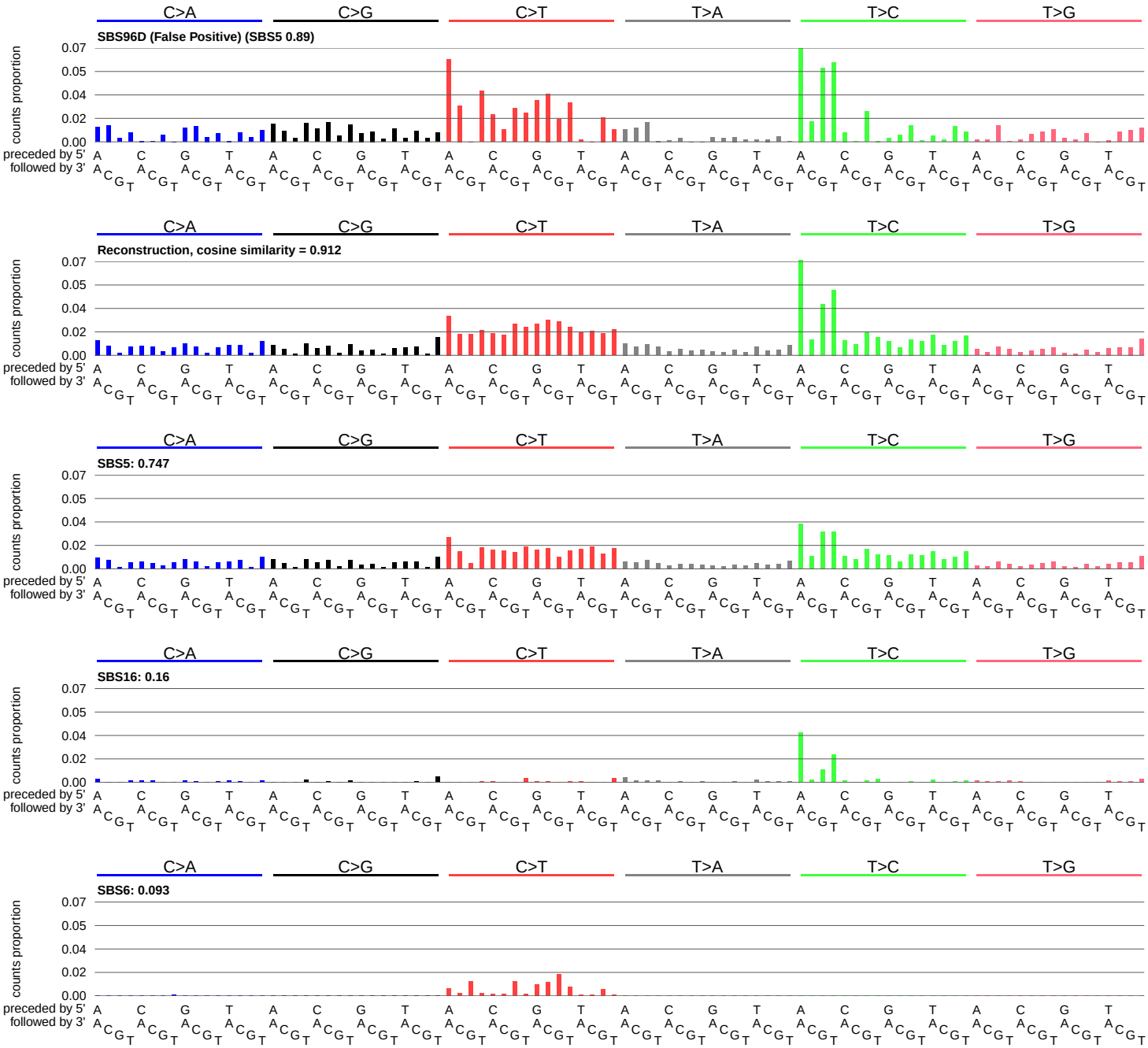
C>A
C>G
C>T
T>A
T>C
T>G
SBS96D (False Positive) (SBS5 0.89)
Reconstruction, cosine similarity = 0.912
SBS5: 0.747
SBS16: 0.16
SBS6: 0.093
counts proportion
0.07
0.05
0.04
0.02
0.00
preceded by 5'
followed by 3'