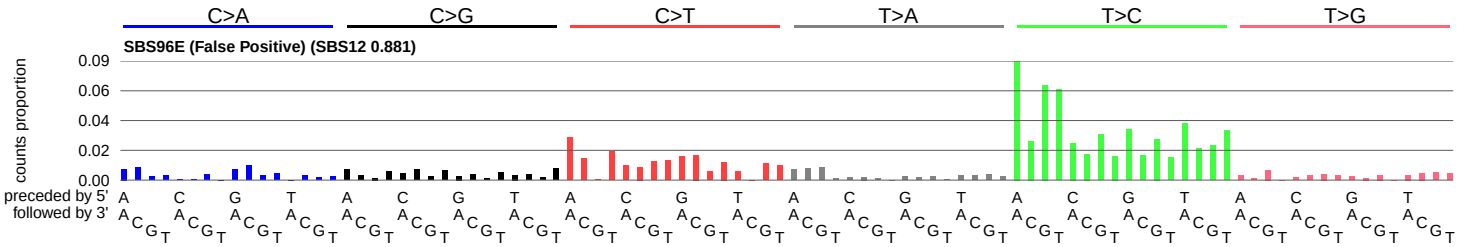
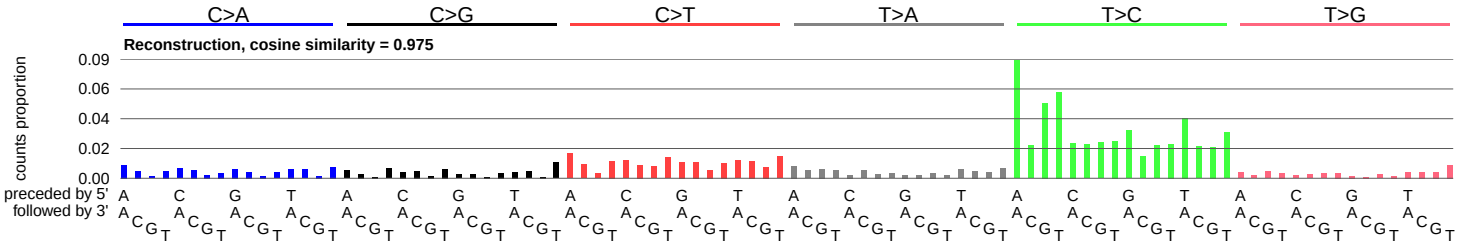
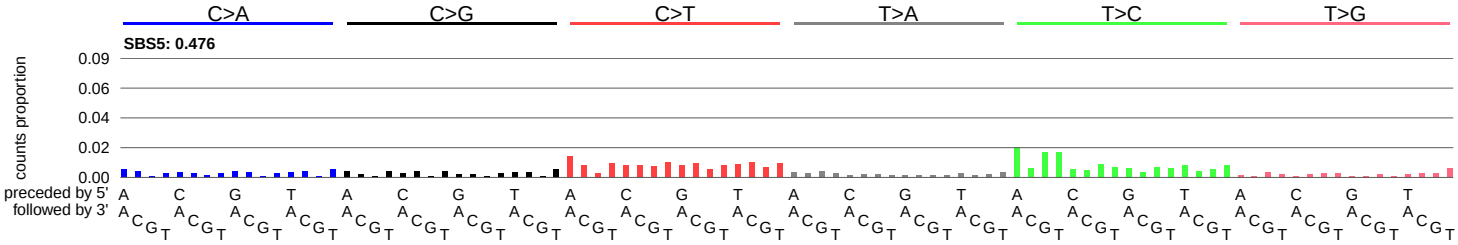
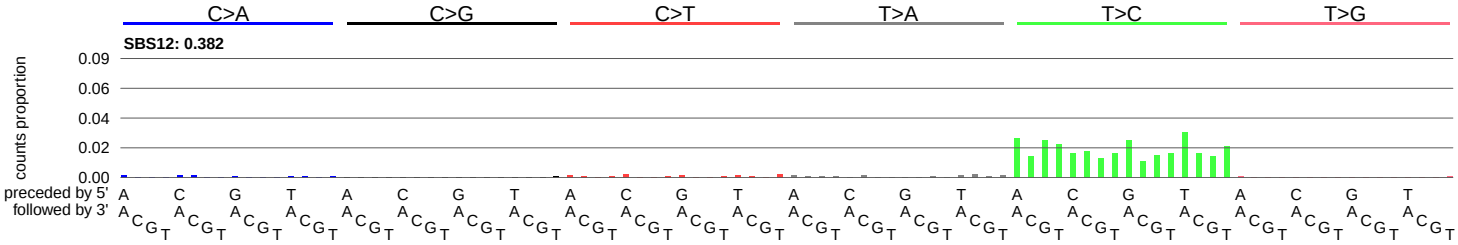
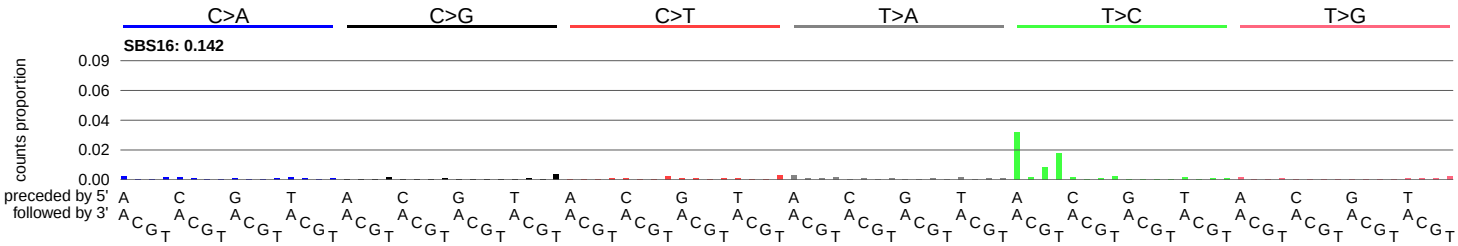

C>A
C>G
C>T
T>A
T>C
T>G
SBS96E (False Positive) (SBS12 0.881)
counts proportion
0.09
0.06
0.04
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.975
counts proportion
0.09
0.06
0.04
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS5: 0.476
counts proportion
0.09
0.06
0.04
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS12: 0.382
counts proportion
0.09
0.06
0.04
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS16: 0.142
counts proportion
0.09
0.06
0.04
0.02
0.00
preceded by 5'
followed by 3'