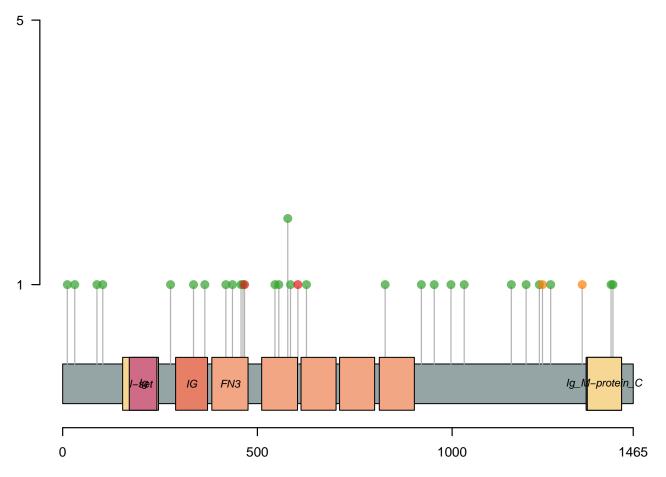
MYOM2: [Somatic Mutation Rate: 3.13%] NM_003970



- Missense_Mutation
- Nonsense_Mutation
- Splice_Site