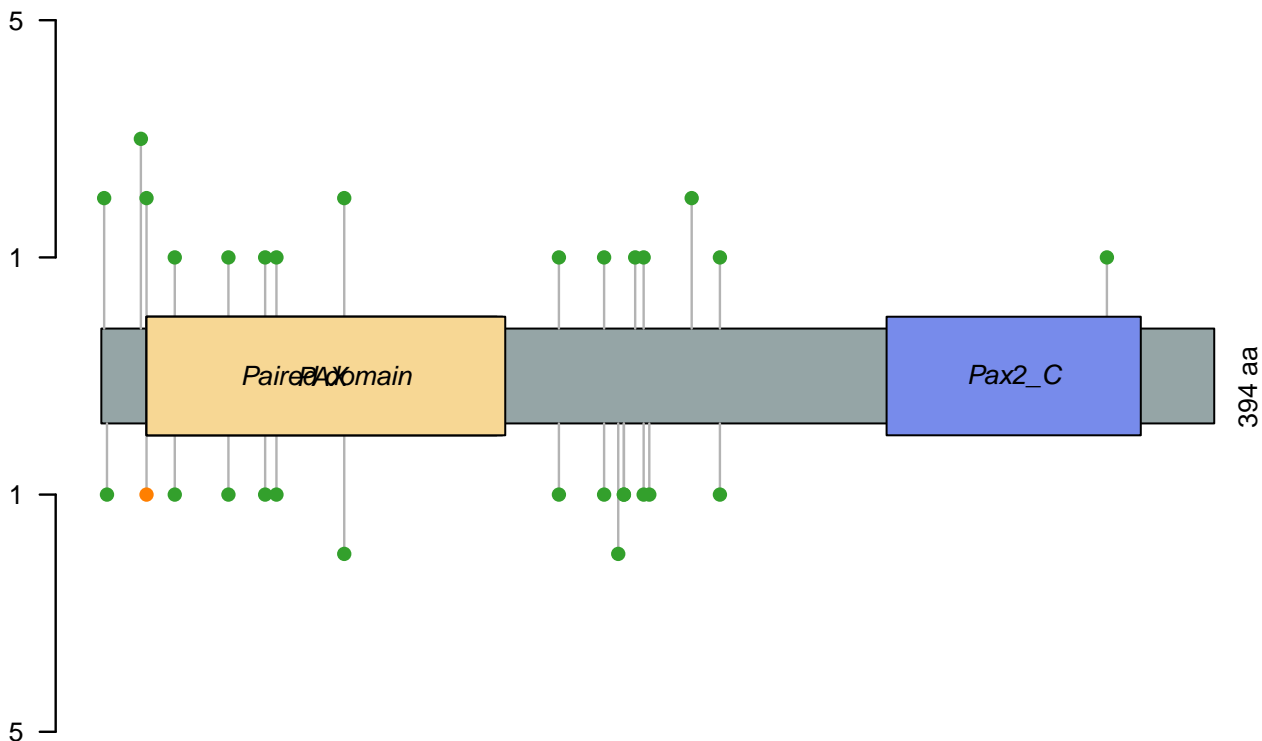


Reddy analysis [2.16%; N = 973]

PAX5: NM_016734



Reanalysis [1.77%; N = 958]

- Missense_Mutation
- Splice_Site