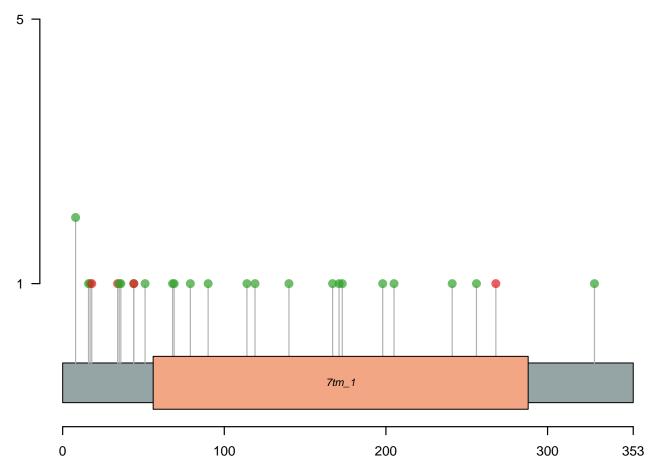
S1PR2: [Somatic Mutation Rate: 1.96%] NM_004230



- Missense_Mutation
- Nonsense_Mutation