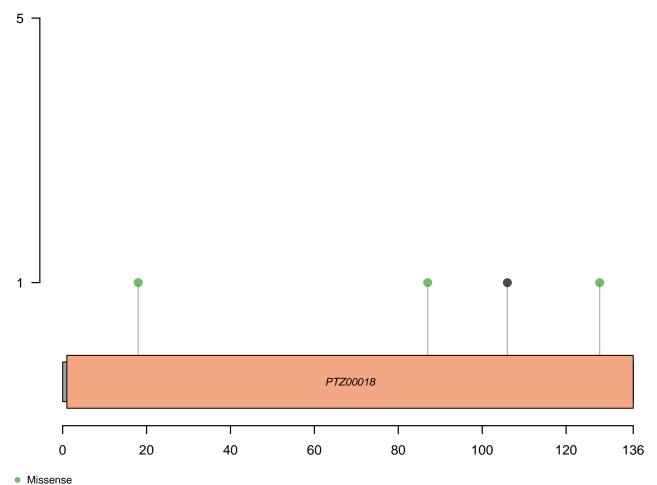
HIST1H3D: [Somatic Mutation Rate: 1.7%] NM_003530



- Truncating