

AMINO ACID DISORDERS		
PANEL	SUBPANEL	GENES
Homocystinuria	Hypermethioninemia	ADK, AHCY, CBS, GNMT, MAT1A, SLC25A13
	Hypomethioninemia	MTHFR, MTR, MTRR
Phenylketonuria	PAH Deficiency	PAH (sequencing + reflex MLPA as needed)
	Biopterin Deficiencies	DNAJC12, GCH1, PCBD1, PTS, QDPR, SPR (for PKU panel, PAH will be done with reflex to these genes)
Tyrosinemia	Elevated Succinylacetone	FAH, GSTZ1
	Elevated Tyrosine	HPD, TAT
Urea Cycle Diseases	High citrulline	ASS1, SLC25A13
	High ASA	ASL
	Low citrulline	CPS1, NAGS, OTC
	Other	ARG1, CA5A, GLUL, GLUD1, OAT, SLC7A7, SLC25A2, SLC25A15
Maple Syrup Urine Disease		BCKDHA, BCKDHB, DBT, DLD
ORGANIC ACID DISORDERS		
PANEL	SUBPANEL	GENES
Multiple carboxylase Deficiency	Biotinidase Deficiency	BTD
	Other	CA5A, HLCS
Propionic / Methylmalonic acidemias	PA	PCCA, PCCB
	MMA	ACSF3, ALDH6A1, MCEE, MLYCD, MMAA, MMAB, MMUT, SUCLA2, SUCLG1
	MMA + Homocysteinemia	ABCD4, AMN, CBLIF, CD320, CUBN, HCF1, LMBRD1, MMACHC, MMADHC, TCN1, TCN2
Isovaleric acidemia		IVD, ACADSB, FLAD1
Glutaric aciduria Type 1		GCDH
Isobutyryl-CoA dehydrogenase deficiency		ACAD8
Succinic semialdehyde dehydrogenase deficiency		ALDH5A1
b-ketothiolase deficiency		ACAT1
Guanidinoacetate Methyltransferase Deficiency		GAMT
FATTY ACID OXIDATION DISORDERS		
PANEL	GENES	
Carnitine Uptake Deficiency	SLC22A5	
MCAD Deficiency	ACADM (c.985A>G common mutation and del/dup +/- reflex sequencing)	
LCHAD/MTP Deficiency	HADHA, HADHB (HADHA c.1528G>C + reflex sequencing)	
VLCAD Deficiency	ACADVL	
MADD/Glutaric Aciduria Type2	ETFA, ETFB, ETFDH, FLAD1, SLC52A2, SLC52A3, SLC52A1	
CPT2 Deficiency	CPT2	
CACT Deficiency	SLC25A20	
CPT1 Deficiency	CPT1A (p.Pro479Leu common mutation + reflex sequencing)	
Other FAOD	ACAA2, ACAD9, ACADL, ACADS, ECHS1, HADH	
CONGENITAL ADRENAL HYPERPLASIA		
21-Hydroxylase Deficiency	CYP21A2 (includes MLPA and long-range PCR analyses for CNVs and common rearrangements)	
Other	ARMC5, CYP11B1, CYP11B2, CYP17A1, HSD3B2, POR, PRKARIA, STAR	
GALACTOSEMIA		
GALT Deficiency	GALT	
Other	GALK1, GALE, GALM, GLUT2 (SLC2A2)	
INBORN ERRORS OF IMMUNITY		
ADA Deficiency	ADA	
Chronic Granulomatous Disease	CYBA, CYBB, CYBC1, NCF1*, NCF2, NCF4, G6PD (*limited coverage due to high homology with duplicated regions in genome)	
Severe Combined/Primary Immune Deficiency Panel (251 genes) (Augmented exome slice with Sanger backfill of underlined genes + reflex MLPA of <u>DOCK8</u> as needed)		
<u>ACD</u> , <u>ACP5</u> , <u>ADA</u> , <u>ADA2</u> , <u>ADAM17</u> , <u>ADAR</u> , <u>AICDA</u> , <u>AIRE</u> , <u>AK2</u> , <u>AP3B1</u> , <u>ARHGEF1</u> , <u>ARPC1B</u> , <u>ATM</u> , <u>ATP6AP1</u> , <u>B2M</u> , <u>BACH2</u> , <u>BCL10</u> , <u>BCL11B</u> , <u>BLM</u> , <u>BLNK</u> , <u>BTK</u> , <u>CYBC1</u> , <u>C1QA</u> , <u>C1QB</u> , <u>C1QC</u> , <u>C1S</u> , <u>C2</u> , <u>C3</u> , <u>CARD11</u> , <u>CARD14</u> , <u>CARD9</u> , <u>CARMIL2</u> , <u>CASP10</u> , <u>CASP8</u> , <u>CD19</u> , <u>CD247</u> , <u>CD27</u> , <u>CD3D</u> , <u>CD3E</u> , <u>CD3G</u> , <u>CD40</u> , <u>CD40LG</u> , <u>CD70</u> , <u>CD79A</u> , <u>CD79B</u> , <u>CD81</u> , <u>CD8A</u> , <u>CDCA7</u> , <u>CFD</u> , <u>CFI</u> , <u>CFP</u> , <u>CHD7</u> , <u>CIITA</u> , <u>COPA</u> , <u>CR2</u> , <u>CTLA4</u> , <u>CTPS1</u> , <u>CTSC</u> , <u>CXCR4</u> , <u>CYBA</u> , <u>CYBB</u> , <u>DBR1</u> , <u>DCLRE1C</u> , <u>DKC1</u> , <u>DNASE2</u> , <u>DNMT3B</u> , <u>DOCK2</u> , <u>DOCK8</u> , <u>EBF1</u> , <u>EPG5</u> , <u>ERCC6L2</u> , <u>EXTL3</u> , <u>FADD</u> , <u>FAS</u> , <u>FASLG</u> , <u>FCHO1</u> , <u>FERMT3</u> , <u>FOXP1</u> , <u>FOXP3</u> , <u>G6PD</u> , <u>GATA2</u> , <u>GFI1</u> , <u>GINS1</u> , <u>HELLS</u> , <u>ICOS</u> , <u>IFIH1</u> , <u>IFNAR2</u> , <u>IFNGR1</u> , <u>IFNGR2</u> , <u>IGHM</u> , <u>IGLL1</u> , <u>IKKB</u> , <u>IKZF1</u> , <u>IL10</u> , <u>IL10RA</u> , <u>IL10RB</u> , <u>IL12B</u> , <u>IL12RB1</u> , <u>IL17RA</u> , <u>IL17RC</u> , <u>IL1RN</u> , <u>IL21</u> , <u>IL21R</u> , <u>IL23R</u> , <u>IL2RA</u> , <u>IL2RB</u> , <u>IL2RG</u> , <u>IL36RN</u> , <u>IL6ST</u> , <u>IL7R</u> , <u>IRAK4</u> , <u>IRF2BP2</u> , <u>IRF8</u> , <u>ISG15</u> , <u>ITGB2</u> , <u>ITK</u> , <u>JAK1</u> , <u>JAK3</u> , <u>KRAS</u> , <u>LAMTOR2</u> , <u>LAT</u> , <u>LCK</u> , <u>LIG1</u> , <u>LIG4</u> , <u>LRBA</u> , <u>LRRC8A</u> , <u>LYST</u> , <u>MAGT1</u> , <u>MALT1</u> , <u>MAP3K14</u> , <u>MEFV</u> , <u>MRTFA</u> , <u>MOGS</u> , <u>MSN</u> , <u>MTHFD1</u> , <u>MVK</u> , <u>MYD88</u> , <u>MYO5A</u> , <u>NBN</u> , <u>NCF2</u> , <u>NCF4</u> , <u>NFKB1</u> , <u>NFKB2</u> , <u>NFKBIA</u> , <u>NHEJ1</u> , <u>NHP2</u> , <u>NLR4</u> , <u>NLRP1</u> , <u>NLRP12</u> , <u>NLRP3</u> , <u>NOD2</u> , <u>NOP10</u> , <u>NRAS</u> , <u>NSMCE3</u> , <u>ORAI1</u> , <u>OTULIN</u> , <u>PARN</u> , <u>PEPD</u> , <u>PGM3</u> , <u>PIK3CD</u> , <u>PIK3R1</u> , <u>PLCG2</u> , <u>PMS2</u> , <u>PNP</u> , <u>POLD1</u> , <u>POLE</u> , <u>POLE2</u> , <u>PRF1</u> , <u>PRKCD</u> , <u>PRKDC</u> , <u>PSMB8</u> , <u>PSTPIP1</u> , <u>PTPRC</u> , <u>RAB27A</u> , <u>RAC2</u> , <u>RAG1</u> , <u>RAG2</u> , <u>RASGRP1</u> , <u>RBCK1</u> , <u>RELA</u> , <u>RELB</u> , <u>RFX5</u> , <u>RFXANK</u> , <u>RFXAP</u> , <u>RHOH</u> , <u>RIPK1</u> , <u>RMRP</u> , <u>RNA5EH2A</u> , <u>RNA5EH2B</u> , <u>RNA5EH2C</u> , <u>RNF168</u> , <u>RNF31</u> , <u>RORC</u> , <u>RTEL1</u> , <u>SAMHD1</u> , <u>SBDS</u> , <u>SEMA3E</u> , <u>SERPING1</u> , <u>SH2D1A</u> , <u>SLC29A3</u> , <u>SLC35C1</u> , <u>SLC39A7</u> , <u>SLC7A7</u> , <u>SMARCAL1</u> , <u>SP110</u> , <u>SPINK5</u> , <u>SPPL2A</u> , <u>STAT1</u> , <u>STAT2</u> , <u>STAT3</u> , <u>STIM1</u> , <u>STING1</u> , <u>STK4</u> , <u>STX11</u> , <u>STXBP2</u> , <u>TAP1</u> , <u>TAP2</u> , <u>TAPBP</u> , <u>TCF3</u> , <u>TCN2</u> , <u>TERC</u> , <u>TERT</u> , <u>TFRC</u> , <u>TGFB1</u> , <u>TINF2</u> , <u>TMC6</u> ,		

TM8, TNFAIP3, TNFRSF13B, TNFRSF1A, TNFRSF4, TNFRSF9, TRAC, TRAF3IP2, TREX1, TRNT1, TRTC37, TRTC7A, TRYK2, UNC13D, UNC93B1, UNG, USP18, WAS, WDR1, WIPF1, WRAP53, XIAP, ZAP70, ZBT24, ZNF341

MITOCHONDRIAL DISEASES (AUGMENTED EXOME SLICES WITH SANGER BACKFILL OF UNDERLINED GENES LISTED)

Full Mitochondrial Disease Nuclear Gene Panel (425 genes)

AARS2, ABAT, ABC7, ACACB, ACAD8, ACAD9, ACADL, ACADM, ACADS, ACADSB, ACADVL, ACAT1, ACLY, ACO2, ACSL5, ACSM3, ADAR, ADSL, AFG3L2, AGK, AGXT2, AIFM1, AK2, AKAP10, AKR7A2, ALDH18A1, ALDH1B1, ALDH5A1, ALDH6A1, ALDH7A1, ALG3, AMPD1, AMT, ANTXR1, AS3MT, ATIC, ATP1A3, ATP10D, ATP5F1A, ATP5F1B, ATP5F1C, ATP5F1D, ATP5F1E, ATP5MC1, ATP5MC2, ATP5MC3, ATP5ME, ATP5ME, ATP5MG, ATP5MGL, ATP5PO, ATP5PB, ATP5PD, ATP5PE, ATPAF1, ATPAF2, AUH, BCKDHA, BCKDHB, BCS1L, BOLA3, BTD, C1QBP, C19orf12, CA5A, CARS2, CCDC88A, CEP89, CHCHD10, CHDH, CHKB, CISD2, CLN3, CLPB, CLPP, CLYBL, COA1, COA3, COA4, COA5, COA6, COA7, COA8, COASY, COMT, COQ2, COQ4, COQ5, COQ6, COQ7, COQ8A, COQ8B, COQ9, COX10, COX11, COX14, COX15, COX16, COX17, COX18, COX19, COX20, COX4I1, COX4I2, COX5A, COX5B, COX6A1, COX6A2, COX6B1, COX6B2, COX6C, COX7A1, COX7A2, COX7B, COX7C, COX8A, CPT1A, CPT1B, CPT2, CYC1, CYCS, CYP11A1, CYP11B1, CYP11B2, D2HGDH, DARS2, DBT, DDAH1, DGUOK, DLAT, DLD, DNA2, DNAJC19, DNAJC30, DNM1L, DNMT1, EARS2, ECHS1, ELAC2, ERAL1, ETFA, ETFB, ETFDH, ETHE1, FA2H, FARS2, FASTKD2, FBXL4, FDX2, FDXR, FH, FLAD1, FOXRED1, FXN, GARS1, GATB, GATC, GATM, GCDH, GCSH, GDAP1, GFER, GFM1, GFM2, GLDC, GLRX5, GLS, GTPBP3, GYG2, HADHA, HADHB, HARS2, HCCS, HIBCH, HLCS, HMGCL, HMGCS2, HSD17B10, HSPA9, HSPD1, IARS2, IBA57, IDH2, IDH3A, IDH3B, ISCA1, ISCA2, ISCU, IVD, KARS1, KIF5A, KIF21A, KLC2, KYNU, L2HGDH, LARS1, LARS2, LIAS, LIPT1, LIPT2, LMBRD1, LONP1, LPIN1, LRPPRC, LYRM4, LYRM7, MARS2, MCEE, MDH2, MECR, MFF, MFN2, MGME1, MICOS13, MICU1, MIPEP, MLYCD, MMAA, MMAB, MMACHC, MPV17, MRM2, MRPL12, MRPL3, MRPL44, MRPS14, MRPS16, MRPS22, MRPS23, MRPS34, MRPS7, MTFMT, MTO1, MTPAP, MTRFR, MTRR, MMUT, NADK2, NARS2, NAXE, NDUFA1, NDUFA10, NDUFA11, NDUFA12, NDUFA13, NDUFA2, NDUFA3, NDUFA5, NDUFA7, NDUFA8, NDUFA9, NDUFAB1, NDUFAF1, NDUFAF2, NDUFAF3, NDUFAF4, NDUFAF5, NDUFAF6, NDUFAF7, NDUFAF8 (C17ORF89), NDUFB1, NDUFB6, NDUFB10, NDUFB11, NDUFB2, NDUFB3, NDUFB4, NDUFB5, NDUFB7, NDUFB8, NDUFB9, NDUFC1, NDUFS1, NDUFS2, NDUFS3, NDUFS4, NDUFS5, NDUFS6, NDUFS7, NDUFS8, NDUFV1, NDUFV2, NDUFV3, NFU1, NFU1, NR2F1, NSUN3, NUBPL, OPA1, OPA3, OXA1L, OXCT1, PANK2, PARS2, PC, PCCA, PCCB, PCK2, PDHA1, PDHB, PDHX, PDK3, PDP1, PDSS1, PDSS2, PET100, PET117, PHOX2A, PITRM1, PLA2G6, PLP1, PMPCA, PMPCB, PNPLA8, PNPT1, POLG, POLG2, PPA2, PRPS1, PTCD3, PTS, PUS1, QARS1, QDPR, QRS1, RARS1, RARS2, RMND1, RNASEH1, ROBO3, RRM2B, RTN4IP1, SACS, SARS2, SCO1, SCO2, SCP2, SDHA, SDHAF1, SDHAF2, SDHAF3, SDHAF4, SDHB, SDHC, SDHD, SERAC1, SFXN4, SLC19A2, SLC19A3, SLC22A5, SLC25A1, SLC25A12, SLC25A13, SLC25A19, SLC25A20, SLC25A21, SLC25A26, SLC25A3, SLC25A32, SLC25A38, SLC25A4, SLC25A42, SLC25A46, SLC52A2, SLC52A3, SNX10, SPATA5, SPG7, SPR, STAR, SUCLA2, SUCLG1, SUOX, SURF1, TACO1, TAFAZZIN, TARS2, TCIRG1, TCN2, TIMM22, TIMM44, TIMM50, TIMM8A, TIMMDC1, TK2, TMEM126A, TMEM126B, TMEM65, TMEM70, TOMM20, TOP3A, TPK1, TRIT1, TRMT10C, TRMT5, TRMU, TRNT1, TSFM, TTC19, TUBB3, TUBB4A, TUFM, TUSC3, TWINK, TXN2, TYMP, UCHL1, UNG, UQCC1, UQCC2, UQCC3, UQCR10, UQCR11, UQCRB, UQCRC1, UQCRC2, UQCRF51, UQCRH, UQCRO, VARS2, WARS2, WDR73, WFS1, YARS2, YME1L1

Mitochondrial Encephalopathy / Leigh Disease Panel (117 genes)

AARS2, ACAD9, ACO2, AFG3L2, AIFM1, ATP5F1E, ATPAF2, BCS1L, BOLA3, COQ2, COQ8A, COQ9, COX7A1, COX10, COX14, COX15, COX20, COX4I1, COX4I2, COX6B1, DARS2, DGUOK, DLAT, DLD, DNM1L, EARS2, ETFDH, ETHE1, FARS2, FASTKD2, FH, FOXRED1, GFER, GFM1, GFM2, HLCS, HSPD1, LARS2, LIAS, LMBRD1, LRPPRC, MARS2, MFN2, MPV17, MRPS16, MTFMT, MTPAP, NDUFA1, NDUFA10, NDUFA11, NDUFA12, NDUFA13, NDUFA2, NDUFA7, NDUFA8, NDUFA9, NDUFAF1, NDUFAF2, NDUFAF3, NDUFAF4, NDUFAF5, NDUFAF6, NDUFAF7, NDUFB6, NDUFS1, NDUFS2, NDUFS3, NDUFS4, NDUFS5, NDUFS6, NDUFS7, NDUFS8, NDUFV1, NDUFV2, NDUFV3, NFU1, NUBPL, PC, PDHA1, PDHB, PDHX, PDP1, PDSS1, PDSS2, PNPT1, POLG, RARS2, RMND1, RRM2B, SCO1, SCO2, SDHA, SDHAF1, SDHAF2, SDHB, SDHD, SERAC1, SLC19A3, SUCLA2, SUCLG1, SURF1, TACO1, TIMM44, TK2, TMEM70, TOMM20, TPK1, TRMU, TSFM, TTC19, TUFM, TUSC3, TWINK, TYMP, UQCRB, UQCRO, YARS2

mtDNA Depletion And Deletion Panel (19 genes)

AGK, DGUOK, DNA2, FBXL4, GFER, MFN2, MGME1, MPV17, OPA1, OPA3, POLG, POLG2, RRM2B, SLC25A4, SUCLA2, SUCLG1, TK2, TYMP, TWINK

Progressive External Ophthalmoplegia (PEO) And Optic Atrophy (77 genes)

ACO2, AFG3L2, ALG3, ANTXR1, ATP1A3, AUH, C19orf12, C1QBP, CCDC88A, CISD2, CLN3, DGUOK, DNA2, DNAJC19, DNAJC30, DNM1L, DNMT1, FA2H, FDX2, FDXR, FH, GYG2, IBA57, ISCA2, KIF21A, KLC2, MECR, MFF, MFN2, MGME1, MICOS13, MTFMT, MTO1, MTPAP, MTRFR, NARS2, NDUFAF3, NDUFS1, NR2F1, OPA1, OPA3, PANK2, PDHX, PDSS1, PHOX2A, PLA2G6, PLP1, POLG, POLG2, PRPS1, RNASEH1, ROBO3, RRM2B, RTN4IP1, SLC19A2, SLC19A3, SLC25A4, SLC25A46, SLC52A2, SLC52A3, SNX10, SPG7, SUCLA2, TACO1, TCIRG1, TIMM8A, TK2, TMEM126A, TSFM, TUBB3, TUBB4A, TWINK, TYMP, UCHL1, WDR73, WFS1, YME1L1

Pyruvate Dehydrogenase Complex Deficiency (16 genes)

BOLA3, DLAT, DLD, LIAS, LIPT1, LIPT2, NFU1, PC, PDHA1, PDHB, PDHX, PDK3, PDP1, SLC19A2, SLC19A3, TPK1