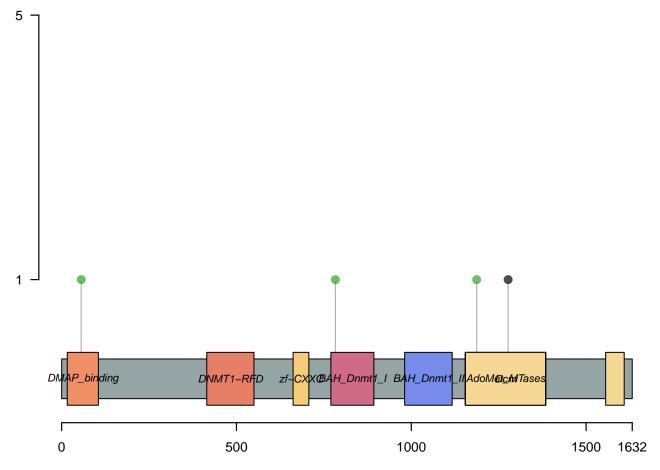
DNMT1: [Somatic Mutation Rate: 1.7%] NM_001130823



- Missense
- Truncating