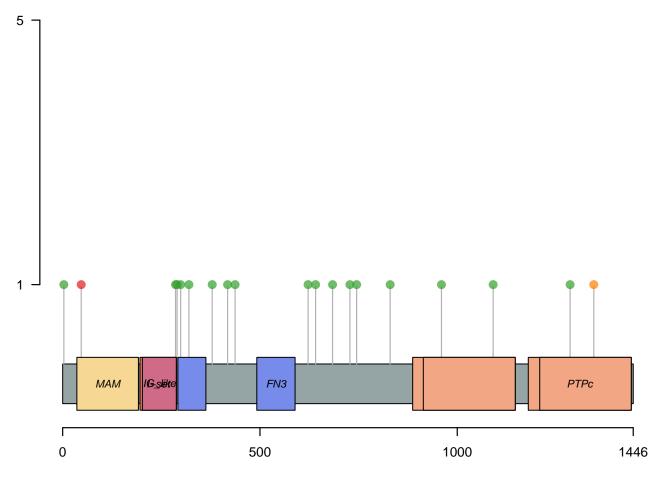
PTPRK: [Somatic Mutation Rate: 3.4%] NM_001135648



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site