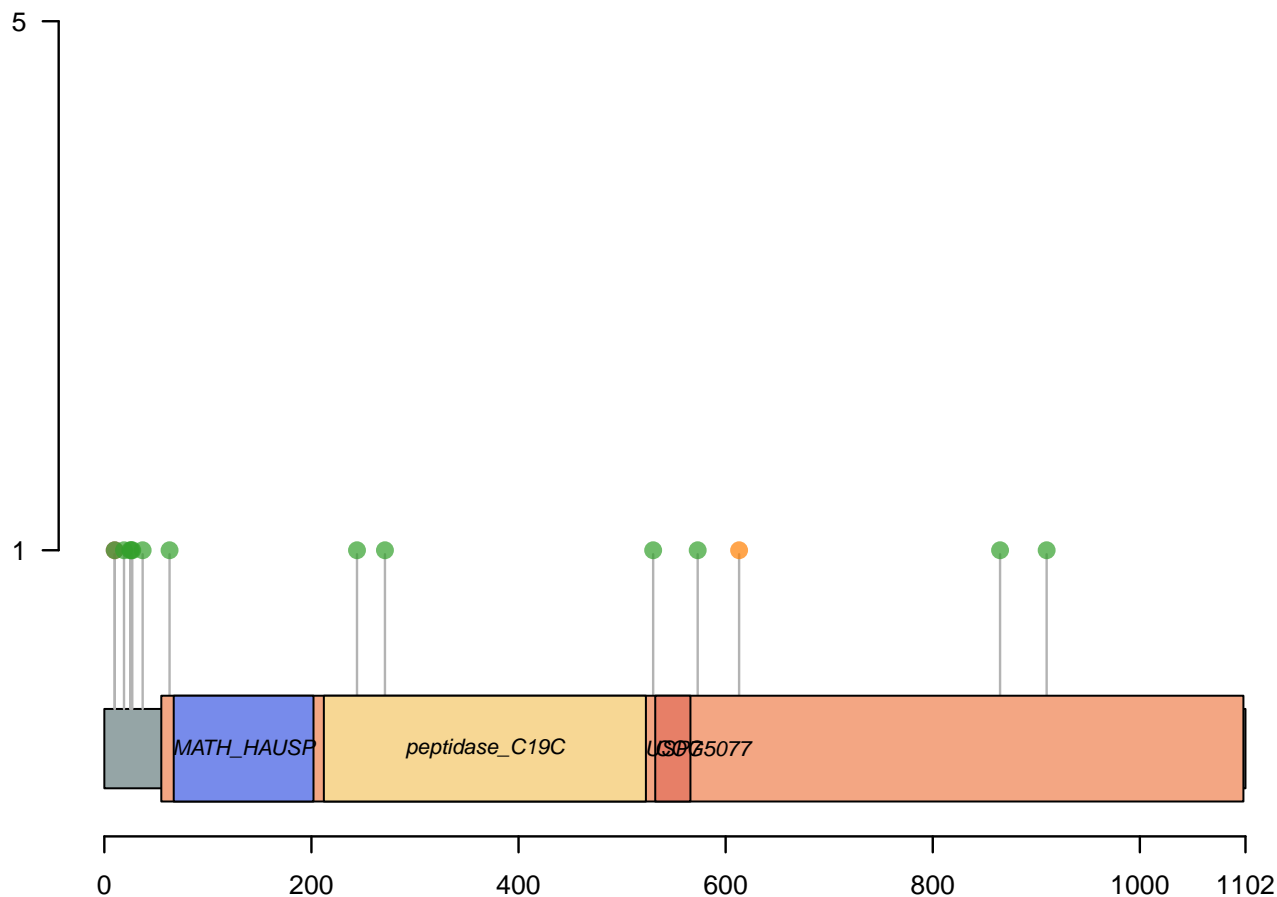


USP7 : [Somatic Mutation Rate: 2.98%]

NM_003470



- In_Frame_Ins
- Missense_Mutation
- Splice_Site