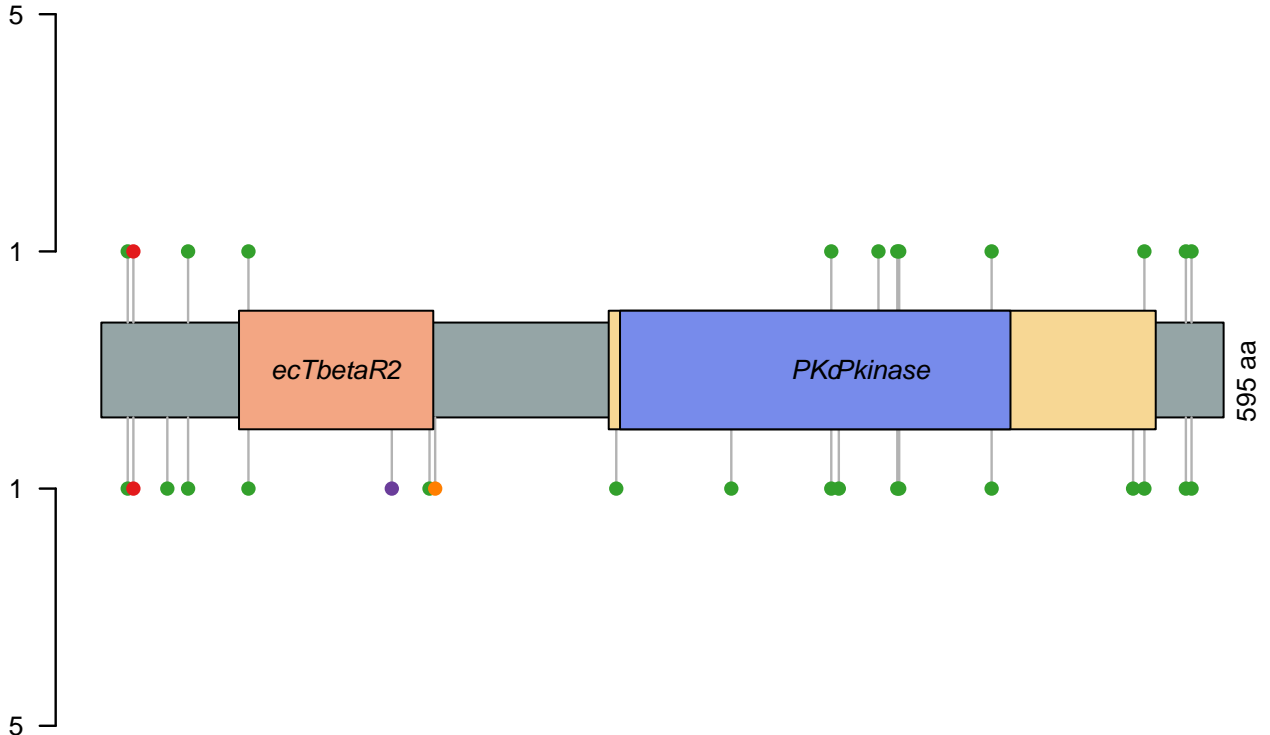


Reddy analysis [1.24%; N = 969]

TGFB2: NM_001024847



Reanalysis [1.88%; N = 958]

- Missense_Mutation
- Nonsense_Mutation
- Frame_Shift_Ins
- Splice_Site