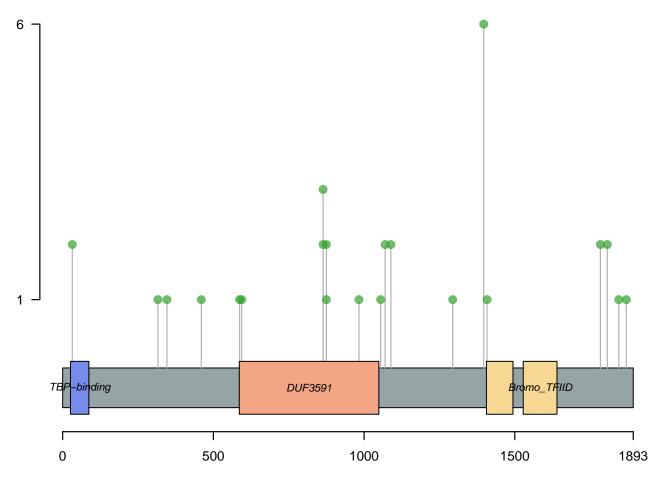
TAF1: [Somatic Mutation Rate: 3.51%] NM_004606



Missense_Mutation