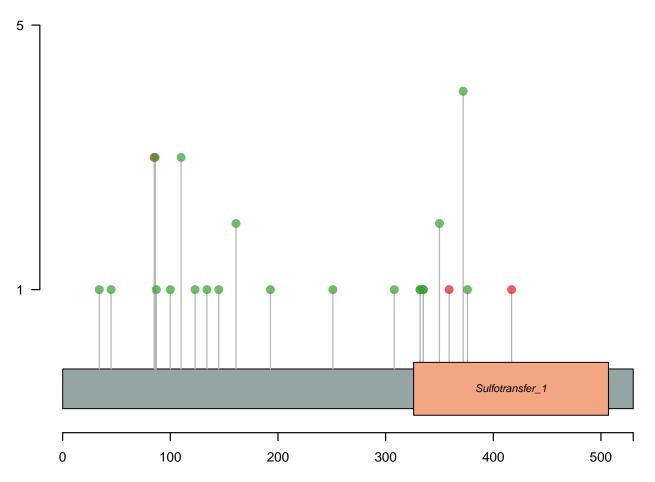
CHST2: [Somatic Mutation Rate: 3.1%] NM_004267



- Missense_Mutation
- Nonsense_Mutation