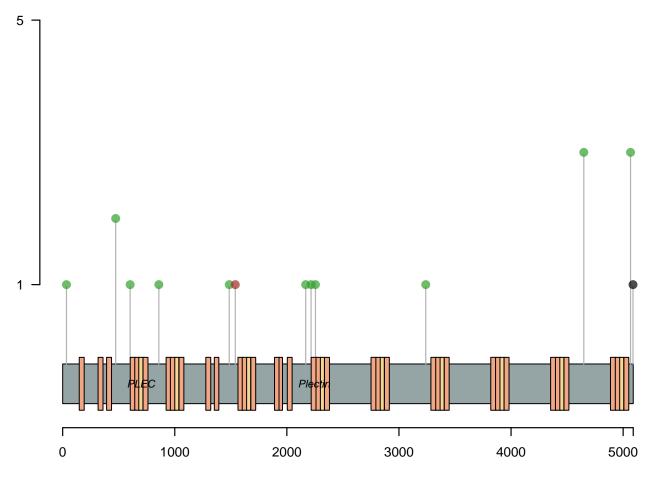
EPPK1: [Somatic Mutation Rate: 7.66%] NM_031308



Missense

In–frame

Truncating