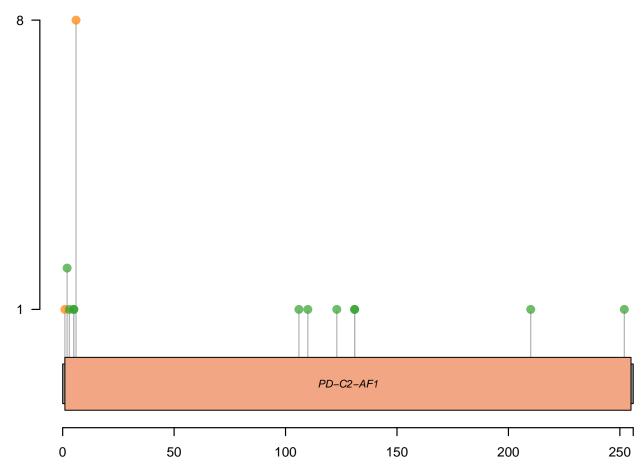
POU2AF1 : [Somatic Mutation Rate: 2.09%] NM_006235



- Splice_Site
- Missense_Mutation