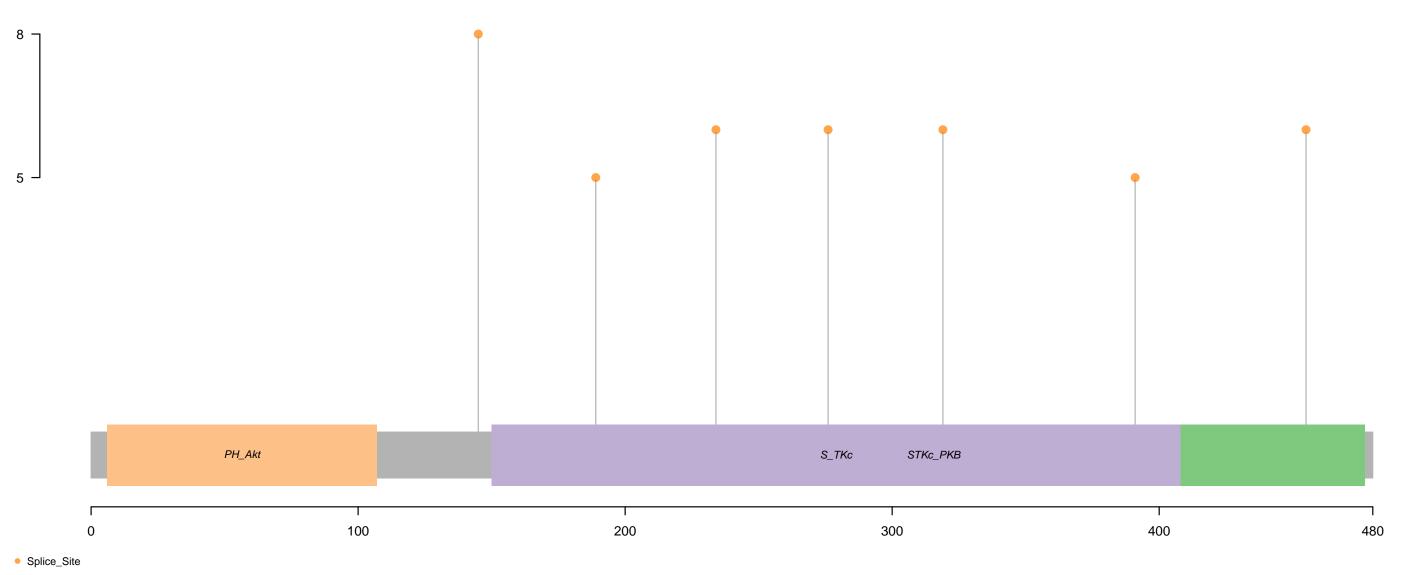
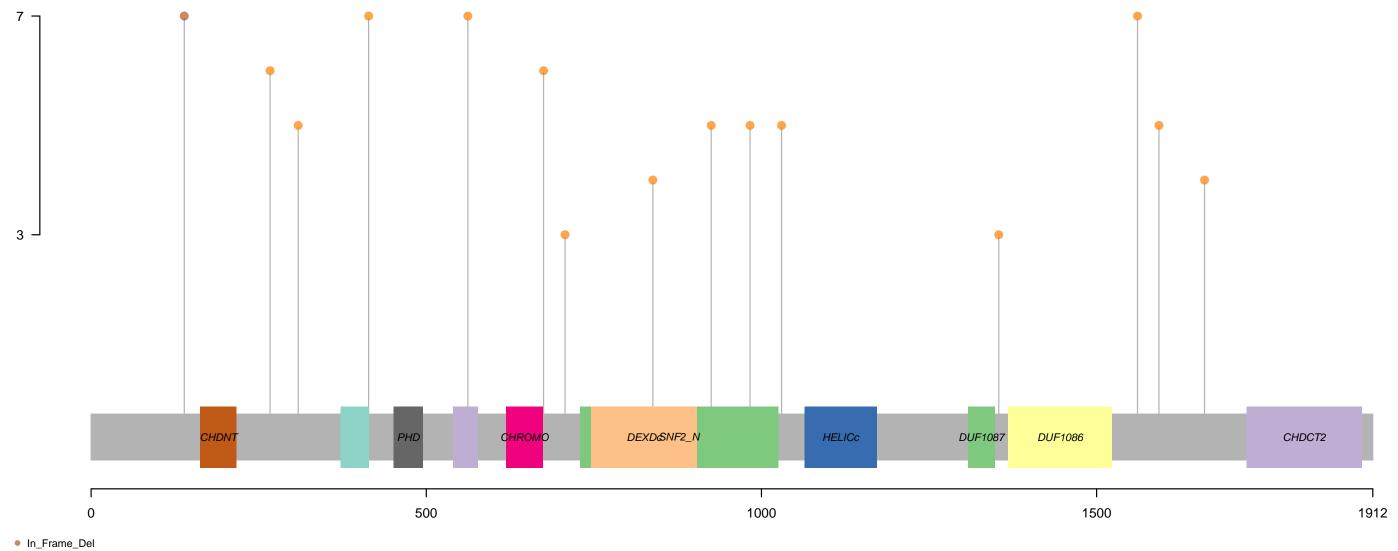
AKT1 : [Somatic Mutation Rate: 2.72%] NM\_005163

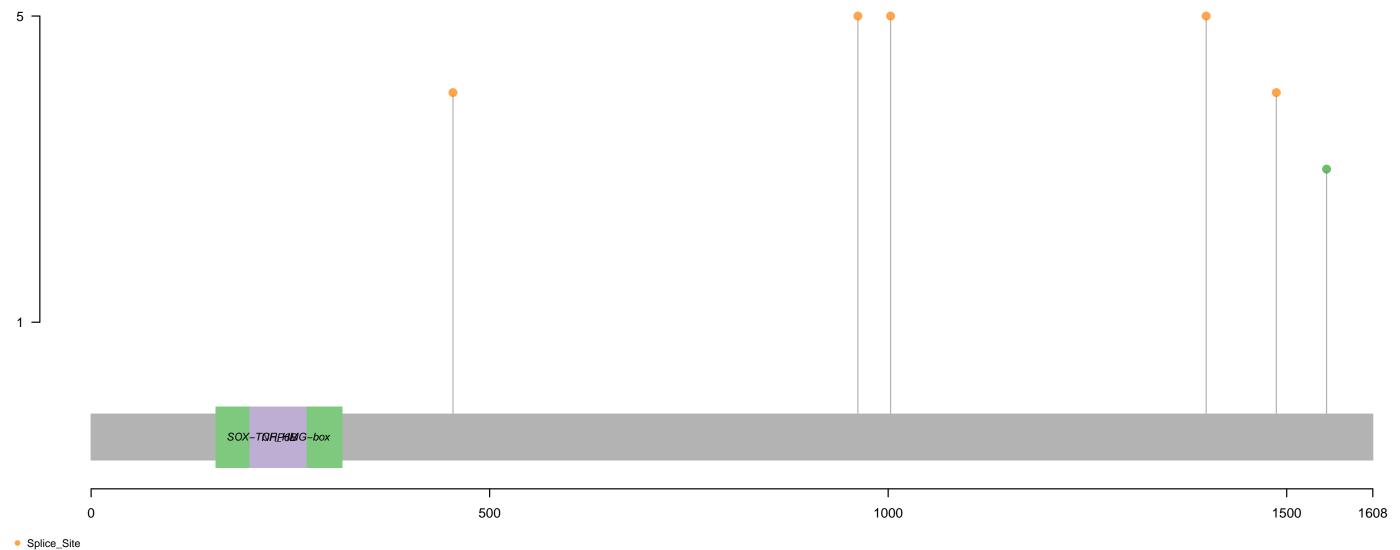


CHD4 : [Somatic Mutation Rate: 5.1%] NM\_001273

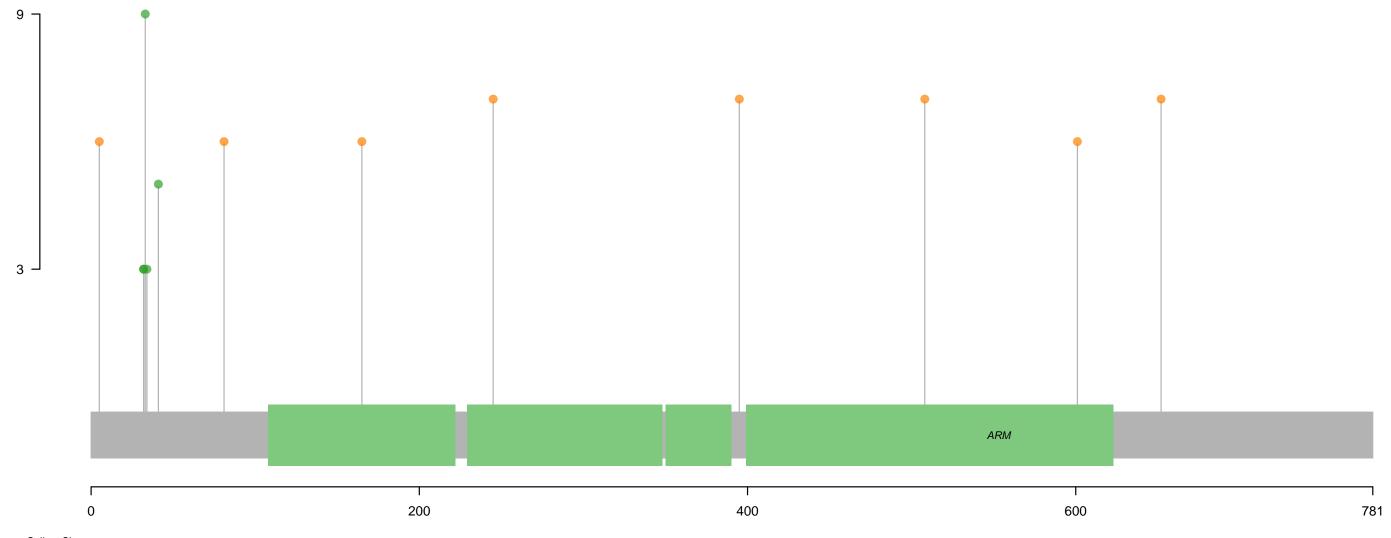


Splice\_Site

CIC: [Somatic Mutation Rate: 3.4%] NM\_015125



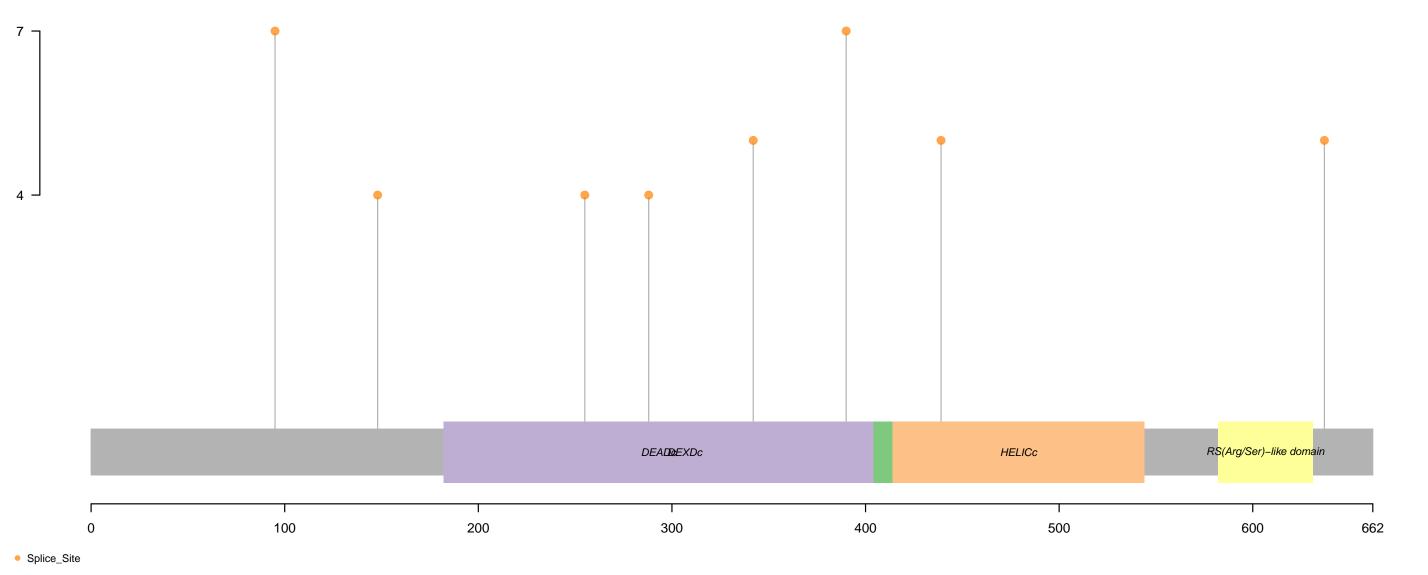
Missense\_Mutation



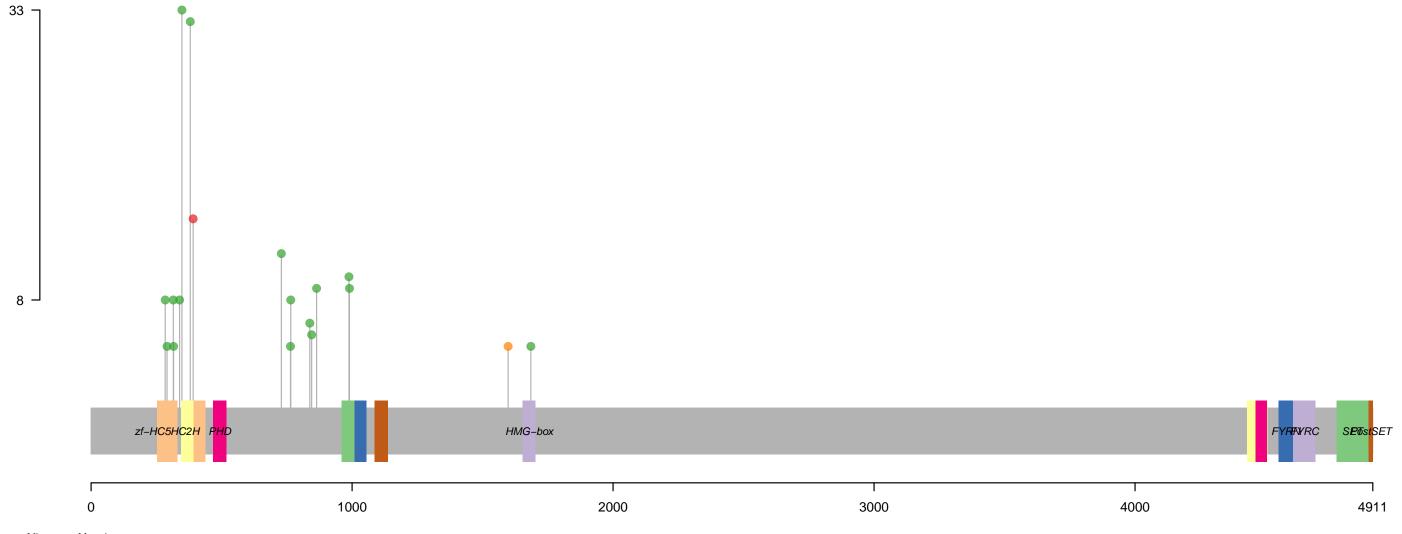
Splice\_Site

Missense\_Mutation

DDX3X : [Somatic Mutation Rate: 3.06%] NM\_001356

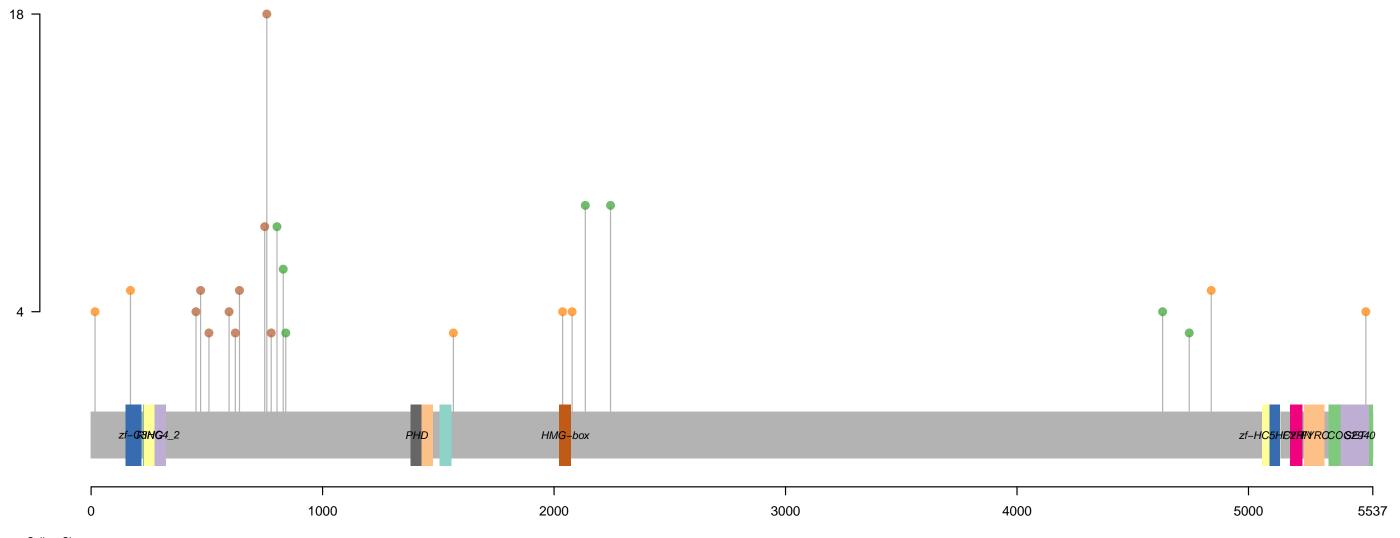


KMT2C : [Somatic Mutation Rate: 52.04%] NM\_170606



- Missense\_Mutation
- Nonsense\_Mutation
- Splice\_Site

KMT2D : [Somatic Mutation Rate: 30.95%] NM\_003482

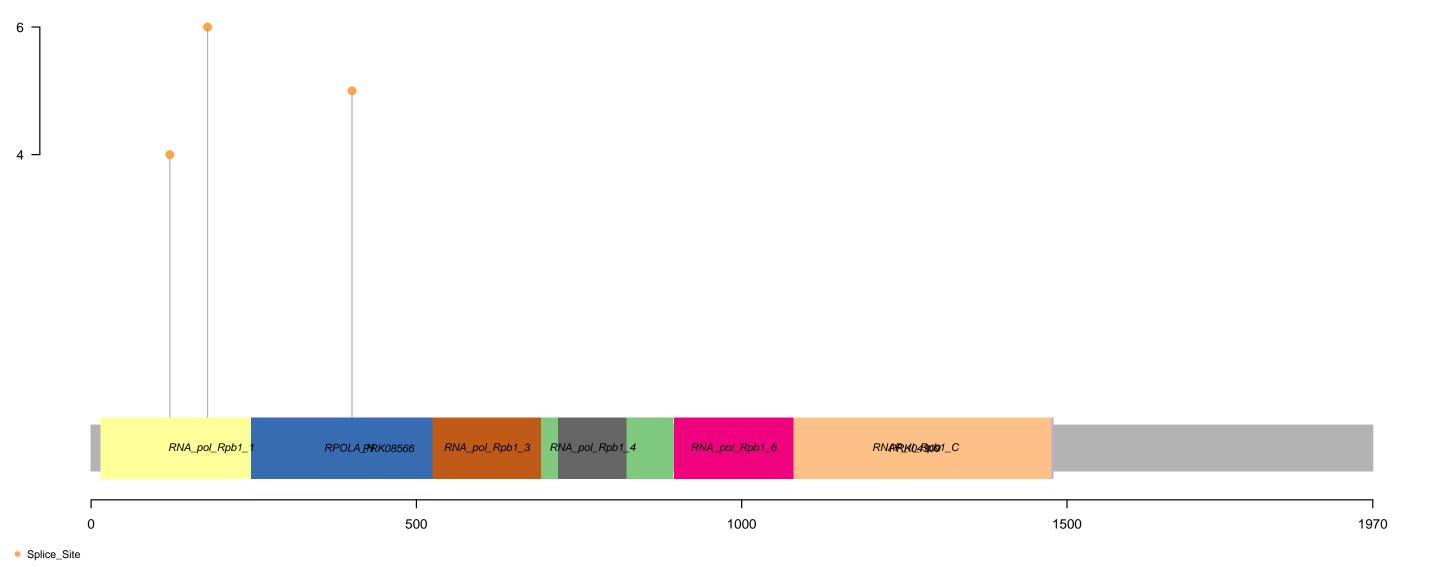


Splice\_Site

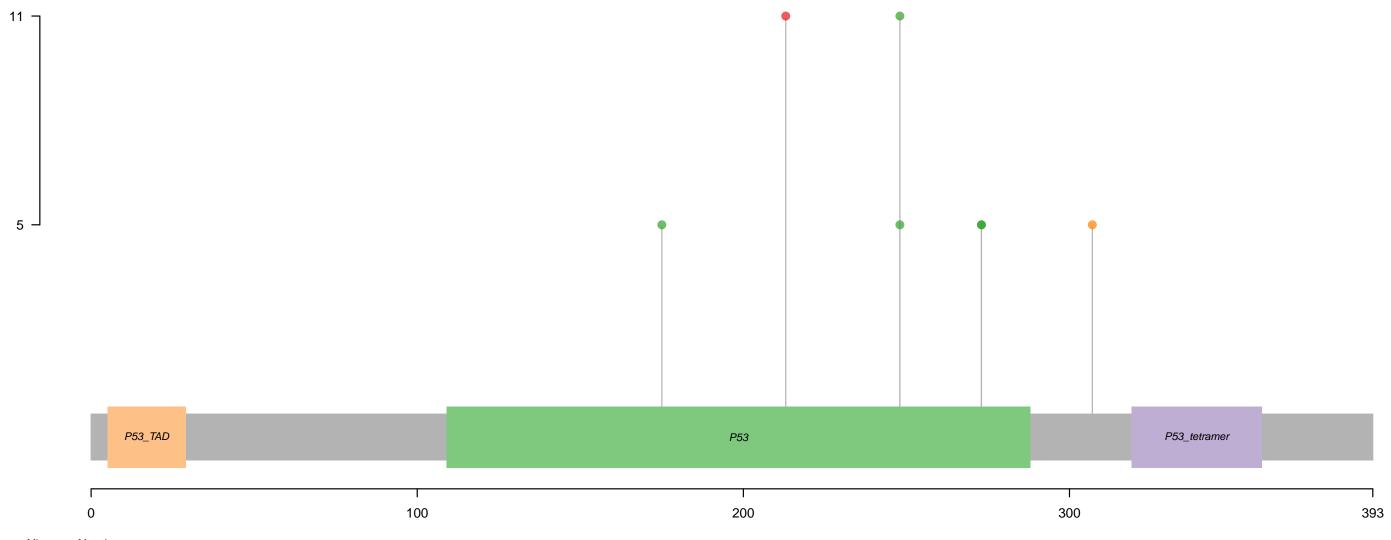
In\_Frame\_Del

Missense\_Mutation

POLR2A: [Somatic Mutation Rate: 2.38%] NM\_000937



*TP53* : [Somatic Mutation Rate: 15.65%] NM\_000546



- Missense\_Mutation
- Nonsense\_Mutation
- Splice\_Site