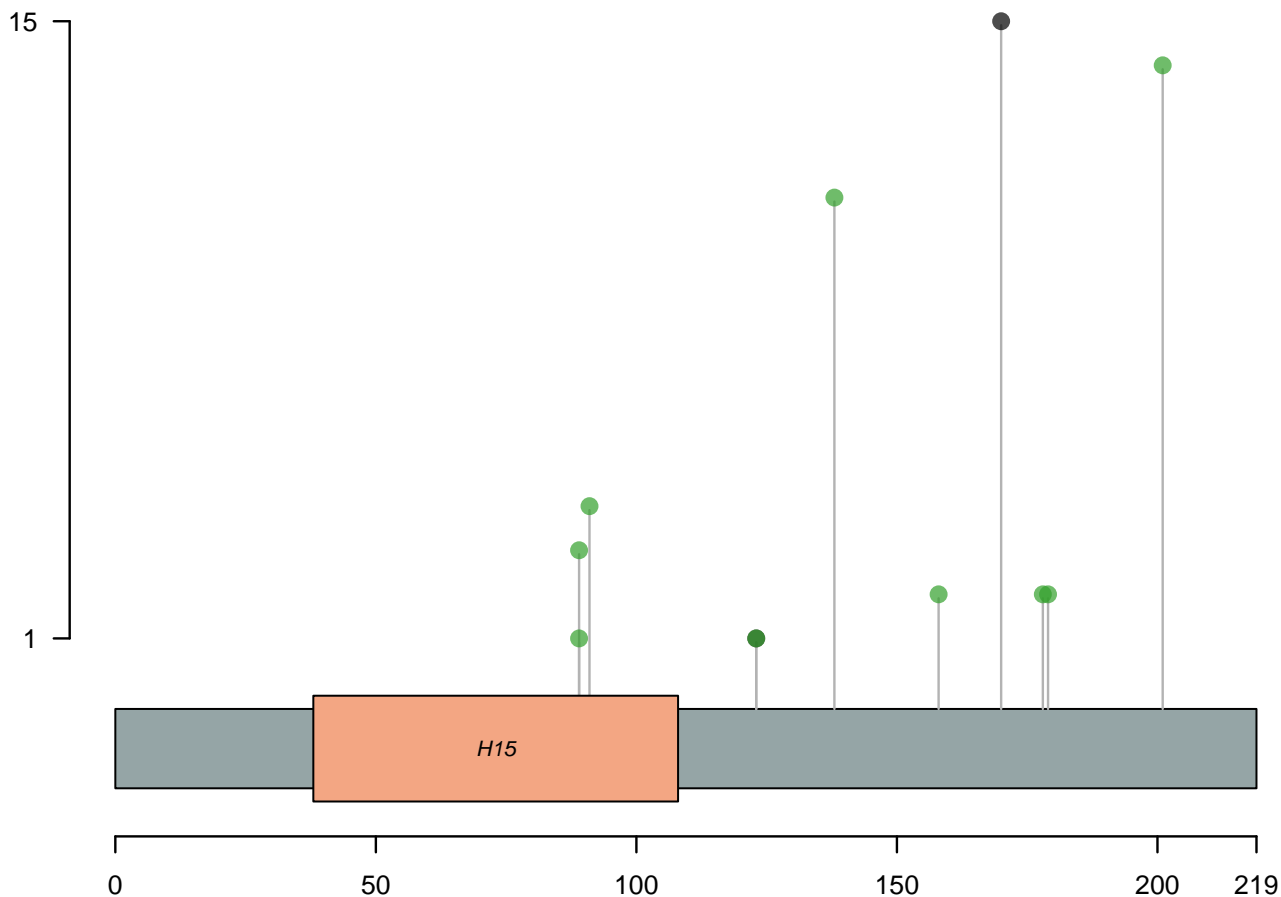


HIST1H1E : [Somatic Mutation Rate: 42.86%]

NM_005321



● Missense

● Truncating