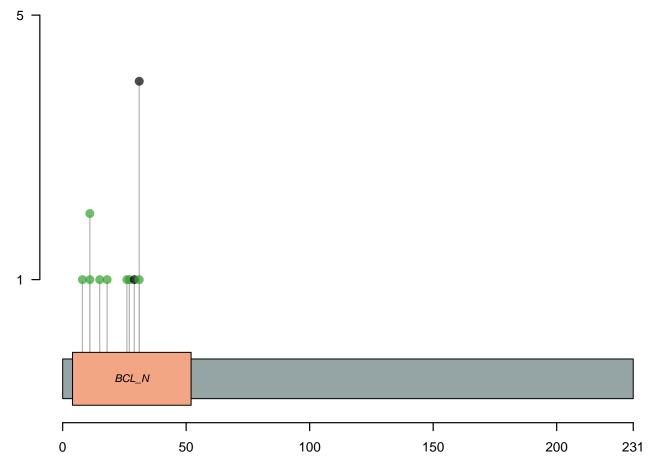
BCL7A: [Somatic Mutation Rate: 5.53%] NM_020993



- Missense
- Truncating