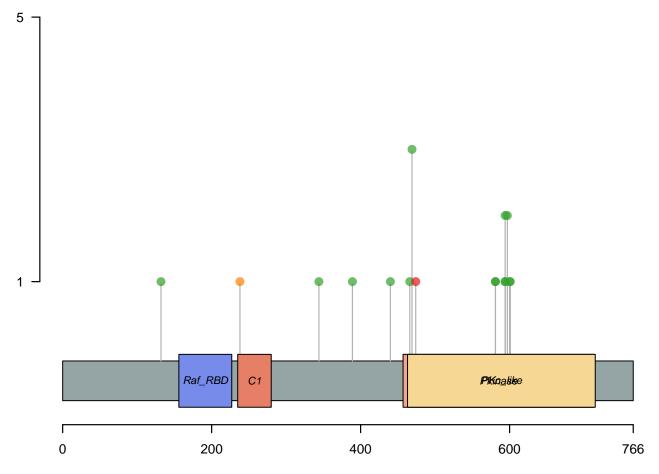
BRAF: [Somatic Mutation Rate: 2.09%] NM_004333



- Missense_Mutation
- Splice_Site
- Nonsense_Mutation