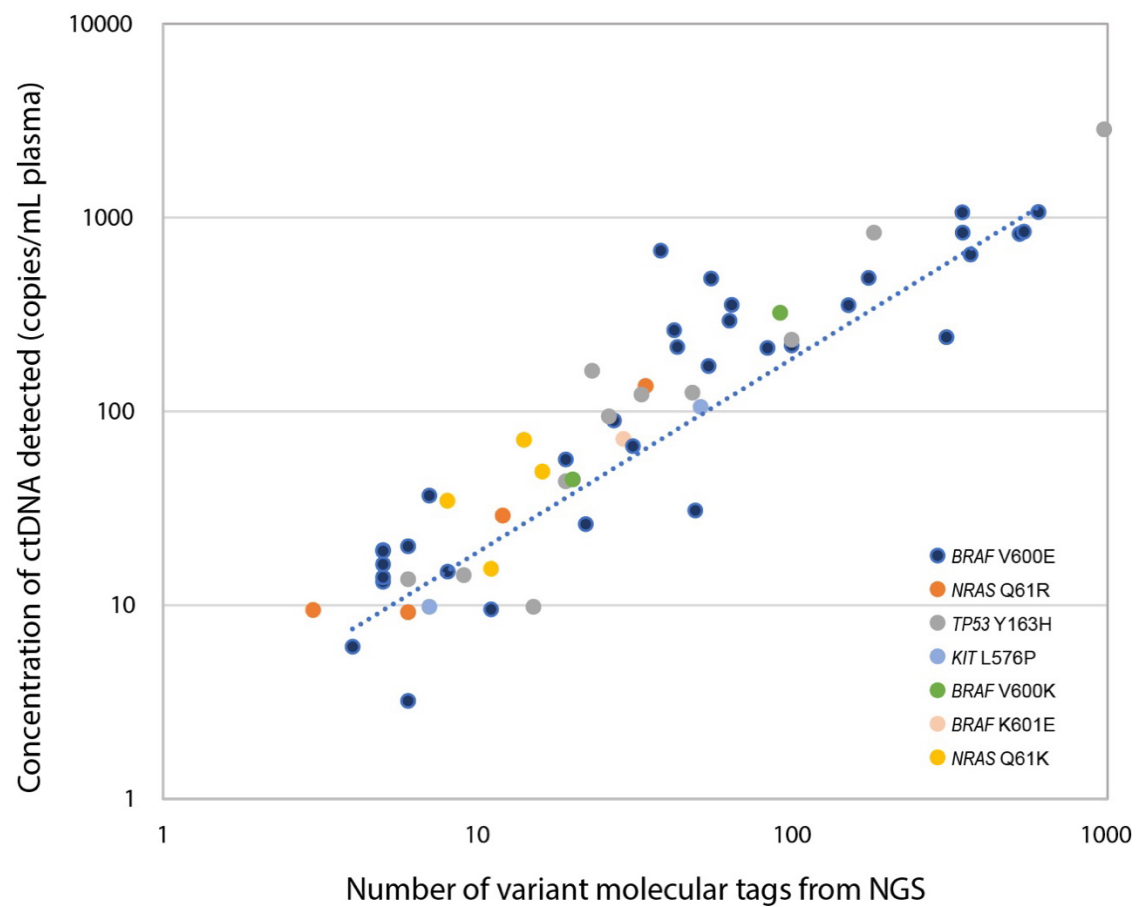




Supplementary Fig. 12 Scatterplot showing the correlation between the number of variant molecular tags (VMT, representative of independent ctDNA molecules sequenced, x-axis) detected by NGS and the quantitation of ctDNA in the same sample by a ddPCR assay (y-axis). The colored circles represent the different mutations identified by NGS and quantitated with ddPCR assays (key to right). The Spearman's Rank correlation is 0.928 and the Pearson correlation coefficient is 0.918. The trendline is shown in blue.