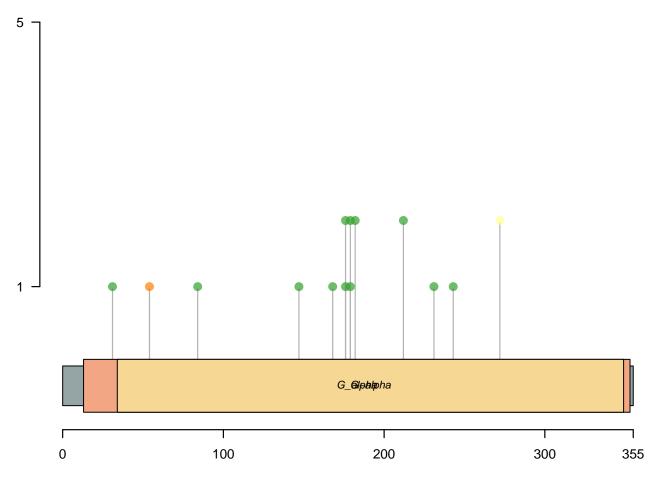
GNAI2: [Somatic Mutation Rate: 1.98%] NM_002070



- Missense_Mutation
- Splice_Site
- In_Frame_Del