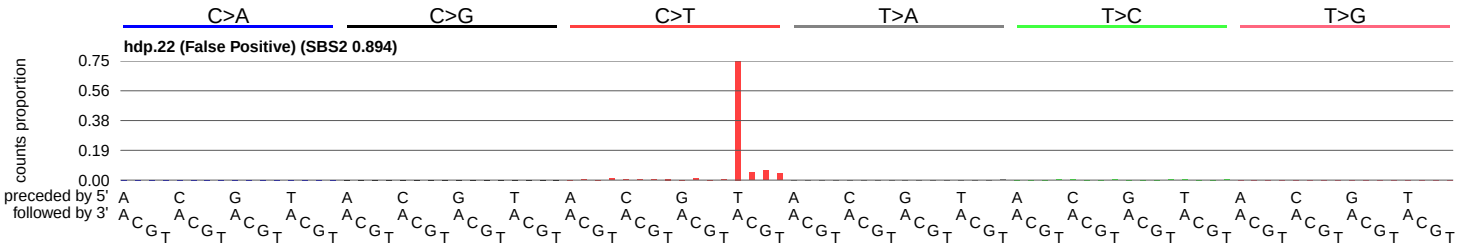
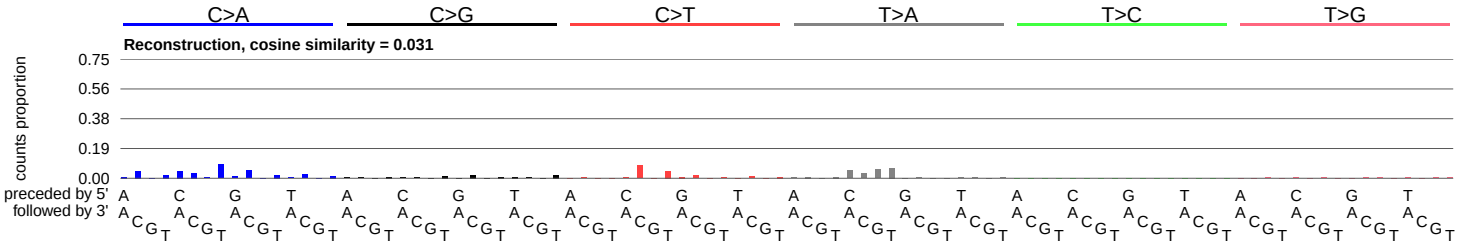
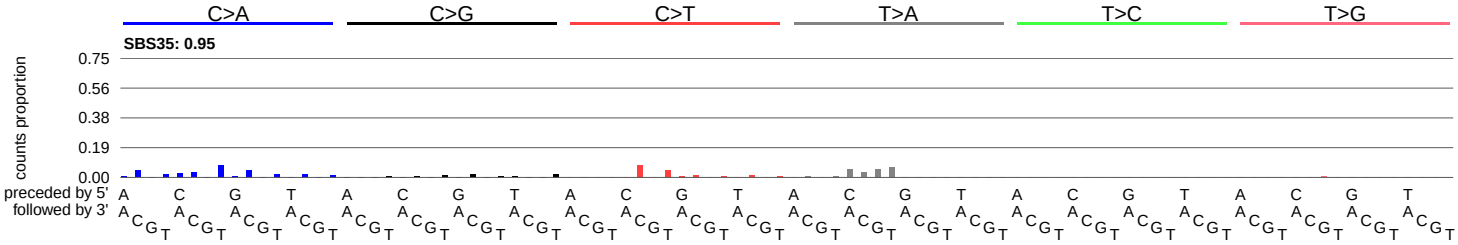
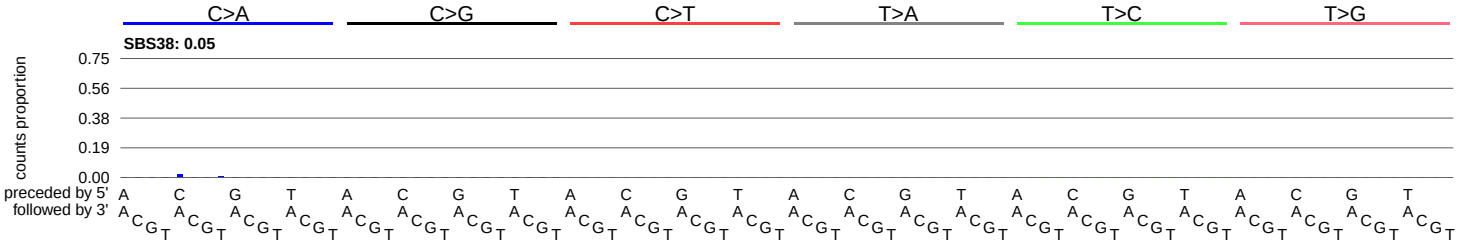

C>A
C>G
C>T
T>A
T>C
T>G
hdp.22 (False Positive) (SBS2 0.894)
counts proportion
0.75
0.56
0.38
0.19
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.031
counts proportion
0.75
0.56
0.38
0.19
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS35: 0.95
counts proportion
0.75
0.56
0.38
0.19
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS38: 0.05
counts proportion
0.75
0.56
0.38
0.19
0.00
preceded by 5'
followed by 3'