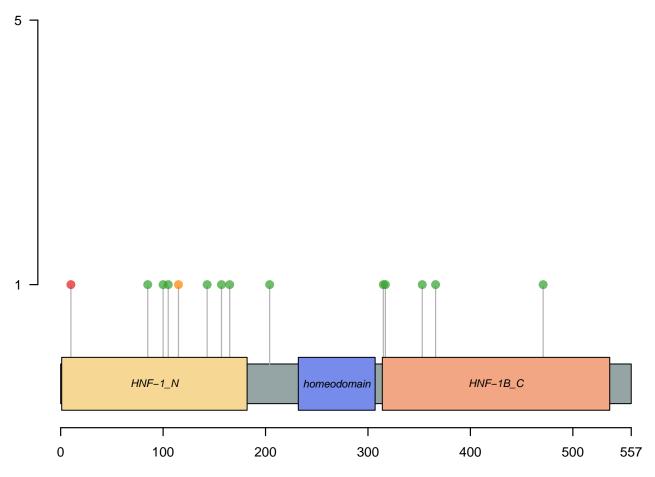
HNF1B: [Somatic Mutation Rate: 2.55%] NM_000458



- Nonsense_Mutation
- Missense_Mutation
- Splice_Site