

Targeted Genes and Methodology Details for Focused Autosomal Dominant Polycystic Kidney Disease Panel

The following applies to ADPKP/Focused Autosomal Dominant Polycystic Kidney Disease Panel. 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 will 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.

This test was updated August 2022. This document is intended to highlight additional evaluations for variants of high clinical interest as well as technical limitations. However, this document does not comprehensively reflect all genomic regions covered by this test. For questions regarding transcripts, genes or regions covered on this version or prior versions of this test, contact the laboratory at 800-533-1710.

Gene	Reference Transcript	Additional Evaluations	Technical Limitations
<i>ALG8</i>	NM_024079.5	-	-
<i>ALG9</i>	NM_024740.2 [GRCh38(hg38)]	-	-
<i>DNAJB11</i>	NM_016306.5	-	-
<i>GANAB</i>	NM_198335.4	-	-
<i>HNF1B</i>	NM_000458.4	-	-
<i>PKD1</i>	NM_001009944.3	-	CNV analysis in exons 1 and 26 is not performed
<i>PKD2</i>	NM_000297.4	-	CNV analysis in exon 9 is not performed
<i>UMOD</i>	NM_003361.3	-	-

Effective Date	Version	Synopsis of Test Change
9/2/2025	V2	Added Technical Limitations and Additional Evaluations