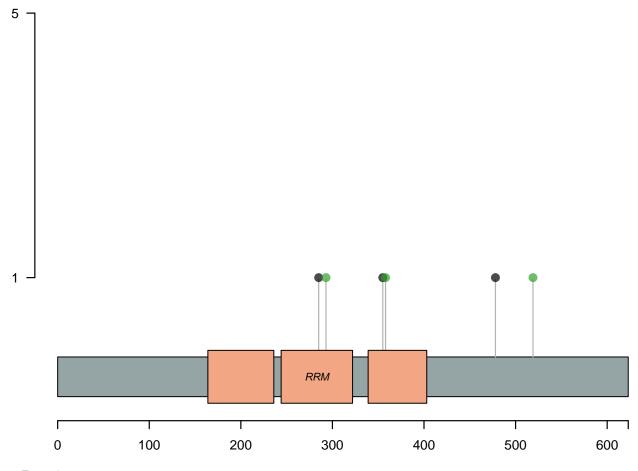
SYNCRIP: [Somatic Mutation Rate: 2.55%] NM_006372



- Truncating
- Missense