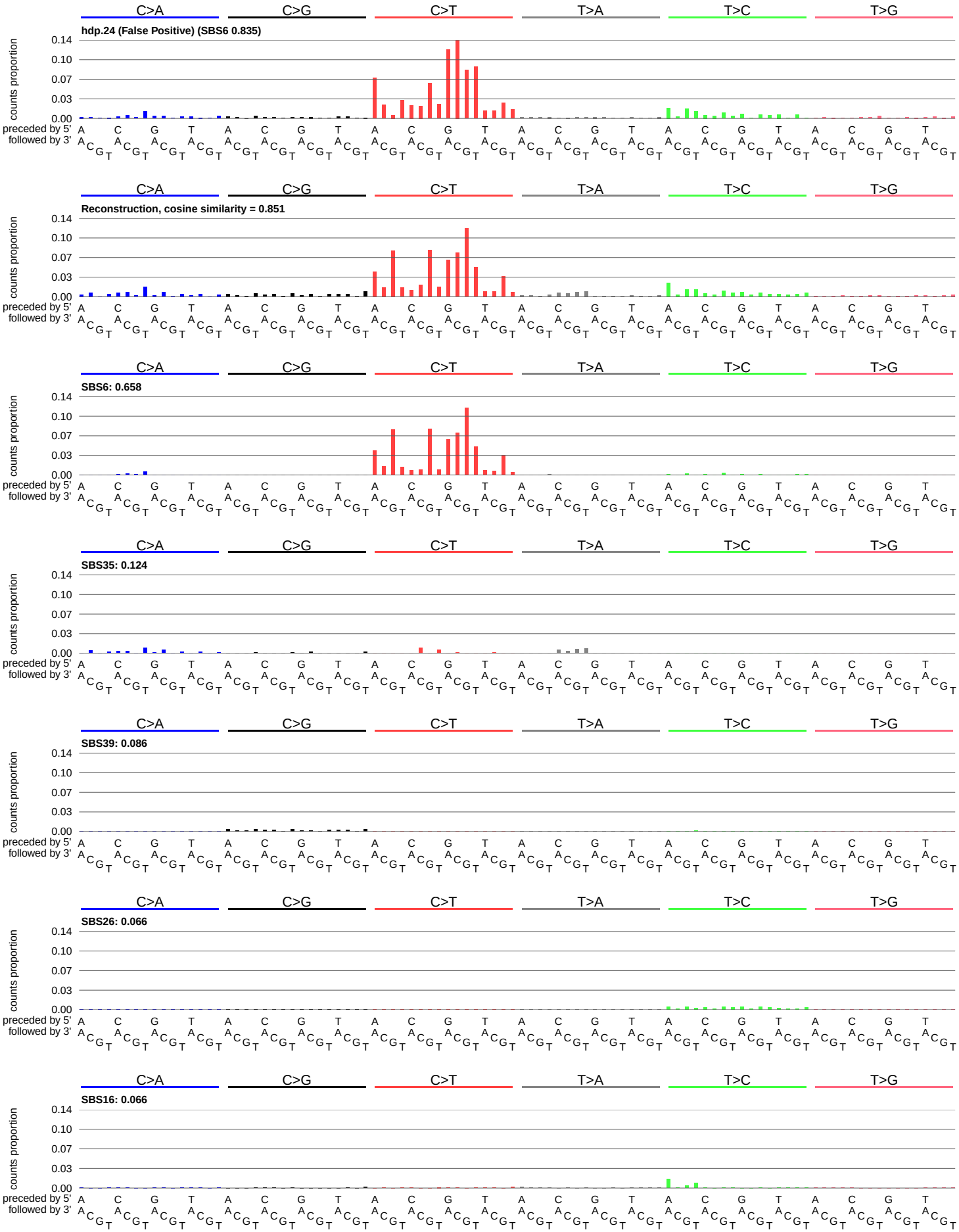
C>A
C>G
C>T
T>A
T>C
T>G
hdp.24 (False Positive) (SBS6 0.835)
Reconstruction, cosine similarity = 0.851
SBS6: 0.658
SBS35: 0.124
SBS39: 0.086
SBS26: 0.066
SBS16: 0.066
counts proportion
0.14
0.10
0.07
0.03
0.00
preceded by 5'
followed by 3'