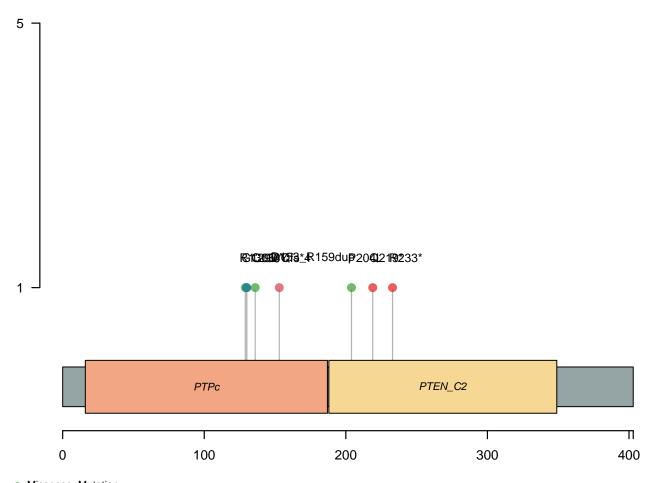
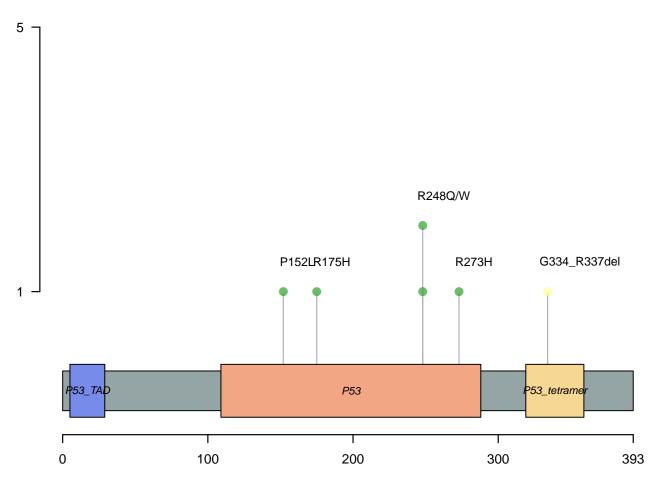
PTEN: [Somatic Mutation Rate: 41.18%] NM_000314



- Missense_Mutation
- Frame_Shift_Del
- In_Frame_Ins
- Nonsense_Mutation

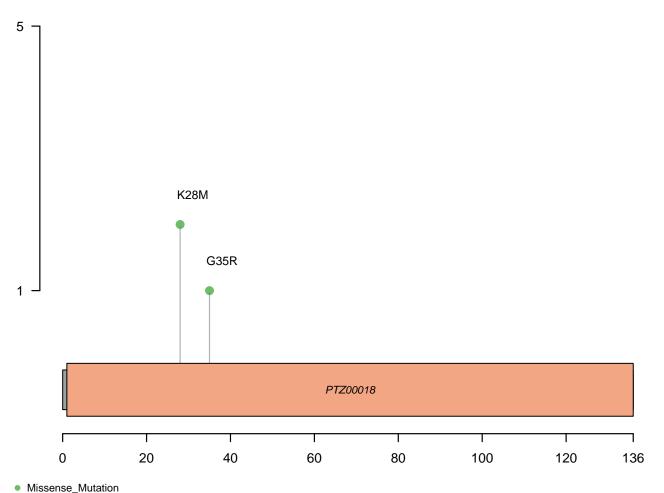
TP53 : [Somatic Mutation Rate: 41.18%] NM_000546



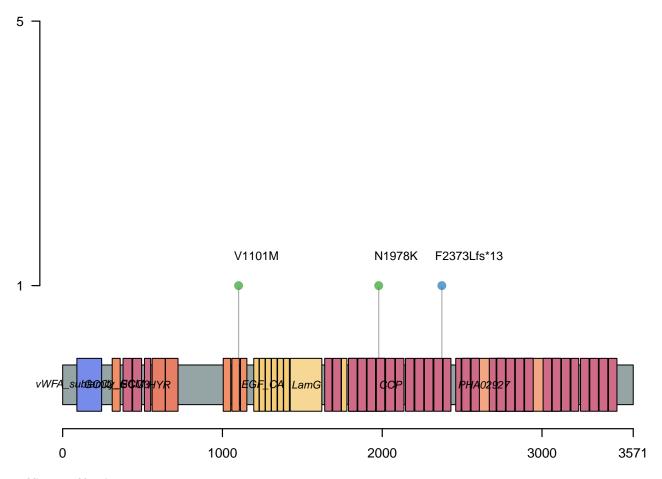
Missense_Mutation

In_Frame_Del

H3F3A: [Somatic Mutation Rate: 17.65%] NM_002107

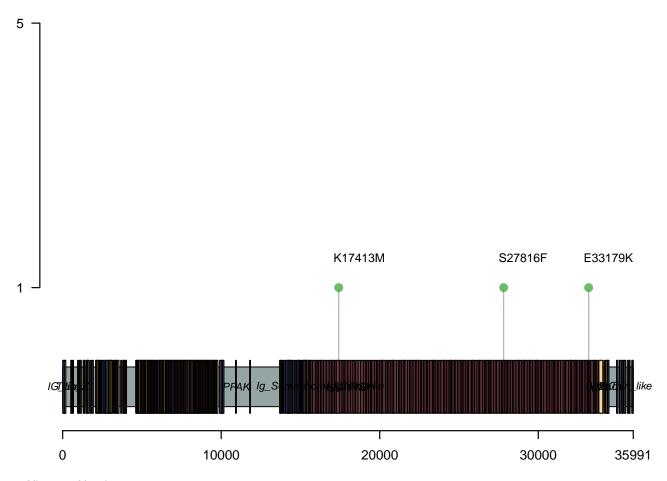


SVEP1 : [Somatic Mutation Rate: 17.65%] NM_153366

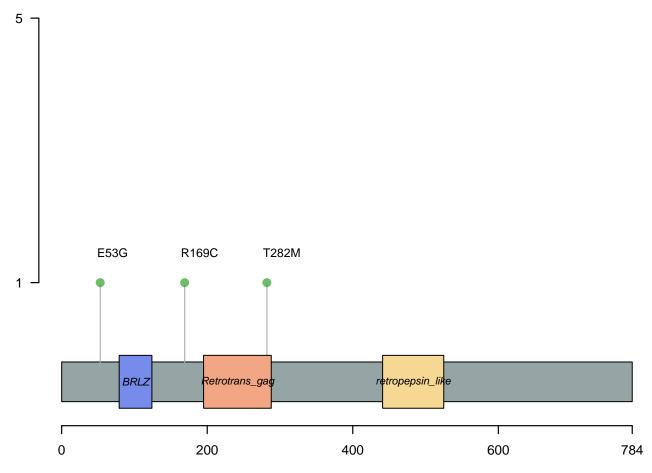


- Missense_Mutation
- Frame_Shift_Del

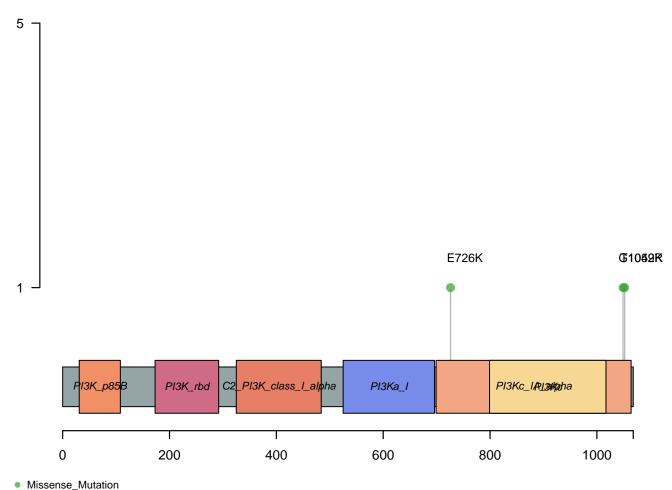
TTN : [Somatic Mutation Rate: 17.65%] NM_001267550



PEG10 : [Somatic Mutation Rate: 11.76%] NM_001172437

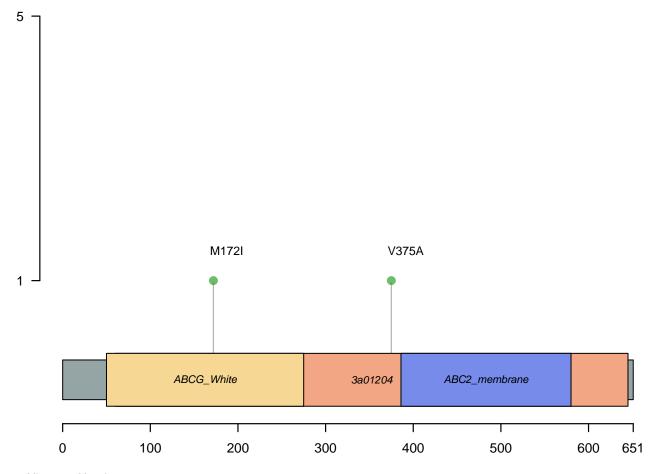


PIK3CA: [Somatic Mutation Rate: 11.76%] NM_006218

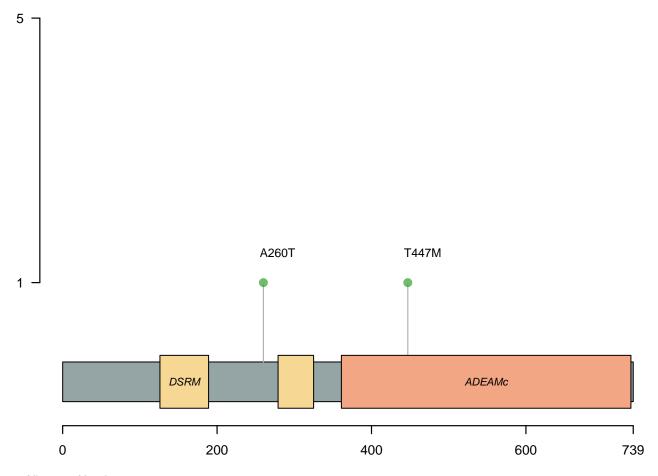


• Wildscride_ividiation

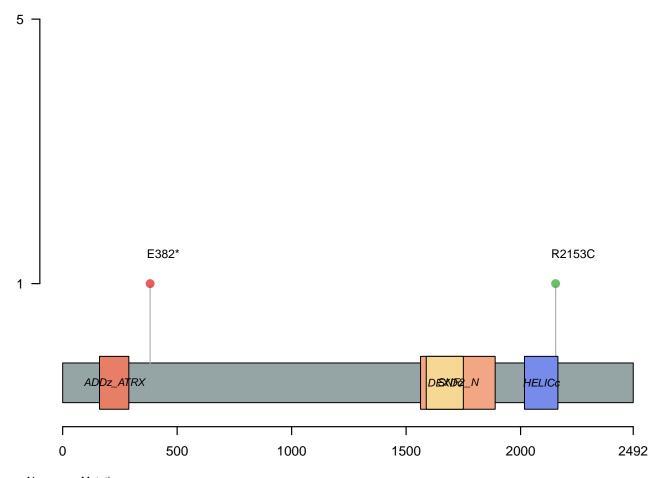
ABCG5 : [Somatic Mutation Rate: 11.76%] NM_022436



ADARB2 : [Somatic Mutation Rate: 11.76%] NM_018702



ATRX : [Somatic Mutation Rate: 11.76%] NM_000489



- Nonsense_Mutation
- Missense_Mutation