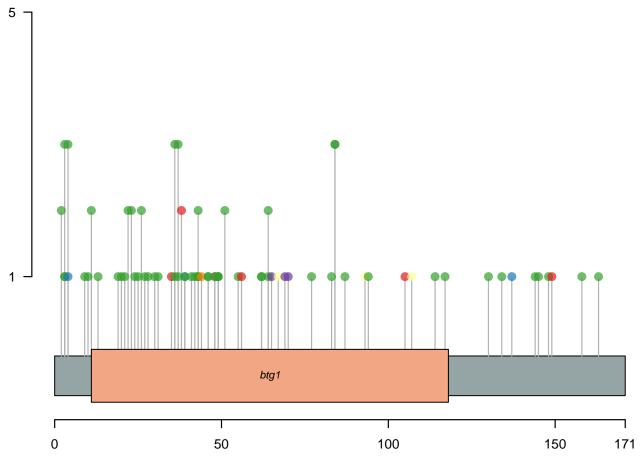
BTG1 : [Somatic Mutation Rate: 13.4%] NM_001731



- Missense_MutationIn_Frame_Ins
- Frame_Shift_Del Frame_Shift_Ins
- Nonsense_Mutation
 In_Frame_Del
- Splice_Site