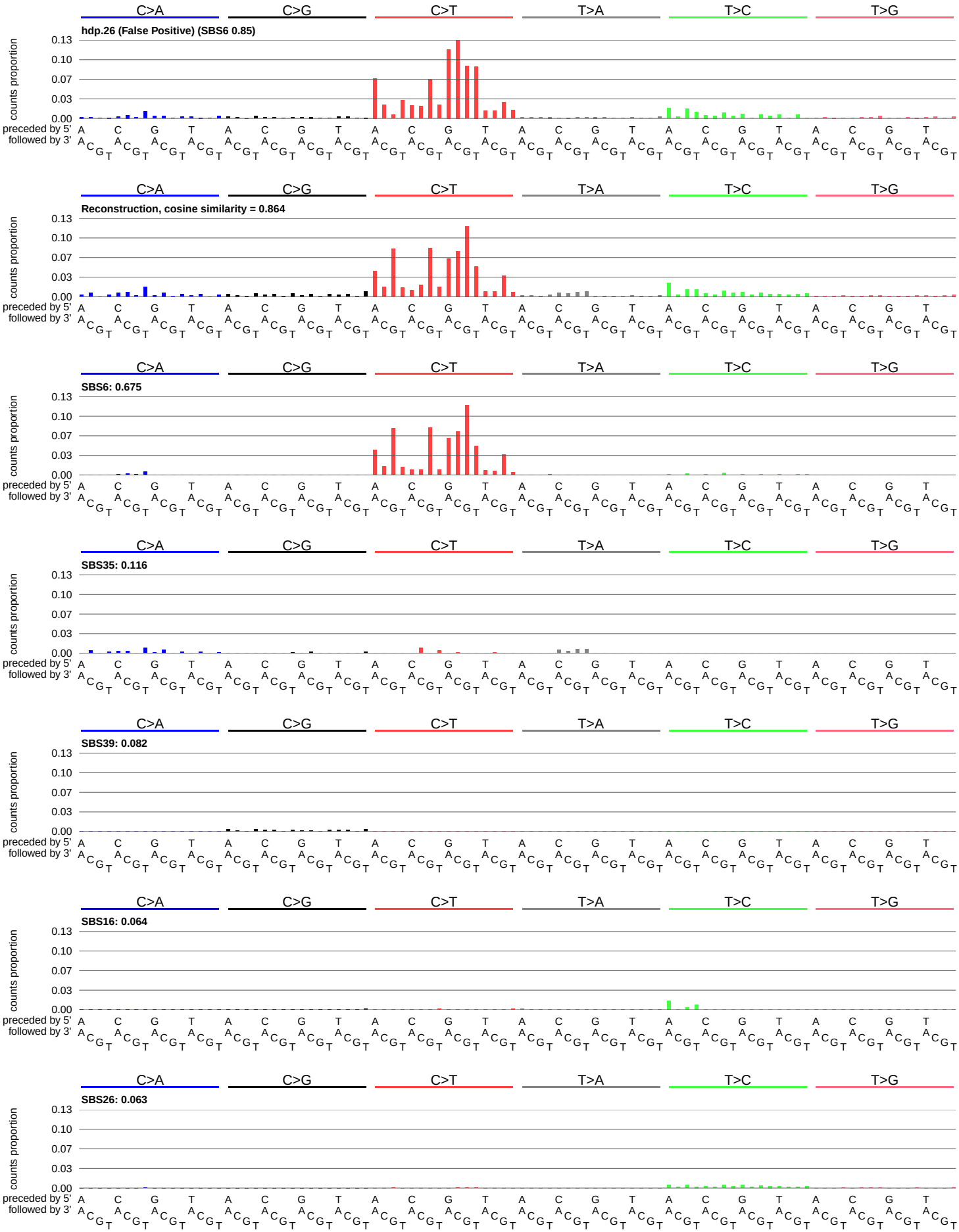
C>A
C>G
C>T
T>A
T>C
T>G
hdp.26 (False Positive) (SBS6 0.85)
Reconstruction, cosine similarity = 0.864
SBS6: 0.675
SBS35: 0.116
SBS39: 0.082
SBS16: 0.064
SBS26: 0.063
counts proportion
0.13
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'