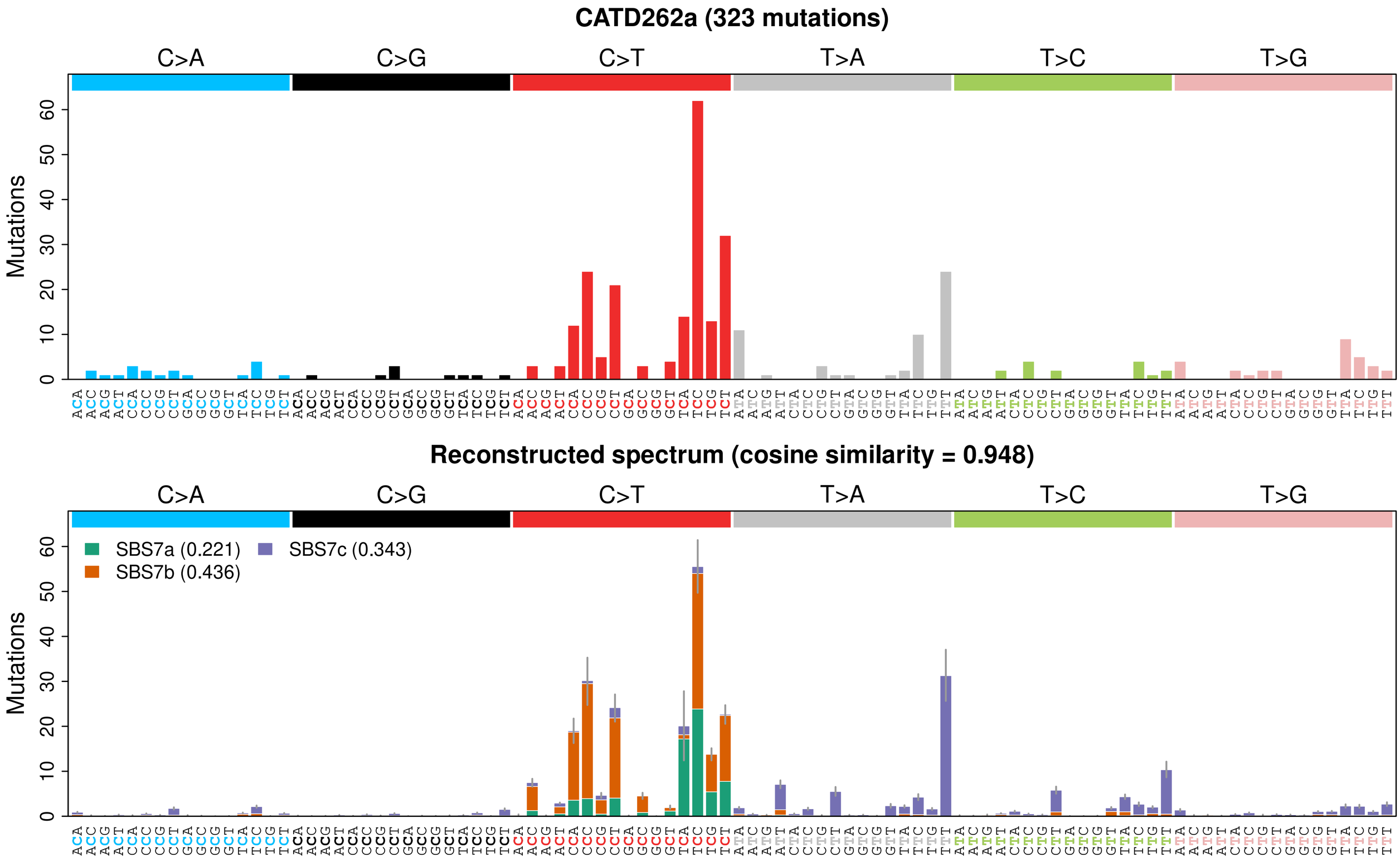
CATD262a (323 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
10
20
30
40
50
60
Reconstructed spectrum (cosine similarity = 0.948)
SBS7a (0.221)
SBS7b (0.436)
SBS7c (0.343)