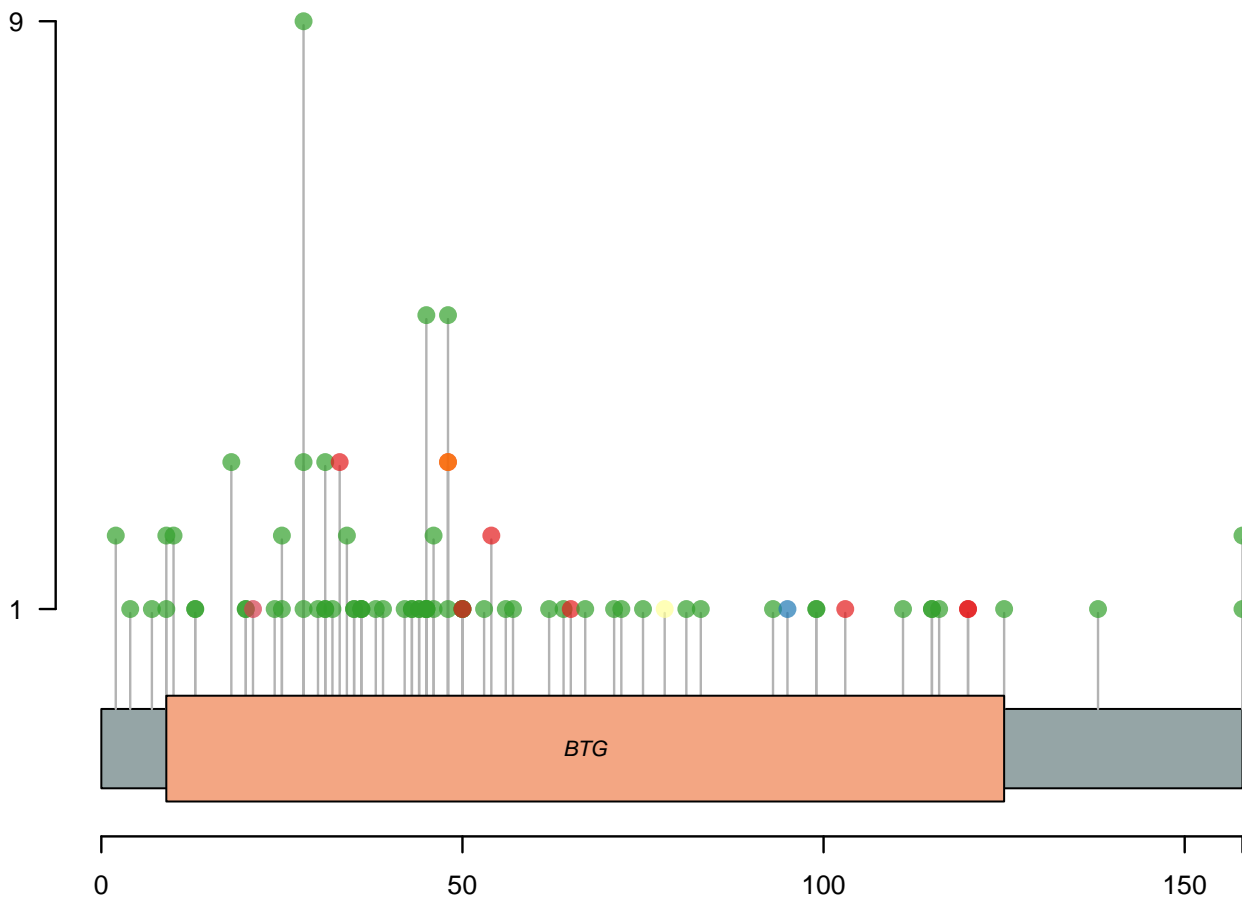


BTG2 : [Somatic Mutation Rate: 16.17%]

NM_006763



- Missense_Mutation
- Splice_Site
- In_Frame_Ins
- In_Frame_Del
- Nonsense_Mutation
- Frame_Shift_Del