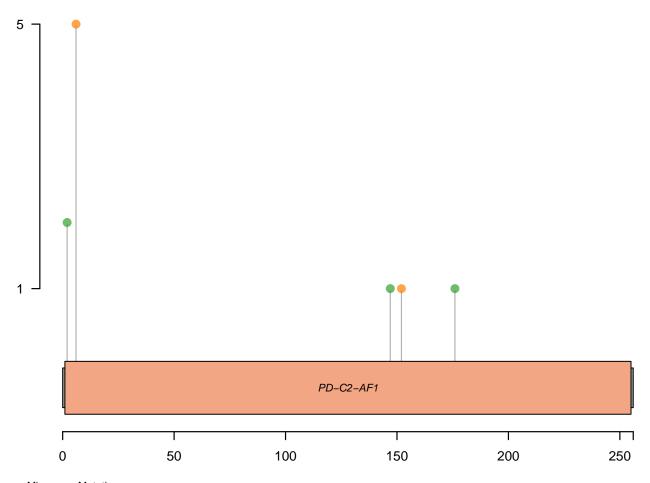
POU2AF1 : [Somatic Mutation Rate: 4.29%] NM_006235



- Missense_Mutation
- Splice_Site