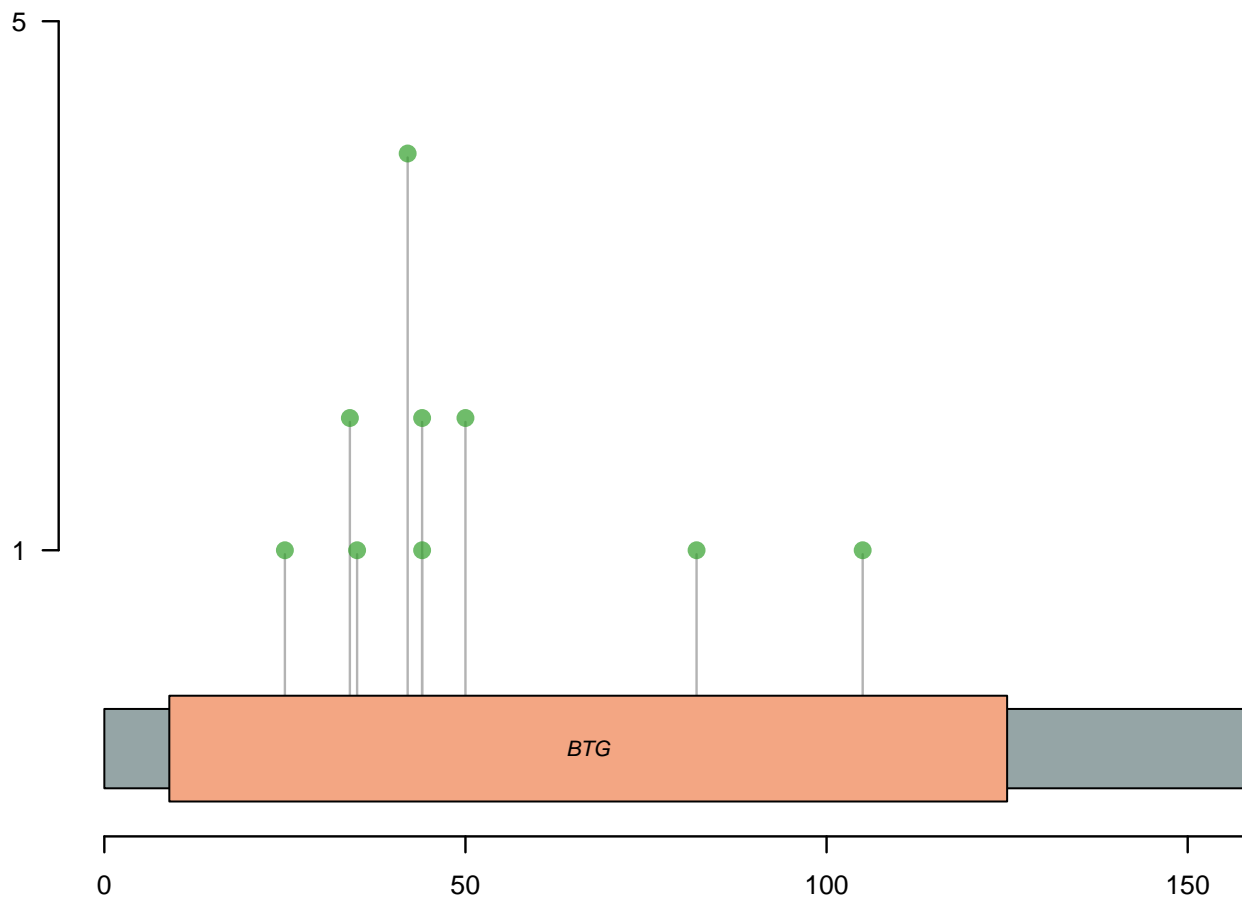


BTG2 : [Somatic Mutation Rate: 14.29%]

NM_006763



● Missense