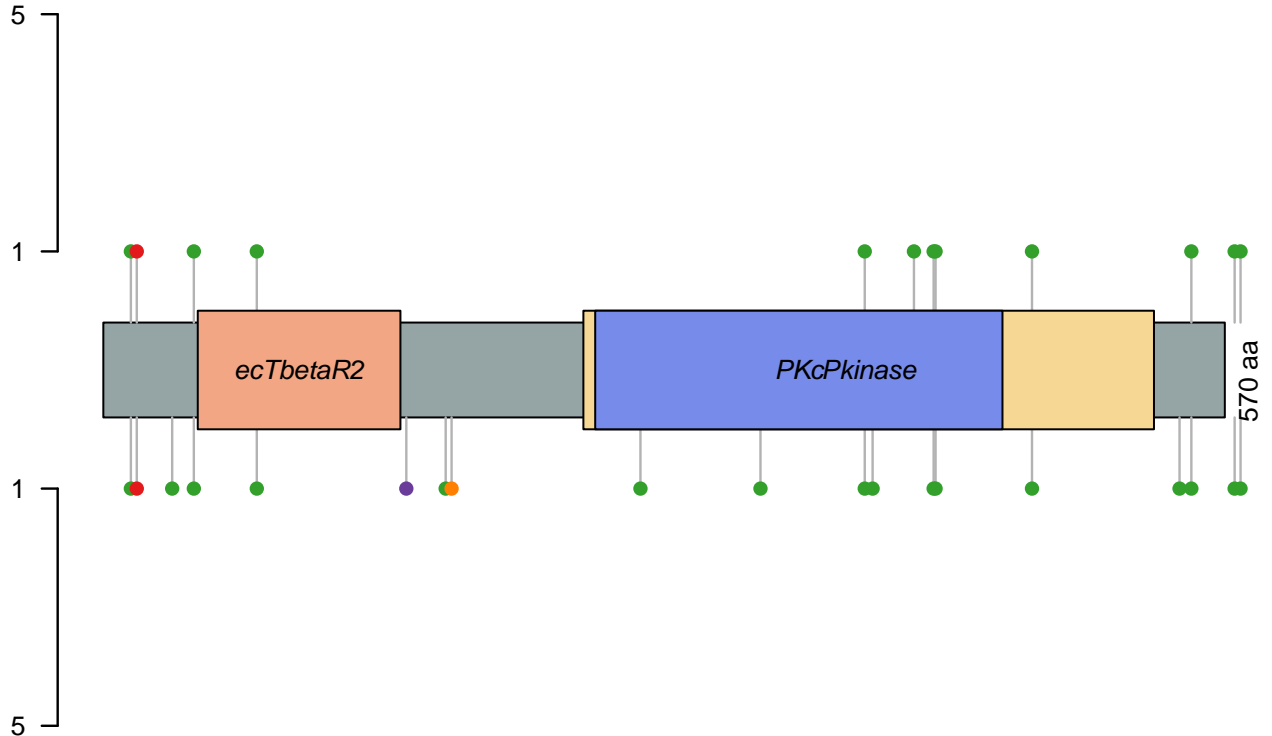


**Reddy analysis [1.23%; N = 973]**

TGFBR2: NM\_003242



**Reanalysis [1.88%; N = 958]**

- Missense\_Mutation
- Nonsense\_Mutation
- Frame\_Shift\_Ins
- Splice\_Site