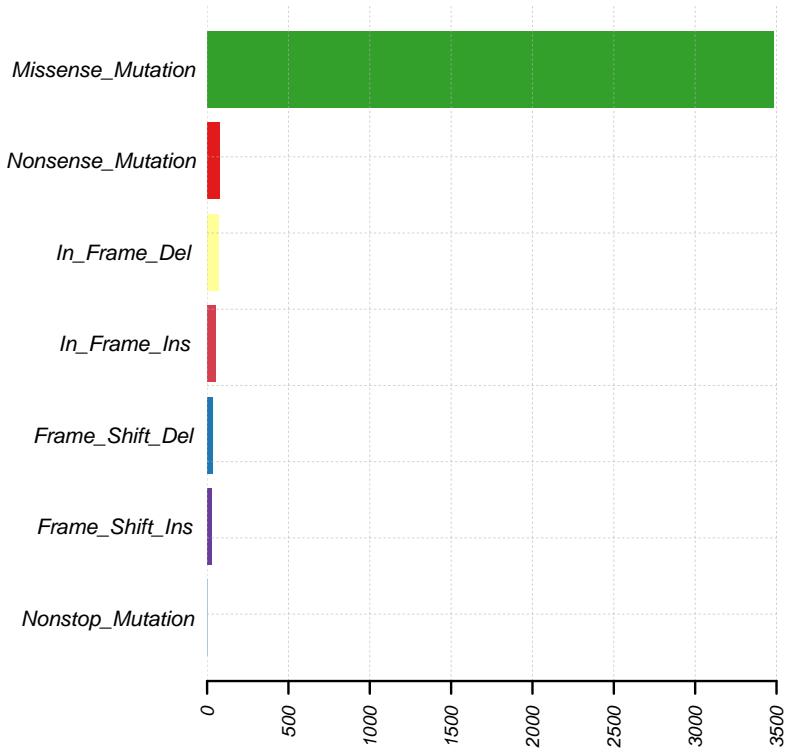
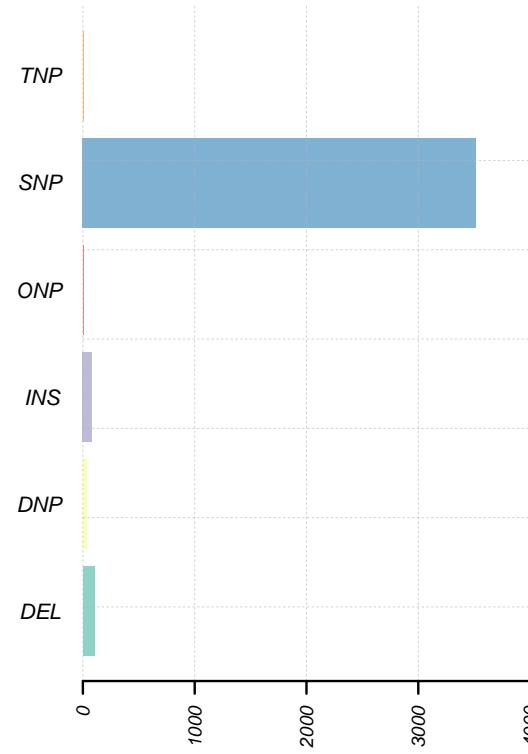
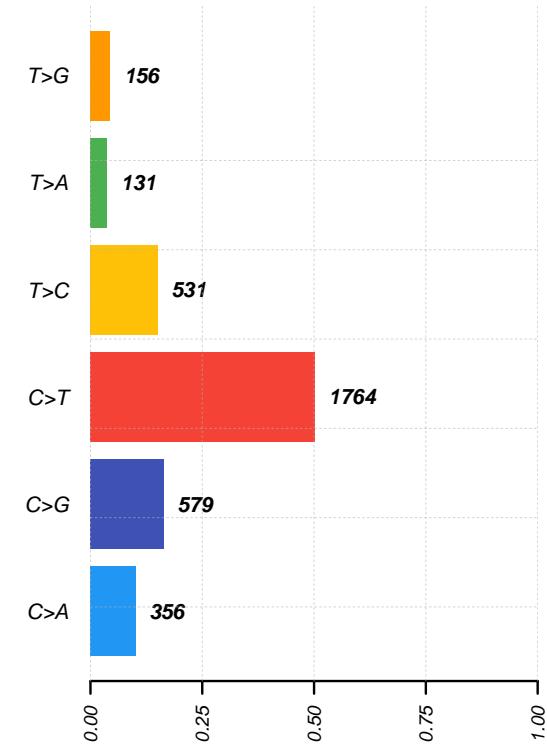
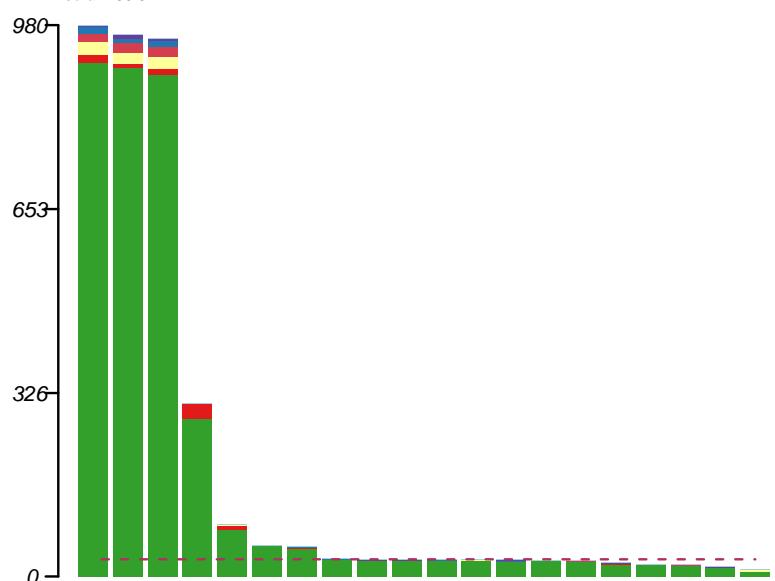
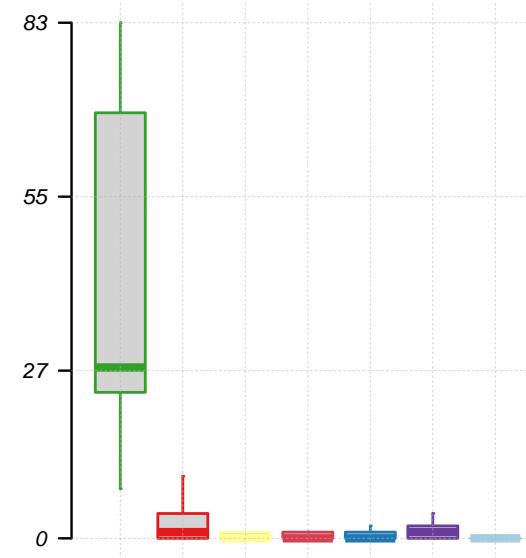


**Variant Classification****Variant Type****SNV Class****Variants per sample**

Median: 30.5

**Variant Classification summary****Top 10 mutated genes**