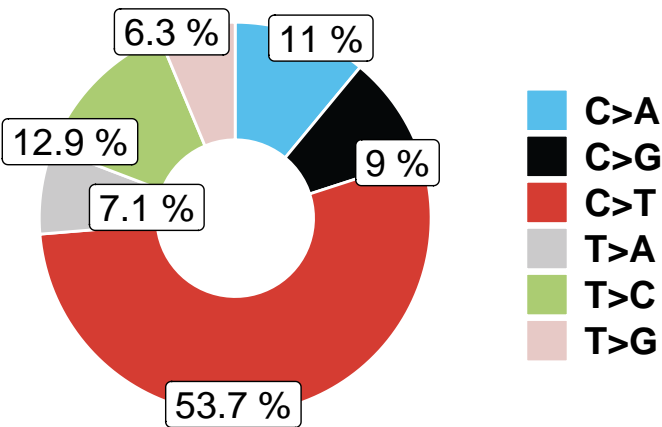
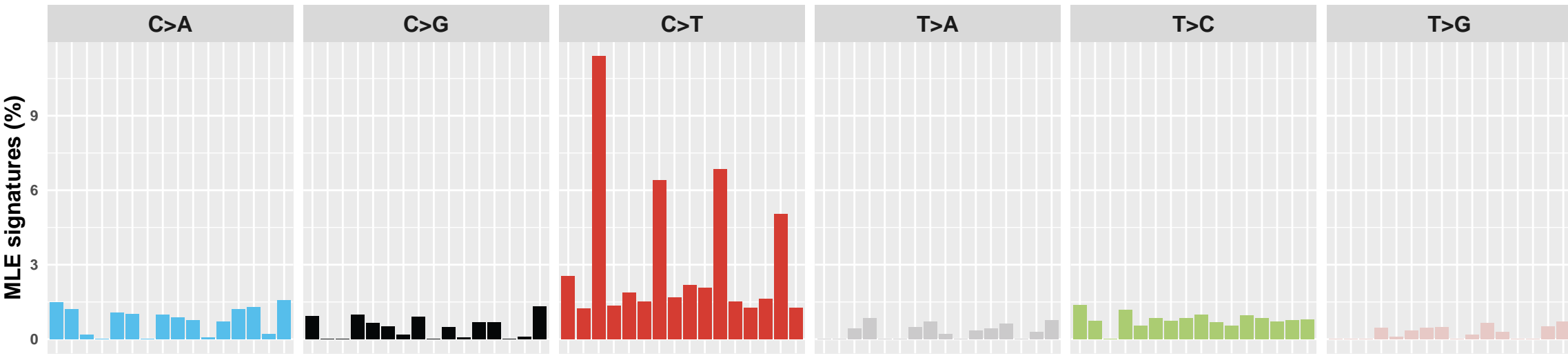
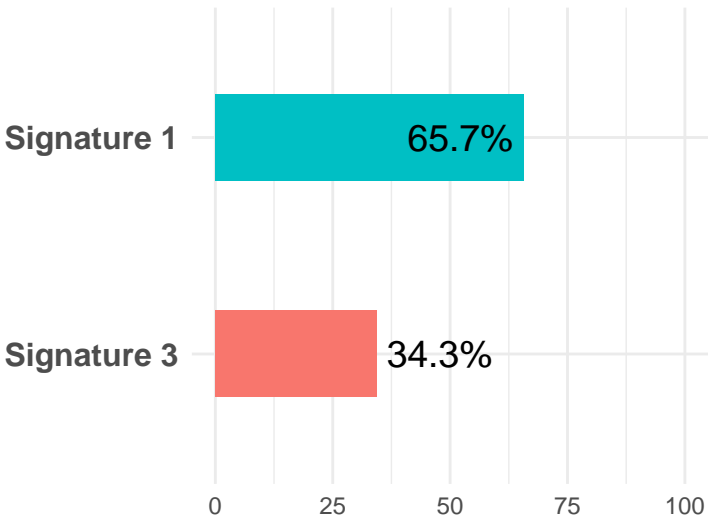
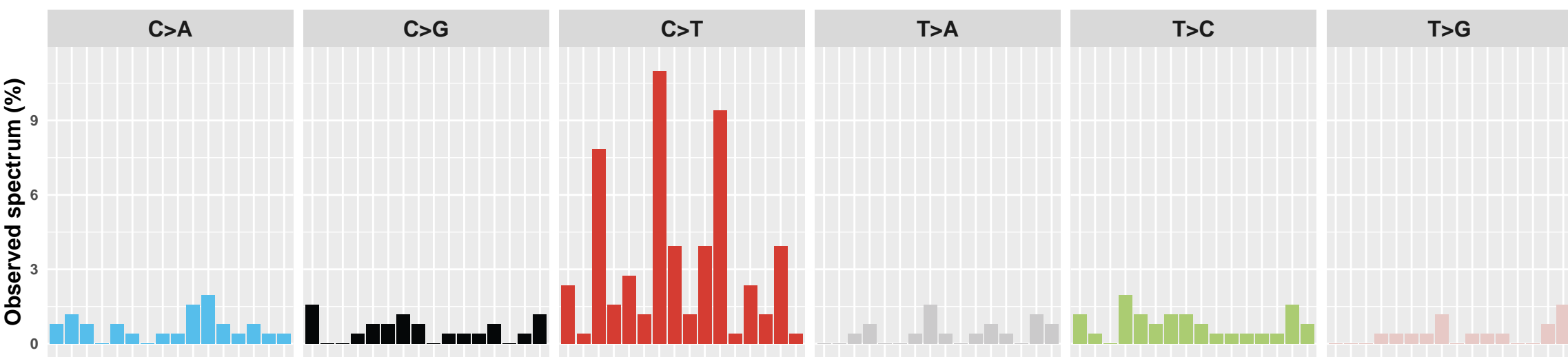


Total mutations: 255



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



Cosine Similarity: 0.901

