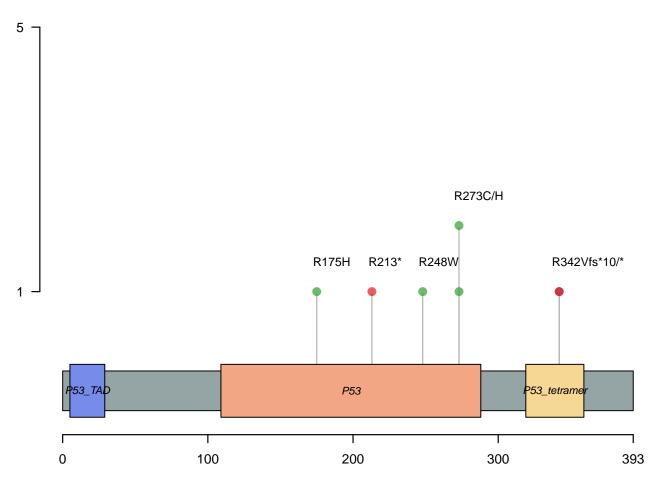
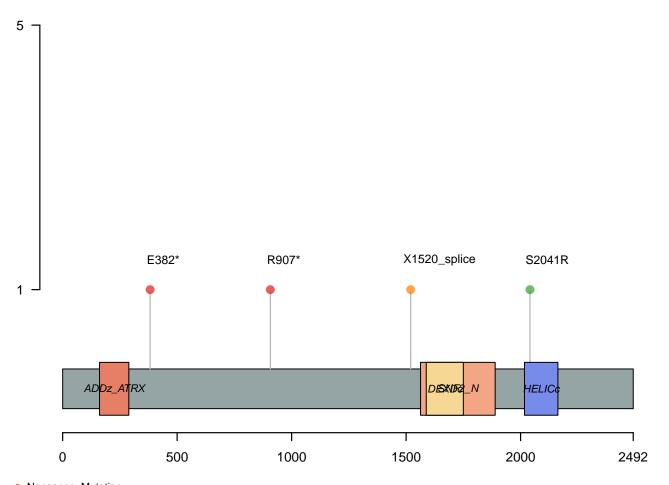
TP53 : [Somatic Mutation Rate: 54.55%] NM_000546



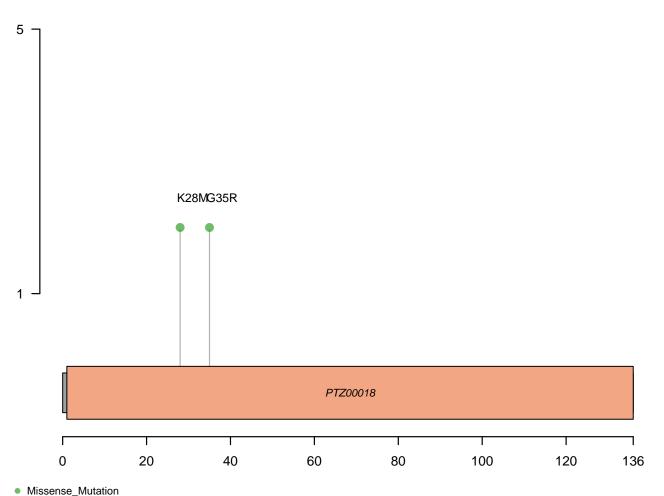
- Missense_Mutation
- Nonsense_Mutation
- Frame_Shift_Ins

ATRX : [Somatic Mutation Rate: 36.36%] NM_000489

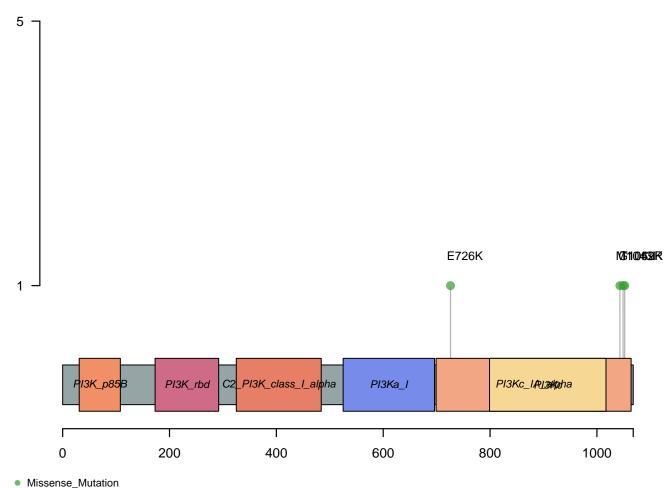


- Nonsense_Mutation
- Splice_Site
- Missense_Mutation

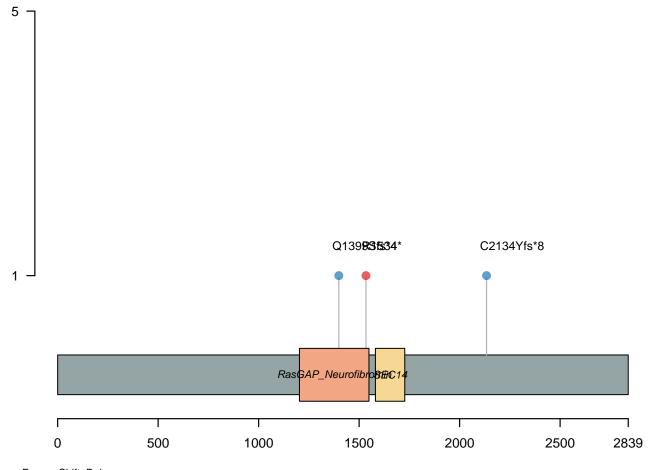
H3F3A : [Somatic Mutation Rate: 36.36%] NM_002107



PIK3CA: [Somatic Mutation Rate: 27.27%] NM_006218

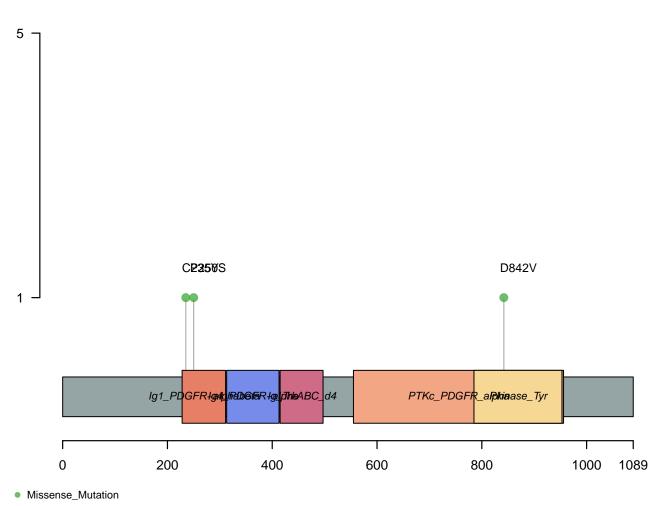


NF1 : [Somatic Mutation Rate: 18.18%] NM_001042492

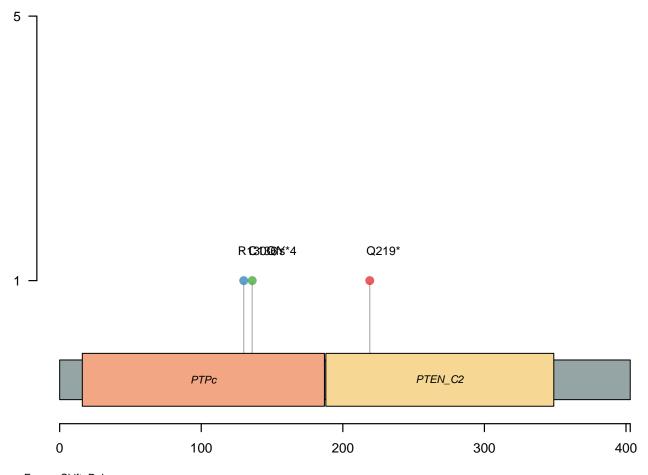


- Frame_Shift_Del
- Nonsense_Mutation

PDGFRA: [Somatic Mutation Rate: 18.18%] NM_006206

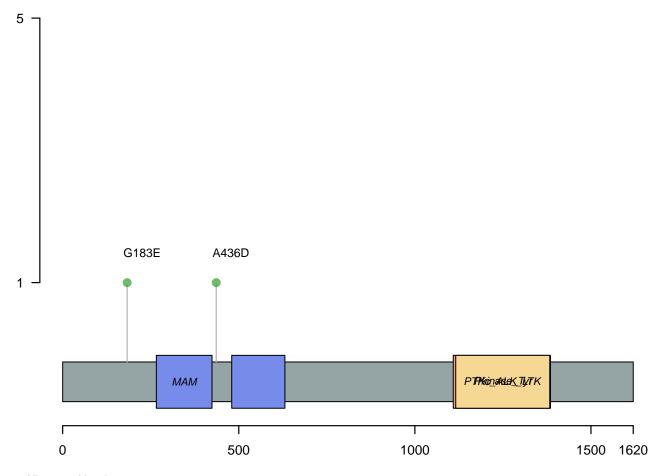


PTEN: [Somatic Mutation Rate: 18.18%] NM_000314



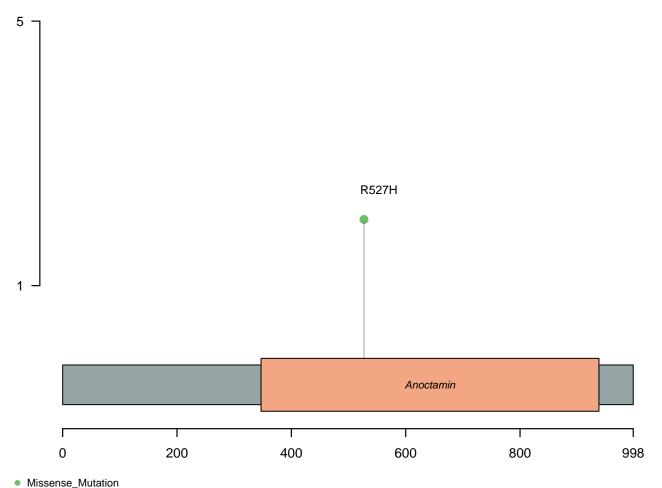
- Frame_Shift_Del
- Missense_Mutation
- Nonsense_Mutation

ALK: [Somatic Mutation Rate: 18.18%] NM_004304

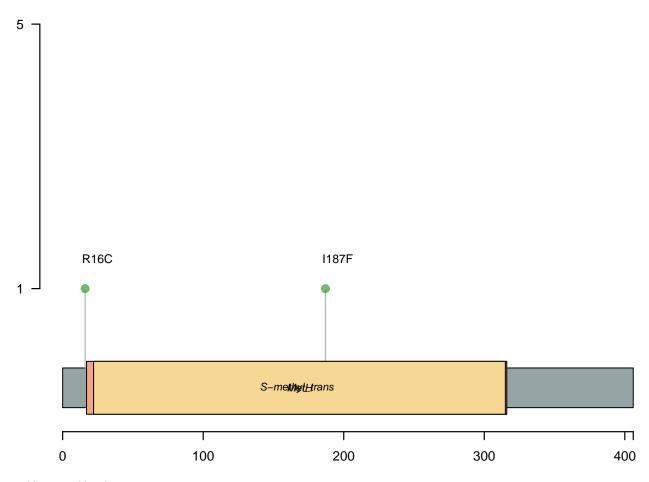


Missense_Mutation

ANO2 : [Somatic Mutation Rate: 18.18%] NM_020373



BHMT: [Somatic Mutation Rate: 18.18%] NM_001713



Missense_Mutation