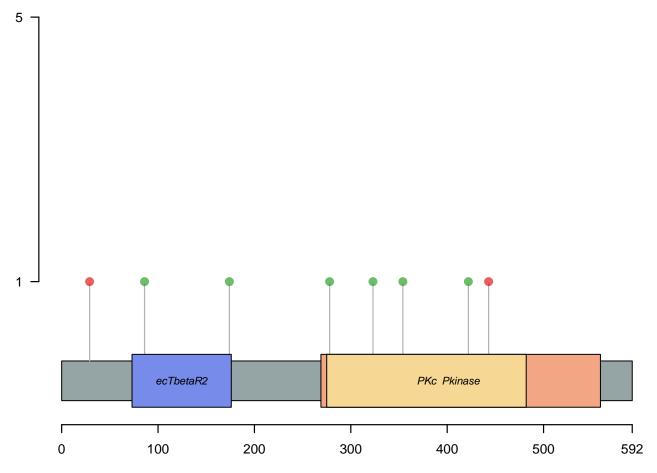
*TGFBR*2 : [Somatic Mutation Rate: 1.7%] NM_001024847



- Nonsense_Mutation
- Missense_Mutation