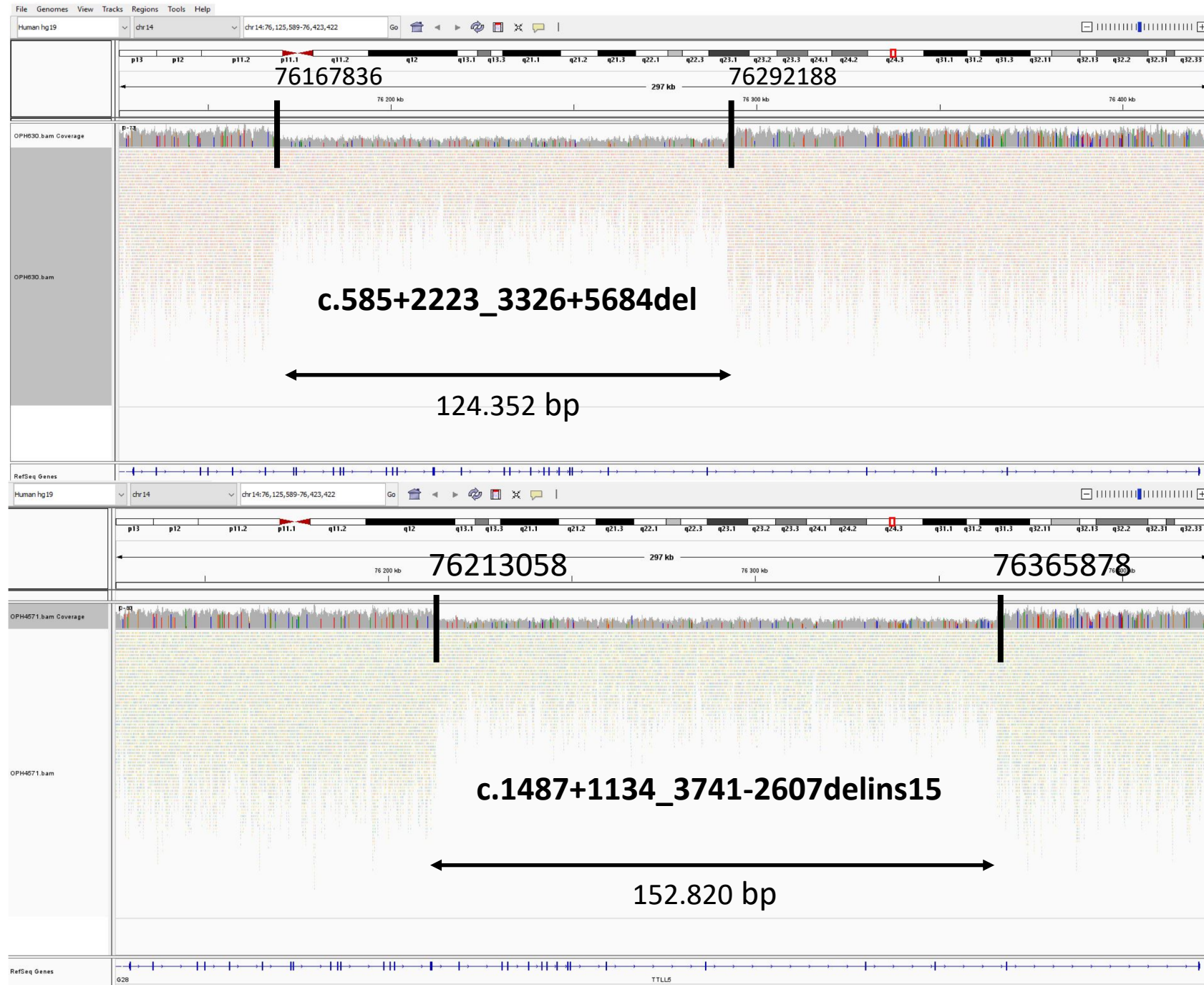


# *TTLL5* gene



Genomic position

Depth of coverage:  
47x

Depth of coverage:  
44x

**Supplemental Figure S3: Large multiexon deletions depicted by IGV screenshots.** Patients XZ-358338 and IM-4476 were heterozygous for large intragenic deletions, encompassing exons 8 to 28 and exons 18 to 30, respectively. Exact breakpoints were detected by whole genome sequencing followed by targeted *TTL5* gene analysis. The mean depth of coverage out of the deletions was 47 and 44, respectively.