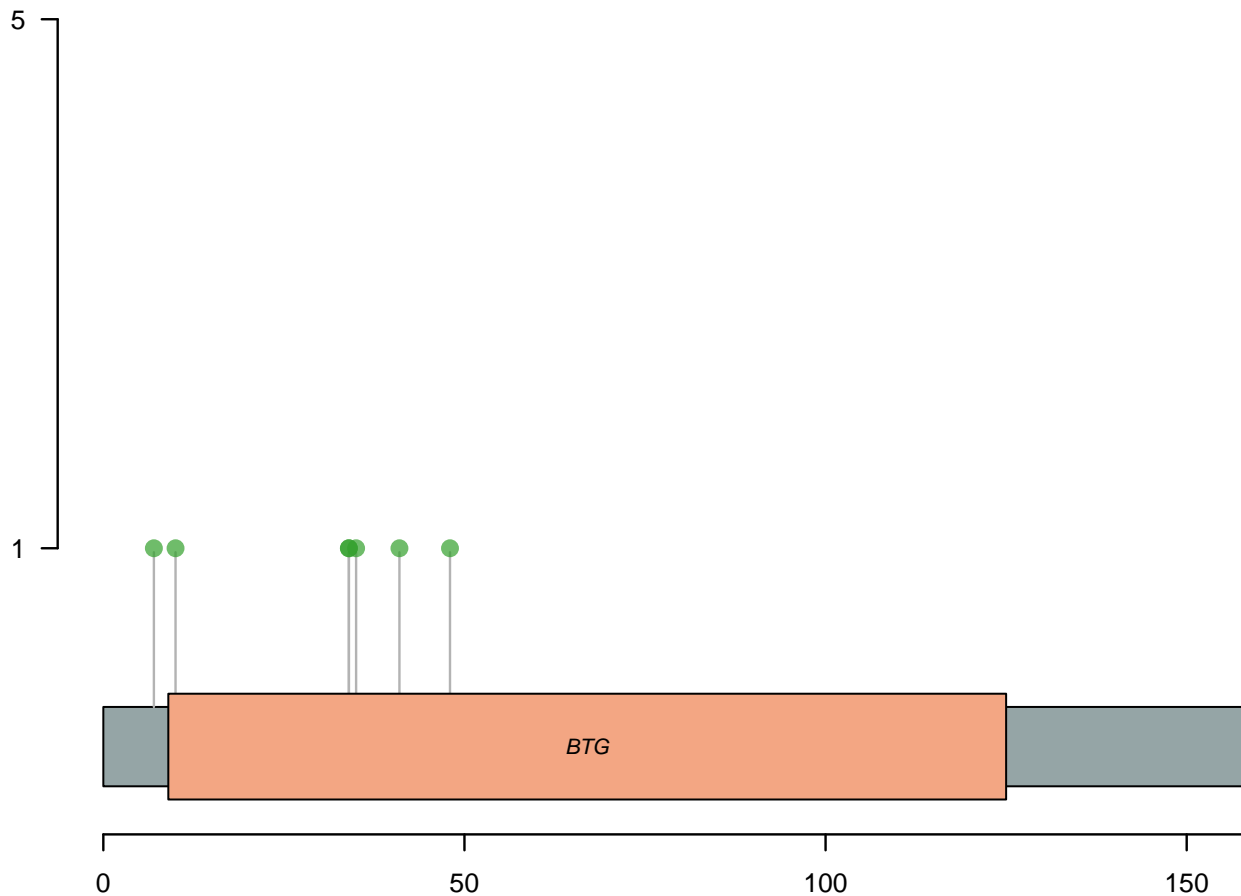


BTG2 : [Somatic Mutation Rate: 2.98%]

NM_006763



● Missense