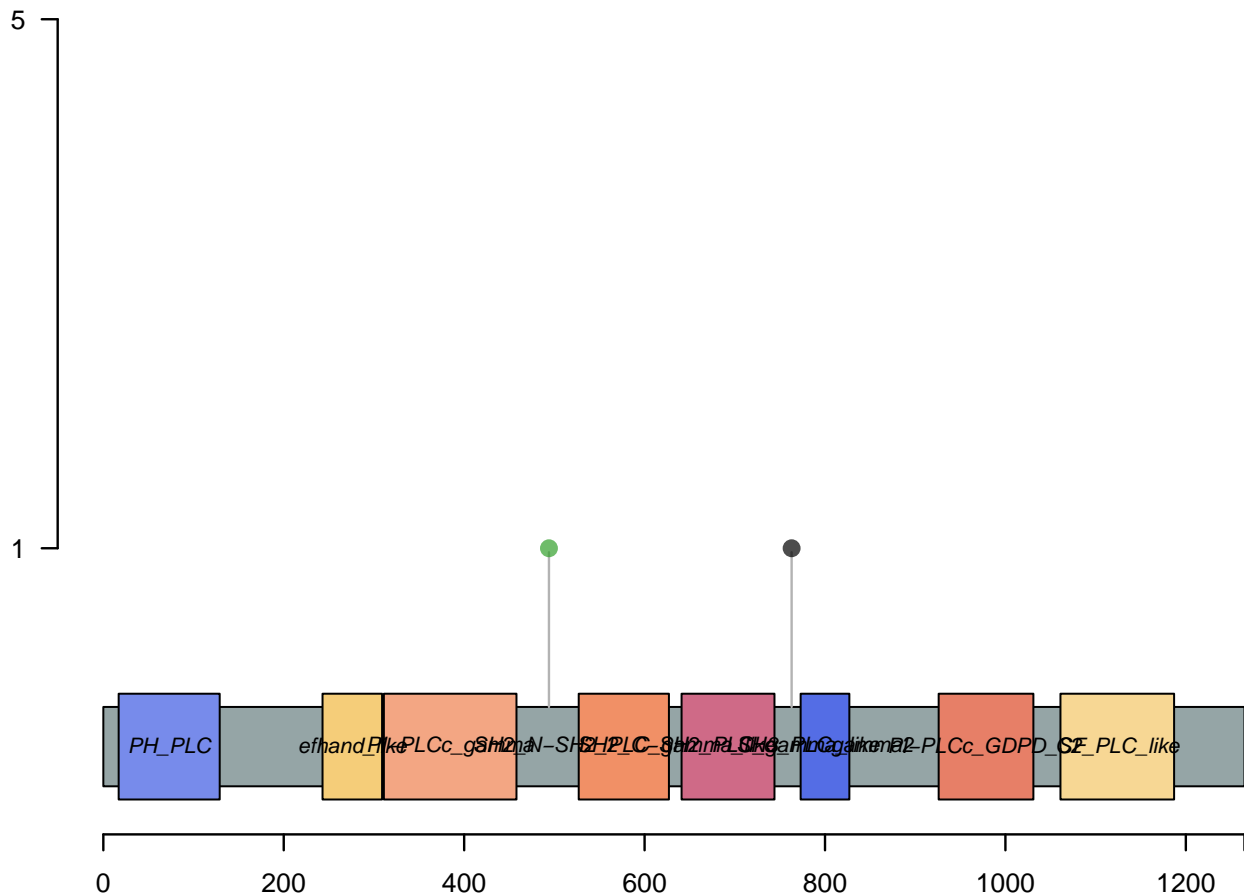


PLCG2 : [Somatic Mutation Rate: 3.06%]

NM_002661



● Missense

● Truncating