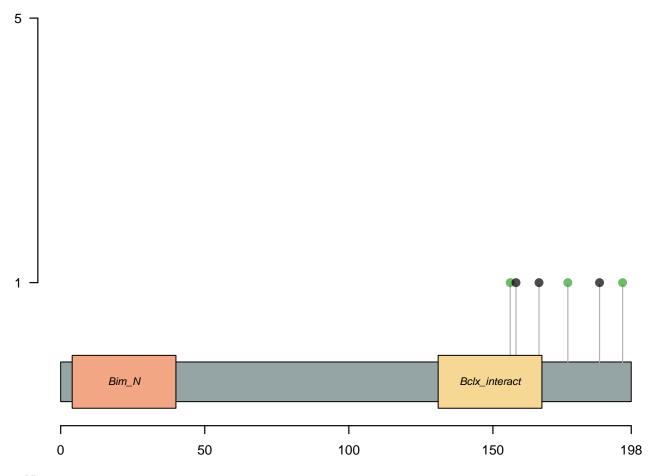
BCL2L11: [Somatic Mutation Rate: 2.13%] NM_138621



- Missense
- Truncating