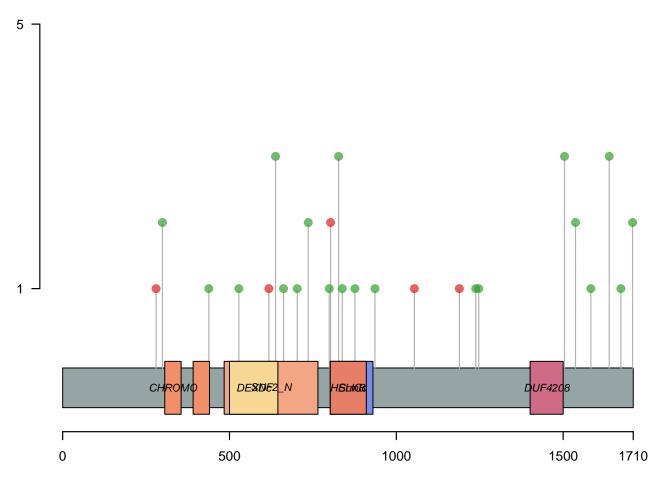
CHD1 : [Somatic Mutation Rate: 3.8%] NM_001270



- Nonsense_Mutation
- Missense_Mutation