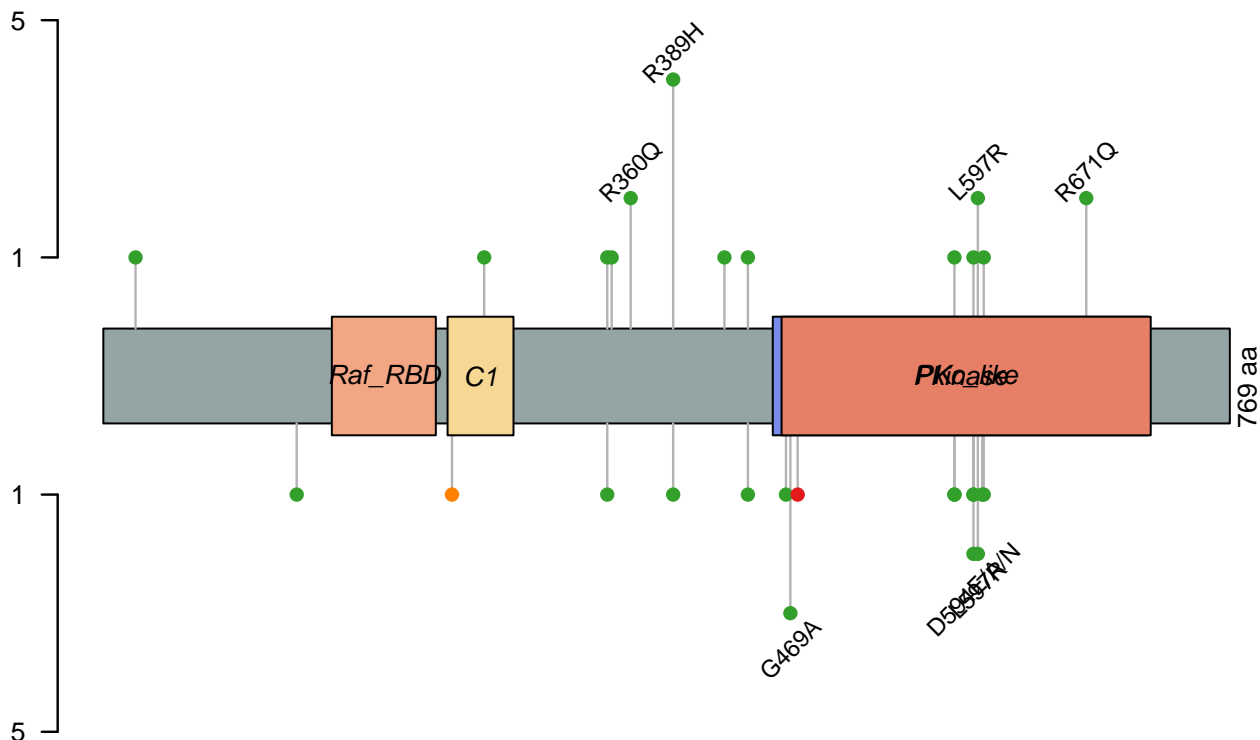


# Reddy analysis [1.96%; N = 969]

BRAF: NM\_004333



# Reanalysis [2.09%; N = 958]

- Missense\_Mutation
- Splice\_Site
- Nonsense\_Mutation