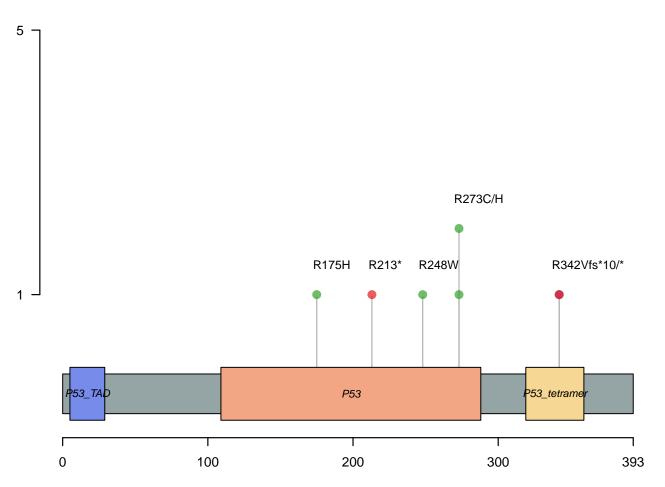
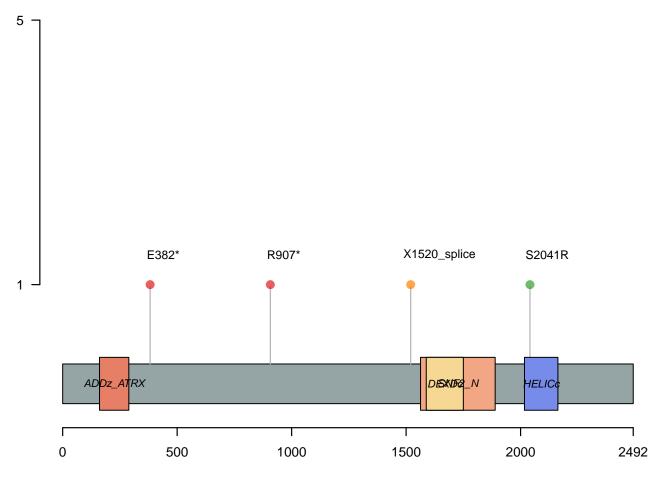
TP53 : [Somatic Mutation Rate: 75%] NM\_000546



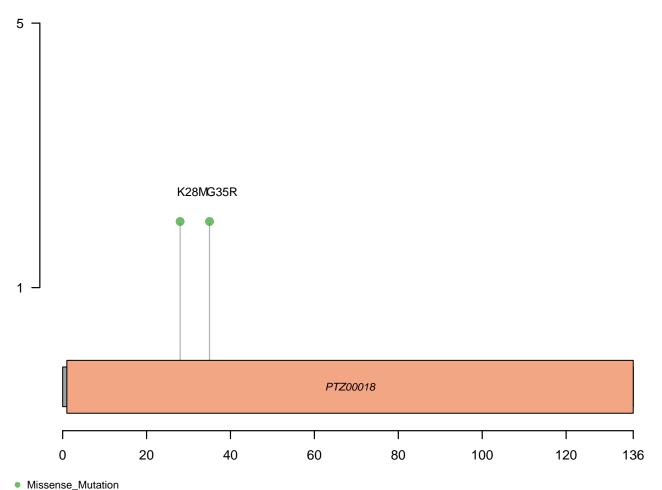
- Missense\_Mutation
- Nonsense\_Mutation
- Frame\_Shift\_Ins

ATRX: [Somatic Mutation Rate: 50%] NM\_000489

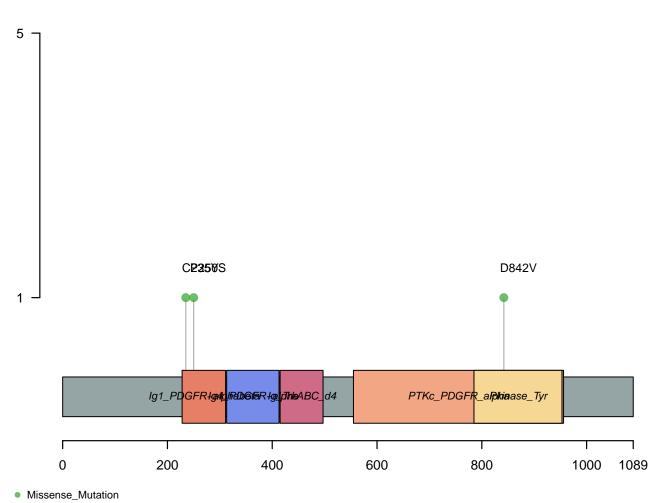


- Nonsense\_Mutation
- Splice\_Site
- Missense\_Mutation

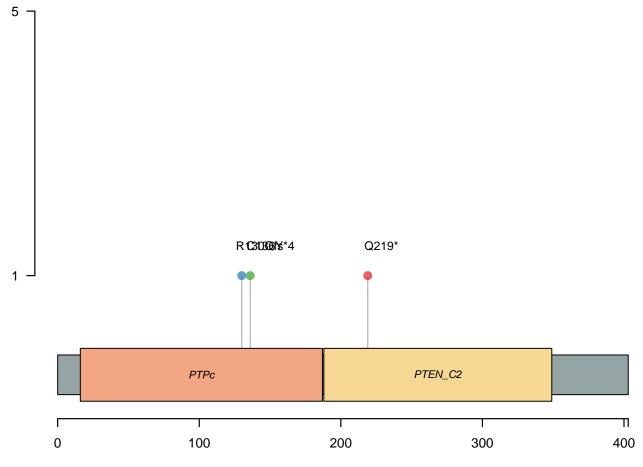
H3F3A: [Somatic Mutation Rate: 50%] NM\_002107



PDGFRA: [Somatic Mutation Rate: 25%] NM\_006206

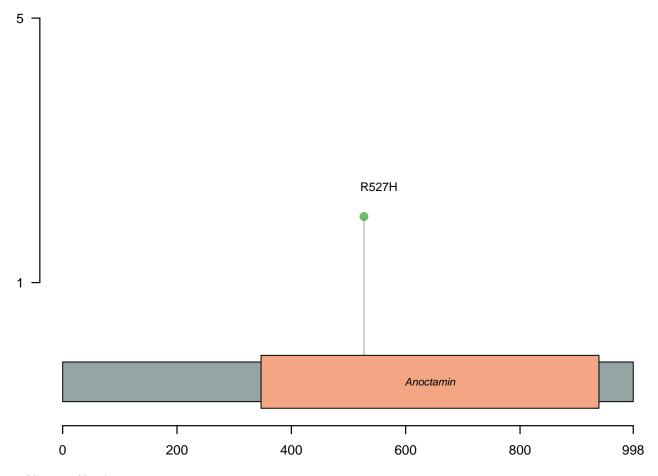


PTEN: [Somatic Mutation Rate: 25%] NM\_000314

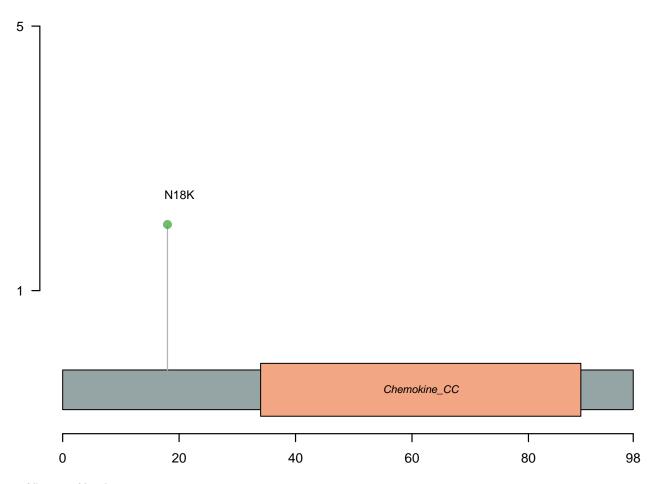


- Frame\_Shift\_Del
- Missense\_Mutation
- Nonsense\_Mutation

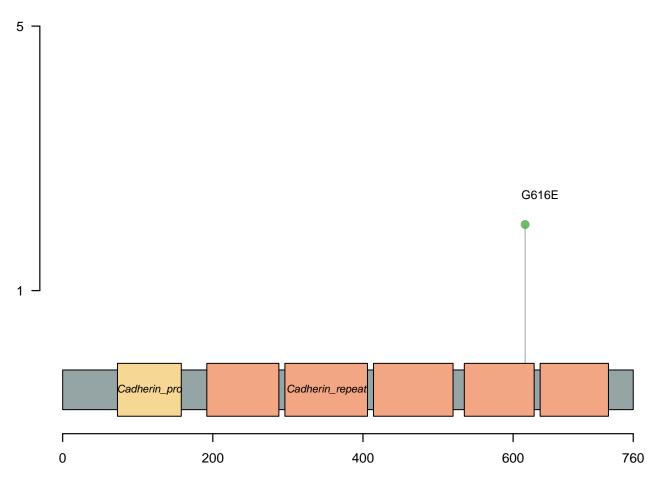
ANO2 : [Somatic Mutation Rate: 25%] NM\_020373



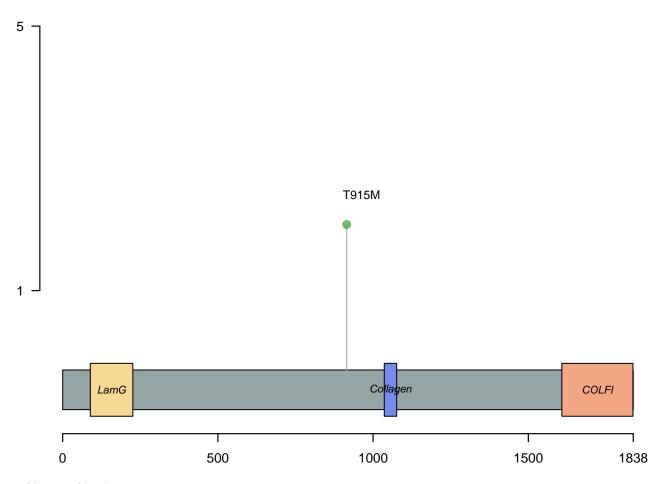
CCL13 : [Somatic Mutation Rate: 25%] NM\_005408



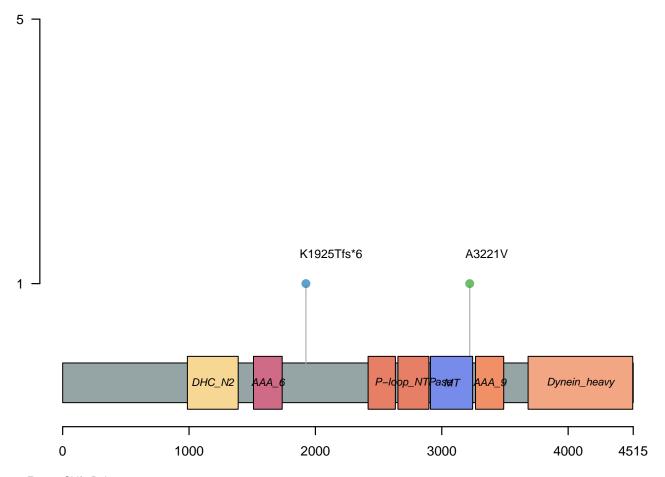
CDH13 : [Somatic Mutation Rate: 25%] NM\_001220488



COL5A1 : [Somatic Mutation Rate: 25%] NM\_000093



DNAH14: [Somatic Mutation Rate: 25%] NM\_001373



- Frame\_Shift\_Del
- Missense\_Mutation