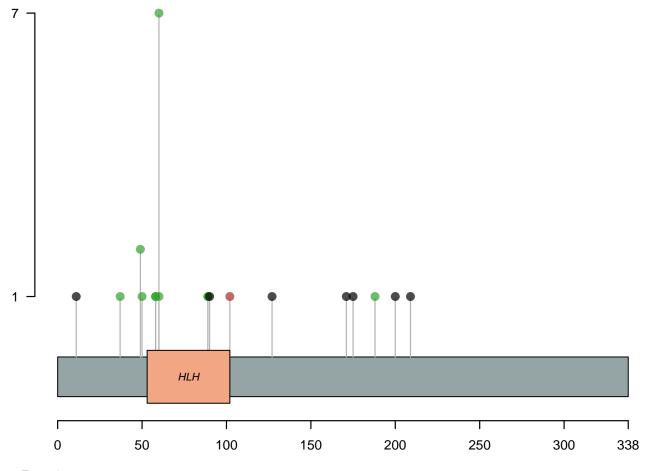
TFAP4: [Somatic Mutation Rate: 10.64%] NM_003223



- Truncating
- Missense
- In–frame