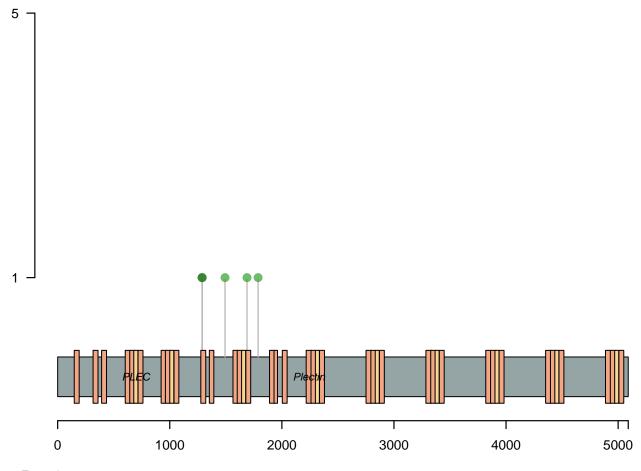
EPPK1: [Somatic Mutation Rate: 5.1%] NM_031308



- Truncating
- Missense