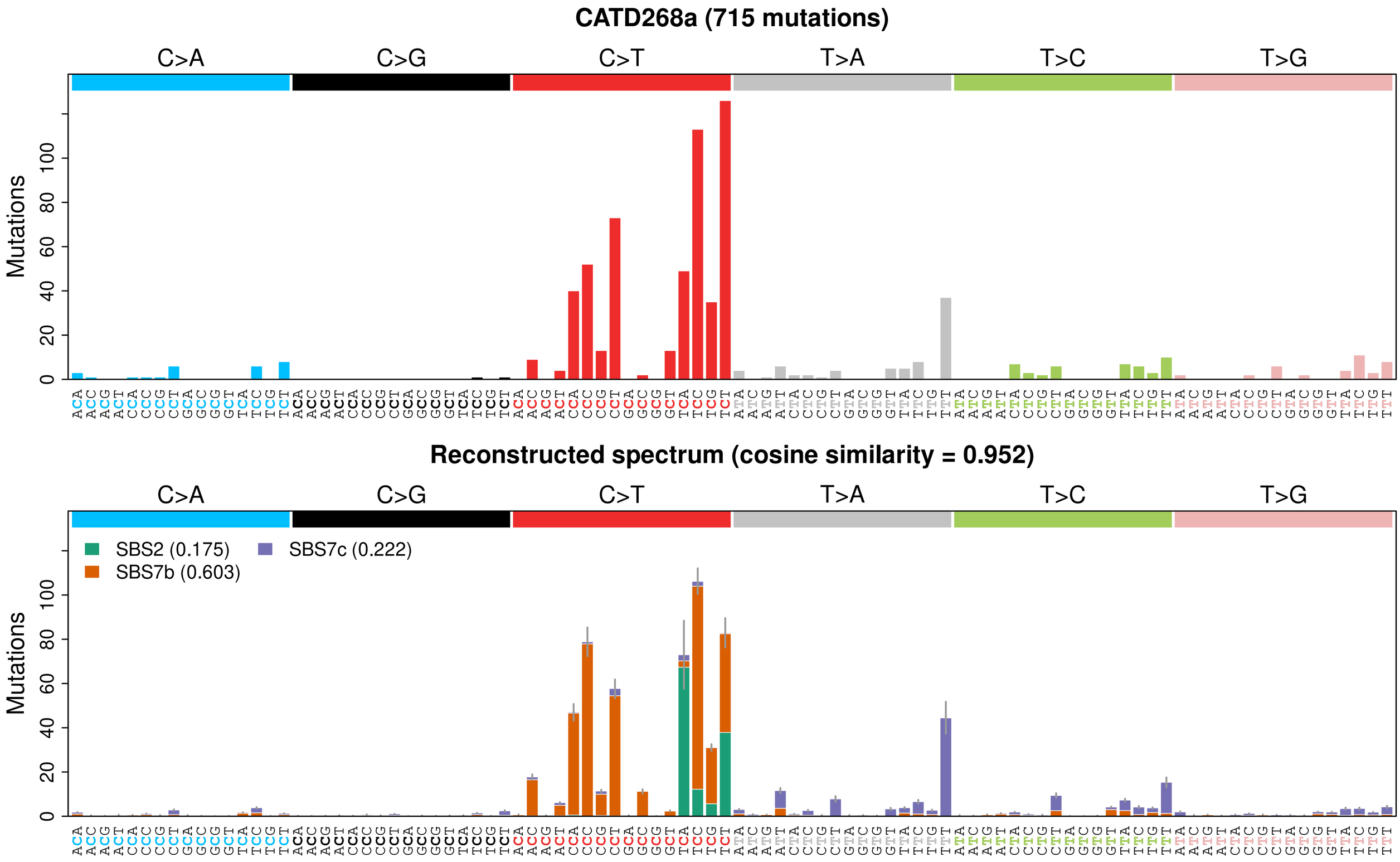
CATD268a (715 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
20
40
60
80
100
Reconstructed spectrum (cosine similarity = 0.952)
SBS2 (0.175)
SBS7b (0.603)
SBS7c (0.222)