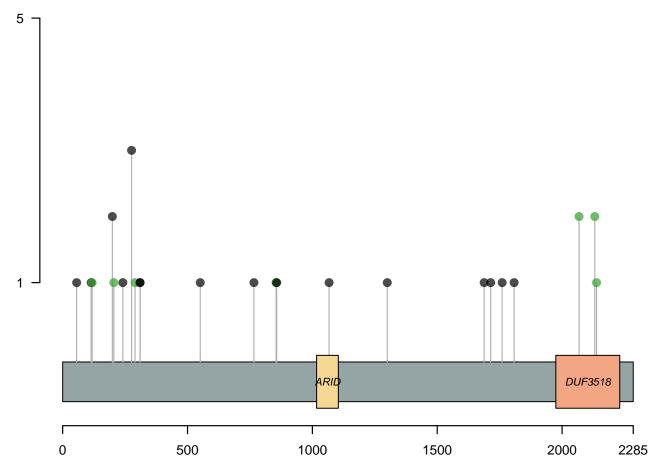
ARID1A: [Somatic Mutation Rate: 24.49%] NM\_006015



- Truncating
- Missense