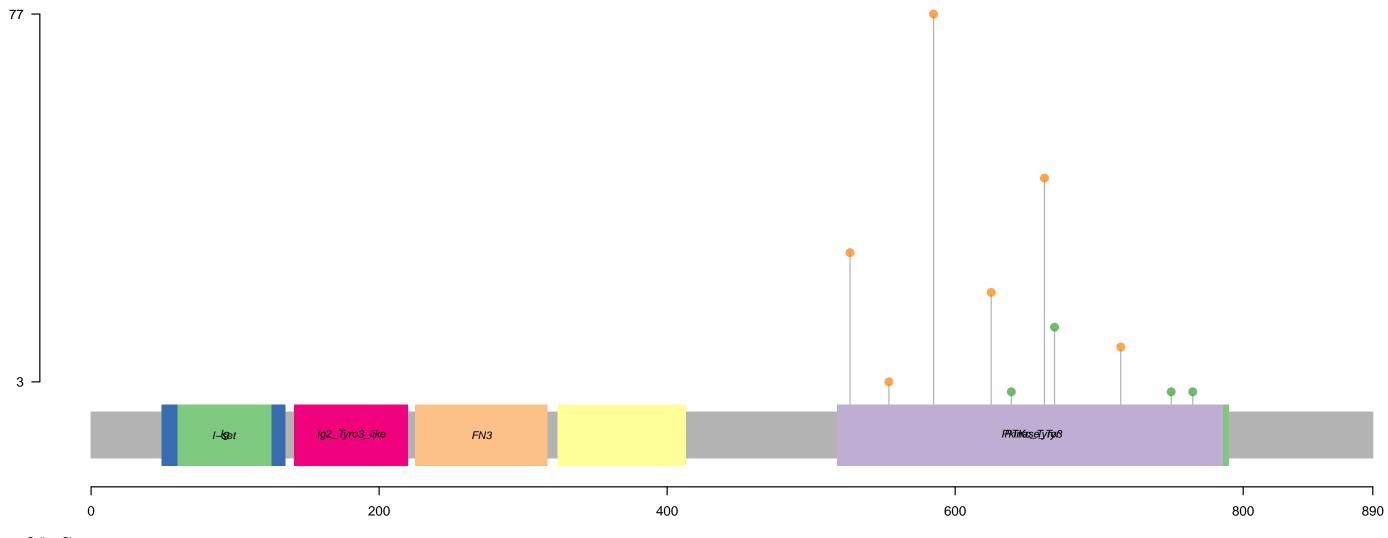
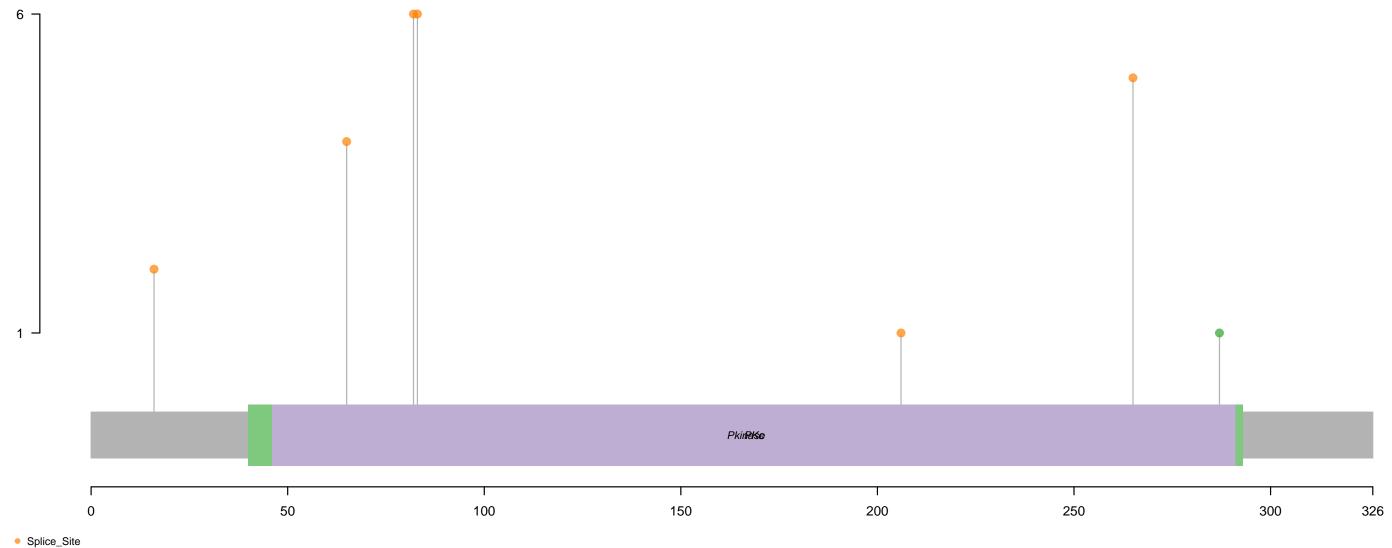
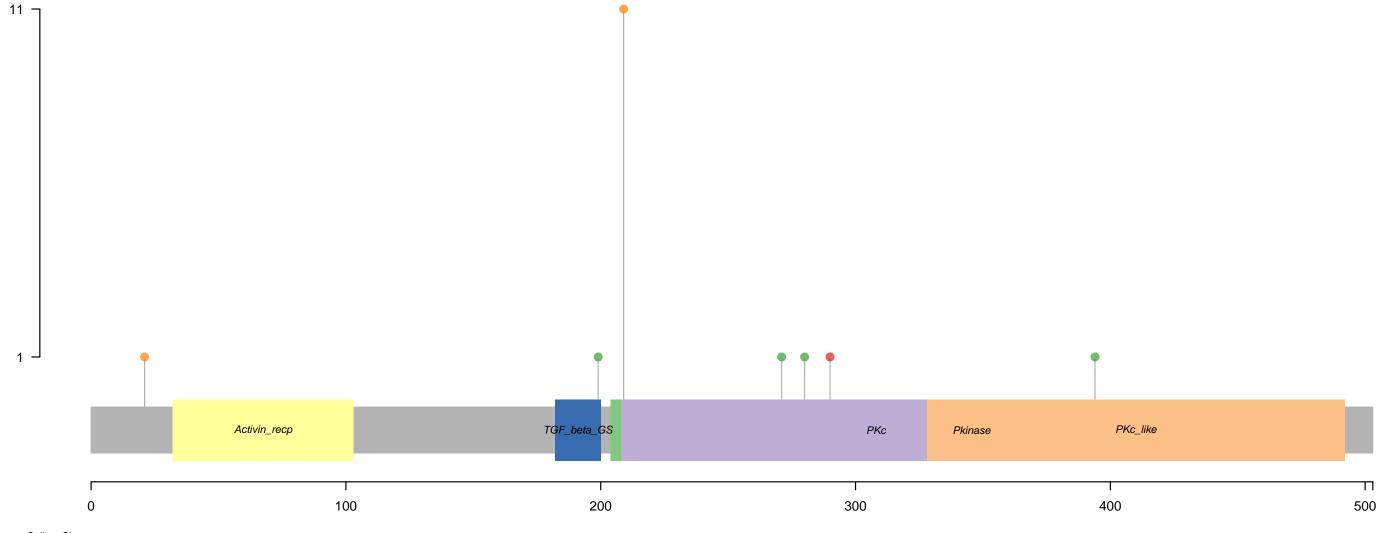
TYRO3 : [Somatic Mutation Rate: 27.16%] NM_006293



PIM3 : [Somatic Mutation Rate: 3.06%] NM_001001852



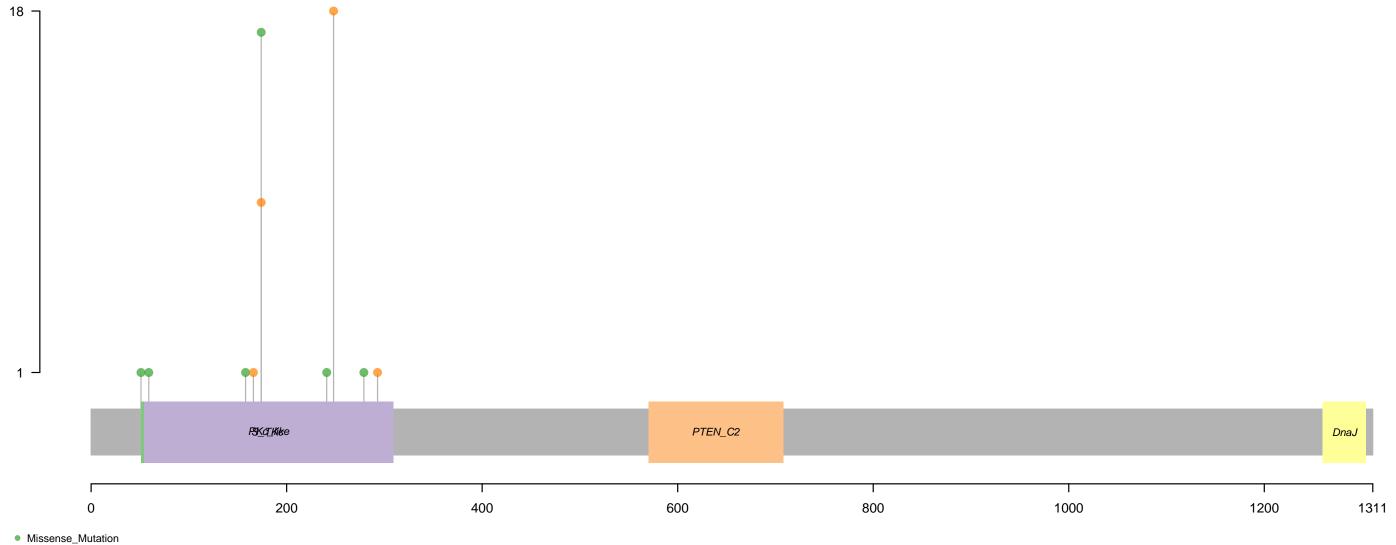
ACVRL1: [Somatic Mutation Rate: 3.06%] NM_000020



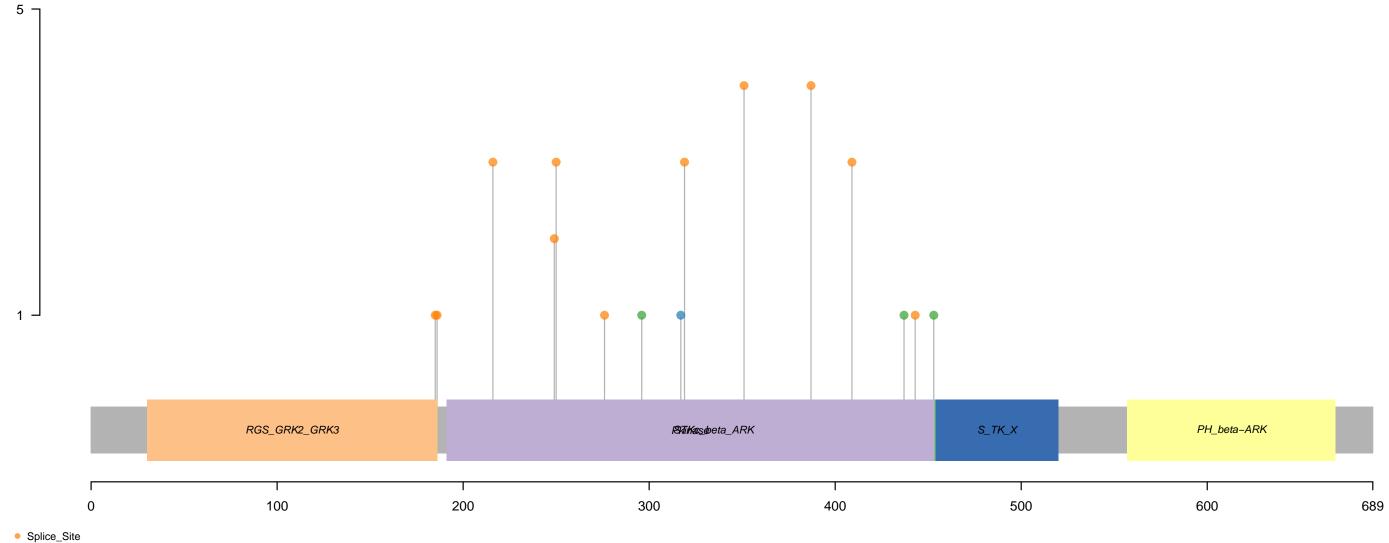
Missense_Mutation

Nonsense_Mutation

GAK: [Somatic Mutation Rate: 8.63%] NM_005255

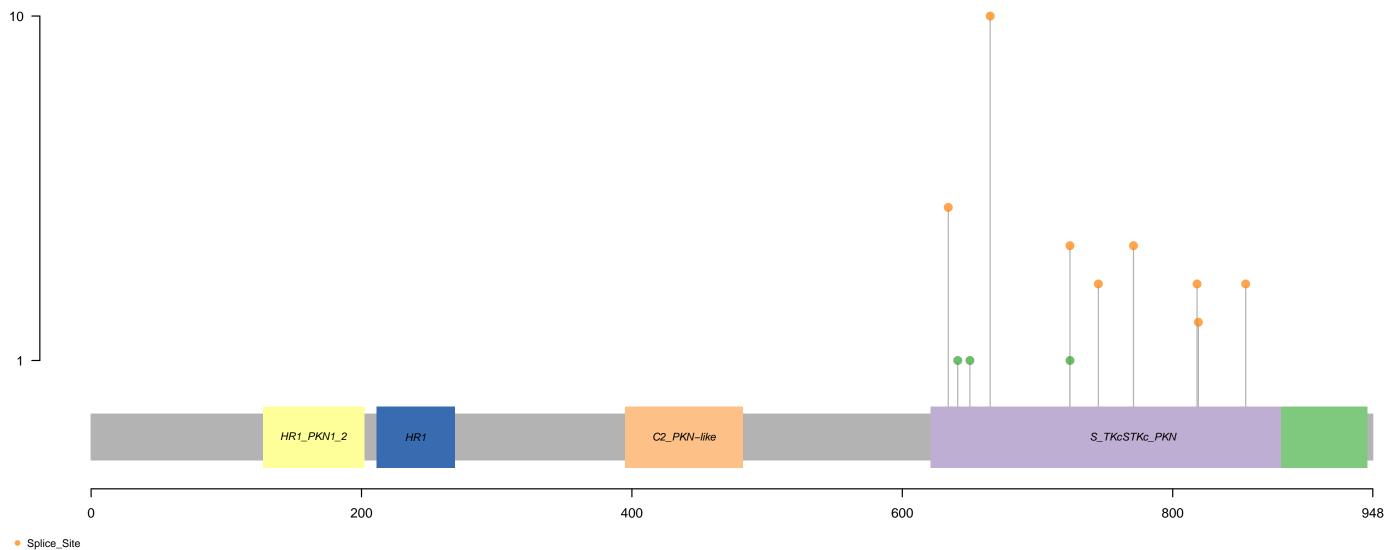


GRK2: [Somatic Mutation Rate: 3.06%] NM_001619

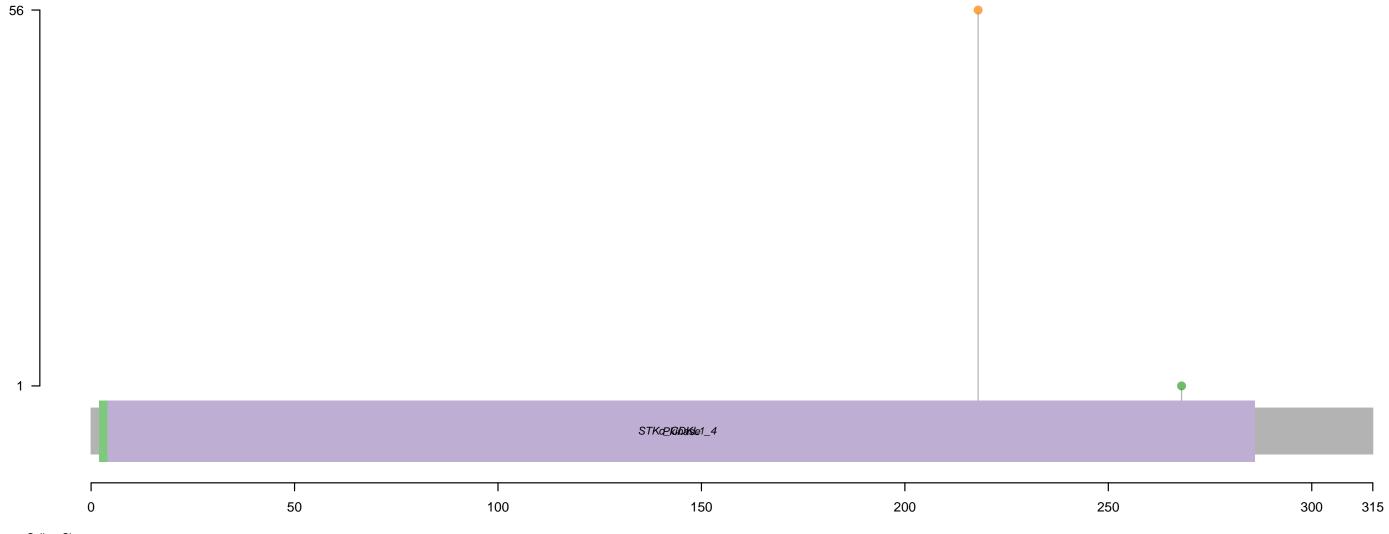


Frame_Shift_Del

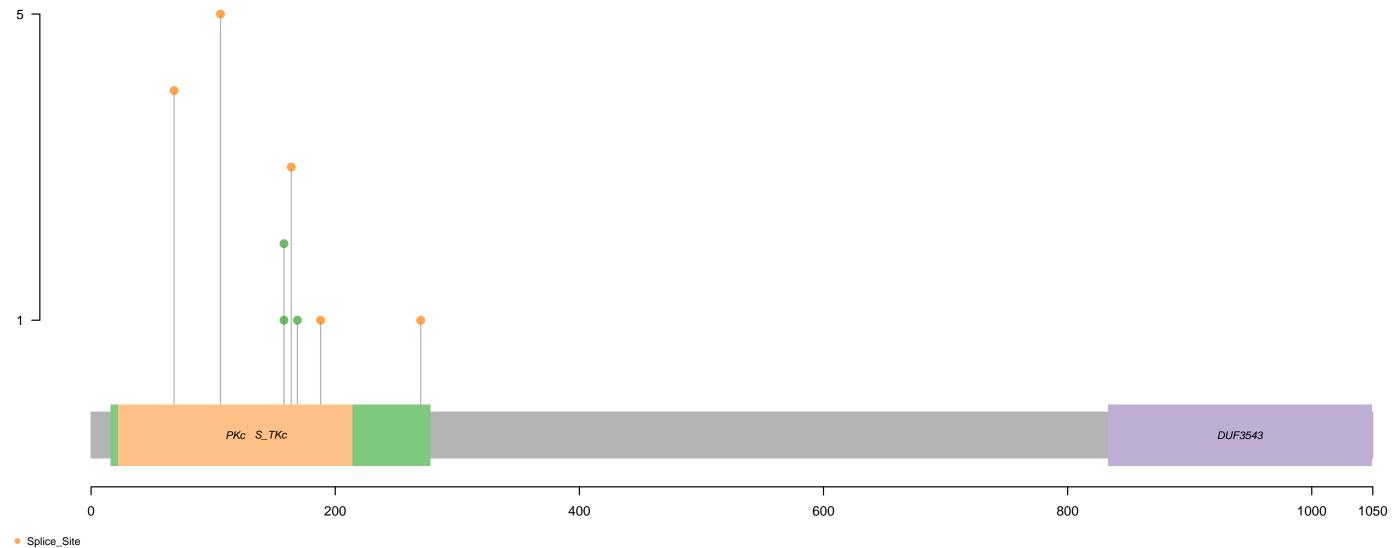
PKN1 : [Somatic Mutation Rate: 3.96%] NM_213560



CDKL4: [Somatic Mutation Rate: 10.07%] NM_001009565



ULK1 : [Somatic Mutation Rate: 3.06%] NM_003565



RPS6KA4 : [Somatic Mutation Rate: 4.32%] NM_003942

