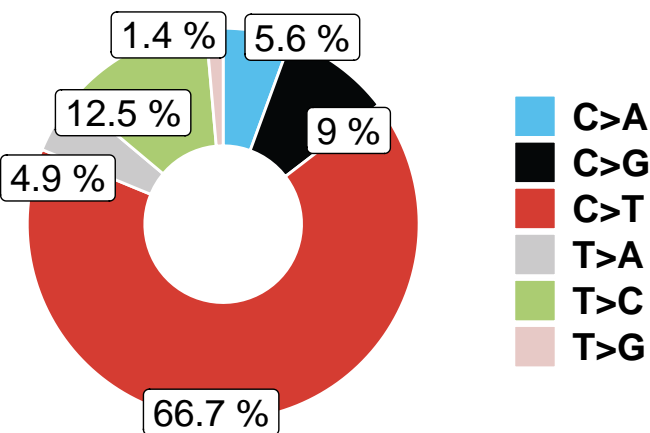
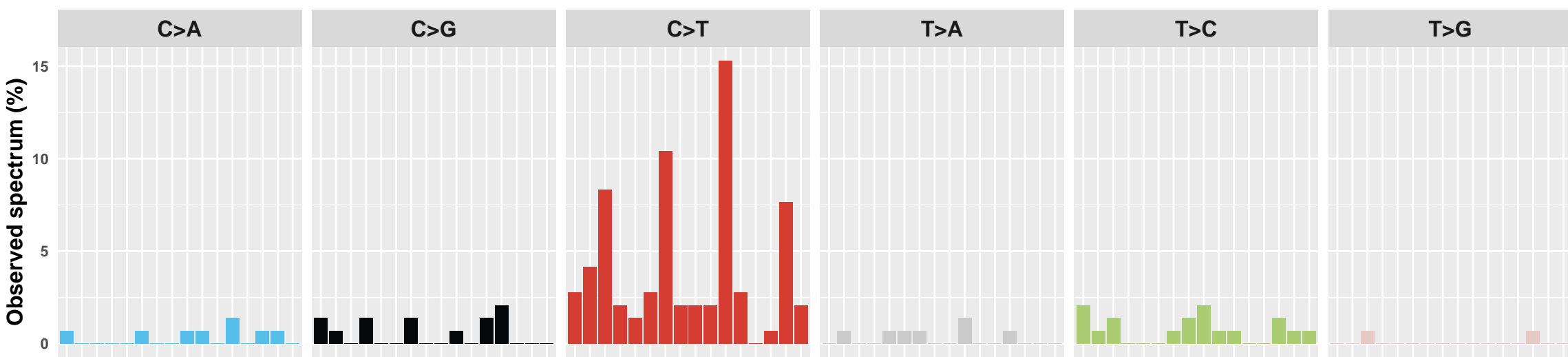


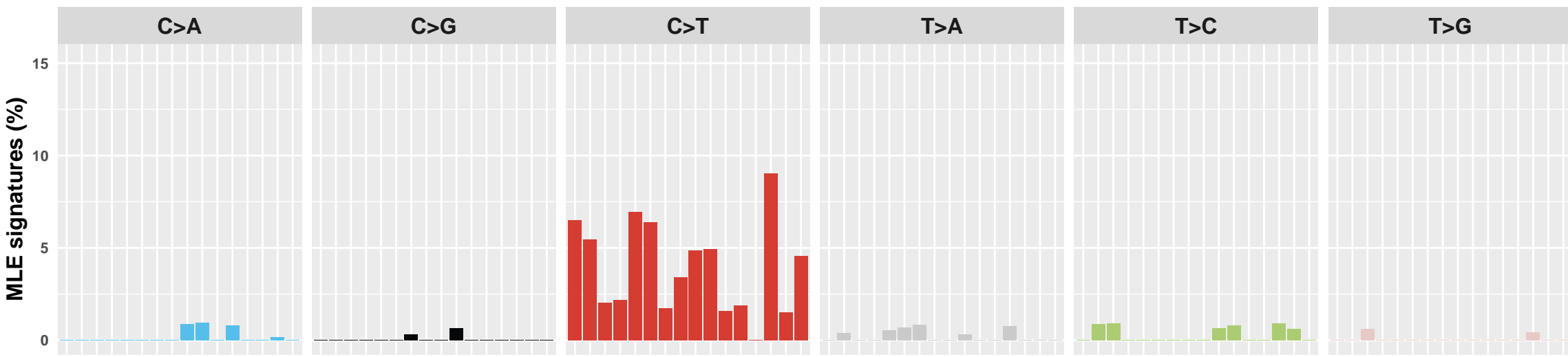
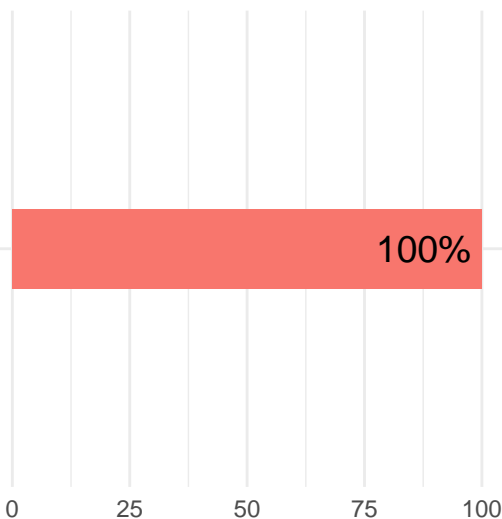
Total mutations: 144



C>A
C>G
C>T
T>A
T>C
T>G



Signature 1



Cosine Similarity: 0.451

