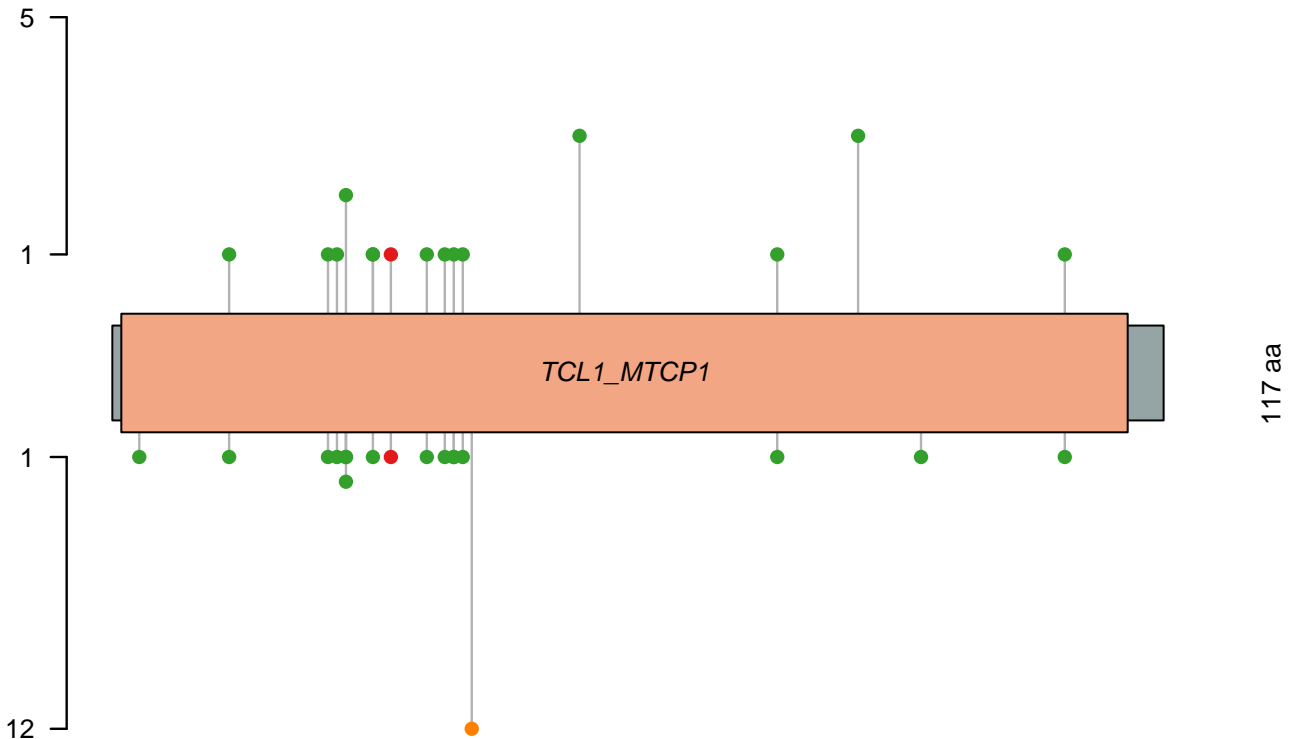


Reddy analysis [1.65%; N = 969]

TCL1A: NM_001098725



Reanalysis [2.51%; N = 958]

- Missense_Mutation
- Nonsense_Mutation
- Splice_Site