ARL13B CLCNKB IQCB1 SLC3A1 MAGI2 SLC22A12 ACE CLDN16 KAL1 SLC4A1 SGPL1 SLC2A2 ACTN4 CLDN19 KCNJ1 SLC4A4 CFH SLC2A9 ADCK4 CNNM2 KIF7 SLC5A2 CFI SLC7A7 AGT COL4A1 KLHL3 SLC7A9 CFHR5 SLC9A3R1 AGTR1 COL4A3 LAMB2 SMARCAL1 CD46 TCTN3 AGXT COL4A4 LMX1B SOX17 C3 THBD AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10
ACTN4 CLDN19 KCNJ1 SLC4A4 CFH SLC2A9 ADCK4 CNNM2 KIF7 SLC5A2 CFI SLC7A7 AGT COL4A1 KLHL3 SLC7A9 CFHR5 SLC9A3R1 AGTR1 COL4A3 LAMB2 SMARCAL1 CD46 TCTN3 AGXT COL4A4 LMX1B SOX17 C3 THBD AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
ADCK4 CNNM2 KIF7 SLC5A2 CFI SLC7A7 AGT COL4A1 KLHL3 SLC7A9 CFHR5 SLC9A3R1 AGTR1 COL4A3 LAMB2 SMARCAL1 CD46 TCTN3 AGXT COL4A4 LMX1B SOX17 C3 THBD AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
AGT COL4A1 KLHL3 SLC7A9 CFHR5 SLC9A3R1 AGTR1 COL4A3 LAMB2 SMARCAL1 CD46 TCTN3 AGXT COL4A4 LMX1B SOX17 C3 THBD AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
AGTR1 COL4A3 LAMB2 SMARCAL1 CD46 TCTN3 AGXT COL4A4 LMX1B SOX17 C3 THBD AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
AGXT COL4A4 LMX1B SOX17 C3 THBD AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
AHI1 COL4A5 LYZ TCTN1 COL4A6 TMEM231 ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
ANLN COQ2 MKKS TCTN2 GANAB TSC1 APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
APOA1 CRB2 MKS1 TMEM138 DZIP1L VIPAS39 APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
APOL1 CSPP1 MYH9 TMEM216 VHL VPS33B APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
APRT CTNS MYO1E TMEM237 ADAMTS13 WDPCP AQP2 CUL3 NEK8 TMEM67 ANKS6 ZMYND10 ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
AQP2CUL3NEK8TMEM67ANKS6ZMYND10ARHGAP24DSTYKNOTCH2TNXBAPOEZNF423ARHGDIAEHHADHNPHP1TRIM32BBS9BICC1
ARHGAP24 DSTYK NOTCH2 TNXB APOE ZNF423 ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
ARHGDIA EHHADH NPHP1 TRIM32 BBS9 BICC1
ADIC EVA1 NDUD3 TDDCC C10A CIV1
ARL6 EYA1 NPHP3 TRPC6 C1QA SIX1
ATP6V0A4 FAH NPHP4 TTC21B C1QB SCARB2
ATP6V1B1 FGF20 NPHS1 TTC8 C1QC CFD
ATP7B FGF23 NPHS2 UMOD C5orf42 CFHR2
AVPR2 FN1 NR3C2 WDR19 CA2 APOA2
B2M FRAS1 OCRL WDR35 CEP83 IFNG
B9D1 FREM2 OFD1 WDR73 CD151 TSC2
B9D2 FXYD2 PAX2 WNK4 CEP164 CCND1
BBS1 GATA3 PDSS2 WNT4 CFB DCDC2
BBS10 GLA PKD1 WT1 CFHR1 MAPKBP1
BBS12 GLI3 PKD2 XPNPEP3 CFHR3 SEC61A1
BBS2 GLIS2 PKHD1 MUC1 ETFA JAG1
BBS4 GPC3 PLCE1 TBX18 ETFB
BBS5 GRHPR PTPRO CHRM3 ETFDH
BBS7 GRIP1 REN FREM1 FAN1
BMP4 HNF1B RET HPSE2 GSN
BSND HNF4A ROBO2 ITGA8 LCAT
CASR HOGA1 RPGRIP1L LRIG2 MASP1
CC2D2A HSD11B2 SALL1 COQ6 MEFV
CD2AP IFT122 SCNN1B DGKE PKHD1L1
CEP290 IFT140 SCNN1G EMP2 PRPS1
CEP41 IFT43 SIX5 ITGA3 RRM2B
CHD1L INF2 SLC12A1 NUP93 SARS2
CLCN5 INPP5E SLC12A3 NUP107 SDCCAG8
CLCNKA INVS SLC34A1 NUP205 SLC1A1

Supplementary Table 1: List of genes included in gene panel