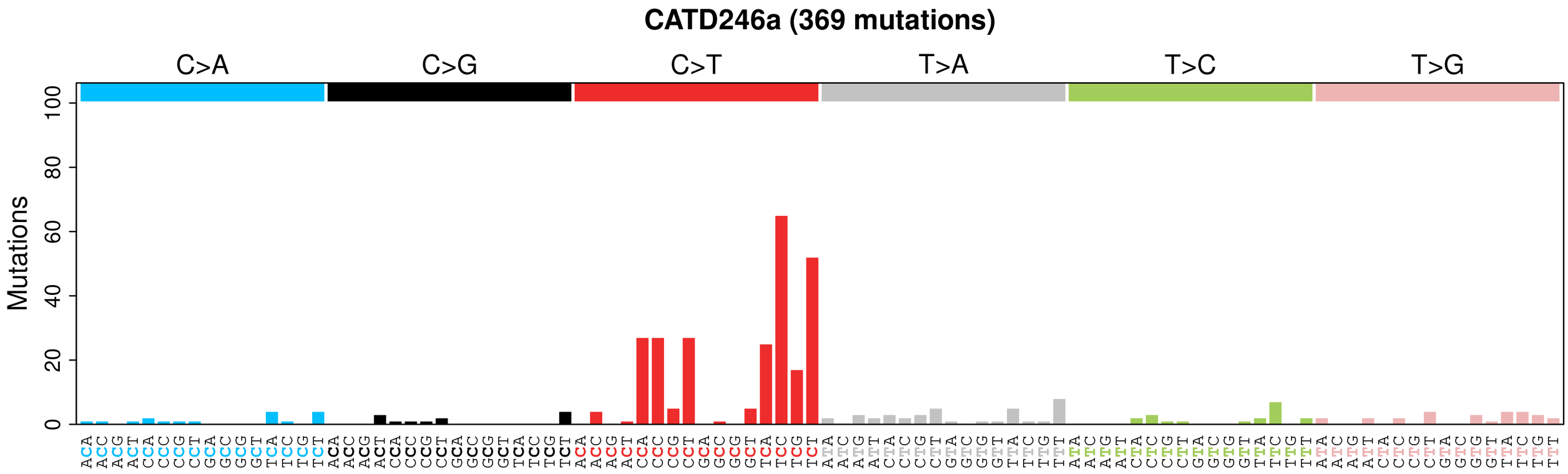
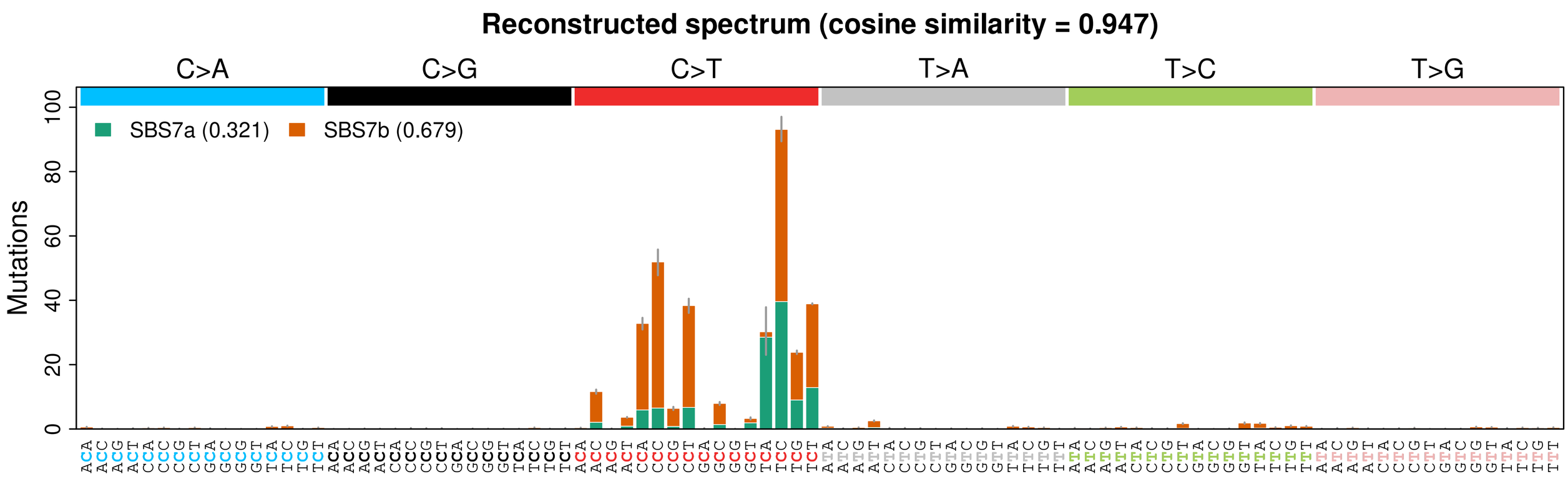

CATD246a (369 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
20
40
60
80
100


Reconstructed spectrum (cosine similarity = 0.947)
C>A
C>G
C>T
T>A
T>C
T>G
SBS7a (0.321)
SBS7b (0.679)
Mutations
0
20
40
60
80
100