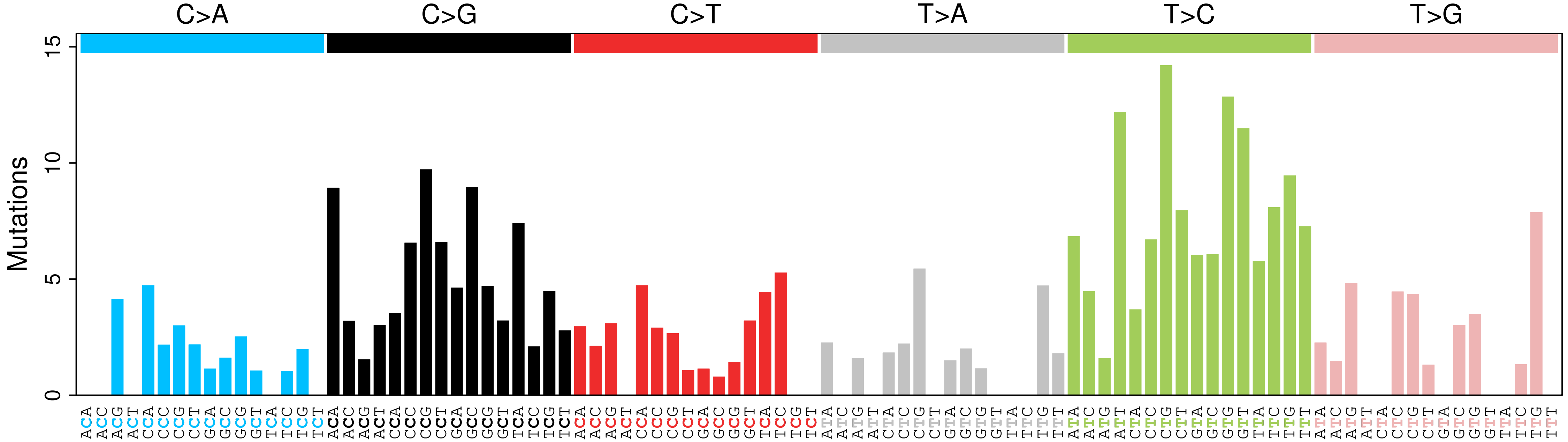


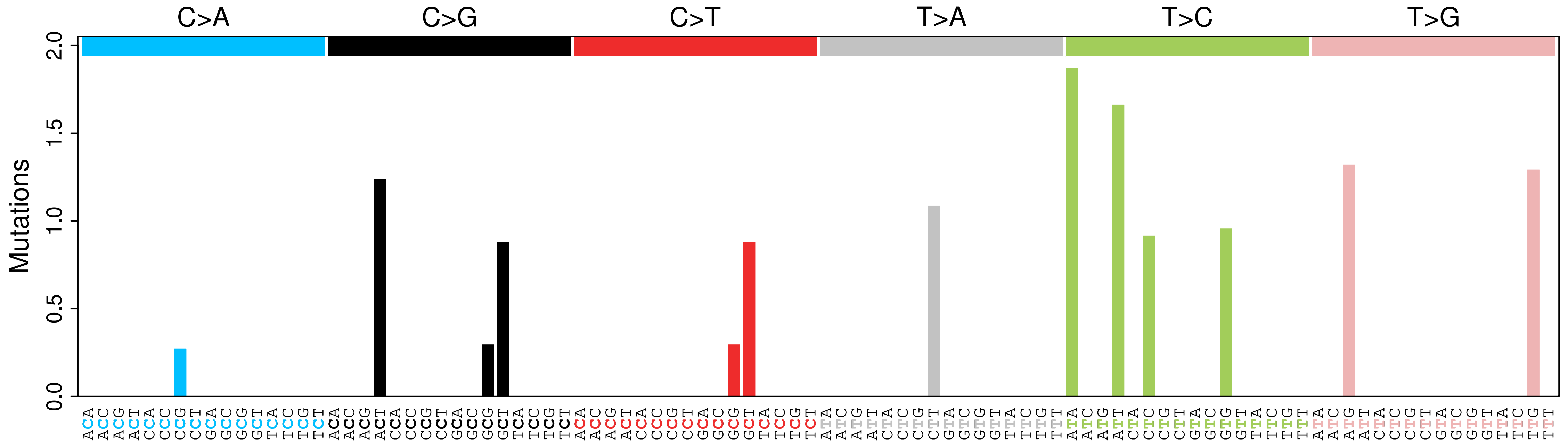
# CATD292a (328 mutations)



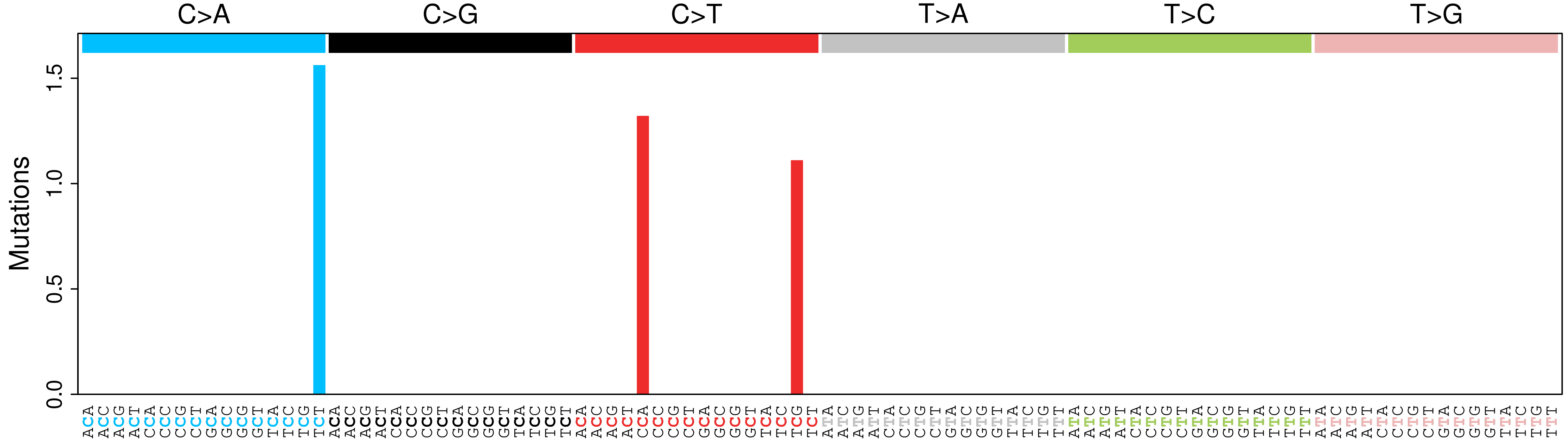




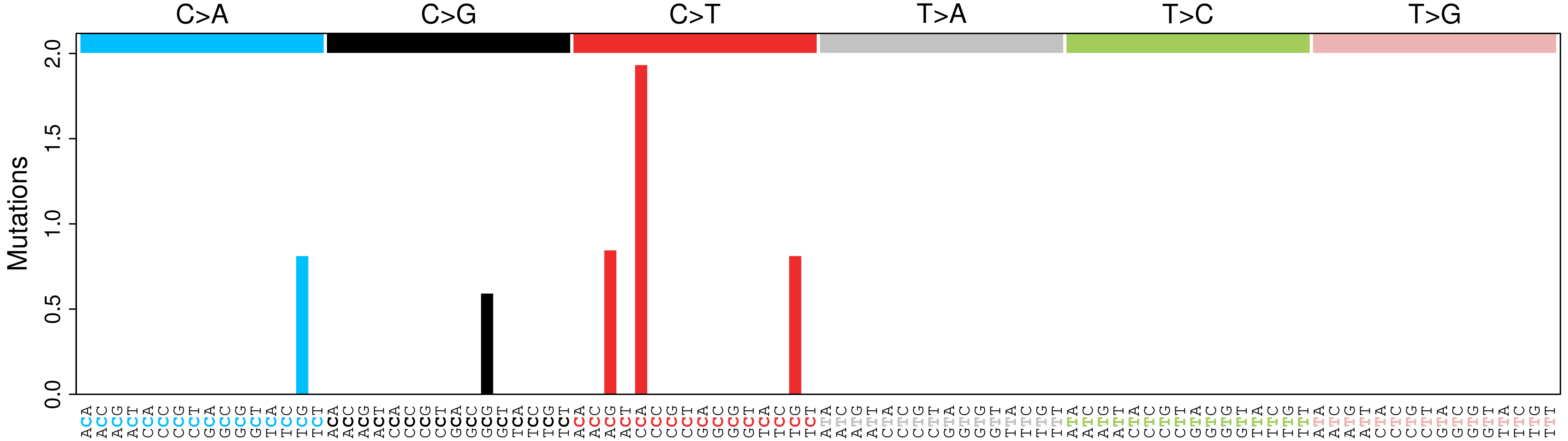
# CATD283a (13 mutations)



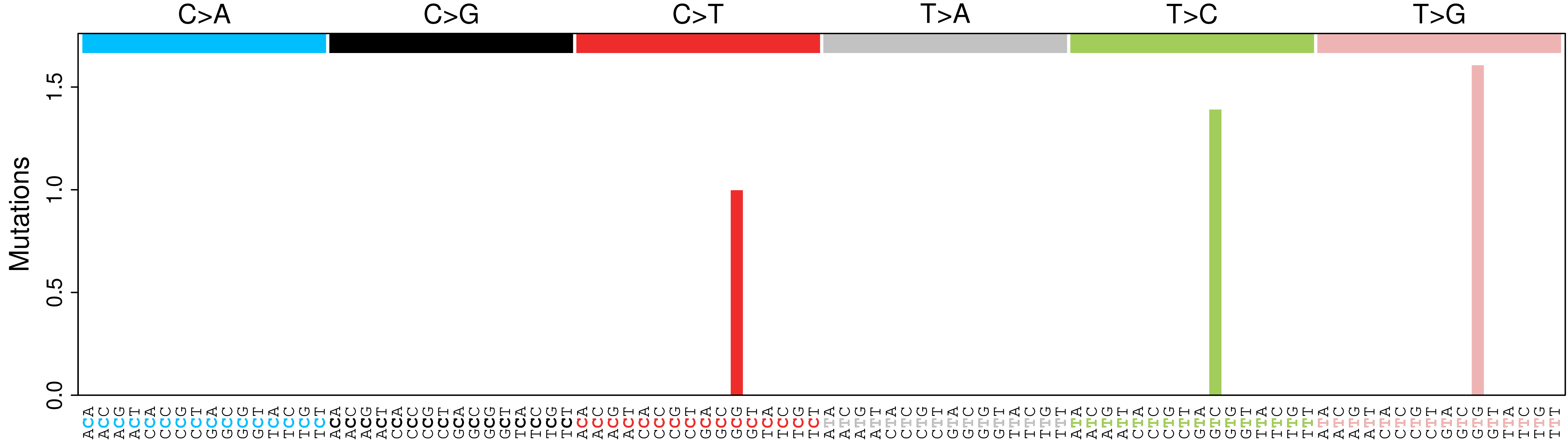
## CATD0645a (4 mutations)



# CATD0670a (5 mutations)



CATD0650a (4 mutations)



## CATD289a (2 mutations)

$C > A$

C > G

$C > T$

**T>A**

 $T > C$ 

T>G

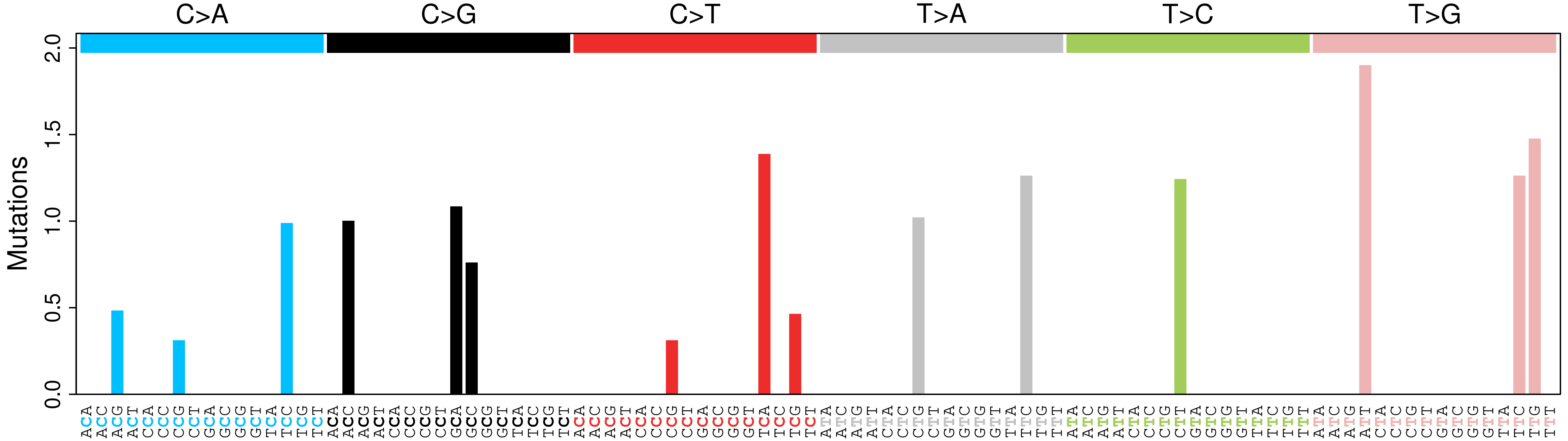
# Mutations

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

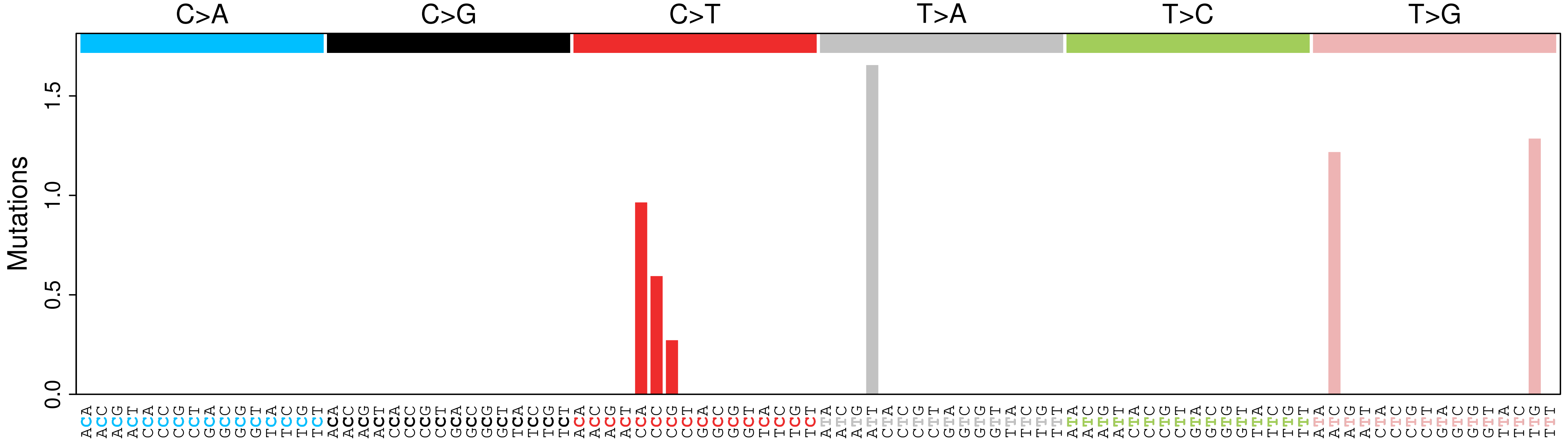
## 2.0

0.5

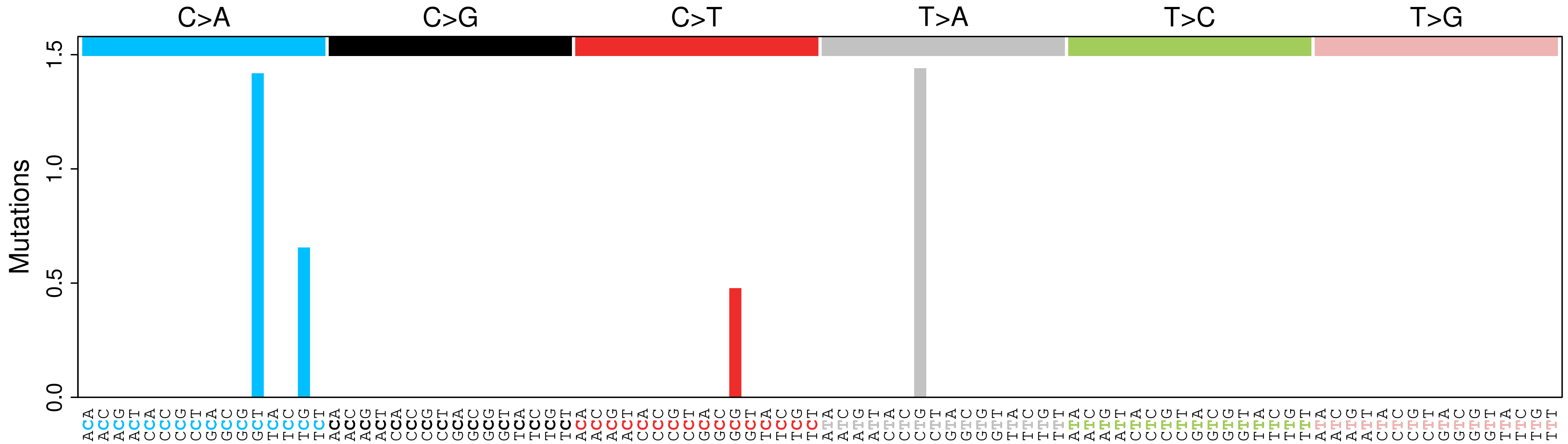
0.0

[illegible]

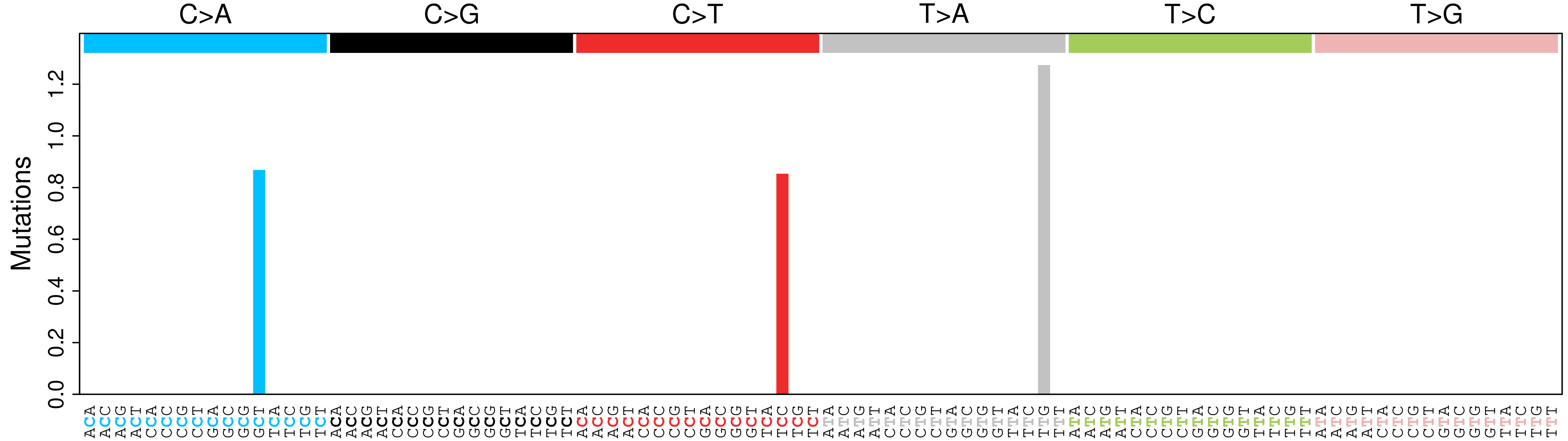
# CATD0668a (6 mutations)



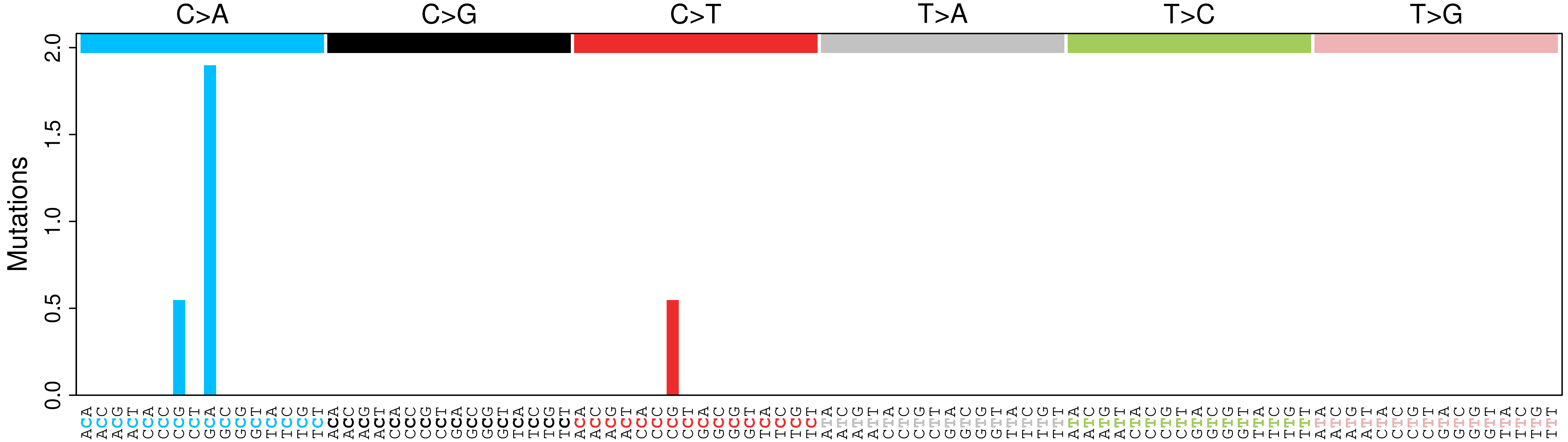
# CATD0146a (4 mutations)



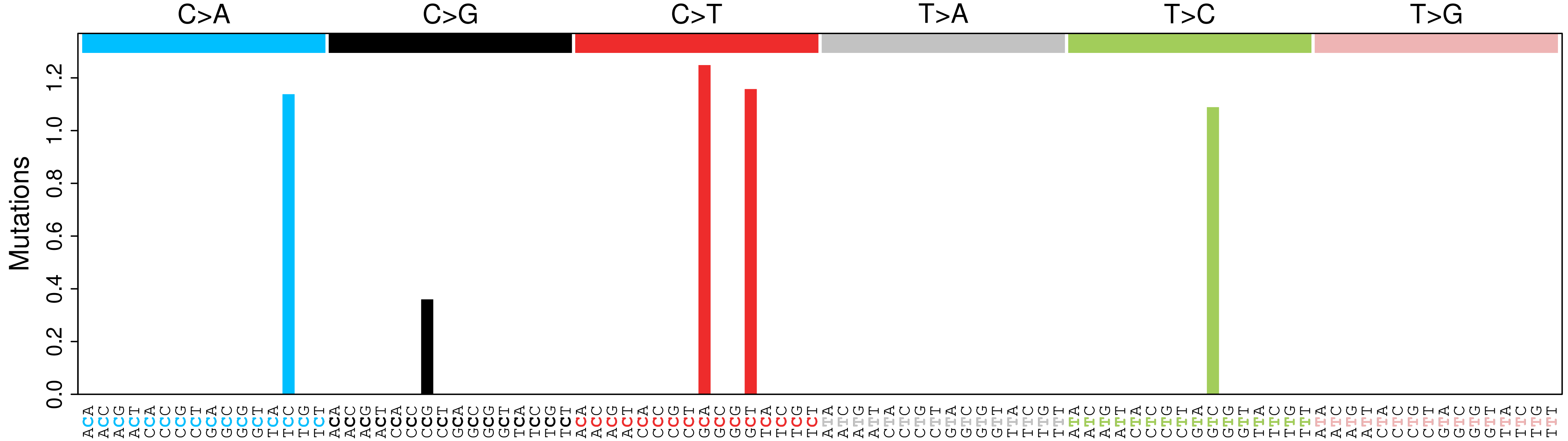
## CATD0641a (3 mutations)



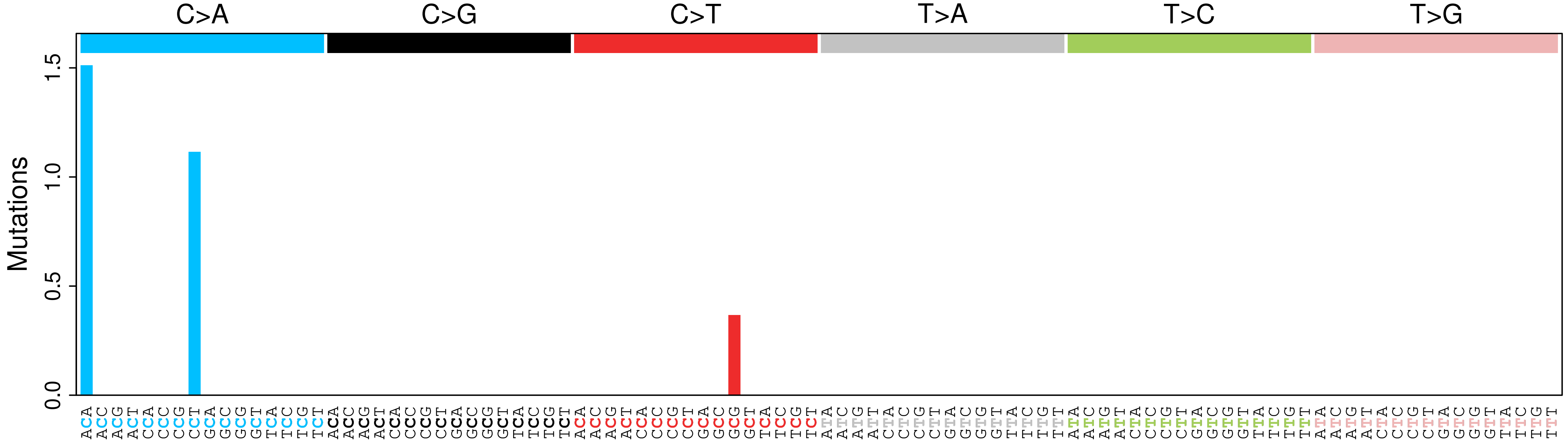
# CATD0143a (3 mutations)



## CATD0632a (5 mutations)

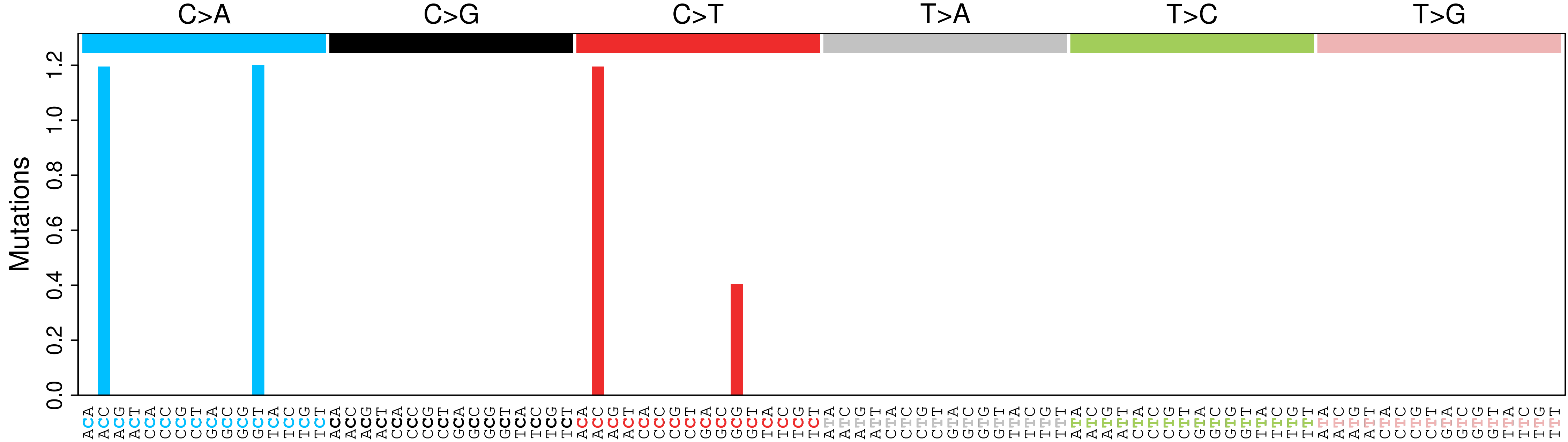


# CATD0633a (3 mutations)

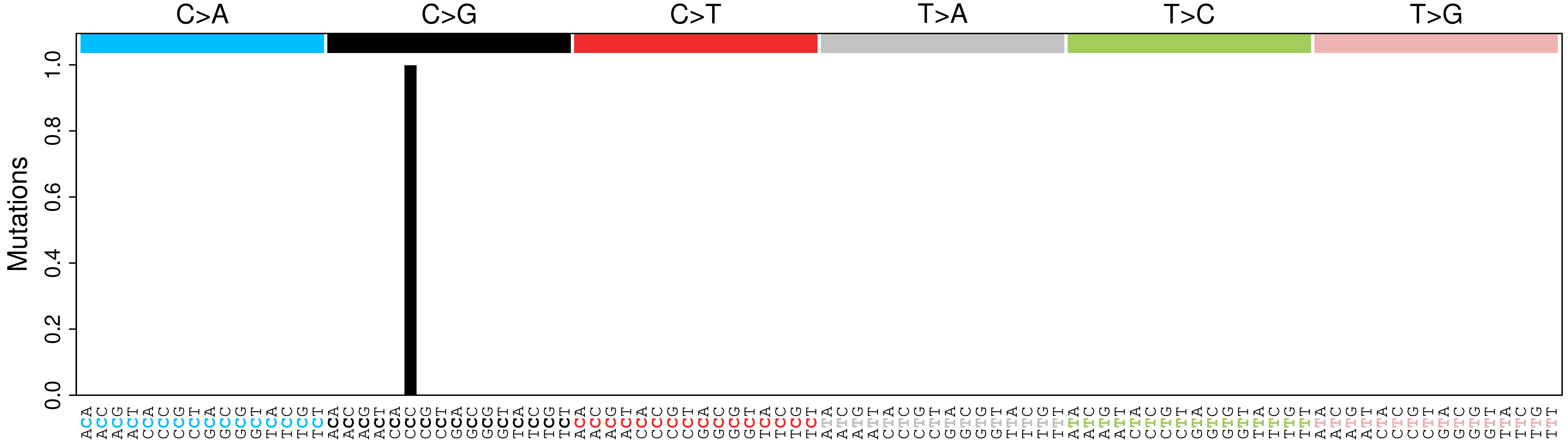




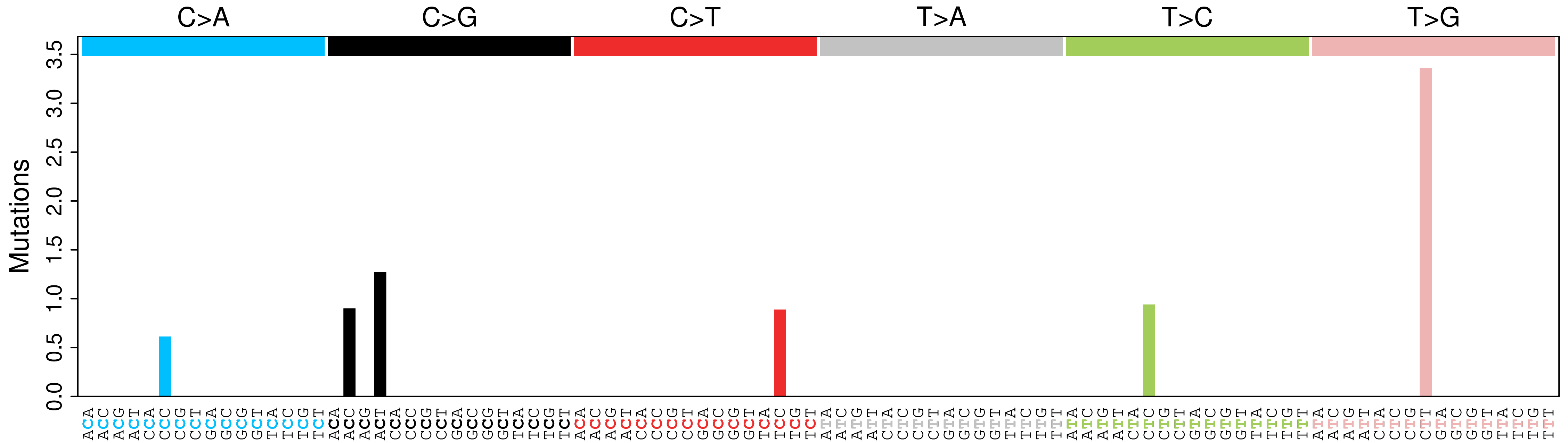
CATD0157a (4 mutations)



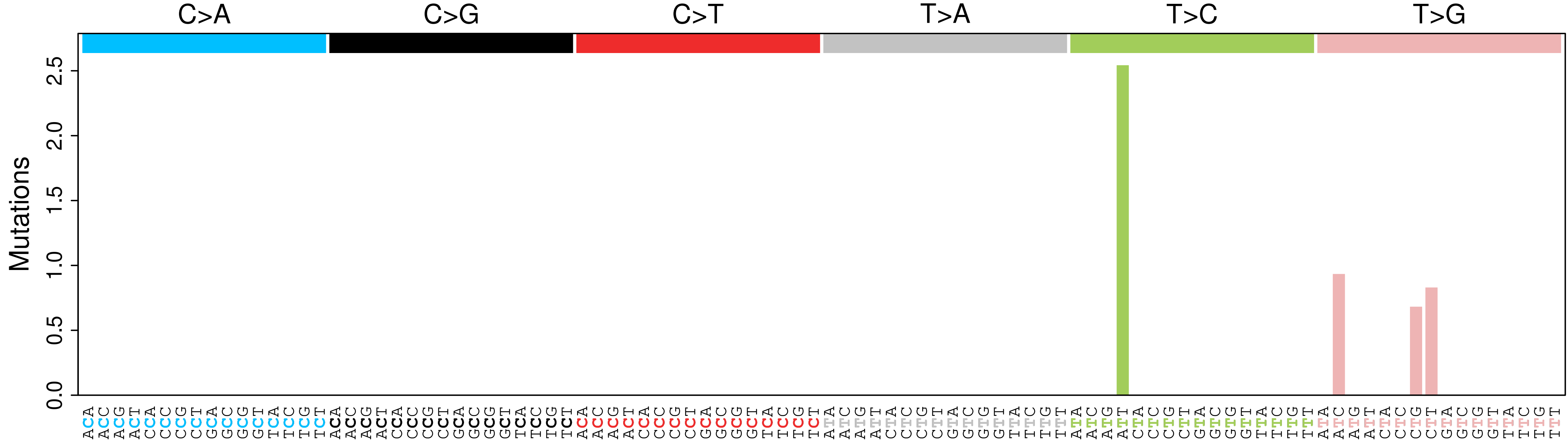
# CATD0147a (1 mutations)



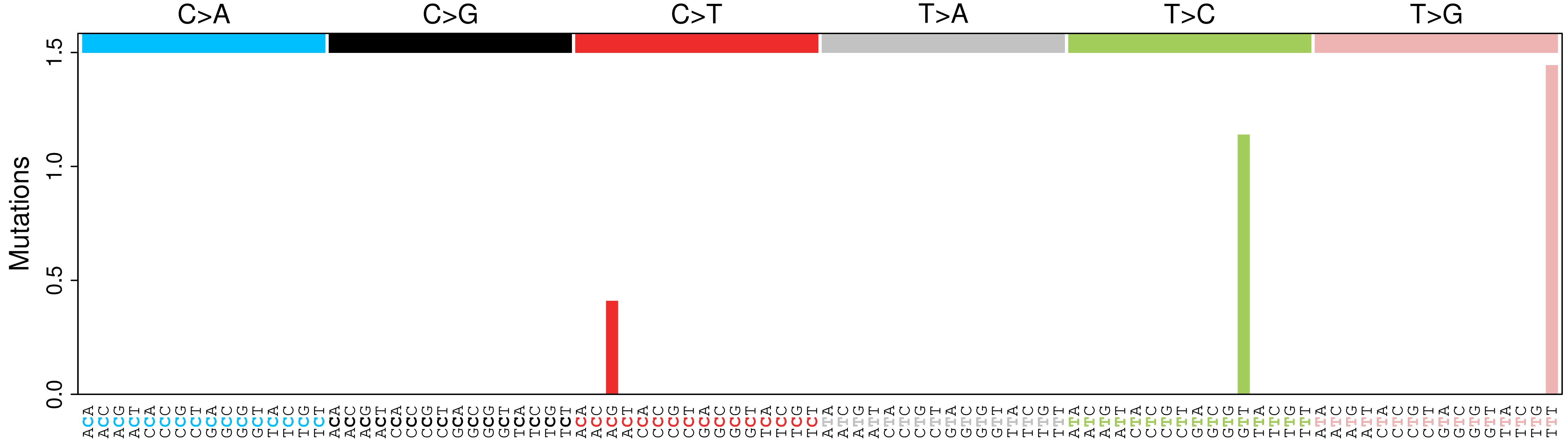
# CATD0145a (8 mutations)



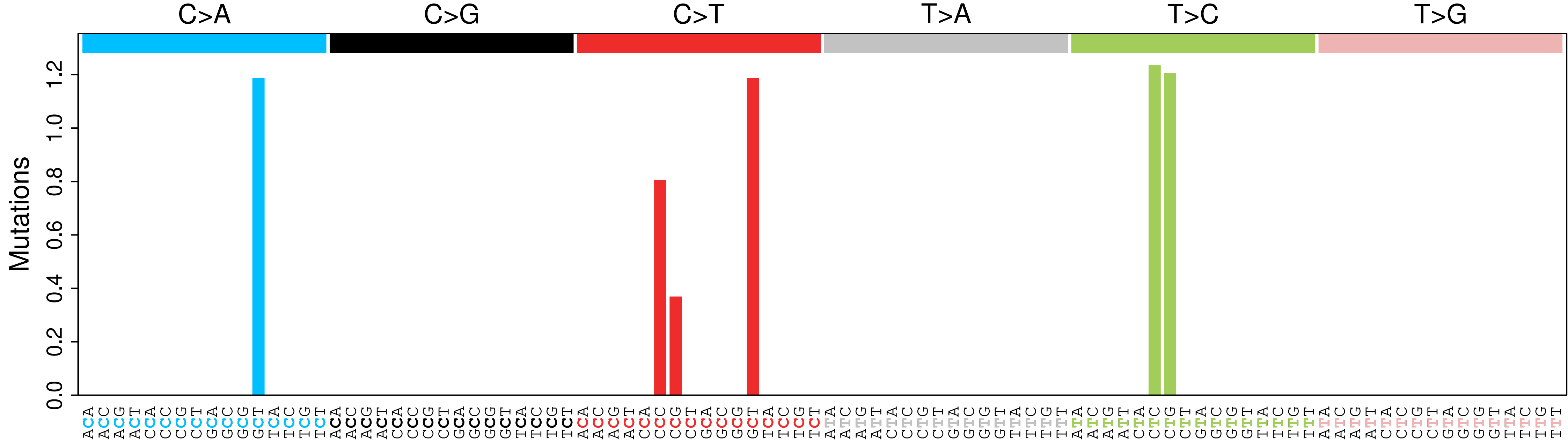
CATD0159a (5 mutations)



## CATD0148a (3 mutations)



## CATD0153a (6 mutations)



## CATD0635a (3 mutations)

**C > A**

C > G

$C > T$

**T > A**

$T > C$

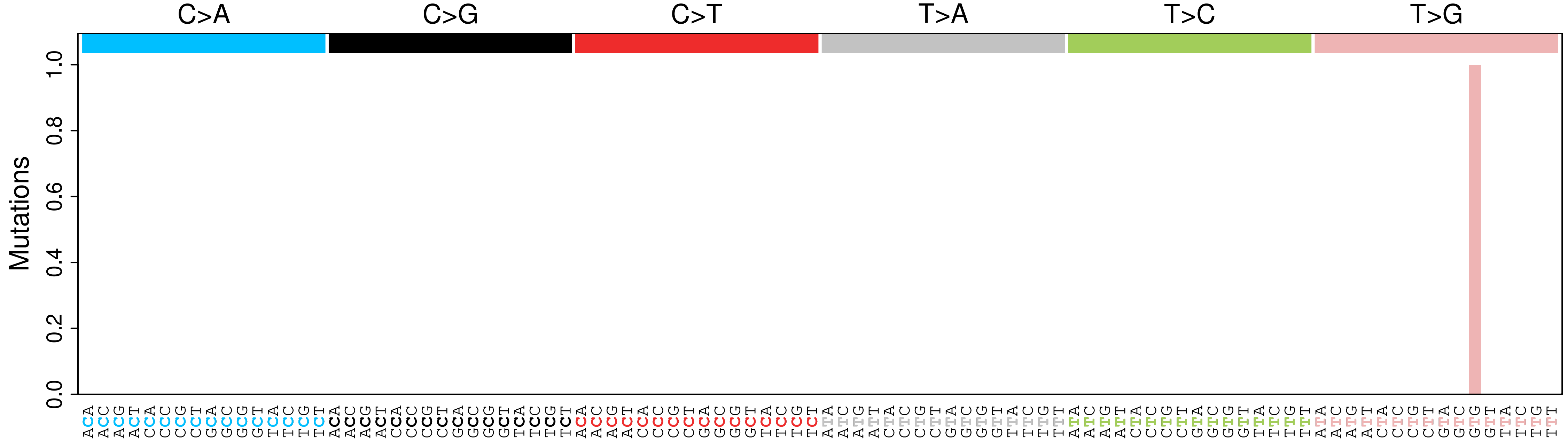
T>G

# Mutations

0.0 0.2 0.4 0.6 0.8 1.0 1.2

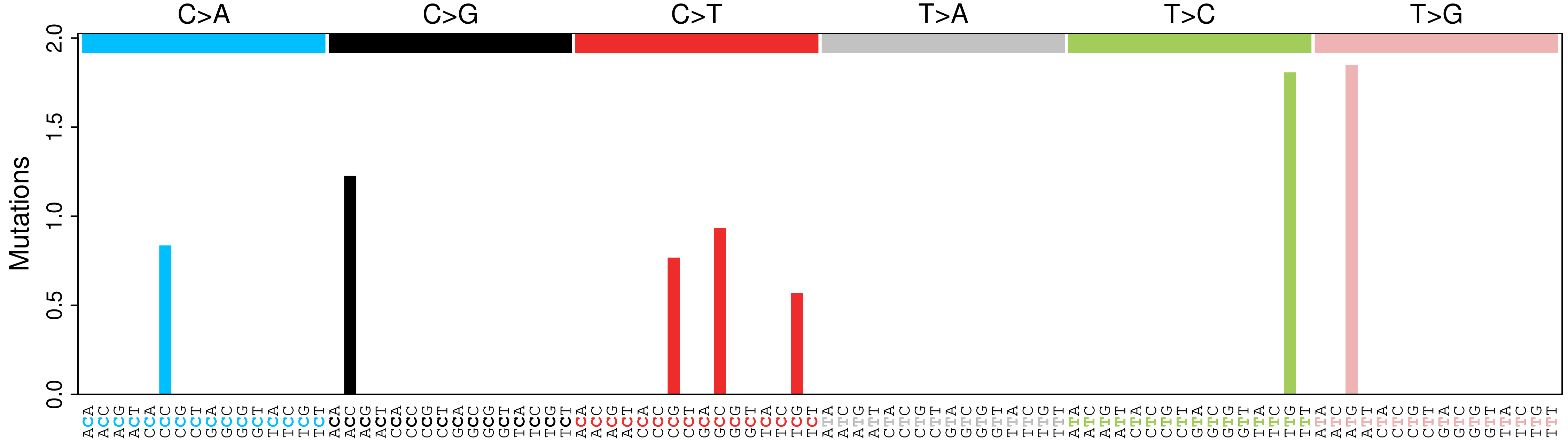
[illegible]

# CATD0646a (1 mutations)





# CATD295a (8 mutations)



## CATD290a (1 mutations)

$C > A$

C > G

$$C > T$$

**T>A**

$$T > C$$

T>G

# Mutations

# 1.0

 $\circ$ 

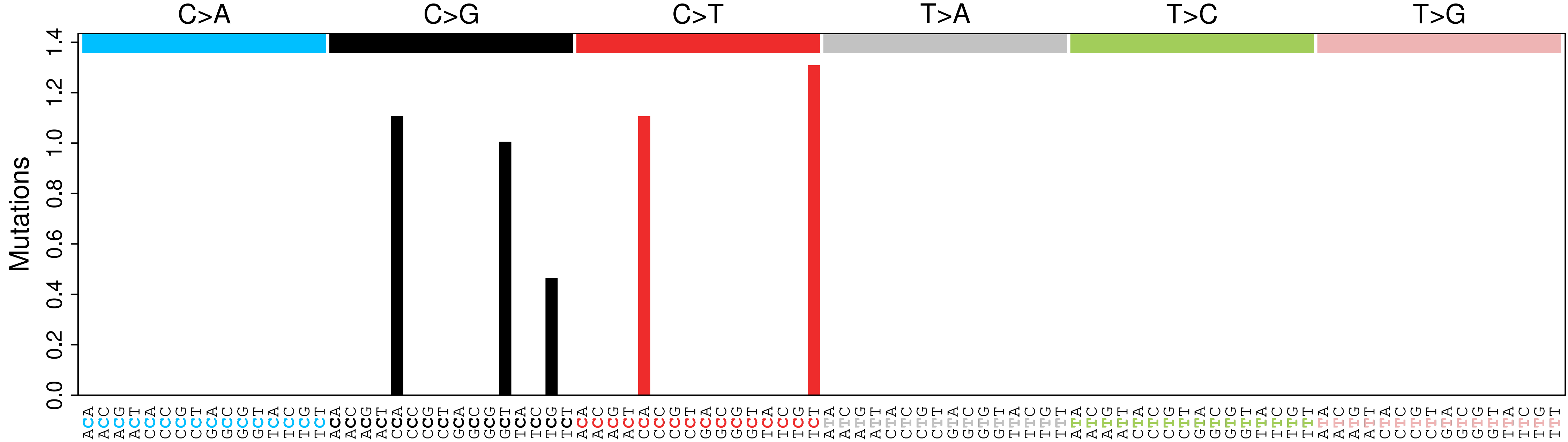
0

0

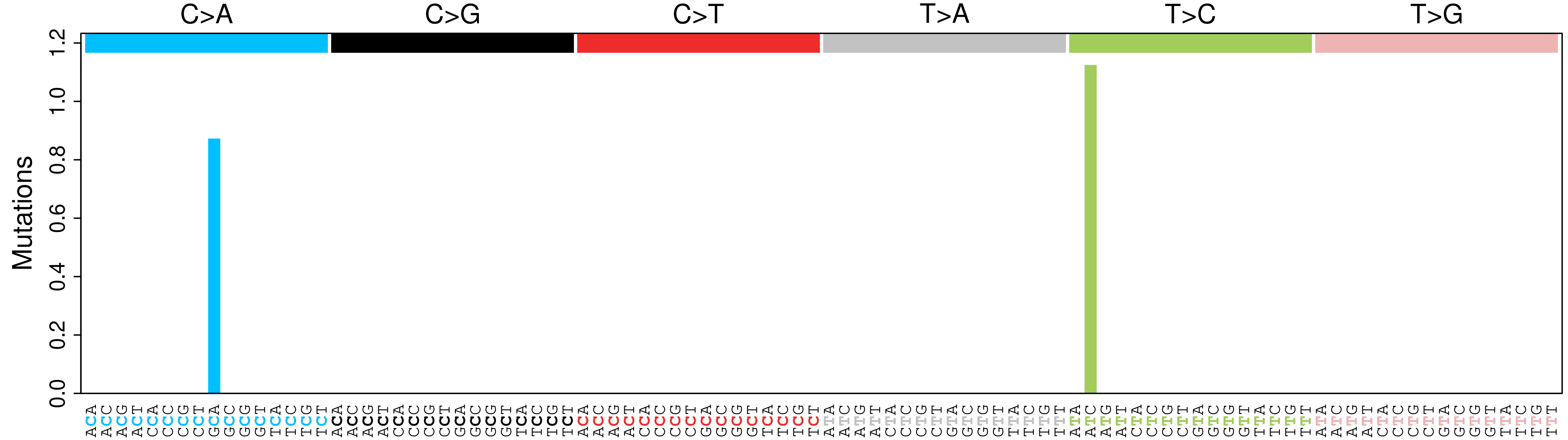
0.0

[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**

$T > C$

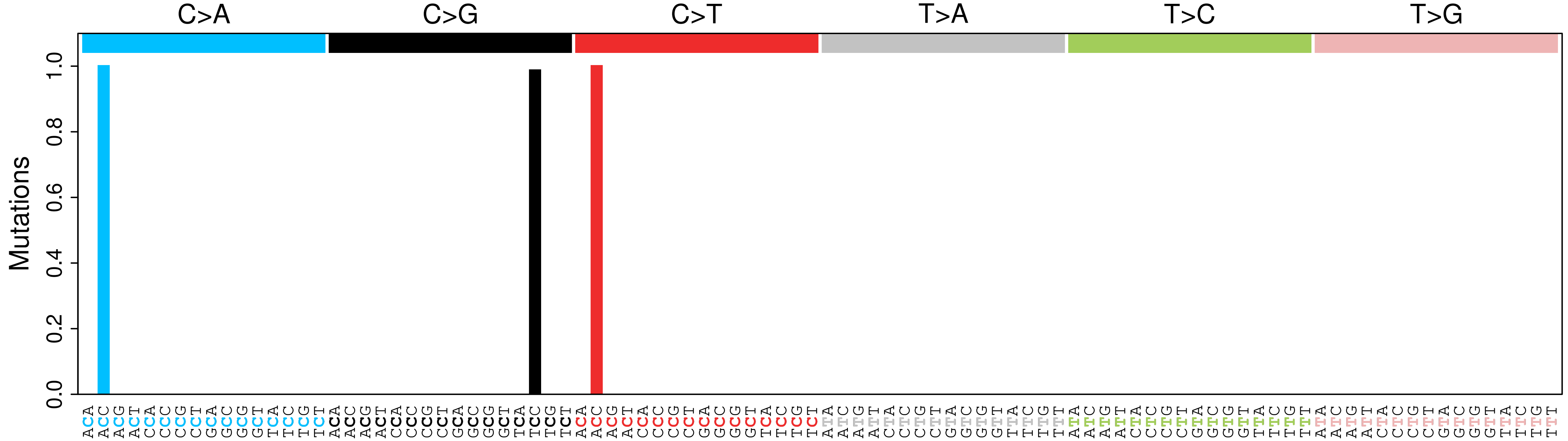
T > G

# Mutations

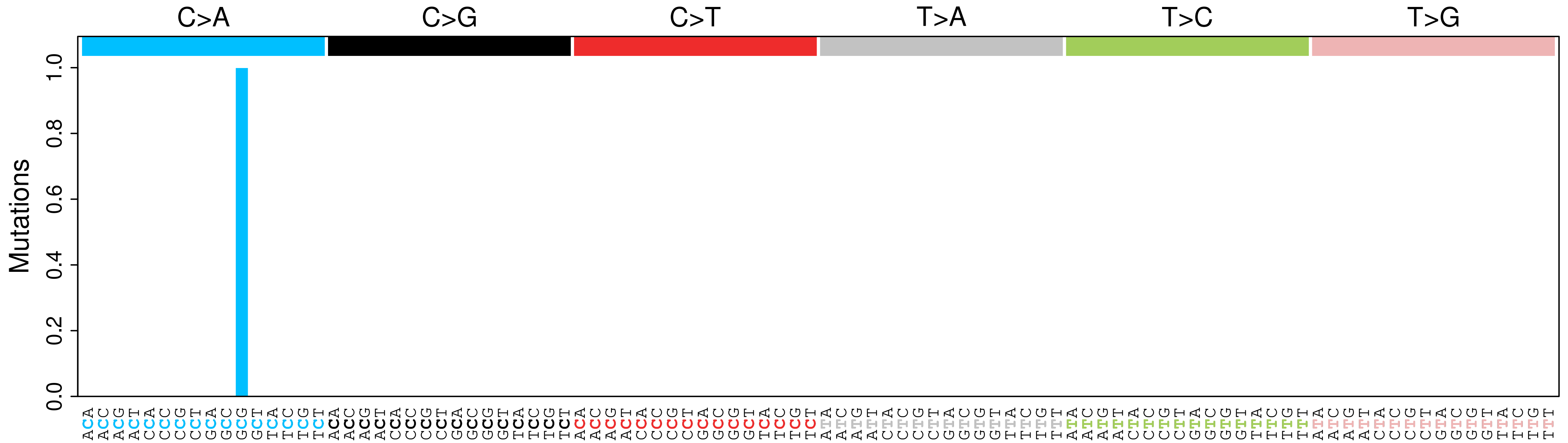
0.0 0.2 0.4 0.6 0.8 1.0 1.2

[illegible]

## 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

0

[illegible]

## CATD0644a (1 mutations)

