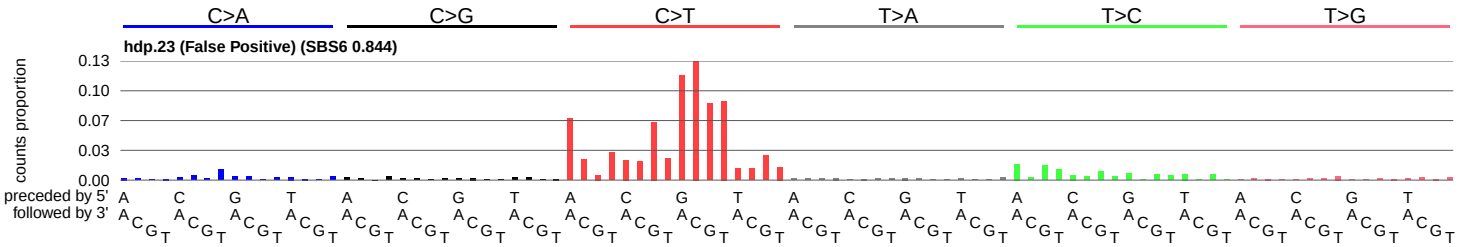
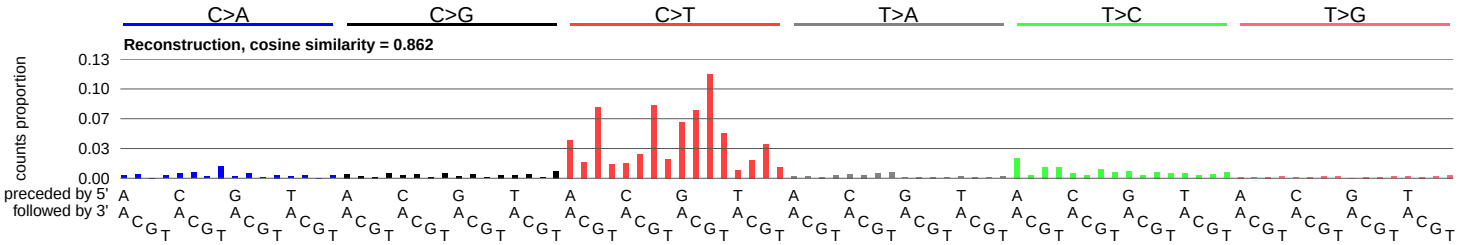
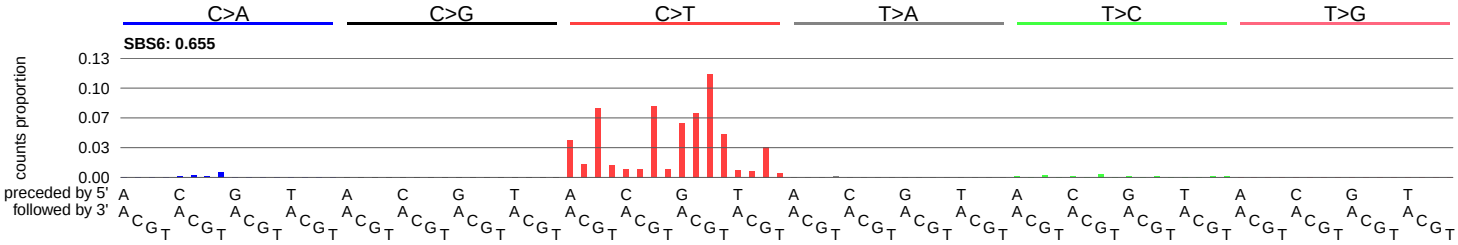
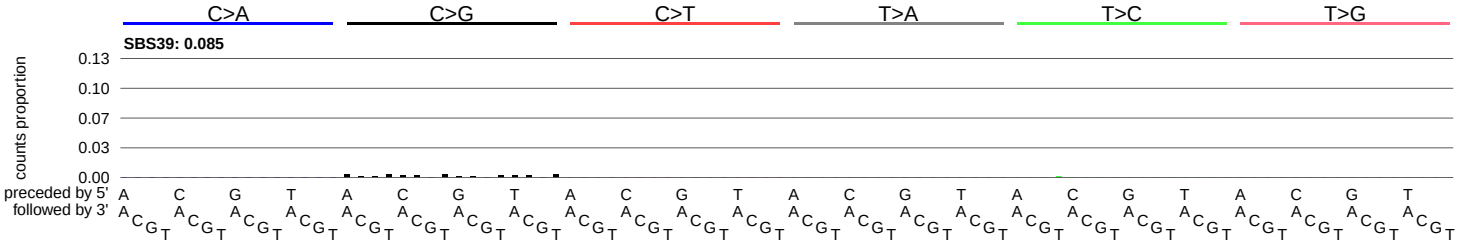
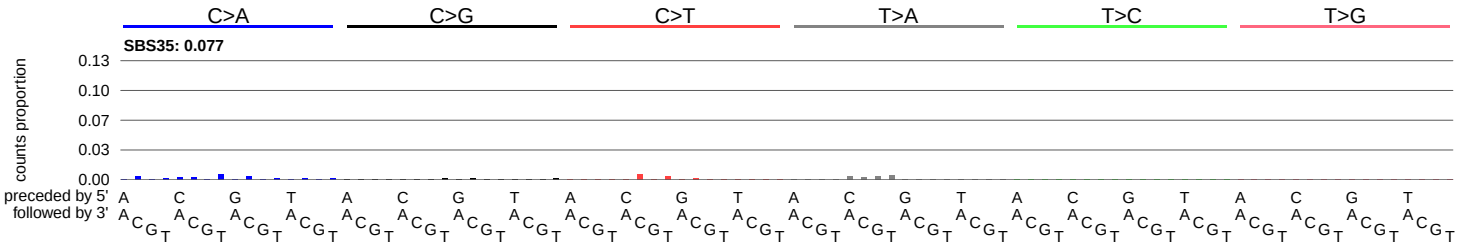
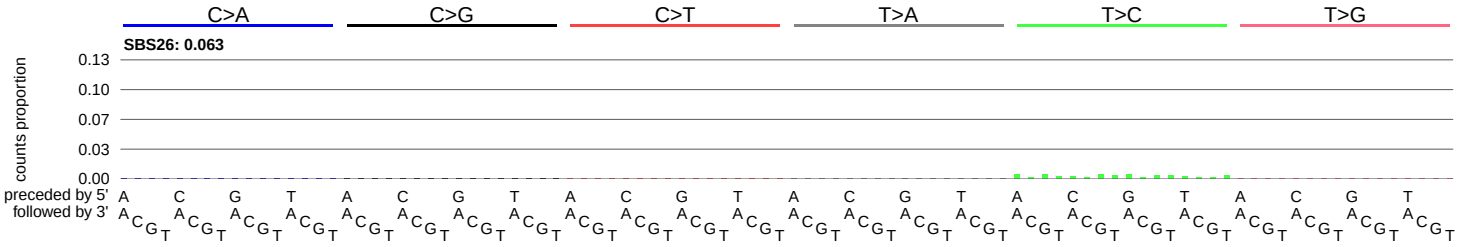

C>A
C>G
C>T
T>A
T>C
T>G
hdp.23 (False Positive) (SBS6 0.844)
counts proportion
0.13
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'

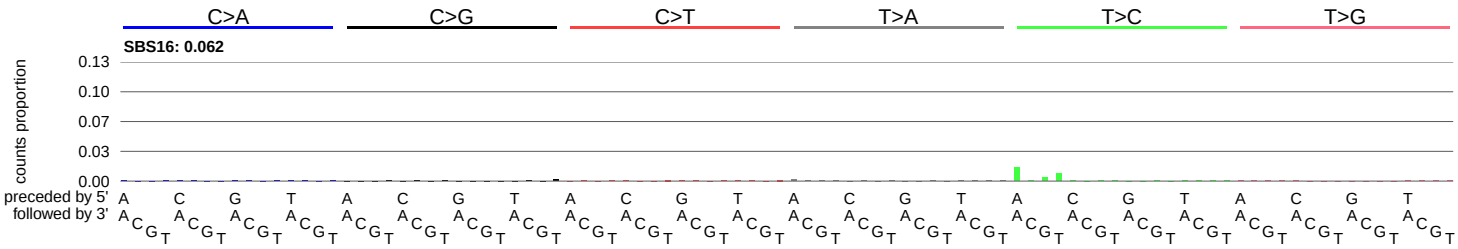

C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.862
counts proportion
0.13
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'

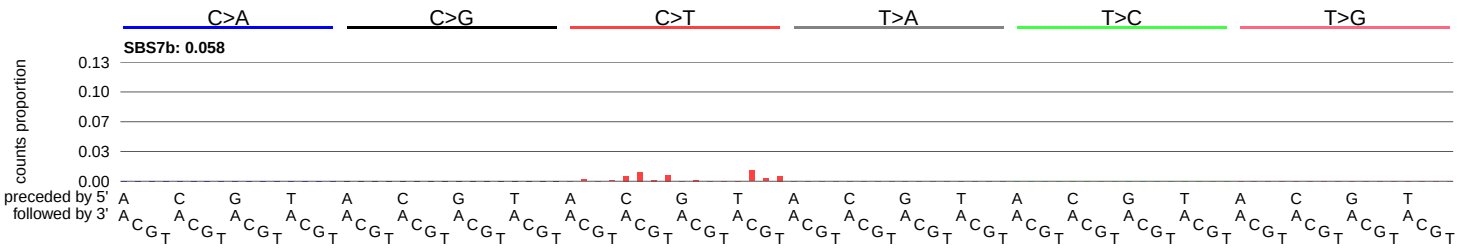

C>A
C>G
C>T
T>A
T>C
T>G
SBS6: 0.655
counts proportion
0.13
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS39: 0.085
counts proportion
0.13
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: 0.077
counts proportion
0.13
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: 0.063
counts proportion
0.13
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: 0.062
counts proportion
0.13
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
SBS7b: 0.058
counts proportion
0.13
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'