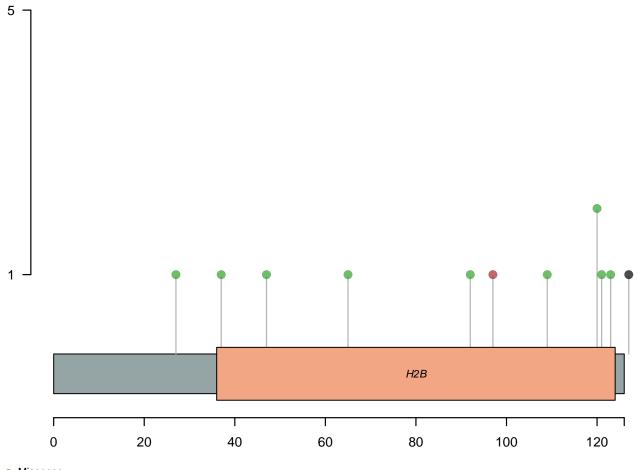
HIST1H2BK: [Somatic Mutation Rate: 4.68%] NM\_080593



- Missense
- In–frame
- Truncating