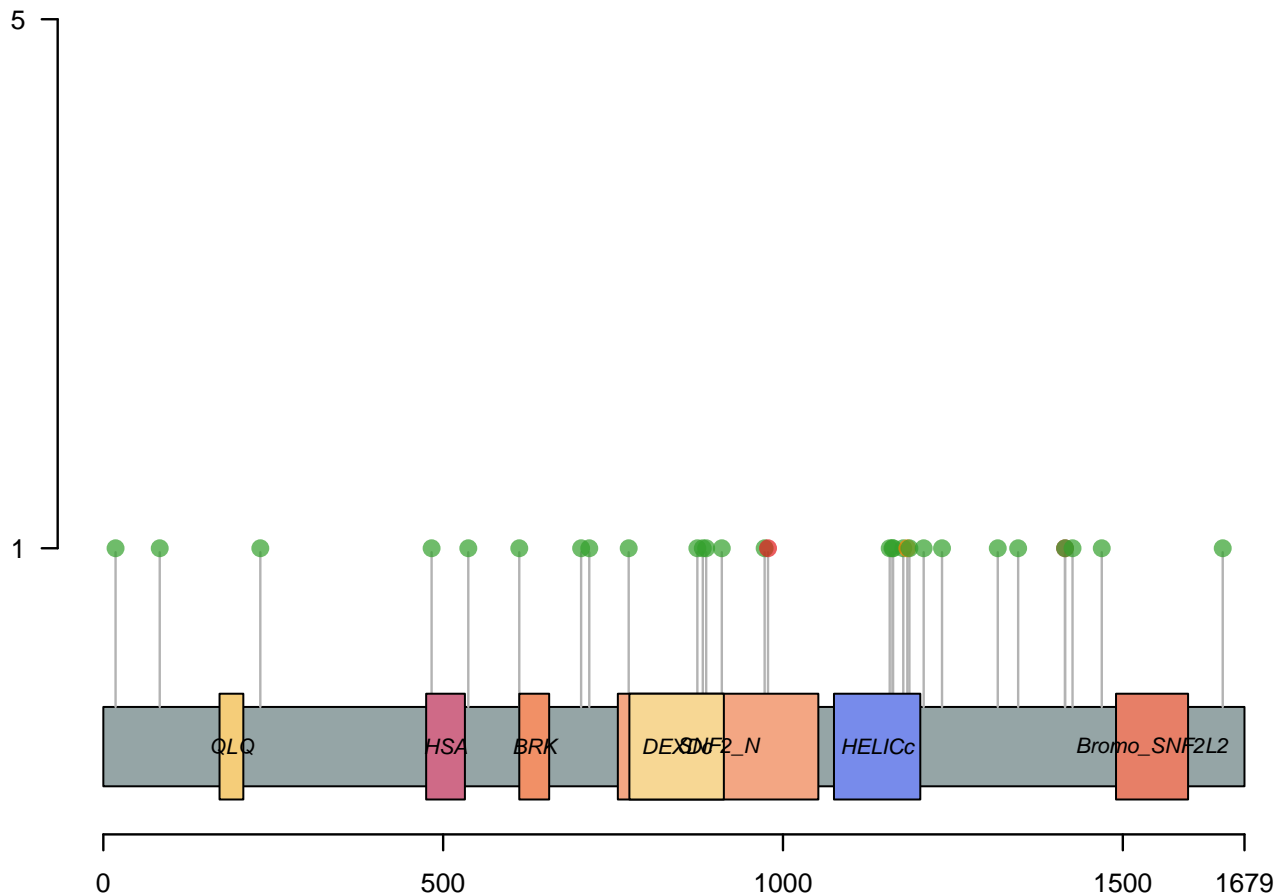


SMARCA4 : [Somatic Mutation Rate: 3.13%]

NM_001128849



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site