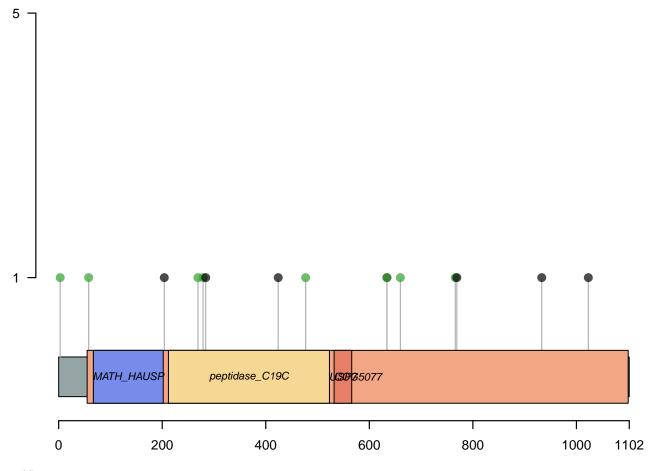
USP7 : [Somatic Mutation Rate: 5.96%] NM_003470



- Missense
- Truncating