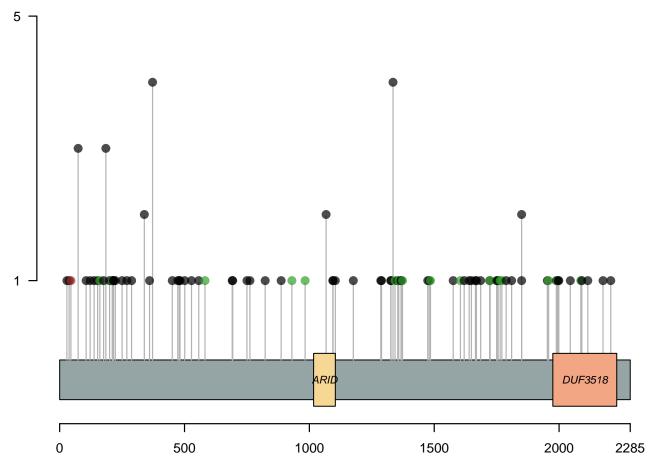
ARID1A: [Somatic Mutation Rate: 36.17%] NM_006015



- Truncating
- In–frame
- Missense