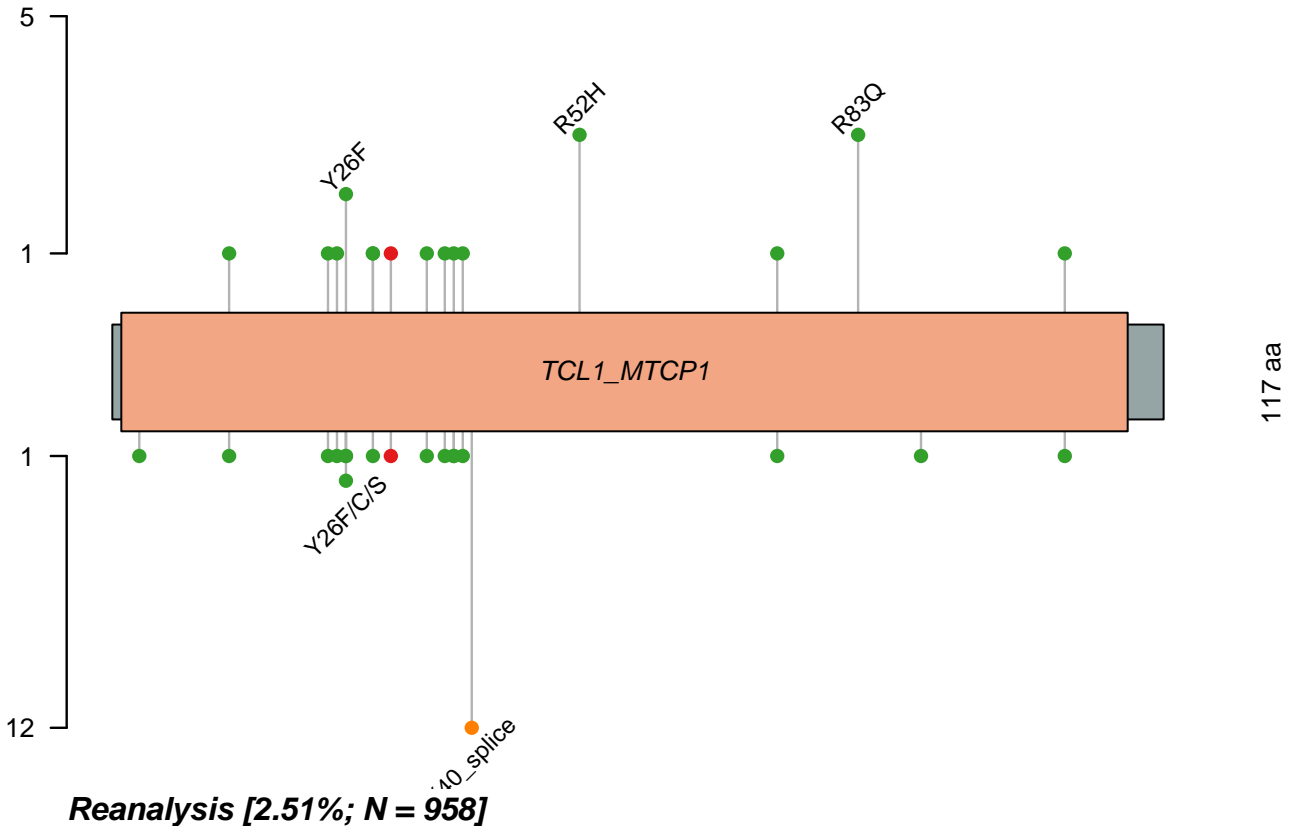


Reddy analysis [1.65%; N = 969]

TCL1A: NM_001098725



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site