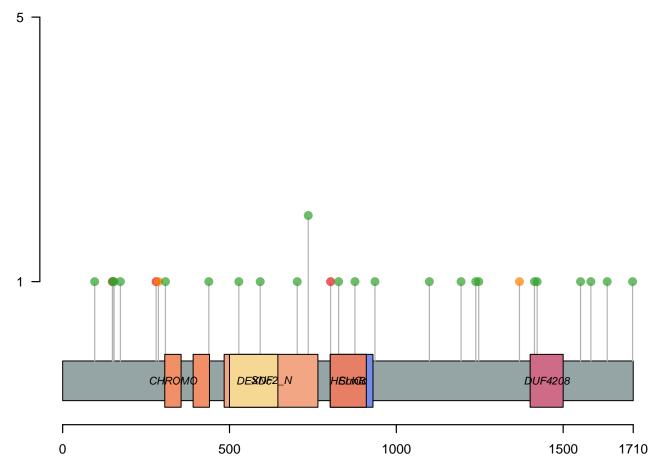
CHD1 : [Somatic Mutation Rate: 2.82%] NM_001270



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site