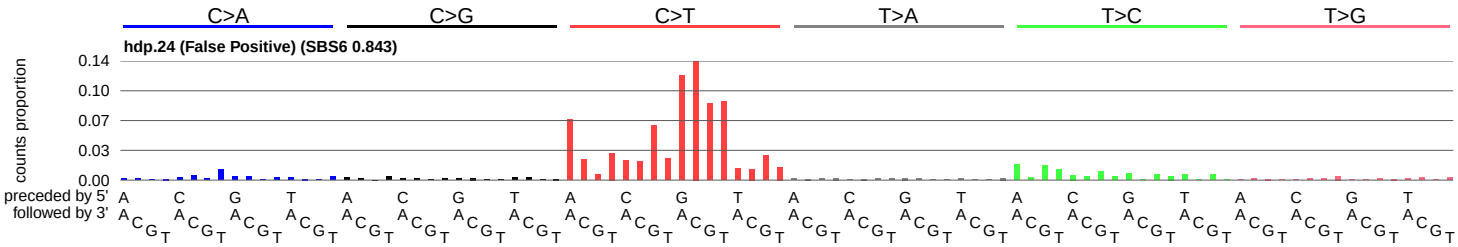
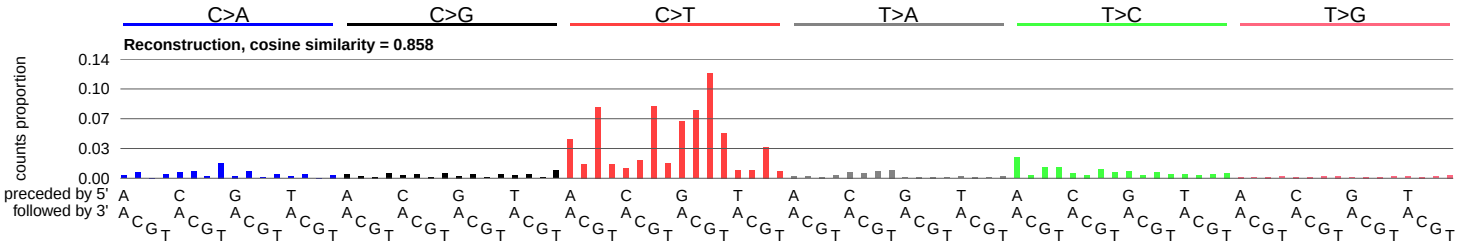
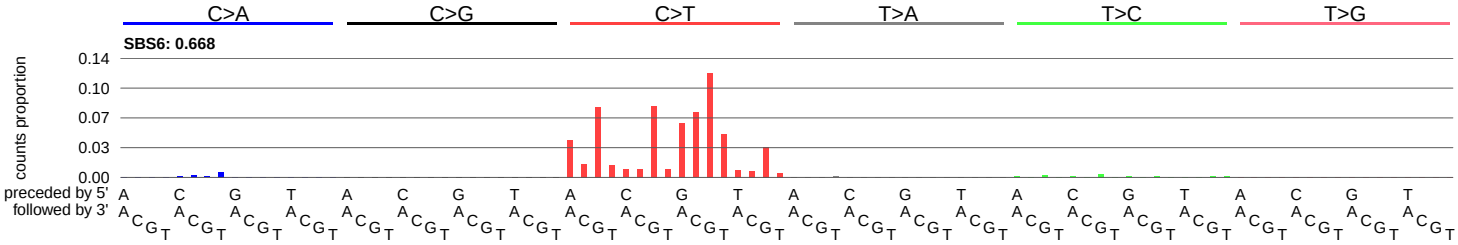
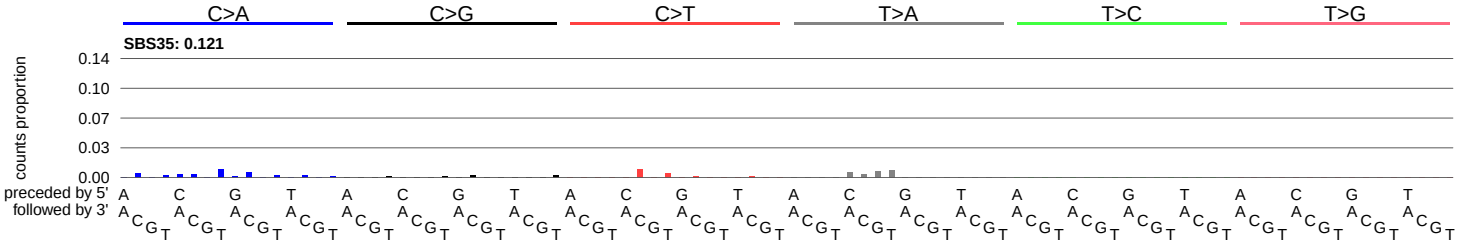
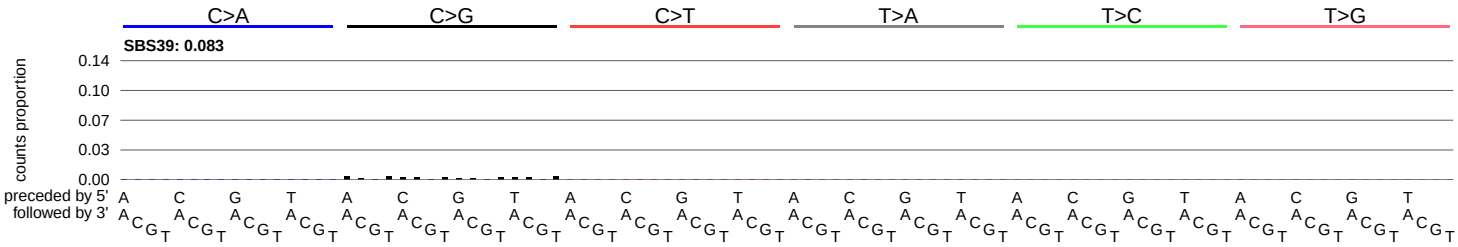
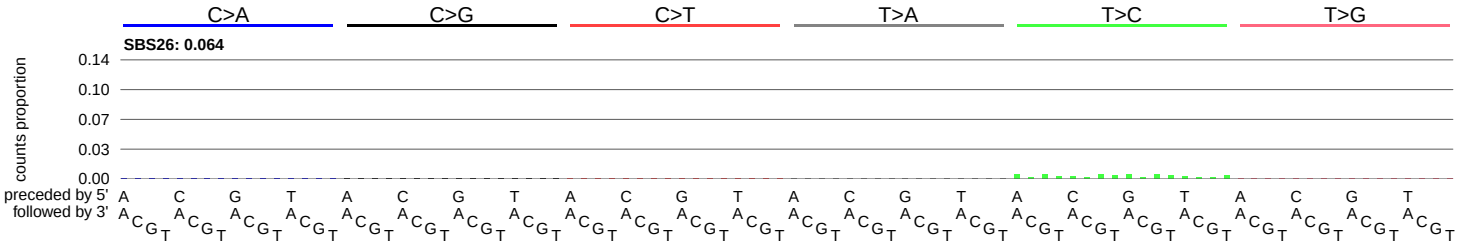

C>A
C>G
C>T
T>A
T>C
T>G
hdp.24 (False Positive) (SBS6 0.843)
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'

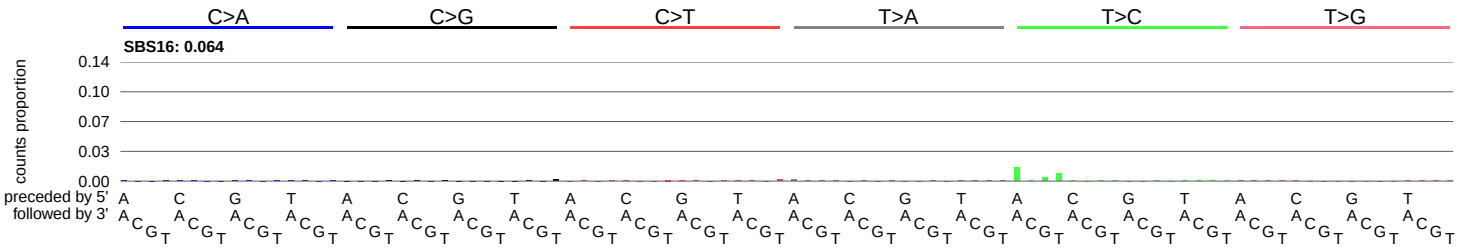

C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.858
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS6: 0.668
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS35: 0.121
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS39: 0.083
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS26: 0.064
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS16: 0.064
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'