

TARGETED GENE PANELS AND REFERENCE SEQUENCE (RefSeq) TRANSCRIPTS BY PHENOTYPE

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<u>Alagille Syndrome</u>	Transcript(s)
JAG1	NM_000214
NOTCH2	NM_024408

<u>Chondrodysplasia punctata</u>	Transcript(s)
AGPS	NM_003659
ARSE	NM_000047
EBP	NM_006579
GNPAT	NM_014236
PEX7	NM_000288

<u>Combined Pituitary Hormone Deficiency</u>	Transcript(s)
HESX1	NM_003865
POU1F1	NM_000306
PROP1	NM_006261
LHX3	NM_014564
LHX4	NM_033343

<u>Congenital Generalised Lipodystrophy</u>	Transcript(s)
AGPAT2	NM_006412
BSCL2	NM_032667
CAV1	NM_001753
PPARG	NM_015869
PTRF	NM_012232

<u>Congenital Hypothyroidism</u>	Transcript(s)
FOXE1	NM_004473
NKX2-1	NM_001079668
PAX8	NM_003466
TSHR	NM_000369
TPO	NM_000547
TG	NM_003235
DUOX2	NM_014080
THRA	NM_199334

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<u>Early-onset Diabetes and Autoimmunity</u>	Transcript(s)
AIRE	NM_000383
CD274	NM_014143
CTLA4	NM_005214
FOXP3	NM_014009
IL2RA	NM_000417
ITCH	NM_001257138
JAK1	NM_002227
LRBA	NM_001199282
SIRT1	NM_012238
STAT1	NM_007315
STAT3	NM_139276
STAT5B	NM_012448
TNFAIP3	NM_001270508

<u>Endocrine Neoplasia Syndromes</u>	Transcript(s)
AIP	NM_003977
CDKN1B	NM_004064
GPR101	NM_054021.1
MEN1	NM_130799
RET	NM_020975
SOS1	NM_005633

<u>Eye Movement disorders (<u>Congenital fibrosis of the extraocular muscles</u>, <u>Duane radial ray syndrome</u>, <u>Duane retraction syndrome</u>)</u>	Transcript(s)
CHN1	NM_001822
HOXA1	NM_005522
HOXB1	NM_002144
KIF21A	NM_001173464
PHOX2A	NM_005169
ROBO3	NM_022370
SALL4	NM_020436
TUBB3	NM_006086

<u>Familial Glucocorticoid Deficiency</u>	Transcript(s)
MC2R	NM_000529
MCM4	NM_005914
MRAP	NM_178817
NNT	NM_012343
STAR	NM_000349

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Familial Hyperparathyroidism	Transcript(s)
<i>CASR</i>	NM_000388
<i>CDC73</i>	NM_024529
<i>CDKN1A</i>	NM_078467
<i>CDKN1B</i>	NM_004064
<i>CDKN2B</i>	NM_004936
<i>CDKN2C</i>	NM_001262
<i>MEN1</i>	NM_130799
<i>RET</i>	NM_020975

Familial Hypocalciuric Hypercalcaemia	Transcript(s)
<i>AP2S1</i>	NM_004069
<i>GNA11</i>	NM_002067
<i>CASR</i>	NM_000388

Familial Hyperparathyroidism/hypercalcaemia	Transcript(s)
<i>AP2S1</i>	NM_004069
<i>CASR</i>	NM_000388
<i>CDC73</i>	NM_024529
<i>CDKN1A</i>	NM_078467
<i>CDKN1B</i>	NM_004064
<i>CDKN2B</i>	NM_004936
<i>CDKN2C</i>	NM_001262
<i>GNA11</i>	NM_002067
<i>MEN1</i>	NM_130799
<i>RET</i>	NM_020975

Familial Hypoparathyroidism	Transcript(s)
<i>CASR</i>	NM_000388
<i>GCM2</i>	NM_004752
<i>GNA11</i>	NM_002067
<i>PTH</i>	NM_000315

Familial Partial Lipodystrophy	Transcript(s)
<i>LMNA</i>	NM_170707
<i>PLIN1</i>	NM_002666
<i>PPARG</i>	NM_015869

Familial Porencephaly and HANAC syndrome	Transcript(s)
<i>COL4A1</i>	NM_001845
<i>COL4A2</i>	NM_001846
<i>JAM3</i>	NM_032801.4

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<u>Familial Tumoral Calcinosis</u>	Transcript(s)
FGF23	NM_020638
GALNT3	NM_004482
KL	NM_004795
SAMD9	NM_017654

<u>Feingold syndrome</u>	Transcript(s)
MYCN	NM_005378
MIR17HG	NR_027350

<u>Gastrointestinal atresia</u>	Transcript(s)
CCDC11 – HGNC approved name CFAP53	NM_145020
CHD7	NM_017780
FANCB	NM_001018113
FANCC	NM_000136
GLI3	NM_000168
MID1	NM_000381
MYCN	NM_005378
RFX6	NM_173560
SOX2	NM_003106
TTC7A	NM_020458
EFTUD2	NM_004247
FOXF1	NM_001451

<u>Generalised Arterial Calcification in Infancy</u>	Transcript(s)
ABCC6	NM_001171
ENPP1	NM_006208
NT5E	NM_002526.3

<u>Holoprosencephaly</u>	Transcript(s)
GLI2	NM_005270
PTCH1	NM_000264
SHH	NM_000193
SIX3	NM_005413
TGIF1	NM_173208
ZIC2	NM_007129

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Hyperinsulinism	Transcript(s)
<i>ABCC8</i>	NM_001287174
<i>AKT2</i>	NM_001626
<i>CACNA1D</i>	NM_000720
<i>GCK</i>	NM_000162
<i>GLUD1</i>	NM_005271
<i>GPC3</i>	NM_001164617
<i>HADH</i>	NM_005327
<i>HNFB1A</i>	NM_000545
<i>HNFB4A</i>	NM_175914
<i>INSR</i>	NM_000208
<i>KCNJ11</i>	NM_000525
<i>KDM6A</i>	NM_001291415
<i>KMT2D</i>	NM_003482
<i>PMM2</i>	NM_00303
<i>SLC16A1</i>	NM_003051
<i>TRMT10A</i>	NM_001134665

Hypophosphatemic Rickets	Transcript(s)
<i>DMP1</i>	NM_004407
<i>ENPP1</i>	NM_006208
<i>FGF23</i>	NM_020638
<i>PHEX</i>	NM_000444
<i>SLC34A3</i>	NM_080877

Isolated Growth Hormone Deficiency	Transcript(s)
<i>GH1</i>	NM_000515
<i>GHRHR</i>	NM_000823

Kabuki syndrome	Transcript(s)
<i>KDM6A</i>	NM_001291415
<i>KMT2D</i>	NM_003482

Kallmann syndrome	Transcript(s)
<i>KAL1</i>	NM_000216.2
<i>FGFR1</i>	NM_023110.2
<i>FGF8</i>	NM_033163.3
<i>PROKR2</i>	NM_144773.2
<i>PROK2</i>	NM_001126128.1

Mandibulofacial Dysostosis with Microcephaly	Transcript(s)
<i>EFTUD2</i>	NM_004247
<i>SF3B4</i>	NM_005850.4

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<u>Moebius syndrome</u>	Transcript(s)
PLXND1	NM_015103.2
REV3L	NM_002912.4

<u>Monogenic Diabetes of the Young (MODY)</u>	Transcript(s)
ABCC8	NM_001287174
APPL1	NM_012096
CEL	NM_001807
CISD2	NM_001008388
DCAF17	NM_025000
DNAJC3	NM_006260
DYRK1B	NM_004714
GATA4	NM_002052
GATA6	NM_005257
GCK	NM_000162
HNF1A	NM_000545
HNF1B	NM_000458
HNF4A	NM_175914
INS	NM_001185098
INSR	NM_000208
KCNJ11	NM_000525
LMNA	NM_170707
mtDNA_3243	NC_012920
NEUROD1	NM_002500
PAX6	NM_001604
PCBD1	NM_000281
PDX1	NM_000209
PIK3R1	NM_181523
PLIN1	NM_002666
POLD1	NM_002691
PPARG	NM_015869
PPP1R15B	NM_032833
RFX6	NM_173560
SLC29A3	NM_018344
TRMT10A	NM_001134665
WFS1	NM_006005
ZBTB20	NM_001164342
ZFP57	NM_001109809

<u>Multiple Exostosis</u>	Transcript(s)
EXT1	NM_000127.2
EXT2	NM_207122

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<u>Neonatal Diabetes</u>	<u>Transcript(s)</u>
<i>ABCC8</i>	NM_001287174
<i>AGPAT2</i>	NM_006412
<i>BSCL2</i>	NM_032667
<i>CISD2</i>	NM_001008388
<i>COQ2</i>	NM_015697
<i>COQ9</i>	NM_020312
<i>EIF2S3</i>	NM_001415
<i>EIF2AK3</i>	NM_004836
<i>FOXP3</i>	NM_014009
<i>GATA4</i>	NM_002052
<i>GATA6</i>	NM_005257
<i>GCK</i>	NM_000162
<i>GLIS3</i>	NM_001042413
<i>HNF1B</i>	NM_000458
<i>IER3IP1</i>	NM_016097
<i>IL2RA</i>	NM_000417
<i>INS</i>	NM_001185098
<i>INSR</i>	NM_000208
<i>KCNJ11</i>	NM_000525
<i>LPL</i>	NM_00237
<i>LRBA</i>	NM_006726
<i>MNX1</i>	NM_005515
<i>NEUROD1</i>	NM_002500
<i>NEUROG3</i>	NM_020999
<i>NKX2-2</i>	NM_002509
<i>PDX1</i>	NM_000209
<i>PTF1A</i>	NM_178161
<i>C10orf115</i>	NR_103721
<i>RFX6</i>	NM_173560
<i>SLC19A2</i>	NM_006996
<i>SLC2A2</i>	NM_000340
<i>STAT3</i>	NM_139276
<i>WFS1</i>	NM_006005
<i>ZFP57</i>	NM_001109809

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Phaeochromocytoma/Paraganglioma	Transcript(s)
<i>FH</i>	NM_000143
<i>MAX</i>	NM_002382
<i>RET</i>	NM_020975
<i>SDHA</i>	NM_004168
<i>SDHAF2</i>	NM_017841
<i>SDHB</i>	NM_003000
<i>SDHC</i>	NM_003001
<i>SDHD</i>	NM_003002
<i>TMEM127</i>	NM_017849
<i>VHL</i>	NM_000551

Pontocerebellar Hypoplasia	Transcript(s)
<i>AMPD2</i>	NM_004037
<i>CASK</i>	NM_003688
<i>CHMP1A</i>	NM_002768
<i>CLP1</i>	NM_006831
<i>EXOSC3</i>	NM_016042
<i>PCLO</i>	NM_033026
<i>RARS2</i>	NM_020320
<i>SEPSECS</i>	NM_016955
<i>TSEN2</i>	NM_025265
<i>TSEN34</i>	NM_024075
<i>TSEN54</i>	NM_207346
<i>VPS53</i>	NM_001128159
<i>VRK1</i>	NM_003384

Primary pigmented nodular adrenocortical disease	Transcript(s)
<i>PDE11A</i>	NM_016953
<i>PDE8B</i>	NM_003719
<i>PRKAR1A</i>	NM_002734

Pseudohypoaldosteronism	Transcript(s)
<i>CUL3</i>	NM_003590
<i>KLHL3</i>	NM_017415
<i>WNK1</i>	NM_018979
<i>WNK4</i>	NM_032387

Spondylocostal Dysostosis	Transcript(s)
<i>DLL3</i>	NM_016941
<i>HES7</i>	NM_032580
<i>LFNG</i>	NM_001040167
<i>MESP2</i>	NM_001039958
<i>TBX6</i>	NM_004608
<i>RIPPLY2</i>	NM_001009994

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<u>Visceral heterotaxy</u>	Transcript(s)
<i>CFC1</i>	NM_032545
<i>ZIC3</i>	NM_003413

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