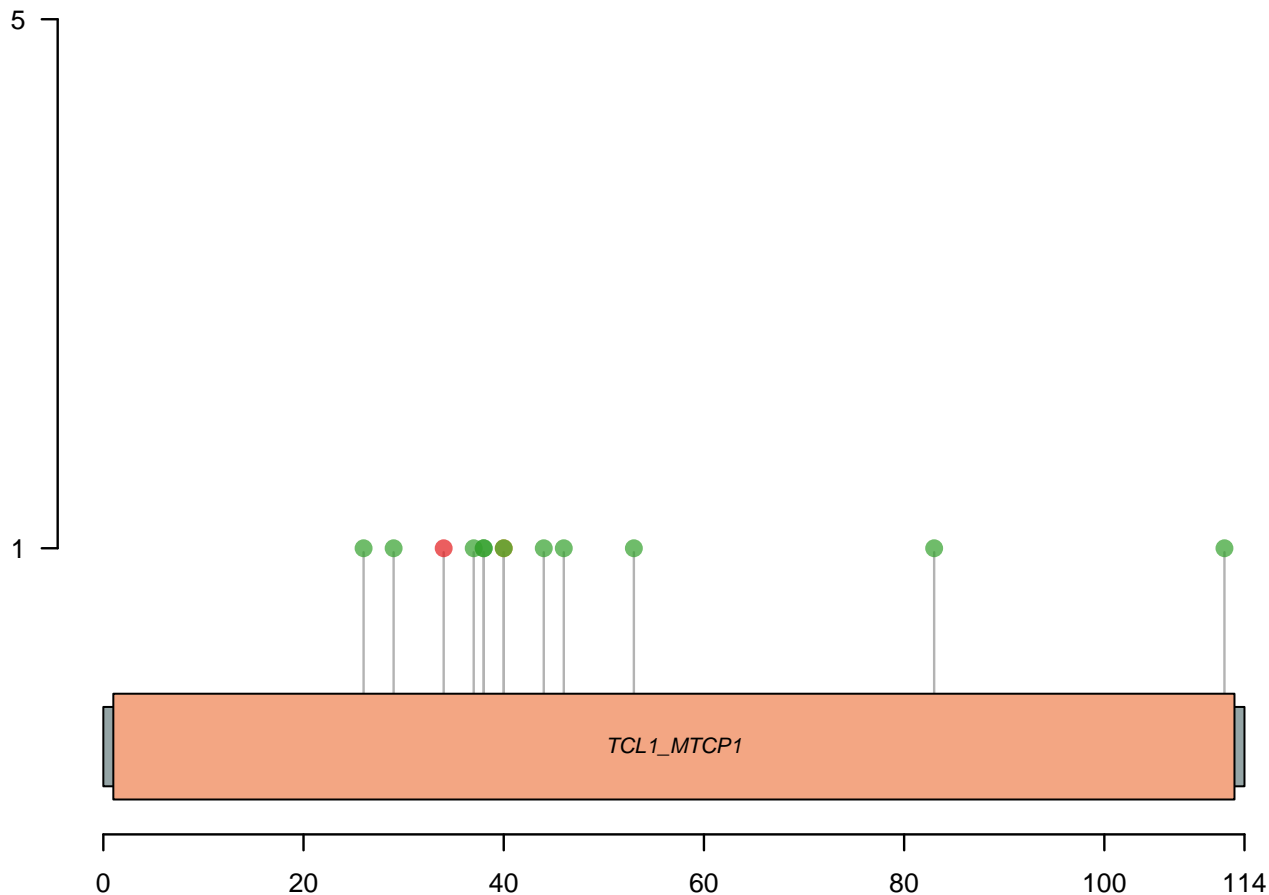


TCL1A : [Somatic Mutation Rate: 2.55%]

NM_001098725



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site