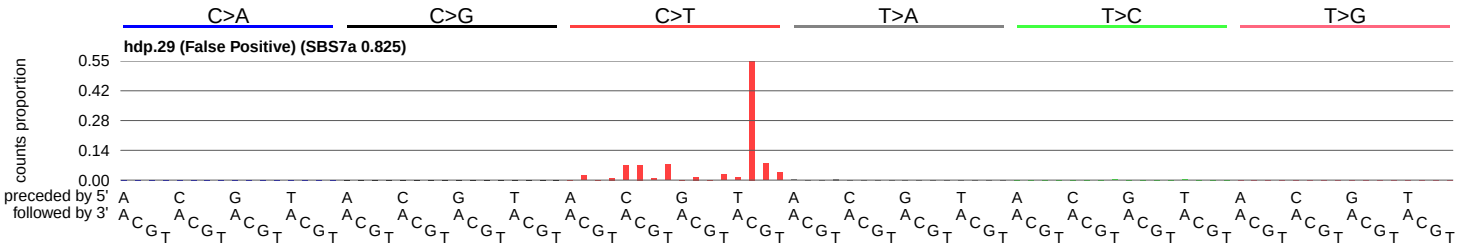
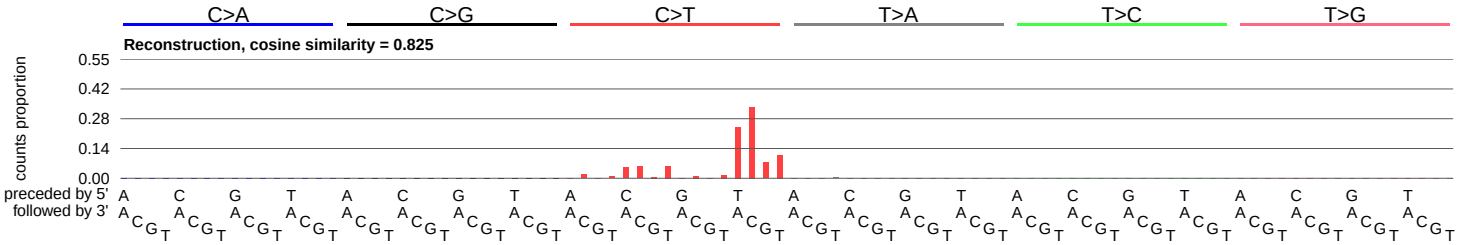
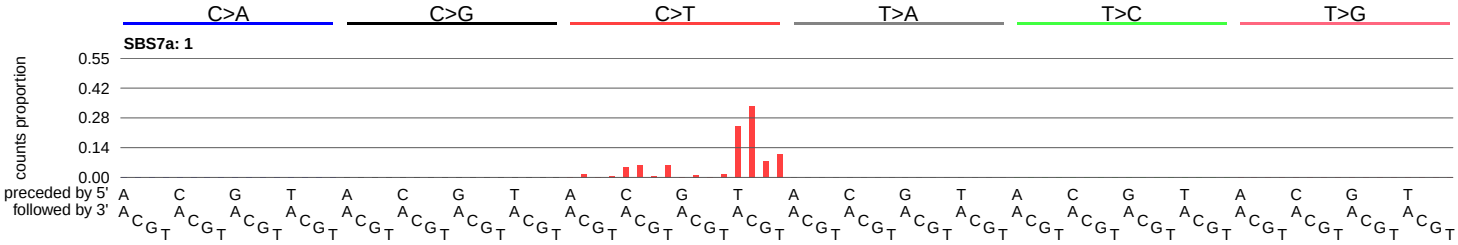

C>A
C>G
C>T
T>A
T>C
T>G
hdp.29 (False Positive) (SBS7a 0.825)
counts proportion
0.55
0.42
0.28
0.14
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
Reconstruction, cosine similarity = 0.825
counts proportion
0.55
0.42
0.28
0.14
0.00
preceded by 5'
followed by 3'


C>A
C>G
C>T
T>A
T>C
T>G
SBS7a: 1
counts proportion
0.55
0.42
0.28
0.14
0.00
preceded by 5'
followed by 3'