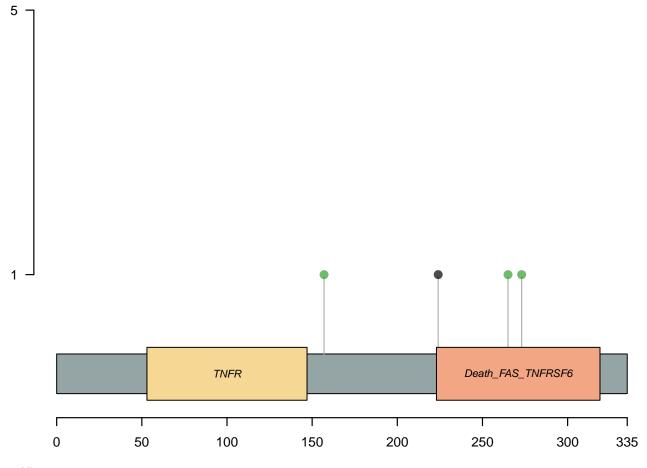
FAS: [Somatic Mutation Rate: 1.7%] NM\_000043



- Missense
- Truncating