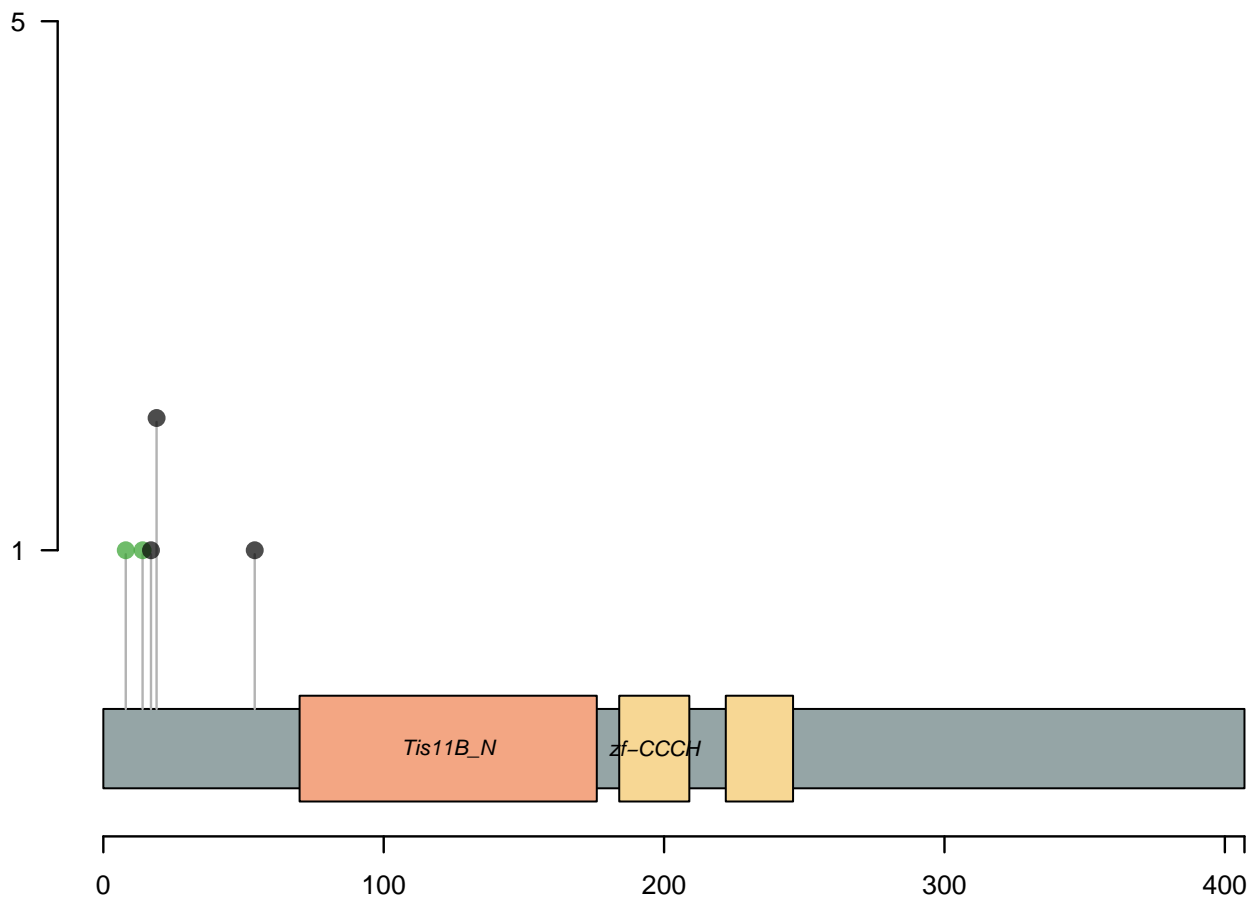


ZFP36L1 : [Somatic Mutation Rate: 2.13%]

NM_001244701



● Missense

● Truncating