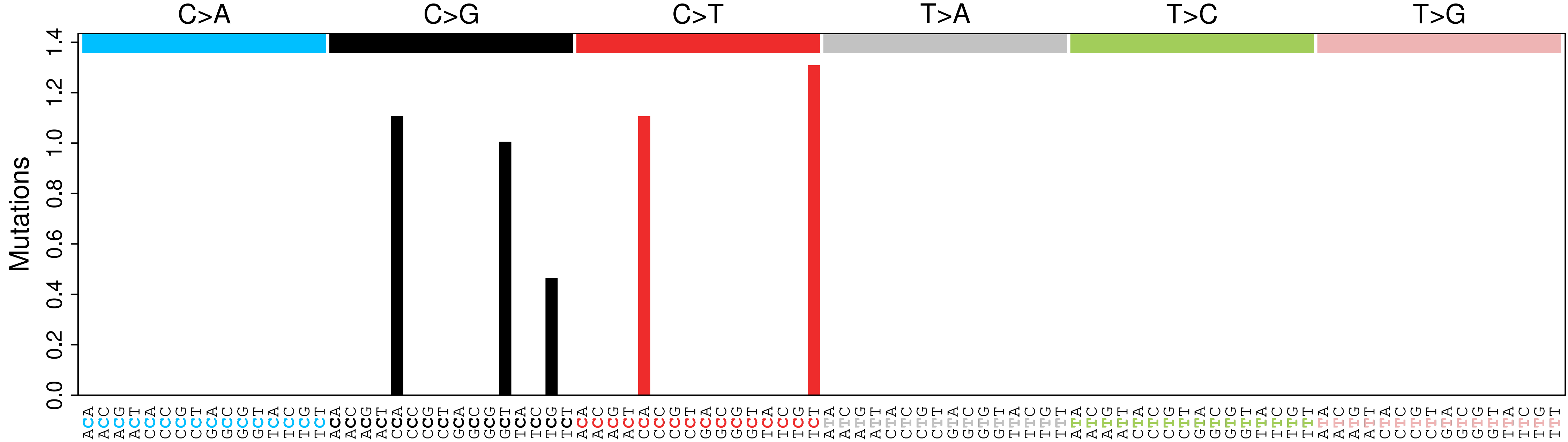
0

2

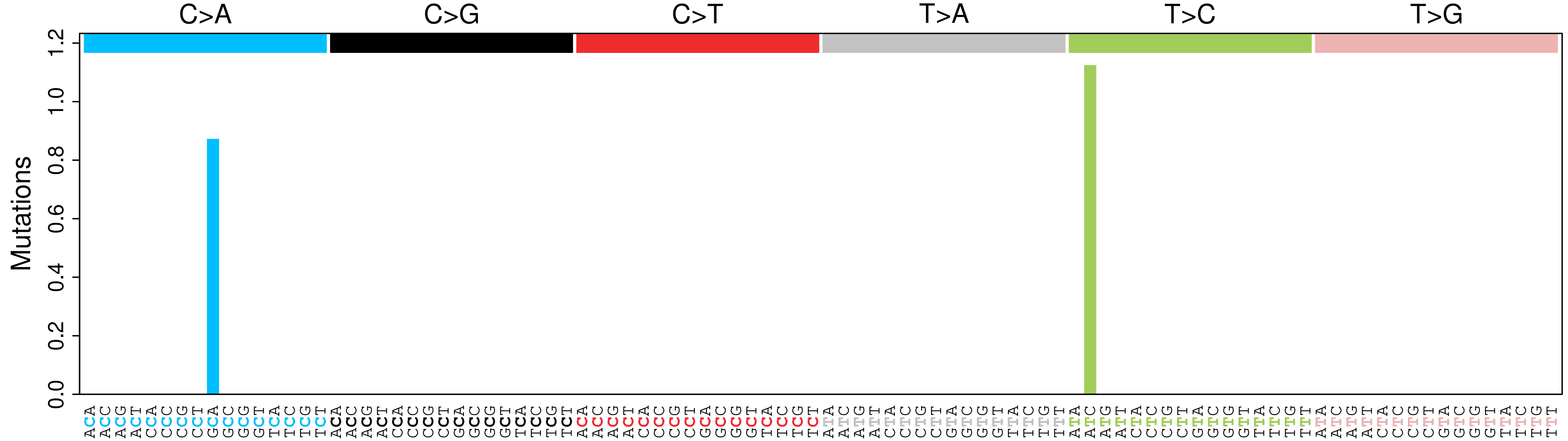
0.

[illegible]

CATD0142a (5 mutations)



CATD291a (2 mutations)



CATD284a (3 mutations)

C>A

C>G

C>T

T>A

T>C

T>G

Mutations

1.2
1.0
0.8
0.6
0.4
0.2
0.0

ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GCG GGT TCA TCC TCG TCT ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GCG GGT TCA TCC TCG TCT ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT



CATD0150a (2 mutations)

C>A

C>G

C>T

T>A

T>C

T>G

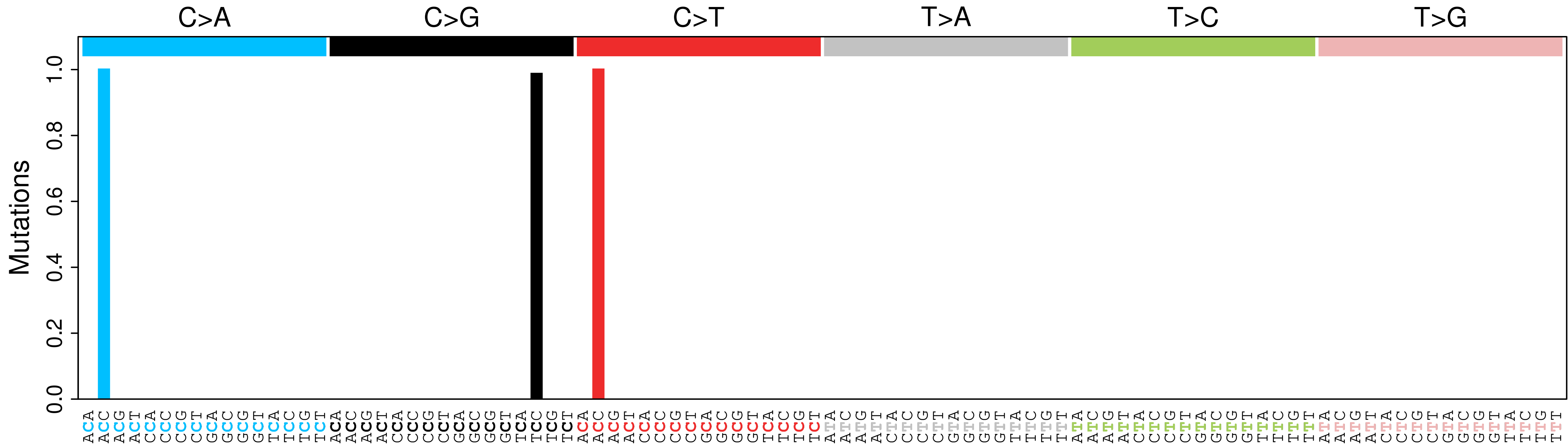
Mutations

1.2
1.0
0.8
0.6
0.4
0.2
0.0

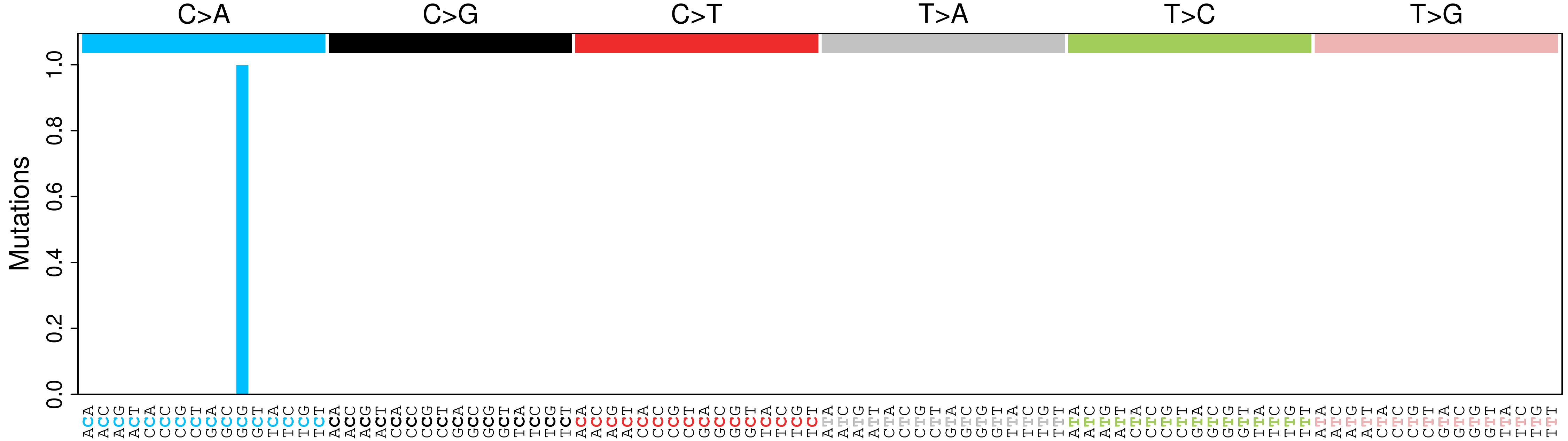
A C A C A C C C G G A G C G T T C A T C T T A C C A C G A C T C C C C G C A G C G G C T T C A T C C T C T A C C A C C A C G T A T C A T G A T T C T C T G T A T T C T T G T T A A T C A T G A T T A C C T C C T G T A G T G T T A T C T T G T T A A T C A T G A T T A C C T G C T A G T C G T T A T C T T T



CATD0156a (3 mutations)



CATD0154a (1 mutations)



CATD0651a (1 mutations)

C > A

C > G

$C > T$

T > A

$$T > C$$

T > G

Mutations

1.0

o

U

21

0.

[illegible]

CATD0152a (1 mutations)

$C > A$

C > G

$C > T$

T > A

$T > C$

T > G

Mutations

1.0

3.0

0

C

[illegible]

CATD0644a (1 mutations)

C > A

C > G

$C > T$

T > A

$$T > C$$

T>G

Mutations

1.0

 \dot{O}

1

2

0.

[illegible]