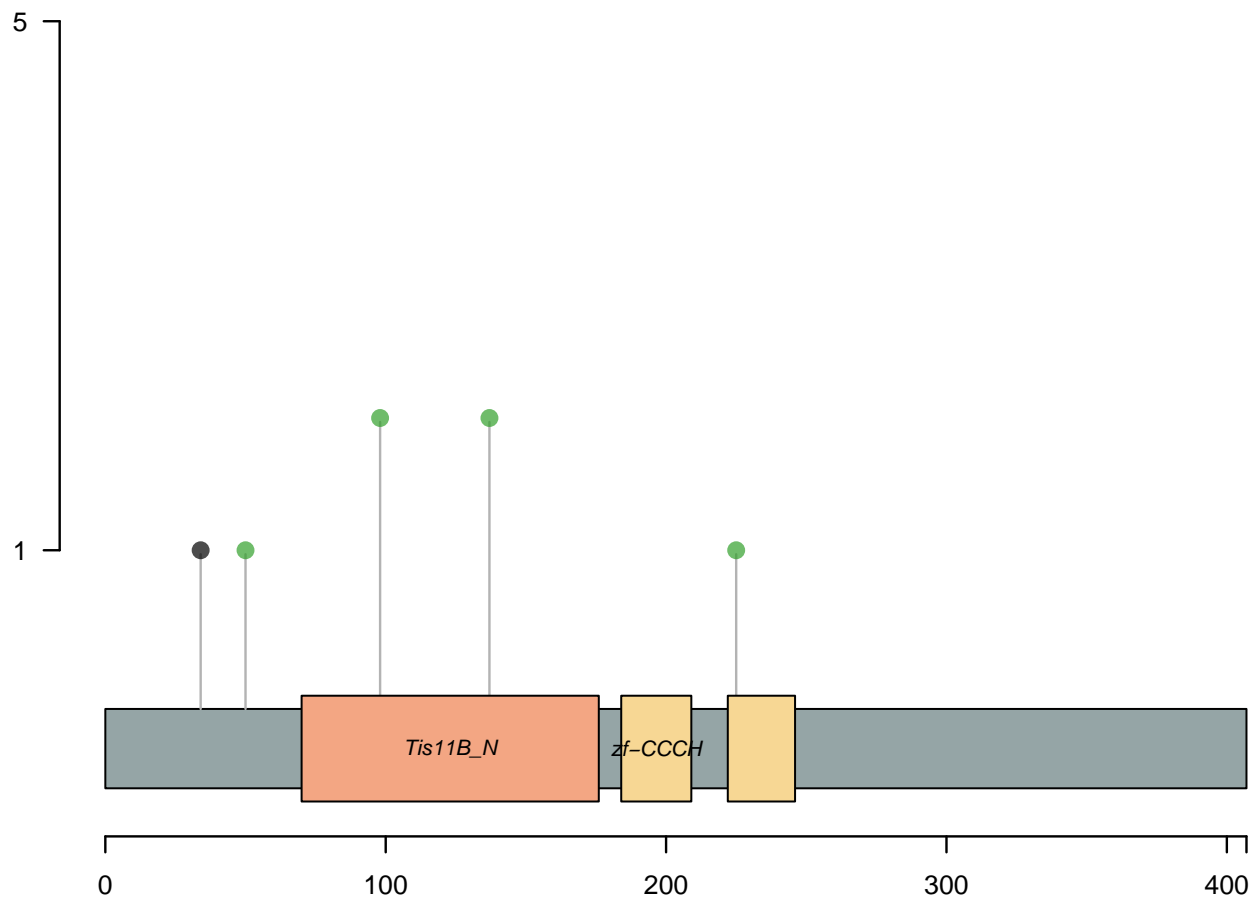


ZFP36L1 : [Somatic Mutation Rate: 3.06%]

NM_001244701



● Truncating

● Missense