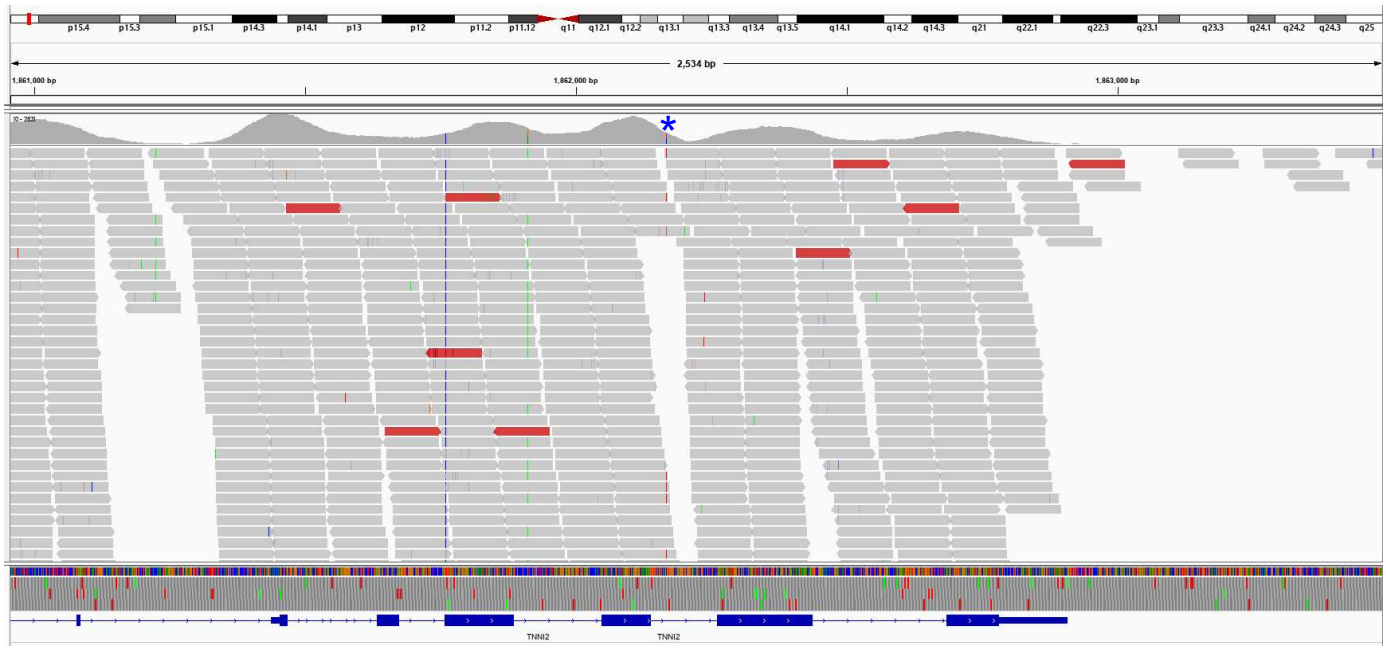
Supplemental Figure S1

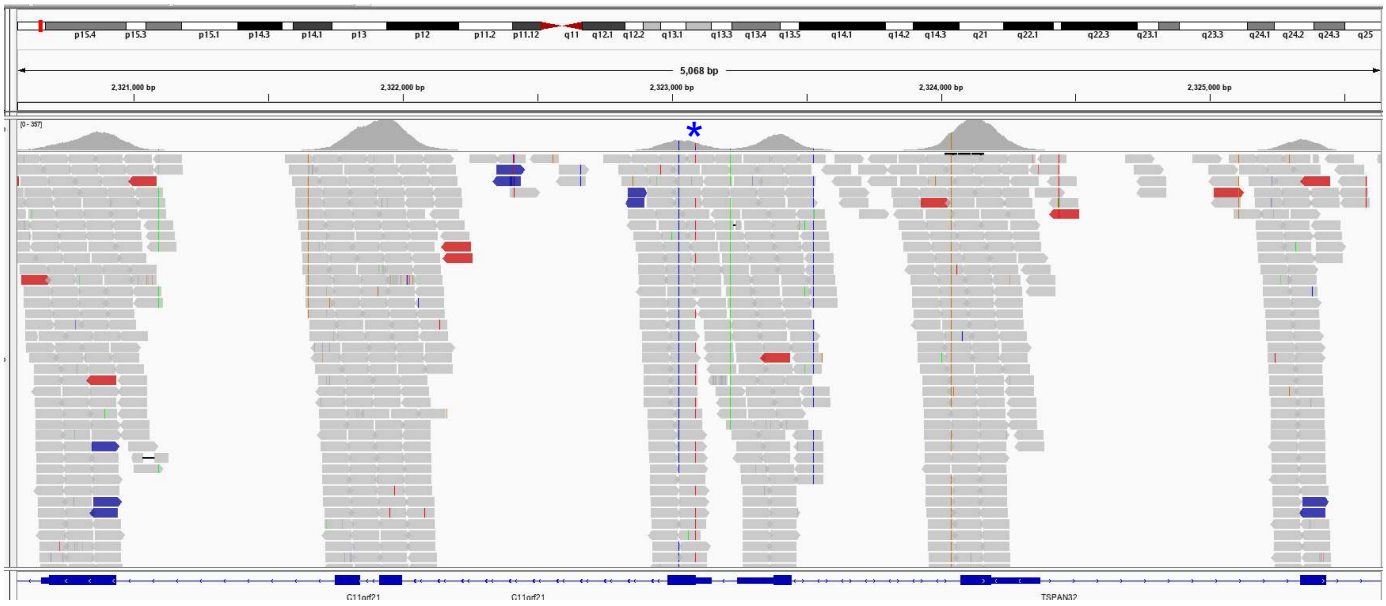
A)

B)

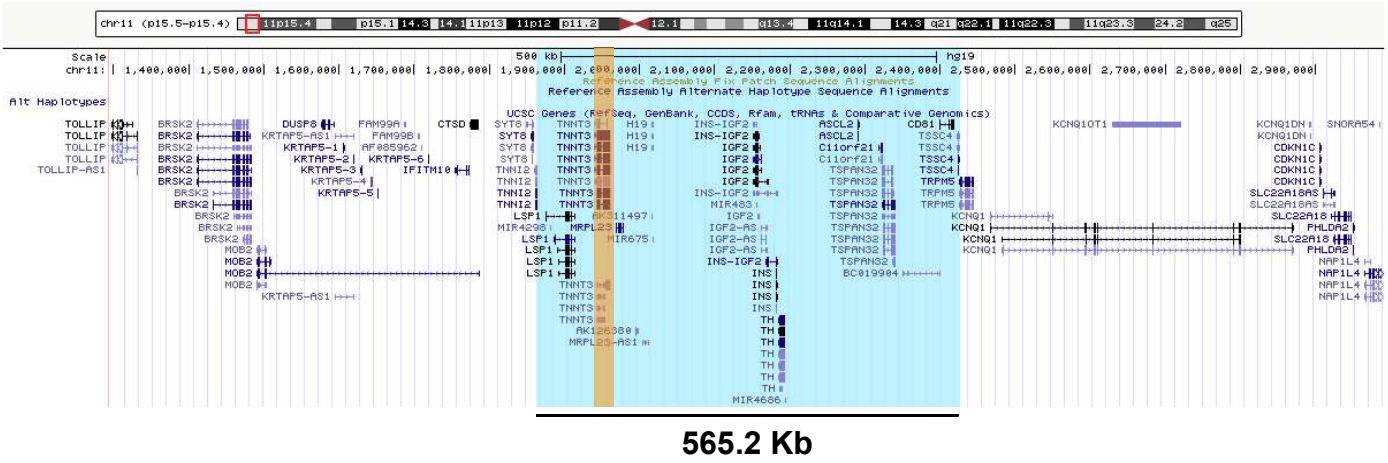
11:1862168



11:2323089



C)



Supplemental Figure S1: Manual inspection of patient's BAM file reveals 565.2 Kb AOH block

(A) Integrative Genomics Viewer (IGV) screenshot of patient's BAM file showing *TNNT3* sequencing data. All polymorphisms are homozygous (indicated by black asterisk). The pathogenic variant c.418-1G>A is indicated by a red asterisk.

(B) Manual inspection revealed a 565.2 Kb AOH block surrounding *TNNT3*:c.418-1G>A. This region was flanked by at least two consecutive heterozygous polymorphisms starting at 11:1862168 and 11:2427364.

(C) Screenshot from UCSC Genome Browser (https://genome.ucsc.edu/) showing AOH block (blue) and *TNNT3* (orange).