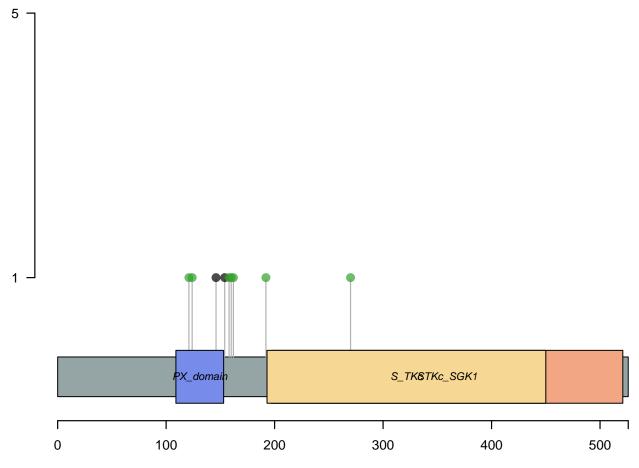
SGK1: [Somatic Mutation Rate: 2.55%] NM_001143676



- Missense
- Truncating