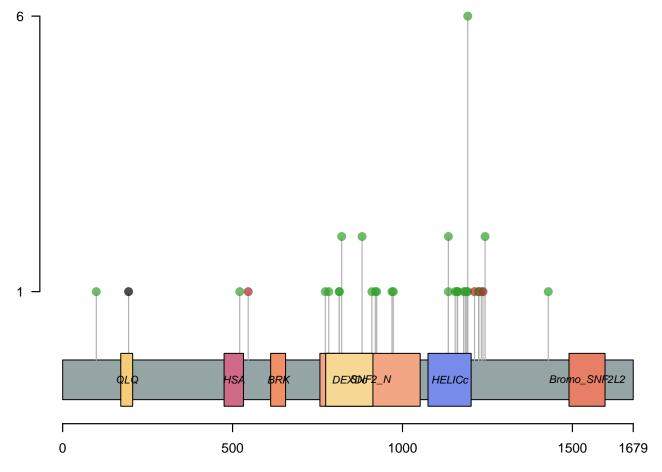
SMARCA4: [Somatic Mutation Rate: 17.45%] NM_001128849



- Missense
- Truncating
- In–frame