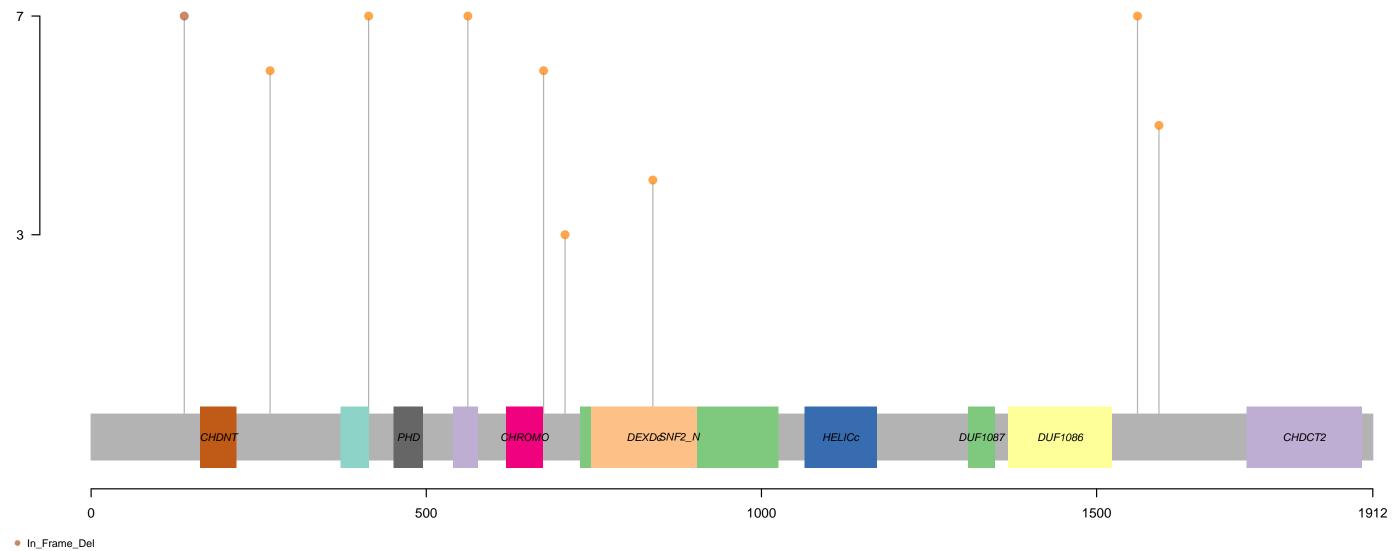
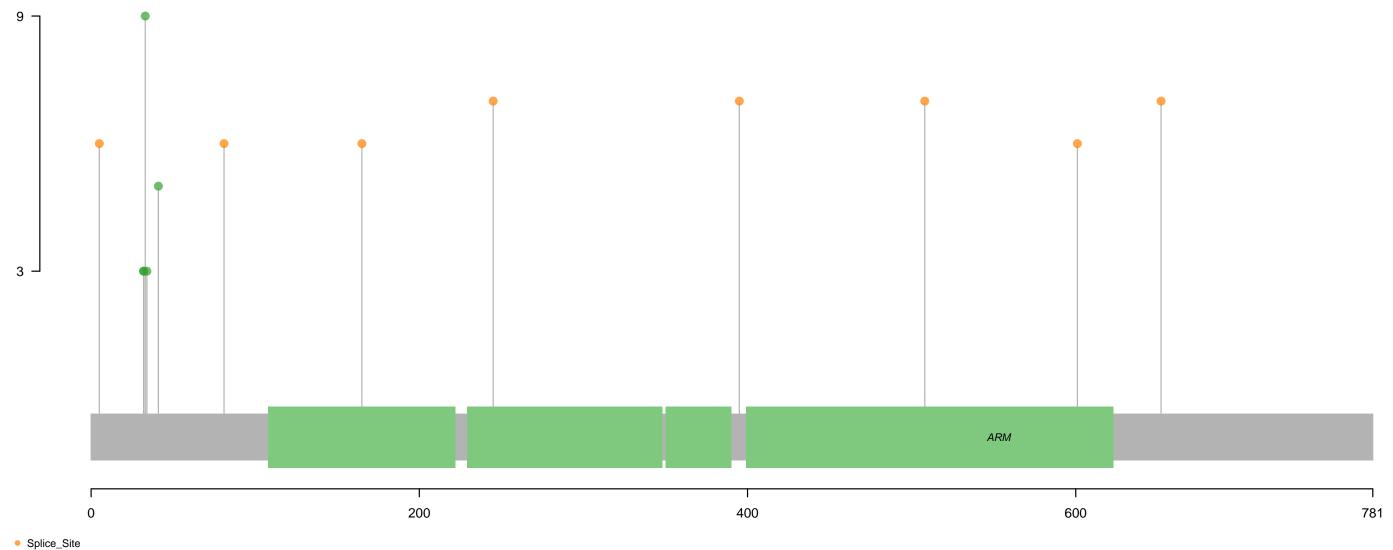
CHD4 : [Somatic Mutation Rate: 5.19%] NM_001273

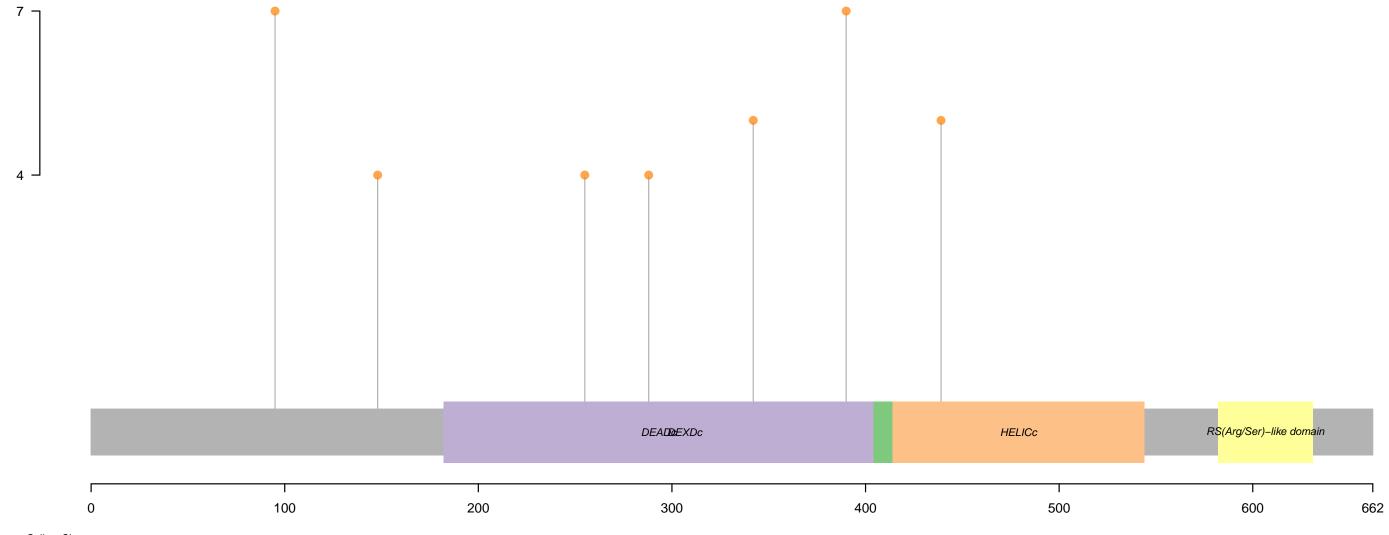


Splice_Site



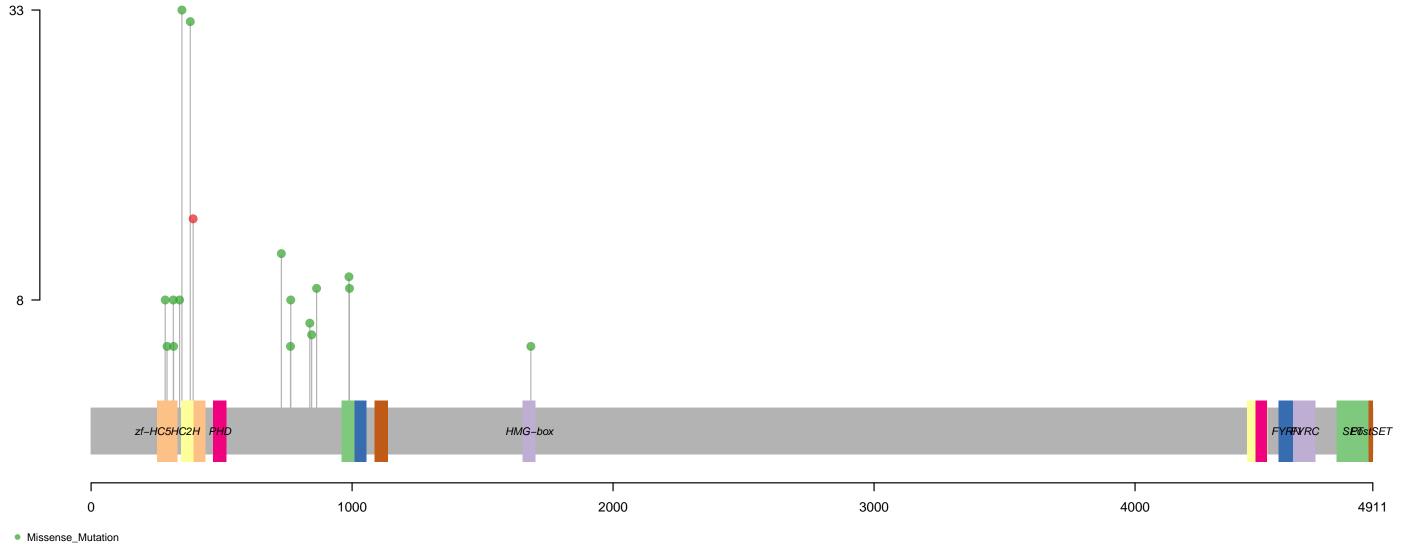
Missense_Mutation

DDX3X : [Somatic Mutation Rate: 3.11%] NM_001356



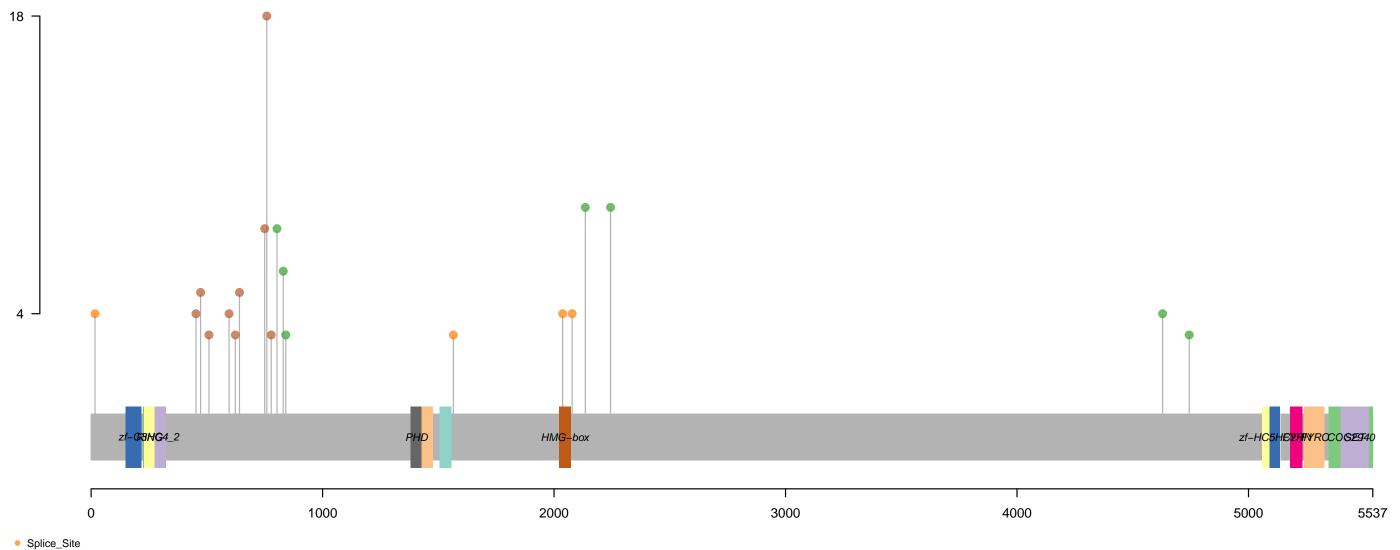
Splice_Site

KMT2C : [Somatic Mutation Rate: 51.56%] NM_170606



Nonsense_Mutation

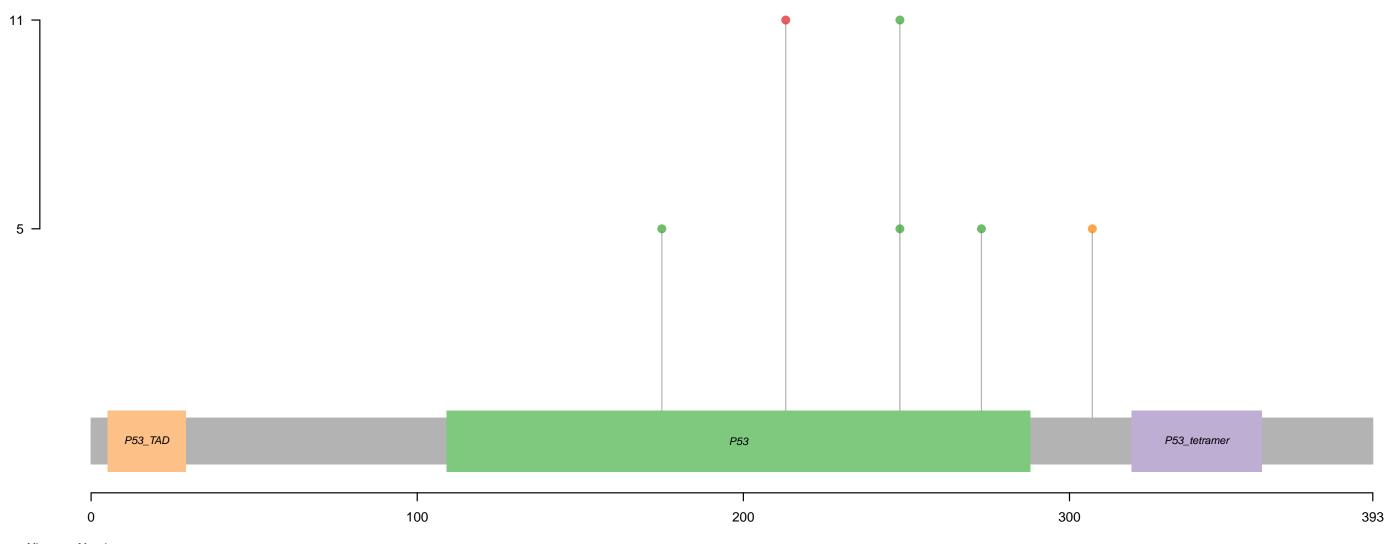
KMT2D: [Somatic Mutation Rate: 31.14%] NM_003482



In_Frame_Del

Missense_Mutation

TP53 : [Somatic Mutation Rate: 14.19%] NM_000546



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site