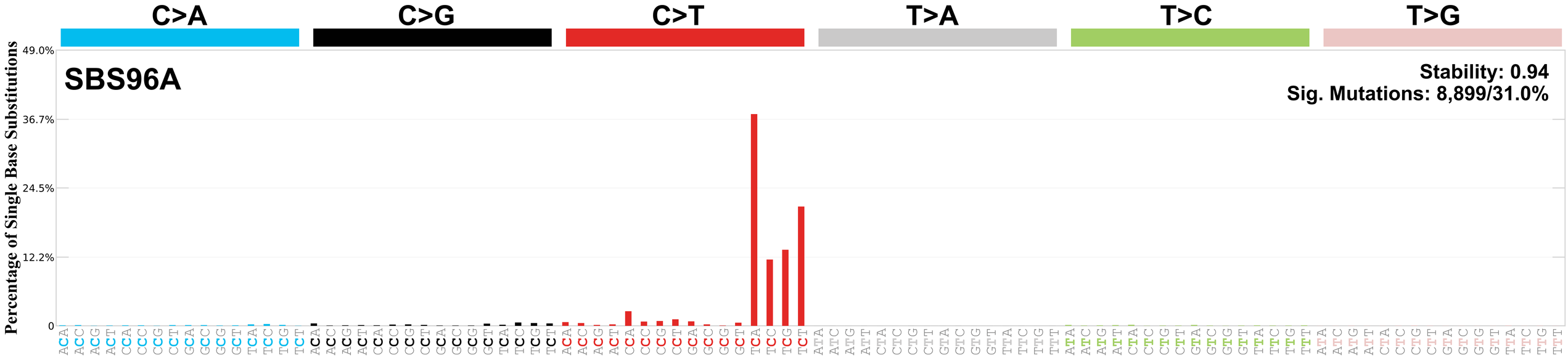
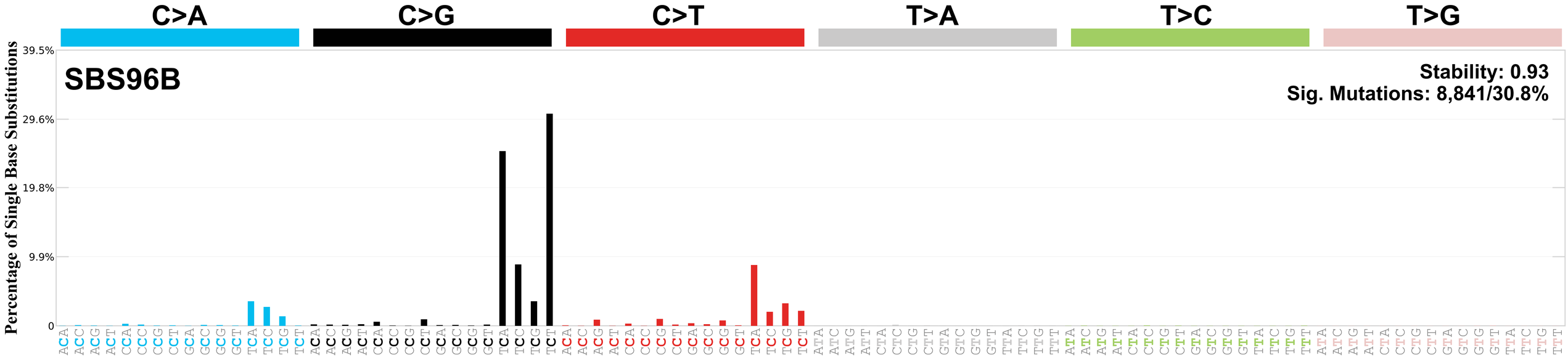
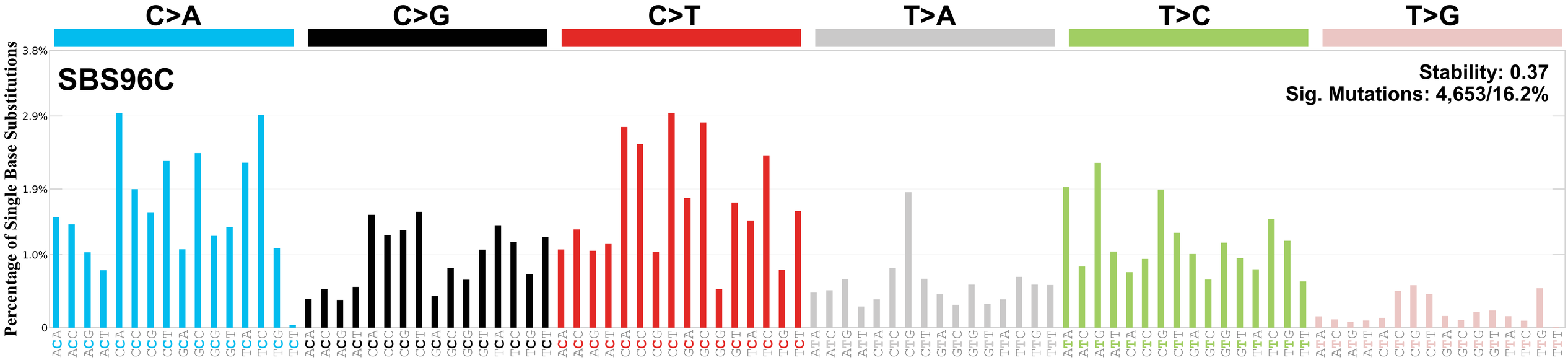
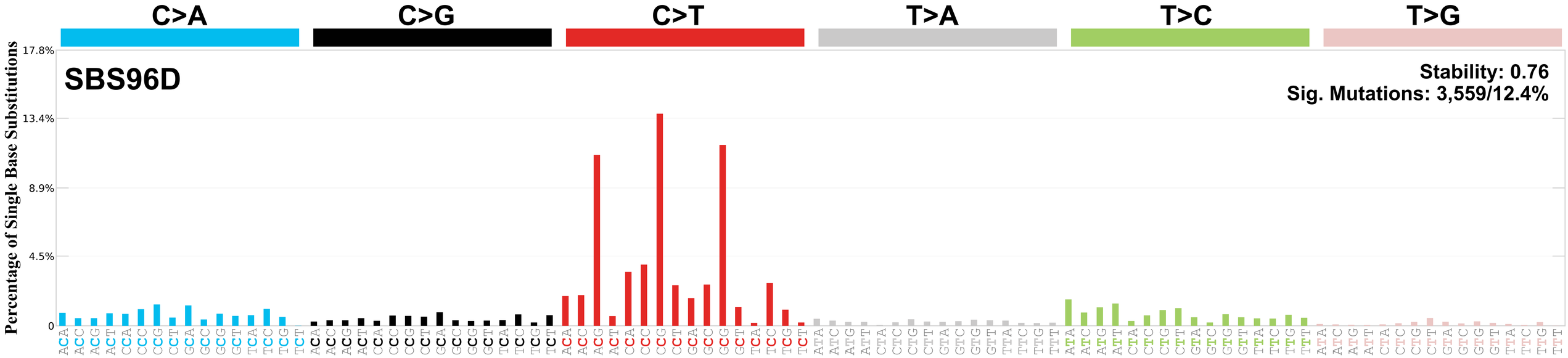
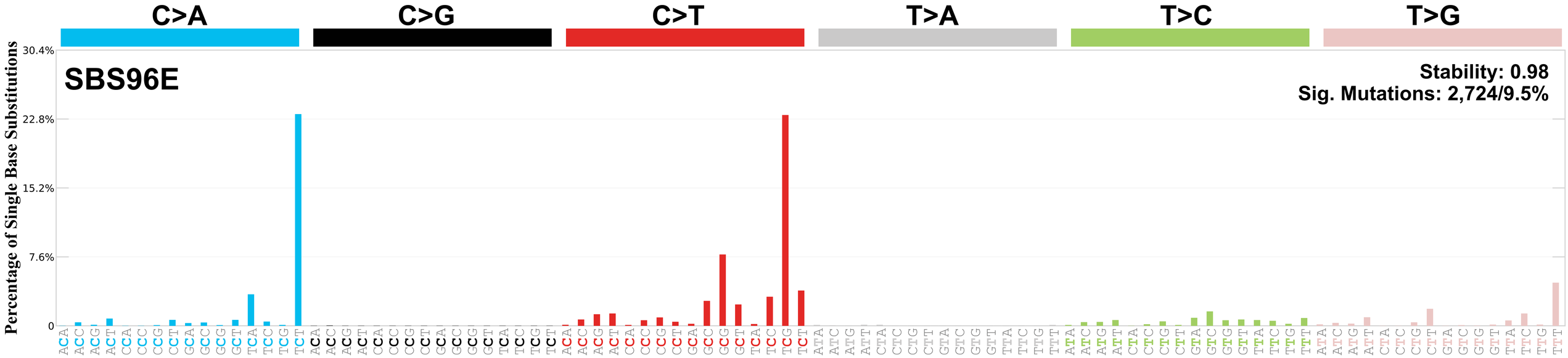

C>A
C>G
C>T
T>A
T>C
T>G
SBS96A
Stability: 0.94
Sig. Mutations: 8,899/31.0%
Percentage of Single Base Substitutions
49.0%
36.7%
24.5%
12.2%
0


C>A
C>G
C>T
T>A
T>C
T>G
SBS96B
Stability: 0.93
Sig. Mutations: 8,841/30.8%
Percentage of Single Base Substitutions
39.5%
29.6%
19.8%
9.9%
0


C>A
C>G
C>T
T>A
T>C
T>G
SBS96C
Stability: 0.37
Sig. Mutations: 4,653/16.2%
Percentage of Single Base Substitutions
3.8%
2.9%
1.9%
1.0%
0
ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GCG GCT TCA TCC TCG TCT
ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GCG GCT TCA TCC TCG TCT
ACA ACC ACG ACT CCA CCC CCG CCT GCA GCC GCG GCT TCA TCC TCG TCT
ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT
ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT
ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT
ATA ATC ATG ATT CTA CTC CTG CTT GTA GTC GTG GTT TTA TTC TTG TTT


C>A
C>G
C>T
T>A
T>C
T>G
SBS96D
Stability: 0.76
Sig. Mutations: 3,559/12.4%
Percentage of Single Base Substitutions
17.8%
13.4%
8.9%
4.5%
0


C>A
C>G
C>T
T>A
T>C
T>G
SBS96E
Stability: 0.98
Sig. Mutations: 2,724/9.5%
Percentage of Single Base Substitutions
30.4%
22.8%
15.2%
7.6%
0