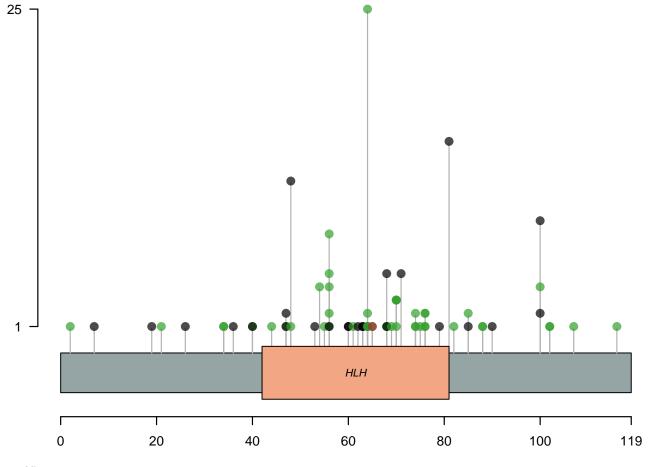
ID3 : [Somatic Mutation Rate: 44.26%] NM_002167



Missense

Truncating

In–frame