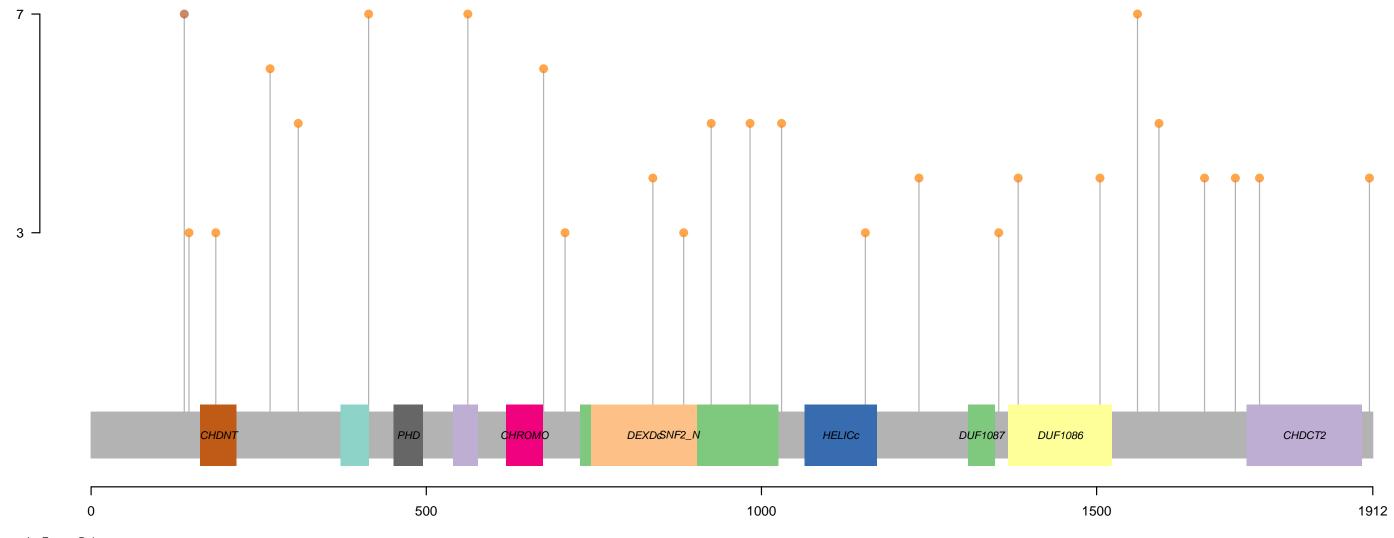
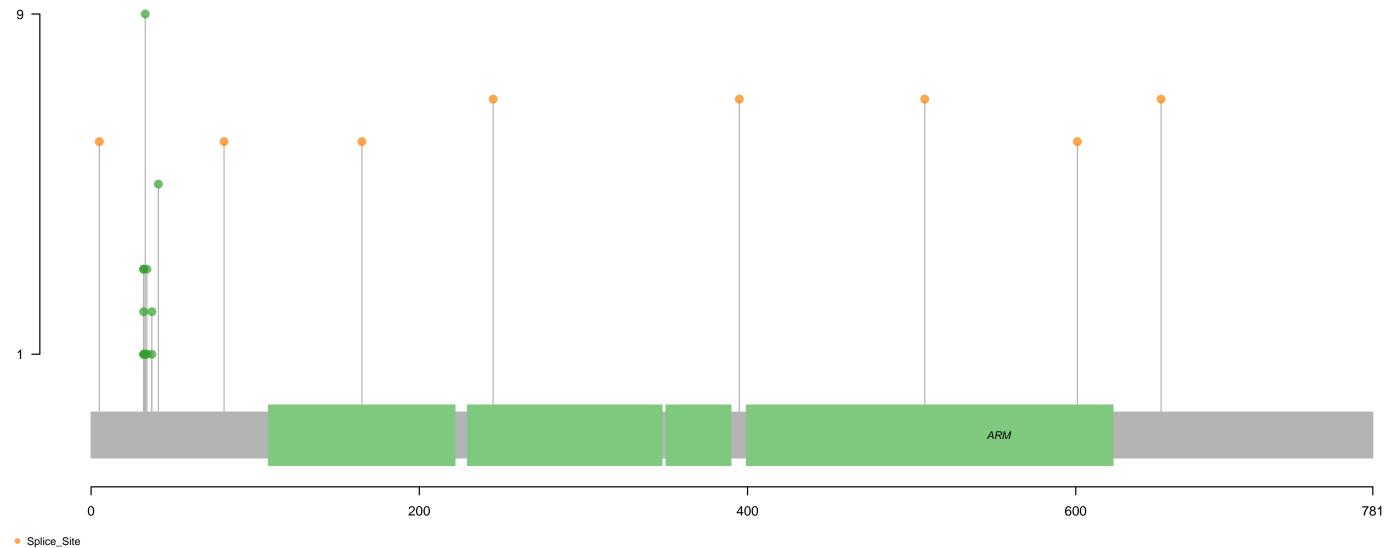
CHD4: [Somatic Mutation Rate: 4.67%] NM\_001273



In\_Frame\_Del

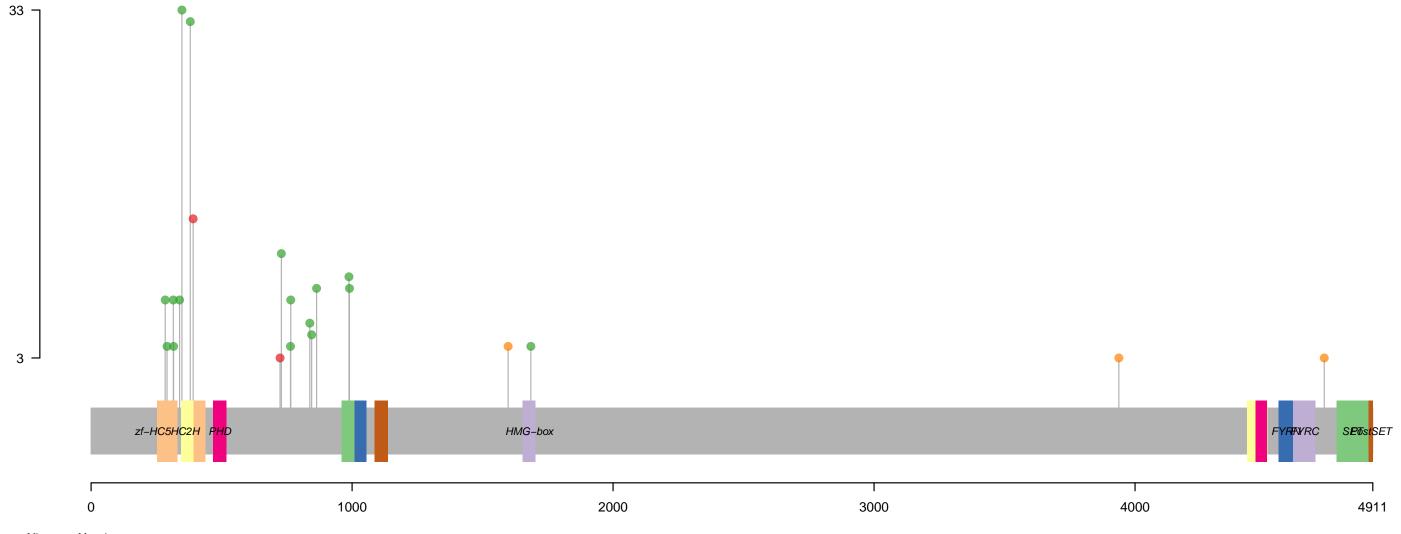
Splice\_Site

CTNNB1 : [Somatic Mutation Rate: 12.77%] NM\_001098210



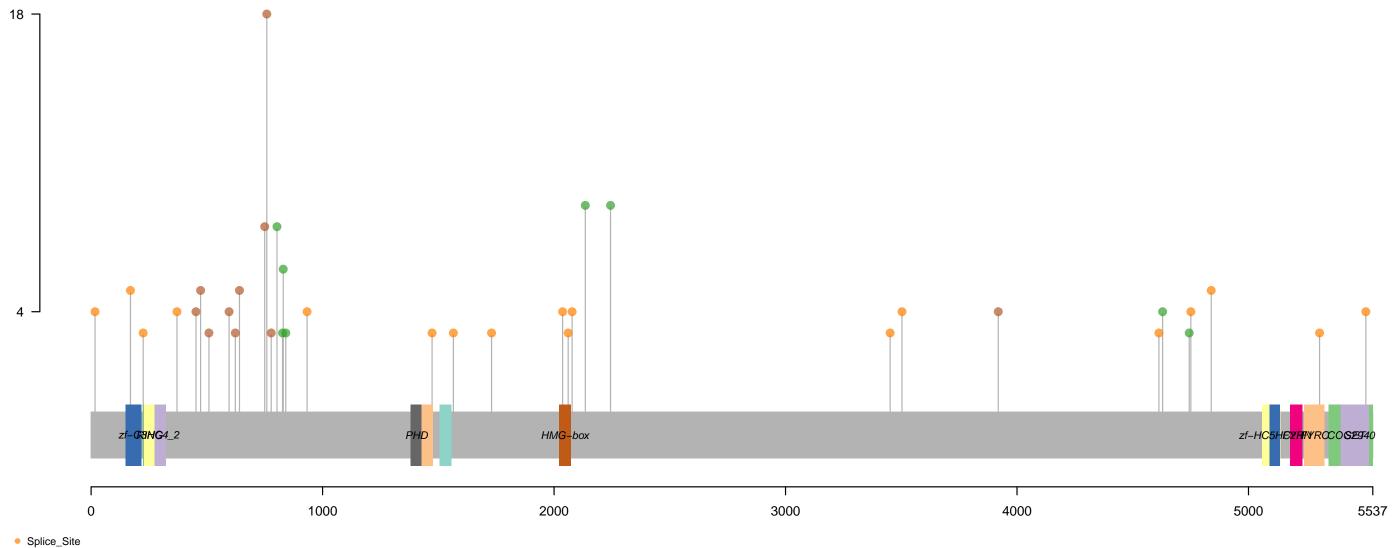
Missense\_Mutation

KMT2C : [Somatic Mutation Rate: 48.6%] NM\_170606



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

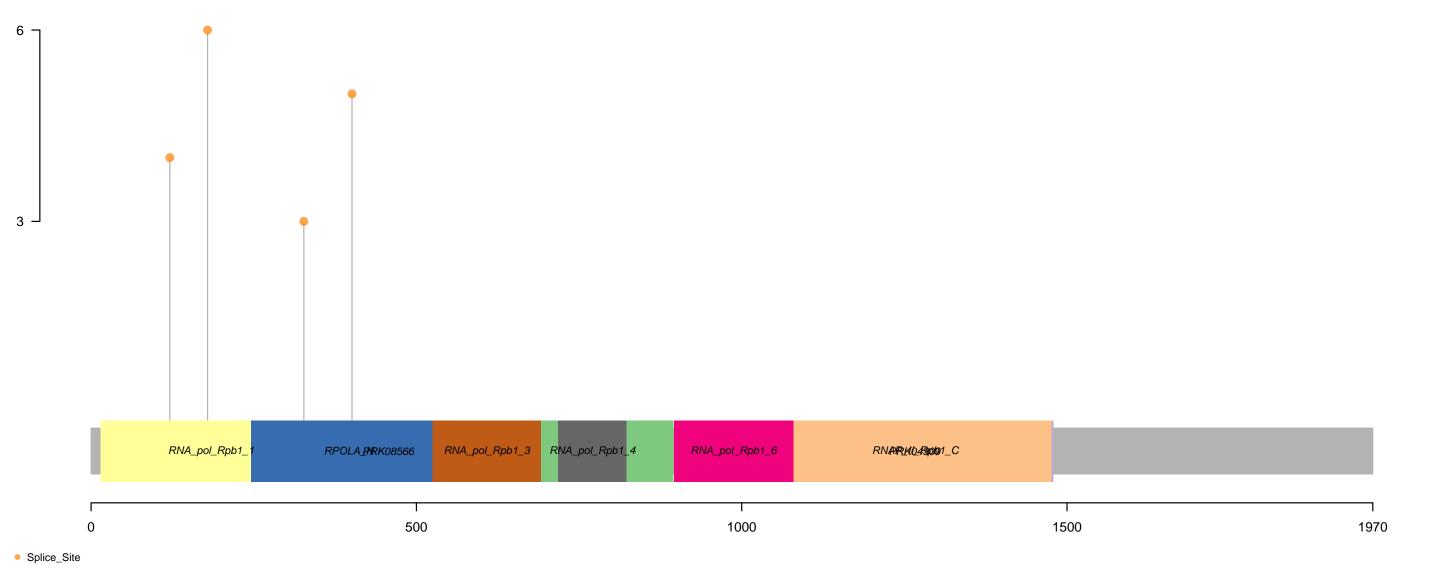
KMT2D : [Somatic Mutation Rate: 30.22%] NM\_003482



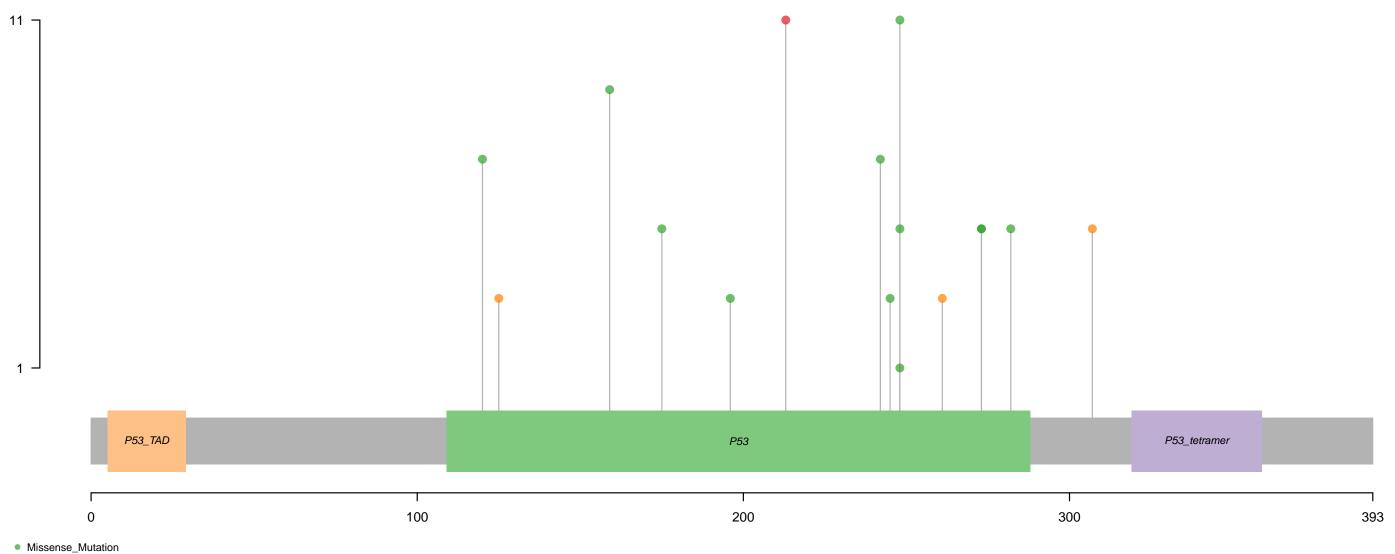
In\_Frame\_Del

Missense\_Mutation

POLR2A: [Somatic Mutation Rate: 2.49%] NM\_000937



*TP53* : [Somatic Mutation Rate: 23.36%] NM\_000546



Splice\_Site

Nonsense\_Mutation