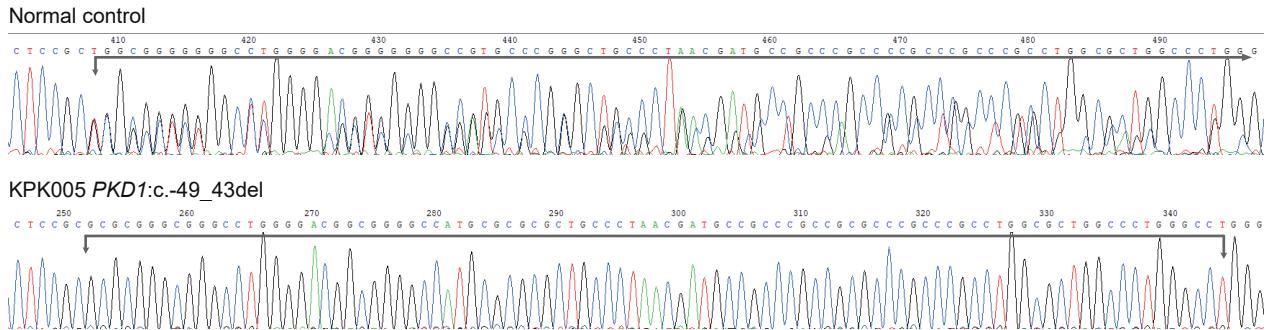
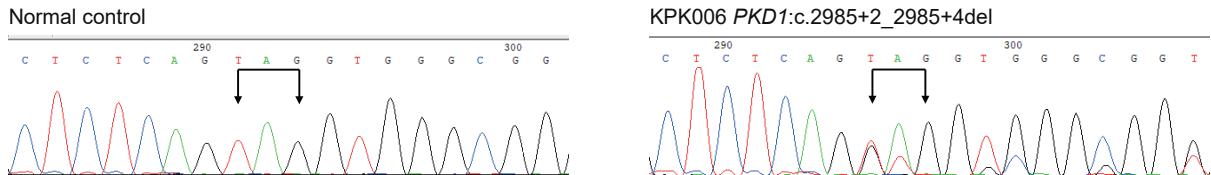


Supplemental Figure 2

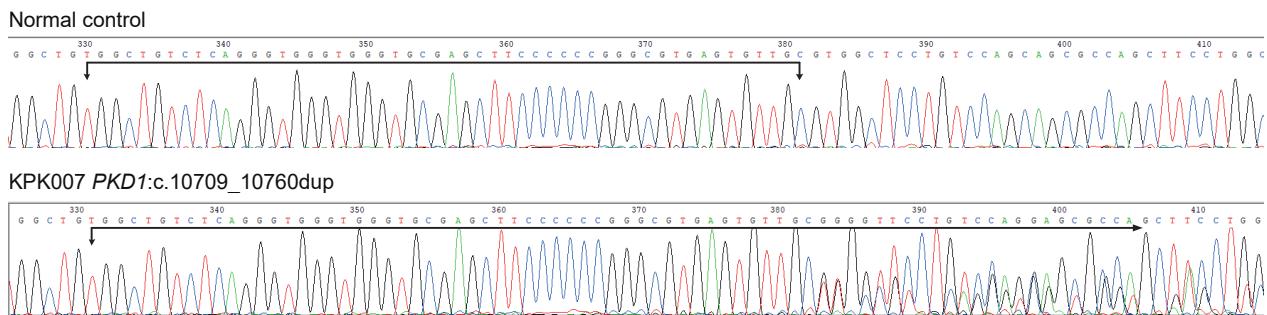
A



B



C



D

PKD1 Intron 42 reference

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CGGGACGGCGTAC
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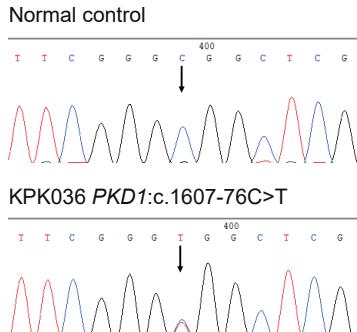
VNTR

PKD1:c.11713-30_11713-63dup

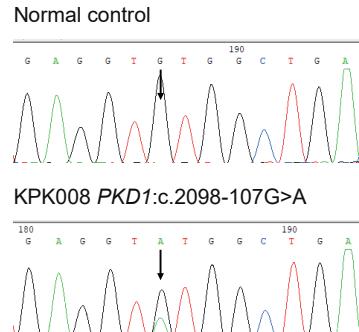
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VNTR

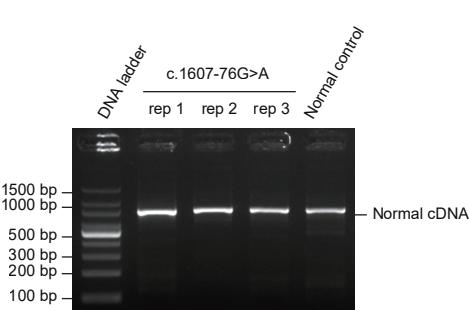
E



F



G



Supplemental Figure 2. Confirmation of variants identified by targeted LRS. (A-C) Confirmation of *PKD1:c.-49_+43del* in KPK005 (A), *PKD1:c.2985+2_2985+4del* in KPK006 (B), and *PKD1:c.10709_10760dup* in KPK007 (C) by Sanger sequencing. (D) Display of *PKD1:c.11713-30_11713-63dup* in an array of variable number of tandem repeats (VNTR) in intron 42. (E-F) Confirmation of *PKD1:c.1607-76C>T* in KPK036 (E), and *PKD1:c.2098-107G>A* in KPK008 (F). (G) Amplification of cDNA showing that mRNA splicing was not affected by *PKD1:c.1607-76C>T* in KPK036.