Investigating associations between cancer driver gene mutations and patient survival
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Network Analysis of HIV Reverse Transcriptase via NAPA to Identify Positions of Mutations Conferring Drug Resistance
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Using an undirected alignment network to find adaptive genetic mutations in HIV-1 RT inhibitor proteins
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Associations between driver gene mutations and mutational signatures in cancer
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Determining Statistical Correlation between Timelines of Phylogenetic Mutation Patterns in HIV Reverse Transcriptase & NRTI/NNRTI FDA Approvals
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Detecting Microsatellite Instability in the Cancer Genome
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Driving Death: The relationship between cancer driver mutations and survival odds
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Identify associations between mutational signatures and immune cell infiltration
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Applying new evaluation protocol of cancer driver gene classifiers to benchmark newer classifiers
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Patient-specific network analysis of protein adaptation in HIV reverse transcriptase
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Detection of association between mutation in driver genes and patient survival in TCGA data
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