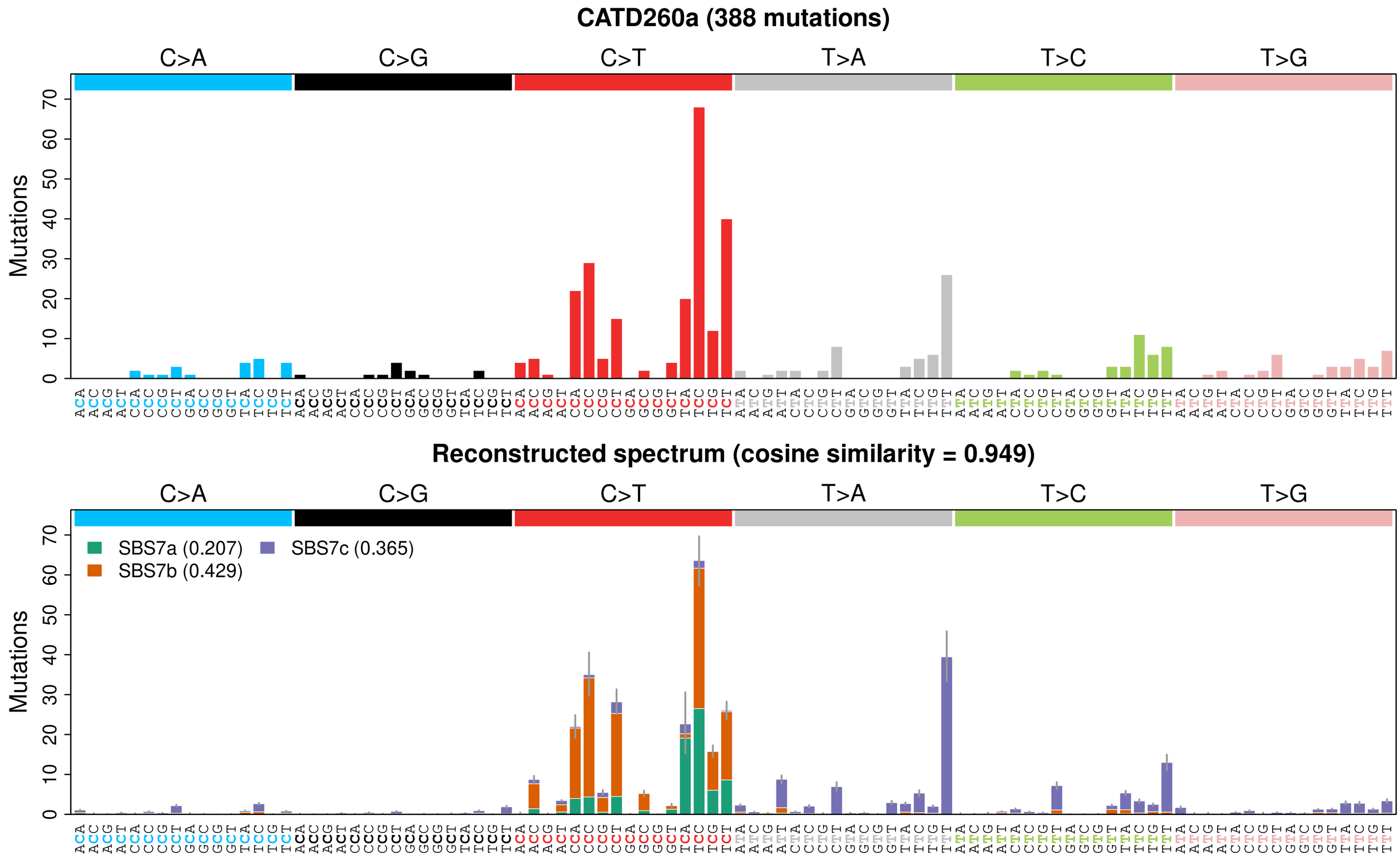
CATD260a (388 mutations)
C>A
C>G
C>T
T>A
T>C
T>G
Mutations
0
10
20
30
40
50
60
70
Reconstructed spectrum (cosine similarity = 0.949)
SBS7a (0.207)
SBS7b (0.429)
SBS7c (0.365)