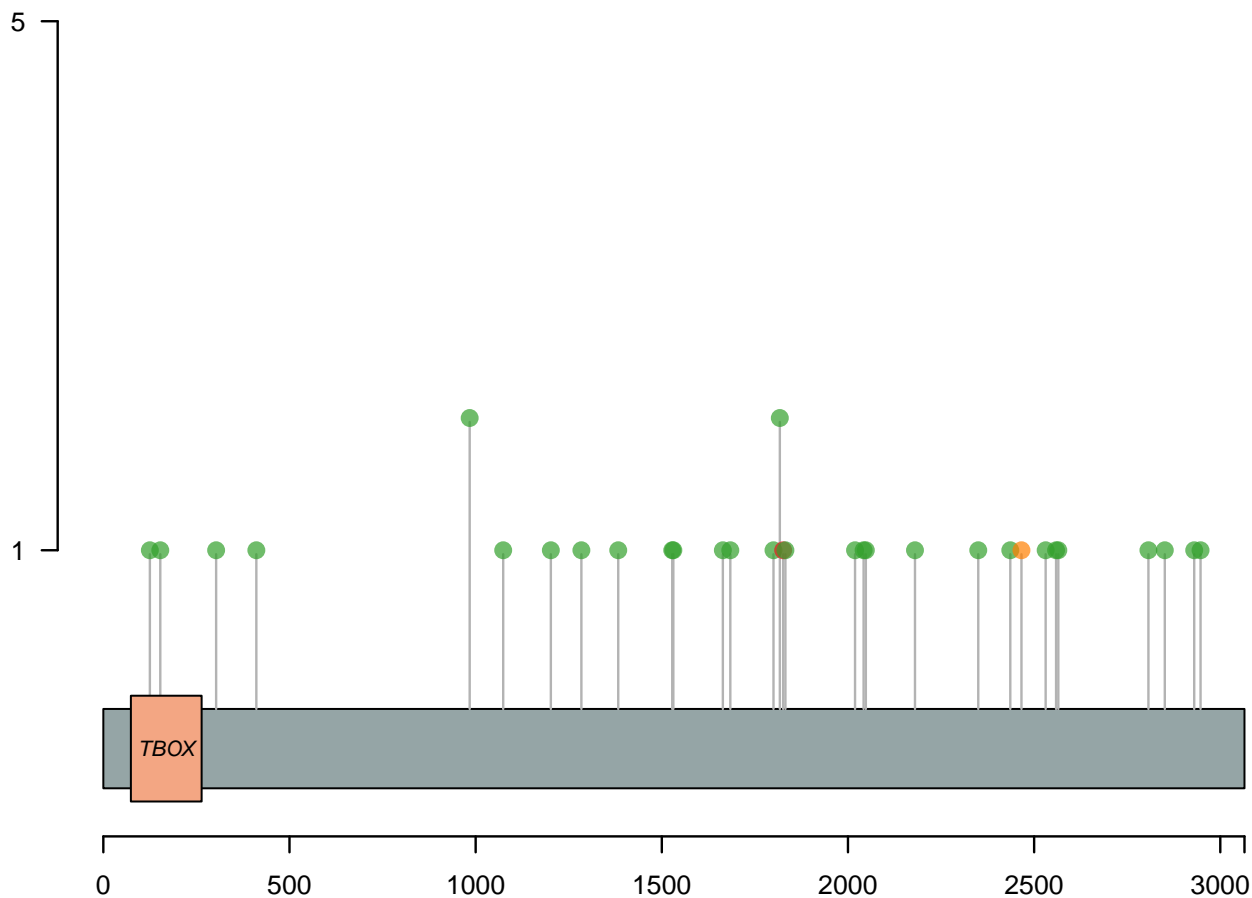


MGA : [Somatic Mutation Rate: 7.02%]

NM_001164273



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site