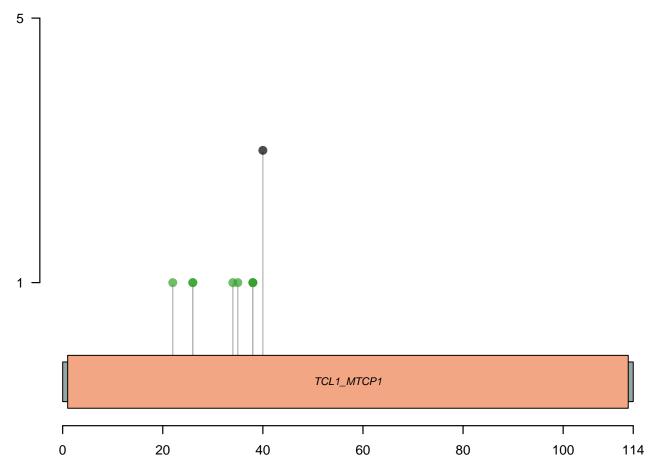
*TCL1A*: [Somatic Mutation Rate: 4.68%] NM\_001098725



- Missense
- Truncating