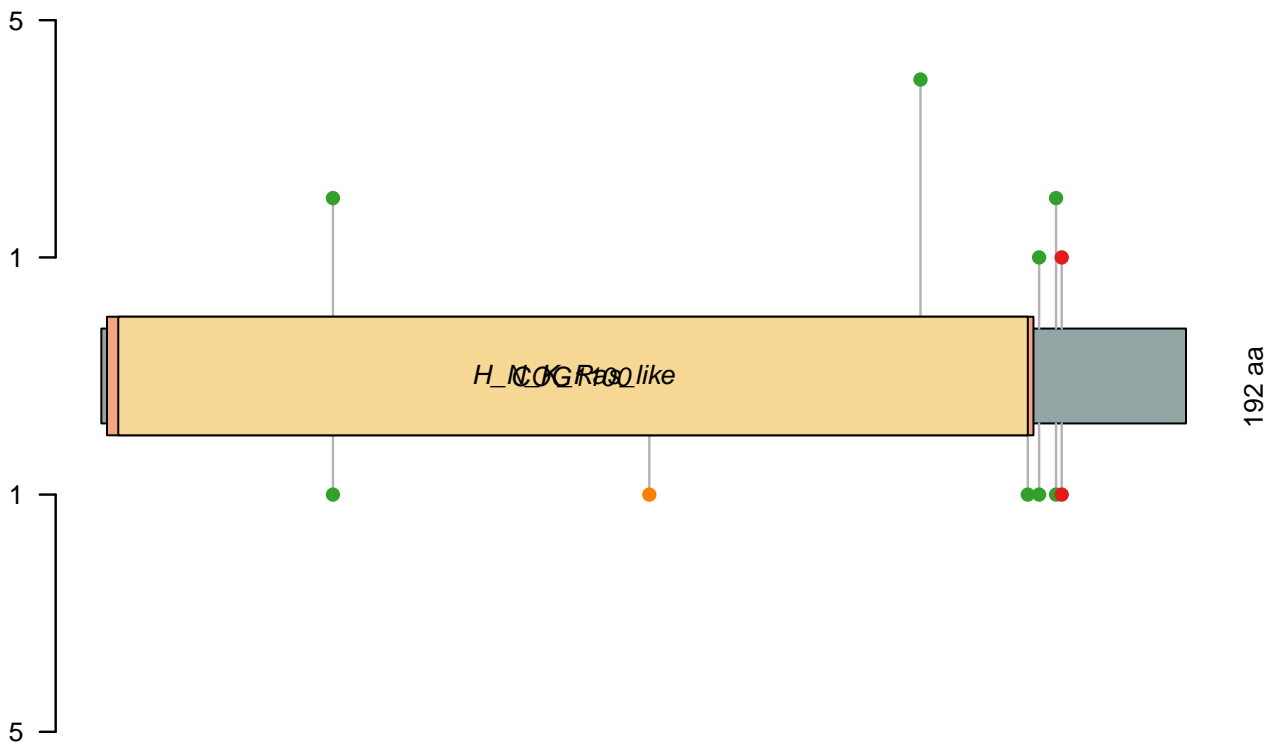


Reddy analysis [1.03%; N = 969]

HRAS: NM_001130442



Reanalysis [0.63%; N = 958]

- Missense_Mutation
- Nonsense_Mutation
- Splice_Site