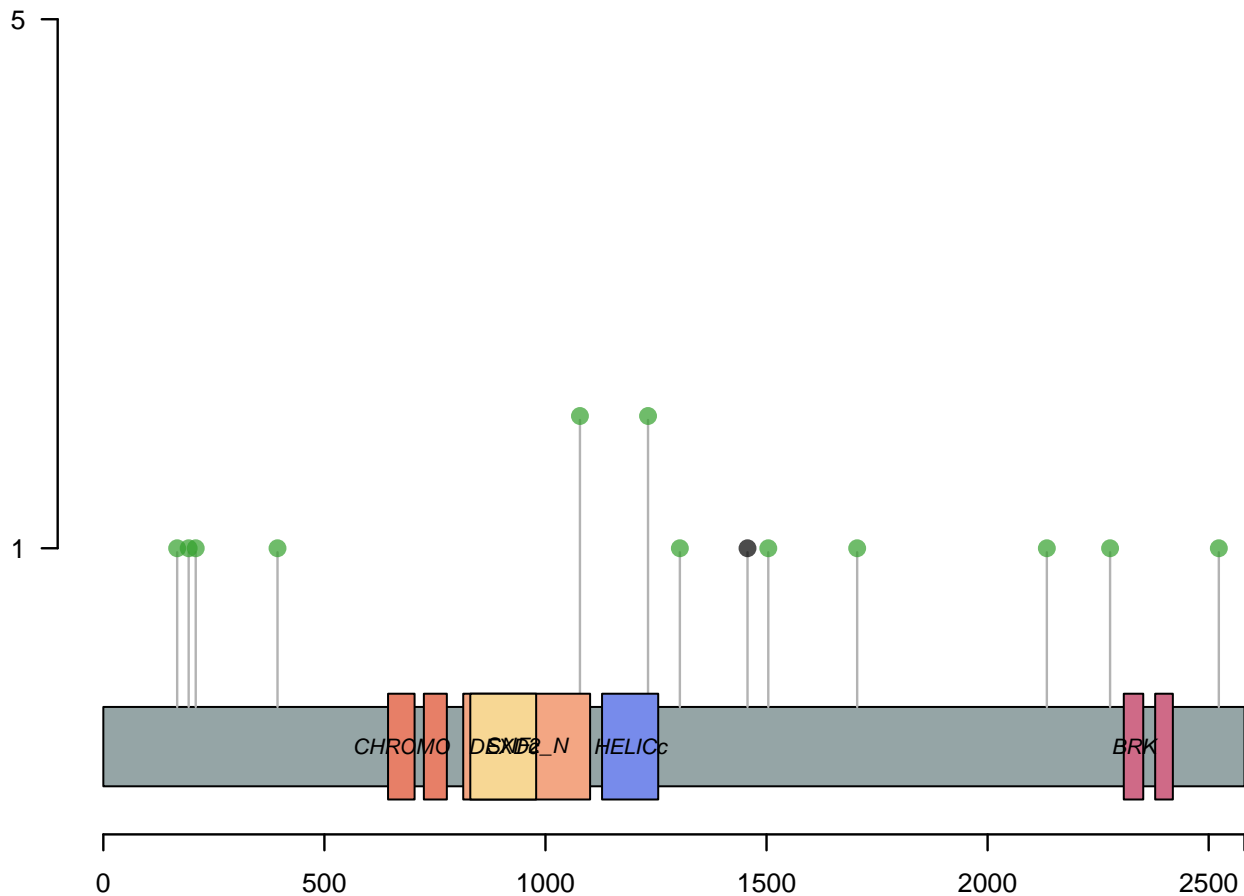


CHD8 : [Somatic Mutation Rate: 10.2%]

NM_001170629



● Missense

● Truncating