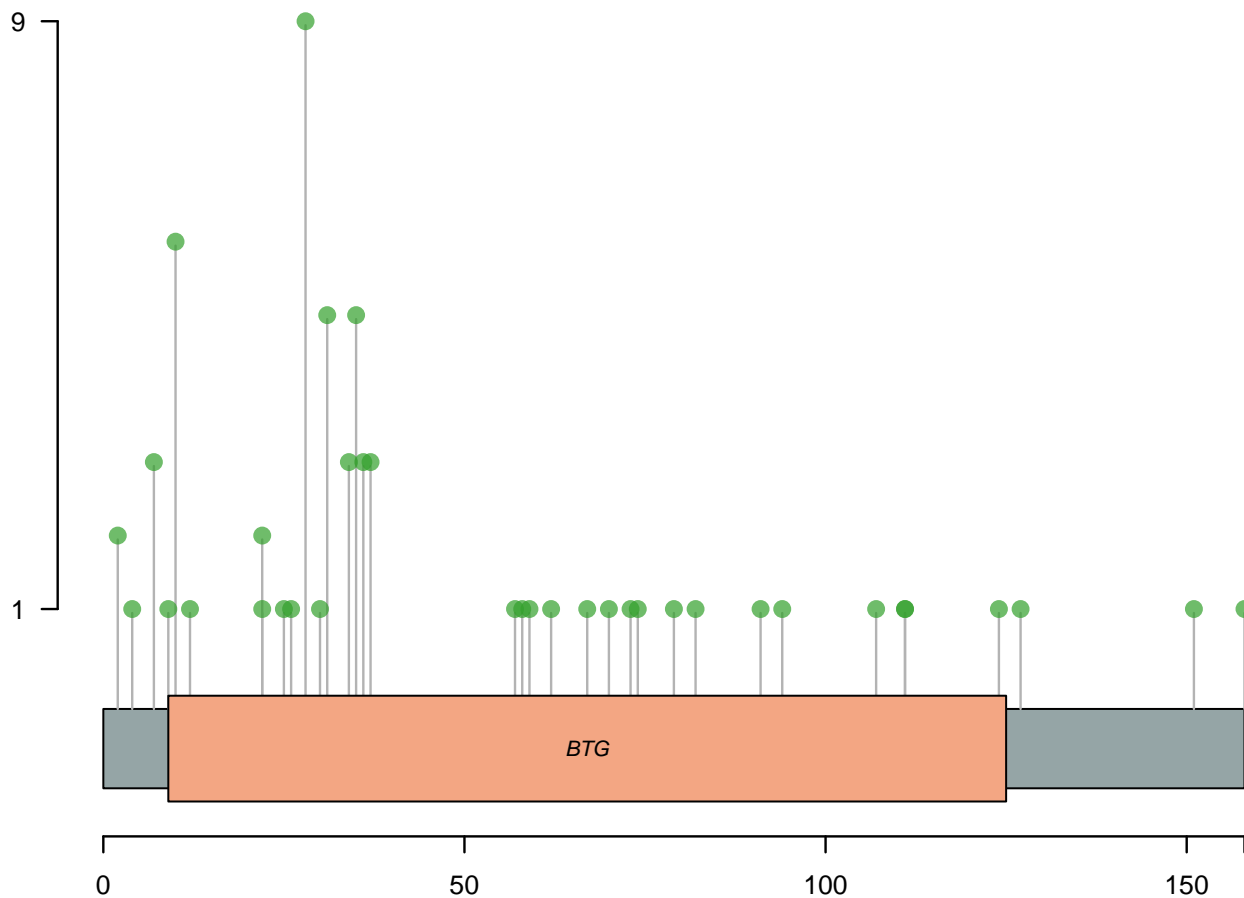


BTG2 : [Somatic Mutation Rate: 5.99%]

NM_006763



● Missense_Mutation