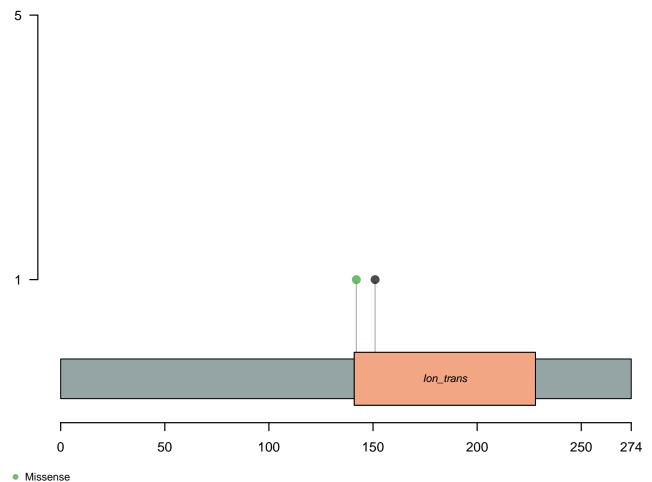
HVCN1 : [Somatic Mutation Rate: 0.85%] NM_001040107



- Truncating