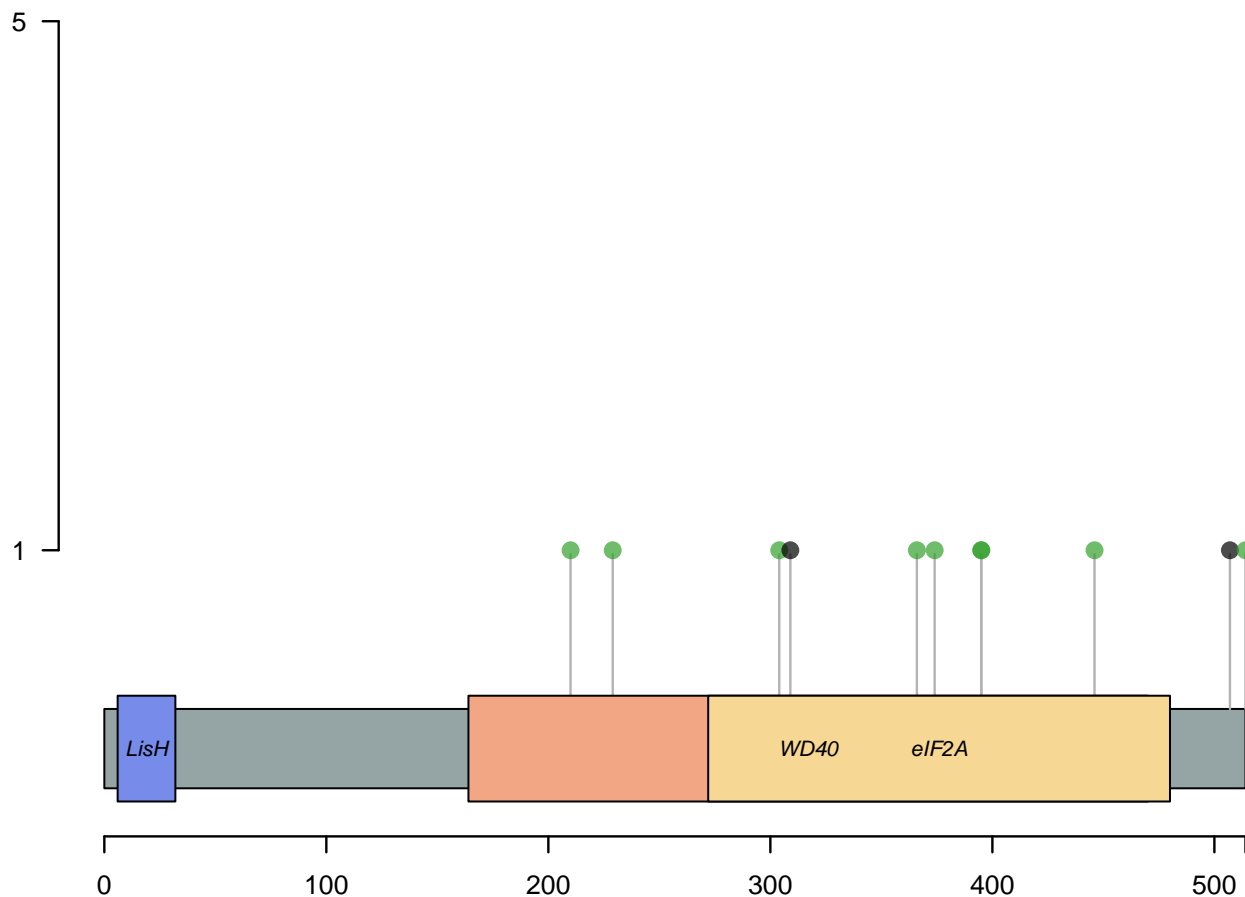


TBL1XR1 : [Somatic Mutation Rate: 4.68%]

NM_024665



● Missense

● Truncating