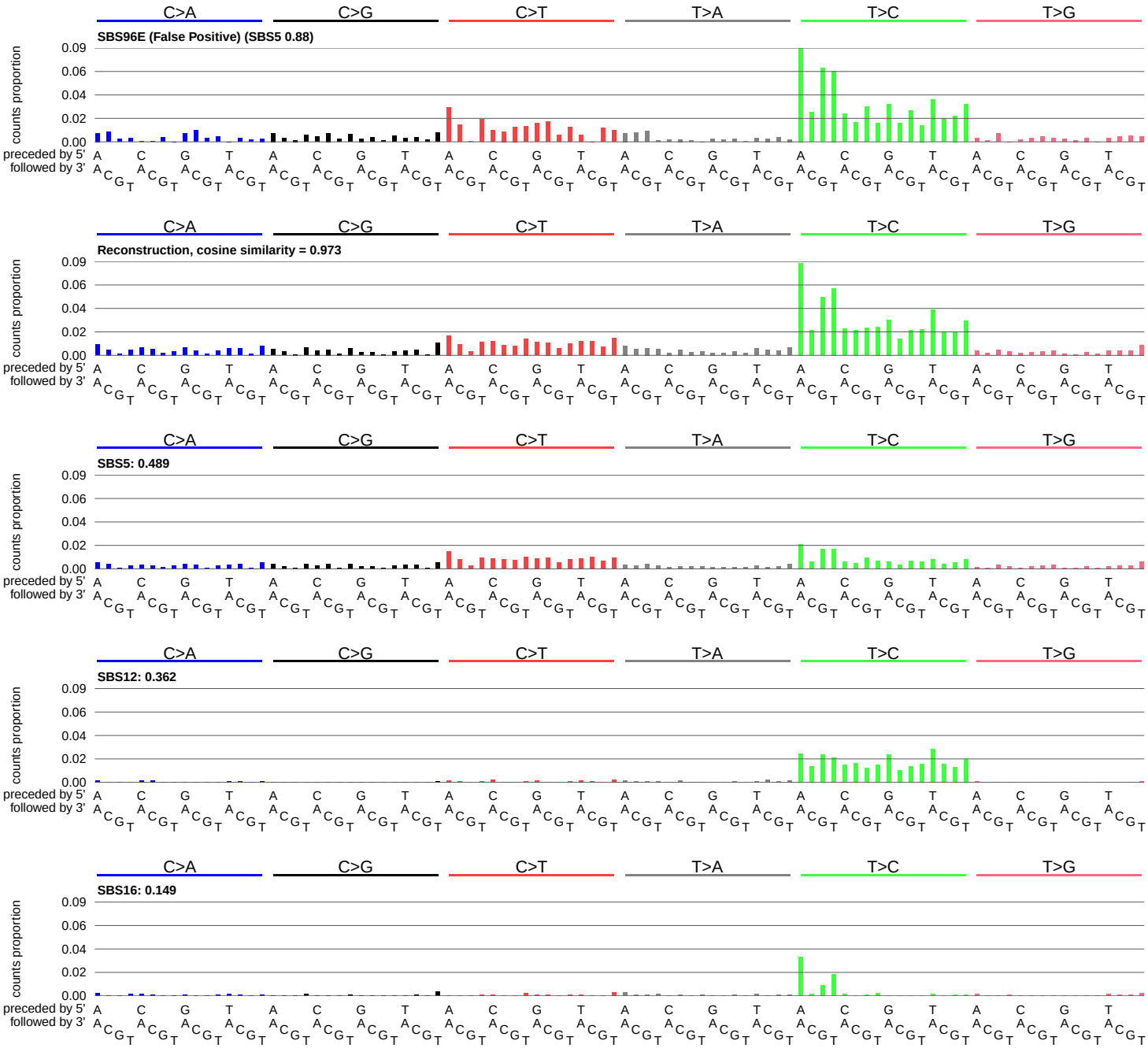
C>A
C>G
C>T
T>A
T>C
T>G
SBS96E (False Positive) (SBS5 0.88)
Reconstruction, cosine similarity = 0.973
SBS5: 0.489
SBS12: 0.362
SBS16: 0.149
counts proportion
0.09
0.06
0.04
0.02
0.00
preceded by 5'
followed by 3'