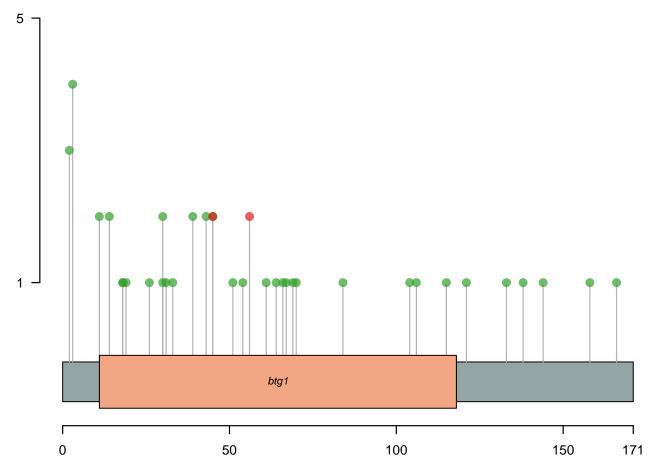
BTG1 : [Somatic Mutation Rate: 4.01%] NM_001731



- Missense_Mutation
- Nonsense_Mutation