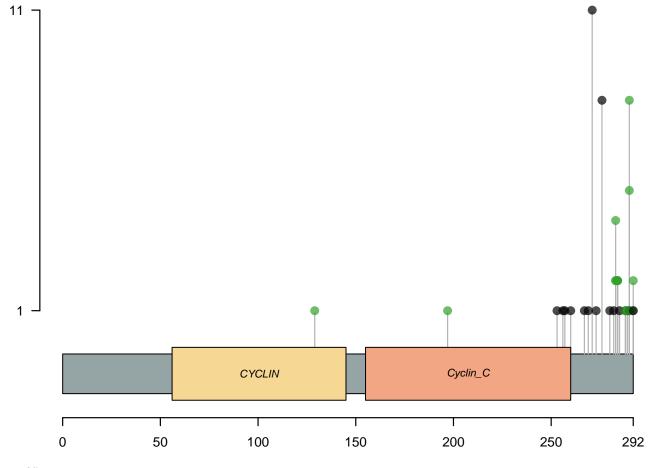
CCND3 : [Somatic Mutation Rate: 28.09%] NM_001760



- Missense
- Truncating