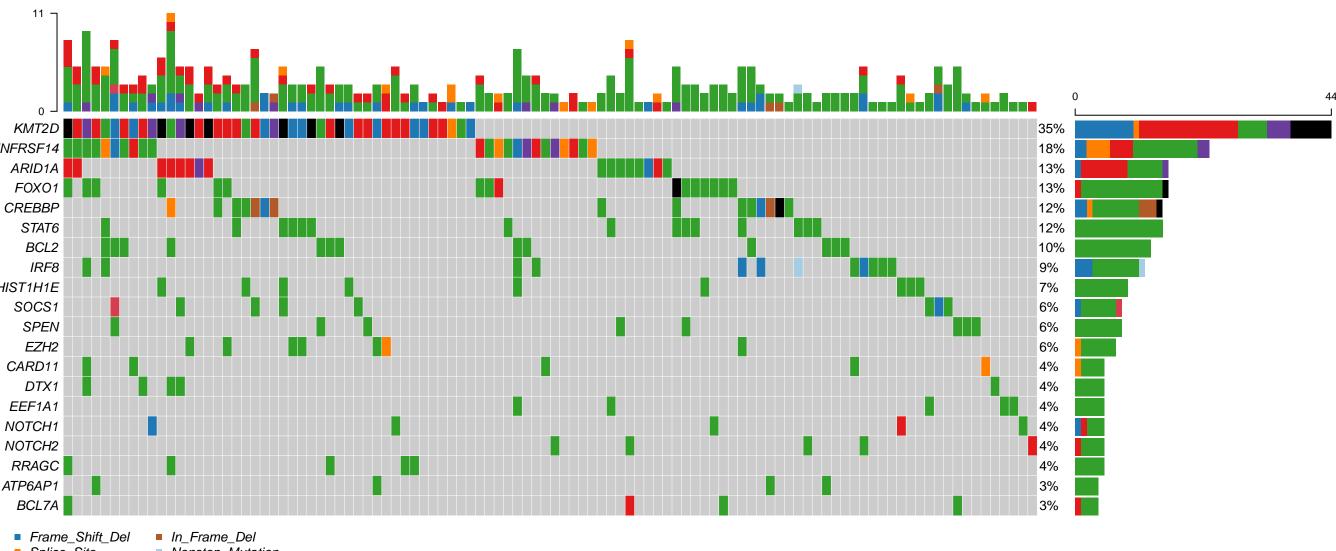
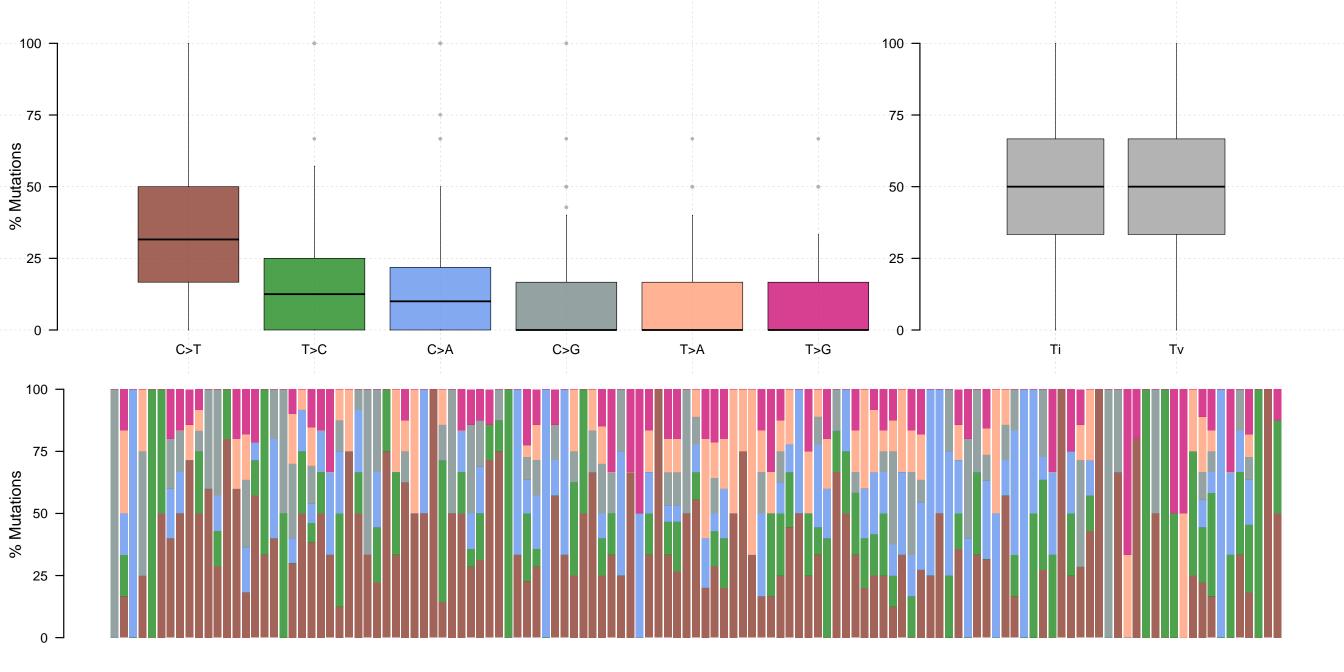


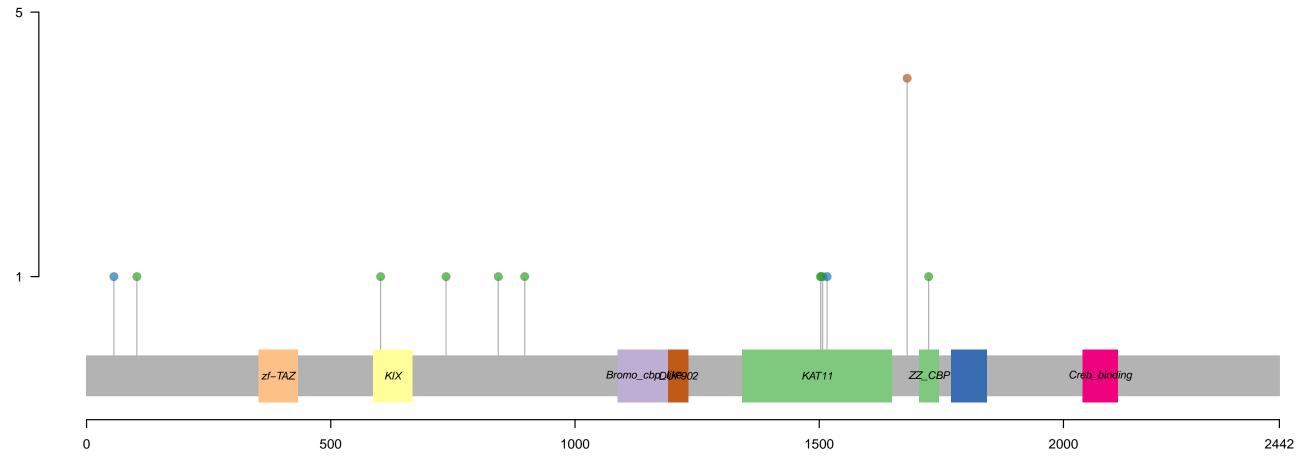
Altered in 104 (81.89%) of 127 samples.



Splice\_Site Nonstop\_Mutation ■ Nonsense\_Mutation ■ In\_Frame\_Ins ■ Missense\_Mutation ■ Multi\_Hit

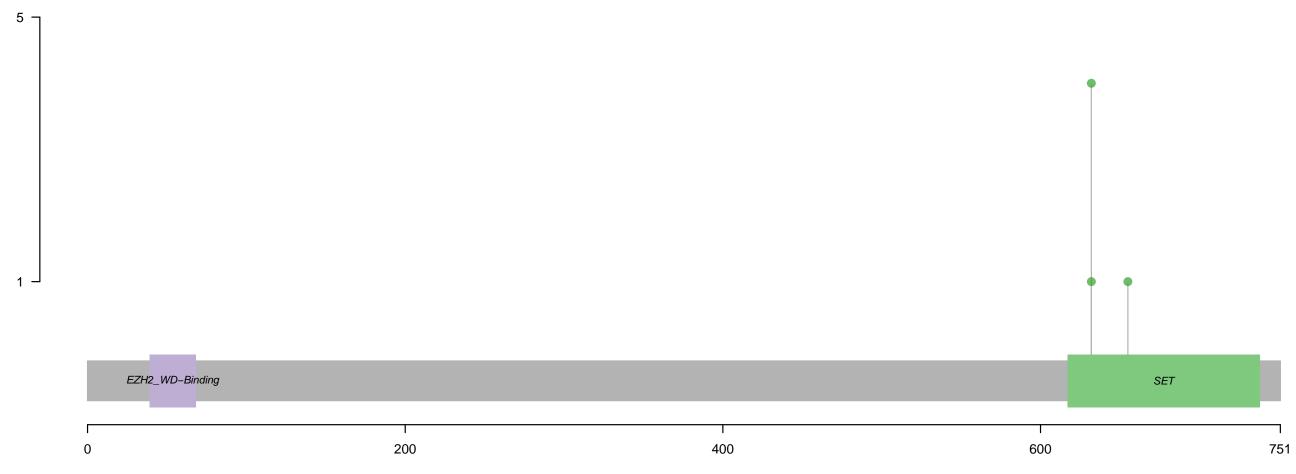
Frame\_Shift\_Ins



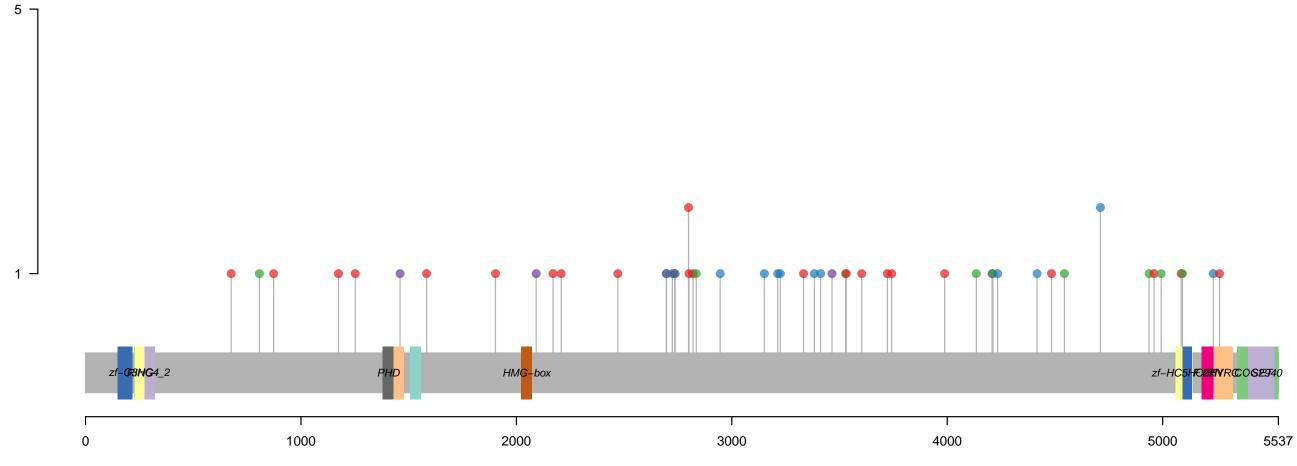


- Frame\_Shift\_Del
- Missense\_Mutation
- In\_Frame\_Del

EZH2: [Somatic Mutation Rate: 5.51%] NM\_004456



Missense\_Mutation



- Nonsense\_Mutation
- Missense\_Mutation
- Frame\_Shift\_Ins
- Frame\_Shift\_Del