

SUPPLEMENTARY FILE

An inversion affecting the *GCHI* gene as a novel finding in dopamine-responsive dystonia

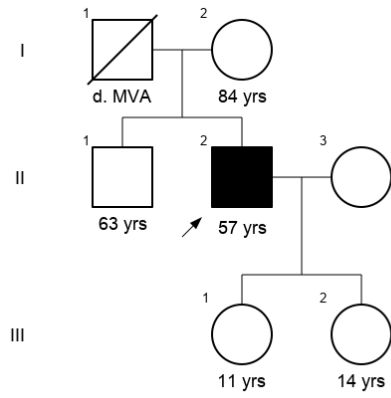
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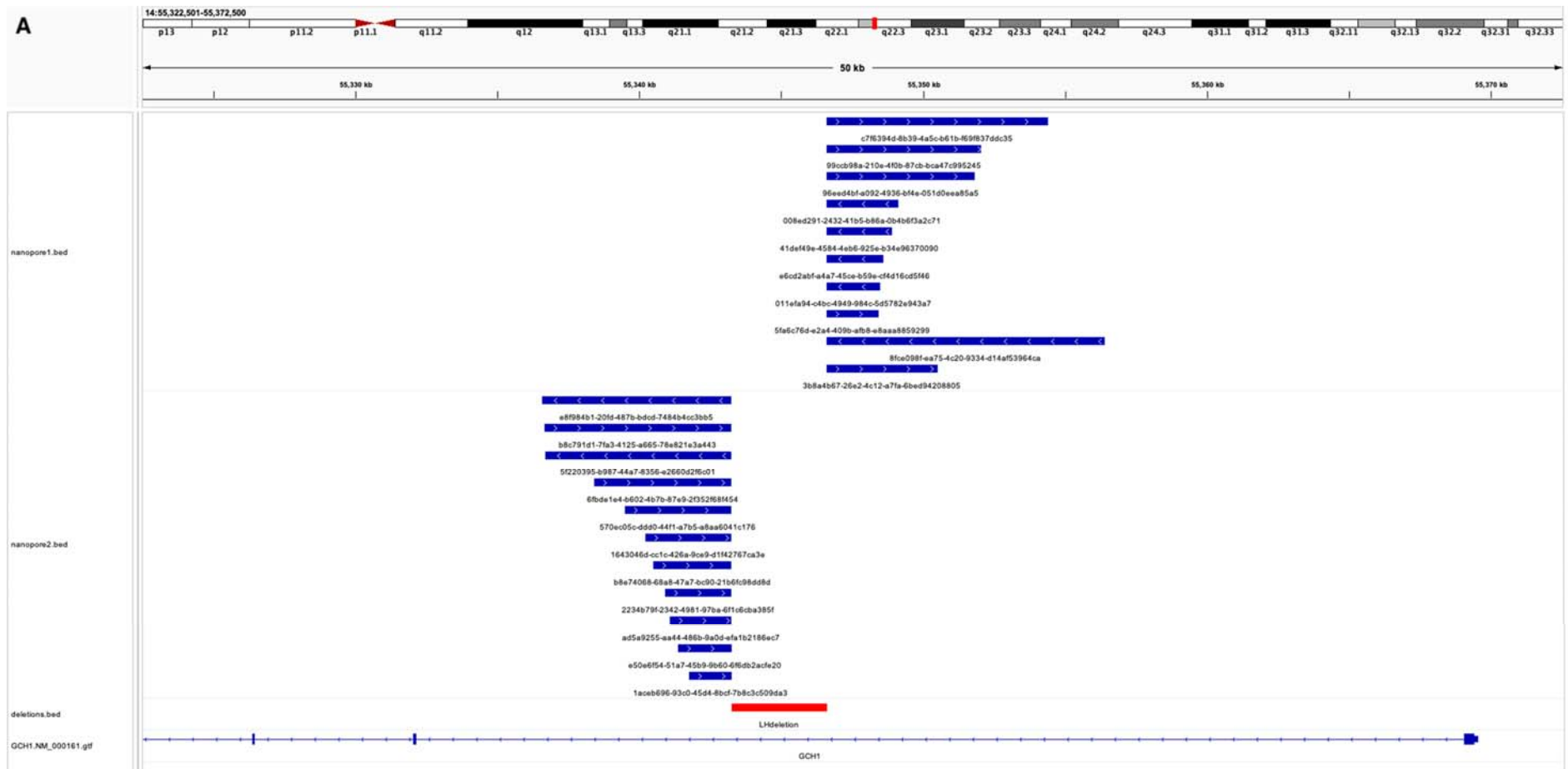
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Supplementary Figure S1. Pedigree of the family. Arrow indicates proband who underwent genetic studies, filled symbol indicates affected, squares represent males and circles represent females. d. MVA = died in motor vehicle accident. Additional family members were not available for testing.



Supplementary Figure S3. Oxford Nanopore long read sequencing (LRS) data supporting the proposed structural variant. Chimaeric nanopore sequences are visualized in the IGV genome browser. The sequence alignments confirm the breakpoints indicated by short read sequencing analysis, and extend wide enough for a high level of confidence in read locations. **(A, B)** Inversion breakpoint regions as in Fig. S2, but in a 50 kb window.



B

