



Testing is performed to evaluate for the presence of variants in coding regions and extending to +/-10 base pairs of adjacent intronic sequence on either side of the coding exons of the genes analyzed. In addition, the analysis may cover select non-coding variants. Next-generation sequencing and/or a polymerase chain reaction-based quantitative method is performed to test for the presence of copy number variants (CNV) in the genes analyzed. Confirmation of select reportable variants may be performed by alternate methodologies based on internal laboratory criteria. To verify if a specific region/exon/variant is covered by this assay or to confirm transcript version used, contact a laboratory genetic counselor at 800-533-1710.

Genomic Build: GRCh37 (hg19) unless otherwise specified

Gene	Reference Transcript	Additional Evaluations	Technical Limitations
<i>ALG8</i>	NM_024079.5	-	-
<i>ALG9</i>	NM_024740.2 [GRCh38(hg38)]	-	-
<i>ANKS6</i>	NM_173551.5	-	CNV analysis in exon 1 is not performed
<i>BICC1</i>	NM_001080512.3	-	-
<i>CEP83</i>	NM_016122.3	-	-
<i>CEP104</i>	NM_014704.4	-	-
<i>CEP164</i>	NM_014956.5	-	-
<i>CEP290</i>	NM_025114.4	chr12:88462434A>T (c.6012-12T>A)	CNV analysis in exons 5, 13, and 53 is not performed
<i>COL4A1</i>	NM_001845.6	-	CNV analysis in exon 16 is not performed
<i>CRB2</i>	NM_173689.7	-	-
<i>DCDC2</i>	NM_016356.5	-	-
<i>DNAJB11</i>	NM_016306.5	-	-
<i>DZIP1L</i>	NM_173543.3	-	-
<i>GANAB</i>	NM_198335.4	-	-
<i>GLIS2</i>	NM_032575.2	-	-
<i>HNF1B</i>	NM_000458.4	-	-
<i>INVS</i>	NM_014425.5	-	-
<i>IQCB1</i>	NM_001023570.4	-	CNV analysis in exon 7 is not performed
<i>JAG1</i>	NM_000214.3	-	-
<i>LRP5</i>	NM_002335.4	-	-
<i>MAPKBP1</i>	NM_001128608.2	-	-
<i>NEK8</i>	NM_178170.3	-	-
<i>NOTCH2</i>	NM_024408.4	-	Analyses for sequence variants and CNV are not performed for exons 1–4
<i>NPHP1</i>	NM_000272.4	-	CNV analysis in exon 3 is not performed
<i>NPHP3</i>	NM_153240.5	-	CNV analysis in exons 8, 22 is not performed
<i>NPHP4</i>	NM_015102.5	-	-
<i>OFD1</i>	NM_003611.3	-	-
<i>PAX2</i>	NM_003987.4	-	-
<i>PAX2</i>	NM_003988.5	-	-

Targeted Genes and Methodology Details
for Cystic Kidney Disease Gene Panel (continued)

Gene	Reference Transcript	Additional Evaluations	Technical Limitations
<i>PKD1</i>	NM_001009944.3	-	CNV analysis in exons 1, 26 is not performed
<i>PKD2</i>	NM_000297.4	-	CNV analysis in exon 9 is not performed
<i>PKHD1</i>	NM_138694.4	-	-
<i>RPGRIP1L</i>	NM_015272.5	-	CNV analysis in exon 23 is not performed
<i>SDCCAG8</i>	NM_006642.5	-	CNV analysis in exon 7 is not performed
<i>SEC61A1</i>	NM_013336.4	-	-
<i>SEC63</i>	NM_007214.5	-	CNV analysis in exon 2 is not performed
<i>TMEM67</i>	NM_153704.6	-	-
<i>TRAF3IP1</i>	NM_015650.4	-	-
<i>TSC1</i>	NM_000368.5	-	-
<i>TSC2</i>	NM_000548.5	-	-
<i>TTC21B</i>	NM_024753.5	-	-
<i>UMOD</i>	NM_003361.3	-	-
<i>WDR19</i>	NM_025132.4	-	CNV analysis in exon 3 is not performed
<i>WDR35</i>	NM_001006657.2	-	-
<i>XPNPEP3</i>	NM_022098.4	-	-