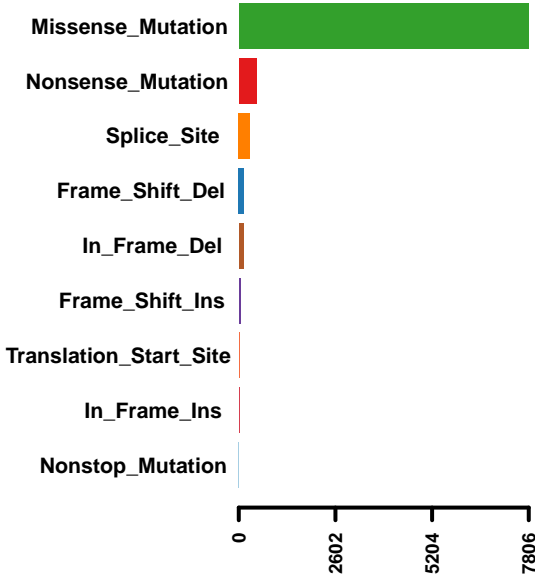
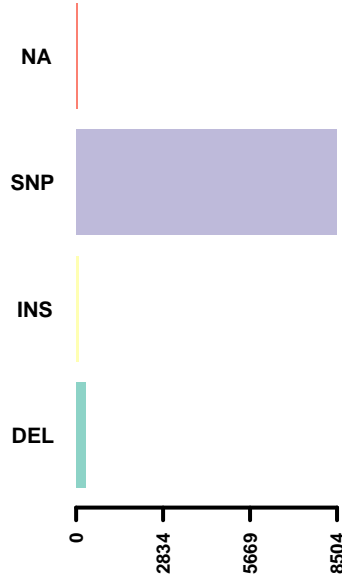


MAF summary

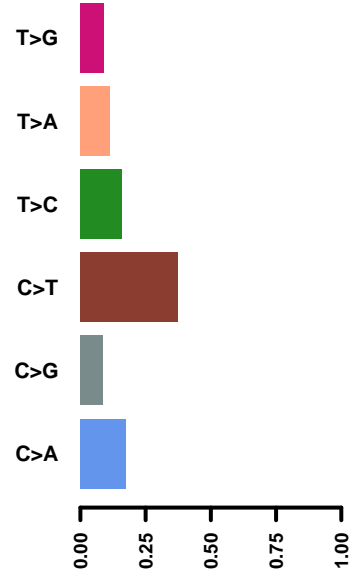
Variant Classification



Variant Type

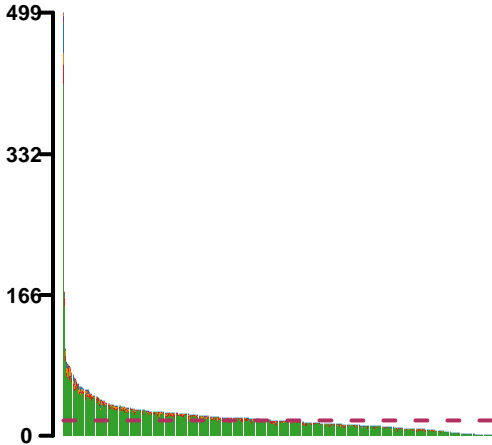


SNV Class

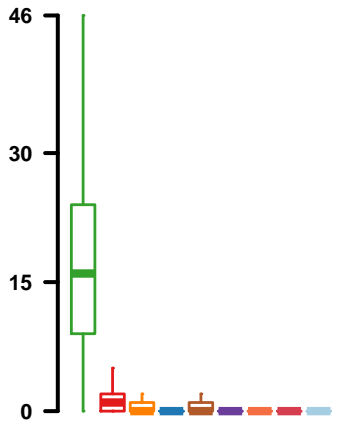


Variants per sample

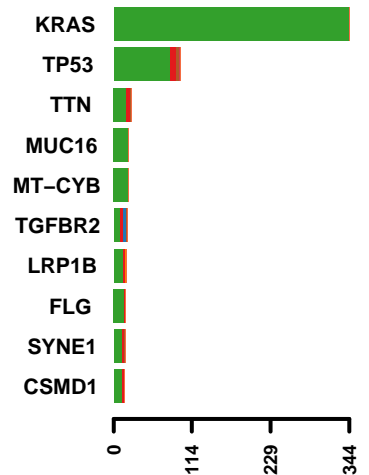
Median: 18



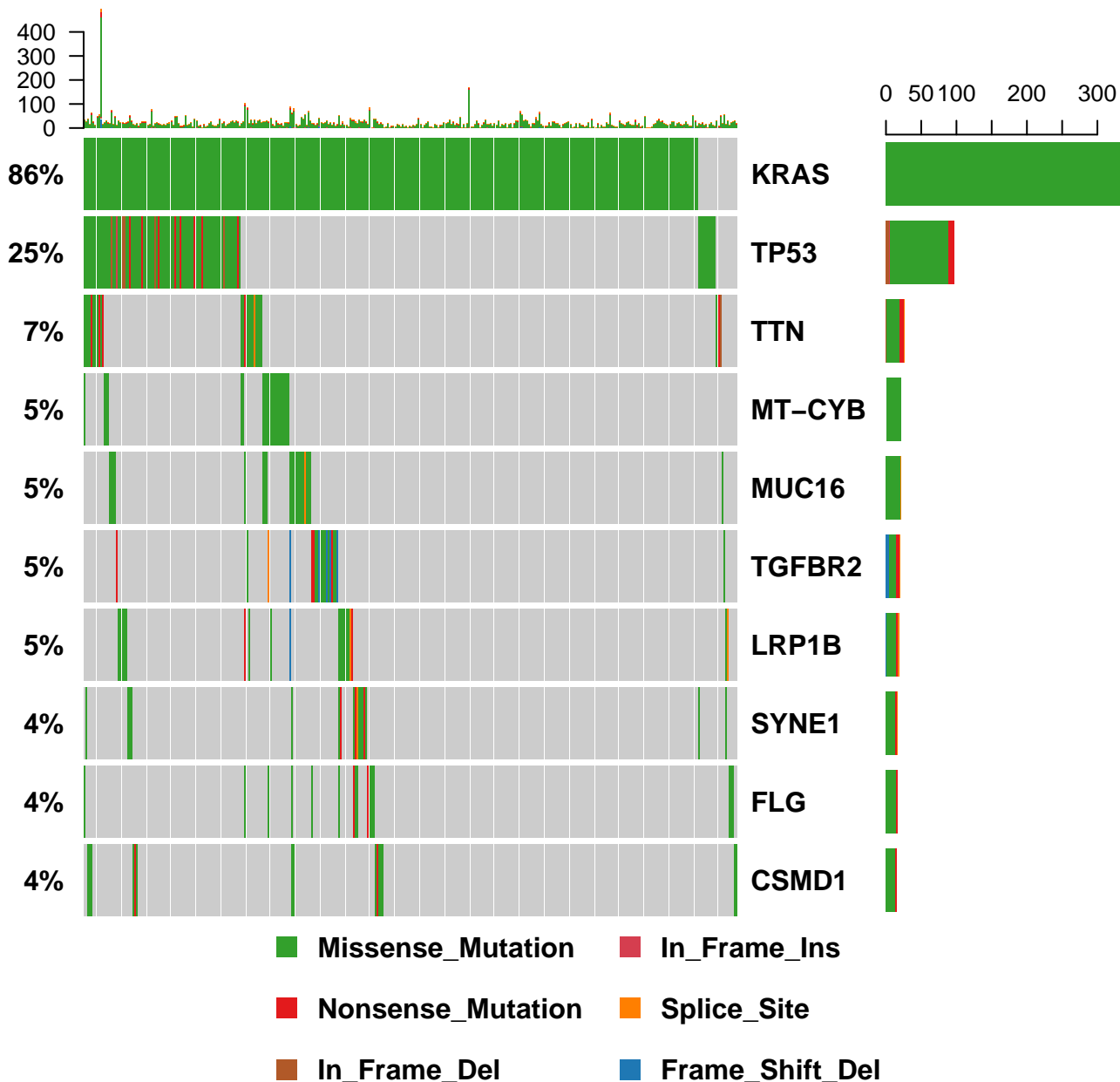
Variant Classification summary



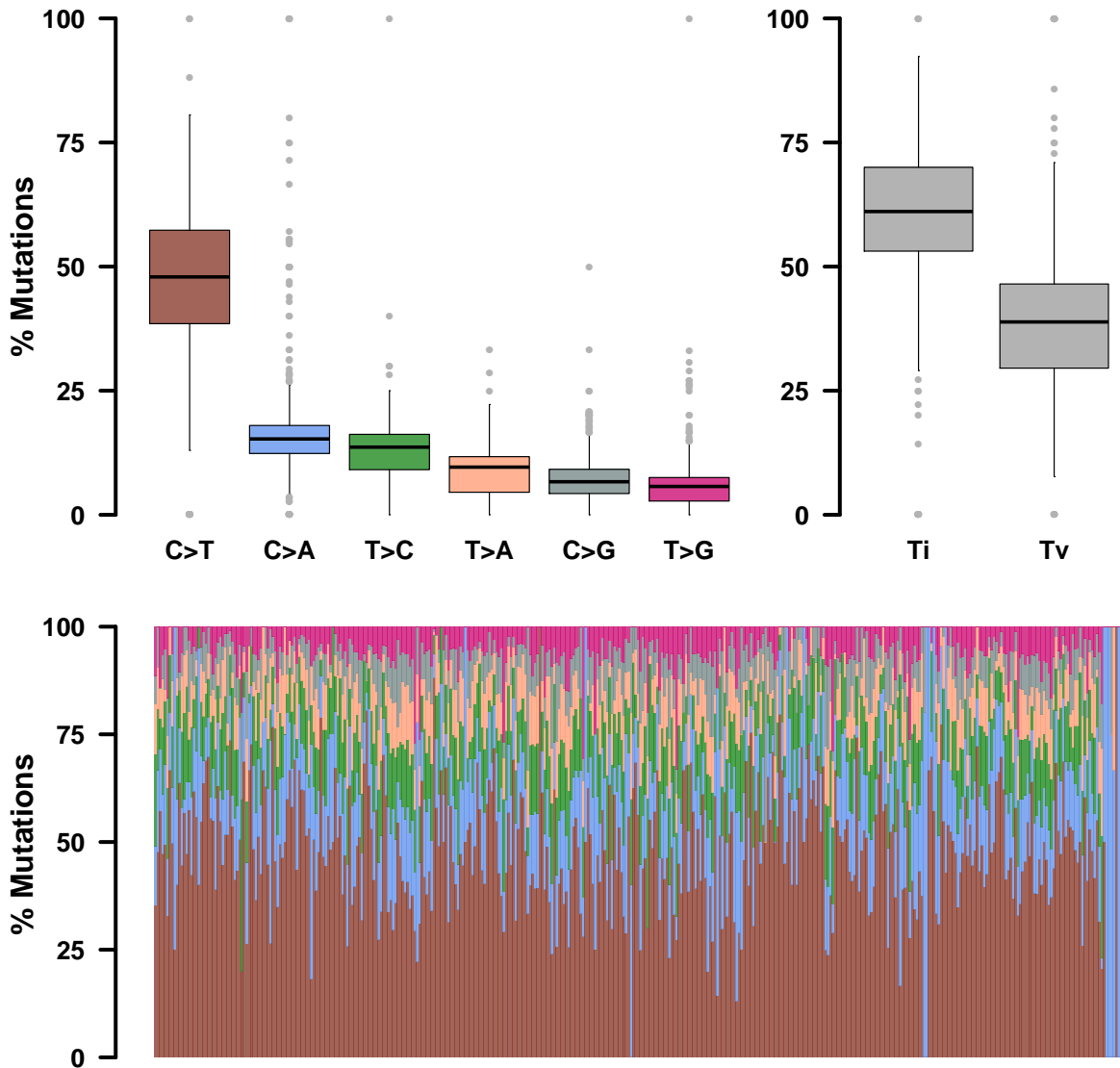
Top 10 mutated genes



Altered in 362 (91.65%) of 395 samples.



Transition and transversions distribution



Mutation load in TCGA cohorts

