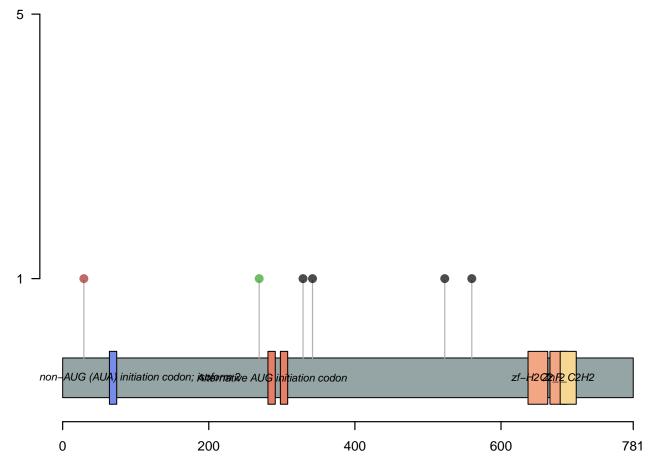
SP3 : [Somatic Mutation Rate: 2.55%] NM_003111



- In–frame
- Missense
- Truncating