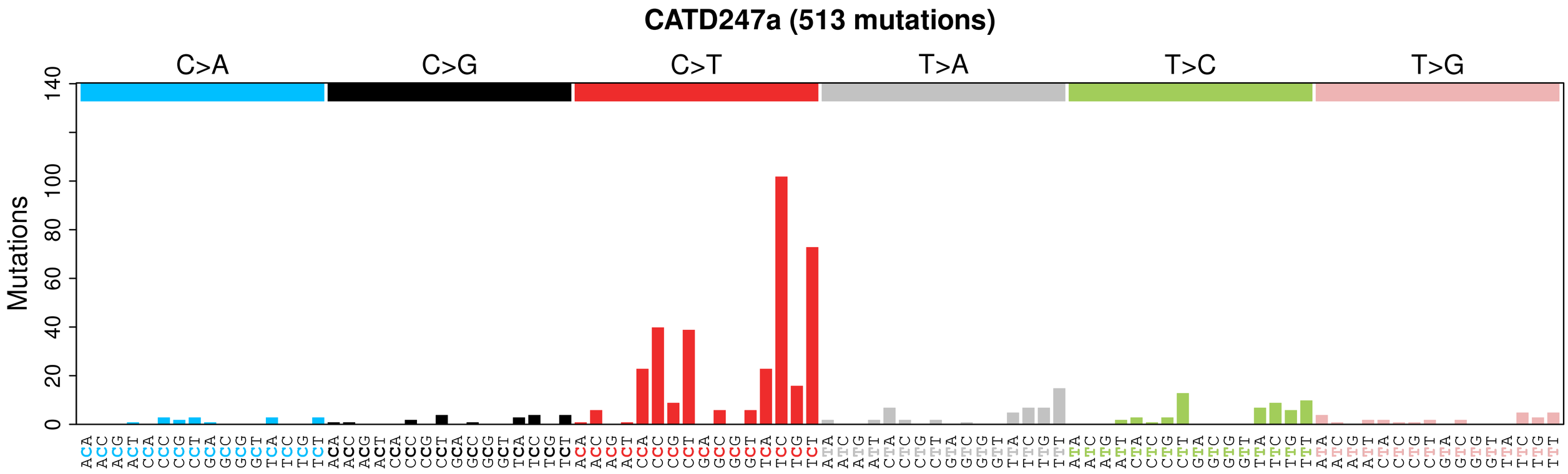
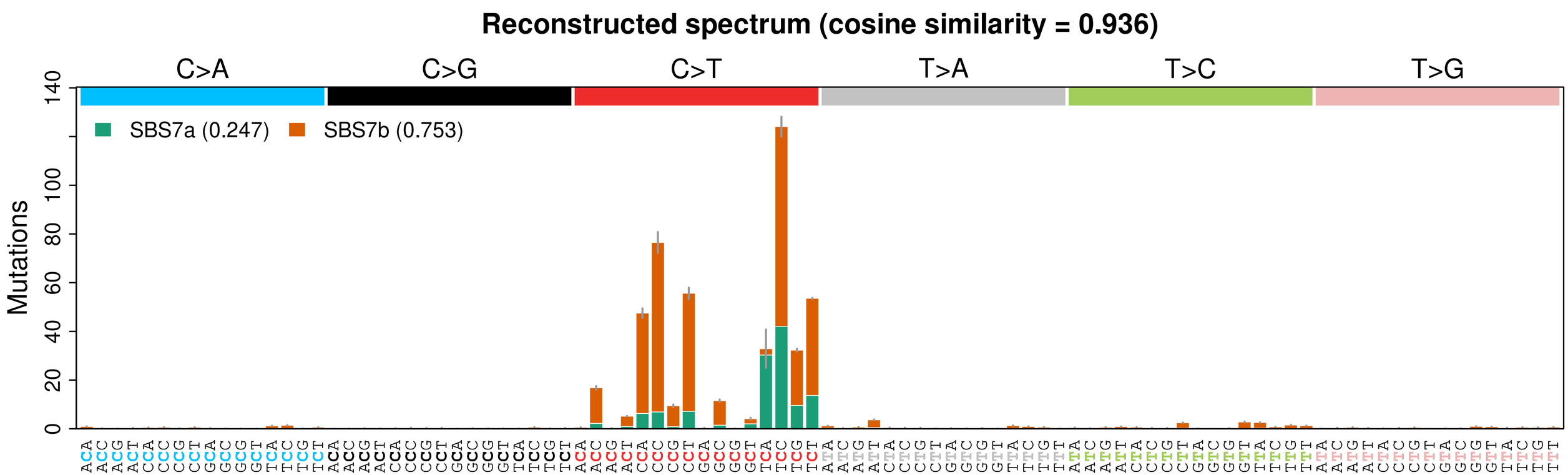

CATD247a (513 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
20
40
60
80
100
140


Reconstructed spectrum (cosine similarity = 0.936)
C>A
C>G
C>T
T>A
T>C
T>G
SBS7a (0.247)
SBS7b (0.753)
Mutations
0
20
40
60
80
100
140