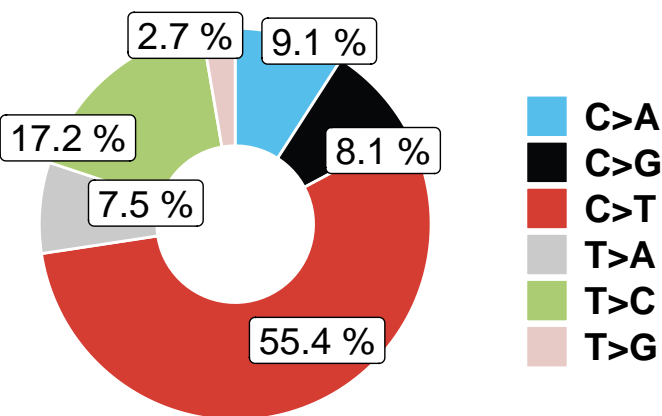
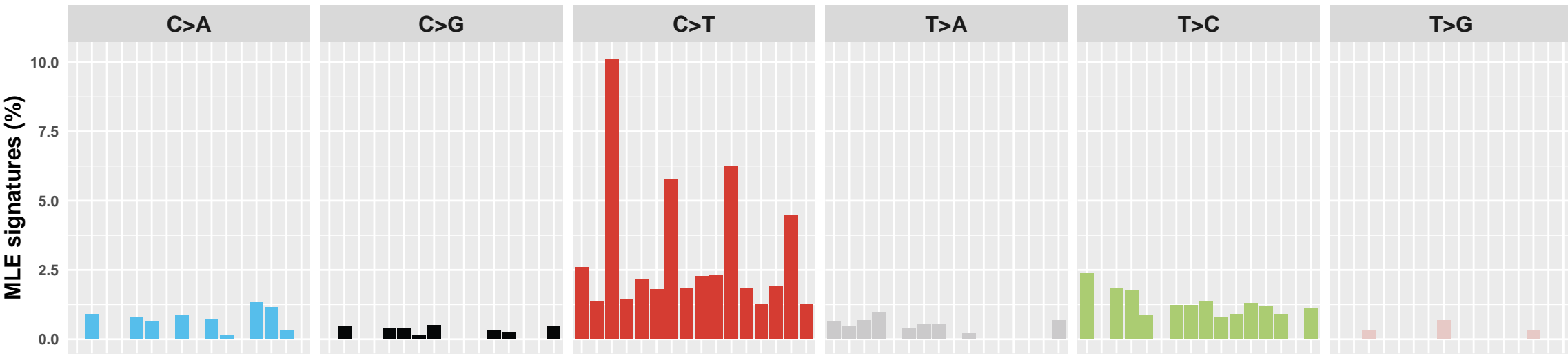
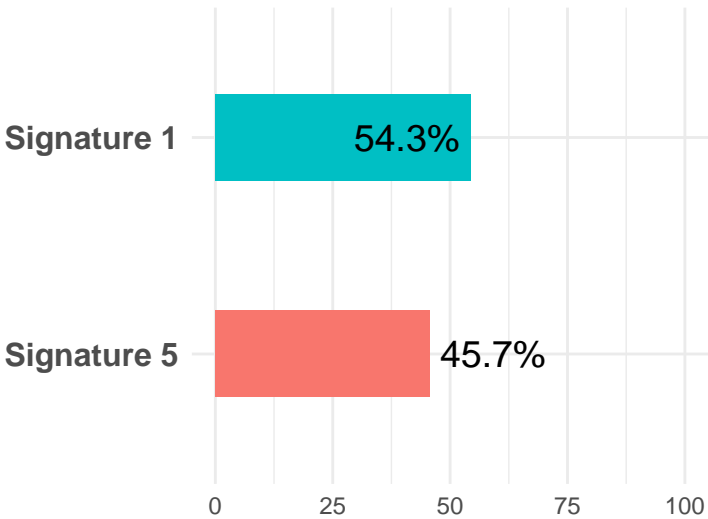
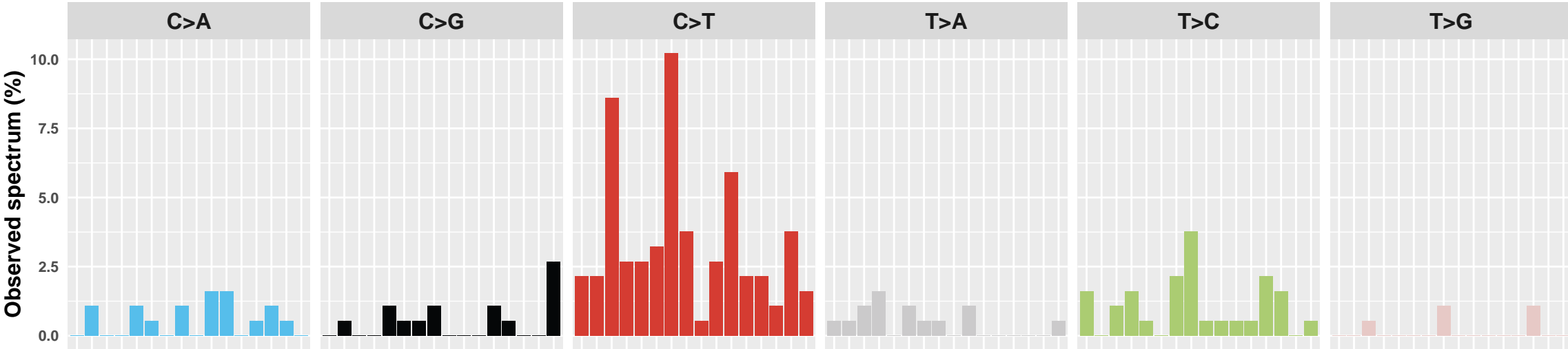


Total mutations: 186



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



Cosine Similarity: 0.915

