FAS : [Somatic Mutation Rate: 4.38%]
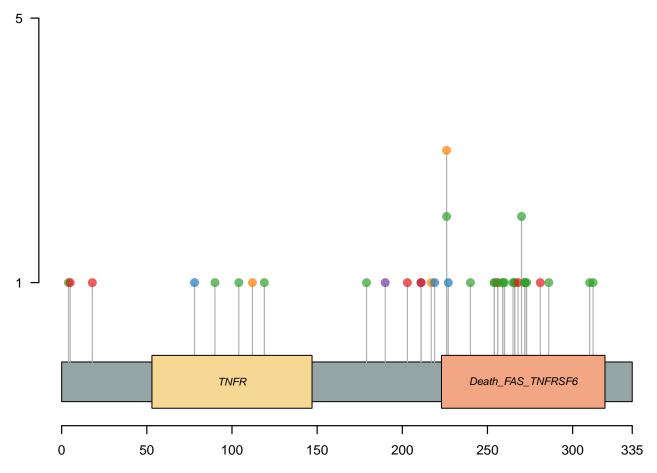NM_000043