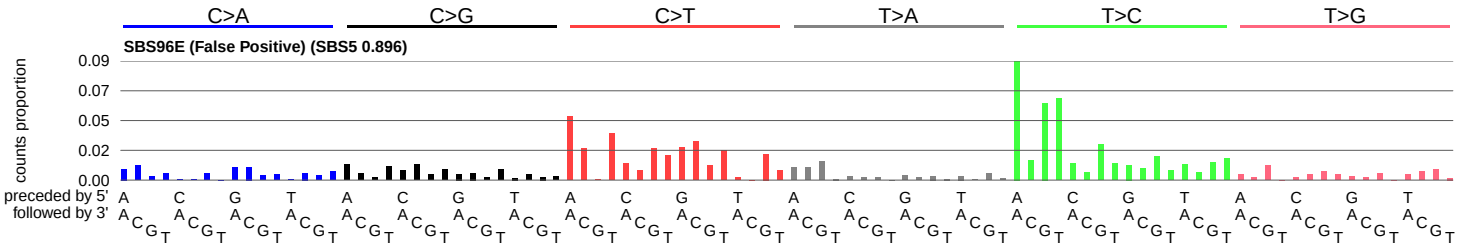
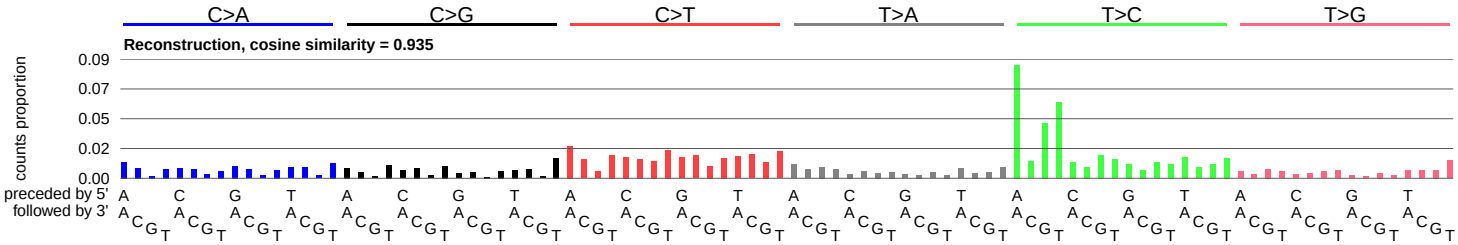
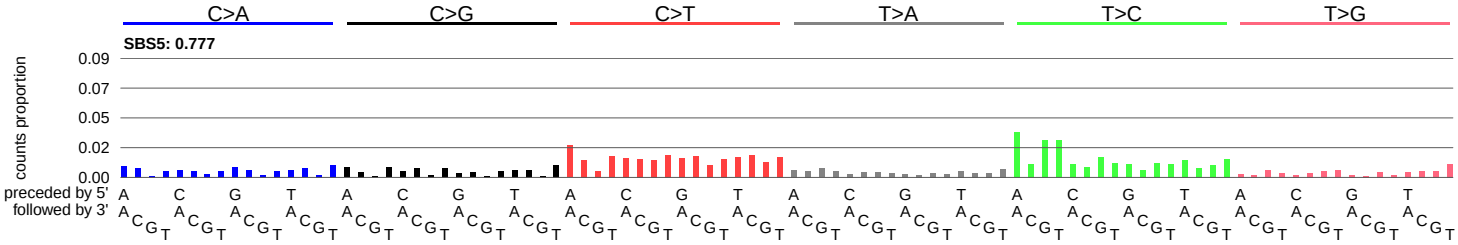
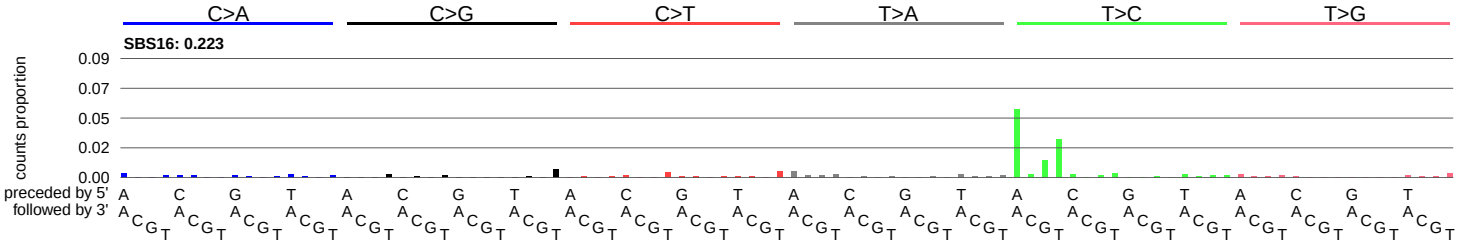

C>A
C>G
C>T
T>A
T>C
T>G
SBS96E (False Positive) (SBS5 0.896)
counts proportion
0.09
0.07
0.05
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.935
counts proportion
0.09
0.07
0.05
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS5: 0.777
counts proportion
0.09
0.07
0.05
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS16: 0.223
counts proportion
0.09
0.07
0.05
0.02
0.00
preceded by 5'
followed by 3'