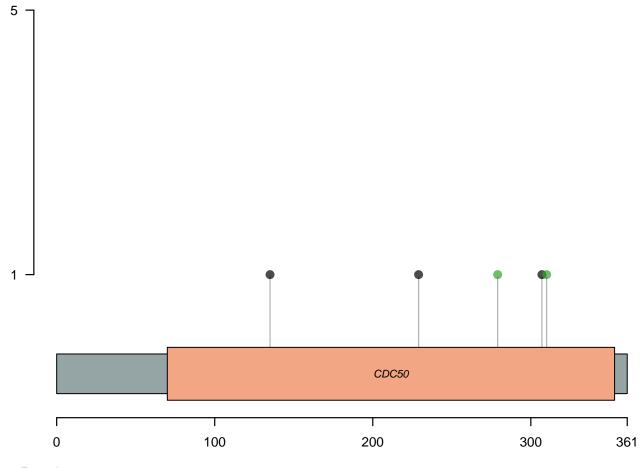
TMEM30A: [Somatic Mutation Rate: 1.28%] NM_018247



- Truncating
- Missense