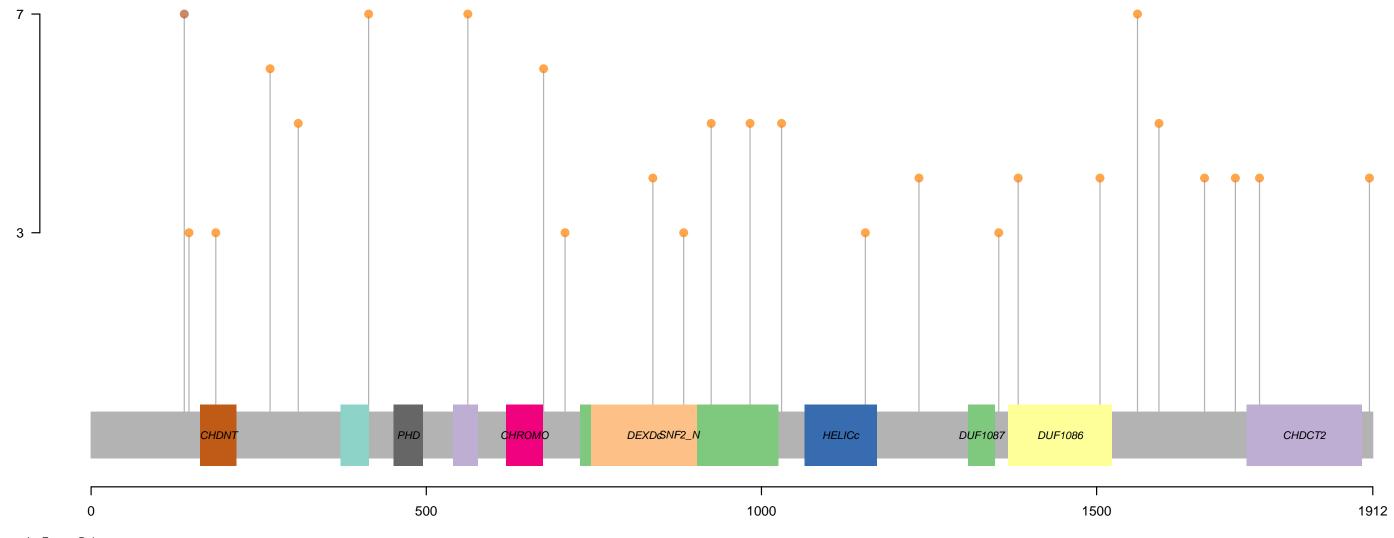
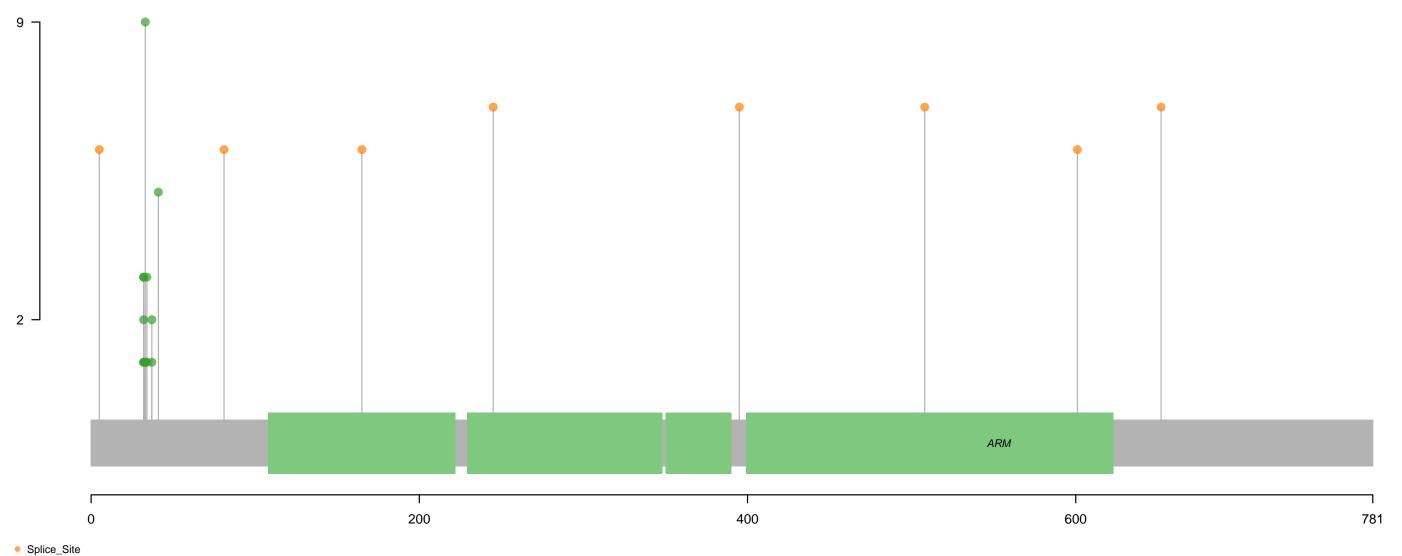
CHD4: [Somatic Mutation Rate: 4.67%] NM_001273



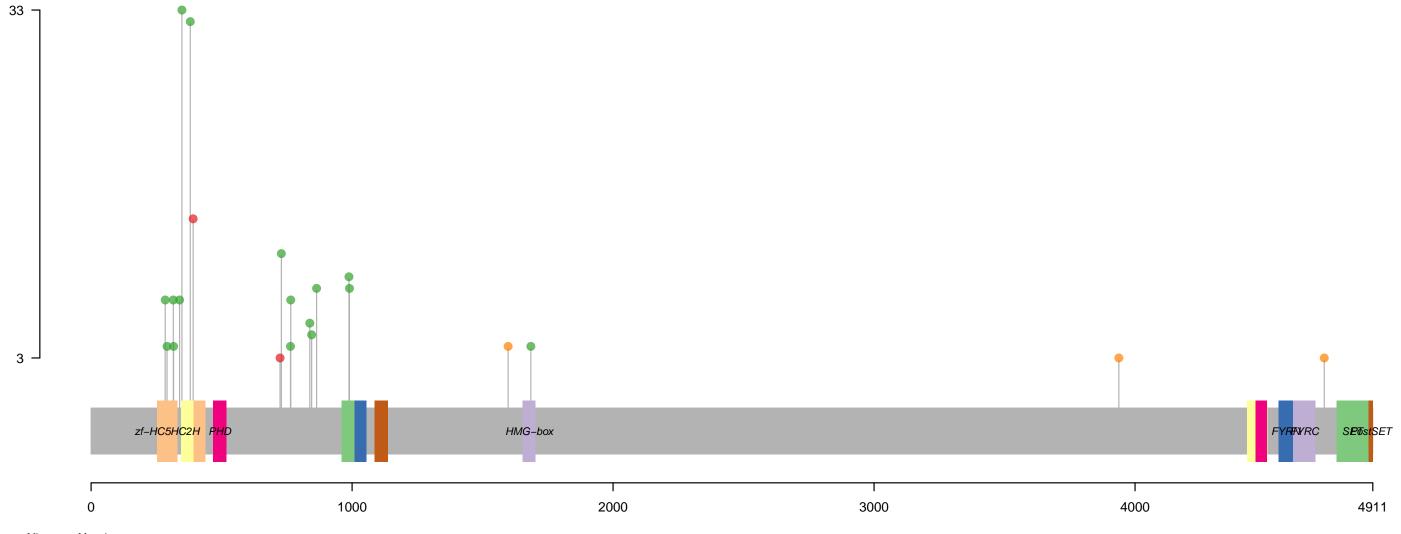
In_Frame_Del

Splice_Site



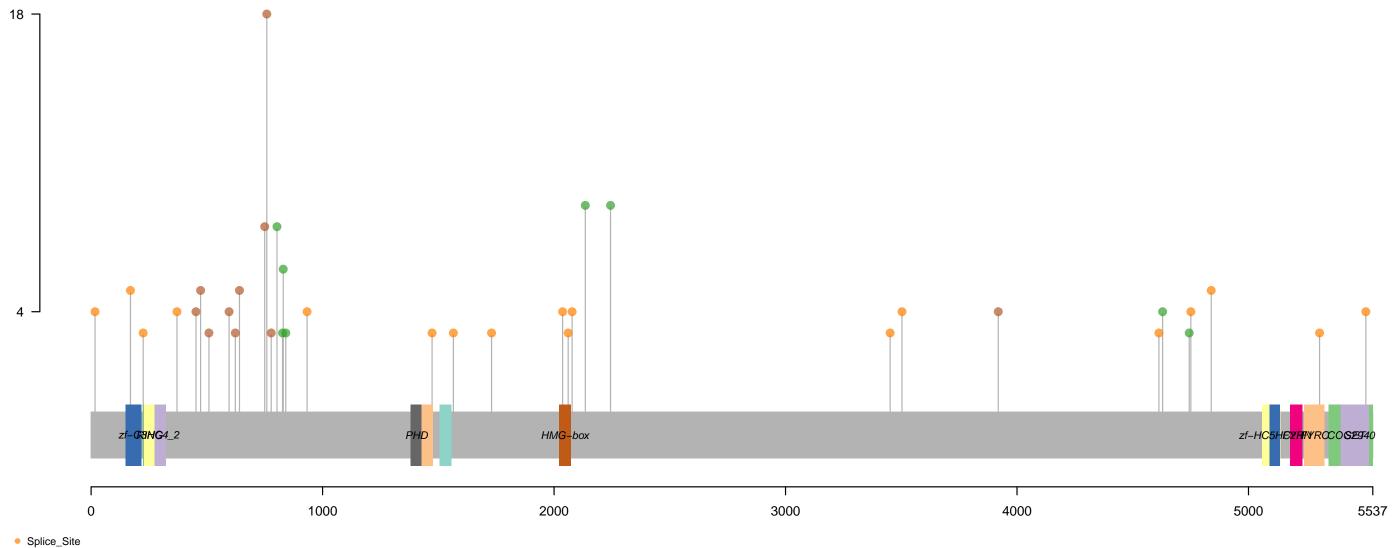
Missense_Mutation

KMT2C : [Somatic Mutation Rate: 48.6%] NM_170606



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site

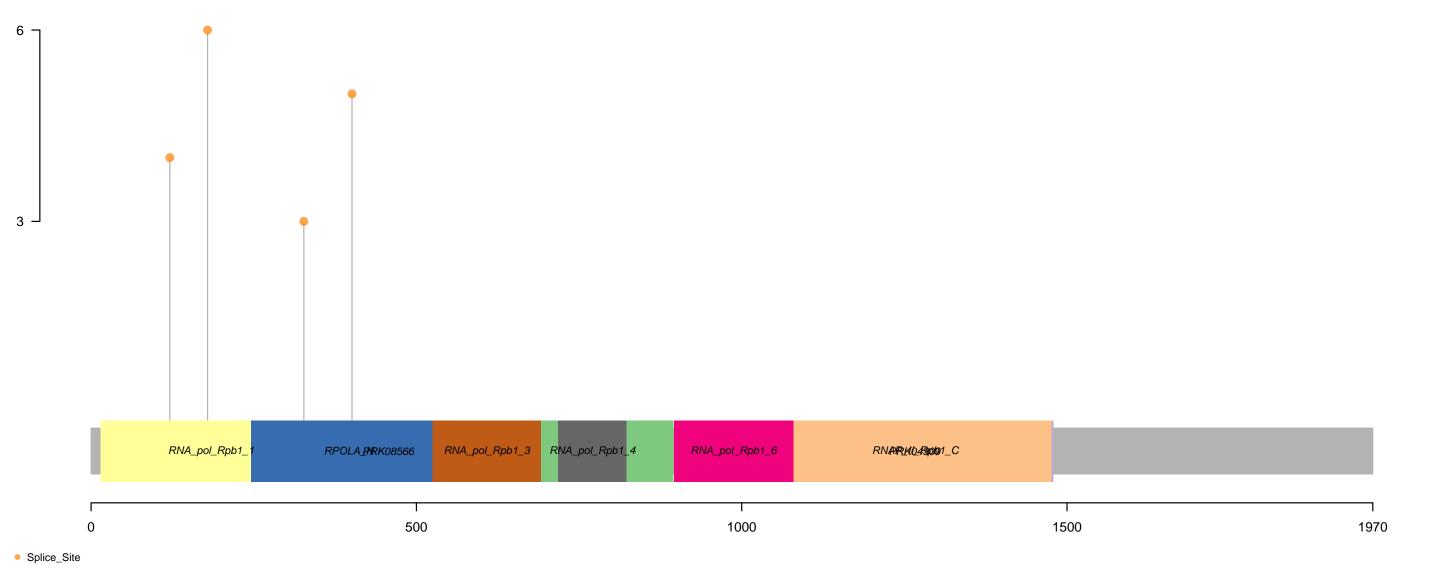
KMT2D : [Somatic Mutation Rate: 30.22%] NM_003482



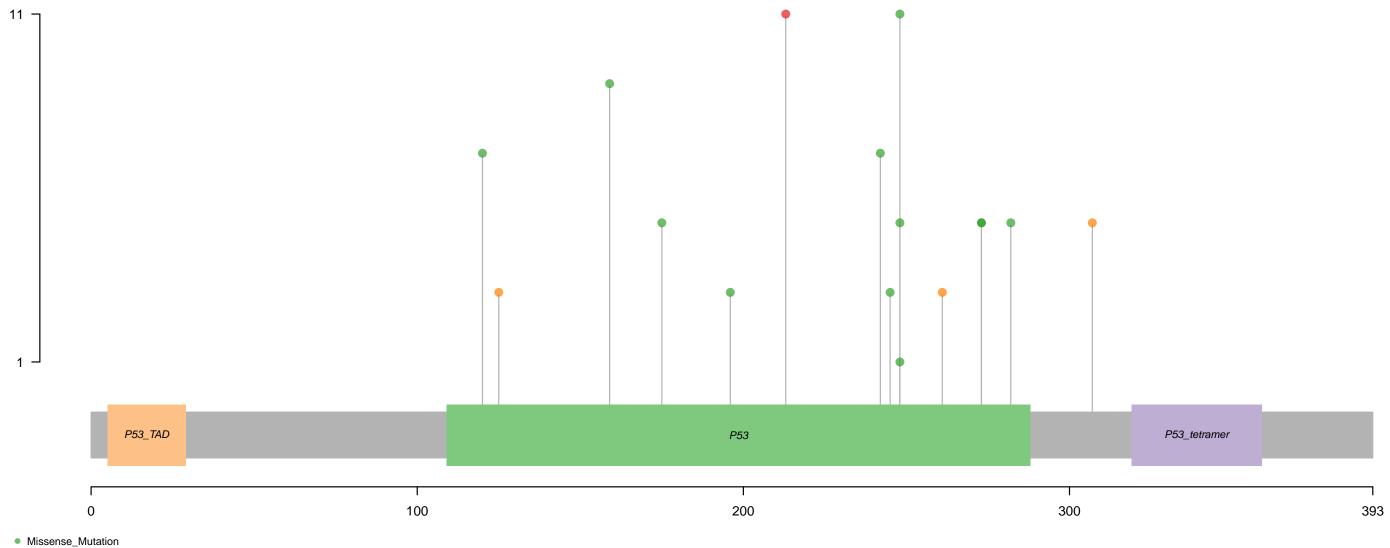
In_Frame_Del

Missense_Mutation

POLR2A: [Somatic Mutation Rate: 2.49%] NM_000937



TP53 : [Somatic Mutation Rate: 23.05%] NM_000546



Splice_Site

Nonsense_Mutation