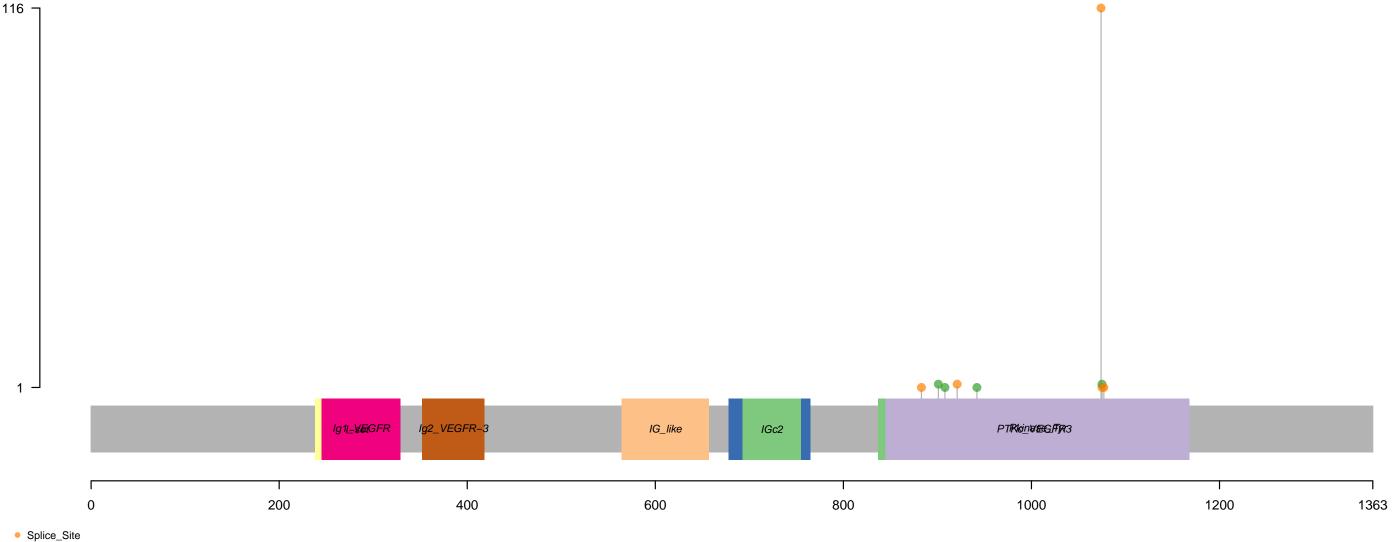
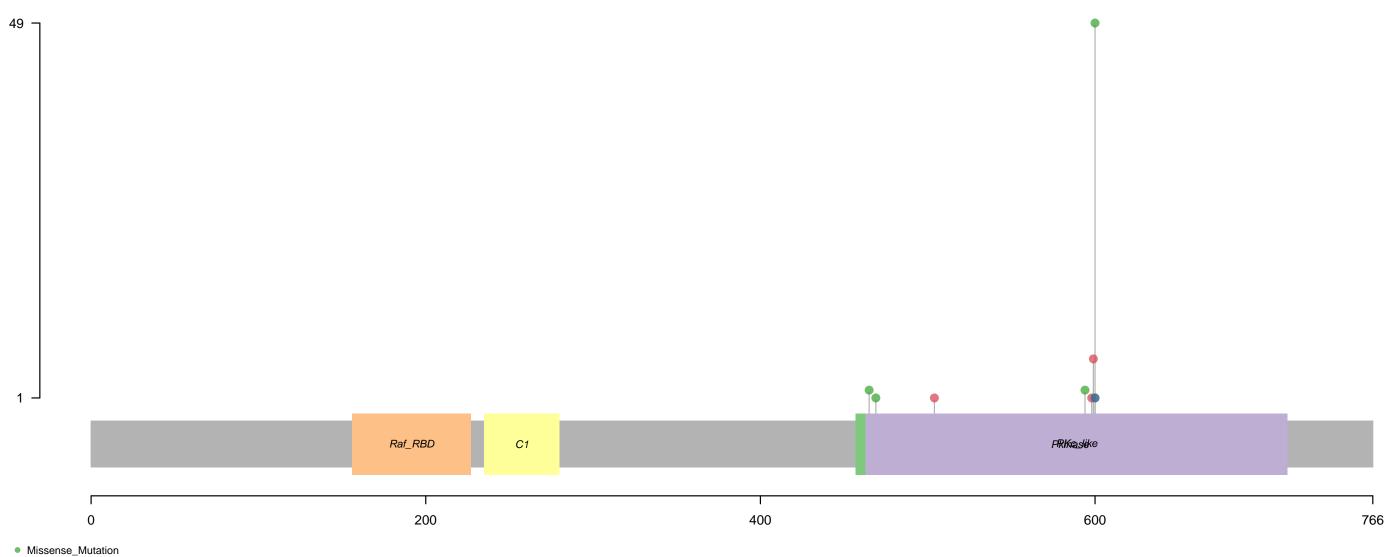
FLT4: [Somatic Mutation Rate: 18.08%] NM_182925

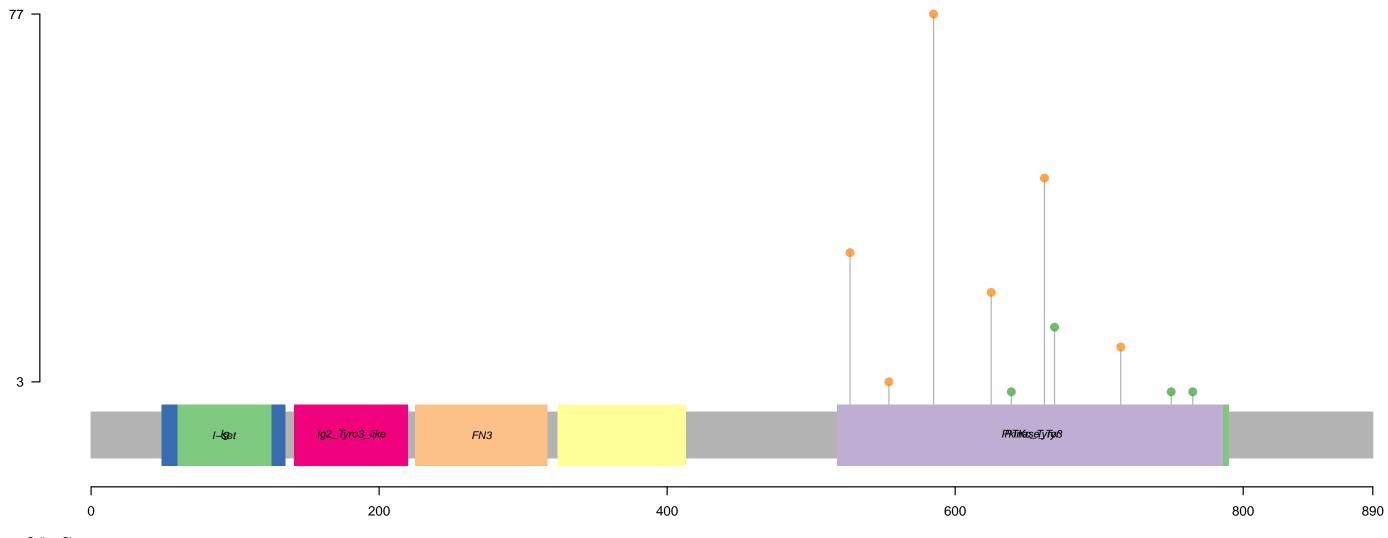


BRAF: [Somatic Mutation Rate: 8.75%] NM_004333



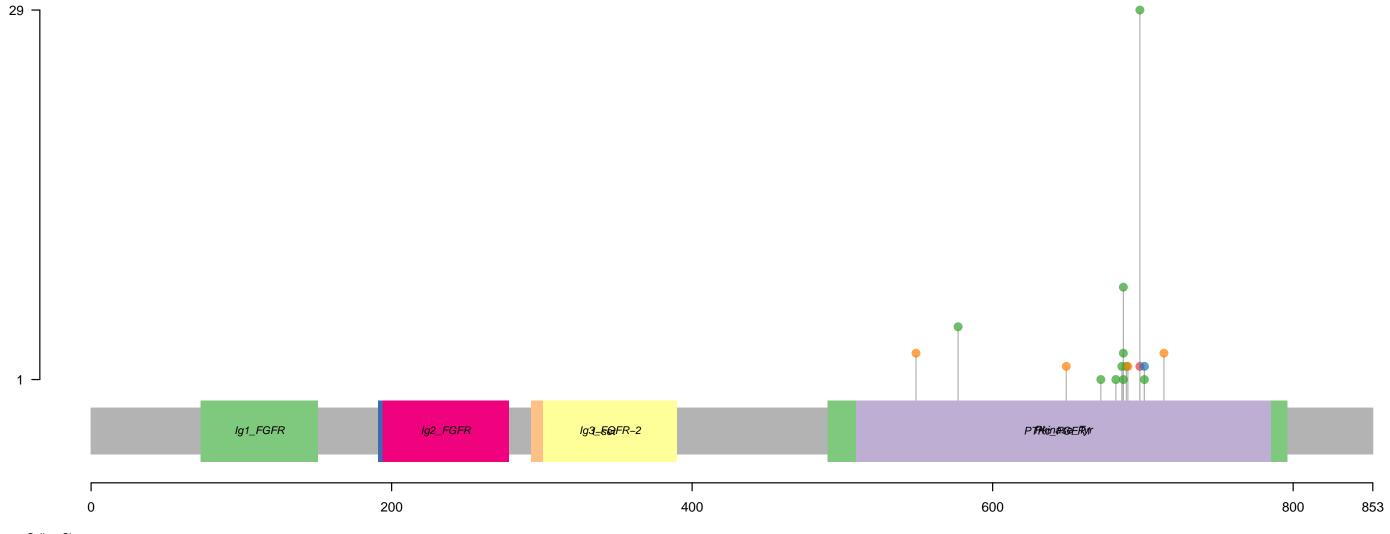
- In_Frame_Ins
- In_Frame_Del
- Frame_Shift_Del

TYRO3 : [Somatic Mutation Rate: 22.01%] NM_006293



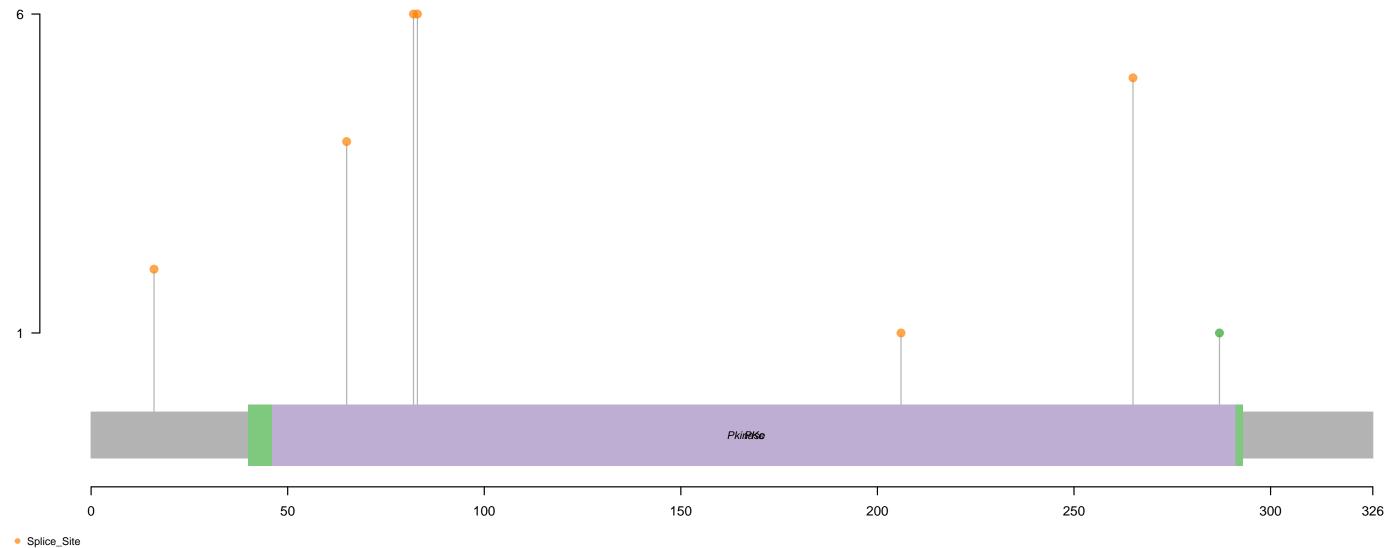
Splice_Site

FGFR1 : [Somatic Mutation Rate: 7.58%] NM_001174067

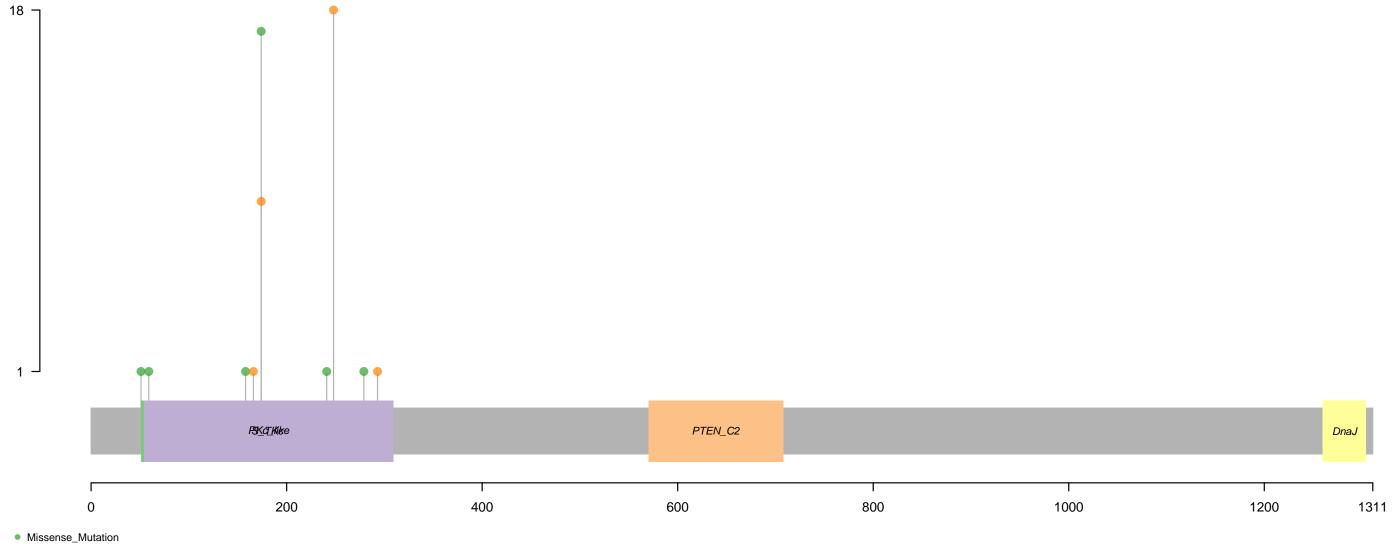


- Splice_Site
- Missense_Mutation
- In_Frame_Ins
- Frame_Shift_Del

PIM3 : [Somatic Mutation Rate: 2.48%] NM_001001852

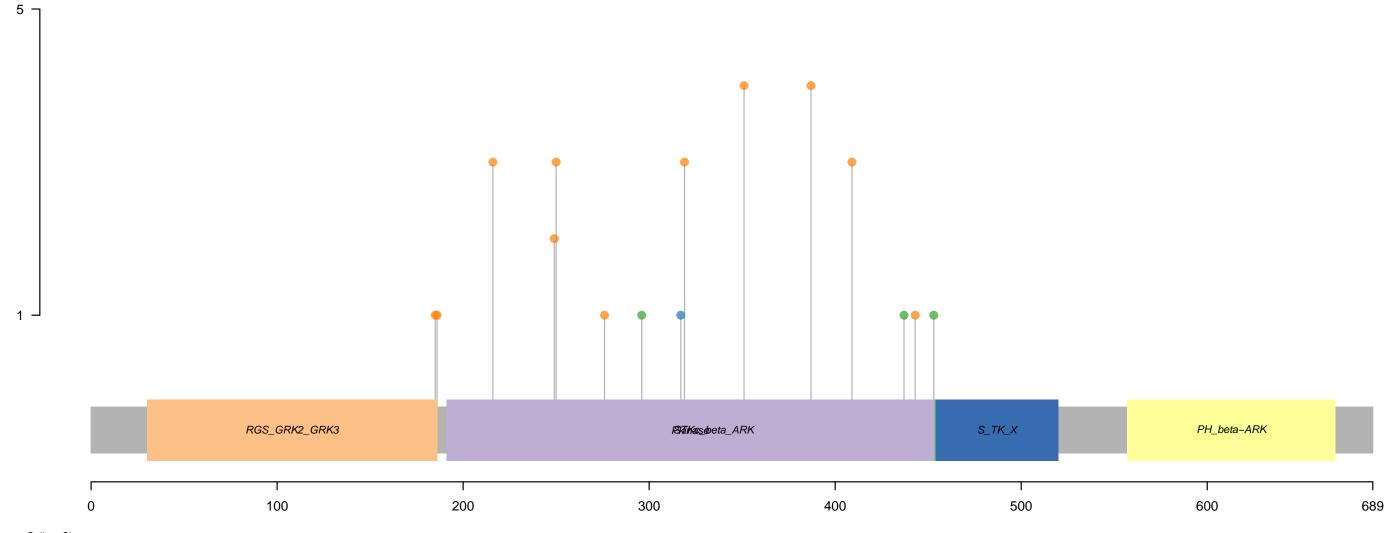


GAK: [Somatic Mutation Rate: 7%] NM_005255



Splice_Site

GRK2: [Somatic Mutation Rate: 2.48%] NM_001619

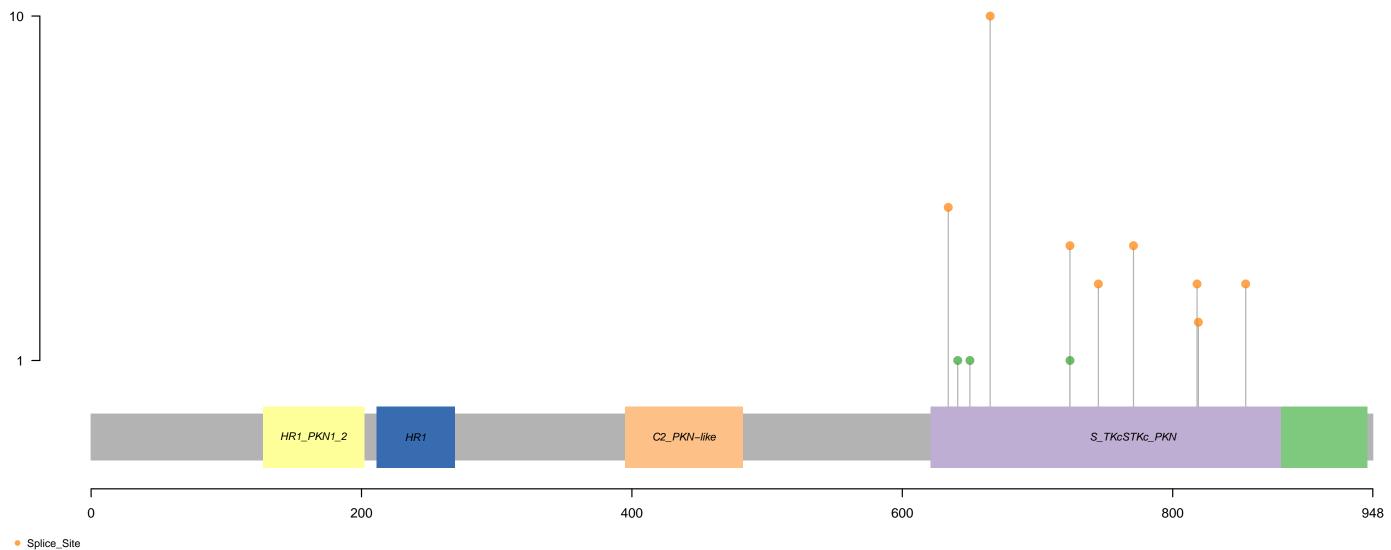


Splice_Site

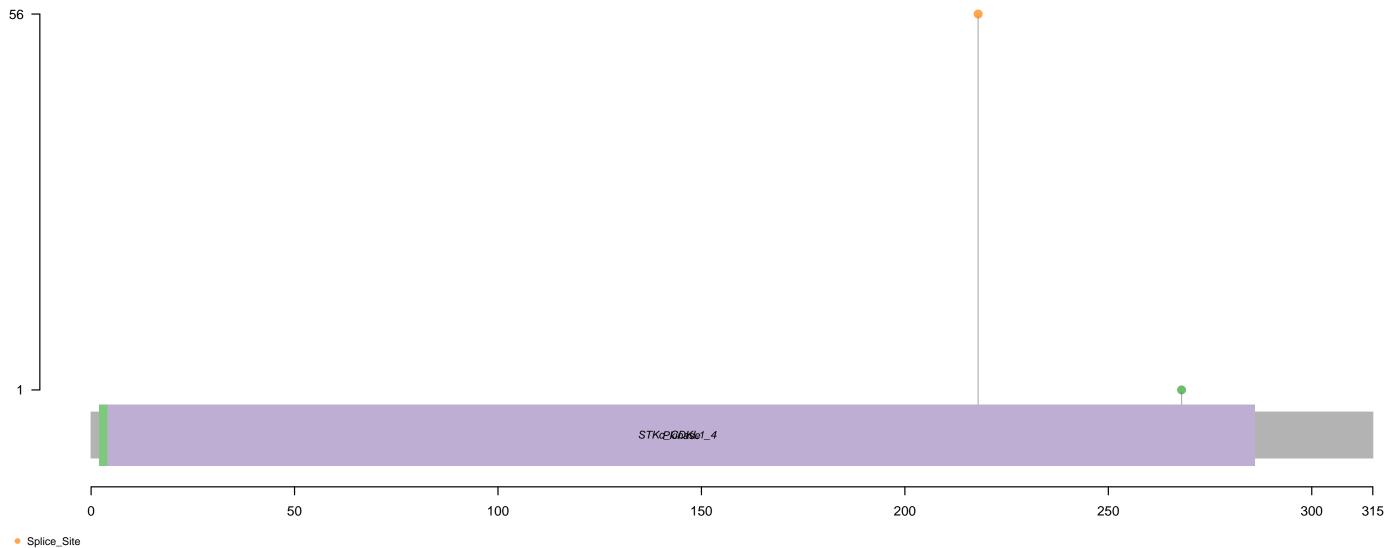
Missense_Mutation

Frame_Shift_Del

PKN1 : [Somatic Mutation Rate: 3.21%] NM_213560



CDKL4: [Somatic Mutation Rate: 8.16%] NM_001009565



RPS6KA4 : [Somatic Mutation Rate: 3.5%] NM_003942

