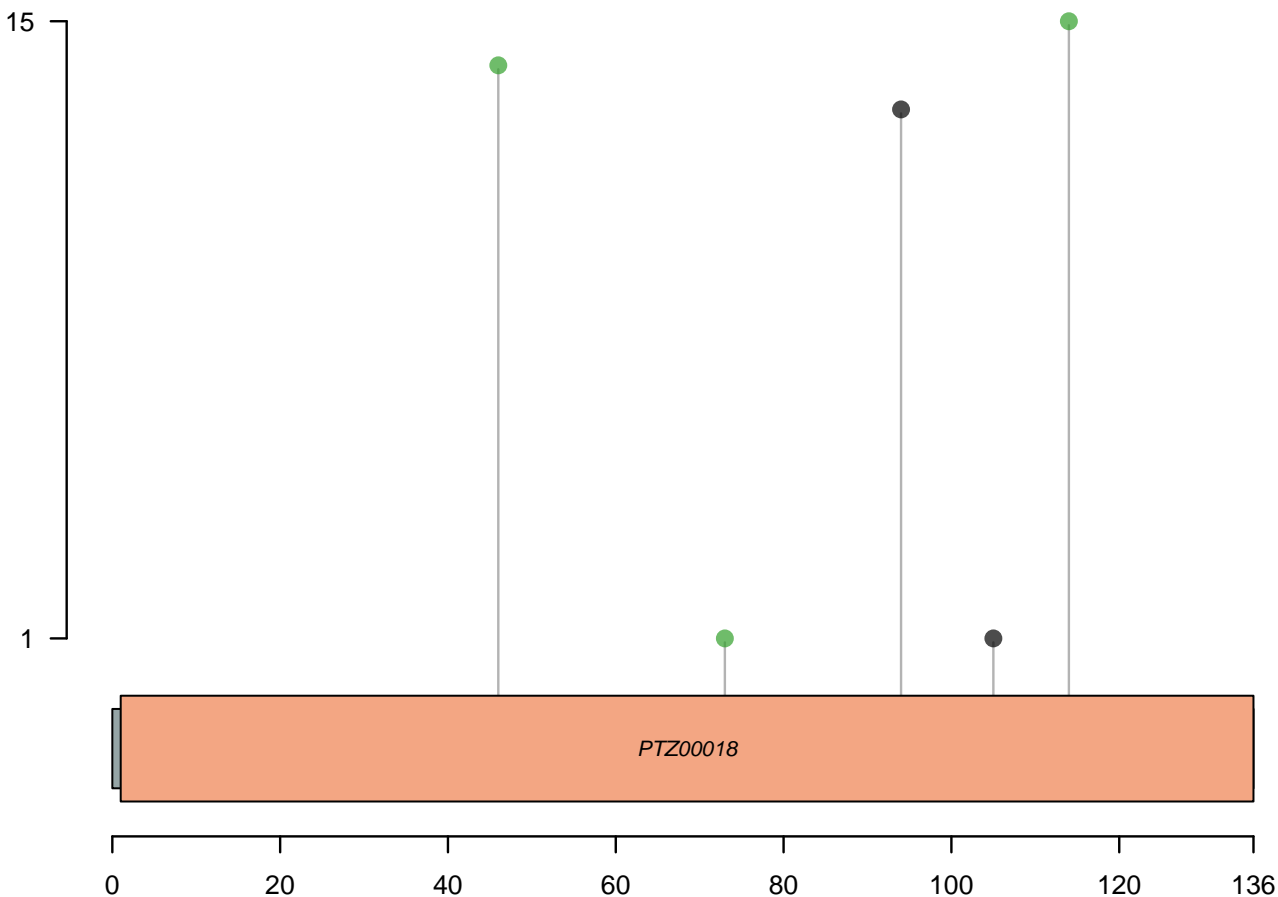


HIST1H3H : [Somatic Mutation Rate: 26.53%]

NM_003536



● Missense

● Truncating