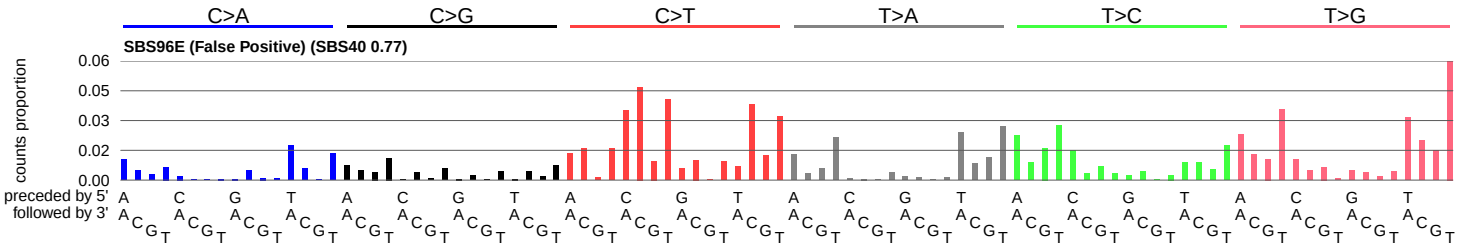
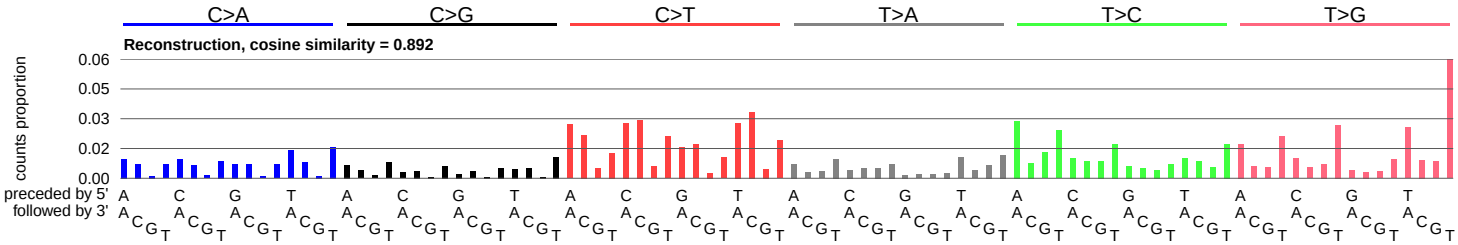
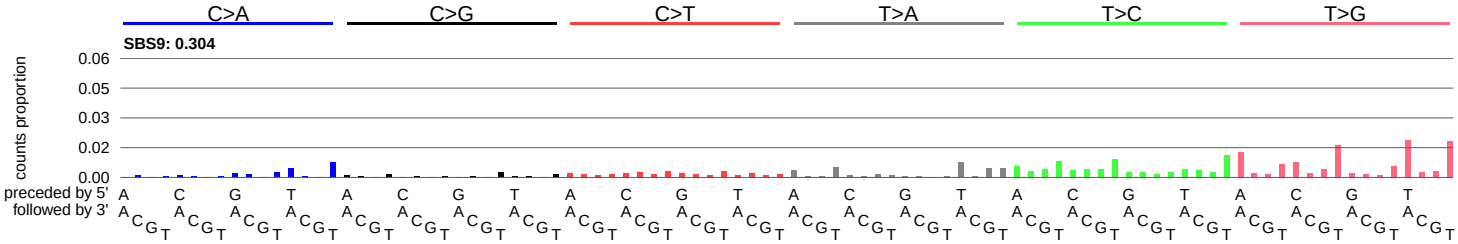
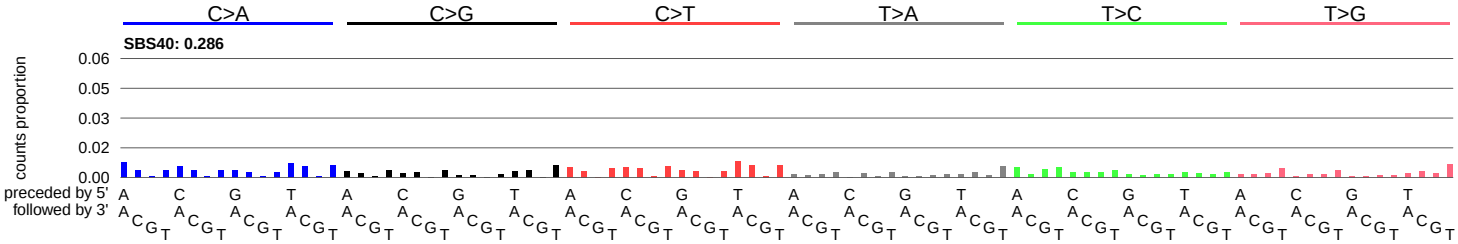
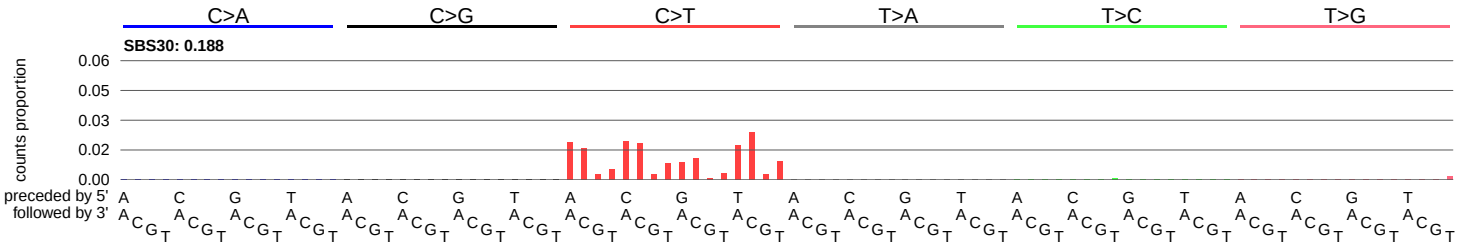
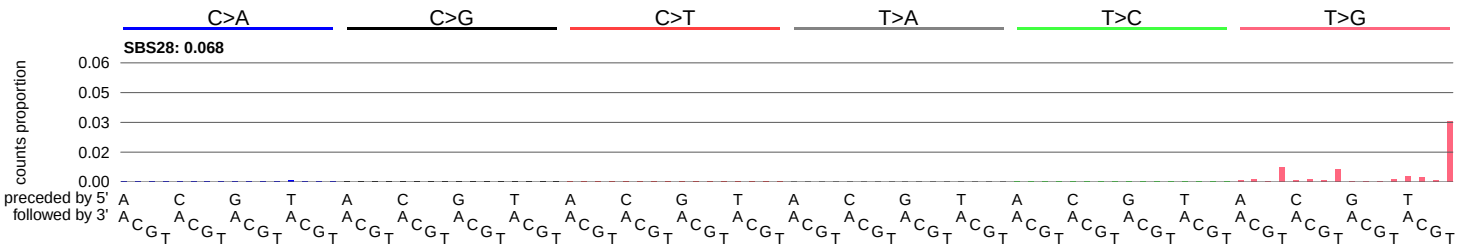

C>A
C>G
C>T
T>A
T>C
T>G
SBS96E (False Positive) (SBS40 0.77)
counts proportion
0.06
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

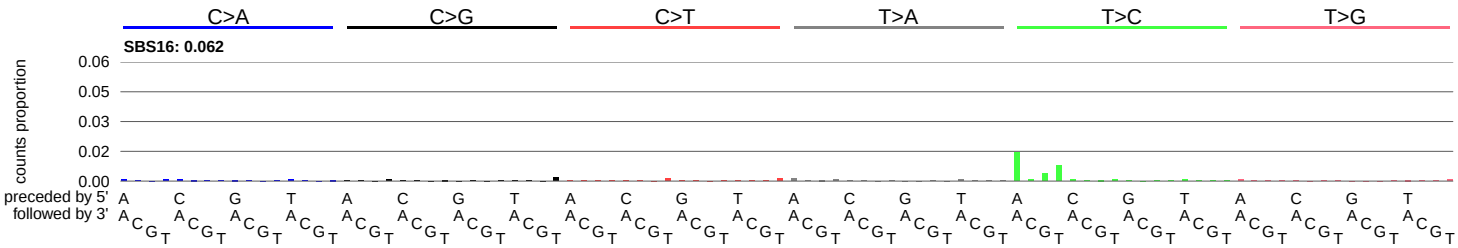

C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.892
counts proportion
0.06
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

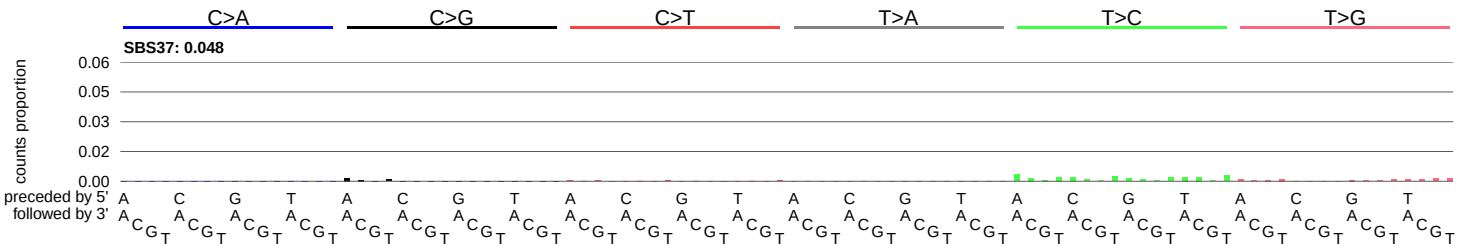

C>A
C>G
C>T
T>A
T>C
T>G
SBS9: 0.304
counts proportion
0.06
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

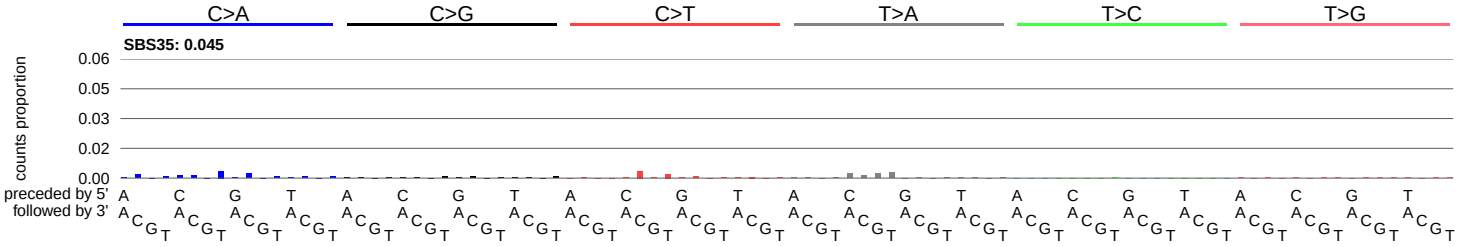

C>A
C>G
C>T
T>A
T>C
T>G
SBS40: 0.286
counts proportion
0.06
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS30: 0.188
counts proportion
0.06
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: 0.068
counts proportion
0.06
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: 0.062
counts proportion
0.06
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: 0.048
counts proportion
0.06
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
SBS35: 0.045
counts proportion
0.06
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'