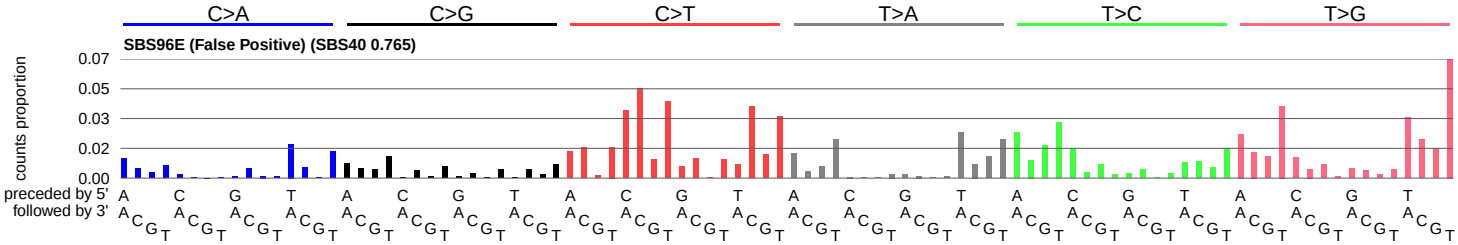
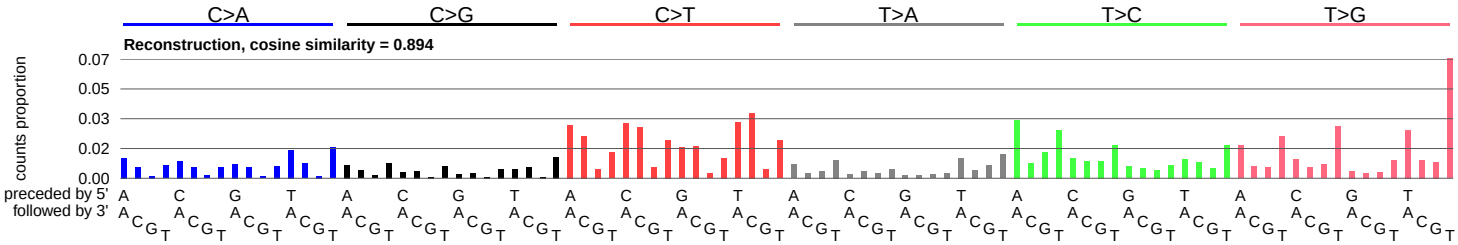
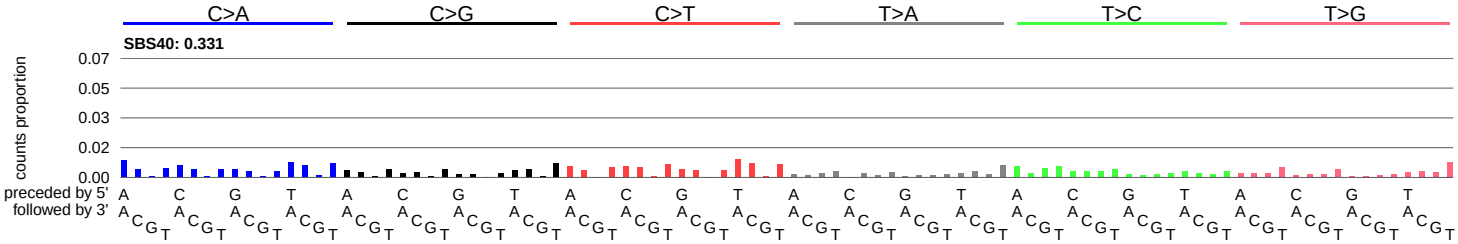
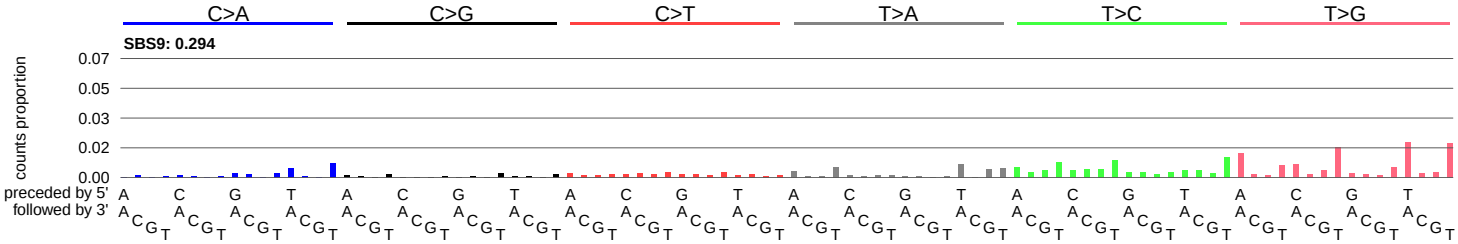
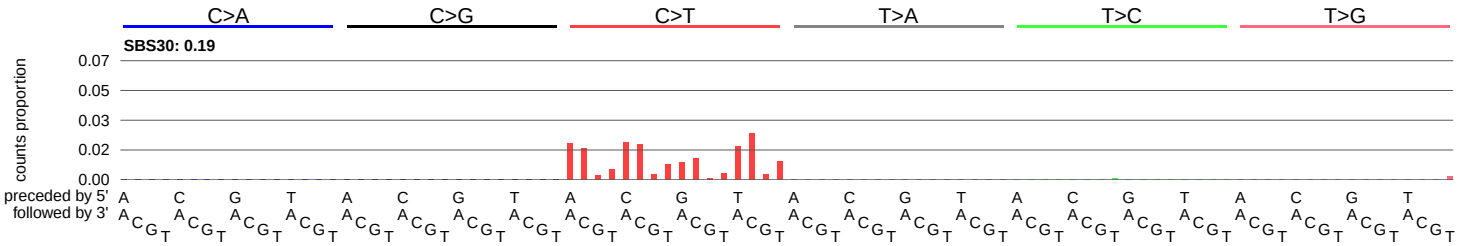
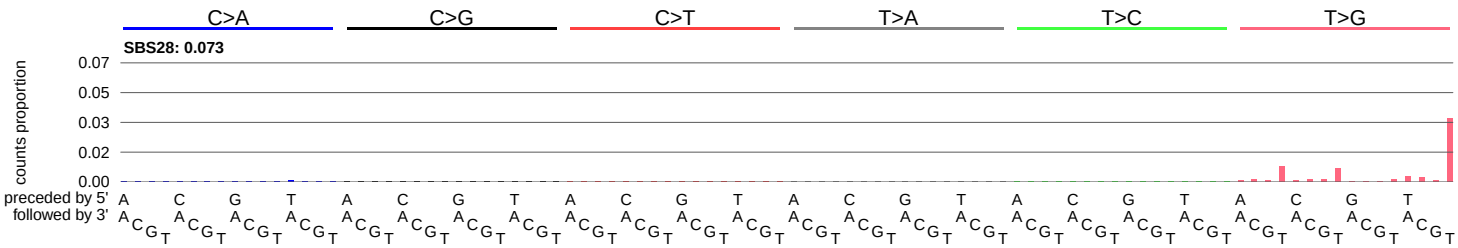

C>A
C>G
C>T
T>A
T>C
T>G
SBS96E (False Positive) (SBS40 0.765)
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

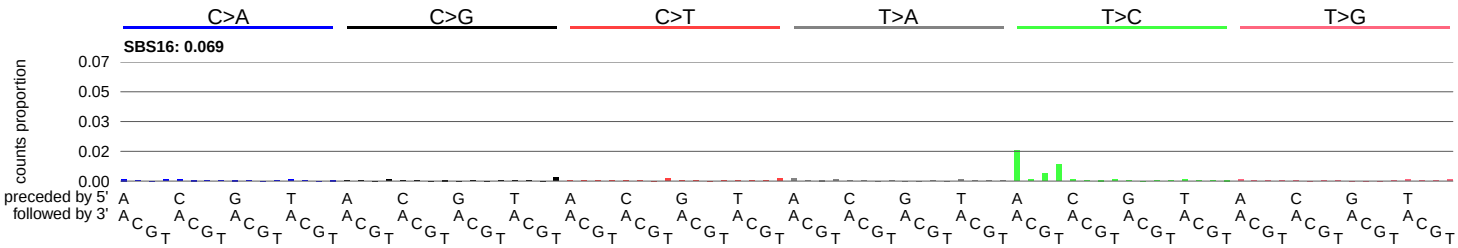

C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.894
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

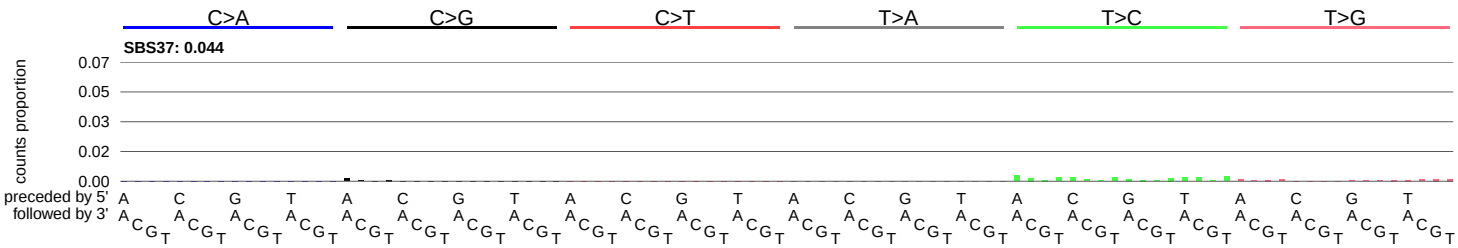

C>A
C>G
C>T
T>A
T>C
T>G
SBS40: 0.331
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS9: 0.294
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS30: 0.19
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS28: 0.073
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS16: 0.069
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS37: 0.044
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'