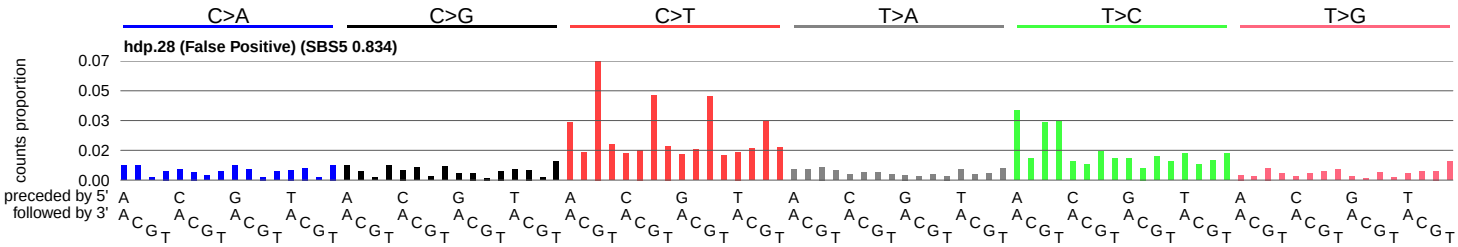
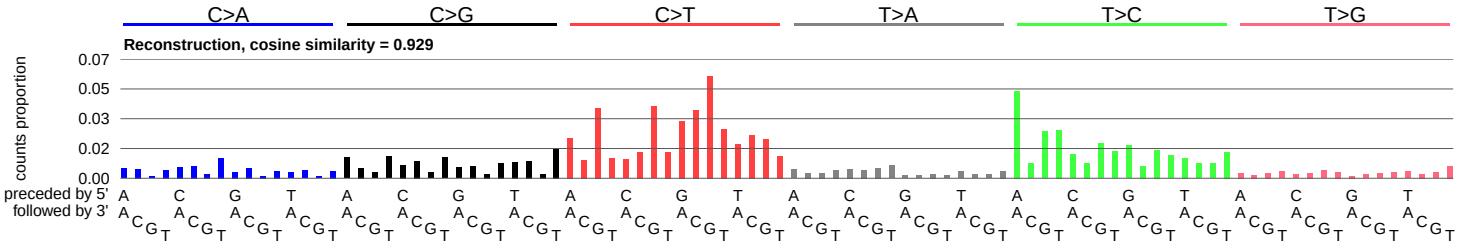
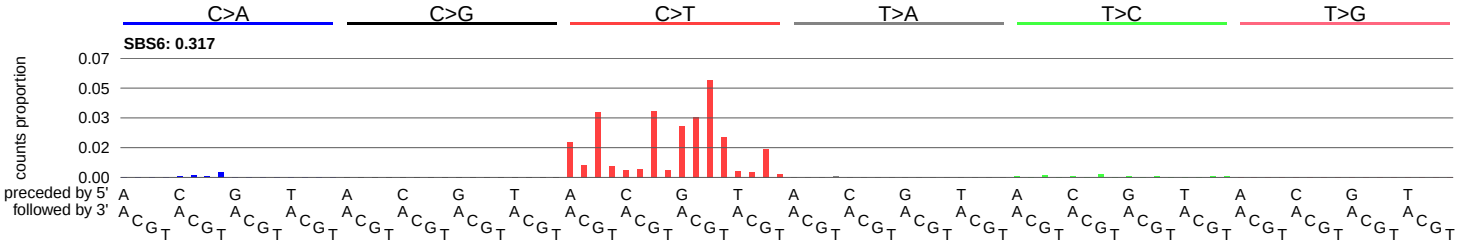
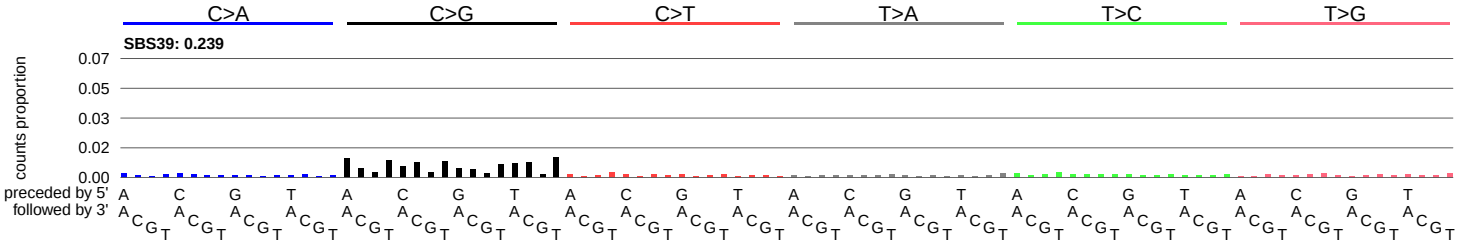
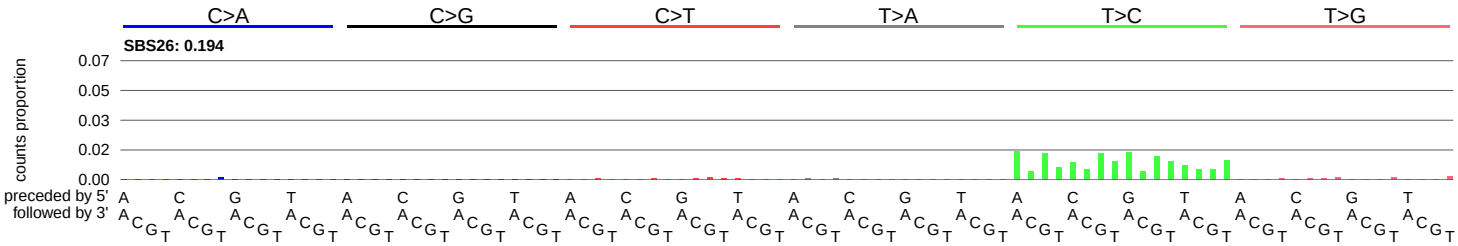
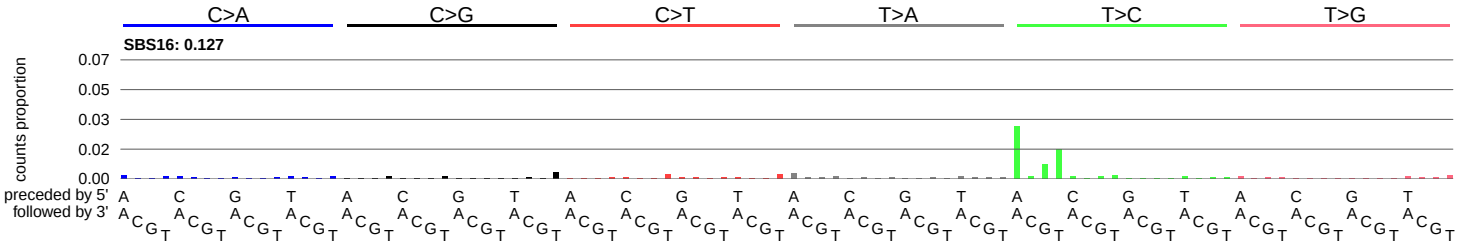

C>A
C>G
C>T
T>A
T>C
T>G
hdp.28 (False Positive) (SBS5 0.834)
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

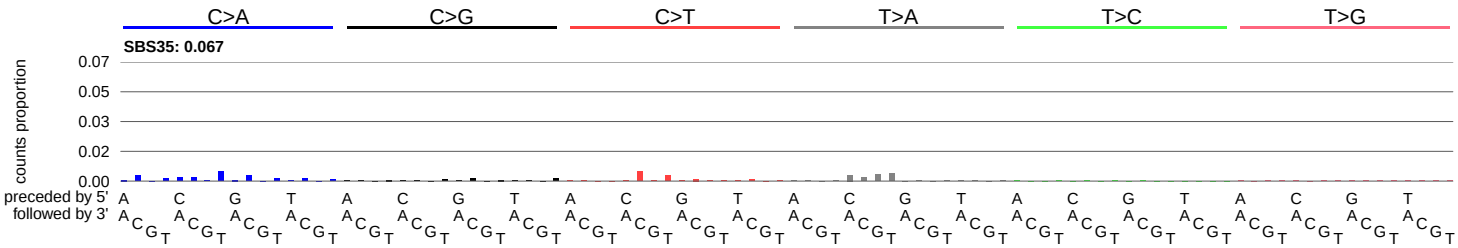

C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.929
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'

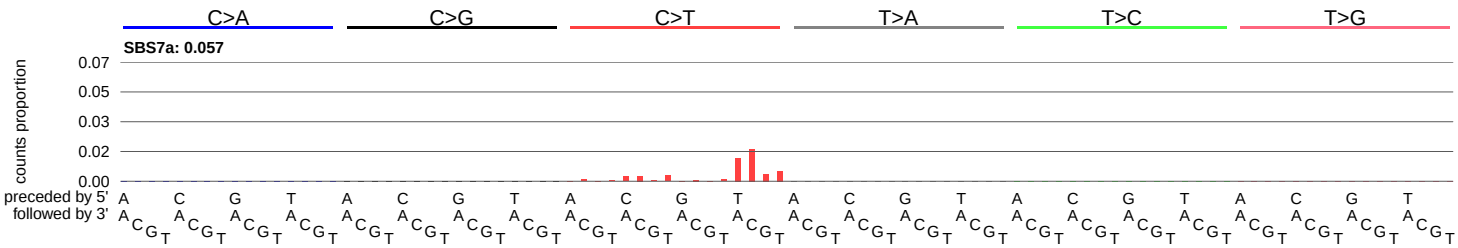

C>A
C>G
C>T
T>A
T>C
T>G
SBS6: 0.317
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS39: 0.239
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
SBS26: 0.194
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: 0.127
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
SBS35: 0.067
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
SBS7a: 0.057
counts proportion
0.07
0.05
0.03
0.02
0.00
preceded by 5'
followed by 3'