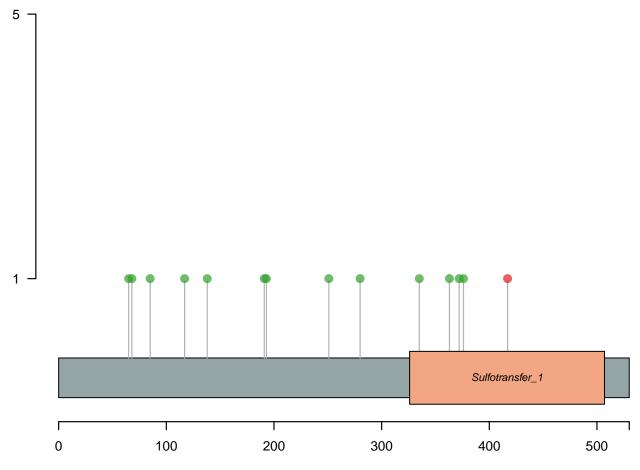
CHST2: [Somatic Mutation Rate: 0.73%] NM_004267



- Missense_Mutation
- Nonsense_Mutation