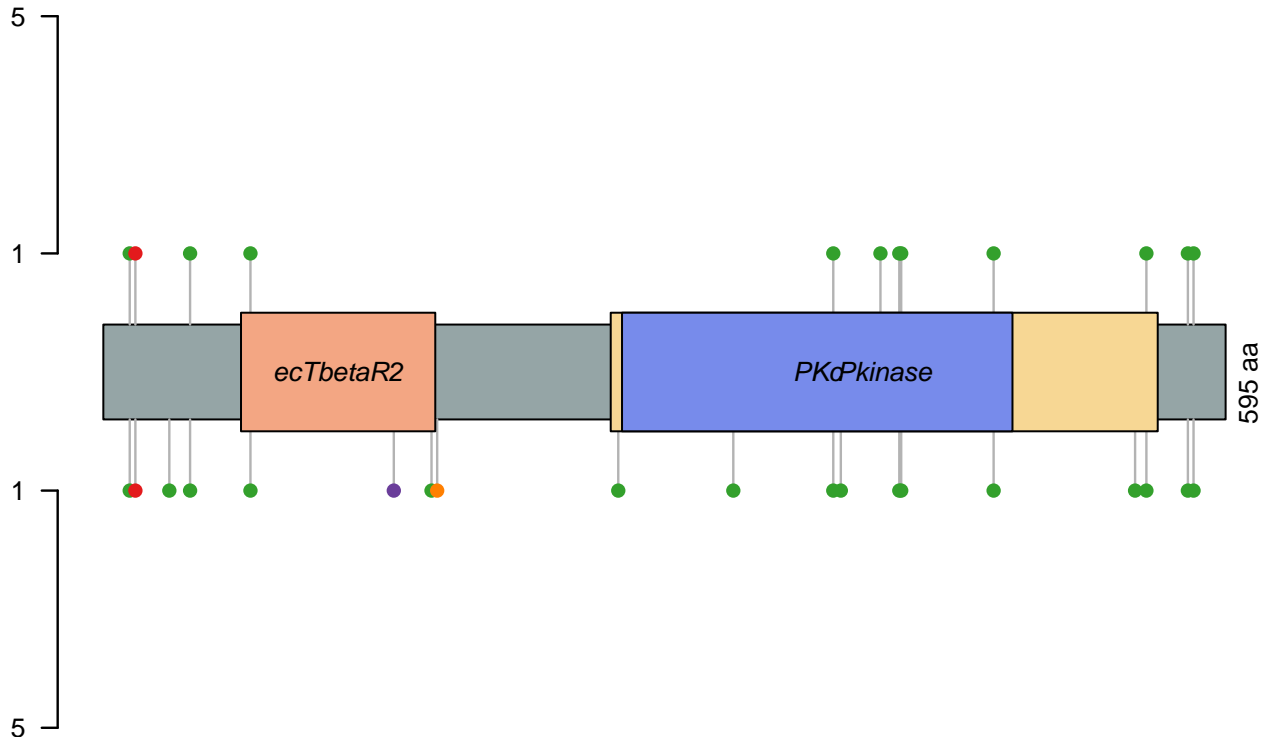**Reddy analysis [1.24%; N = 969]**

TGFBR2: NM_001024847

595 aa

Reanalysis [1.88%; N = 958]

- ● Missense_Mutation
- ● Nonsense_Mutation
- ● Frame_Shift_Ins
- ● Splice_Site