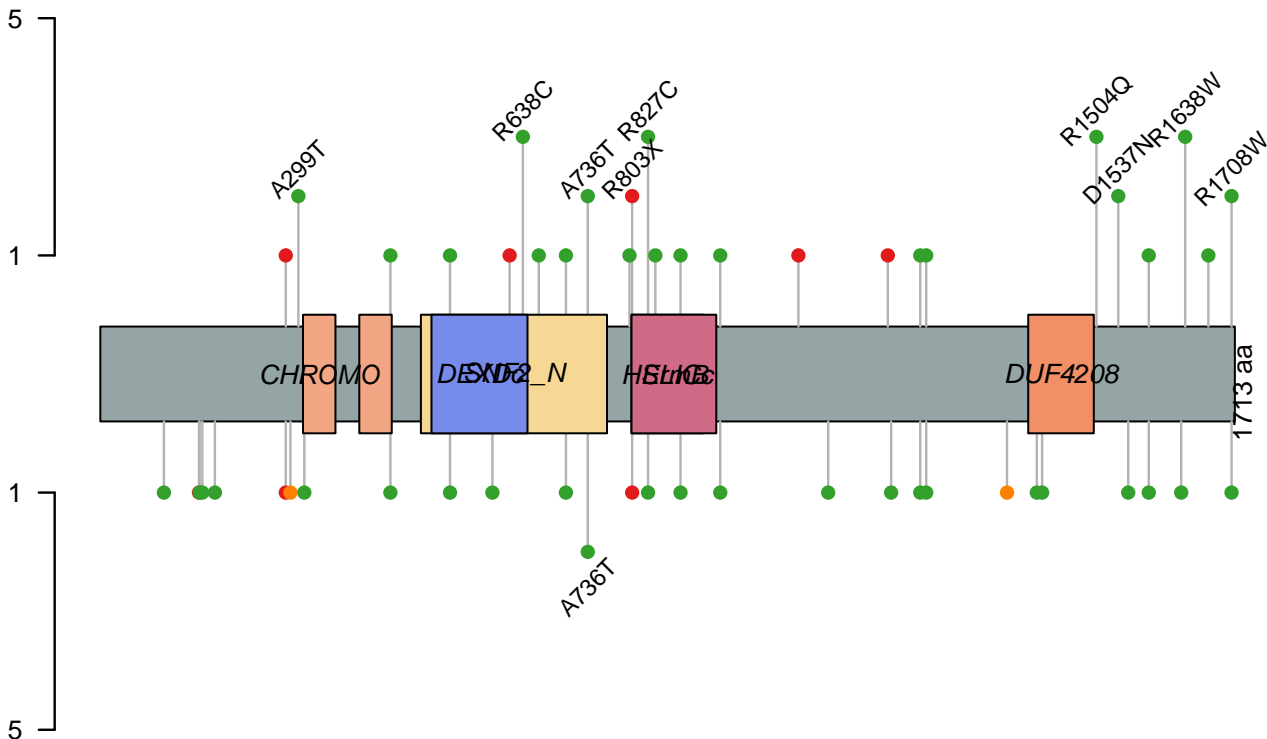


# Reddy analysis [3.82%; N = 969]

CHD1: NM\_001270



# Reanalysis [2.82%; N = 958]

- Nonsense\_Mutation
- Missense\_Mutation
- Splice\_Site