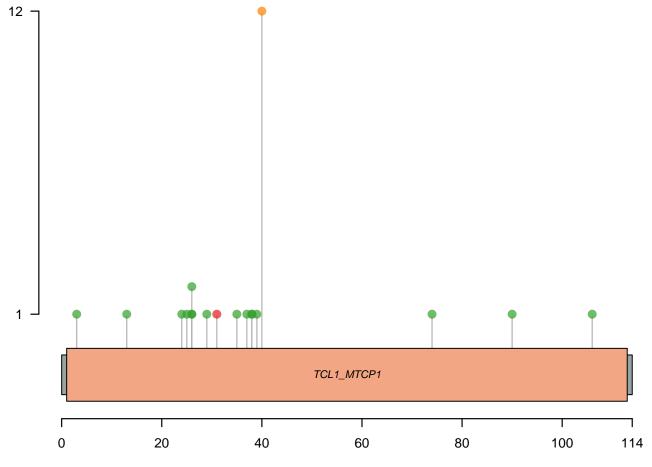
TCL1A: [Somatic Mutation Rate: 2.51%] NM_001098725



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site