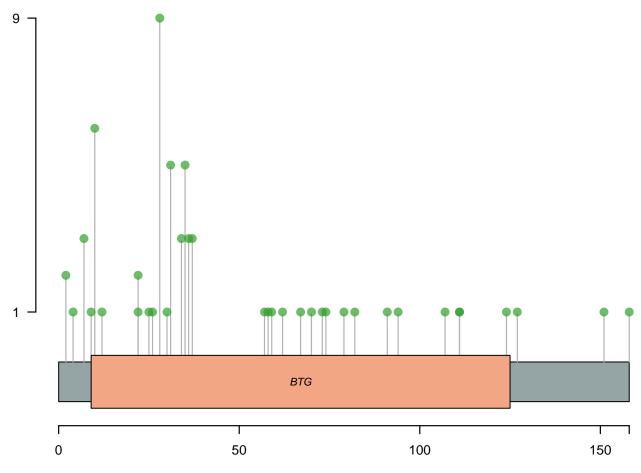
BTG2 : [Somatic Mutation Rate: 5.96%] NM_006763



Missense_Mutation