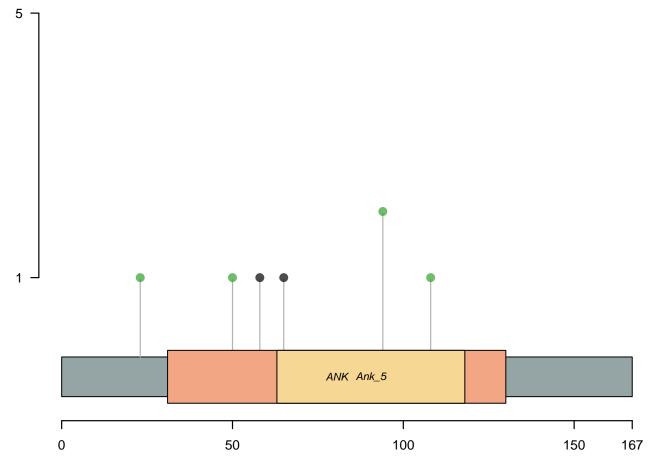
CDKN2A: [Somatic Mutation Rate: 2.98%] NM_001195132



- Missense
- Truncating