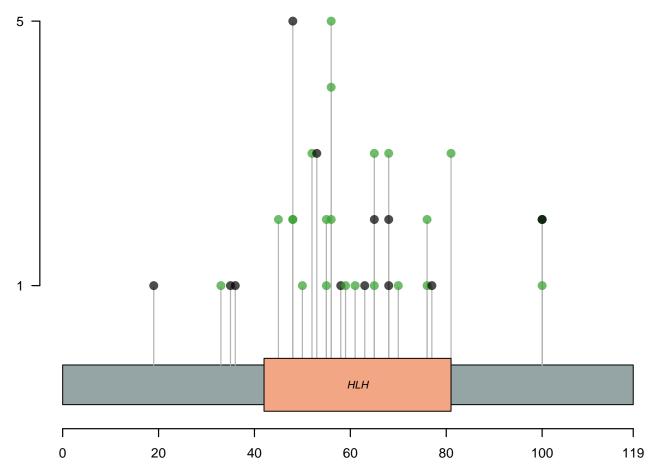
ID3 : [Somatic Mutation Rate: 33.67%] NM_002167



Truncating

Missense