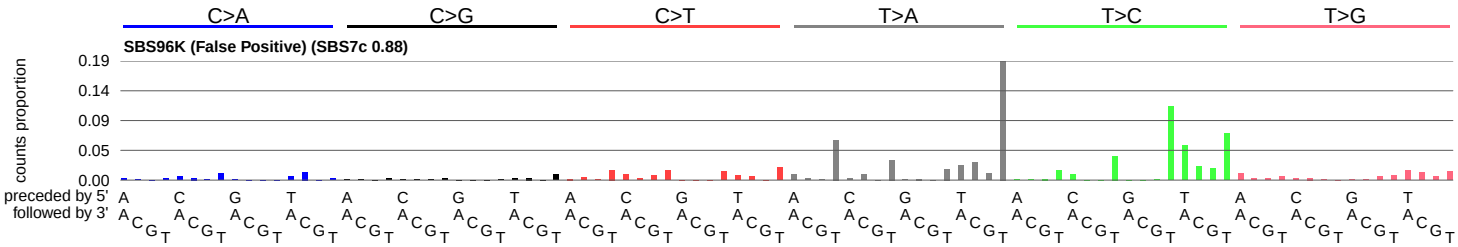
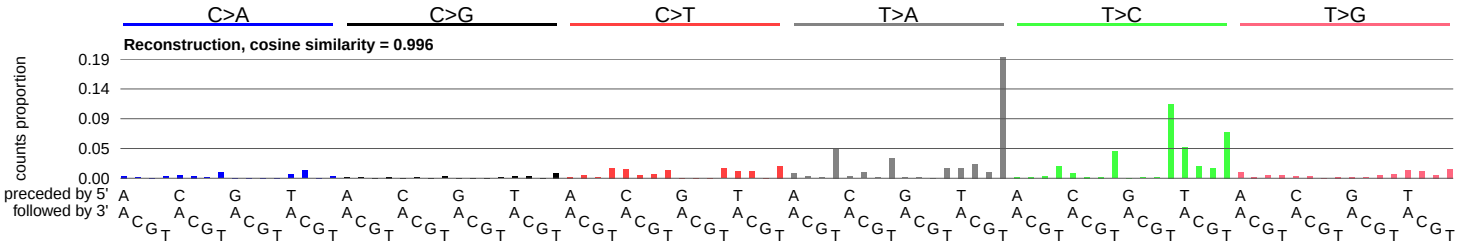
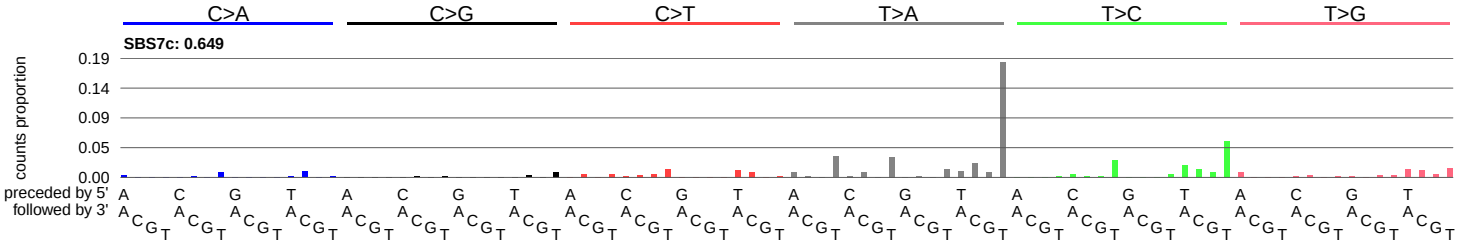
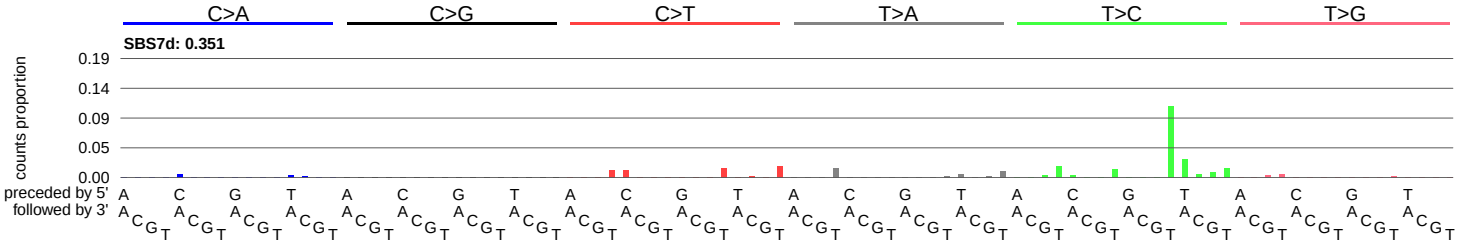

C>A
C>G
C>T
T>A
T>C
T>G
SBS96K (False Positive) (SBS7c 0.88)
counts proportion
0.19
0.14
0.09
0.05
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.996
counts proportion
0.19
0.14
0.09
0.05
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS7c: 0.649
counts proportion
0.19
0.14
0.09
0.05
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS7d: 0.351
counts proportion
0.19
0.14
0.09
0.05
0.00
preceded by 5'
followed by 3'