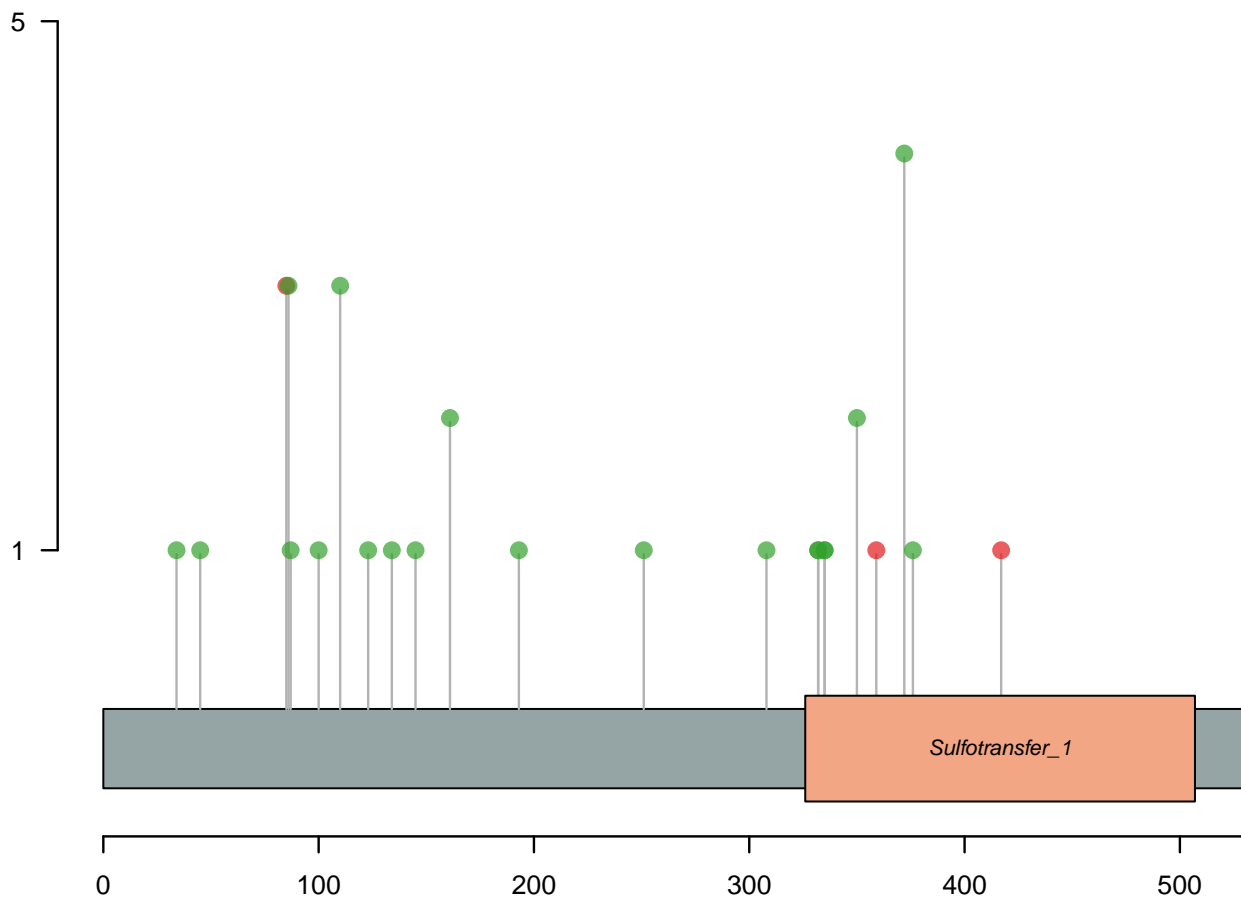


CHST2 : [Somatic Mutation Rate: 3.1%]
NM_004267



● Missense_Mutation
● Nonsense_Mutation