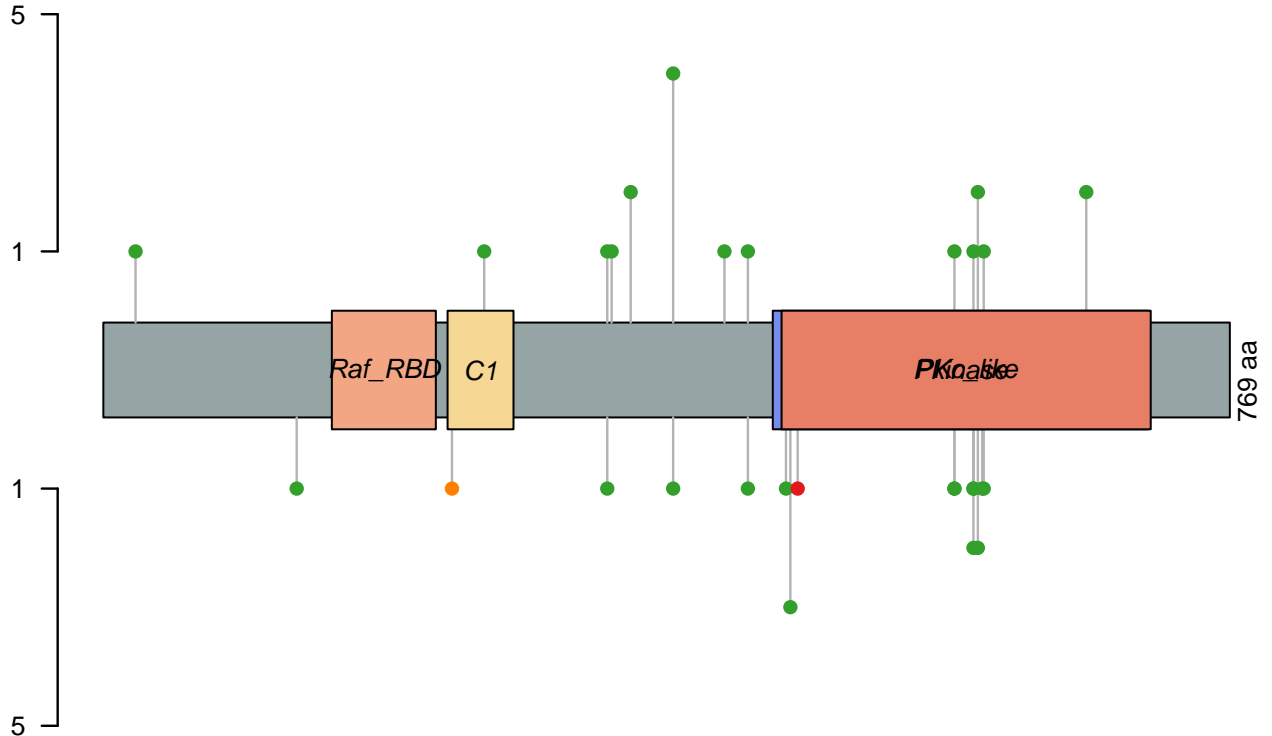


Reddy analysis [1.96%; N = 969]

BRAF: NM_004333



Reanalysis [2.09%; N = 958]

- Missense_Mutation
- Splice_Site
- Nonsense_Mutation