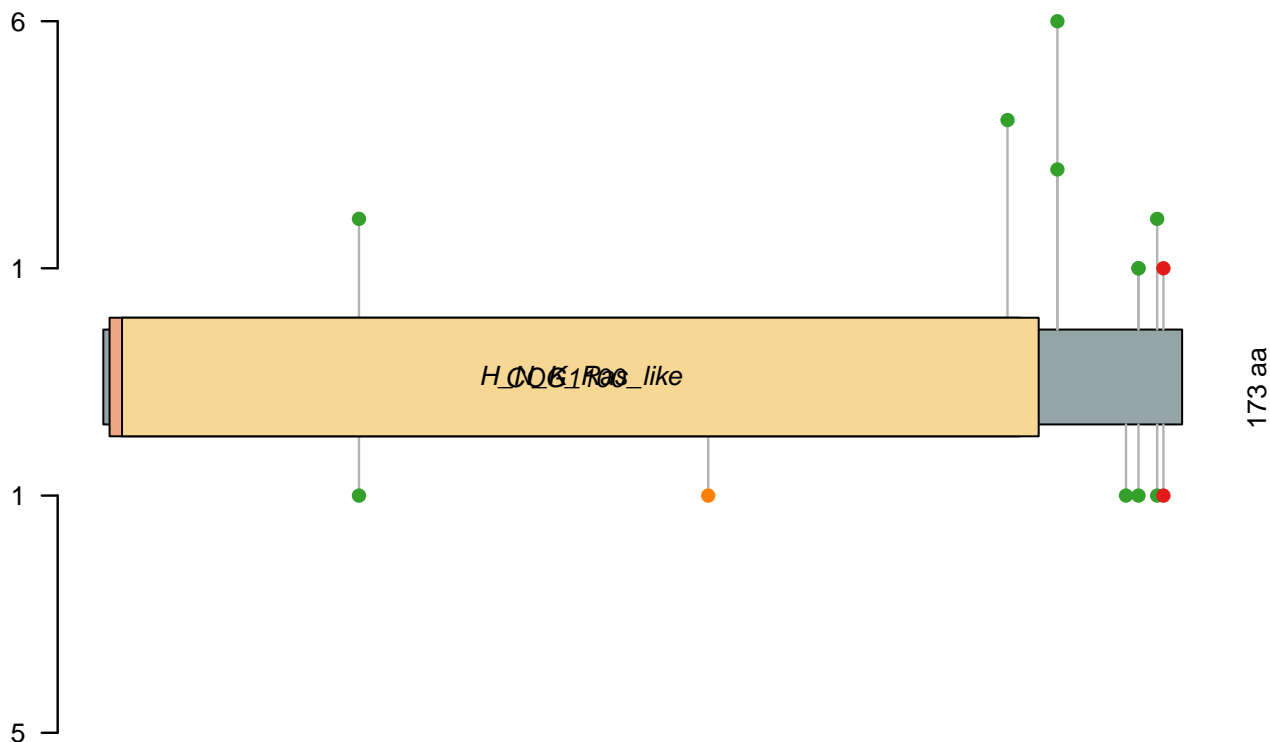


Reddy analysis [2.06%; N = 973]

HRAS: NM_176795



Reanalysis [0.63%; N = 958]

- Missense_Mutation
- Nonsense_Mutation
- Splice_Site