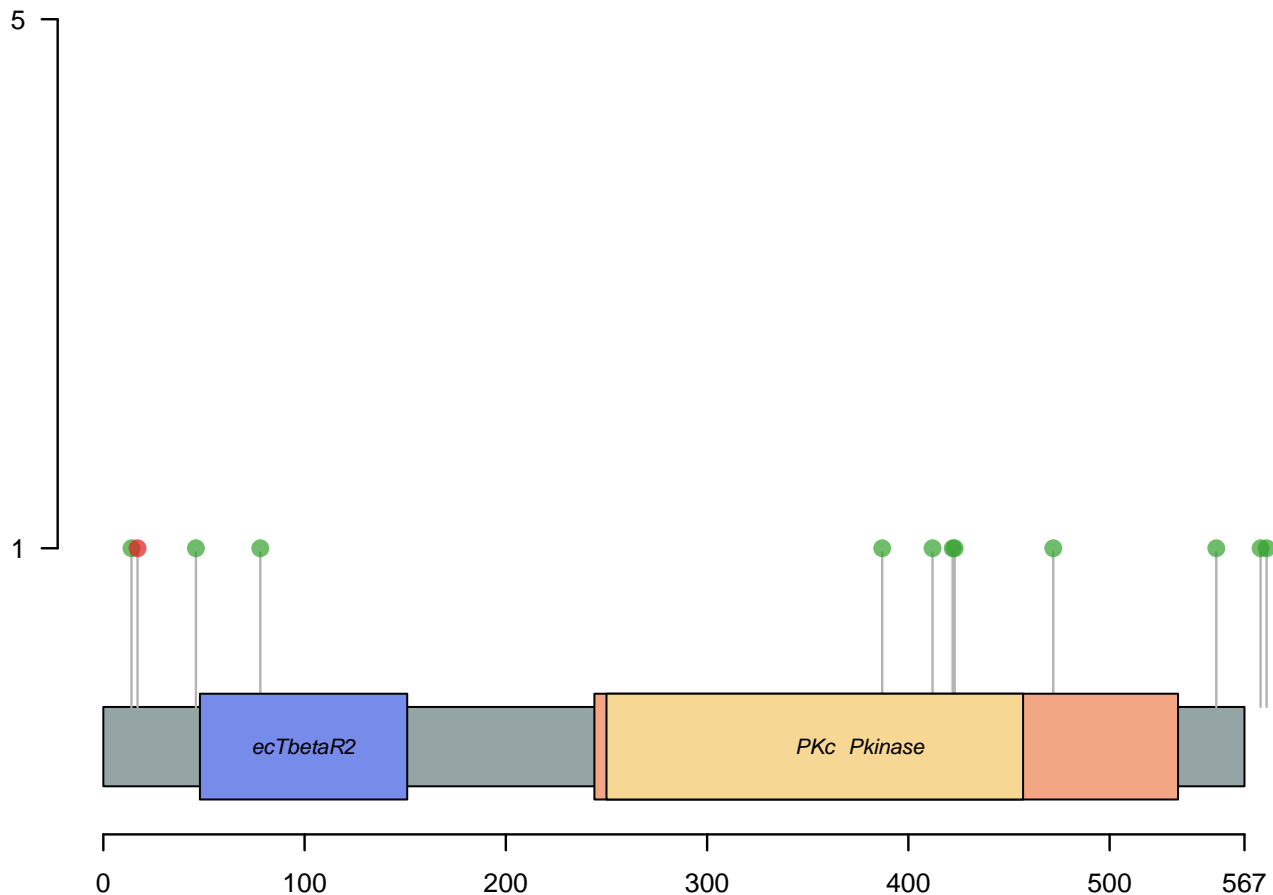


TGFBFR2 : [Somatic Mutation Rate: 1.23%]

NM_003242



● Missense_Mutation

● Nonsense_Mutation