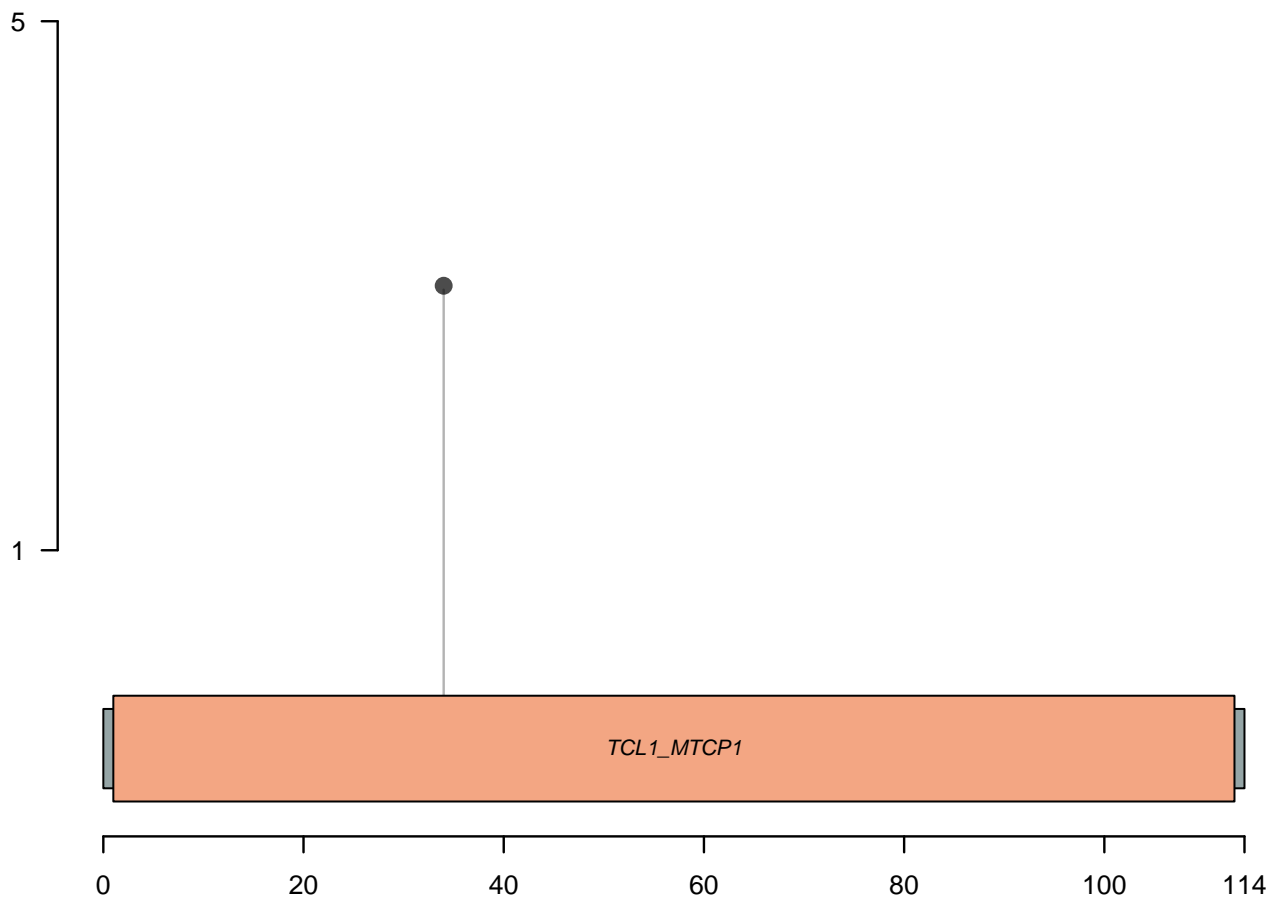


TCL1A : [Somatic Mutation Rate: 3.06%]

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● Truncating