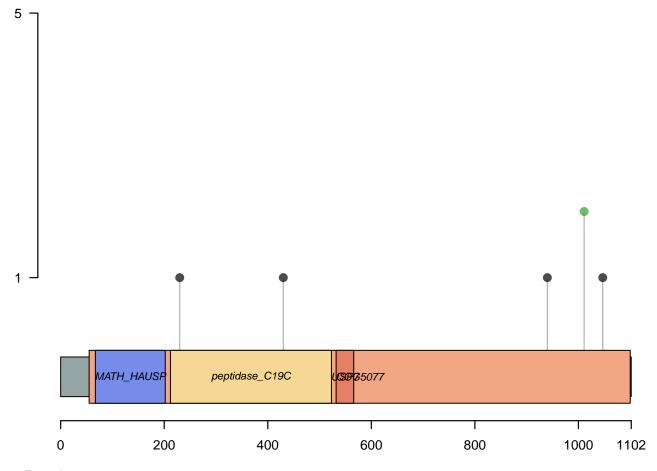
USP7 : [Somatic Mutation Rate: 6.12%] NM_003470



- Truncating
- Missense