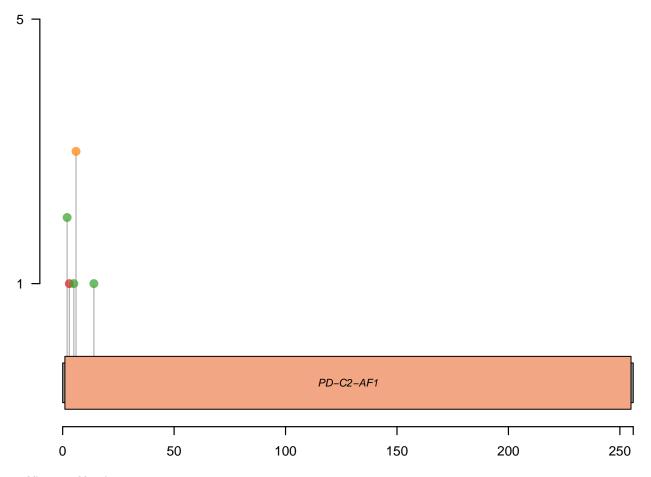
POU2AF1: [Somatic Mutation Rate: 1.91%] NM_006235



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site