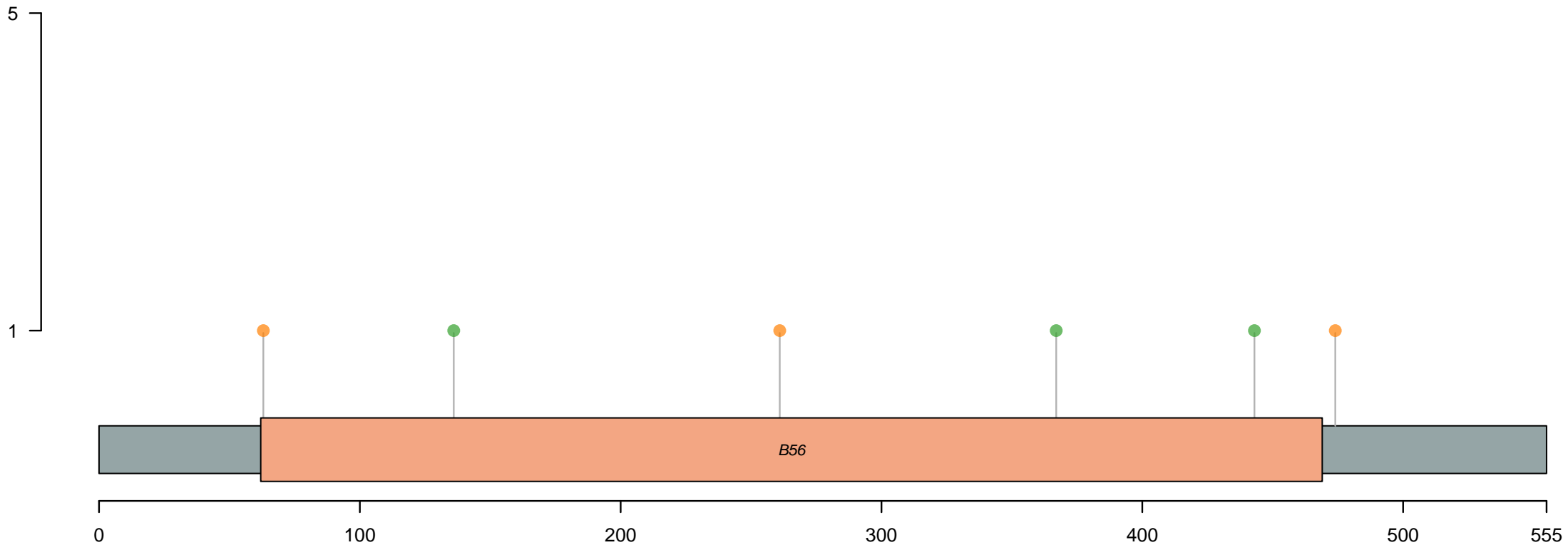


PPP2R5C : [Somatic Mutation Rate: 0.64%]

NM_001161725



● Splice_Site

● Missense_Mutation