

Information regarding the percentage of transitions and transversions across the samples:

The percentage of transitions is slightly higher than transversions across all the samples. Across all the samples the highest mutations are C > T, which is also the most commonly present point mutation across variety of cancer genomes . Of the Transversion mutations C>A is higher than the other types of mutations in any given sample and also over all across all samples. Together C>T (>35%) and C>A (25%) forms the major bulk of the point mutations across all the samples. T>C transition is also observed in higher percentage (<25%).