

METABOLIC DISORDERS GENE PANEL DG 2.18 (683 genes)

Releasedate: 20-04-2020

Gene	Agilent V5 covered >10x	Agilent V5 covered > 20x	TWIST covered >10x	TWIST covered >20x	Associated Phenotype description and OMIM disease ID
AASS	100%	99,70%	100%	100%	Hyperlysinemia, 238700
ABAT	100%	99,40%	100%	100%	GABA-transaminase deficiency, 613163
ABCC8	100%	99,80%	100%	100%	Diabetes mellitus, permanent neonatal, 606176 Diabetes mellitus, noninsulin-dependent, 125853 Diabetes mellitus, transient neonatal 2, 610374 Hyperinsulinemic hypoglycemia, familial, 1, 256450 Hypoglycemia of infancy, leucine-sensitive, 240800
ABCD1	75,80%	71,60%	100%	100%	Adrenomyeloneuropathy, adult, 300100 Adrenoleukodystrophy, 300100
ABCD2	100%	99,80%	100%	100%	No OMIM disease ID
ABCD3	99,80%	97,70%	100%	100%	?Bile acid synthesis defect, congenital, 5, 616278
ABCD4	99,90%	98,60%	100%	100%	Methylmalonic aciduria and homocystinuria, cblJ type, 614857
ABCG5	100%	100%	100%	100%	Sitosterolemia 2, 618666
ABCG8	99,10%	97,30%	100%	100%	Sitosterolemia 1, 210250
ABHD12	98,70%	92,30%	100%	99,30%	Polyneuropathy, hearing loss, ataxia, retinitis pigmentosa, and cataract, 612674
ABHD5	100%	100%	100%	100%	Chanarin-Dorfman syndrome, 275630
ACACA	98,40%	98,10%	100%	100%	No OMIM disease ID
ACAD8	100%	100%	100%	100%	Isobutyryl-CoA dehydrogenase deficiency, 611283
ACAD9	100%	99,90%	100%	100%	Mitochondrial complex I deficiency, nuclear type 20, 611126
ACADM	99,80%	99,00%	100%	100%	Acyl-CoA dehydrogenase, medium chain, deficiency of, 201450
ACADS	99,90%	98,20%	100%	100%	Acyl-CoA dehydrogenase, short-chain, deficiency of, 201470
ACADSB	100%	99,20%	100%	100%	2-methylbutyrylglucosuria, 610006
ACADVL	99,40%	97,30%	100%	100%	VLCAD deficiency, 201475
ACAT1	99,90%	97,50%	100%	100%	Alpha-methylacetoacetic aciduria, 203750
ACAT2	100%	100%	100%	100%	No OMIM disease ID
ACBD5	100%	99,20%	100%	100%	Retinal dystrophy with leukodystrophy, 618863
ACO2	96,30%	90,30%	100%	100%	?Optic atrophy 9, 616289 Infantile cerebellar-retinal degeneration, 614559
ACOX1	100%	99,90%	100%	100%	Peroxisomal acyl-CoA oxidase deficiency, 264470

ACOX2	100%	99,20%	100%	100%	Bile acid synthesis defect, congenital, 6, 617308
ACSF3	100%	99,90%	100%	100%	Combined malonic and methylmalonic aciduria, 614265
ACSL4	98,70%	94,60%	100%	100%	Mental retardation, X-linked 63, 300387
ACY1	100%	98,80%	100%	100%	Aminoacylase 1 deficiency, 609924
ADA	100%	99,70%	100%	100%	Adenosine deaminase deficiency, partial, 102700 Severe combined immunodeficiency due to ADA deficiency, 102700
ADCK5	100%	99,90%	100%	100%	No OMIM disease ID
ADCY5	95,10%	91,20%	99,20%	98,00%	Dyskinesia, familial, with facial myokymia, 606703
ADK	99,50%	95,80%	100%	100%	Hypermethioninemia due to adenosine kinase deficiency, 614300
ADSL	99,20%	98,70%	100%	100%	Adenylosuccinase deficiency, 103050
AGA	100%	100%	100%	100%	Aspartylglucosaminuria, 208400
AGK	99,90%	97,60%	100%	100%	Sengers syndrome, 212350 Cataract 38, autosomal recessive, 614691
AGL	100%	99,40%	100%	100%	Glycogen storage disease IIIb, 232400 Glycogen storage disease IIIa, 232400
AGPAT2	99,60%	96,10%	100%	100%	Lipodystrophy, congenital generalized, type 1, 608594
AGPS	99,30%	95,40%	100%	99,90%	Rhizomelic chondrodysplasia punctata, type 3, 600121
AGXT	100%	100%	100%	100%	Hyperoxaluria, primary, type 1, 259900
AHCY	100%	99,20%	100%	100%	Hypermethioninemia with deficiency of S-adenosylhomocysteine hydrolase, 613752
AK1	100%	100%	100%	100%	Hemolytic anemia due to adenylate kinase deficiency, 612631
AK2	98,80%	94,50%	100%	100%	Reticular dysgenesis, 267500
AKR1D1	100%	99,40%	100%	100%	Bile acid synthesis defect, congenital, 2, 235555
ALAD	99,30%	94,10%	100%	100%	Porphyria, acute hepatic, 612740
ALAS2	98,90%	94,90%	100%	100%	Protoporphyrinemia, erythropoietic, X-linked, 300752 Anemia, sideroblastic, 1, 300751
ALDH18A1	100%	99,90%	100%	100%	Cutis laxa, autosomal recessive, type IIIA, 219150 Cutis laxa, autosomal dominant 3, 616603 Spastic paraplegia 9B, autosomal recessive, 616586 Spastic paraplegia 9A, autosomal dominant, 601162
ALDH1A3	97,20%	94,50%	100%	99,90%	Microphthalmia, isolated 8, 615113
ALDH2	100%	100%	100%	100%	Alcohol sensitivity, acute, 610251
ALDH3A2	95,30%	94,60%	100%	100%	Sjogren-Larsson syndrome, 270200
ALDH4A1	100%	99,40%	100%	100%	Hyperprolinemia, type II, 239510
ALDH5A1	91,00%	81,50%	100%	100%	Succinic semialdehyde dehydrogenase deficiency, 271980
ALDH6A1	100%	99,90%	100%	100%	Methylmalonate semialdehyde dehydrogenase deficiency, 614105
ALDH7A1	94,40%	88,80%	100%	100%	Epilepsy, pyridoxine-dependent, 266100
ALDOA	75,50%	74,00%	100%	100%	Glycogen storage disease XII, 611881

ALDOB	100%	99,10%	100%	100%	Fructose intolerance, hereditary, 229600
ALG1	53,00%	45,80%	100%	100%	Congenital disorder of glycosylation, type Ik, 608540
ALG10	100%	100%	100%	100%	No OMIM disease ID
ALG11	96,80%	96,80%	96,80%	96,80%	Congenital disorder of glycosylation, type Ip, 613661
ALG12	100%	100%	100%	100%	Congenital disorder of glycosylation, type Ig, 607143
ALG13	98,40%	92,60%	100%	99,60%	Epileptic encephalopathy, early infantile, 36, 300884 ?Congenital disorder of glycosylation, type Is, 300884
ALG14	100%	99,90%	100%	100%	?Myasthenic syndrome, congenital, 15, without tubular aggregates, 616227
ALG2	100%	100%	100%	100%	Myasthenic syndrome, congenital, 14, with tubular aggregates, 616228 ?Congenital disorder of glycosylation, type Ii, 607906
ALG3	100%	99,70%	100%	100%	Congenital disorder of glycosylation, type Id, 601110
ALG6	98,60%	94,80%	100%	100%	Congenital disorder of glycosylation, type Ic, 603147
ALG8	97,20%	95,60%	96,60%	96,60%	Congenital disorder of glycosylation, type Ih, 608104 Polycystic liver disease 3 with or without kidney cysts, 617874
ALG9	100%	99,70%	100%	100%	Gillessen-Kaesbach-Nishimura syndrome, 263210 Congenital disorder of glycosylation, type II, 608776
ALOX12B	100%	100%	100%	100%	Ichthyosis, congenital, autosomal recessive 2, 242100
ALPL	100%	100%	100%	100%	Hypophosphatasia, adult, 146300 Odontohypophosphatasia, 146300 Hypophosphatasia, childhood, 241510 Hypophosphatasia, infantile, 241500
AMACR	100%	100%	100%	100%	Bile acid synthesis defect, congenital, 4, 214950 Alpha-methylacyl-CoA racemase deficiency, 614307
AMN	89,70%	80,00%	100%	100%	Megaloblastic anemia-1, Norwegian type, 261100
AMPD1	99,90%	98,60%	100%	100%	Myopathy due to myoadenylate deaminase deficiency, 615511
AMPD3	99,90%	98,50%	100%	100%	No OMIM disease ID
AMT	100%	100%	100%	100%	Glycine encephalopathy, 605899
AP1S1	99,90%	99,50%	100%	100%	MEDNIK syndrome, 609313
AP3B2	99,40%	95,10%	100%	100%	Epileptic encephalopathy, early infantile, 48, 617276
APOC2	100%	100%	100%	100%	Hyperlipoproteinemia, type Ib, 207750
APRT	100%	99,50%	100%	100%	Adenine phosphoribosyltransferase deficiency, 614723
ARG1	100%	100%	100%	100%	Argininemia, 207800
ARSA	100%	99,80%	100%	100%	Metachromatic leukodystrophy, 250100
ARSB	96,90%	88,30%	100%	100%	Mucopolysaccharidosis type VI (Maroteaux-Lamy), 253200
ASAH1	99,70%	98,60%	100%	100%	Farber lipogranulomatosis, 228000 Spinal muscular atrophy with progressive myoclonic epilepsy, 159950
ASL	100%	99,60%	100%	100%	Argininosuccinic aciduria, 207900

ASNS	99,40%	95,20%	100%	100%	Asparagine synthetase deficiency, 615574
ASPA	99,90%	98,30%	100%	100%	Canavan disease, 271900
ASS1	95,40%	87,90%	100%	100%	Citrullinemia, 215700
ATIC	99,90%	99,30%	100%	100%	AICA-ribosiduria due to ATIC deficiency, 608688
ATP1A1	100%	100%	100%	100%	Charcot-Marie-Tooth disease, axonal, type 2DD, 618036 Hypomagnesemia, seizures, and mental retardation 2, 618314
ATP6AP1	98,20%	92,10%	100%	100%	Immunodeficiency 47, 300972
ATP6VOA2	100%	99,50%	100%	100%	Wrinkly skin syndrome, 278250 Cutis laxa, autosomal recessive, type IIA, 219200
ATP6V1A	99,90%	98,70%	100%	100%	Epileptic encephalopathy, infantile or early childhood, 3, 618012 Cutis laxa, autosomal recessive, type IID, 617403
ATP6V1E1	93,10%	88,30%	100%	100%	Cutis laxa, autosomal recessive, type IIC, 617402
ATP7A	99,70%	97,50%	100%	100%	Occipital horn syndrome, 304150 Menkes disease, 309400 Spinal muscular atrophy, distal, X-linked 3, 300489
ATP7B	99,90%	99,20%	100%	100%	Wilson disease, 277900
ATP8B1	96,50%	94,00%	100%	100%	Cholestasis, intrahepatic, of pregnancy, 1, 147480 Cholestasis, progressive familial intrahepatic 1, 211600 Cholestasis, benign recurrent intrahepatic, 243300
AUH	100%	99,80%	100%	100%	3-methylglutaconic aciduria, type I, 250950
B3GALNT1	100%	99,80%	100%	100%	No OMIM disease ID
B3GALNT2	93,80%	89,40%	92,50%	92,50%	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies, type A, 11, 615181
B3GALT6	75,70%	69,70%	89,80%	81,60%	Al-Gazali syndrome, 609465 Ehlers-Danlos syndrome, spondylodysplastic type, 2, 615349 Spondyloepimetaphyseal dysplasia with joint laxity, type 1, with or without fractures, 271640
B3GAT3	99,90%	98,20%	94,80%	94,80%	Multiple joint dislocations, short stature, craniofacial dysmorphism, with or without congenital heart defects, 245600
B3GLCT	99,60%	96,30%	99,90%	99,20%	Peters-plus syndrome, 261540
B4GALT1	100%	99,80%	100%	100%	Congenital disorder of glycosylation, type IId, 607091
B4GALT7	99,80%	97,40%	99,90%	98,60%	Ehlers-Danlos syndrome, spondylodysplastic type, 1, 130070
B4GAT1	100%	100%	100%	100%	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 13, 615287
BAAT	99,80%	98,40%	100%	100%	Hypercholanemia, familial, 607748
BCAT1	100%	100%	100%	100%	No OMIM disease ID
BCAT2	100%	100%	100%	100%	?Hypervalinemia or hyperleucine-isoleucinemia, 618850
BCKDHA	99,90%	99,20%	100%	100%	Maple syrup urine disease, type Ia, 248600
BCKDHB	99,50%	94,40%	100%	100%	Maple syrup urine disease, type Ib, 248600
BCKDK	100%	100%	100%	100%	Branched-chain ketoacid dehydrogenase kinase deficiency, 614923

<i>BCO1</i>	100%	100%	100%	100%	?Hypercarotenemia and vitamin A deficiency, autosomal dominant, 115300
<i>BLVRA</i>	100%	99,40%	100%	100%	Hyperbiliverdinemia, 614156
<i>BMP2</i>	100%	100%	100%	100%	Brachydactyly, type A2, 112600 Short stature, facial dysmorphism, and skeletal anomalies with or without cardiac anomalies, 617877
<i>BPGM</i>	100%	100%	100%	100%	Erythrocytosis, familial, 8, 222800
<i>BTB</i>	100%	99,90%	100%	100%	Biotinidase deficiency, 253260
<i>C1GALT1C1</i>	100%	99,50%	100%	100%	Tn polyagglutination syndrome, somatic, 300622
<i>CA5A</i>	99,70%	97,10%	100%	100%	Hyperammonemia due to carbonic anhydrase VA deficiency, 615751
<i>CAD</i>	100%	99,20%	100%	100%	Epileptic encephalopathy, early infantile, 50, 616457
<i>CANT1</i>	100%	99,90%	100%	100%	Desbuquois dysplasia 1, 251450 Epiphyseal dysplasia, multiple, 7, 617719
<i>CAT</i>	100%	100%	100%	100%	Acatlasemia, 614097
<i>CBS</i>	99,80%	98,30%	100%	100%	Homocystinuria, B6-responsive and nonresponsive types, 236200 Thrombosis, hyperhomocysteinemic, 236200
<i>CCDC115</i>	95,30%	90,00%	100%	100%	Congenital disorder of glycosylation, type Ilo, 616828
<i>CEL</i>	89,70%	88,00%	100%	99,80%	Maturity-onset diabetes of the young, type VIII, 609812
<i>CERKL</i>	99,50%	96,90%	100%	100%	Retinitis pigmentosa 26, 608380
<i>CERS3</i>	99,90%	98,90%	100%	100%	Ichthyosis, congenital, autosomal recessive 9, 615023
<i>CFTR</i>	99,60%	97,90%	100%	100%	Cystic fibrosis, 219700 Congenital bilateral absence of vas deferens, 277180 Sweat chloride elevation without CF, 0
<i>CHIT1</i>	99,70%	98,10%	100%	100%	No OMIM disease ID
<i>CHKB</i>	100%	99,70%	100%	100%	Muscular dystrophy, congenital, megaconial type, 602541
<i>CHST14</i>	99,90%	98,90%	100%	100%	Ehlers-Danlos syndrome, musculocontractural type 1, 601776
<i>CHST3</i>	100%	99,40%	100%	100%	Spondyloepiphyseal dysplasia with congenital joint dislocations, 143095
<i>CHST6</i>	100%	100%	100%	100%	Macular corneal dystrophy, 217800
<i>CHSY1</i>	97,20%	95,70%	99,70%	98,00%	Temtamy preaxial brachydactyly syndrome, 605282
<i>CLCN7</i>	99,70%	98,40%	100%	100%	Osteopetrosis, autosomal recessive 4, 611490 Osteopetrosis, autosomal dominant 2, 166600 Hypopigmentation, organomegaly, and delayed myelination and development, 618541
<i>CLN3</i>	92,50%	91,80%	92,50%	92,50%	Ceroid lipofuscinosis, neuronal, 3, 204200
<i>CLN5</i>	99,40%	95,50%	100%	100%	Ceroid lipofuscinosis, neuronal, 5, 256731
<i>CLN6</i>	99,90%	97,10%	100%	100%	Ceroid lipofuscinosis, neuronal, Kufs type, adult onset, 204300 Ceroid lipofuscinosis, neuronal, 6, 601780
<i>CLN8</i>	83,50%	83,50%	100%	100%	Ceroid lipofuscinosis, neuronal, 8, 600143 Ceroid lipofuscinosis, neuronal, 8, Northern epilepsy variant, 610003
<i>CLPB</i>	100%	99,90%	100%	100%	3-methylglutaconic aciduria, type VII, with cataracts, neurologic involvement and neutropenia, 616271

<i>CMAS</i>	99,90%	97,80%	100%	100%	No OMIM disease ID
<i>COG1</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type IIg, 611209
<i>COG2</i>	99,90%	98,50%	100%	100%	?Congenital disorder of glycosylation, type IIq, 617395
<i>COG4</i>	100%	99,90%	100%	100%	Saul-Wilson syndrome, 618150 Congenital disorder of glycosylation, type IIj, 613489
<i>COG5</i>	99,70%	97,60%	100%	100%	Congenital disorder of glycosylation, type IIi, 613612
<i>COG6</i>	99,10%	93,90%	100%	100%	Shaheen syndrome, 615328 Congenital disorder of glycosylation, type III, 614576
<i>COG7</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type IIe, 608779
<i>COG8</i>	100%	99,60%	100%	100%	Congenital disorder of glycosylation, type IIh, 611182
<i>COMT</i>	100%	99,90%	100%	100%	No OMIM disease ID
<i>COQ2</i>	98,00%	95,30%	97,20%	97,20%	Coenzyme Q10 deficiency, primary, 1, 607426
<i>COQ4</i>	90,90%	89,30%	100%	100%	Coenzyme Q10 deficiency, primary, 7, 616276
<i>COQ5</i>	100%	100%	100%	100%	No OMIM disease ID
<i>COQ6</i>	99,90%	98,40%	100%	100%	Coenzyme Q10 deficiency, primary, 6, 614650
<i>COQ7</i>	100%	99,80%	100%	100%	?Coenzyme Q10 deficiency, primary, 8, 616733
<i>COQ8A</i>	100%	99,50%	100%	100%	Coenzyme Q10 deficiency, primary, 4, 612016
<i>COQ8B</i>	100%	99,30%	100%	100%	Nephrotic syndrome, type 9, 615573
<i>COQ9</i>	100%	97,90%	100%	100%	Coenzyme Q10 deficiency, primary, 5, 614654
<i>CP</i>	94,80%	88,90%	100%	100%	Hemosiderosis, systemic, due to aceruloplasminemia, 604290 Cerebellar ataxia, 604290
<i>CPOX</i>	99,90%	95,40%	100%	100%	Harderoporphyria, 121300 Coproporphyria, 121300
<i>CPS1</i>	100%	99,90%	100%	100%	Carbamoylphosphate synthetase I deficiency, 237300
<i>CPT1A</i>	100%	98,90%	100%	100%	CPT deficiency, hepatic, type IA, 255120
<i>CPT2</i>	98,20%	97,80%	100%	100%	CPT II deficiency, myopathic, stress-induced, 255110 CPT II deficiency, infantile, 600649 CPT II deficiency, lethal neonatal, 608836
<i>CRAT</i>	100%	99,80%	100%	100%	?Neurodegeneration with brain iron accumulation 8, 617917
<i>CTH</i>	100%	100%	100%	100%	Cystathioninuria, 219500 Homocysteine, total plasma, elevated, 0
<i>CTNS</i>	100%	99,80%	100%	100%	Cystinosis, nephropathic, 219800 Cystinosis, ocular nonnephropathic, 219750 Cystinosis, late-onset juvenile or adolescent nephropathic, 219900 Cystinosis, atypical nephropathic, 219800
<i>CTSA</i>	100%	100%	100%	100%	Galactosialidosis, 256540

CTSC	100%	100%	100%	100%	Periodontitis 1, juvenile, 170650 Papillon-Lefevre syndrome, 245000 Haim-Munk syndrome, 245010
CTSD	98,40%	95,00%	100%	100%	Ceroid lipofuscinosis, neuronal, 10, 610127
CTSK	100%	99,90%	100%	100%	Pycnodysostosis, 265800
CUBN	99,70%	98,30%	100%	100%	Megaloblastic anemia-1, Finnish type, 261100
CYB561	92,80%	92,60%	100%	99,90%	Orthostatic hypotension 2, 618182
CYB5R3	98,40%	98,00%	99,80%	98,90%	Methemoglobinemia, type I, 250800 Methemoglobinemia, type II, 250800
CYP11A1	99,30%	96,10%	100%	100%	Adrenal insufficiency, congenital, with 46XY sex reversal, partial or complete, 613743
CYP11B1	100%	100%	100%	100%	Adrenal hyperplasia, congenital, due to 11-beta-hydroxylase deficiency, 202010 Aldosteronism, glucocorticoid-remediable, 103900
CYP11B2	100%	100%	100%	100%	Hypoaldosteronism, congenital, due to CMO II deficiency, 610600 Hypoaldosteronism, congenital, due to CMO I deficiency, 203400 Aldosterone to renin ratio raised, 0
CYP17A1	100%	99,50%	100%	100%	17-alpha-hydroxylase/17,20-lyase deficiency, 202110 17,20-lyase deficiency, isolated, 202110
CYP19A1	98,80%	96,80%	100%	100%	Aromatase deficiency, 613546 Aromatase excess syndrome, 139300
CYP1B1	100%	100%	100%	100%	Glaucoma 3A, primary open angle, congenital, juvenile, or adult onset, 231300 Anterior segment dysgenesis 6, multiple subtypes, 617315
CYP21A2	97,80%	88,40%	100%	100%	Adrenal hyperplasia, congenital, due to 21-hydroxylase deficiency, 201910 Hyperandrogenism, nonclassic type, due to 21-hydroxylase deficiency, 201910
CYP27A1	98,90%	96,70%	100%	100%	Cerebrotendinous xanthomatosis, 213700
CYP27B1	99,90%	99,30%	100%	100%	Vitamin D-dependent rickets, type I, 264700
CYP2R1	99,40%	95,60%	100%	100%	Rickets due to defect in vitamin D 25-hydroxylation, 600081
CYP2U1	94,80%	91,50%	100%	99,90%	Spastic paraplegia 56, autosomal recessive, 615030
CYP7B1	98,00%	92,80%	100%	100%	Spastic paraplegia 5A, autosomal recessive, 270800 Bile acid synthesis defect, congenital, 3, 613812
D2HGDH	99,20%	97,20%	100%	100%	D-2-hydroxyglutaric aciduria, 600721
DAO	100%	100%	100%	100%	No OMIM disease ID
DBH	100%	100%	100%	100%	Orthostatic hypotension 1, due to DBH deficiency, 223360
DBT	99,80%	98,00%	100%	100%	Maple syrup urine disease, type II, 248600
DCXR	98,60%	93,60%	100%	100%	No OMIM disease ID
DDC	99,70%	96,40%	100%	100%	Aromatic L-amino acid decarboxylase deficiency, 608643
DDHD1	97,90%	95,80%	100%	100%	Spastic paraplegia 28, autosomal recessive, 609340
DDOST	100%	99,90%	100%	100%	?Congenital disorder of glycosylation, type I _r , 614507

<i>DEGS1</i>	100%	100%	100%	100%	Leukodystrophy, hypomyelinating, 18, 618404
<i>DGAT1</i>	91,90%	87,60%	99,70%	98,60%	?Diarrhea 7, protein-losing enteropathy type, 615863
<i>DGKE</i>	99,80%	98,10%	100%	100%	Nephrotic syndrome, type 7, 615008
<i>DGUOK</i>	100%	99,40%	100%	100%	Progressive external ophthalmoplegia with mitochondrial DNA deletions, autosomal recessive 4, 617070 Portal hypertension, noncirrhotic, 617068 Mitochondrial DNA depletion syndrome 3 (hepatocerebral type), 251880
<i>DHCR24</i>	100%	100%	100%	100%	Desmosterolosis, 602398
<i>DHCR7</i>	100%	100%	100%	100%	Smith-Lemli-Opitz syndrome, 270400
<i>DHDDS</i>	99,00%	95,00%	95,20%	95,20%	Retinitis pigmentosa 59, 613861 ?Congenital disorder of glycosylation, type 1bb, 613861 Developmental delay and seizures with or without movement abnormalities, 617836
<i>DHFR</i>	92,10%	78,90%	100%	100%	Megaloblastic anemia due to dihydrofolate reductase deficiency, 613839
<i>DHODH</i>	100%	100%	100%	100%	Miller syndrome, 263750
<i>DLD</i>	100%	99,70%	100%	100%	Dihydrolipoamide dehydrogenase deficiency, 246900
<i>DMGDH</i>	100%	99,70%	100%	100%	Dimethylglycine dehydrogenase deficiency, 605850
<i>DNAJC12</i>	87,40%	87,40%	100%	100%	Hyperphenylalaninemia, mild, non-BH4-deficient, 617384
<i>DNAJC19</i>	98,90%	96,20%	100%	100%	3-methylglutaconic aciduria, type V, 610198
<i>DNM1L</i>	99,90%	98,50%	100%	100%	Encephalopathy, lethal, due to defective mitochondrial peroxisomal fission 1, 614388 Optic atrophy 5, 610708
<i>DNM2</i>	98,10%	94,50%	100%	100%	Lethal congenital contracture syndrome 5, 615368 Charcot-Marie-Tooth disease, axonal type 2M, 606482 Centronuclear myopathy 1, 160150 Charcot-Marie-Tooth disease, dominant intermediate B, 606482
<i>DNMT1</i>	99,20%	99,00%	99,70%	99,20%	Neuropathy, hereditary sensory, type IE, 614116 Cerebellar ataxia, deafness, and narcolepsy, autosomal dominant, 604121
<i>DNMT3B</i>	100%	100%	100%	100%	Immunodeficiency-centromeric instability-facial anomalies syndrome 1, 242860
<i>DOLK</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type Im, 610768
<i>DPAGT1</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type Ij, 608093 Myasthenic syndrome, congenital, 13, with tubular aggregates, 614750
<i>DPM1</i>	98,20%	91,30%	99,70%	97,10%	Congenital disorder of glycosylation, type Ie, 608799
<i>DPM2</i>	100%	98,70%	100%	100%	Congenital disorder of glycosylation, type Iu, 615042
<i>DPM3</i>	100%	100%	100%	100%	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 15, 612937
<i>DPYD</i>	99,70%	97,70%	100%	100%	Dihydropyrimidine dehydrogenase deficiency, 274270 5-fluorouracil toxicity, 274270
<i>DPYS</i>	100%	99,90%	100%	100%	Dihydropyrimidinuria, 222748
<i>DTYMK</i>	100%	99,80%	100%	100%	No OMIM disease ID
<i>EBP</i>	99,70%	95,80%	100%	100%	Chondrodysplasia punctata, X-linked dominant, 302960 MEND syndrome, 300960

<i>ECHS1</i>	99,90%	99,00%	100%	100%	Mitochondrial short-chain enoyl-CoA hydratase 1 deficiency, 616277
<i>ELOVL1</i>	99,80%	97,60%	100%	100%	Ichthyotic keratoderma, spasticity, hypomyelination, and dysmorphic facies, 618527
<i>ELOVL4</i>	100%	99,50%	100%	100%	Spinocerebellar ataxia 34, 133190 Stargardt disease 3, 600110 Ichthyosis, spastic quadriplegia, and mental retardation, 614457
<i>ENO3</i>	100%	99,90%	100%	100%	?Glycogen storage disease XIII, 612932
<i>EOGT</i>	79,40%	78,40%	91,90%	89,00%	Adams-Oliver syndrome 4, 615297
<i>EPHX1</i>	99,90%	98,80%	100%	100%	?Hypercholanemia, familial, 607748
<i>EPHX2</i>	99,50%	96,20%	100%	100%	No OMIM disease ID
<i>ETFA</i>	100%	100%	100%	100%	Glutaric acidemia IIA, 231680
<i>ETFB</i>	100%	99,80%	100%	100%	Glutaric acidemia IIB, 231680
<i>ETFDH</i>	100%	99,80%	100%	100%	Glutaric acidemia IIC, 231680
<i>ETHE1</i>	99,90%	97,40%	100%	100%	Ethylmalonic encephalopathy, 602473
<i>EXT1</i>	99,90%	98,40%	100%	100%	Exostoses, multiple, type 1, 133700 Chondrosarcoma, 215300
<i>EXT2</i>	100%	99,30%	100%	100%	Exostoses, multiple, type 2, 133701 Seizures, scoliosis, and macrocephaly syndrome, 616682
<i>FA2H</i>	92,00%	83,10%	100%	100%	Spastic paraplegia 35, autosomal recessive, 612319
<i>FAH</i>	100%	100%	100%	100%	Tyrosinemia, type I, 276700
<i>FAR1</i>	97,60%	92,80%	100%	100%	Peroxisomal fatty acyl-CoA reductase 1 disorder, 616154
<i>FBP1</i>	100%	99,50%	100%	100%	Fructose-1,6-bisphosphatase deficiency, 229700
<i>FDFT1</i>	97,70%	96,00%	100%	100%	Squalene synthase deficiency, 618156
<i>FECH</i>	100%	100%	100%	100%	Protoporphyrinemia, erythropoietic, 1, 177000
<i>FH</i>	92,10%	88,30%	100%	100%	Fumarase deficiency, 606812 Leiomyomatosis and renal cell cancer, 150800
<i>FKRP</i>	100%	100%	100%	99,90%	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 5, 607155 Muscular dystrophy-dystroglycanopathy (congenital with or without mental retardation), type B, 5, 606612 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 5, 613153
<i>FKTN</i>	99,70%	97,00%	100%	100%	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 4, 611588 Cardiomyopathy, dilated, 1X, 611615 Muscular dystrophy-dystroglycanopathy (congenital without mental retardation), type B, 4, 613152 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 4, 253800
<i>FLAD1</i>	100%	99,80%	100%	100%	Lipid storage myopathy due to flavin adenine dinucleotide synthetase deficiency, 255100
<i>FMO3</i>	99,90%	99,70%	100%	100%	Trimethylaminuria, 602079
<i>FOLR1</i>	100%	100%	100%	100%	Neurodegeneration due to cerebral folate transport deficiency, 613068
<i>FTCD</i>	95,70%	91,00%	100%	100%	Glutamate formiminotransferase deficiency, 229100

<i>FUCA1</i>	100%	99,90%	100%	100%	Fucosidosis, 230000
<i>FUK</i>	97,70%	95,40%	100%	100%	Congenital disorder of glycosylation with defective fucosylation 2, 618324
<i>FUT2</i>	100%	100%	100%	100%	No OMIM disease ID
<i>FUT6</i>	100%	100%	100%	100%	Fucosyltransferase 6 deficiency, 613852
<i>FUT8</i>	100%	99,20%	100%	100%	Congenital disorder of glycosylation with defective fucosylation 1, 618005
<i>G6PC</i>	100%	100%	100%	100%	Glycogen storage disease Ia, 232200
<i>G6PC3</i>	100%	99,90%	100%	100%	Dursun syndrome, 612541 Neutropenia, severe congenital 4, autosomal recessive, 612541
<i>G6PD</i>	99,30%	98,10%	100%	99,30%	Hemolytic anemia, G6PD deficient (favism), 300908
<i>GAA</i>	100%	99,90%	100%	100%	Glycogen storage disease II, 232300
<i>GAD1</i>	100%	99,90%	100%	100%	?Cerebral palsy, spastic quadriplegic, 1, 603513
<i>GALC</i>	99,80%	98,30%	100%	100%	Krabbe disease, 245200
<i>GALE</i>	100%	100%	100%	100%	Galactose epimerase deficiency, 230350
<i>GALK1</i>	100%	99,10%	100%	100%	Galactokinase deficiency with cataracts, 230200
<i>GALM</i>	100%	99,90%	100%	100%	No OMIM disease ID
<i>GALNS</i>	100%	99,80%	100%	100%	Mucopolysaccharidosis IVA, 253000
<i>GALNT3</i>	99,80%	99,00%	100%	100%	Tumoral calcinosis, hyperphosphatemic, familial, 1, 211900
<i>GALT</i>	100%	99,70%	100%	100%	Galactosemia, 230400
<i>GAMT</i>	93,10%	82,70%	100%	100%	Cerebral creatine deficiency syndrome 2, 612736
<i>GANAB</i>	99,90%	99,00%	100%	100%	Polycystic kidney disease 3, 600666
<i>GATM</i>	100%	100%	100%	100%	Cerebral creatine deficiency syndrome 3, 612718
<i>GBA</i>	100%	100%	100%	100%	Gaucher disease, type III, 231000 Gaucher disease, type IIIC, 231005 Gaucher disease, type I, 230800 Gaucher disease, perinatal lethal, 608013 Gaucher disease, type II, 230900
<i>GBA2</i>	100%	99,70%	100%	100%	Spastic paraplegia 46, autosomal recessive, 614409
<i>GBE1</i>	100%	99,60%	100%	100%	Polyglucosan body disease, adult form, 263570 Glycogen storage disease IV, 232500
<i>GCDH</i>	100%	99,20%	100%	100%	Glutaricaciduria, type I, 231670
<i>GCH1</i>	99,90%	95,50%	100%	100%	Hyperphenylalaninemia, BH4-deficient, B, 233910 Dystonia, DOPA-responsive, with or without hyperphenylalaninemia, 128230
<i>GCK</i>	100%	100%	100%	100%	Diabetes mellitus, noninsulin-dependent, late onset, 125853 Diabetes mellitus, permanent neonatal, 606176 MODY, type II, 125851 Hyperinsulinemic hypoglycemia, familial, 3, 602485
<i>GCLC</i>	99,80%	98,00%	100%	100%	Hemolytic anemia due to gamma-glutamylcysteine synthetase deficiency, 230450

<i>GCLM</i>	99,60%	95,80%	100%	100%	No OMIM disease ID
<i>GCSH</i>	75,70%	68,90%	100%	100%	?Glycine encephalopathy, 605899
<i>GFPT1</i>	100%	99,40%	100%	100%	Myasthenia, congenital, 12, with tubular aggregates, 610542
<i>GIF</i>	100%	99,70%	100%	100%	Intrinsic factor deficiency, 261000
<i>GK</i>	88,90%	70,40%	100%	99,90%	Glycerol kinase deficiency, 307030
<i>GLA</i>	99,80%	96,60%	100%	100%	Fabry disease, 301500 Fabry disease, cardiac variant, 301500
<i>GLB1</i>	99,90%	97,40%	100%	100%	GM1-gangliosidosis, type III, 230650 GM1-gangliosidosis, type I, 230500 Mucopolysaccharidosis type IVB (Morquio), 253010 GM1-gangliosidosis, type II, 230600
<i>GLDC</i>	89,90%	82,00%	100%	99,90%	Glycine encephalopathy, 605899
<i>GLRA1</i>	100%	99,80%	100%	100%	Hyperekplexia 1, 149400
<i>GLRX5</i>	97,30%	89,10%	99,60%	95,40%	Anemia, sideroblastic, 3, pyridoxine-refractory, 616860 Spasticity, childhood-onset, with hyperglycinemia, 616859
<i>GLS</i>	96,30%	87,20%	100%	99,90%	?Infantile cataract, skin abnormalities, glutamate excess, and impaired intellectual development, 618339 Global developmental delay, progressive ataxia, and elevated glutamine, 618412 Epileptic encephalopathy, early infantile, 71, 618328
<i>GLUD1</i>	94,20%	82,90%	100%	100%	Hyperinsulinism-hyperammonemia syndrome, 606762
<i>GLUL</i>	99,90%	98,20%	100%	100%	Glutamine deficiency, congenital, 610015
<i>GLYCK</i>	98,80%	97,30%	100%	100%	D-glyceric aciduria, 220120
<i>GM2A</i>	100%	100%	100%	100%	GM2-gangliosidosis, AB variant, 272750
<i>GMPPA</i>	100%	100%	100%	100%	Alacrima, achalasia, and mental retardation syndrome, 615510
<i>GMPPB</i>	100%	100%	100%	100%	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 14, 615350 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 14, 615351 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 14, 615352
<i>GMPS</i>	99,10%	96,10%	100%	100%	No OMIM disease ID
<i>GNE</i>	100%	99,70%	100%	100%	Sialuria, 269921 Nonaka myopathy, 605820
<i>GNMT</i>	100%	100%	100%	100%	Glycine N-methyltransferase deficiency, 606664
<i>GNPAT</i>	99,70%	97,30%	100%	100%	Rhizomelic chondrodysplasia punctata, type 2, 222765
<i>GNPTAB</i>	100%	99,90%	100%	100%	Mucopolysaccharidosis II alpha/beta, 252500 Mucopolysaccharidosis III alpha/beta, 252600
<i>GNPTG</i>	99,10%	94,30%	100%	99,90%	Mucopolysaccharidosis III gamma, 252605
<i>GNS</i>	98,40%	94,80%	100%	100%	Mucopolysaccharidosis type IIID, 252940
<i>GOT1</i>	100%	99,30%	100%	100%	Aspartate aminotransferase, serum level of, QTL1, 614419
<i>GOT2</i>	97,50%	90,90%	100%	100%	Epileptic encephalopathy, early infantile, 82, 618721

<i>GPD1</i>	100%	100%	100%	100%	Hypertriglyceridemia, transient infantile, 614480
<i>GPD1L</i>	100%	99,80%	100%	100%	Brugada syndrome 2, 611777
<i>GPHN</i>	100%	99,50%	100%	100%	Molybdenum cofactor deficiency C, 615501
<i>GPI</i>	100%	100%	100%	100%	Hemolytic anemia, nonspherocytic, due to glucose phosphate isomerase deficiency, 613470
<i>GPT2</i>	99,20%	93,60%	100%	99,80%	Mental retardation, autosomal recessive 49, 616281
<i>GPX1</i>	95,40%	84,60%	100%	100%	No OMIM disease ID
<i>GRHPR</i>	84,20%	81,30%	100%	99,30%	Hyperoxaluria, primary, type II, 260000
<i>GSS</i>	100%	99,90%	100%	100%	Glutathione synthetase deficiency, 266130 Hemolytic anemia due to glutathione synthetase deficiency, 231900
<i>GUSB</i>	92,90%	91,70%	100%	100%	Mucopolysaccharidosis VII, 253220
<i>GYG1</i>	99,90%	99,20%	100%	100%	?Glycogen storage disease XV, 613507 Polyglucosan body myopathy 2, 616199
<i>GYS1</i>	100%	98,60%	100%	100%	Glycogen storage disease 0, muscle, 611556
<i>GYS2</i>	99,80%	99,00%	100%	100%	Glycogen storage disease 0, liver, 240600
<i>H6PD</i>	99,00%	99,00%	100%	100%	Cortisone reductase deficiency 1, 604931
<i>HADH</i>	99,00%	97,50%	100%	100%	3-hydroxyacyl-CoA dehydrogenase deficiency, 231530 Hyperinsulinemic hypoglycemia, familial, 4, 609975
<i>HADHA</i>	97,10%	91,30%	100%	100%	LCHAD deficiency, 609016 HELLP syndrome, maternal, of pregnancy, 609016 Fatty liver, acute, of pregnancy, 609016 Trifunctional protein deficiency, 609015
<i>HADHB</i>	98,80%	89,70%	100%	100%	Trifunctional protein deficiency, 609015
<i>HAGH</i>	100%	99,70%	98,70%	96,10%	No OMIM disease ID
<i>HEXA</i>	93,80%	93,30%	100%	100%	GM2-gangliosidosis, several forms, 272800 Tay-Sachs disease, 272800
<i>HEXB</i>	99,60%	96,90%	100%	99,90%	Sandhoff disease, infantile, juvenile, and adult forms, 268800
<i>HFE</i>	100%	99,70%	100%	100%	Hemochromatosis, 235200
<i>HGD</i>	100%	100%	100%	100%	Alkaptonuria, 203500
<i>HGSNAT</i>	86,40%	86,30%	91,20%	89,30%	Retinitis pigmentosa 73, 616544 Mucopolysaccharidosis type IIIC (Sanfilippo C), 252930
<i>HIBADH</i>	94,40%	91,20%	100%	100%	No OMIM disease ID
<i>HIBCH</i>	98,20%	88,50%	100%	100%	3-hydroxyisobutyryl-CoA hydrolase deficiency, 250620
<i>HK1</i>	100%	100%	100%	100%	Hemolytic anemia due to hexokinase deficiency, 235700 Neuropathy, hereditary motor and sensory, Russe type, 605285 Neurodevelopmental disorder with visual defects and brain anomalies, 618547 Retinitis pigmentosa 79, 617460
<i>HLCS</i>	100%	100%	100%	100%	Holocarboxylase synthetase deficiency, 253270

<i>HMBS</i>	99,90%	99,40%	100%	100%	Porphyria, acute intermittent, 176000 Porphyria, acute intermittent, nonerythroid variant, 176000
<i>HMGCL</i>	100%	99,80%	100%	100%	HMG-CoA lyase deficiency, 246450
<i>HMGCS2</i>	100%	99,60%	100%	100%	HMG-CoA synthase-2 deficiency, 605911
<i>HMOX1</i>	98,40%	89,90%	100%	100%	Heme oxygenase-1 deficiency, 614034
<i>HNF1A</i>	100%	99,80%	100%	100%	MODY, type III, 600496 Hepatic adenoma, somatic, 142330 Renal cell carcinoma, 144700 Diabetes mellitus, insulin-dependent, 20, 612520
<i>HNF4A</i>	99,90%	99,00%	100%	100%	MODY, type I, 125850 Fanconi renotubular syndrome 4, with maturity-onset diabetes of the young, 616026
<i>HOGA1</i>	100%	96,40%	100%	100%	Hyperoxaluria, primary, type III, 613616
<i>HPD</i>	100%	100%	100%	100%	Tyrosinemia, type III, 276710 Hawkinsinuria, 140350
<i>HPRT1</i>	99,30%	91,80%	100%	99,30%	HPRT-related gout, 300323 Lesch-Nyhan syndrome, 300322
<i>HS6ST1</i>	92,90%	84,50%	100%	100%	No OMIM disease ID
<i>HSD11B1</i>	100%	99,60%	100%	100%	Cortisone reductase deficiency 2, 614662
<i>HSD11B2</i>	86,00%	82,70%	99,90%	98,10%	Apparent mineralocorticoid excess, 218030
<i>HSD17B10</i>	100%	99,10%	100%	100%	HSD10 mitochondrial disease, 300438
<i>HSD17B3</i>	100%	100%	100%	100%	Pseudohermaphroditism, male, with gynecomastia, 264300
<i>HSD17B4</i>	96,00%	93,70%	96,60%	96,60%	D-bifunctional protein deficiency, 261515 Perrault syndrome 1, 233400
<i>HSD3B2</i>	100%	99,70%	100%	100%	Adrenal hyperplasia, congenital, due to 3-beta-hydroxysteroid dehydrogenase 2 deficiency, 201810
<i>HSD3B7</i>	99,10%	95,50%	100%	100%	Bile acid synthesis defect, congenital, 1, 607765
<i>HTRA2</i>	100%	99,90%	100%	100%	3-methylglutaconic aciduria, type VIII, 617248
<i>HYAL1</i>	100%	100%	100%	100%	?Mucopolysaccharidosis type IX, 601492
<i>IDH2</i>	99,70%	97,40%	100%	99,80%	D-2-hydroxyglutaric aciduria 2, 613657
<i>IDH3B</i>	95,40%	95,40%	100%	100%	Retinitis pigmentosa 46, 612572
<i>IDI1</i>	99,80%	97,60%	100%	100%	No OMIM disease ID
<i>IDS</i>	99,90%	98,00%	100%	100%	Mucopolysaccharidosis II, 309900
<i>IDUA</i>	93,70%	86,80%	100%	100%	Mucopolysaccharidosis Ih/s, 607015 Mucopolysaccharidosis Ih, 607014 Mucopolysaccharidosis Is, 607016
<i>IMPAD1</i>	100%	100%	100%	100%	Chondrodysplasia with joint dislocations, GPAPP type, 614078
<i>IMPDH1</i>	87,90%	80,20%	100%	100%	Leber congenital amaurosis 11, 613837 Retinitis pigmentosa 10, 180105

<i>INPP5E</i>	97,10%	92,70%	100%	100%	Mental retardation, truncal obesity, retinal dystrophy, and micropenis, 610156 Joubert syndrome 1, 213300
<i>INPPL1</i>	98,40%	94,50%	99,90%	99,70%	Opsismodysplasia, 258480
<i>INSR</i>	97,80%	94,70%	99,90%	99,20%	Hyperinsulinemic hypoglycemia, familial, 5, 609968 Rabson-Mendenhall syndrome, 262190 Diabetes mellitus, insulin-resistant, with acanthosis nigricans, 610549 Leprechaunism, 246200
<i>MR PL44</i>	99,90%	98,70%	100%	100%	?Combined oxidative phosphorylation deficiency 16, 615395
<i>MR PS36</i>	95,20%	77,60%	100%	100%	No OMIM disease ID
<i>IREB2</i>	100%	99,80%	100%	100%	Neurodegeneration, early-onset, with choreoathetoid movements and microcytic anemia, 618451
<i>ISPD</i>	98,50%	94,80%	100%	99,40%	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 7, 616052 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 7, 614643
<i>ITCH</i>	95,40%	95,20%	100%	99,00%	Autoimmune disease, multisystem, with facial dysmorphism, 613385
<i>ITPA</i>	100%	100%	100%	100%	Epileptic encephalopathy, early infantile, 35, 616647
<i>IVD</i>	100%	100%	100%	100%	Isovaleric acidemia, 243500
<i>KCNA2</i>	100%	99,60%	100%	100%	Epileptic encephalopathy, early infantile, 32, 616366
<i>KCNJ11</i>	100%	100%	100%	100%	Maturity-onset diabetes of the young, type 13, 616329 Diabetes, permanent neonatal, with or without neurologic features, 606176 Diabetes mellitus, transient neonatal, 3, 610582 Hyperinsulinemic hypoglycemia, familial, 2, 601820
<i>KMT2A</i>	100%	99,90%	99,90%	99,40%	Wiedemann-Steiner syndrome, 605130
<i>KMT2D</i>	100%	99,40%	100%	100%	Kabuki syndrome 1, 147920
<i>L2HGDH</i>	99,00%	97,20%	100%	100%	L-2-hydroxyglutaric aciduria, 236792
<i>LAMP2</i>	99,20%	95,60%	100%	100%	Danon disease, 300257
<i>LARGE1</i>	100%	99,60%	100%	100%	Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 6, 608840 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 6, 613154
<i>LCAT</i>	99,00%	93,80%	100%	100%	Norum disease, 245900 Fish-eye disease, 136120
<i>LCT</i>	99,80%	98,50%	100%	100%	Lactase deficiency, congenital, 223000
<i>LDHA</i>	95,00%	91,70%	100%	100%	Glycogen storage disease XI, 612933
<i>LDHB</i>	96,30%	86,00%	100%	100%	No OMIM disease ID
<i>LFNG</i>	87,90%	86,40%	92,20%	87,70%	Spondylocostal dysostosis 3, autosomal recessive, 609813
<i>LIAS</i>	100%	99,10%	100%	100%	Hyperglycinemia, lactic acidosis, and seizures, 614462
<i>LIPA</i>	99,20%	95,20%	95,20%	95,20%	Wolman disease, 278000 Cholesteryl ester storage disease, 278000
<i>LIPC</i>	100%	99,40%	100%	100%	Hepatic lipase deficiency, 614025

<i>LIPT1</i>	100%	99,90%	100%	100%	Lipoyltransferase 1 deficiency, 616299
<i>LIPT2</i>	94,90%	75,20%	100%	100%	Encephalopathy, neonatal severe, with lactic acidosis and brain abnormalities, 617668
<i>LMBRD1</i>	98,50%	93,90%	100%	100%	Methylmalonic aciduria and homocystinuria, cb1F type, 277380
<i>LPIN1</i>	99,60%	97,30%	100%	100%	Myoglobinuria, acute recurrent, autosomal recessive, 268200
<i>LPIN2</i>	100%	100%	100%	100%	Majeed syndrome, 609628
<i>LPL</i>	100%	100%	100%	100%	Lipoprotein lipase deficiency, 238600 Combined hyperlipidemia, familial, 144250
<i>LRAT</i>	100%	100%	100%	100%	Retinal dystrophy, early-onset severe, 613341 Leber congenital amaurosis 14, 613341 Retinitis pigmentosa, juvenile, 613341
<i>LTC4S</i>	74,20%	68,50%	100%	100%	No OMIM disease ID
<i>LYST</i>	99,60%	98,30%	100%	100%	Chediak-Higashi syndrome, 214500
<i>MAN1B1</i>	100%	99,70%	100%	99,90%	Mental retardation, autosomal recessive 15, 614202
<i>MAN2B1</i>	99,80%	97,90%	100%	100%	Mannosidosis, alpha-, types I and II, 248500
<i>MANBA</i>	99,80%	98,40%	100%	100%	Mannosidosis, beta, 248510
<i>MAOA</i>	100%	99,70%	99,80%	98,50%	Brunner syndrome, 300615
<i>MAT1A</i>	99,70%	97,70%	100%	100%	Methionine adenosyltransferase deficiency, autosomal recessive, 250850 Hypermethioninemia, persistent, autosomal dominant, due to methionine adenosyltransferase I/III deficiency, 250850
<i>MBOAT7</i>	100%	99,50%	100%	100%	Mental retardation, autosomal recessive 57, 617188
<i>MCCC1</i>	100%	99,80%	100%	100%	3-Methylcrotonyl-CoA carboxylase 1 deficiency, 210200
<i>MCCC2</i>	99,90%	98,40%	100%	100%	3-Methylcrotonyl-CoA carboxylase 2 deficiency, 210210
<i>MCEE</i>	100%	100%	100%	100%	Methylmalonyl-CoA epimerase deficiency, 251120
<i>MCOLN1</i>	99,80%	98,40%	100%	100%	Mucopolidosis IV, 252650
<i>MDH1</i>	100%	99,40%	100%	100%	No OMIM disease ID
<i>MFSD2A</i>	99,70%	98,50%	100%	100%	Microcephaly 15, primary, autosomal recessive, 616486
<i>MFSD8</i>	100%	99,70%	100%	100%	Macular dystrophy with central cone involvement, 616170 Ceroid lipofuscinosis, neuronal, 7, 610951
<i>MGAT2</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type IIa, 212066
<i>MINPP1</i>	100%	99,50%	100%	100%	No OMIM disease ID
<i>MLYCD</i>	96,00%	90,40%	100%	98,90%	Malonyl-CoA decarboxylase deficiency, 248360
<i>MMAA</i>	100%	100%	100%	100%	Methylmalonic aciduria, vitamin B12-responsive, 251100
<i>MMAB</i>	100%	99,60%	100%	100%	Methylmalonic aciduria, vitamin B12-responsive, due to defect in synthesis of adenosylcobalamin, cb1B complementation type, 251110
<i>MMACHC</i>	100%	100%	100%	100%	Methylmalonic aciduria and homocystinuria, cb1C type, 277400

<i>MMADHC</i>	94,40%	83,50%	89,70%	89,70%	Homocystinuria, cbID type, variant 1, 277410 Methylmalonic aciduria and homocystinuria, cbID type, 277410 Methylmalonic aciduria, cbID type, variant 2, 277410
<i>MOCOS</i>	99,80%	97,70%	100%	100%	Xanthinuria, type II, 603592
<i>MOCS1</i>	99,20%	95,40%	100%	100%	Molybdenum cofactor deficiency A, 252150
<i>MOCS2</i>	99,60%	99,50%	100%	100%	Molybdenum cofactor deficiency B, 252160
<i>MOGS</i>	100%	99,90%	100%	100%	Congenital disorder of glycosylation, type IIb, 606056
<i>MPDU1</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type If, 609180
<i>MPI</i>	100%	99,90%	100%	100%	Congenital disorder of glycosylation, type Ib, 602579
<i>MSMO1</i>	96,30%	88,90%	100%	100%	Microcephaly, congenital cataract, and psoriasiform dermatitis, 616834
<i>MTHFD1</i>	100%	99,50%	100%	100%	Combined immunodeficiency and megaloblastic anemia with or without hyperhomocysteinemia, 617780
<i>MTHFR</i>	97,30%	96,00%	100%	100%	Homocystinuria due to MTHFR deficiency, 236250
<i>MTMR2</i>	100%	99,00%	100%	100%	Charcot-Marie-Tooth disease, type 4B1, 601382
<i>MTM1</i>	99,00%	93,30%	100%	100%	Myotubular myopathy, X-linked, 310400
<i>MTR</i>	100%	100%	100%	100%	Homocystinuria-megaloblastic anemia, cblG complementation type, 250940
<i>MTRR</i>	100%	99,60%	100%	100%	Homocystinuria-megaloblastic anemia, cbl E type, 236270
<i>MUT</i>	99,80%	98,30%	100%	100%	Methylmalonic aciduria, mut(0) type, 251000
<i>MVK</i>	90,90%	90,50%	90,50%	90,50%	Hyper-IgD syndrome, 260920 Porokeratosis 3, multiple types, 175900 Mevalonic aciduria, 610377
<i>NADK2</i>	99,90%	97,20%	99,00%	96,30%	2,4-dienoyl-CoA reductase deficiency, 616034
<i>NAGA</i>	100%	100%	100%	100%	Kanzaki disease, 609242 Schindler disease, type I, 609241 Schindler disease, type III, 609241
<i>NAGLU</i>	92,90%	89,90%	99,90%	99,20%	Mucopolysaccharidosis type IIIB (Sanfilippo B), 252920 ?Charcot-Marie-Tooth disease, axonal, type 2V, 616491
<i>NAGS</i>	99,70%	95,00%	100%	100%	N-acetylglutamate synthase deficiency, 237310
<i>NANS</i>	100%	99,90%	100%	100%	Spondyloepimetaphyseal dysplasia, Camera-Genevieve type, 610442
<i>NAXE</i>	100%	99,80%	100%	100%	Encephalopathy, progressive, early-onset, with brain edema and/or leukoencephalopathy, 617186
<i>NBAS</i>	100%	99,60%	100%	100%	Infantile liver failure syndrome 2, 616483 Short stature, optic nerve atrophy, and Pelger-Huet anomaly, 614800
<i>NEU1</i>	99,70%	97,70%	100%	100%	Sialidosis, type II, 256550 Sialidosis, type I, 256550
<i>NGLY1</i>	100%	99,80%	100%	100%	Congenital disorder of deglycosylation, 615273
<i>NMNAT1</i>	100%	99,20%	98,30%	95,60%	Leber congenital amaurosis 9, 608553
<i>NNT</i>	100%	99,40%	100%	100%	Glucocorticoid deficiency 4, with or without mineralocorticoid deficiency, 614736

<i>NPC1</i>	99,60%	98,70%	100%	100%	Niemann-Pick disease, type D, 257220 Niemann-Pick disease, type C1, 257220
<i>NPC2</i>	100%	99,60%	100%	100%	Niemann-pick disease, type C2, 607625
<i>NPL</i>	100%	99,90%	100%	100%	No OMIM disease ID
<i>NSD1</i>	100%	99,90%	100%	100%	Sotos syndrome 1, 117550
<i>NSDHL</i>	100%	98,70%	100%	100%	CHILD syndrome, 308050 CK syndrome, 300831
<i>NT5C3A</i>	97,90%	88,30%	100%	100%	Anemia, hemolytic, due to UMPH1 deficiency, 266120
<i>NT5E</i>	100%	99,90%	100%	100%	Calcification of joints and arteries, 211800
<i>NUS1</i>	60,00%	44,50%	100%	100%	Mental retardation, autosomal dominant 55, with seizures, 617831 ?Congenital disorder of glycosylation, type 1aa, 617082
<i>OAT</i>	85,20%	76,30%	100%	100%	Gyrate atrophy of choroid and retina with or without ornithinemia, 258870
<i>OCRL</i>	99,90%	98,60%	100%	99,90%	Lowe syndrome, 309000 Dent disease 2, 300555
<i>OPA3</i>	100%	99,00%	100%	100%	Optic atrophy 3 with cataract, 165300 3-methylglutaconic aciduria, type III, 258501
<i>OPLAH</i>	100%	99,80%	100%	100%	5-oxoprolinase deficiency, 260005
<i>OTC</i>	100%	100%	100%	100%	Ornithine transcarbamylase deficiency, 311250
<i>OXCT1</i>	99,80%	98,10%	100%	100%	Succinyl CoA:3-oxoacid CoA transferase deficiency, 245050
<i>PAH</i>	100%	100%	100%	100%	Phenylketonuria, 261600
<i>PANK2</i>	100%	99,30%	100%	100%	HARP syndrome, 607236 Neurodegeneration with brain iron accumulation 1, 234200
<i>PC</i>	99,80%	97,30%	100%	100%	Pyruvate carboxylase deficiency, 266150
<i>PCBD1</i>	100%	99,60%	100%	99,70%	Hyperphenylalaninemia, BH4-deficient, D, 264070
<i>PCCA</i>	99,50%	96,70%	100%	100%	Propionicacidemia, 606054
<i>PCCB</i>	97,90%	96,00%	98,70%	96,20%	Propionicacidemia, 606054
<i>PCK1</i>	100%	100%	100%	100%	?Phosphoenolpyruvate carboxykinase deficiency, cytosolic, 261680
<i>PCK2</i>	100%	100%	100%	100%	No OMIM disease ID
<i>PCYT1A</i>	98,90%	95,50%	100%	100%	Spondylometaphyseal dysplasia with cone-rod dystrophy, 608940
<i>PCYT2</i>	99,80%	97,10%	100%	98,80%	Spastic paraplegia 82, autosomal recessive, 618770
<i>PDSS1</i>	94,70%	87,60%	97,30%	96,60%	Coenzyme Q10 deficiency, primary, 2, 614651
<i>PDSS2</i>	99,80%	97,10%	100%	100%	Coenzyme Q10 deficiency, primary, 3, 614652
<i>PEPD</i>	100%	98,80%	100%	100%	Prolidase deficiency, 170100
<i>PEX1</i>	99,90%	99,40%	100%	100%	Heimler syndrome 1, 234580 Peroxisome biogenesis disorder 1B (NALD/IRD), 601539 Peroxisome biogenesis disorder 1A (Zellweger), 214100

<i>PEX10</i>	96,80%	89,70%	100%	99,90%	Peroxisome biogenesis disorder 6B, 614871 Peroxisome biogenesis disorder 6A (Zellweger), 614870
<i>PEX11B</i>	100%	99,60%	100%	100%	?Peroxisome biogenesis disorder 14B, 614920
<i>PEX12</i>	100%	100%	100%	100%	Peroxisome biogenesis disorder 3A (Zellweger), 614859 Peroxisome biogenesis disorder 3B, 266510
<i>PEX13</i>	100%	100%	100%	100%	Peroxisome biogenesis disorder 11A (Zellweger), 614883 Peroxisome biogenesis disorder 11B, 614885
<i>PEX14</i>	96,70%	90,80%	100%	100%	Peroxisome biogenesis disorder 13A (Zellweger), 614887
<i>PEX16</i>	97,90%	94,20%	100%	100%	Peroxisome biogenesis disorder 8A (Zellweger), 614876 Peroxisome biogenesis disorder 8B, 614877
<i>PEX19</i>	99,90%	98,50%	100%	100%	Peroxisome biogenesis disorder 12A (Zellweger), 614886
<i>PEX2</i>	100%	100%	100%	100%	Peroxisome biogenesis disorder 5A (Zellweger), 614866 Peroxisome biogenesis disorder 5B, 614867
<i>PEX26</i>	100%	100%	100%	100%	Peroxisome biogenesis disorder 7A (Zellweger), 614872 Peroxisome biogenesis disorder 7B, 614873
<i>PEX3</i>	100%	99,30%	100%	100%	Peroxisome biogenesis disorder 10A (Zellweger), 614882 ?Peroxisome biogenesis disorder 10B, 617370
<i>PEX5</i>	99,90%	99,00%	100%	100%	Peroxisome biogenesis disorder 2B, 202370 Rhizomelic chondrodysplasia punctata, type 5, 616716 Peroxisome biogenesis disorder 2A (Zellweger), 214110
<i>PEX6</i>	94,50%	86,70%	100%	100%	Peroxisome biogenesis disorder 4B, 614863 Heimler syndrome 2, 616617 Peroxisome biogenesis disorder 4A (Zellweger), 614862
<i>PEX7</i>	87,80%	80,70%	91,30%	91,30%	Peroxisome biogenesis disorder 9B, 614879 Rhizomelic chondrodysplasia punctata, type 1, 215100
<i>PFKM</i>	100%	99,50%	100%	100%	Glycogen storage disease VII, 232800
<i>PGAM2</i>	100%	100%	100%	100%	Glycogen storage disease X, 261670
<i>PGAP1</i>	99,00%	94,40%	100%	100%	Mental retardation, autosomal recessive 42, 615802
<i>PGAP2</i>	100%	99,90%	100%	100%	Hyperphosphatasia with mental retardation syndrome 3, 614207
<i>PGAP3</i>	63,50%	59,60%	100%	100%	Hyperphosphatasia with mental retardation syndrome 4, 615716
<i>PGK1</i>	92,80%	79,30%	100%	100%	Phosphoglycerate kinase 1 deficiency, 300653
<i>PGM1</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type It, 614921
<i>PGM3</i>	100%	99,80%	100%	100%	Immunodeficiency 23, 615816
<i>PHGDH</i>	99,90%	98,80%	100%	100%	Neu-Laxova syndrome 1, 256520 Phosphoglycerate dehydrogenase deficiency, 601815
<i>PHKA1</i>	99,20%	95,30%	100%	99,90%	Muscle glycogenosis, 300559
<i>PHKA2</i>	100%	99,70%	100%	99,60%	Glycogen storage disease, type IXa2, 306000 Glycogen storage disease, type IXa1, 306000

<i>PHKB</i>	99,90%	99,20%	100%	100%	Phosphorylase kinase deficiency of liver and muscle, autosomal recessive, 261750
<i>PHKG1</i>	99,90%	97,80%	100%	100%	No OMIM disease ID
<i>PHKG2</i>	100%	99,90%	100%	100%	Glycogen storage disease IXc, 613027 Cirrhosis due to liver phosphorylase kinase deficiency, 0
<i>PHYH</i>	100%	99,60%	100%	100%	Refsum disease, 266500
<i>TALDO1</i>	100%	97,90%	100%	100%	Transaldolase deficiency, 606003
<i>TANGO2</i>	100%	99,30%	100%	100%	metabolic encephalomyopathic crises, recurrent, with rhabdomyolysis, cardiac arrhythmias, and neurodegeneration, 616878
<i>TAT</i>	100%	100%	100%	100%	Tyrosinemia, type II, 276600
<i>TAZ</i>	99,20%	96,50%	100%	100%	Barth syndrome, 302060
<i>PIGA</i>	93,80%	86,70%	100%	100%	Multiple congenital anomalies-hypotonia-seizures syndrome 2, 300868 Paroxysmal nocturnal hemoglobinuria, somatic, 300818
<i>PIGB</i>	99,90%	97,80%	100%	100%	Epileptic encephalopathy, early infantile, 80, 618580
<i>PIGC</i>	99,20%	90,90%	100%	100%	Glycosylphosphatidylinositol biosynthesis defect 16, 617816
<i>PIGL</i>	100%	100%	100%	100%	CHIME syndrome, 280000
<i>PIGM</i>	100%	100%	100%	100%	Glycosylphosphatidylinositol deficiency, 610293
<i>PIGN</i>	93,80%	91,50%	98,80%	98,80%	Multiple congenital anomalies-hypotonia-seizures syndrome 1, 614080
<i>PIGO</i>	100%	99,90%	100%	100%	Hyperphosphatasia with mental retardation syndrome 2, 614749
<i>PIGP</i>	95,80%	87,30%	100%	100%	Epileptic encephalopathy, early infantile, 55, 617599
<i>PIGQ</i>	92,80%	90,80%	100%	100%	Epileptic encephalopathy, early infantile, 77, 618548
<i>PIGT</i>	98,10%	98,10%	100%	100%	?Paroxysmal nocturnal hemoglobinuria 2, 615399 Multiple congenital anomalies-hypotonia-seizures syndrome 3, 615398
<i>PIGV</i>	100%	100%	100%	100%	Hyperphosphatasia with mental retardation syndrome 1, 239300
<i>PIGW</i>	100%	99,80%	100%	100%	Glycosylphosphatidylinositol biosynthesis defect 11, 616025
<i>PIGY</i>	100%	99,90%	100%	100%	Hyperphosphatasia with mental retardation syndrome 6, 616809
<i>PIK3CA</i>	100%	99,80%	100%	100%	Ovarian cancer, somatic, 167000 Colorectal cancer, somatic, 114500 CLAPO syndrome, somatic, 613089 Cowden syndrome 5, 615108 Hepatocellular carcinoma, somatic, 114550 Breast cancer, somatic, 114480 Macrodactyly, somatic, 155500 Keratosis, seborrheic, somatic, 182000 Gastric cancer, somatic, 613659 Megalencephaly-capillary malformation-polymicrogyria syndrome, somatic, 602501 Nevus, epidermal, somatic, 162900 CLOVE syndrome, somatic, 612918 Nonsmall cell lung cancer, somatic, 211980

<i>PIK3R1</i>	99,80%	99,00%	100%	100%	SHORT syndrome, 269880 Immunodeficiency 36, 616005 ?Agammaglobulinemia 7, autosomal recessive, 615214
<i>PIK3R2</i>	90,70%	89,60%	99,30%	96,10%	Megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome 1, 603387
<i>PIK3R5</i>	100%	99,90%	100%	100%	Ataxia-oculomotor apraxia 3, 615217
<i>PIKFYVE</i>	99,90%	99,40%	100%	100%	Corneal fleck dystrophy, 121850
<i>PIP5K1C</i>	98,00%	95,80%	99,90%	99,80%	Lethal congenital contractural syndrome 3, 611369
<i>PKLR</i>	100%	99,20%	100%	100%	Pyruvate kinase deficiency, 266200 Adenosine triphosphate, elevated, of erythrocytes, 102900
<i>PLA2G5</i>	100%	100%	100%	100%	No OMIM disease ID
<i>PLA2G6</i>	99,90%	98,30%	100%	100%	Infantile neuroaxonal dystrophy 1, 256600 Parkinson disease 14, autosomal recessive, 612953 Neurodegeneration with brain iron accumulation 2B, 610217
<i>PLA2G7</i>	99,90%	99,00%	100%	100%	Platelet-activating factor acetylhydrolase deficiency, 614278
<i>PLCB1</i>	100%	99,80%	100%	100%	Epileptic encephalopathy, early infantile, 12, 613722
<i>PLCB4</i>	99,90%	98,80%	100%	100%	Auriculocondylar syndrome 2, 614669
<i>PLCD1</i>	99,90%	97,80%	100%	100%	Nail disorder, nonsyndromic congenital, 3, (leukonychia), 151600
<i>PLCE1</i>	99,90%	99,30%	100%	100%	Nephrotic syndrome, type 3, 610725
<i>PLCG2</i>	100%	99,80%	100%	100%	Familial cold autoinflammatory syndrome 3, 614468 Autoinflammation, antibody deficiency, and immune dysregulation syndrome, 614878
<i>PLIN1</i>	99,60%	94,90%	100%	99,50%	Lipodystrophy, familial partial, type 4, 613877
<i>PLOD1</i>	100%	98,40%	100%	100%	Ehlers-Danlos syndrome, kyphoscoliotic type, 1, 225400
<i>PLOD2</i>	99,30%	97,30%	100%	100%	Bruck syndrome 2, 609220
<i>PLOD3</i>	99,80%	98,00%	100%	100%	Lysyl hydroxylase 3 deficiency, 612394
<i>PLPBP</i>	98,20%	90,10%	100%	100%	Epilepsy, early-onset, vitamin B6-dependent, 617290
<i>PMM2</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type Ia, 212065
<i>PNLIP</i>	100%	99,80%	100%	100%	?Pancreatic lipase deficiency, 614338
<i>PNMT</i>	99,60%	96,70%	100%	100%	No OMIM disease ID
<i>PNP</i>	99,80%	98,90%	100%	100%	Immunodeficiency due to purine nucleoside phosphorylase deficiency, 613179
<i>PNPLA2</i>	99,70%	96,10%	100%	100%	Neutral lipid storage disease with myopathy, 610717
<i>PNPLA6</i>	100%	99,70%	100%	100%	Spastic paraplegia 39, autosomal recessive, 612020 Boucher-Neuhauser syndrome, 215470 Oliver-McFarlane syndrome, 275400 ?Laurence-Moon syndrome, 245800
<i>PNPO</i>	99,90%	97,70%	100%	100%	Pyridoxamine 5'-phosphate oxidase deficiency, 610090
<i>POFUT1</i>	100%	99,00%	100%	100%	Dowling-Degos disease 2, 615327

<i>POGLUT1</i>	99,40%	94,60%	100%	100%	?Muscular dystrophy, limb-girdle, autosomal recessive 21, 617232 Dowling-Degos disease 4, 615696
<i>POLR3A</i>	100%	99,70%	100%	100%	Leukodystrophy, hypomyelinating, 7, with or without oligodontia and/or hypogonadotropic hypogonadism, 607694 Wiedemann-Rautenstrauch syndrome, 264090
<i>POLR3B</i>	99,90%	98,60%	100%	100%	Leukodystrophy, hypomyelinating, 8, with or without oligodontia and/or hypogonadotropic hypogonadism, 614381
<i>POMGNT1</i>	100%	99,90%	100%	100%	Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 3, 613151 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 3, 613157 Retinitis pigmentosa 76, 617123 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 3, 253280
<i>POMGNT2</i>	100%	100%	100%	100%	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 8, 614830 Muscular dystrophy-dystroglycanopathy (limb-girdle) type C, 8, 618135
<i>POMK</i>	100%	100%	100%	100%	?Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 12, 616094 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 12, 615249
<i>POMT1</i>	99,30%	97,50%	100%	100%	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 1, 609308 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 1, 236670 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 1, 613155
<i>POMT2</i>	99,40%	96,40%	100%	100%	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 2, 613158 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 2, 613150 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 2, 613156
<i>PPCS</i>	99,80%	99,50%	100%	100%	Cardiomyopathy, dilated, 2C, 618189
<i>PPM1K</i>	100%	100%	100%	100%	?Maple syrup urine disease, mild variant, 615135
<i>PPOX</i>	99,70%	96,80%	100%	100%	Porphyria variegata, 176200
<i>PPT1</i>	90,30%	90,30%	82,50%	82,50%	Ceroid lipofuscinosis, neuronal, 1, 256730
<i>PRKAG2</i>	99,10%	97,50%	100%	100%	Glycogen storage disease of heart, lethal congenital, 261740 Wolff-Parkinson-White syndrome, 194200 Cardiomyopathy, hypertrophic 6, 600858
<i>PRKCSH</i>	99,80%	95,40%	100%	100%	Polycystic liver disease 1, 174050
<i>PRODH</i>	85,00%	80,60%	100%	100%	Hyperprolinemia, type I, 239500
<i>PRPS1</i>	100%	100%	100%	100%	Charcot-Marie-Tooth disease, X-linked recessive, 5, 311070 Phosphoribosylpyrophosphate synthetase superactivity, 300661 Deafness, X-linked 1, 304500 Arts syndrome, 301835 Gout, PRPS-related, 300661
<i>PSAP</i>	100%	100%	100%	100%	Gaucher disease, atypical, 610539 Krabbe disease, atypical, 611722

					Combined SAP deficiency, 611721 Metachromatic leukodystrophy due to SAP-b deficiency, 249900
<i>PSAT1</i>	95,30%	81,60%	100%	100%	Neu-Laxova syndrome 2, 616038 ?Phosphoserine aminotransferase deficiency, 610992
<i>PSPH</i>	100%	100%	100%	100%	Phosphoserine phosphatase deficiency, 614023
<i>PTEN</i>	99,50%	97,00%	100%	100%	Prostate cancer, somatic, 176807 Cowden syndrome 1, 158350 Lhermitte-Duclos syndrome, 158350 Macrocephaly/autism syndrome, 605309
<i>PTGIS</i>	98,20%	95,10%	100%	100%	Hypertension, essential, 145500
<i>PTPN11</i>	99,10%	93,70%	100%	100%	LEOPARD syndrome 1, 151100 Metachondromatosis, 156250 Noonan syndrome 1, 163950 Leukemia, juvenile myelomonocytic, somatic, 607785
<i>PTS</i>	99,90%	99,10%	100%	100%	Hyperphenylalaninemia, BH4-deficient, A, 261640
<i>PUS3</i>	100%	100%	100%	100%	Mental retardation, autosomal recessive 55, 617051
<i>PYCR1</i>	99,90%	97,70%	100%	100%	Cutis laxa, autosomal recessive, type IIIB, 614438 Cutis laxa, autosomal recessive, type IIB, 612940
<i>PYCR2</i>	100%	99,10%	100%	100%	Leukodystrophy, hypomyelinating, 10, 616420
<i>PYGL</i>	100%	100%	100%	100%	Glycogen storage disease VI, 232700
<i>PYGM</i>	100%	99,90%	100%	100%	McArdle disease, 232600
<i>QDPR</i>	100%	99,70%	100%	100%	Hyperphenylalaninemia, BH4-deficient, C, 261630
<i>RBCK1</i>	99,90%	98,20%	100%	100%	Polyglucosan body myopathy 1 with or without immunodeficiency, 615895
<i>RDH12</i>	100%	98,60%	100%	100%	Leber congenital amaurosis 13, 612712
<i>RDH5</i>	100%	99,90%	100%	100%	Fundus albipunctatus, 136880
<i>RFT1</i>	99,80%	99,60%	100%	100%	Congenital disorder of glycosylation, type In, 612015
<i>RINT1</i>	99,90%	98,60%	100%	100%	Infantile liver failure syndrome 3, 618641
<i>RPE65</i>	99,80%	97,80%	100%	100%	Leber congenital amaurosis 2, 204100 Retinitis pigmentosa 87 with choroidal involvement, 618697 Retinitis pigmentosa 20, 613794
<i>RPIA</i>	98,60%	94,90%	100%	100%	Ribose 5-phosphate isomerase deficiency, 608611
<i>SARDH</i>	93,70%	91,70%	91,40%	91,40%	No OMIM disease ID
<i>SAT1</i>	99,90%	98,50%	100%	99,90%	No OMIM disease ID
<i>SC5D</i>	100%	99,50%	100%	100%	Lathosterolosis, 607330
<i>SCARB2</i>	100%	99,80%	100%	100%	Epilepsy, progressive myoclonic 4, with or without renal failure, 254900
<i>SCP2</i>	100%	99,20%	100%	100%	?Leukoencephalopathy with dystonia and motor neuropathy, 613724
<i>SCYL1</i>	100%	100%	100%	100%	Spinocerebellar ataxia, autosomal recessive 21, 616719

<i>SEC23B</i>	99,90%	99,30%	100%	100%	?Cowden syndrome 7, 616858 Dyserythropoietic anemia, congenital, type II, 224100
<i>SELENBP1</i>	100%	100%	100%	100%	Extraoral halitosis due to MTO deficiency, 618148
<i>SEPSECS</i>	100%	100%	100%	100%	Pontocerebellar hypoplasia type 2D, 613811
<i>SERAC1</i>	99,90%	99,50%	100%	100%	3-methylglutaconic aciduria with deafness, encephalopathy, and Leigh-like syndrome, 614739
<i>SGSH</i>	94,40%	94,10%	100%	100%	Mucopolysaccharidosis type IIIA (Sanfilippo A), 252900
<i>SI</i>	99,20%	96,10%	100%	100%	Sucrase-isomaltase deficiency, congenital, 222900
<i>SLC10A7</i>	99,70%	98,00%	100%	100%	Short stature, amelogenesis imperfecta, and skeletal dysplasia with scoliosis, 618363
<i>SLC12A1</i>	100%	99,90%	100%	100%	Bartter syndrome, type 1, 601678
<i>SLC13A3</i>	99,40%	97,50%	100%	100%	Leukoencephalopathy, acute reversible, with increased urinary alpha-ketoglutarate, 618384
<i>SLC16A1</i>	100%	99,30%	100%	100%	Monocarboxylate transporter 1 deficiency, 616095 Hyperinsulinemic hypoglycemia, familial, 7, 610021 Erythrocyte lactate transporter defect, 245340
<i>SLC17A5</i>	99,60%	97,00%	100%	100%	Sialic acid storage disorder, infantile, 269920 Salla disease, 604369
<i>SLC18A2</i>	100%	99,70%	100%	100%	?Parkinsonism-dystonia, infantile, 2, 618049
<i>SLC22A12</i>	100%	99,80%	100%	100%	Hypouricemia, renal, 220150
<i>SLC22A5</i>	100%	100%	100%	100%	Carnitine deficiency, systemic primary, 212140
<i>SLC25A1</i>	95,80%	88,60%	99,50%	97,80%	Myasthenic syndrome, congenital, 23, presynaptic, 618197 Combined D-2- and L-2-hydroxyglutaric aciduria, 615182
<i>SLC25A13</i>	100%	99,70%	100%	100%	Citrullinemia, adult-onset type II, 603471 Citrullinemia, type II, neonatal-onset, 605814
<i>SLC25A15</i>	99,80%	98,10%	100%	100%	Hyperornithinemia-hyperammonemia-homocitrullinemia syndrome, 238970
<i>SLC25A19</i>	100%	98,50%	100%	100%	Thiamine metabolism dysfunction syndrome 4 (progressive polyneuropathy type), 613710 Microcephaly, Amish type, 607196
<i>SLC25A20</i>	100%	100%	100%	100%	Carnitine-acylcarnitine translocase deficiency, 212138
<i>SLC25A21</i>	100%	99,70%	100%	100%	?Mitochondrial DNA depletion syndrome 18, 618811
<i>SLC25A32</i>	100%	100%	100%	100%	?Exercise intolerance, riboflavin-responsive, 616839
<i>SLC25A38</i>	99,70%	97,10%	100%	100%	Anemia, sideroblastic, 2, pyridoxine-refractory, 205950
<i>SLC25A42</i>	96,50%	93,20%	100%	100%	metabolic crises, recurrent, with variable encephalomyopathic features and neurologic regression, 618416
<i>SLC28A1</i>	100%	98,80%	100%	100%	No OMIM disease ID
<i>SLC2A1</i>	92,80%	92,80%	100%	100%	Dystonia 9, 601042 GLUT1 deficiency syndrome 1, infantile onset, severe, 606777 Stomatin-deficient cryohydrocytosis with neurologic defects, 608885 GLUT1 deficiency syndrome 2, childhood onset, 612126
<i>SLC2A2</i>	100%	100%	100%	100%	Fanconi-Bickel syndrome, 227810

SLC2A9	99,80%	96,10%	100%	100%	Hypouricemia, renal, 2, 612076
SLC30A10	100%	100%	100%	100%	Hypermannesemia with dystonia 1, 613280
SLC33A1	99,90%	98,90%	100%	100%	Spastic paraplegia 42, autosomal dominant, 612539 Congenital cataracts, hearing loss, and neurodegeneration, 614482
SLC35A1	100%	99,70%	100%	100%	Congenital disorder of glycosylation, type If, 603585
SLC35A2	99,90%	98,40%	100%	100%	Congenital disorder of glycosylation, type Im, 300896
SLC35A3	80,70%	78,60%	81,10%	81,00%	?Arthrogyrosis, mental retardation, and seizures, 615553
SLC35C1	99,90%	98,70%	100%	100%	Congenital disorder of glycosylation, type Ic, 266265
SLC35D1	100%	97,70%	100%	100%	Schneckenbecken dysplasia, 269250
SLC37A4	100%	99,20%	100%	100%	Glycogen storage disease Ic, 232240 Glycogen storage disease Ib, 232220
SLC39A14	100%	99,40%	93,50%	93,50%	?Hyperostosis cranialis interna, 144755 Hypermannesemia with dystonia 2, 617013
SLC39A4	99,50%	95,50%	100%	100%	Acrodermatitis enteropathica, 201100
SLC39A8	100%	99,70%	100%	100%	Congenital disorder of glycosylation, type In, 616721
SLC3A1	100%	99,80%	96,60%	96,60%	Cystinuria, 220100
SLC44A1	98,20%	98,20%	100%	100%	No OMIM disease ID
SLC46A1	99,90%	98,50%	100%	100%	Folate malabsorption, hereditary, 229050
SLC52A1	100%	100%	100%	100%	Riboflavin deficiency, 615026
SLC52A2	100%	100%	100%	100%	Brown-Vialetto-Van Laere syndrome 2, 614707
SLC52A3	100%	100%	100%	100%	Brown-Vialetto-Van Laere syndrome 1, 211530 ?Fazio-Londe disease, 211500
SLC5A1	100%	100%	100%	100%	Glucose/galactose malabsorption, 606824
SLC5A2	100%	100%	100%	100%	Renal glucosuria, 233100
SLC6A19	100%	100%	100%	100%	Iminoglycinuria, digenic, 242600 HEARTnup disorder, 234500 Hyperglycinuria, 138500
SLC6A8	93,50%	81,60%	100%	99,80%	Cerebral creatine deficiency syndrome 1, 300352
SLC7A7	100%	99,90%	100%	100%	Lysinuric protein intolerance, 222700
SLC7A9	100%	99,90%	100%	100%	Cystinuria, 220100
SLCO1B1	99,20%	93,70%	100%	100%	Hyperbilirubinemia, Rotor type, digenic, 237450
SLCO1B3	98,80%	90,80%	100%	100%	Hyperbilirubinemia, Rotor type, digenic, 237450
SMPD1	100%	100%	100%	100%	Niemann-Pick disease, type A, 257200 Niemann-Pick disease, type B, 607616
SMS	91,50%	78,50%	100%	99,90%	Mental retardation, X-linked, Snyder-Robinson type, 309583
SNX14	99,60%	95,90%	100%	100%	Spinocerebellar ataxia, autosomal recessive 20, 616354

<i>SOD1</i>	100%	99,90%	100%	100%	Amyotrophic lateral sclerosis 1, 105400 Spastic tetraplegia and axial hypotonia, progressive, 618598
<i>SOD2</i>	100%	100%	100%	100%	No OMIM disease ID
<i>SPR</i>	99,80%	96,30%	100%	100%	Dystonia, dopa-responsive, due to sepiapterin reductase deficiency, 612716
<i>SPTLC1</i>	99,20%	95,40%	100%	100%	Neuropathy, hereditary sensory and autonomic, type IA, 162400
<i>SPTLC2</i>	100%	100%	100%	100%	Neuropathy, hereditary sensory and autonomic, type IC, 613640
<i>SQOR</i>	100%	97,80%	100%	100%	No OMIM disease ID
<i>SRD5A2</i>	99,90%	99,00%	100%	100%	Pseudovaginal perineoscrotal hypospadias, 264600
<i>SRD5A3</i>	99,90%	99,10%	100%	100%	Kahrizi syndrome, 612713 Congenital disorder of glycosylation, type Iq, 612379
<i>SSR4</i>	100%	99,70%	100%	100%	Congenital disorder of glycosylation, type Iy, 300934
<i>ST3GAL3</i>	100%	99,80%	100%	100%	Mental retardation, autosomal recessive 12, 611090 ?Epileptic encephalopathy, early infantile, 15, 615006
<i>ST3GAL5</i>	85,00%	84,20%	98,70%	98,40%	Salt and pepper developmental regression syndrome, 609056
<i>STAR</i>	100%	100%	100%	100%	Lipoid adrenal hyperplasia, 201710
<i>STS</i>	99,70%	98,10%	100%	99,90%	Ichthyosis, X-linked, 308100
<i>STT3A</i>	100%	100%	100%	100%	?Congenital disorder of glycosylation, type Iw, 615596
<i>STT3B</i>	100%	99,60%	100%	100%	?Congenital disorder of glycosylation, type Ix, 615597
<i>SUCLA2</i>	94,30%	86,60%	100%	100%	Mitochondrial DNA depletion syndrome 5 (encephalomyopathic with or without methylmalonic aciduria), 612073
<i>SUCLG1</i>	99,90%	99,80%	100%	100%	Mitochondrial DNA depletion syndrome 9 (encephalomyopathic type with methylmalonic aciduria), 245400
<i>SUCLG2</i>	96,70%	86,30%	100%	100%	No OMIM disease ID
<i>SUGCT</i>	99,90%	98,50%	100%	100%	Glutaric aciduria III, 231690
<i>SUMF1</i>	97,50%	90,80%	100%	100%	Multiple sulfatase deficiency, 272200
<i>SUOX</i>	100%	100%	100%	100%	Sulfite oxidase deficiency, 272300
<i>TBXAS1</i>	100%	100%	100%	100%	Ghosal hematodiaphyseal syndrome, 231095
<i>TCIRG1</i>	97,60%	90,10%	100%	100%	Osteopetrosis, autosomal recessive 1, 259700
<i>TCN2</i>	100%	100%	100%	100%	Transcobalamin II deficiency, 275350
<i>TECR</i>	100%	98,90%	100%	100%	Mental retardation, autosomal recessive 14, 614020
<i>TH</i>	99,30%	96,10%	100%	100%	Segawa syndrome, recessive, 605407
<i>TIMM50</i>	98,30%	94,40%	100%	100%	3-methylglutaconic aciduria, type IX, 617698
<i>TK2</i>	99,20%	96,30%	100%	100%	Mitochondrial DNA depletion syndrome 2 (myopathic type), 609560 ?Progressive external ophthalmoplegia with mitochondrial DNA deletions, autosomal recessive 3, 617069
<i>TKFC</i>	100%	99,50%	100%	100%	Triokinase and FMN cyclase deficiency syndrome, 618805
<i>TKT</i>	98,70%	97,80%	98,70%	98,70%	Short stature, developmental delay, and congenital heart defects, 617044
<i>TMEM165</i>	100%	100%	100%	100%	Congenital disorder of glycosylation, type IIk, 614727

<i>TMEM199</i>	100%	99,90%	100%	100%	Congenital disorder of glycosylation, type IIp, 616829
<i>TMEM5</i>	99,50%	96,80%	100%	99,90%	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 10, 615041
<i>TMEM70</i>	98,00%	93,90%	100%	100%	Mitochondrial complex V (ATP synthase) deficiency, nuclear type 2, 614052
<i>TMLHE</i>	99,50%	97,10%	100%	99,90%	No OMIM disease ID
<i>TPI1</i>	99,80%	97,50%	100%	100%	Hemolytic anemia due to triosephosphate isomerase deficiency, 615512
<i>TPK1</i>	99,80%	99,00%	100%	100%	Thiamine metabolism dysfunction syndrome 5 (episodic encephalopathy type), 614458
<i>TPMT</i>	99,10%	90,10%	100%	100%	No OMIM disease ID
<i>TPP1</i>	100%	100%	100%	100%	Spinocerebellar ataxia, autosomal recessive 7, 609270 Ceroid lipofuscinosis, neuronal, 2, 204500
<i>TRAK1</i>	100%	99,60%	100%	100%	Epileptic encephalopathy, early infantile, 68, 618201
<i>TRAPPC11</i>	100%	99,20%	100%	100%	Muscular dystrophy, limb-girdle, autosomal recessive 18, 615356
<i>TRAPPC2L</i>	100%	100%	100%	100%	Encephalopathy, progressive, early-onset, with episodic rhabdomyolysis, 618331
<i>TREH</i>	96,90%	92,10%	100%	100%	Trehalase deficiency, 612119
<i>TUSC3</i>	100%	99,50%	100%	100%	Mental retardation, autosomal recessive 7, 611093
<i>TYMP</i>	100%	97,00%	100%	100%	Mitochondrial DNA depletion syndrome 1 (MNGIE type), 603041
<i>TYMS</i>	99,90%	99,60%	100%	100%	No OMIM disease ID
<i>TYR</i>	100%	100%	100%	100%	Waardenburg syndrome/albinism, digenic, 103470 Albinism, oculocutaneous, type IB, 606952 Albinism, oculocutaneous, type IA, 203100
<i>TYRP1</i>	100%	99,80%	100%	100%	Albinism, oculocutaneous, type III, 203290
<i>UGT1A1</i>	100%	100%	100%	100%	Hyperbilirubinemia, familial transient neonatal, 237900 Crigler-Najjar syndrome, type I, 218800 Crigler-Najjar syndrome, type II, 606785
<i>UMPS</i>	100%	99,40%	97,00%	97,00%	Orotic aciduria, 258900
<i>UPB1</i>	100%	100%	100%	100%	Beta-ureidopropionase deficiency, 613161
<i>UROC1</i>	100%	100%	100%	100%	?Urocanase deficiency, 276880
<i>UROD</i>	98,90%	96,10%	100%	100%	Porphyria, hepatoerythropoietic, 176100 Porphyria cutanea tarda, 176100
<i>UROS</i>	100%	99,90%	100%	100%	Porphyria, congenital erythropoietic, 263700
<i>VMA21</i>	99,00%	94,60%	100%	98,60%	Myopathy, X-linked, with excessive autophagy, 310440
<i>VPS13B</i>	99,50%	98,20%	99,50%	99,40%	Cohen syndrome, 216550
<i>VPS33A</i>	97,30%	95,70%	95,80%	95,80%	Mucopolysaccharidosis-plus syndrome, 617303
<i>XDH</i>	100%	99,90%	100%	100%	Xanthinuria, type I, 278300
<i>XYLT1</i>	97,40%	89,60%	98,10%	94,80%	Desbuquois dysplasia 2, 615777
<i>XYLT2</i>	100%	98,30%	96,70%	96,70%	Spondyloocular syndrome, 605822

Gene symbols used follow HGNC guidelines: Gray KA, Yates B, Seal RL, Wright MW, Bruford EA. Nucleic Acids Res. 2015 Jan 43(Database issue):D1079-85.

Agilent V5 is the default chemistry, and used for all exome analyses apart from the (in-house) TURBO/RAPID WES route.

TWIST is the chemistry used for (in-house) TURBO/RAPID WES analysis.

Covered 10x describes the percentage of a gene's coding sequence that is covered at least 10x.

Covered 20x describes the percentage of a gene's coding sequence that is covered at least 20x.

Genes with coverage denoting NC are non-DNA coding genes.

non-DNA coding genes are covered, but as coverage statistics are based on DNA coding regions, statistics could not be generated.

OMIM release used for OMIM disease identifiers and descriptions : April 20th , 2020.

This list is accurate for panel version DG 2.18

Ad 1. "No OMIM Disease ID" signifies a gene without a current OMIM association Ad 2. OMIM phenotype descriptions between {} signify risk factors
