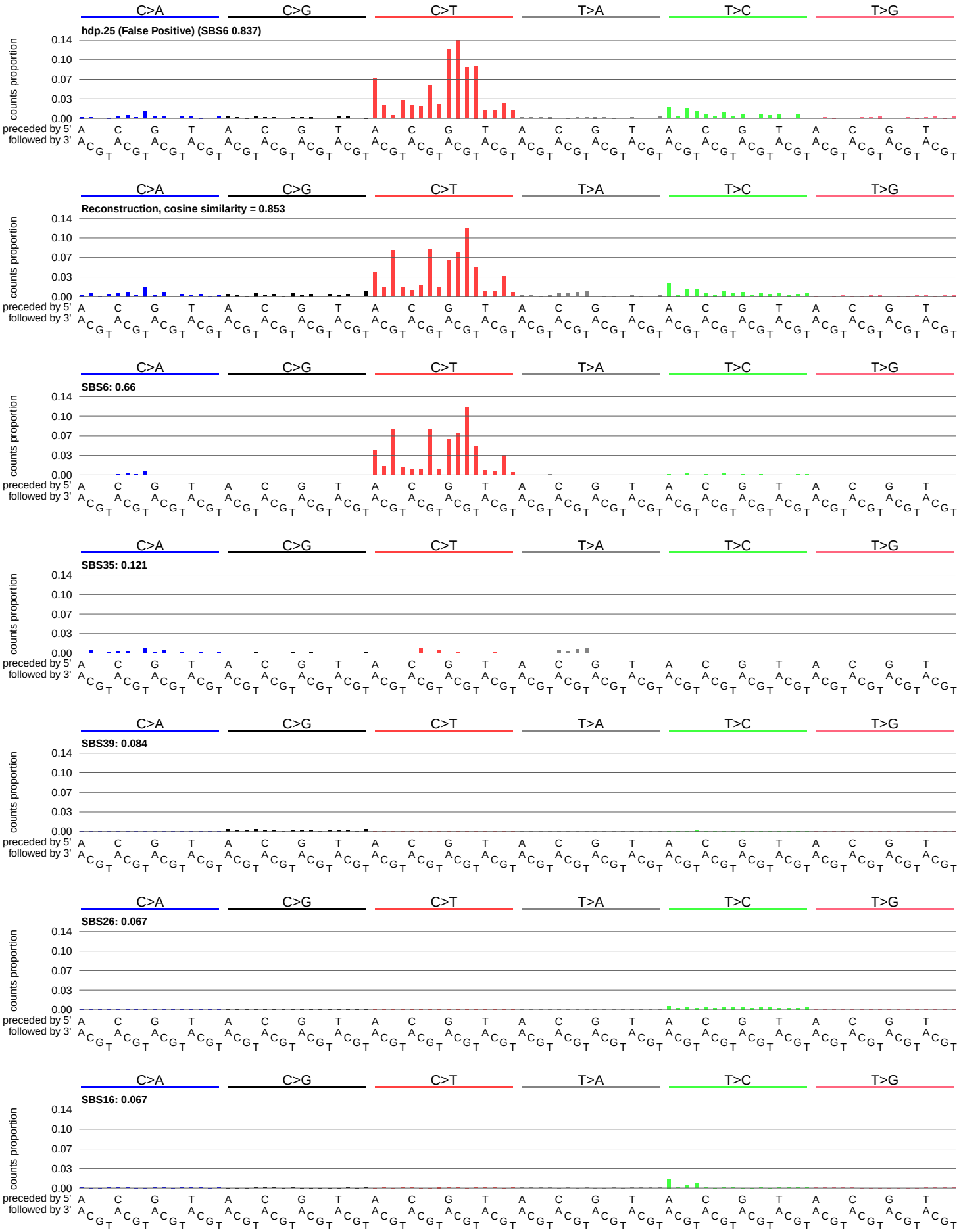
C>A
C>G
C>T
T>A
T>C
T>G
hdp.25 (False Positive) (SBS6 0.837)
Reconstruction, cosine similarity = 0.853
SBS6: 0.66
SBS35: 0.121
SBS39: 0.084
SBS26: 0.067
SBS16: 0.067
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'