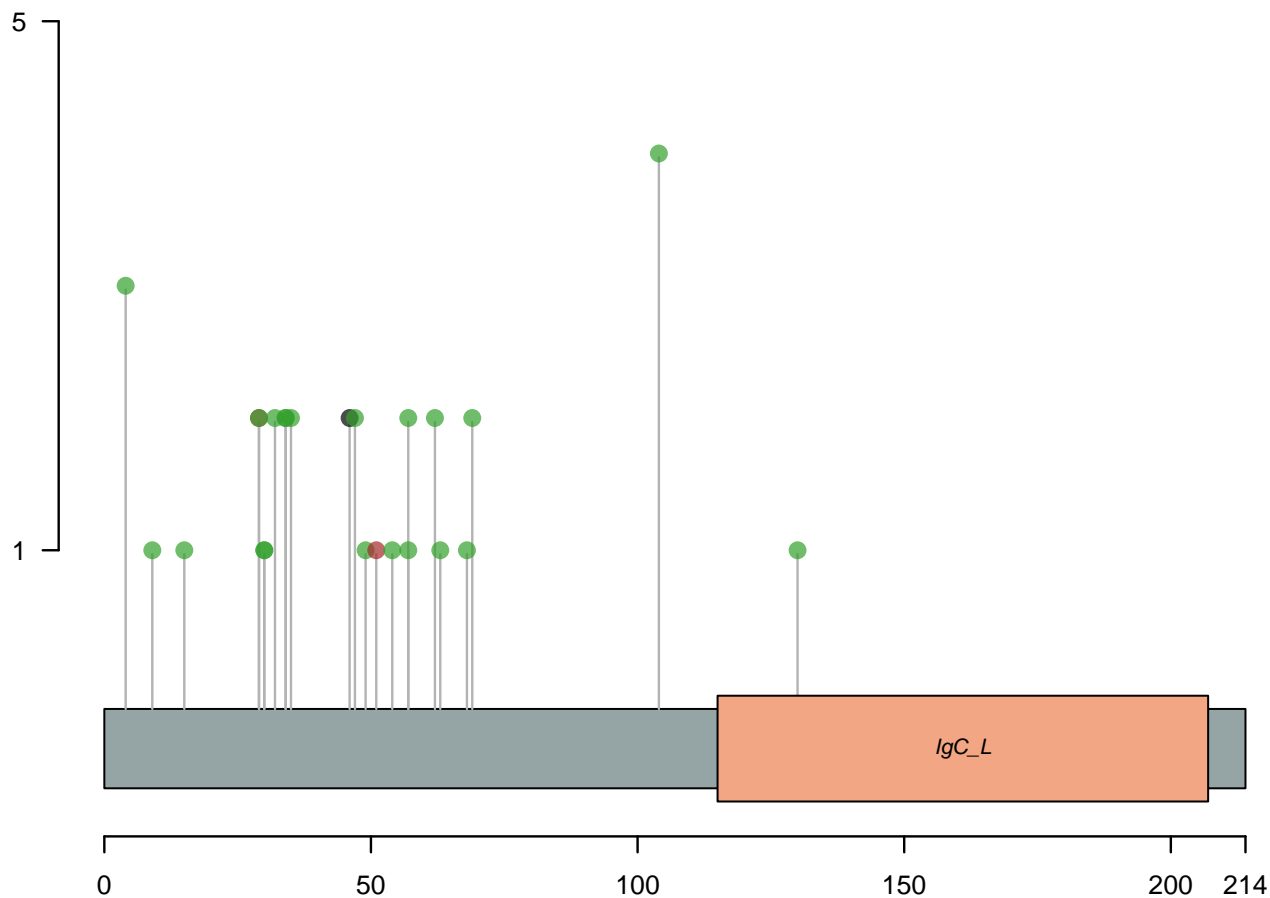


IGLL5 : [Somatic Mutation Rate: 26.53%]

NM_001178126



- Missense
- In-frame
- Truncating