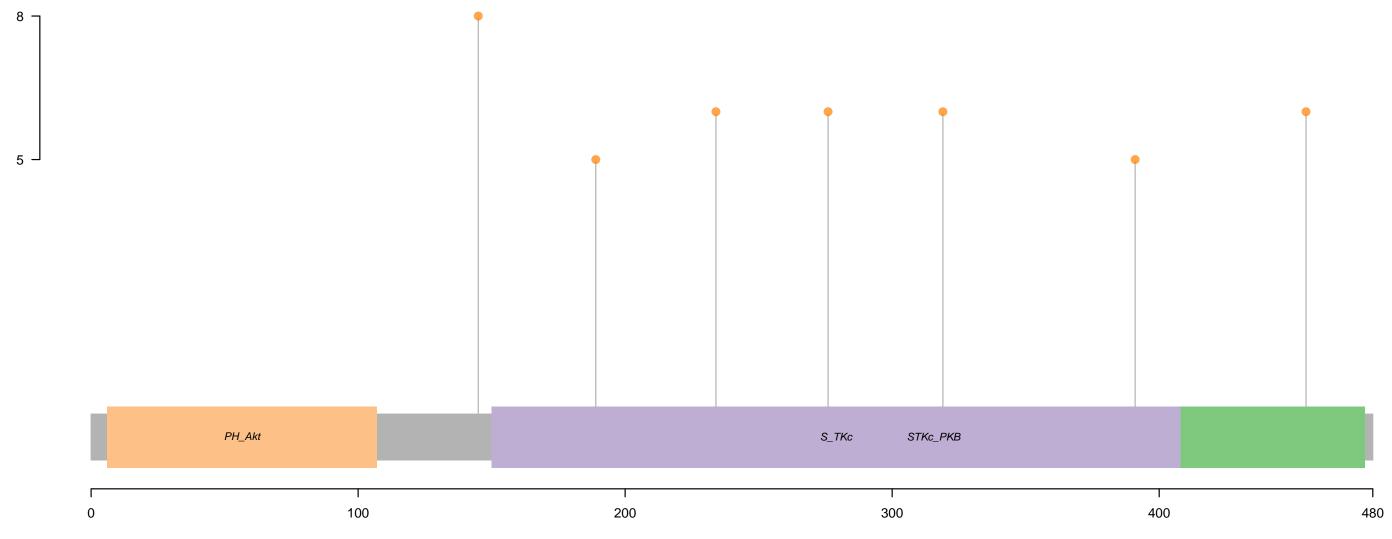
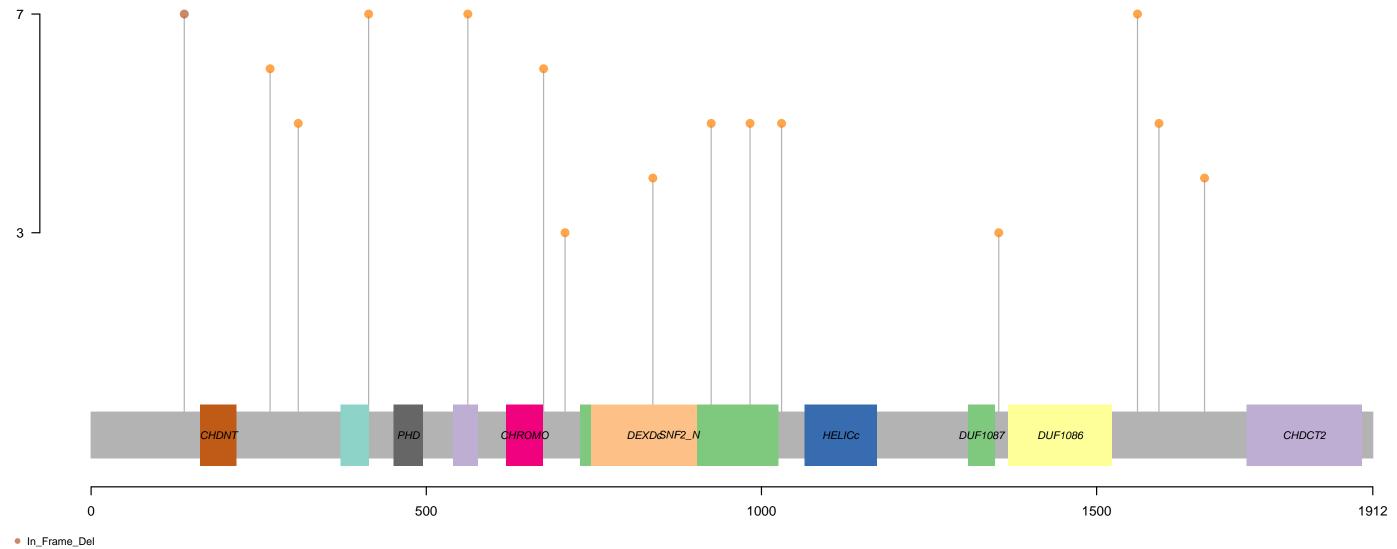
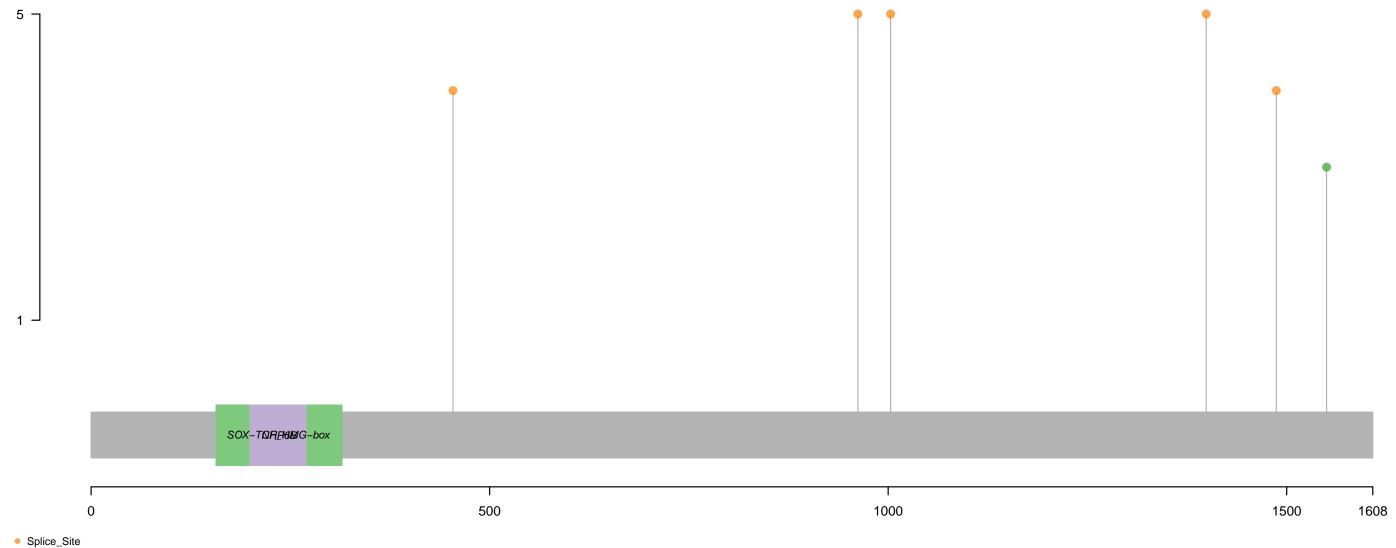
AKT1: [Somatic Mutation Rate: 4.3%] NM_005163



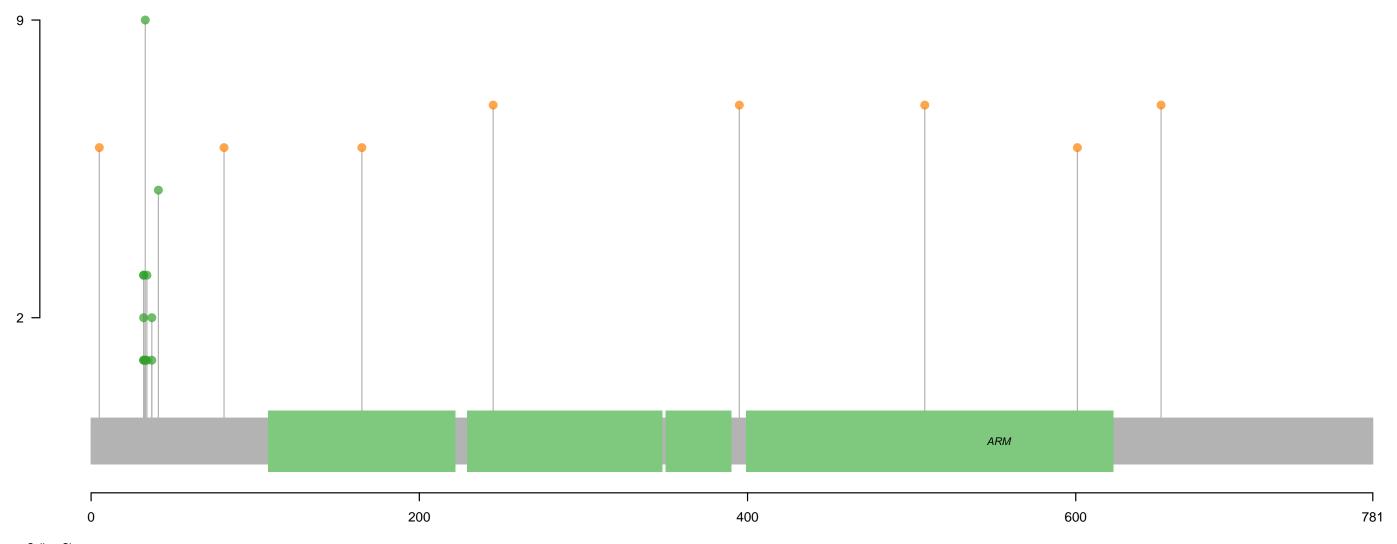
CHD4: [Somatic Mutation Rate: 4.97%] NM_001273



CIC: [Somatic Mutation Rate: 3.31%] NM_015125

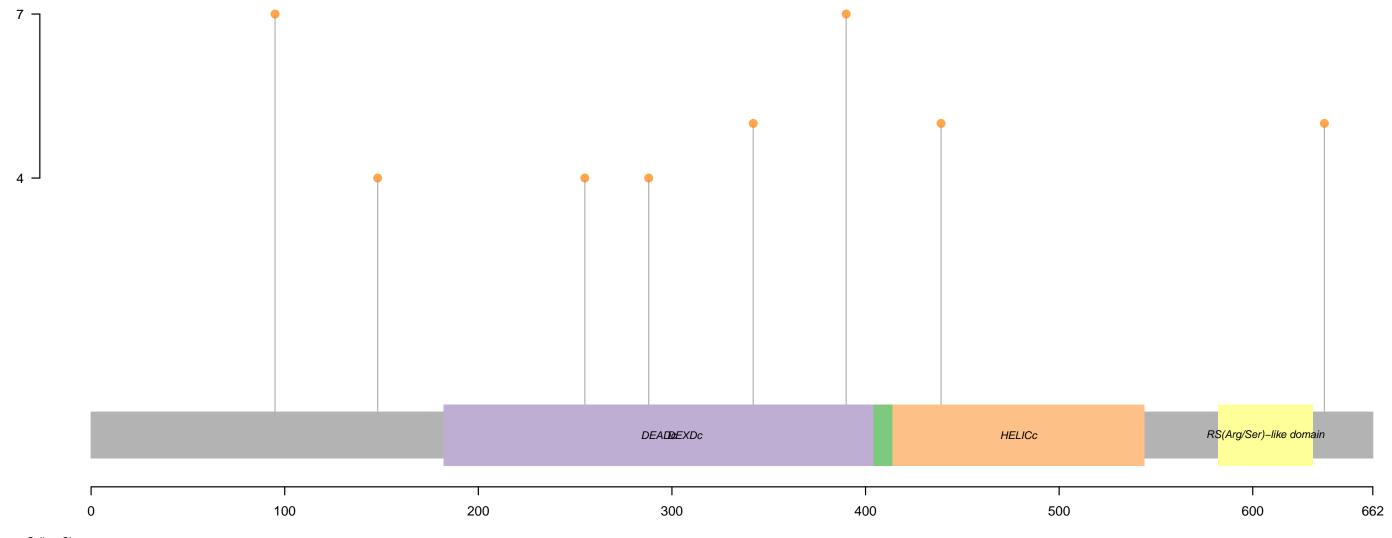


Missense_Mutation

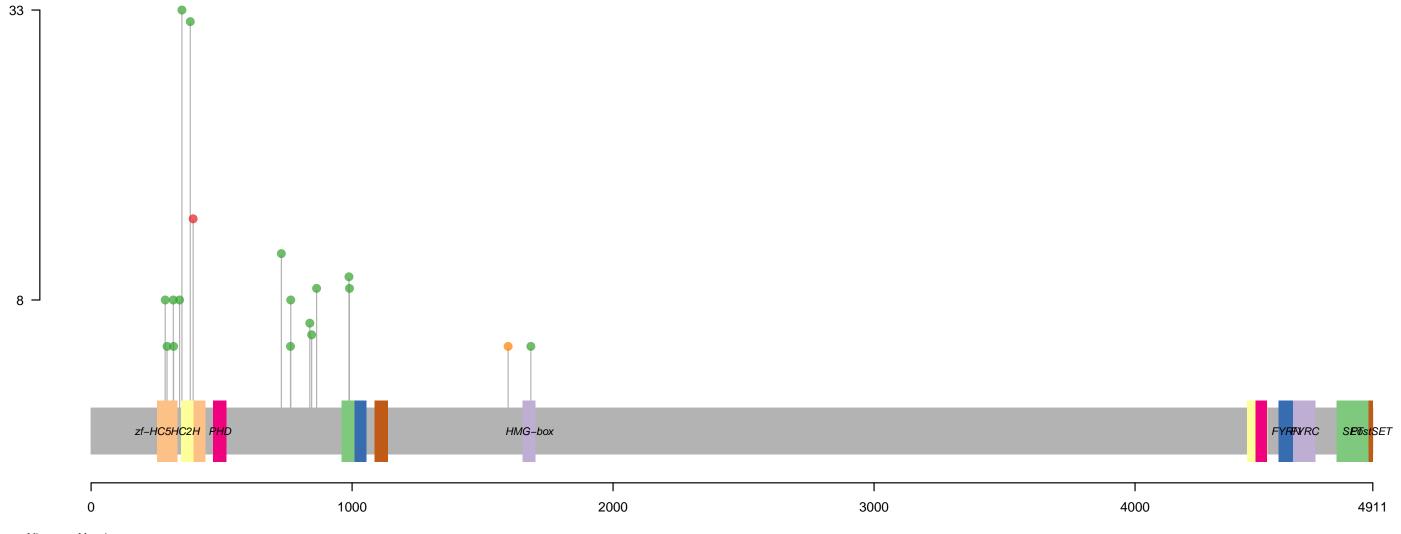


Missense_Mutation

DDX3X : [Somatic Mutation Rate: 2.98%] NM_001356

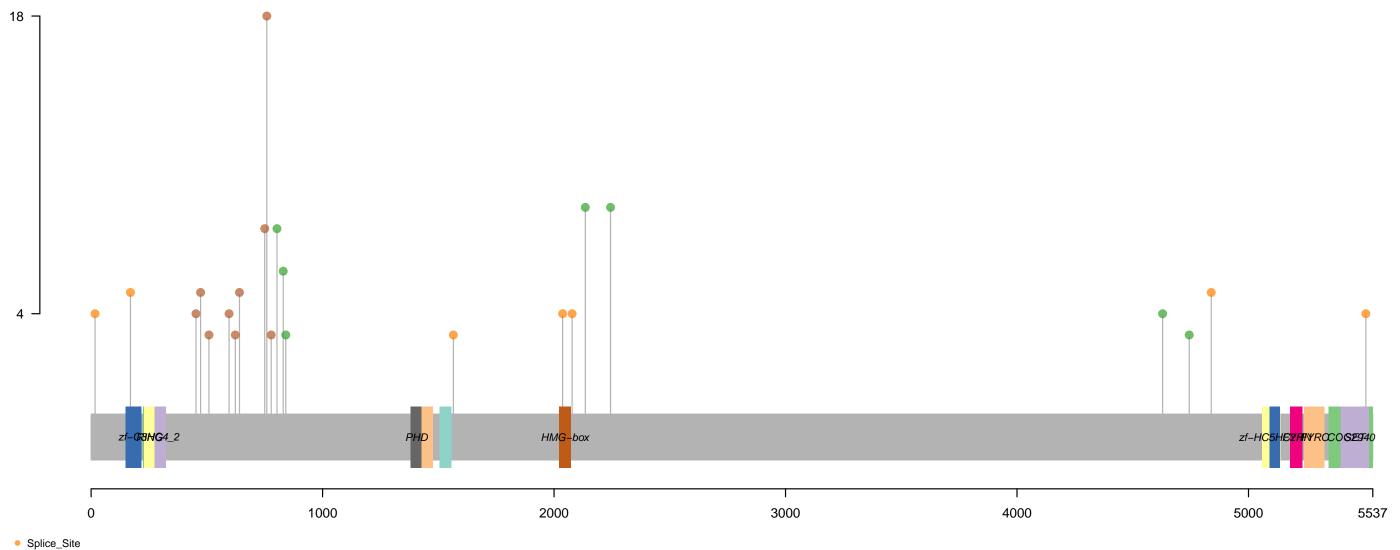


KMT2C : [Somatic Mutation Rate: 50.66%] NM_170606



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site

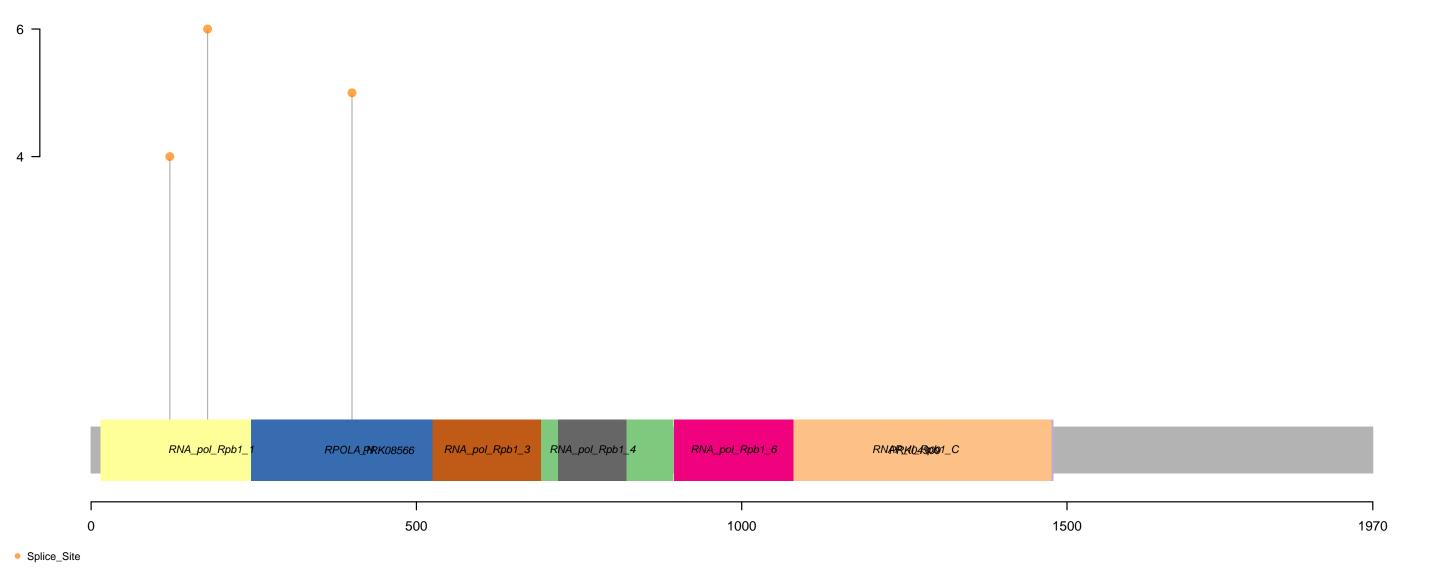
KMT2D : [Somatic Mutation Rate: 30.13%] NM_003482



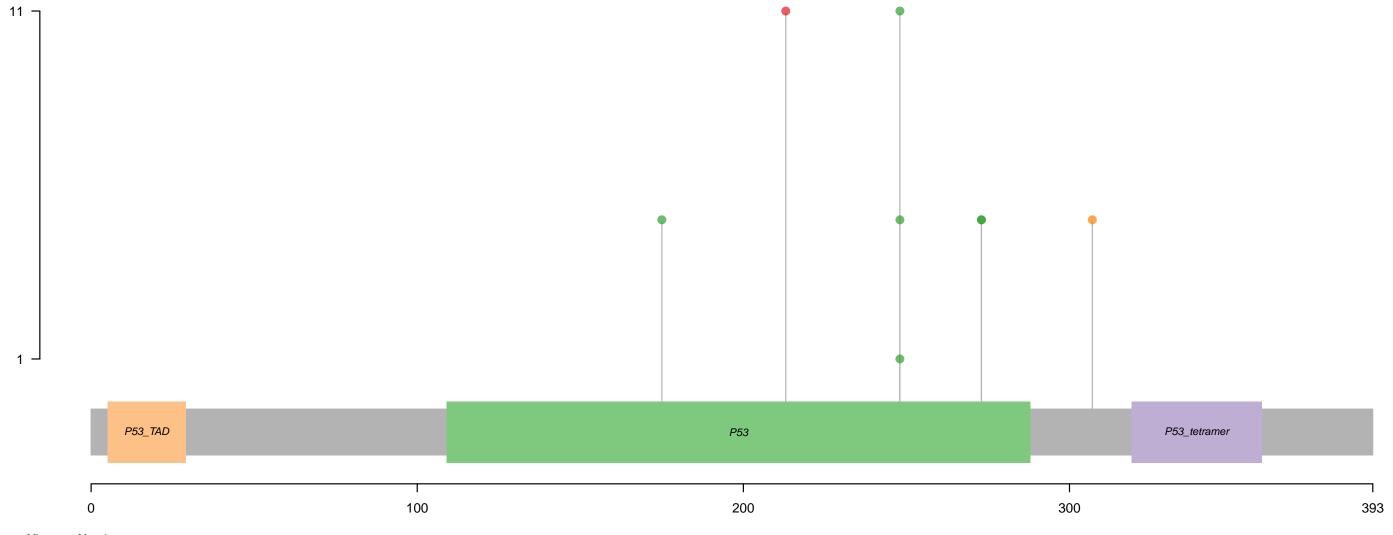
In_Frame_Del

Missense_Mutation

POLR2A: [Somatic Mutation Rate: 2.32%] NM_000937



TP53 : [Somatic Mutation Rate: 15.56%] NM_000546



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site