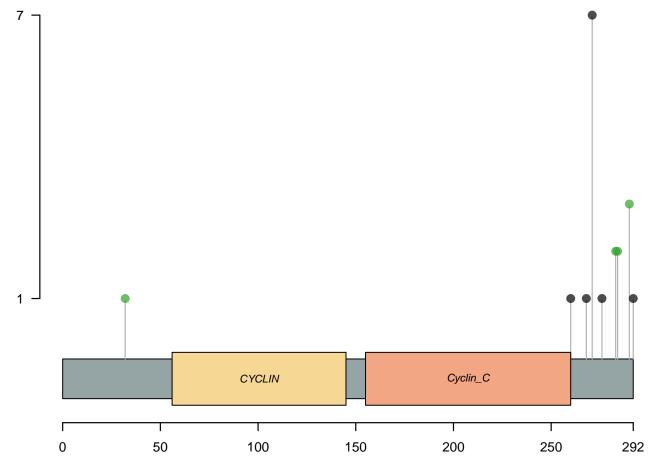
CCND3 : [Somatic Mutation Rate: 19.39%] NM_001760



- Missense
- Truncating