

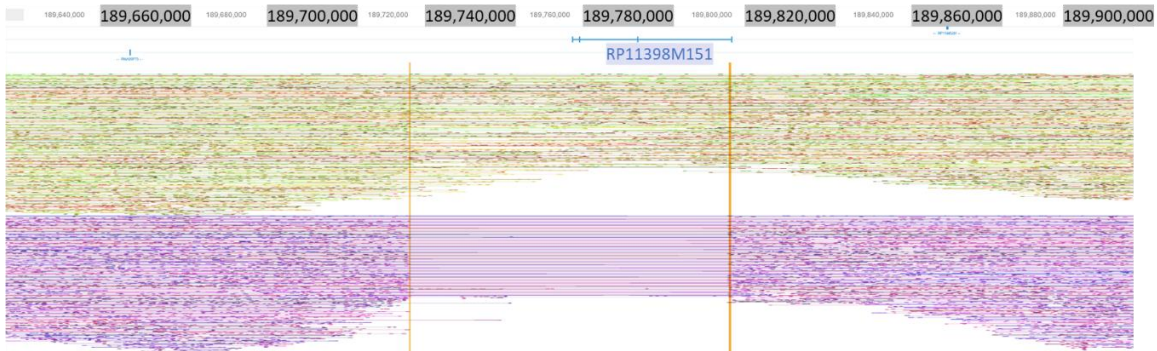
A

hg19: Chr5:104432113-104503673 -> hg38:Chr5:105096412-105167972



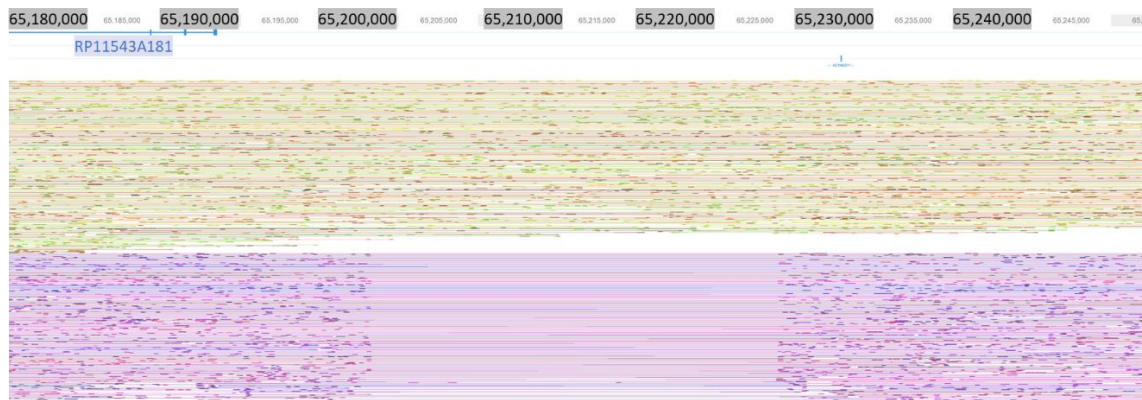
B

hg19: Chr1:189704509-189783359 -> hg38: Chr1:189735379-189814229



C

hg19: Chr3:65189000-65213999 -> hg38: Chr3:65203325-65228324



Supplemental Figure S3. Phased read graphs from TELL-Seq data on the NA12878 sample. **(A)** TELL-Seq data showed a heterozygous deletion on Chromosome 5: 104,432,113-104,503,673. 10x data (Zheng et al. 2016) called this region as a heterozygous deletion, while stLFR data (Wang et al. 2019) called this region as a homozygous deletion. **(B)** A heterozygous deletion on Chromosome 1: 189,704,509-189,783,359. **(C)** A heterozygous deletion on Chromosome 3: 65,189,000-65,213,999. GRCh38(hg38) coordinates were used for visualization data.