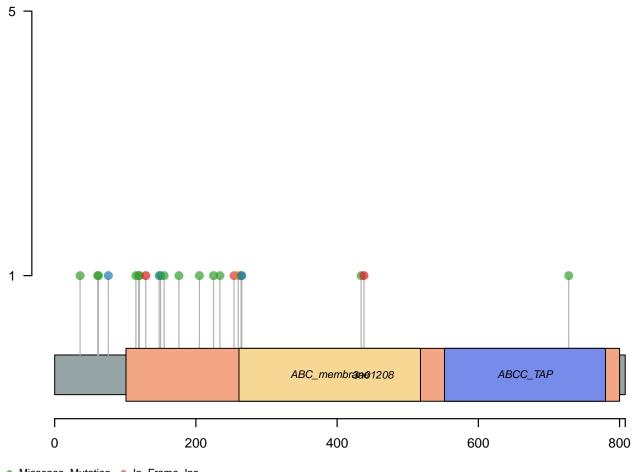
TAP1: [Somatic Mutation Rate: 3.62%] NM_000593



- Missense_MutationIn_Frame_Ins
- Frame_Shift_Del Splice_Site
- Nonsense_Mutation