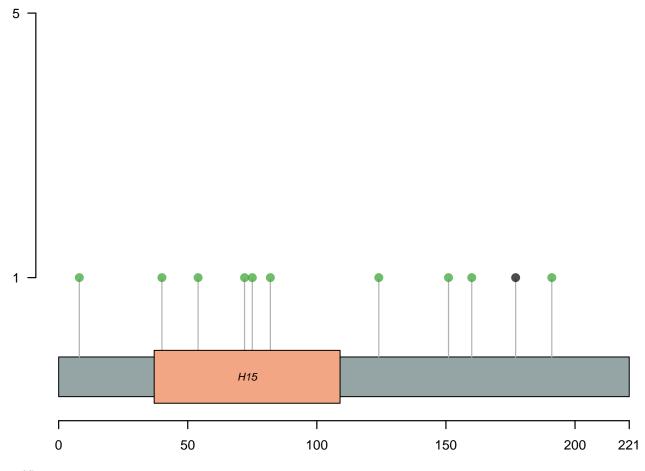
HIST1H1D: [Somatic Mutation Rate: 4.68%] NM_005320



- Missense
- Truncating