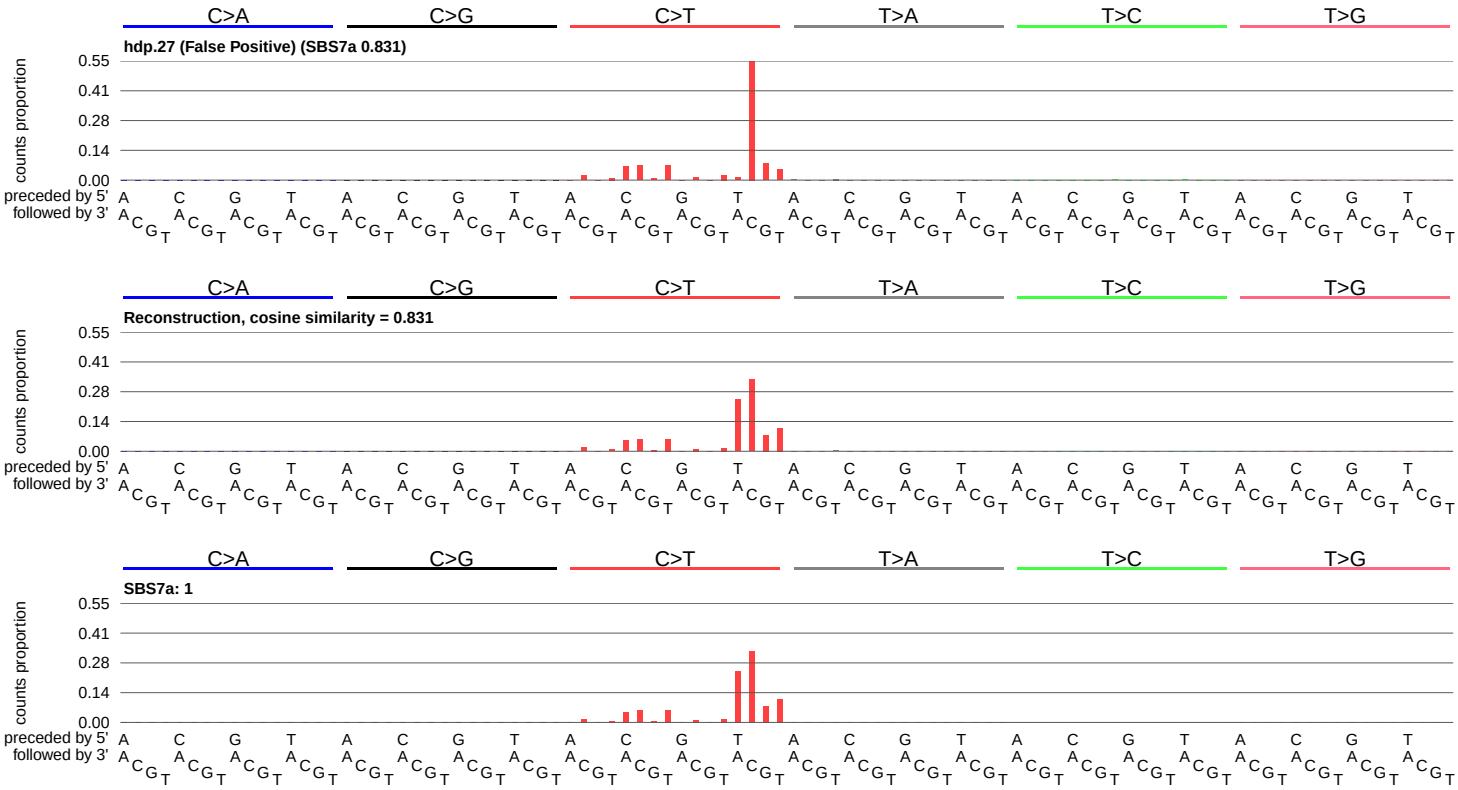
C>A
C>G
C>T
T>A
T>C
T>G
hdp.27 (False Positive) (SBS7a 0.831)
counts proportion
0.55
0.41
0.28
0.14
0.00
preceded by 5'
followed by 3'
Reconstruction, cosine similarity = 0.831
SBS7a: 1