

Whole Exome Sequencing

Gene package Multiple congenital anomaly, version 8.1, 31-1-2020



Technical information

DNA was enriched using Agilent SureSelect DNA + SureSelect OneSeq 300kb CNV Backbone + Human All Exon V7 capture and paired-end sequenced on the Illumina platform (outsourced). The aim is to obtain 10 Giga base pairs per exome with a mapped fraction of 0.99. The average coverage of the exome is ~50x. Duplicate and non-unique reads are excluded. Data are demultiplexed with bcl2fastq Conversion Software from Illumina. Reads are mapped to the genome using the BWA-MEM algorithm (reference: <http://bio-bwa.sourceforge.net/>). Variant detection is performed by the Genome Analysis Toolkit HaplotypeCaller (reference: <http://www.broadinstitute.org/gatk/>). The detected variants are filtered and annotated with Cartagenia software and classified with Alamut Visual. It is not excluded that pathogenic mutations are being missed using this technology. At this moment, there is not enough information about the sensitivity of this technique with respect to the detection of deletions and duplications of more than 5 nucleotides and of somatic mosaic mutations (all types of sequence changes).



Dept. Clinical Genetics

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
A4GALT	[Blood group, P1Pk system, P(2) phenotype], 111400 [Blood group, P1Pk system, p phenotype], 111400	607922	146	100	100	99
AAAS	Achalasia-addisonianism-alacrimia syndrome, 231550	605378	102	100	100	100
AAGAB	Keratoderma, palmoplantar, punctate type IA, 148600	614888	59	100	100	97
AARS1	Charcot-Marie-Tooth disease, axonal, type 2N, 613287	601065	89	100	100	96
	Epileptic encephalopathy, early infantile, 29, 616339					
AARS2	Combined oxidative phosphorylation deficiency 8, 614096	612035	122	100	100	100
	Leukoencephalopathy, progressive, with ovarian failure, 615889					
AASS	Hyperlysinemia, 238700	605113	63	100	97	90
	Saccharopinuria, 268700					
ABAT	GABA-transaminase deficiency, 613163	137150	100	100	100	96
ABC A1	HDL deficiency, familial, 1, 604091	600046	92	100	100	97
	Tangier disease, 205400					
ABC A12	Ichthyosis, congenital 4A, 601277	607800	68	100	100	95
	Ichthyosis, congenital 4B (harlequin), 242500					
ABC A3	Surfactant metabolism dysfunction, pulmonary, 3, 610921	601615	107	100	100	98

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ABCA4	Cone-rod dystrophy 3, 604116 Fundus flavimaculatus, 248200 {Macular degeneration, age-related, 2}, 153800 Retinal dystrophy, early-onset severe, 248200 Retinitis pigmentosa 19, 601718 Stargardt disease 1, 248200	601691	94	100	100	97
ABCB11	Cholestasis, benign recurrent intrahepatic, 2, 605479 Cholestasis, progressive familial intrahepatic 2, 601847	603201	67	100	100	96
ABCB4	Cholestasis, intrahepatic, of pregnancy, 3, 614972 Cholestasis, progressive familial intrahepatic 3, 602347 Gallbladder disease 1, 600803	171060	69	100	99	92
ABCB6	[Blood group, Langereis system], 111600 Dyschromatosis universalis hereditaria 3, 615402 Microphthalmia, isolated, with coloboma 7, 614497 Pseudohyperkalemia, familial, 2, due to red cell leak, 609153	605452	110	100	100	100
ABCB7	Anemia, sideroblastic, with ataxia, 301310	300135	59	100	98	87
ABCC2	Dubin-Johnson syndrome, 237500	601107	76	100	99	95
ABCC6	Arterial calcification, generalized, of infancy, 2, 614473 Pseudoxanthoma elasticum, 264800 Pseudoxanthoma elasticum, forme fruste, 177850	603234	118	100	100	99
ABCC8	Diabetes mellitus, noninsulin-dependent, 125853 Diabetes mellitus, permanent neonatal, 606176 Diabetes mellitus, transient neonatal 2, 610374 Hyperinsulinemic hypoglycemia, familial, 1, 256450 Hypoglycemia of infancy, leucine-sensitive, 240800	600509	105	100	100	100
ABCC9	Atrial fibrillation, familial, 12, 614050 Cardiomyopathy, dilated, 10, 608569 Hypertrichotic osteochondrodysplasia, 239850	601439	65	100	100	95
ABCD1	Adrenoleukodystrophy, 300100 Adrenomyeloneuropathy, adult, 300100	300371	86	84	78	75
ABCD4	Methylmalonic aciduria and homocystinuria, cblJ type, 614857	603214	108	100	100	100
ABCG5	Sitosterolemia, 210250	605459	80	100	100	99
ABCG8	{Gallbladder disease 4}, 611465 Sitosterolemia, 210250	605460	149	100	98	95
ABHD12	Polyneuropathy, hearing loss, ataxia, retinitis pigmentosa, and cataract, 612674	613599	61	100	100	94

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ABHD5	Chanarin-Dorfman syndrome, 275630	604780	74	100	100	98
ABL1	Congenital heart defects and skeletal malformations syndrome, 617602 Leukemia, Philadelphia chromosome-positive, resistant to imatinib, 608232	189980	141	100	100	100
ACAD8	Isobutyryl-CoA dehydrogenase deficiency, 611283	604773	159	100	100	100
ACAD9	Mitochondrial complex I deficiency, nuclear type 20, 611126	611103	95	100	100	100
ACADM	Acyl-CoA dehydrogenase, medium chain, deficiency of, 201450	607008	66	100	100	96
ACADS	Acyl-CoA dehydrogenase, short-chain, deficiency of, 201470	606885	121	100	100	100
ACADSB	2-methylbutyrylglycinuria, 610006	600301	67	100	100	94
ACADVL	VLCAD deficiency, 201475	609575	114	100	100	98
ACAN	Short stature and advanced bone age, with or without early-onset osteoarthritis and/or osteochondritis dissecans, 165800 Spondyloepimetaphyseal dysplasia, aggrecan type, 612813 ?Spondyloepiphyseal dysplasia, Kimberley type, 608361	155760	150	93	93	92
ACAT1	Alpha-methylacetoadipic aciduria, 203750	607809	71	100	100	93
ACE	[Angiotensin I-converting enzyme, benign serum increase] {Microvascular complications of diabetes 3}, 612624 {Myocardial infarction, susceptibility to} Renal tubular dysgenesis, 267430 {SARS, progression of} {Stroke, hemorrhagic}, 614519	106180	108	100	100	100
ACO2	Infantile cerebellar-retinal degeneration, 614559 ?Optic atrophy 9, 616289	100850	146	100	98	94
ACOX1	Peroxisomal acyl-CoA oxidase deficiency, 264470	609751	103	100	100	99
ACP5	Spondyloenchondrodysplasia with immune dysregulation, 607944	171640	143	100	100	100
ACSF3	Combined malonic and methylmalonic aciduria, 614265	614245	149	100	100	100
ACSL4	Mental retardation 63, 300387	300157	54	100	98	88
ACSL6	Myelodysplastic syndrome Myelogenous leukemia, acute	604443	83	100	100	98
ACTA1	Myopathy, actin, congenital, with cores, 161800 Myopathy, actin, congenital, with excess of thin myofilaments, 161800 Myopathy, congenital, with fiber-type disproportion 1, 255310 ?Myopathy, scapulohumeroperoneal, 616852 Nemaline myopathy 3 or recessive, 161800	102610	138	100	100	100

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ACTA2	Aortic aneurysm, familial thoracic 6, 611788 Moyamoya disease 5, 614042 Multisystemic smooth muscle dysfunction syndrome, 613834	102620	167	100	100	100
ACTB	Baraitser-Winter syndrome 1, 243310 ?Dystonia, juvenile-onset, 607371	102630	192	100	100	100
ACTC1	Atrial septal defect 5, 612794 Cardiomyopathy, dilated, 1R, 613424 Cardiomyopathy, hypertrophic, 11, 612098 Left ventricular noncompaction 4, 613424	102540	194	100	100	100
ACTG1	Baraitser-Winter syndrome 2, 614583 Deafness 20/26, 604717	102560	188	100	100	100
ACTG2	Visceral myopathy, 155310	102545	95	100	100	100
ACTN1	Bleeding disorder, platelet-type, 15, 615193	102575	118	100	100	100
ACTN4	Glomerulosclerosis, focal segmental, 1, 603278	604638	119	100	100	98
ACVR1	Fibrodysplasia ossificans progressiva, 135100	102576	73	100	100	99
ACVR1B	Pancreatic cancer, somatic	601300	81	100	100	94
ACVR2B	Heterotaxy, visceral, 4, autosomal, 613751	602730	105	100	100	98
ACVRL1	Telangiectasia, hereditary hemorrhagic, type 2, 600376	601284	102	100	100	95
ACY1	Aminoacylase 1 deficiency, 609924	104620	109	100	100	100
ADA	Adenosine deaminase deficiency, partial, 102700, Somatic mosaicism Severe combined immunodeficiency due to ADA deficiency, 102700, Somatic mosaicism	608958	85	100	100	96
ADA2	?Sneddon syndrome, 182410 Vasculitis, autoinflammation, immunodeficiency, and hematologic defects syndrome, 615688	607575	96	100	100	99
ADAM10	{Alzheimer disease 18, susceptibility to}, 615590 Reticulate acropigmentation of Kitamura, 615537	602192	57	100	99	91
ADAM17	?Inflammatory skin and bowel disease, neonatal, 1, 614328	603639	67	100	99	94
ADAM9	Cone-rod dystrophy 9, 612775	602713	57	100	99	92
ADAMTS10	Weill-Marchesani syndrome 1, recessive, 277600	608990	112	100	100	100
ADAMTS13	Thrombotic thrombocytopenic purpura, familial, 274150	604134	92	97	97	95
ADAMTS17	Weill-Marchesani 4 syndrome, recessive, 613195	607511	100	97	95	92
ADAMTS18	Microcornea, myopic chorioretinal atrophy, and telecanthus, 615458	607512	79	100	100	98
ADAMTS2	Ehlers-Danlos syndrome, dermatosparaxis type, 225410	604539	111	100	100	99
ADAMTSL2	Geleophysic dysplasia 1, 231050	612277	51	44	41	39
ADAMTSL4	Ectopia lentis et pupillae, 225200 Ectopia lentis, isolated, 225100	610113	115	100	100	99

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ADAR	Aicardi-Goutieres syndrome 6, 615010 Dyschromatosis symmetrica hereditaria, 127400	146920	90	100	100	100
ADAT3	Mental retardation 36, 615286	615302	152	100	100	100
ADCY5	Dyskinesia, familial, with facial myokymia, 606703	600293	111	98	95	93
ADGRG1	Polymicrogyria, bilateral frontoparietal, 606854 Polymicrogyria, bilateral perisylvian, 615752	604110	120	100	100	100
ADGRV1	?Febrile seizures, familial, 4, 604352 Usher syndrome, type 2C, 605472 Usher syndrome, type 2C, GPR98/PDZD7 digenic, 605472	602851	80	100	100	97
ADIPOQ	Adiponectin deficiency, 612556	605441	87	100	100	100
ADK	Hypermethioninemia due to adenosine kinase deficiency, 614300	102750	53	100	93	80
ADRB2	{Asthma, nocturnal, susceptibility to}, 600807 Beta-2-adrenoreceptor agonist, reduced response to {Obesity, susceptibility to}, 601665	109690	117	100	100	100
ADSL	Adenylosuccinate deficiency, 103050	608222	89	100	100	96
AFF2	Mental retardation, FRADE type, 309548	300806	57	100	99	92
AFG3L2	Spastic ataxia 5, 614487 Spinocerebellar atrophy 28, 610246	604581	77	99	95	90
AGA	Aspartylglucosaminuria, 208400	613228	73	100	100	96
AGBL1	Corneal dystrophy, Fuchs endothelial, 8, 615523	615496	74	98	98	97
AGK	Cataract 38, 614691 Sengers syndrome, 212350	610345	57	100	100	93
AGL	Glycogen storage disease IIIa, 232400 Glycogen storage disease IIIB, 232400	610860	72	100	100	95
AGPAT2	Lipodystrophy, congenital generalized, type 1, 608594	603100	142	100	100	96
AGPS	Rhizomelic chondrodysplasia punctata, type 3, 600121	603051	61	100	99	90
AGRN	Myasthenic syndrome, congenital, 8, with pre- and postsynaptic defects, 615120	103320	140	100	98	95
AGT	{Hypertension, essential, susceptibility to}, 145500 {Preeclampsia, susceptibility to} Renal tubular dysgenesis, 267430	106150	129	100	100	100
AGTR1	{Hypertension, essential}, 145500 Renal tubular dysgenesis, 267430	106165	55	100	100	100
AGXT	Hyperoxaluria, primary, type 1, 259900	604285	133	100	100	100
AHCY	Hypermethioninemia with deficiency of S-adenosylhomocysteine hydrolase, 613752	180960	151	96	96	96
AHI1	Joubert syndrome 3, 608629	608894	67	100	98	90

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AHNAK2	No OMIM phenotype	608570	284	100	100	99
AICDA	Immunodeficiency with hyper-IgM, type 2, 605258	605257	134	100	100	95
AIFM1	Combined oxidative phosphorylation deficiency 6, 300816 Cowchock syndrome, 310490 Deafness 5, 300614	300169	55	100	96	82
AIMP1	Leukodystrophy, hypomyelinating, 3, 260600	603605	78	100	100	96
AIP	Pituitary adenoma 1, multiple types, 102200 Pituitary adenoma predisposition, 102200	605555	134	100	100	100
AIPL1	Cone-rod dystrophy, 604393 Leber congenital amaurosis 4, 604393 Retinitis pigmentosa, juvenile, 604393	604392	141	100	100	100
AIRE	Autoimmune polyendocrinopathy syndrome , type I, with or without reversible metaphyseal dysplasia, 240300	607358	100	100	100	100
AK1	Hemolytic anemia due to adenylate kinase deficiency, 612631	103000	124	100	100	100
AK2	Reticular dysgenesis, 267500	103020	68	100	100	97
AKAP9	?Long QT syndrome-11, 611820	604001	76	100	100	96
AKR1C2	46XY sex reversal 8, 614279	600450	71	91	91	85
AKR1D1	Bile acid synthesis defect, congenital, 2, 235555	604741	64	100	100	97
AKT1	Breast cancer, somatic, 114480 Colorectal cancer, somatic, 114500 Cowden syndrome 6, 615109 Ovarian cancer, somatic, 167000 Proteus syndrome, somatic, 176920 {Schizophrenia, susceptibility to}, 181500	164730	147	100	100	100
AKT2	Diabetes mellitus, type II, 125853 Hypoinsulinemic hypoglycemia with hemihypertrophy, 240900	164731	112	100	100	100
AKT3	Megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome 2, 615937	611223	70	100	99	90
ALAD	{Lead poisoning, susceptibility to}, 612740 Porphyria, acute hepatic, 612740	125270	104	100	100	100
ALAS2	Anemia, sideroblastic, 1, 300751 Protoporphyrinia, erythropoietic, 300752	301300	60	100	98	90
ALB	Analbuminemia, 616000 [Dysalbuminemic hyperthyroxinemia], 615999	103600	63	100	100	95

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ALDH18A1	Cutis laxa 3, 616603 Cutis laxa, type IIIA, 219150 Spastic paraplegia 9A, 601162 Spastic paraplegia 9B, 616586	138250	80	100	100	98
ALDH1A3	Microphthalmia, isolated 8, 615113	600463	88	100	100	97
ALDH2	Alcohol sensitivity, acute, 610251 {Esophageal cancer, alcohol-related, susceptibility to} {Hangover, susceptibility to}, 610251 {Sublingual nitroglycerin, susceptibility to poor response to}	100650	127	100	100	100
ALDH3A2	Sjogren-Larsson syndrome, 270200	609523	64	100	99	93
ALDH4A1	Hyperprolinemia, type II, 239510	606811	118	100	100	99
ALDH5A1	Succinic semialdehyde dehydrogenase deficiency, 271980	610045	66	100	97	91
ALDH6A1	Methylmalonate semialdehyde dehydrogenase deficiency, 614105	603178	98	100	100	98
ALDH7A1	Epilepsy, pyridoxine-dependent, 266100	107323	69	100	100	91
ALDOA	Glycogen storage disease XII, 611881	103850	144	100	100	100
ALDOB	Fructose intolerance, hereditary, 229600	612724	102	100	100	100
ALG1	Congenital disorder of glycosylation, type I κ , 608540	605907	61	91	79	73
ALG11	Congenital disorder of glycosylation, type I ρ , 613661	613666	73	100	100	99
ALG12	Congenital disorder of glycosylation, type I \gamm , 607143	607144	158	100	100	100
ALG13	?Congenital disorder of glycosylation, type I σ , 300884 Epileptic encephalopathy, early infantile, 36, 300884	300776	50	100	97	83
ALG2	?Congenital disorder of glycosylation, type I ι , 607906 Myasthenic syndrome, congenital, 14, with tubular aggregates, 616228	607905	81	100	100	99
ALG3	Congenital disorder of glycosylation, type I δ , 601110	608750	95	100	100	100
ALG6	Congenital disorder of glycosylation, type I ζ , 603147	604566	73	100	100	96
ALG8	Congenital disorder of glycosylation, type I η , 608104 Polycystic liver disease 3 with or without kidney cysts, 617874	608103	62	100	100	94
ALG9	Congenital disorder of glycosylation, type I ι , 608776 Gillessen-Kaesbach-Nishimura syndrome, 263210	606941	64	100	100	94
ALMS1	Alstrom syndrome, 203800	606844	100	100	100	99
ALOX12B	Ichthyosis, congenital 2, 242100	603741	113	100	100	99
ALOXE3	Ichthyosis, congenital 3, 606545	607206	111	100	100	98
ALPK1	No OMIM phenotype	607347	73	100	99	94

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ALPL	Hypophosphatasia, adult, 146300 Hypophosphatasia, childhood, 241510 Hypophosphatasia, infantile, 241500 Odontohypophosphatasia, 146300	171760	130	100	100	100
ALS2	Amyotrophic lateral sclerosis 2, juvenile, 205100 Primary lateral sclerosis, juvenile, 606353 Spastic paralysis, infantile onset ascending, 607225	606352	72	100	100	96
ALX1	?Frontonasal dysplasia 3, 613456	601527	78	100	100	100
ALX3	Frontonasal dysplasia 1, 136760	606014	116	98	91	86
ALX4	{Craniosynostosis 5, susceptibility to}, 615529 Frontonasal dysplasia 2, 613451 Parietal foramina 2, 609597	605420	139	100	100	100
AMACR	Alpha-methylacyl-CoA racemase deficiency, 614307 Bile acid synthesis defect, congenital, 4, 214950	604489	96	100	100	98
AMELX	Amelogenesis imperfecta, type 1E, 301200	300391	137	100	100	100
AMER1	Osteopathia striata with cranial sclerosis, 300373	300647	73	100	99	97
AMH	Persistent Mullerian duct syndrome, type I, 261550	600957	104	100	100	100
AMHR2	Persistent Mullerian duct syndrome, type II, 261550	600956	123	100	100	99
AMN	Megaloblastic anemia-1, Norwegian type, 261100	605799	89	100	93	83
AMPD1	Myopathy due to myoadenylate deaminase deficiency, 615511	102770	67	100	99	95
AMT	Glycine encephalopathy, 605899	238310	130	100	100	100
ANG	Amyotrophic lateral sclerosis 9, 611895	105850	126	100	100	100
ANGPTL3	Hypobetalipoproteinemia, familial, 2, 605019	604774	59	100	100	95
ANK1	Spherocytosis, type 1, 182900	612641	113	100	100	98
ANK2	Cardiac arrhythmia, ankyrin-B-related, 600919 Long QT syndrome 4, 600919	106410	74	100	100	98
ANKH	Chondrocalcinosis 2, 118600 Craniometaphyseal dysplasia, 123000	605145	91	100	100	99
ANKK1	No OMIM phenotype	608774	110	100	100	100
ANKRD11	KBG syndrome, 148050	611192	112	100	98	96
ANKRD26	Thrombocytopenia 2, 188000	610855	70	100	98	90
ANKS6	Nephronophthisis 16, 615382	615370	89	95	93	90
ANO10	Spinocerebellar ataxia 10, 613728	613726	57	100	99	88
ANO3	Dystonia 24, 615034	610110	61	93	92	87

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ANO5	Gnathodiaphyseal dysplasia, 166260 Miyoshi muscular dystrophy 3, 613319 Muscular dystrophy, limb-girdle 12, 611307	608662	77	100	100	96
ANO6	Scott syndrome, 262890	608663	70	100	100	97
ANOS1	Hypogonadotropic hypogonadism 1 with or without anosmia (Kallmann syndrome 1), 308700	300836	57	100	95	84
ANTXR1	GAPO syndrome, 230740 {?Hemangioma, capillary infantile, susceptibility to}, 602089	606410	60	99	97	92
ANTXR2	Hyaline fibromatosis syndrome, 228600	608041	65	100	100	96
AP1S1	MEDNIK syndrome, 609313	603531	84	100	100	96
AP1S2	Mental retardation syndromic 5, 304340	300629	40	100	87	62
AP2S1	Hypocalciuric hypercalcemia, type III, 600740	602242	100	100	100	100
AP3B1	Hermansky-Pudlak syndrome 2, 608233	603401	66	100	98	87
AP4B1	Spastic paraplegia 47, 614066	607245	81	100	100	100
AP4E1	Spastic paraplegia 51, 613744 Stuttering, familial persistent, 1, 184450	607244	64	100	100	96
AP4M1	Spastic paraplegia 50, 612936	602296	130	100	100	99
AP4S1	Spastic paraplegia 52, 614067	607243	44	100	99	87
AP5Z1	Spastic paraplegia 48, 613647	613653	111	100	100	99
APCDD1	Hypotrichosis 1, 605389	607479	125	100	100	100
APOA1	Amyloidosis, 3 or more types, 105200 ApoA-I and apoC-III deficiency, combined, 618463 Hypoalphalipoproteinemia, primary, 2, with or without corneal clouding, 618463	107680	130	100	100	100
APOA2	Apolipoprotein A-II deficiency {Hypercholesterolemia, familial, modifier of}, 143890	107670	90	100	100	100
APOA5	Hyperchylomicronemia, late-onset, 144650 {Hypertriglyceridemia, susceptibility to}, 145750	606368	179	100	100	100
APOB	Hypercholesterolemia, familial, 2, 144010 Hypobetalipoproteinemia, 615558	107730	159	100	100	100
APOC2	Hyperlipoproteinemia, type Ib, 207750	608083	91	100	100	100
APOC3	Apolipoprotein C-III deficiency, 614028	107720	136	100	100	100
APRT	Adenine phosphoribosyltransferase deficiency, 614723	102600	81	100	100	100
APTX	Ataxia, early-onset, with oculomotor apraxia and hypoalbuminemia, 208920	606350	76	100	99	92
AQP2	Diabetes insipidus, nephrogenic, 125800	107777	113	100	100	100
AQP5	Palmoplantar keratoderma, Bothnian type, 600231	600442	116	100	100	100

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AR	Androgen insensitivity, 300068 Androgen insensitivity, partial, with or without breast cancer, 312300 Hypospadias 1, 300633 {Prostate cancer, susceptibility to}, 176807 Spinal and bulbar muscular atrophy of Kennedy, 313200	313700	81	100	98	94
ARFGEF2	Periventricular heterotopia with microcephaly, 608097	605371	83	100	100	96
ARG1	Argininemia, 207800	608313	75	100	100	98
ARHGAP26	Leukemia, juvenile myelomonocytic, somatic, 607785	605370	88	100	100	99
ARHGAP31	Adams-Oliver syndrome 1, 100300	610911	103	100	100	99
ARHGEF10	?Slowed nerve conduction velocity, AD, 608236	608136	106	100	100	98
ARHGEF12	No OMIM phenotype	604763	67	100	99	94
ARHGEF6	No OMIM phenotype	300267	47	100	97	83
ARHGEF9	Epileptic encephalopathy, early infantile, 8, 300607	300429	50	100	99	85
ARID1A	Coffin-Siris syndrome 2, 614607	603024	121	100	99	97
ARID1B	Coffin-Siris syndrome 1, 135900	614556	102	100	100	98
ARL13B	Joubert syndrome 8, 612291	608922	63	100	100	94
ARL2BP	Retinitis pigmentosa with or without situs inversus, 615434	615407	57	100	100	96
ARL6	{Bardet-Biedl syndrome 1, modifier of}, 209900 Bardet-Biedl syndrome 3, 600151 ?Retinitis pigmentosa 55, 613575	608845	43	96	95	83
ARMC4	Ciliary dyskinesia, primary, 23, 615451	615408	67	98	96	92
ARNT	No OMIM phenotype	126110	53	100	99	91
ARSA	Metachromatic leukodystrophy, 250100	607574	136	100	100	100
ARSB	Mucopolysaccharidosis type VI (Maroteaux-Lamy), 253200	611542	73	100	100	98
ARSL	Chondrodysplasia punctata recessive, 302950	300180	81	100	99	92
ARX	Epileptic encephalopathy, early infantile, 1, 308350 Hydranencephaly with abnormal genitalia, 300215 Lissencephaly 2, 300215 Mental retardation 29 and others, 300419 Partington syndrome, 309510 Proud syndrome, 300004	300382	46	89	79	68
ASAHI	Farber lipogranulomatosis, 228000 Spinal muscular atrophy with progressive myoclonic epilepsy, 159950	613468	67	100	100	93
ASB10	Glaucoma 1, open angle, F, 603383	615054	98	100	100	98

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ASCC1	Barrett esophagus/esophageal adenocarcinoma, 614266 ?Spinal muscular atrophy with congenital bone fractures 2, 616867	614215	52	100	93	81
ASCL1	Central hypoventilation syndrome, congenital, 209880 Haddad syndrome, 209880	100790	259	100	100	100
ASL	Argininosuccinic aciduria, 207900	608310	117	100	100	99
GET3	No OMIM phenotype	601913	114	100	100	100
ASNS	Asparagine synthetase deficiency, 615574	108370	73	100	100	92
ASPA	Canavan disease, 271900	608034	55	100	99	91
ASPM	Microcephaly 5, primary, 608716	605481	71	100	100	97
ASPSCR1	Alveolar soft-part sarcoma, 606243	606236	106	100	98	95
ASS1	Citrullinemia, 215700	603470	115	100	98	90
ASXL1	Bohring-Opitz syndrome, 605039 Myelodysplastic syndrome, somatic, 614286	612990	107	100	98	98
ASXL3	Bainbridge-Ropers syndrome, 615485	615115	69	99	99	97
ATCAY	Ataxia, cerebellar, Cayman type, 601238	608179	116	100	100	96
ATIC	AICA-ribosiduria due to ATIC deficiency, 608688	601731	62	100	100	96
ATL1	Neuropathy, hereditary sensory, type ID, 613708 Spastic paraparesis 3A, 182600	606439	72	100	100	93
ATL3	Neuropathy, hereditary sensory, type IF, 615632	609369	56	100	99	93
ATM	Ataxia-telangiectasia, 208900 {Breast cancer, susceptibility to}, 114480 Lymphoma, B-cell non-Hodgkin, somatic Lymphoma, mantle cell, somatic T-cell prolymphocytic leukemia, somatic	607585	75	100	99	95
ATN1	Congenital hypotonia, epilepsy, developmental delay, and digital anomalies, 618494 Dentatorubral-pallidoluysian atrophy, 125370	607462	155	100	100	99
ATP13A2	Kufor-Rakeb syndrome, 606693 Spastic paraparesis 7B, 617225	610513	117	100	100	98
ATP1A2	Alternating hemiplegia of childhood 1, 104290 Migraine, familial basilar, 602481 Migraine, familial hemiplegic, 2, 602481	182340	130	100	100	100
ATP1A3	Alternating hemiplegia of childhood 2, 614820 CAPOS syndrome, 601338 Dystonia-12, 128235	182350	144	100	100	100
ATP2A1	Brody myopathy, 601003	108730	115	100	100	99

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ATP2A2	Acrokeratosis verruciformis, 101900 Darier disease, 124200	108740	99	100	100	100
ATP2C1	Hailey-Hailey disease, 169600	604384	61	100	98	93
ATP5F1C	No OMIM phenotype	108729	52	100	93	74
ATP5F1E	?Mitochondrial complex V (ATP synthase) deficiency, nuclear type 3, 614053	606153	88	100	100	100
ATP6V0A2	Cutis laxa, type IIA, 219200 Wrinkly skin syndrome, 278250	611716	81	100	100	97
ATP6V0A4	Renal tubular acidosis, distal, 602722	605239	77	100	100	96
ATP6V1B1	Renal tubular acidosis with deafness, 267300	192132	136	100	100	100
ATP7A	Menkes disease, 309400 Occipital horn syndrome, 304150 Spinal muscular atrophy, distal 3, 300489	300011	49	100	98	84
ATP7B	Wilson disease, 277900	606882	99	100	100	99
ATP8B1	Cholestasis, benign recurrent intrahepatic, 243300 Cholestasis, intrahepatic, of pregnancy, 1, 147480 Cholestasis, progressive familial intrahepatic 1, 211600	602397	70	100	100	95
ATPAF2	?Mitochondrial complex V (ATP synthase) deficiency, nuclear type 1, 604273	608918	73	100	100	100
ATR	?Cutaneous telangiectasia and cancer syndrome, familial, 614564 Seckel syndrome 1, 210600	601215	91	100	99	92
ATRX	Alpha-thalassemia myelodysplasia syndrome, somatic, 300448 Alpha-thalassemia/mental retardation syndrome, 301040 Mental retardation-hypotonic facies syndrome, 309580	300032	40	100	94	78
ATXN1	Spinocerebellar ataxia 1, 164400	601556	116	100	100	100
ATXN10	Spinocerebellar ataxia 10, 603516	611150	63	100	100	95
ATXN2	{Amyotrophic lateral sclerosis, susceptibility to, 13}, 183090 {Parkinson disease, late-onset, susceptibility to}, 168600 Spinocerebellar ataxia 2, 183090	601517	75	97	93	86
ATXN3	Machado-Joseph disease, 109150	607047	87	100	99	92
ATXN7	Spinocerebellar ataxia 7, 164500	607640	90	100	97	93
ATXN8OS	{Parkinson disease, susceptibility to}, 168600 Spinocerebellar ataxia 8, 608768	603680	No coverage data			
AUH	3-methylglutaconic aciduria, type I, 250950	600529	94	100	100	98
AURKC	Spermatogenic failure 5, 243060	603495	76	100	100	100
AVP	Diabetes insipidus, neurohypophyseal, 125700	192340	64	100	97	86

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AVPR2	Diabetes insipidus, nephrogenic, 304800 Nephrogenic syndrome of inappropriate antidiuresis, 300539	300538	85	100	100	99
AXIN1	?Caudal duplication anomaly, 607864 Hepatocellular carcinoma, somatic, 114550	603816	127	100	99	97
AXIN2	Colorectal cancer, somatic, 114500 Oligodontia-colorectal cancer syndrome, 608615	604025	108	100	100	98
B2M	?Amyloidosis, familial visceral, 105200 Immunodeficiency 43, 241600	109700	94	100	100	100
B3GALNT2	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies, type A, 11, 615181	610194	59	100	100	97
B3GALT6	Ehlers-Danlos syndrome, spondylodysplastic type, 2, 615349 Spondyloepimetaphyseal dysplasia with joint laxity, type 1, with or without fractures, 271640	615291	64	79	75	72
B3GAT3	Multiple joint dislocations, short stature, craniofacial dysmorphism, with or without congenital heart defects, 245600	606374	125	100	100	100
B3GLCT	Peters-plus syndrome, 261540	610308	67	100	99	85
B4GALNT1	Spastic paraparesis 26, 609195	601873	111	100	100	100
B4GALT1	Congenital disorder of glycosylation, type IIId, 607091	137060	95	100	100	100
B4GALT7	Ehlers-Danlos syndrome, spondylodysplastic type, 1, 130070	604327	126	100	100	98
B4GAT1	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 13, 615287	605517	154	100	100	100
B9D1	Joubert syndrome 27, 617120 ?Meckel syndrome 9, 614209	614144	90	100	100	100
B9D2	Joubert syndrome 34, 614175 ?Meckel syndrome 10, 614175	611951	82	100	100	100
BAAT	Hypercholanemia, familial, 607748	602938	73	100	100	87
BAG3	Cardiomyopathy, dilated, 1HH, 613881 Myopathy, myofibrillar, 6, 612954	603883	196	100	100	100
BANF1	Nestor-Guillermo progeria syndrome, 614008	603811	115	100	100	100
BAP1	Tumor predisposition syndrome, 614327	603089	121	100	100	100
BAX	Colorectal cancer, somatic, 114500 T-cell acute lymphoblastic leukemia, somatic, 613065	600040	93	100	100	93
BBS1	Bardet-Biedl syndrome 1, 209900	209901	126	100	100	100
BBS10	Bardet-Biedl syndrome 10, 615987	610148	66	100	100	100
BBS12	Bardet-Biedl syndrome 12, 615989	610683	59	100	100	99
BBS2	Bardet-Biedl syndrome 2, 615981 Retinitis pigmentosa 74, 616562	606151	77	100	100	97
BBS4	Bardet-Biedl syndrome 4, 615982	600374	76	100	100	95

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BBS5	Bardet-Biedl syndrome 5, 615983	603650	65	100	98	89
BBS7	Bardet-Biedl syndrome 7, 615984	607590	64	100	99	95
BBS9	Bardet-Biedl syndrome 9, 615986	607968	59	96	95	90
BCAP31	Deafness, dystonia, and cerebral hypomyelination, 300475	300398	83	100	100	97
BCHE	{Apnea, postanesthetic, susceptibility to, due to BCHE deficiency}, 617936 Butyrylcholinesterase deficiency, 617936	177400	71	100	100	99
BCKDHA	Maple syrup urine disease, type Ia, 248600	608348	160	100	100	100
BCKDHB	Maple syrup urine disease, type Ib, 248600	248611	64	100	99	94
BCKDK	Branched-chain ketoacid dehydrogenase kinase deficiency, 614923	614901	167	100	100	100
BCL10	?Immunodeficiency 37, 616098 Lymphoma, MALT, somatic, 137245 {Lymphoma, follicular, somatic}, 605027 {Male germ cell tumor, somatic}, 273300 {Mesothelioma, somatic}, 156240 {Sezary syndrome, somatic}	603517	66	100	100	96
BCL2	Leukemia/lymphoma, B-cell, 2	151430	152	100	100	100
BCL7A	B-cell non-Hodgkin lymphoma, high-grade	601406	110	100	100	100
BCO1	?Hypercarotenemia and vitamin A deficiency, 115300	605748	74	100	100	97
BCOR	Microphthalmia, syndromic 2, 300166	300485	78	100	99	94
BCR	Leukemia, acute lymphocytic, Philadelphia chromosome positive, somatic, 613065 Leukemia, chronic myeloid, Philadelphia chromosome positive, somatic, 608232	151410	144	100	100	100
BCS1L	Bjornstad syndrome, 262000 GRACILE syndrome, 603358 Leigh syndrome, 256000 Mitochondrial complex III deficiency, nuclear type 1, 124000	603647	170	100	100	100
BDNF	No OMIM phenotype	113505	104	100	100	100
BEAN1	Spinocerebellar ataxia 31, 117210	612051	159	100	100	100
BEST1	Bestrophinopathy, 611809 Macular dystrophy, vitelliform, 2, 153700 Microcornea, rod-cone dystrophy, cataract, and posterior staphyloma, 193220 Retinitis pigmentosa, concentric, 613194 Retinitis pigmentosa-50, 613194 Vitreoretinochoroidopathy, 193220	607854	94	100	99	98
BFSP1	Cataract 33, multiple types, 611391	603307	98	100	100	96
BFSP2	Cataract 12, multiple types, 611597	603212	73	100	99	92

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BGN	Meester-Loeys syndrome, 300989 Spondyloepimetaphyseal dysplasia, 300106	301870	87	100	100	98
BICD2	Spinal muscular atrophy, lower extremity-predominant, 2A, 615290 Spinal muscular atrophy, lower extremity-predominant, 2B, 618291	609797	133	100	100	99
BIN1	Centronuclear myopathy 2, 255200	601248	105	100	100	94
BLK	Maturity-onset diabetes of the young, type 11, 613375	191305	113	100	100	100
BLM	Bloom syndrome, 210900	604610	82	100	100	96
BLNK	?Agammaglobulinemia 4, 613502	604515	60	100	98	88
BLOC1S3	Hermansky-Pudlak syndrome 8, 614077	609762	104	100	100	100
BLOC1S6	?Hermansky-pudlak syndrome 9, 614171	604310	46	100	100	84
BLVRA	Hyperbiliverdinemia, 614156	109750	78	100	100	97
BMP1	Osteogenesis imperfecta, type XIII, 614856	112264	118	100	100	100
BMP15	Ovarian dysgenesis 2, 300510 Premature ovarian failure 4, 300510	300247	56	100	100	96
BMP2	Brachydactyly, type A2, 112600 {HFE hemochromatosis, modifier of}, 235200 Short stature, facial dysmorphism, and skeletal anomalies with or without cardiac anomalies, 617877	112261	123	100	100	99
BMP4	Microphthalmia, syndromic 6, 607932 Orofacial cleft 11, 600625	112262	103	100	96	93
BMPER	Diaphanospondylodysostosis, 608022	608699	99	100	100	97
BMPR1A	Juvenile polyposis syndrome, infantile form, 174900 Polyposis syndrome, hereditary mixed, 2, 610069 Polyposis, juvenile intestinal, 174900	601299	76	100	100	100
BMPR1B	Acromesomelic dysplasia, Demirhan type, 609441 Brachydactyly, type A1, D, 616849 Brachydactyly, type A2, 112600	603248	72	100	100	100
BMPR2	Pulmonary hypertension, familial primary, 1, with or without HHT, 178600 Pulmonary hypertension, primary, fenfluramine or dexfenfluramine-associated, 178600 Pulmonary venoocclusive disease 1, 265450	600799	65	100	100	98
BOLA3	Multiple mitochondrial dysfunctions syndrome 2 with hyperglycinemia, 614299	613183	69	100	100	91
PGM	Erythrocytosis, familial, 8, 222800	613896	71	100	100	99

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BRAF	Adenocarcinoma of lung, somatic, 211980 Cardiofaciocutaneous syndrome, 115150 Colorectal cancer, somatic LEOPARD syndrome 3, 613707 Melanoma, malignant, somatic Nonsmall cell lung cancer, somatic Noonan syndrome 7, 613706	164757	68	100	100	94
BRAT1	Neurodevelopmental disorder with cerebellar atrophy and with or without seizures, 618056 Rigidity and multifocal seizure syndrome, lethal neonatal, 614498	614506	125	100	100	100
BRIP1	{Breast cancer, early-onset, susceptibility to}, 114480 Fanconi anemia, complementation group J, 609054	605882	55	100	99	91
BRWD3	Mental retardation 93, 300659	300553	45	100	95	81
BSCL2	Encephalopathy, progressive, with or without lipodystrophy, 615924 Lipodystrophy, congenital generalized, type 2, 269700 Neuropathy, distal hereditary motor, type VA, 600794 Silver spastic paraplegia syndrome, 270685	606158	101	100	100	100
BSND	Bartter syndrome, type 4a, 602522 Sensorineural deafness with mild renal dysfunction, 602522	606412	115	100	100	100
BTD	Biotinidase deficiency, 253260	609019	89	100	100	100
BTK	Agammaglobulinemia 1, 300755 Isolated growth hormone deficiency, type III, with agammaglobulinemia, 307200	300300	50	100	98	83
BUB1	Colorectal cancer with chromosomal instability, somatic	602452	79	100	100	97
BUB1B	Colorectal cancer, somatic, 114500 Mosaic variegated aneuploidy syndrome 1, 257300 [Premature chromatid separation trait], 176430	602860	70	100	99	94
C12orf57	Temptany syndrome, 218340	615140	141	100	100	100
C12orf65	Combined oxidative phosphorylation deficiency 7, 613559 Spastic paraplegia 55, 615035	613541	78	100	100	100
C15orf41	Dyserythropoietic anemia, congenital, type Ib, 615631	615626	63	100	100	99
C19orf12	Neurodegeneration with brain iron accumulation 4, 614298 ?Spastic paraplegia 43, 615043	614297	186	100	100	100
C1GALT1C1	Tn polyagglutination syndrome, somatic, 300622	300611	47	100	100	95
C1QA	C1q deficiency, 613652	120550	155	100	100	100
C1QB	C1q deficiency, 613652	120570	121	100	100	100
C1QC	C1q deficiency, 613652	120575	150	100	100	100

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C1QTNF5	Retinal degeneration, late-onset, 605670	608752	102	100	100	93
C1S	C1s deficiency, 613783 Ehlers-Danlos syndrome, periodontal type, 2, 617174	120580	96	100	100	98
C2	C2 deficiency, 217000 {Macular degeneration, age-related, 14, reduced risk of}, 615489	613927	106	100	100	100
C3	C3 deficiency, 613779 {Hemolytic uremic syndrome, atypical, susceptibility to, 5}, 612925 {Macular degeneration, age-related, 9}, 611378	120700	127	100	100	99
C4A	[Blood group, Rodgers], 614374 C4a deficiency, 614380	120810	38	22	20	19
C4B	C4B deficiency, 614379	120820	36	24	22	21
C5	C5 deficiency, 609536 [Eculizumab, poor response to], 615749	120900	72	100	98	94
C6	C6 deficiency, 612446 Combined C6/C7 deficiency	217050	65	100	99	94
C7	C7 deficiency, 610102	217070	79	100	99	96
C8A	C8 deficiency, type I, 613790	120950	69	100	100	98
C8B	C8 deficiency, type II, 613789	120960	100	100	100	98
C8orf37	Bardet-Biedl syndrome 21, 617406 Cone-rod dystrophy 16, 614500 Retinitis pigmentosa 64, 614500	614477	78	100	100	96
C9	C9 deficiency, 613825 {Macular degeneration, age-related, 15, susceptibility to}, 615591	120940	66	100	100	97
C9orf72	Frontotemporal dementia and/or amyotrophic lateral sclerosis 1, 105550	614260	56	100	100	96
CA12	Hyperchlorhidrosis, isolated, 143860	603263	73	100	100	99
CA2	Osteopetrosis 3, with renal tubular acidosis, 259730	611492	99	100	100	100
CA4	Retinitis pigmentosa 17, 600852	114760	100	100	100	100
CA5A	Hyperammonemia due to carbonic anhydrase VA deficiency, 615751	114761	175	100	100	100
CA8	Cerebellar ataxia and mental retardation with or without quadrupedal locomotion 3, 613227	114815	79	100	100	94
CABP2	Deafness 93, 614899	607314	94	100	100	96
CABP4	Cone-rod synaptic disorder, congenital nonprogressive, 610427	608965	149	100	100	100

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CACNA1A	Epileptic encephalopathy, early infantile, 42, 617106 Episodic ataxia, type 2, 108500 Migraine, familial hemiplegic, 1, 141500 Migraine, familial hemiplegic, 1, with progressive cerebellar ataxia, 141500 Spinocerebellar ataxia 6, 183086	601011	87	100	98	93
CACNA1C	Brugada syndrome 3, 611875 Long QT syndrome 8, 618447 Timothy syndrome, 601005	114205	123	100	100	99
CACNA1D	Primary aldosteronism, seizures, and neurologic abnormalities, 615474 Sinoatrial node dysfunction and deafness, 614896	114206	102	100	100	98
CACNA1F	Aland Island eye disease, 300600 Cone-rod dystrophy, 3, 300476 Night blindness, congenital stationary (incomplete), 2A, 300071	300110	76	100	100	96
CACNA1G	Spinocerebellar ataxia 42, 616795 Spinocerebellar ataxia 42, early-onset, severe, with neurodevelopmental deficits, 618087	604065	146	100	100	100
CACNA1S	Hypokalemic periodic paralysis, type 1, 170400 {Malignant hyperthermia susceptibility 5}, 601887 {Thyrotoxic periodic paralysis, susceptibility to, 1}, 188580	114208	119	100	100	100
CACNA2D4	Retinal cone dystrophy 4, 610478	608171	86	100	100	98
CACNB2	Brugada syndrome 4, 611876	600003	100	100	100	100
CACNB4	{Epilepsy, idiopathic generalized, susceptibility to, 9}, 607682 {Epilepsy, juvenile myoclonic, susceptibility to, 6}, 607682 Episodic ataxia, type 5, 613855	601949	57	100	100	94
CACNG2	?Mental retardation 10, 614256	602911	107	100	100	99
CALCOCO1	No OMIM phenotype	No ID	85	100	100	98
CALM1	Long QT syndrome 14, 616247 Ventricular tachycardia, catecholaminergic polymorphic, 4, 614916	114180	64	100	100	100
CALR	Myelofibrosis, somatic, 254450 Thrombocythemia, somatic, 187950	109091	136	100	100	100
CALR3	No OMIM phenotype	611414	67	100	100	97
CAMTA1	Cerebellar ataxia, nonprogressive, with mental retardation, 614756	611501	161	100	99	97
CANT1	Desbuquois dysplasia 1, 251450 Epiphyseal dysplasia, multiple, 7, 617719	613165	128	100	100	100
CAPN3	Muscular dystrophy, limb-girdle 4, 618129 Muscular dystrophy, limb-girdle 1, 253600	114240	91	100	100	99

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CAPN5	Vitreoretinopathy, neovascular inflammatory, 193235	602537	143	100	100	99
CARD11	B-cell expansion with NFKB and T-cell anergy, 616452 Immunodeficiency 11A, 615206 Immunodeficiency 11B with atopic dermatitis, 617638	607210	108	100	100	99
CARD14	Pityriasis rubra pilaris, 173200 Psoriasis 2, 602723	607211	112	100	99	99
CARD9	Candidiasis, familial, 2, 212050	607212	122	100	100	99
CARS1	No OMIM phenotype	123859	83	100	98	92
CASK	FG syndrome 4, 300422 Mental retardation and microcephaly with pontine and cerebellar hypoplasia, 300749 Mental retardation, with or without nystagmus, 300422	300172	51	100	96	80
CASP10	Autoimmune lymphoproliferative syndrome, type II, 603909 Gastric cancer, somatic, 613659 Lymphoma, non-Hodgkin, somatic, 605027	601762	75	100	100	97
CASP8	?Autoimmune lymphoproliferative syndrome, type IIB, 607271 {Breast cancer, protection against}, 114480 Hepatocellular carcinoma, somatic, 114550 {Lung cancer, protection against}, 211980	601763	87	100	100	100
CASQ2	Ventricular tachycardia, catecholaminergic polymorphic, 2, 611938	114251	65	100	100	97
CASR	{Epilepsy idiopathic generalized, susceptibility to, 8}, 612899 Hyperparathyroidism, neonatal, 239200 Hypocalcemia, 601198 Hypocalcemia, with Bartter syndrome, 601198 Hypocalciuric hypercalcemia, type I, 145980	601199	146	100	100	100
CAT	Acatalasemia, 614097	115500	70	100	100	93
CATSPER1	Spermatogenic failure 7, 612997	606389	109	100	100	99
CAV1	?Lipodystrophy, congenital generalized, type 3, 612526 Lipodystrophy, familial partial, type 7, 606721 Pulmonary hypertension, primary, 3, 615343	601047	95	100	100	100
CAV3	Cardiomyopathy, familial hypertrophic, 192600 Creatine phosphokinase, elevated serum, 123320 Long QT syndrome 9, 611818 Myopathy, distal, Tateyama type, 614321 Rippling muscle disease 2, 606072	601253	150	100	100	100
CAVIN1	Lipodystrophy, congenital generalized, type 4, 613327	603198	172	100	100	100

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CBL	?Juvenile myelomonocytic leukemia, 607785 Noonan syndrome-like disorder with or without juvenile myelomonocytic leukemia, 613563	165360	78	100	100	100
CBLIF	Intrinsic factor deficiency, 261000	609342	No coverage data			
CBS	Homocystinuria, B6-responsive and nonresponsive types, 236200 Thrombosis, hyperhomocysteinemic, 236200	613381	127	100	100	100
CBX2	?46XY sex reversal 5, 613080	602770	176	100	100	100
CC2D1A	Mental retardation 3, 608443	610055	132	100	100	100
CC2D2A	COACH syndrome, 216360 Joubert syndrome 9, 612285 Meckel syndrome 6, 612284	612013	66	100	100	95
CCBE1	Hennekam lymphangiectasia-lymphedema syndrome 1, 235510	612753	87	100	99	95
CCDC103	Ciliary dyskinesia, primary, 17, 614679	614677	83	100	100	100
CCDC114	Ciliary dyskinesia, primary, 20, 615067	615038	125	100	100	100
CCDC39	Ciliary dyskinesia, primary, 14, 613807	613798	80	100	99	94
CCDC40	Ciliary dyskinesia, primary, 15, 613808	613799	100	100	100	100
CCDC50	?Deafness 44, 607453	611051	73	100	100	98
CCDC65	Ciliary dyskinesia, primary, 27, 615504	611088	57	100	100	94
CCDC78	?Centronuclear myopathy 4, 614807	614666	123	100	100	100
CCDC8	3-M syndrome 3, 614205	614145	194	100	100	100
CCDC88C	Hydrocephalus, congenital, 1, 236600 ?Spinocerebellar ataxia 40, 616053	611204	104	100	100	97
CCM2	Cerebral cavernous malformations-2, 603284	607929	144	100	100	96
CCN6	Arthropathy, progressive pseudorheumatoid, of childhood, 208230 Spondyloepiphyseal dysplasia tarda with progressive arthropathy, 208230	603400	83	100	100	100
CCNQ	STAR syndrome, 300707	300708	62	81	81	81
CCT5	Neuropathy, hereditary sensory, with spastic paraplegia, 256840	610150	85	100	100	91
CD151	[Blood group, Raph], 179620 Nephropathy with pretibial epidermolysis bullosa and deafness, 609057	602243	89	100	100	100
CD19	Immunodeficiency, common variable, 3, 613493	107265	107	100	100	99
CD247	?Immunodeficiency 25, 610163	186780	74	100	100	100
CD27	Lymphoproliferative syndrome 2, 615122	186711	95	100	100	100
CD2AP	Glomerulosclerosis, focal segmental, 3, 607832	604241	68	100	100	96
CD320	Methylmalonic aciduria, transient, due to transcobalamin receptor defect, 613646	606475	95	100	100	100

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CD36	{Coronary heart disease, susceptibility to}, 7, 610938 [Macrothrombocytopenia] {Malaria, cerebral, reduced risk of}, 611162 {Malaria, cerebral, susceptibility to}, 611162 Platelet glycoprotein IV deficiency, 608404	173510	67	100	99	93
CD3D	Immunodeficiency 19, 615617	186790	100	100	100	99
CD3E	Immunodeficiency 18, 615615 Immunodeficiency 18, SCID variant, 615615	186830	113	100	100	100
CD3G	Immunodeficiency 17, CD3 gamma deficient, 615607	186740	76	100	100	99
CD4	OKT4 epitope deficiency, 613949	186940	103	100	100	100
CD40	Immunodeficiency with hyper-IgM, type 3, 606843	109535	104	100	100	100
CD40LG	Immunodeficiency, with hyper-IgM, 308230	300386	67	100	100	97
CD59	Hemolytic anemia, CD59-mediated, with or without immune-mediated polyneuropathy, 612300	107271	86	100	100	100
CD79A	Agammaglobulinemia 3, 613501	112205	83	100	91	85
CD79B	Agammaglobulinemia 6, 612692	147245	146	100	100	100
CD81	Immunodeficiency, common variable, 6, 613496	186845	146	97	90	90
CD8A	CD8 deficiency, familial, 608957	186910	100	100	100	97
CD96	C syndrome, 211750	606037	74	100	99	96
CDAN1	Dyserythropoietic anemia, congenital, type Ia, 224120	607465	95	100	100	98
CDC6	?Meier-Gorlin syndrome 5, 613805	602627	65	100	100	98
CDC73	Hyperparathyroidism, familial primary, 145000 Hyperparathyroidism-jaw tumor syndrome, 145001 Parathyroid adenoma with cystic changes, 145001 Parathyroid carcinoma, 608266	607393	75	100	100	97
CDH1	Blepharocheilodontic syndrome 1, 119580 {Breast cancer, lobular}, 114480 Endometrial carcinoma, somatic, 608089 Gastric cancer, hereditary diffuse, with or without cleft lip and/or palate, 137215 Ovarian cancer, somatic, 167000 {Prostate cancer, susceptibility to}, 176807	192090	92	100	100	99
CDH15	Mental retardation 3, 612580	114019	125	100	100	100
CDH23	Deafness 12, 601386 {Pituitary adenoma 5, multiple types}, 617540 Usher syndrome, type 1D, 601067 Usher syndrome, type 1D/F digenic, 601067	605516	135	100	100	100

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CDH3	Ectodermal dysplasia, ectrodactyly, and macular dystrophy, 225280 Hypotrichosis, congenital, with juvenile macular dystrophy, 601553	114021	124	100	100	100
CDHR1	Cone-rod dystrophy 15, 613660 Retinitis pigmentosa 65, 613660	609502	116	100	100	98
CDK5RAP2	Microcephaly 3, primary, 604804	608201	69	100	99	94
CDKL5	Epileptic encephalopathy, early infantile, 2, 300672	300203	58	100	97	87
CDKN1B	Multiple endocrine neoplasia, type IV, 610755	600778	163	100	100	100
CDKN1C	Beckwith-Wiedemann syndrome, 130650 IMAGE syndrome, 614732	600856	74	90	83	76
CDKN2A	Melanoma and neural system tumor syndrome, 155755 {Melanoma, cutaneous malignant, 2}, 155601 Orolaryngeal cancer, multiple Pancreatic cancer/melanoma syndrome, 606719	600160	139	100	100	100
CDON	Holoprosencephaly 11, 614226	608707	89	100	100	99
CDSN	Hypotrichosis 2, 146520 Peeling skin syndrome 1, 270300	602593	103	100	100	100
CDT1	Meier-Gorlin syndrome 4, 613804	605525	125	100	98	95
CEACAM16	Deafness 4B, 614614 Deafness 113, 618410	614591	94	100	100	100
CEBPA	?Leukemia, acute myeloid, 601626 Leukemia, acute myeloid, somatic, 601626	116897	130	100	93	75
CEBPE	Specific granule deficiency, 245480	600749	57	100	100	99
CEL	Maturity-onset diabetes of the young, type VIII, 609812	114840	151	94	90	87
CENPJ	Microcephaly 6, primary, 608393 ?Seckel syndrome 4, 613676	609279	74	100	100	98
CEP135	Microcephaly 8, primary, 614673	611423	83	100	99	91
CEP152	Microcephaly 9, primary, 614852 Seckel syndrome 5, 613823	613529	67	100	98	93
CEP164	Nephronophthisis 15, 614845	614848	93	100	100	98
CEP19	Morbid obesity and spermatogenic failure, 615703	615586	77	100	100	100
CEP290	?Bardet-Biedl syndrome 14, 615991 Joubert syndrome 5, 610188 Leber congenital amaurosis 10, 611755 Meckel syndrome 4, 611134 Senior-Loken syndrome 6, 610189	610142	71	100	98	89

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CEP41	Joubert syndrome 15, 614464	610523	69	100	100	94
CEP57	Mosaic variegated aneuploidy syndrome 2, 614114	607951	62	100	100	95
CERKL	Retinitis pigmentosa 26, 608380	608381	73	100	99	94
CERS3	Ichthyosis, congenital 9, 615023	615276	65	100	100	94
CES1	Drug metabolism, altered, CES1-related, 618057	114835	126	77	77	76
CETP	[High density lipoprotein cholesterol level QTL 10], 143470 Hyperalphalipoproteinemia, 143470	118470	95	100	100	100
CFAP298	Ciliary dyskinesia, primary, 26, 615500	615494	75	100	100	96
CFAP53	Heterotaxy, visceral, 6, 614779	614759	98	100	100	98
CFC1	Heterotaxy, visceral, 2, autosomal, 605376	605194	70	41	34	28
CFD	Complement factor D deficiency, 613912	134350	99	100	98	90
CFH	Basal laminar drusen, 126700 Complement factor H deficiency, 609814 {Hemolytic uremic syndrome, atypical, susceptibility to, 1}, 235400 {Macular degeneration, age-related, 4}, 610698	134370	78	100	100	96
CFHR5	Nephropathy due to CFHR5 deficiency, 614809	608593	75	100	100	97
CFI	Complement factor I deficiency, 610984 {Hemolytic uremic syndrome, atypical, susceptibility to, 3}, 612923 {Macular degeneration, age-related, 13, susceptibility to}, 615439	217030	67	100	99	93
CFL2	Nemaline myopathy 7, 610687	601443	45	100	97	77
CFP	Properdin deficiency, 312060	300383	68	100	95	86
CFTR	{Bronchiectasis with or without elevated sweat chloride 1, modifier of}, 211400 Congenital bilateral absence of vas deferens, 277180 Cystic fibrosis, 219700 {Hypertrypsinemia, neonatal} {Pancreatitis, hereditary}, 167800 Sweat chloride elevation without CF	602421	93	100	100	97
CHAT	Myasthenic syndrome, congenital, 6, presynaptic, 254210	118490	130	99	92	88
CHD2	Epileptic encephalopathy, childhood-onset, 615369	602119	72	100	100	96
CHD7	CHARGE syndrome, 214800 Hypogonadotropic hypogonadism 5 with or without anosmia, 612370	608892	87	100	100	97
CHKB	Muscular dystrophy, congenital, megaconial type, 602541	612395	111	100	100	100
CHM	Choroideremia, 303100	300390	50	100	94	80
CHMP1A	Pontocerebellar hypoplasia, type 8, 614961	164010	102	100	100	100

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CHMP2B	Amyotrophic lateral sclerosis 17, 614696 Dementia, familial, nonspecific, 600795	609512	83	100	100	97
CHMP4B	Cataract 31, multiple types, 605387	610897	178	100	100	100
CHN1	Duane retraction syndrome 2, 604356	118423	62	100	100	92
CHRDL1	Megalocornea 1, 309300	300350	48	100	100	92
CHRM3	?Prune belly syndrome, 100100	118494	110	100	100	100
CHRNA1	Multiple pterygium syndrome, lethal type, 253290 Myasthenic syndrome, congenital, 1A, slow-channel, 601462 Myasthenic syndrome, congenital, 1B, fast-channel, 608930	100690	86	100	100	98
CHRNA2	Epilepsy, nocturnal frontal lobe, type 4, 610353	118502	170	100	100	100
CHRNA4	Epilepsy, nocturnal frontal lobe, 1, 600513 {Nicotine addiction, susceptibility to}, 188890	118504	138	100	100	96
CHRNBT1	Myasthenic syndrome, congenital, 2A, slow-channel, 616313 ?Myasthenic syndrome, congenital, 2C, associated with acetylcholine receptor deficiency, 616314	100710	124	100	100	100
CHRNBT2	Epilepsy, nocturnal frontal lobe, 3, 605375	118507	144	100	100	100
CHRND	Multiple pterygium syndrome, lethal type, 253290 ?Myasthenic syndrome, congenital, 3A, slow-channel, 616321 Myasthenic syndrome, congenital, 3B, fast-channel, 616322 ?Myasthenic syndrome, congenital, 3C, associated with acetylcholine receptor deficiency, 616323	100720	135	100	100	100
CHRNE	Myasthenic syndrome, congenital, 4A, slow-channel, 605809 Myasthenic syndrome, congenital, 4B, fast-channel, 616324 Myasthenic syndrome, congenital, 4C, associated with acetylcholine receptor deficiency, 608931	100725	138	100	100	100
CHRNG	Escobar syndrome, 265000 Multiple pterygium syndrome, lethal type, 253290	100730	116	100	100	99
CHST14	Ehlers-Danlos syndrome, musculocontractural type 1, 601776	608429	133	100	98	97
CHST3	Spondyloepiphyseal dysplasia with congenital joint dislocations, 143095	603799	129	100	100	100
CHST6	Macular corneal dystrophy, 217800	605294	246	100	100	100
CHSY1	Temptamy preaxial brachydactyly syndrome, 605282	608183	79	98	95	92
CHUK	Cocoon syndrome, 613630	600664	66	100	99	92
CIB2	Deafness 48, 609439 Usher syndrome, type IJ, 614869	605564	195	100	100	100
CIITA	Bare lymphocyte syndrome, type II, complementation group A, 209920 {Rheumatoid arthritis, susceptibility to}, 180300	600005	117	100	100	99
CISD2	Wolfram syndrome 2, 604928	611507	137	100	100	100

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CITED2	Atrial septal defect 8, 614433 Ventricular septal defect 2, 614431	602937	93	100	100	100
CLCF1	Cold-induced sweating syndrome 2, 610313	607672	58	100	99	93
CLCN1	Myotonia congenita, dominant, 160800 Myotonia congenita, recessive, 255700 Myotonia levior, recessive	118425	101	100	100	100
CLCN2	{Epilepsy, idiopathic generalized, susceptibility to, 11}, 607628 {Epilepsy, juvenile absence, susceptibility to, 2}, 607628 {Epilepsy, juvenile myoclonic, susceptibility to, 8}, 607628 Hyperaldosteronism, familial, type II, 605635 Leukoencephalopathy with ataxia, 615651	600570	100	100	100	100
CLCN5	Dent disease, 300009 Hypophosphatemic rickets, 300554 Nephrolithiasis, type I, 310468 Proteinuria, low molecular weight, with hypercalciuric nephrocalcinosis, 308990	300008	59	100	100	96
CLCN7	Hypopigmentation, organomegaly, and delayed myelination and development, 618541 Osteopetrosis 2, 166600 Osteopetrosis 4, 611490	602727	128	100	100	99
CLCNKA	Bartter syndrome, type 4b, digenic, 613090	602024	151	100	100	99
CLCNKB	Bartter syndrome, type 3, 607364 Bartter syndrome, type 4b, digenic, 613090	602023	138	100	100	100
CLDN1	Ichthyosis, leukocyte vacuoles, alopecia, and sclerosing cholangitis, 607626	603718	115	100	100	100
CLDN14	Deafness 29, 614035	605608	66	100	98	91
CLDN16	Hypomagnesemia 3, renal, 248250	603959	73	100	100	97
CLDN19	Hypomagnesemia 5, renal, with ocular involvement, 248190	610036	149	100	100	100
CLEC7A	{Aspergillosis, susceptibility to}, 614079 Candidiasis, familial, 4, 613108	606264	64	100	100	91
CLIC2	?Mental retardation, syndromic 32, 300886	300138	40	100	90	69
CLMP	Congenital short bowel syndrome, 615237	611693	73	100	100	100
CLN3	Ceroid lipofuscinosi, neuronal, 3, 204200	607042	111	100	100	100
CLN5	Ceroid lipofuscinosi, neuronal, 5, 256731	608102	70	100	100	100
CLN6	Ceroid lipofuscinosi, neuronal, 6, 601780 Ceroid lipofuscinosi, neuronal, Kufs type, adult onset, 204300	606725	120	100	100	99
CLN8	Ceroid lipofuscinosi, neuronal, 8, 600143 Ceroid lipofuscinosi, neuronal, 8, Northern epilepsy variant, 610003	607837	129	100	100	100

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CLPP	Perrault syndrome 3, 614129	601119	125	100	100	100
CLRN1	Retinitis pigmentosa 61, 614180 Usher syndrome, type 3A, 276902	606397	83	100	100	97
CNBP	Myotonic dystrophy 2, 602668	116955	47	100	100	94
CNGA1	Retinitis pigmentosa 49, 613756	123825	62	96	91	86
CNGA3	Achromatopsia 2, 216900	600053	118	100	100	100
CNGB1	Retinitis pigmentosa 45, 613767	600724	112	100	100	99
CNGB3	Achromatopsia 3, 262300 Macular degeneration, juvenile, 248200	605080	62	100	100	95
CNNM2	Hypomagnesemia 6, renal, 613882 Hypomagnesemia, seizures, and mental retardation, 616418	607803	161	100	100	99
CNNM4	Jalili syndrome, 217080	607805	140	100	100	99
CNTN1	?Myopathy, congenital, Compton-North, 612540	600016	70	100	100	97
CNTNAP2	{Autism susceptibility 15}, 612100 Cortical dysplasia-focal epilepsy syndrome, 610042 Pitt-Hopkins like syndrome 1, 610042	604569	73	100	100	99
CNTNAP4	No OMIM phenotype	610518	86	100	99	95
COA5	?Cardioencephalomyopathy, fatal infantile, due to cytochrome c oxidase deficiency 3, 616500	613920	92	100	100	89
COASY	Neurodegeneration with brain iron accumulation 6, 615643 Pontocerebellar hypoplasia, type 12, 618266	609855	139	100	100	100
COCH	Deafness 9, 601369 ?Deafness 110, 618094	603196	71	98	95	93
COG1	Congenital disorder of glycosylation, type IIg, 611209	606973	96	100	100	98
COG4	Congenital disorder of glycosylation, type IIj, 613489 Saul-Wilson syndrome, 618150	606976	81	100	100	99
COG5	Congenital disorder of glycosylation, type III, 613612	606821	66	100	99	91
COG6	Congenital disorder of glycosylation, type III, 614576 Shaheen syndrome, 615328	606977	65	100	99	90
COG7	Congenital disorder of glycosylation, type Ile, 608779	606978	83	100	100	98
COG8	Congenital disorder of glycosylation, type IIh, 611182	606979	125	100	100	100
COL10A1	Metaphyseal chondrodysplasia, Schmid type, 156500	120110	90	100	100	99

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COL11A1	?Deafness 37, 618533 Fibrochondrogenesis 1, 228520 {Lumbar disc herniation, susceptibility to}, 603932 Marshall syndrome, 154780 Stickler syndrome, type II, 604841	120280	62	100	99	93
COL11A2	Deafness 13, 601868 Deafness 53, 609706 Fibrochondrogenesis 2, 614524 Otospondylomegaepiphyseal dysplasia, 184840 Otospondylomegaepiphyseal dysplasia, 215150	120290	110	100	100	100
COL17A1	Epidermolysis bullosa, junctional, localisata variant, 226650 Epidermolysis bullosa, junctional, non-Herlitz type, 226650 Epithelial recurrent erosion dystrophy, 122400	113811	98	100	100	99
COL18A1	Knobloch syndrome, type 1, 267750	120328	143	100	100	97
COL1A1	{Bone mineral density variation QTL, osteoporosis}, 166710 Caffey disease, 114000 Ehlers-Danlos syndrome, arthrochalasia type, 1, 130060 Osteogenesis imperfecta, type I, 166200 Osteogenesis imperfecta, type II, 166210 Osteogenesis imperfecta, type III, 259420 Osteogenesis imperfecta, type IV, 166220	120150	130	100	100	100
COL1A2	Ehlers-Danlos syndrome, arthrochalasia type, 2, 617821 Ehlers-Danlos syndrome, cardiac valvular type, 225320 Osteogenesis imperfecta, type II, 166210 Osteogenesis imperfecta, type III, 259420 Osteogenesis imperfecta, type IV, 166220 {Osteoporosis, postmenopausal}, 166710	120160	70	100	100	97

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COL2A1	Achondrogenesis, type II or hypochondrogenesis, 200610 Avascular necrosis of the femoral head, 608805 Czech dysplasia, 609162 Epiphyseal dysplasia, multiple, with myopia and deafness, 132450 Kniest dysplasia, 156550 Legg-Calve-Perthes disease, 150600 Osteoarthritis with mild chondrodysplasia, 604864 Platyspondylic skeletal dysplasia, Torrance type, 151210 SED congenita, 183900 SMED Strudwick type, 184250 Spondyloepiphyseal dysplasia, Stanescu type, 616583 Spondyloperipheral dysplasia, 271700 Stickler syndrome, type I, nonsyndromic ocular, 609508 Stickler syndrome, type I, 108300 Vitreoretinopathy with phalangeal epiphyseal dysplasia	120140	97	100	100	99
COL3A1	Ehlers-Danlos syndrome, vascular type, 130050 Polymicrogyria with or without vascular-type EDS, 618343	120180	122	100	100	100
COL4A1	Angiopathy, hereditary, with nephropathy, aneurysms, and muscle cramps, 611773 Brain small vessel disease with or without ocular anomalies, 175780 {Hemorrhage, intracerebral, susceptibility to}, 614519 ?Retinal arteries, tortuosity of, 180000	120130	88	100	100	97
COL4A2	Brain small vessel disease 2, 614483 {Hemorrhage, intracerebral, susceptibility to}, 614519	120090	102	100	100	99
COL4A3	Alport syndrome 2, 203780 Alport syndrome 3, 104200 Hematuria, benign familial, 141200	120070	63	100	97	90
COL4A4	Alport syndrome 2, 203780 Hematuria, familial benign, 141200	120131	70	100	100	96
COL4A5	Alport syndrome 1, 301050	303630	42	100	88	67
COL5A1	Ehlers-Danlos syndrome, classic type, 1, 130000	120215	136	100	100	98
COL5A2	Ehlers-Danlos syndrome, classic type, 2, 130010	120190	65	100	100	94
COL6A1	Bethlem myopathy 1, 158810 Ullrich congenital muscular dystrophy 1, 254090	120220	147	100	100	100

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COL6A2	Bethlem myopathy 1, 158810 ?Myosclerosis, congenital, 255600 Ullrich congenital muscular dystrophy 1, 254090	120240	148	100	100	99
COL6A3	Bethlem myopathy 1, 158810 Dystonia 27, 616411 Ullrich congenital muscular dystrophy 1, 254090	120250	104	100	100	99
COL7A1	EBD inversa, 226600 EBD, Bart type, 132000 EBD, localisata variant Epidermolysis bullosa dystrophica, AD, 131750 Epidermolysis bullosa dystrophica, AR, 226600 Epidermolysis bullosa pruriginosa, 604129 Epidermolysis bullosa, pretibial, 131850 Toenail dystrophy, isolated, 607523 Transient bullous of the newborn, 131705	120120	129	100	100	100
COL8A2	Corneal dystrophy, Fuchs endothelial, 1, 136800 Corneal dystrophy, posterior polymorphous 2, 609140	120252	95	100	100	98
COL9A1	?Epiphyseal dysplasia, multiple, 6, 614135 Stickler syndrome, type IV, 614134	120210	65	100	99	91
COL9A2	Epiphyseal dysplasia, multiple, 2, 600204 ?Stickler syndrome, type V, 614284	120260	100	100	99	96
COL9A3	Epiphyseal dysplasia, multiple, 3, with or without myopathy, 600969 {Intervertebral disc disease, susceptibility to}, 603932	120270	88	98	96	92
COLEC11	3MC syndrome 2, 265050	612502	168	100	100	100
COLQ	Myasthenic syndrome, congenital, 5, 603034	603033	83	100	100	98
COMP	Epiphyseal dysplasia, multiple, 1, 132400 Pseudoachondroplasia, 177170	600310	112	100	97	93
COQ2	Coenzyme Q10 deficiency, primary, 1, 607426 {Multiple system atrophy, susceptibility to}, 146500	609825	66	100	100	94
COQ6	Coenzyme Q10 deficiency, primary, 6, 614650	614647	81	100	99	93
COQ8A	Coenzyme Q10 deficiency, primary, 4, 612016	606980	129	100	100	100
COQ8B	Nephrotic syndrome, type 9, 615573	615567	93	100	98	96
COQ9	Coenzyme Q10 deficiency, primary, 5, 614654	612837	105	100	100	100
CORIN	Preeclampsia/eclampsia 5, 614595	605236	78	100	100	99
CORO1A	Immunodeficiency 8, 615401	605000	122	92	92	91

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COX10	Leigh syndrome due to mitochondrial COX4 deficiency, 256000 Mitochondrial complex IV deficiency, 220110	602125	139	100	100	99
COX14	?Mitochondrial complex IV deficiency, 220110	614478	138	100	100	100
COX15	Cardioencephalomyopathy, fatal infantile, due to cytochrome c oxidase deficiency 2, 615119 Leigh syndrome due to cytochrome c oxidase deficiency, 256000	603646	65	100	98	92
COX20	Mitochondrial complex IV deficiency, 220110	614698	53	100	97	83
COX4I2	Exocrine pancreatic insufficiency, dyserythropoietic anemia, and calvarial hyperostosis, 612714	607976	97	100	100	100
COX7B	Linear skin defects with multiple congenital anomalies 2, 300887	300885	61	100	100	100
CP	Cerebellar ataxia, 604290 Hemosiderosis, systemic, due to aceruloplasminemia, 604290 [Hypoceruloplasminemia, hereditary], 604290	117700	63	100	100	96
CPA6	Epilepsy, familial temporal lobe, 5, 614417 Febrile seizures, familial, 11, 614418	609562	79	100	99	94
CPLANE1	Joubert syndrome 17, 614615 Orofaciodigital syndrome VI, 277170	614571	71	100	99	95
CPN1	Carboxypeptidase N deficiency, 212070	603103	87	100	100	97
CPOX	Coproporphyrin, 121300 Harderoporphyrin, 121300	612732	82	100	100	98
CPS1	Carbamoylphosphate synthetase I deficiency, 237300 {Pulmonary hypertension, neonatal, susceptibility to}, 615371	608307	61	100	100	96
CPT1A	CPT deficiency, hepatic, type IA, 255120	600528	109	100	100	98
CPT2	CPT II deficiency, infantile, 600649 CPT II deficiency, lethal neonatal, 608836 CPT II deficiency, myopathic, stress-induced, 255110 {Encephalopathy, acute, infection-induced, 4, susceptibility to}, 614212	600650	106	100	100	99
CR2	Immunodeficiency, common variable, 7, 614699 {Systemic lupus erythematosus, susceptibility to, 9}, 610927	120650	63	100	100	97
CRADD	Mental retardation 34, with variant lissencephaly, 614499	603454	139	100	100	100
CRB1	Leber congenital amaurosis 8, 613835 Pigmented paravenous chorioretinal atrophy, 172870 Retinitis pigmentosa-12, 600105	604210	71	100	100	99
CRBN	Mental retardation 2, 607417	609262	67	100	99	91
CREB1	Histiocytoma, angiomyoid fibrous, somatic, 612160	123810	77	100	100	100
CREBBP	Menke-Hennekam syndrome 1, 618332 Rubinstein-Taybi syndrome 1, 180849	600140	85	100	99	94

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CRELD1	Atrioventricular septal defect, partial, with heterotaxy syndrome, 606217 {Atrioventricular septal defect, susceptibility to, 2}, 606217	607170	100	100	100	100
CRLF1	Cold-induced sweating syndrome 1, 272430	604237	105	96	91	89
CRPPA	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 7, 614643 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 7, 616052	614631	81	100	99	93
CRTAP	Osteogenesis imperfecta, type VII, 610682	605497	86	100	100	99
CRTC1	Mucoepidermoid salivary gland carcinoma	607536	146	100	100	99
CRX	Cone-rod retinal dystrophy-2, 120970 Leber congenital amaurosis 7, 613829	602225	196	100	100	100
CRYAA	Cataract 9, multiple types, 604219	123580	136	100	100	100
CRYAB	Cardiomyopathy, dilated, 1II, 615184 Cataract 16, multiple types, 613763 Myopathy, myofibrillar, 2, 608810 Myopathy, myofibrillar, fatal infantile hypertonic, alpha-B crystallin-related, 613869	123590	79	100	100	100
CRYBA1	Cataract 10, multiple types, 600881	123610	74	100	100	100
CRYBA4	Cataract 23, 610425	123631	106	100	100	100
CRYBB1	Cataract 17, multiple types, 611544	600929	97	100	100	100
CRYBB2	Cataract 3, multiple types, 601547	123620	181	100	100	100
CRYBB3	Cataract 22, 609741	123630	105	100	100	100
CRYGB	Cataract 39, multiple types, 615188	123670	87	100	100	100
CRYGC	Cataract 2, multiple types, 604307	123680	86	100	100	100
CRYGD	Cataract 4, multiple types, 115700	123690	97	100	100	100
CRYGS	Cataract 20, multiple types, 116100	123730	104	100	100	100
CRYM	Deafness 40, 616357	123740	76	100	100	99
CSF1R	Brain abnormalities, neurodegeneration, and dysosteosclerosis, 618476 Leukoencephalopathy, diffuse hereditary, with spheroids, 221820	164770	99	100	100	100
CSF2RA	Surfactant metabolism dysfunction, pulmonary, 4, 300770	306250	0	0	0	0
CSF2RB	Surfactant metabolism dysfunction, pulmonary, 5, 614370	138981	132	100	100	100
CSF3R	Neutropenia, severe congenital, 7, 617014	138971	98	100	100	100
CSNK1D	Advanced sleep-phase syndrome, familial, 2, 615224	600864	101	100	100	100
CSPP1	Joubert syndrome 21, 615636	611654	83	100	100	97
CSRP3	?Cardiomyopathy, dilated, 1M, 607482 Cardiomyopathy, hypertrophic, 12, 612124	600824	102	100	100	100
CST3	Cerebral amyloid angiopathy, 105150 {Macular degeneration, age-related, 11}, 611953	604312	97	100	100	92

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CSTA	Peeling skin syndrome 4, 607936	184600	60	100	100	96
CSTB	Epilepsy, progressive myoclonic 1A (Unverricht and Lundborg), 254800	601145	84	100	100	100
CTC1	Cereboretinal microangiopathy with calcifications and cysts, 612199	613129	100	100	100	100
CTCF	Mental retardation 21, 615502	604167	98	100	100	100
CTDP1	Congenital cataracts, facial dysmorphism, and neuropathy, 604168	604927	125	98	89	87
CTH	Cystathioninuria, 219500 Homocysteine, total plasma, elevated	607657	76	100	100	98
CTHRC1	Barrett esophagus/esophageal adenocarcinoma, 614266	610635	80	100	97	91
CTNNA3	Arrhythmogenic right ventricular dysplasia, familial, 13, 615616	607667	61	100	100	96
CTNNB1	Colorectal cancer, somatic, 114500	116806	62	100	100	97
	Exudative vitreoretinopathy 7, 617572					
	Hepatocellular carcinoma, somatic, 114550					
	Medulloblastoma, somatic, 155255					
	Neurodevelopmental disorder with spastic diplegia and visual defects, 615075					
	Ovarian cancer, somatic, 167000					
	Pilomatricoma, somatic, 132600					
CTNS	Cystinosis, atypical nephropathic, 219800	606272	118	100	100	100
	Cystinosis, late-onset juvenile or adolescent nephropathic, 219900					
	Cystinosis, nephropathic, 219800					
	Cystinosis, ocular nonnephropathic, 219750					
CTSA	Galactosialidosis, 256540	613111	124	100	100	100
CTSC	Haim-Munk syndrome, 245010	602365	75	100	100	97
	Papillon-Lefevre syndrome, 245000					
	Periodontitis 1, juvenile, 170650					
CTSD	Ceroid lipofuscinosi, neuronal, 10, 610127	116840	137	100	100	100
CTSF	Ceroid lipofuscinosi, neuronal, 13, Kufs type, 615362	603539	94	100	94	91
CTSK	Pycnodysostosis, 265800	601105	65	100	100	98
CUBN	Megaloblastic anemia-1, Finnish type, 261100	602997	82	100	100	96
CUL3	Pseudohypoaldosteronism, type II E, 614496	603136	73	100	97	88
CUL4B	Mental retardation, syndromic 15 (Cabezas type), 300354	300304	52	100	96	82
CUL7	3-M syndrome 1, 273750	609577	111	100	100	100
CXCR4	Myelokathexis, isolated	162643	82	83	83	82
	WHIM syndrome, 193670					
CYB5A	Methemoglobinemia and ambiguous genitalia, 250790	613218	77	100	100	100

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CYB5R3	Methemoglobinemia, type I, 250800 Methemoglobinemia, type II, 250800	613213	127	100	100	100
CYBA	Chronic granulomatous disease, autosomal, due to deficiency of CYBA, 233690	608508	106	94	81	66
CYBB	Chronic granulomatous disease, 306400 Immunodeficiency 34, mycobacteriosis, 300645	300481	49	100	98	91
CYC1	Mitochondrial complex III deficiency, nuclear type 6, 615453	123980	151	100	100	94
CYCS	Thrombocytopenia 4, 612004	123970	55	100	100	99
CYLD	Brooke-Spiegler syndrome, 605041 Cylindromatosis, familial, 132700 Trichoepithelioma, multiple familial, 1, 601606	605018	56	100	98	89
CYP11A1	Adrenal insufficiency, congenital, with 46XY sex reversal, partial or complete, 613743	118485	108	100	100	100
CYP11B1	Adrenal hyperplasia, congenital, due to 11-beta-hydroxylase deficiency, 202010 Aldosteronism, glucocorticoid-remediable, 103900	610613	209	100	100	100
CYP11B2	Aldosterone to renin ratio raised Hypoaldosteronism, congenital, due to CMO I deficiency, 203400 Hypoaldosteronism, congenital, due to CMO II deficiency, 610600 {Low renin hypertension, susceptibility to}	124080	212	100	100	100
CYP17A1	17-alpha-hydroxylase/17,20-lyase deficiency, 202110 17,20-lyase deficiency, isolated, 202110	609300	113	100	100	100
CYP19A1	Aromatase deficiency, 613546 Aromatase excess syndrome, 139300	107910	66	100	99	93
CYP1B1	Anterior segment dysgenesis 6, multiple subtypes, 617315 Glaucoma 3A, primary open angle, congenital, juvenile, or adult onset, 231300	601771	97	100	100	100
CYP21A2	Adrenal hyperplasia, congenital, due to 21-hydroxylase deficiency, 201910 Hyperandrogenism, nonclassic type, due to 21-hydroxylase deficiency, 201910	613815	201	100	100	100
CYP24A1	Hypercalcemia, infantile, 1, 143880	126065	90	100	100	98
CYP26B1	Craniosynostosis with radiohumeral fusions and other skeletal and craniofacial anomalies, 614416	605207	159	100	100	100
CYP26C1	Focal facial dermal dysplasia 4, 614974	608428	130	100	100	100
CYP27A1	Cerebrotendinous xanthomatosis, 213700	606530	131	100	100	100
CYP27B1	Vitamin D-dependent rickets, type I, 264700	609506	122	100	100	100
CYP2A6	Coumarin resistance, 122700 {Lung cancer, resistance to}, 211980 {Nicotine addiction, protection from}, 188890	122720	198	100	100	100
CYP2B6	{Efavirenz central nervous system toxicity, susceptibility to}, 614546 Efavirenz, poor metabolism of, 614546	123930	109	100	100	100

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CYP2C19	Clopidogrel, impaired responsiveness to, 609535 Mephenytoin poor metabolizer, 609535 Omeprazole poor metabolizer, 609535 Proguanil poor metabolizer, 609535	124020	101	100	100	99
CYP2C8	{Drug metabolism, altered, CYP2C8-related}, 618018	601129	86	100	97	94
CYP2C9	Tolbutamide poor metabolizer Warfarin sensitivity, 122700	601130	101	100	99	96
CYP2R1	Rickets due to defect in vitamin D 25-hydroxylation, 600081	608713	74	100	100	98
CYP2U1	Spastic paraplegia 56, 615030	610670	66	100	100	96
CYP46A1	No OMIM phenotype	604087	67	100	97	89
CYP4F22	Ichthyosis, congenital 5, 604777	611495	113	100	100	100
CYP4V2	Bietti crystalline corneoretinal dystrophy, 210370	608614	82	100	100	98
CYP7B1	Bile acid synthesis defect, congenital, 3, 613812 Spastic paraplegia 5A, 270800	603711	67	100	100	98
D2HGDH	D-2-hydroxyglutaric aciduria, 600721	609186	140	100	100	100
DAG1	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 9, 616538 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 9, 613818	128239	164	100	100	100
DARS1	Hypomyelination with brainstem and spinal cord involvement and leg spasticity, 615281	603084	60	100	99	89
DARS2	Leukoencephalopathy with brain stem and spinal cord involvement and lactate elevation, 611105	610956	63	100	100	97
DBH	Orthostatic hypotension 1, due to DBH deficiency, 223360	609312	136	100	100	100
DBT	Maple syrup urine disease, type II, 248600	248610	115	100	100	99
DCAF17	Woodhouse-Sakati syndrome, 241080	612515	75	100	100	94
DCC	Colorectal cancer, somatic, 114500 Esophageal carcinoma, somatic, 133239 Gaze palsy, familial horizontal, with progressive scoliosis, 2, 617542 Mirror movements 1 and/or agenesis of the corpus callosum, 157600	120470	70	100	100	98
DCHS1	Mitral valve prolapse 2, 607829 Van Maldergem syndrome 1, 601390	603057	128	100	100	100
DCLRE1C	Omenn syndrome, 603554 Severe combined immunodeficiency, Athabascan type, 602450	605988	64	100	100	92
DCN	Corneal dystrophy, congenital stromal, 610048	125255	69	100	100	100
DCTN1	{Amyotrophic lateral sclerosis, susceptibility to}, 105400 Neuropathy, distal hereditary motor, type VIIB, 607641 Perry syndrome, 168605	601143	95	100	100	99

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DCX	Lissencephaly, 300067 Subcortical laminal heterotopia, 300067	300121	57	100	98	92
DDB2	Xeroderma pigmentosum, group E, DDB-negative subtype, 278740	600811	113	100	100	99
DDC	Aromatic L-amino acid decarboxylase deficiency, 608643	107930	84	100	99	92
DDHD1	Spastic paraplegia 28, 609340	614603	107	100	98	92
DDHD2	Spastic paraplegia 54, 615033	615003	76	100	100	97
DDOST	?Congenital disorder of glycosylation, type Ir, 614507	602202	92	100	100	100
DDR2	Spondylometaepiphyseal dysplasia, short limb-hand type, 271665 Warburg-Cinotti syndrome, 618175	191311	89	100	100	99
DDX11	Warsaw breakage syndrome, 613398	601150	254	100	100	100
DDX3X	Mental retardation 102, 300958 dominant	300160	76	100	100	98
DDX59	Orofaciodigital syndrome V, 174300	615464	62	100	98	92
DEPDC5	Epilepsy, familial focal, with variable foci 1, 604364	614191	83	100	100	97
DES	Cardiomyopathy, dilated, 1I, 604765 Myopathy, myofibrillar, 1, 601419 Scapuloperoneal syndrome, neurogenic, Kaeser type, 181400	125660	106	100	100	100
DGKE	{Hemolytic uremic syndrome, atypical, susceptibility to, 7}, 615008 Nephrotic syndrome, type 7, 615008	601440	106	100	100	98
DGUOK	Mitochondrial DNA depletion syndrome 3 (hepatocerebral type), 251880 Portal hypertension, noncirrhotic, 617068 Progressive external ophthalmoplegia with mitochondrial DNA deletions 4, 617070	601465	73	100	100	92
DHCR24	Desmosterolosis, 602398	606418	119	100	100	100
DHCR7	Smith-Lemli-Opitz syndrome, 270400	602858	112	100	100	100
DHDDS	?Congenital disorder of glycosylation, type 1bb, 613861 Developmental delay and seizures with or without movement abnormalities, 617836 Retinitis pigmentosa 59, 613861	608172	69	100	100	99
DHFR	Megaloblastic anemia due to dihydrofolate reductase deficiency, 613839	126060	76	100	100	98
DHH	46XY partial gonadal dysgenesis, with minifascicular neuropathy, 607080 46XY sex reversal 7, 233420	605423	138	100	100	100
DHODH	Miller syndrome, 263750	126064	87	100	100	96
DHTKD1	2-amino adipic 2-oxoadipic aciduria, 204750 ?Charcot-Marie-Tooth disease, axonal, type 2Q, 615025	614984	85	100	100	98
DIABLO	Deafness 64, 614152	605219	105	100	100	99
DIAPH1	Deafness 1, 124900 Seizures, cortical blindness, microcephaly syndrome, 616632	602121	84	100	99	93

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DIAPH2	?Premature ovarian failure 2A, 300511	300108	44	100	92	75
DIAPH3	Auditory neuropathy, 1, 609129	614567	61	100	99	93
DICER1	GLOW syndrome, somatic mosaic, 618272 Goiter, multinodular 1, with or without Sertoli-Leydig cell tumors, 138800 Pleuropulmonary blastoma, 601200 Rhabdomyosarcoma, embryonal, 2, 180295	606241	77	100	99	95
DIP2B	Mental retardation, FRA12A type, 136630	611379	65	100	100	97
DIS3L2	Perlman syndrome, 267000	614184	100	100	100	97
DKC1	Dyskeratosis congenita, 305000	300126	53	100	98	89
DLAT	Pyruvate dehydrogenase E2 deficiency, 245348	608770	74	100	98	90
DLC1	Colorectal cancer, somatic, 114500	604258	98	100	100	99
DLD	Dihydrolipoamide dehydrogenase deficiency, 246900	238331	77	100	100	98
DLG3	Mental retardation 90, 300850	300189	61	100	97	88
DLL3	Spondylocostal dysostosis 1, 277300	602768	100	100	96	89
DLX3	Amelogenesis imperfecta, type IV, 104510 Trichodontosseous syndrome, 190320	600525	137	100	100	84
DLX5	?Split-hand/foot malformation 1 with sensorineural hearing loss, 220600	600028	109	100	100	100
DMD	Becker muscular dystrophy, 300376 Cardiomyopathy, dilated, 3B, 302045 Duchenne muscular dystrophy, 310200	300377	46	100	96	81
DMGDH	Dimethylglycine dehydrogenase deficiency, 605850	605849	75	100	100	99
DMP1	Hypophosphatemic rickets, AR, 241520	600980	80	100	100	97
DMPK	Myotonic dystrophy 1, 160900	605377	117	100	100	99
DNA2	Progressive external ophthalmoplegia with mitochondrial DNA deletions 6, 615156 ?Seckel syndrome 8, 615807	601810	93	100	100	96
DNAAF1	Ciliary dyskinesia, primary, 13, 613193	613190	131	100	100	99
DNAAF2	Ciliary dyskinesia, primary, 10, 612518	612517	124	100	100	100
DNAAF3	Ciliary dyskinesia, primary, 2, 606763	614566	98	100	100	99
DNAAF4	Ciliary dyskinesia, primary, 25, 615482 {Dyslexia, susceptibility to, 1}, 127700	608706	54	100	99	89
DNAAF5	Ciliary dyskinesia, primary, 18, 614874	614864	104	96	86	79
DNAH11	Ciliary dyskinesia, primary, 7, with or without situs inversus, 611884	603339	74	100	100	96
DNAH5	Ciliary dyskinesia, primary, 3, with or without situs inversus, 608644	603335	74	100	100	96
DNAI1	Ciliary dyskinesia, primary, 1, with or without situs inversus, 244400	604366	91	100	98	97
DNAI2	Ciliary dyskinesia, primary, 9, with or without situs inversus, 612444	605483	131	100	100	99

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DNAJB2	Spinal muscular atrophy, distal, 5, 614881	604139	103	100	100	100
DNAJB6	Muscular dystrophy, limb-girdle 1, 603511	611332	66	100	99	93
DNAJC19	3-methylglutaconic aciduria, type V, 610198	608977	75	100	100	96
DNAJC5	Ceroid lipofuscinoses, neuronal, 4, Parry type, 162350	611203	217	100	100	100
DNAJC6	Parkinson disease 19a, juvenile-onset, 615528	608375	84	100	100	98
	Parkinson disease 19b, early-onset, 615528					
DNAL1	Ciliary dyskinesia, primary, 16, 614017	610062	77	100	100	95
DNASE1L3	Systemic lupus erythematosus 16, 614420	602244	67	100	100	98
DNHD1	No OMIM phenotype	617277	117	100	100	100
DNM1L	Encephalopathy, lethal, due to defective mitochondrial peroxisomal fission 1, 614388	603850	70	100	99	94
	Optic atrophy 5, 610708					
DNM2	Centronuclear myopathy 1, 160150	602378	111	100	100	99
	Charcot-Marie-Tooth disease, axonal type 2M, 606482					
	Charcot-Marie-Tooth disease, dominant intermediate B, 606482					
	Lethal congenital contracture syndrome 5, 615368					
DNMT1	Cerebellar ataxia, deafness, and narcolepsy, 604121	126375	93	100	100	97
	Neuropathy, hereditary sensory, type Ie, 614116					
DNMT3B	Immunodeficiency-centromeric instability-facial anomalies syndrome 1, 242860	602900	115	100	100	99
DOCK3	Neurodevelopmental disorder with impaired intellectual development, hypotonia, and ataxia, 618292	603123	84	100	100	98
DOCK6	Adams-Oliver syndrome 2, 614219	614194	108	100	98	96
DOCK8	Hyper-IgE recurrent infection syndrome, 243700	611432	74	100	99	94
DOK7	?Fetal akinesia deformation sequence 3, 618389	610285	119	96	93	92
	Myasthenic syndrome, congenital, 10, 254300					
DOLK	Congenital disorder of glycosylation, type Im, 610768	610746	136	100	100	100
DPAGT1	Congenital disorder of glycosylation, type Ij, 608093	191350	76	100	100	100
	Myasthenic syndrome, congenital, 13, with tubular aggregates, 614750					
DPM1	Congenital disorder of glycosylation, type Ie, 608799	603503	79	94	89	85
DPM2	Congenital disorder of glycosylation, type Iu, 615042	603564	76	100	100	100
DPM3	Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 15, 612937	605951	227	100	100	100
DPP6	Mental retardation 33, 616311	126141	82	100	97	90
	{Ventricular fibrillation, paroxysmal familial, 2}, 612956					
DPY19L2	Spermatogenic failure 9, 613958	613893	90	98	95	88
DPYD	Dihydropyrimidine dehydrogenase deficiency, 274270	612779	66	100	99	95
	5-fluorouracil toxicity, 274270					
DPYS	Dihydropyrimidinuria, 222748	613326	75	100	100	98

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DRC1	Ciliary dyskinesia, primary, 21, 615294	615288	81	100	100	98
DRD2	No OMIM phenotype	126450	113	100	100	100
DRD4	{Attention deficit-hyperactivity disorder}, 143465 Autonomic nervous system dysfunction [Novelty seeking personality], 601696	126452	107	100	90	82
DRD5	{Attention deficit-hyperactivity disorder, susceptibility to}, 143465 {Blepharospasm, primary benign}, 606798	126453	239	100	100	100
DSC2	Arrhythmogenic right ventricular dysplasia 11, 610476 Arrhythmogenic right ventricular dysplasia 11 with mild palmoplantar keratoderma and woolly hair, 610476	125645	184	100	100	100
DSC3	?Hypotrichosis and recurrent skin vesicles, 613102	600271	73	100	99	89
DSG1	Erythroderma, congenital, with palmoplantar keratoderma, hypotrichosis, and hyper IgE, 615508 Keratosis palmoplantaris striata I, AD, 148700	125670	80	100	99	95
DSG2	Arrhythmogenic right ventricular dysplasia 10, 610193 Cardiomyopathy, dilated, 1BB, 612877	125671	158	100	100	99
DSG4	Hypotrichosis 6, 607903	607892	77	100	100	98
DSP	Arrhythmogenic right ventricular dysplasia 8, 607450 Cardiomyopathy, dilated, with woolly hair and keratoderma, 605676 Dilated cardiomyopathy with woolly hair, keratoderma, and tooth agenesis, 615821 Epidermolysis bullosa, lethal acantholytic, 609638 Keratosis palmoplantaris striata II, 612908 Skin fragility-woolly hair syndrome, 607655	125647	156	100	100	100
DSPP	Deafness 39, with dentinogenesis, 605594 Dentin dysplasia, type II, 125420 Dentinogenesis imperfecta, Shields type II, 125490 Dentinogenesis imperfecta, Shields type III, 125500	125485	43	95	70	55
DST	Epidermolysis bullosa simplex 2, 615425 ?Neuropathy, hereditary sensory and autonomic, type VI, 614653	113810	67	100	99	95
DTNA	Left ventricular noncompaction 1, with or without congenital heart defects, 604169	601239	84	100	100	100
DTNBP1	Hermansky-Pudlak syndrome 7, 614076	607145	94	100	100	96
DUOX2	Thyroid dyshormonogenesis 6, 607200	606759	113	98	97	96
DUOXA2	Thyroid dyshormonogenesis 5, 274900	612772	121	100	100	100
DUSP6	Hypogonadotropic hypogonadism 19 with or without anosmia, 615269	602748	144	100	100	100
DYM	Dyggve-Melchior-Clausen disease, 223800 Smith-McCort dysplasia, 607326	607461	67	100	98	88

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DYNC1H1	Charcot-Marie-Tooth disease, axonal, type 20, 614228 Mental retardation 13, 614563 Spinal muscular atrophy, lower extremity-predominant 1, AD, 158600	600112	96	100	100	99
DYNC2H1	Short-rib thoracic dysplasia 3 with or without polydactyly, 613091	603297	71	100	99	91
DYRK1A	Mental retardation 7, 614104	600855	75	100	100	98
DYSF	Miyoshi muscular dystrophy 1, 254130 Muscular dystrophy, limb-girdle 2, 253601 Myopathy, distal, with anterior tibial onset, 606768	603009	108	100	100	99
EARS2	Combined oxidative phosphorylation deficiency 12, 614924	612799	91	100	100	99
EBP	Chondrodysplasia punctata dominant, 302960 MEND syndrome, 300960	300205	93	100	100	100
ECE1	?Hirschsprung disease, cardiac defects, and autonomic dysfunction, 613870 {Hypertension, essential, susceptibility to}, 145500	600423	114	97	97	97
ECEL1	Arthrogryposis, distal, type 5D, 615065	605896	98	100	98	93
ECM1	Urbach-Wiethe disease, 247100	602201	125	100	100	100
EDA	Ectodermal dysplasia 1, hypohidrotic, 305100 Tooth agenesis, selective 1, 313500	300451	85	100	99	95
EDAR	Ectodermal dysplasia 10A, hypohidrotic/hair/nail type, 129490 Ectodermal dysplasia 10B, hypohidrotic/hair/tooth type, 224900 [Hair morphology 1, hair thickness], 612630	604095	103	100	100	100
EDARADD	Ectodermal dysplasia 11A, hypohidrotic/hair/tooth type, 614940 Ectodermal dysplasia 11B, hypohidrotic/hair/tooth type, 614941	606603	65	100	100	98
EDN1	Auriculocondylar syndrome 3, 615706 {High density lipoprotein cholesterol level QTL 7} Question mark ears, isolated, 612798	131240	92	100	100	93
EDN3	Central hypoventilation syndrome, congenital, 209880 {Hirschsprung disease, susceptibility to}, 4, 613712 Waardenburg syndrome, type 4B, 613265	131242	90	100	100	98
EDNRA	Mandibulofacial dysostosis with alopecia, 616367 {Migraine, resistance to}, 157300	131243	74	100	100	98
EDNRB	ABCD syndrome, 600501 {Hirschsprung disease, susceptibility to}, 2, 600155 Waardenburg syndrome, type 4A, 277580	131244	96	100	100	100
EFEMP1	Doyne honeycomb degeneration of retina, 126600	601548	99	100	100	99
EFEMP2	Cutis laxa, type IB, 614437	604633	103	100	100	100

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EFNB1	Craniofrontonasal dysplasia, 304110	300035	68	100	100	100
EFTUD2	Mandibulofacial dysostosis, Guion-Almeida type, 610536	603892	85	100	100	99
EGF	Hypomagnesemia 4, renal, 611718	131530	74	100	100	97
EGFR	Adenocarcinoma of lung, response to tyrosine kinase inhibitor in, 211980 ?Inflammatory skin and bowel disease, neonatal, 2, 616069	131550	87	100	100	98
	Non-small cell lung cancer, response to tyrosine kinase inhibitor in, 211980 {Non-small cell lung cancer, susceptibility to}, 211980					
EGLN1	Erythrocytosis, familial, 3, 609820 [Hemoglobin, high altitude adaptation], 609070	606425	114	100	92	83
EGR2	Charcot-Marie-Tooth disease, type 1D, 607678	129010	99	100	100	100
	Dejerine-Sottas disease, 145900					
	Hypomyelinating neuropathy, congenital, 1, 605253					
EHMT1	Kleefstra syndrome 1, 610253	607001	126	99	99	99
EIF2AK3	Wolcott-Rallison syndrome, 226980	604032	72	100	99	94
EIF2AK4	Pulmonary venoocclusive disease 2, 234810	609280	73	100	99	95
EIF2B1	Leukoencephalopathy with vanishing white matter, 603896	606686	79	100	100	100
EIF2B2	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896	606454	77	100	98	91
EIF2B3	Leukoencephalopathy with vanishing white matter, 603896	606273	58	100	100	94
EIF2B4	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896	606687	89	100	100	100
EIF2B5	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896	603945	72	100	100	98
EIF4A3	Robin sequence with cleft mandible and limb anomalies, 268305	608546	68	100	100	97
EIF4G1	{Parkinson disease 18}, 614251	600495	100	100	100	100
ELAC2	Combined oxidative phosphorylation deficiency 17, 615440 {Prostate cancer, hereditary, 2, susceptibility to}, 614731	605367	87	100	100	98
ELANE	Neutropenia, cyclic, 162800	130130	127	100	100	100
	Neutropenia, severe congenital 1, 202700					
ELF4	No OMIM phenotype	300775	67	100	100	100
ELN	Cutis laxa, 123700	130160	85	100	100	98
	Supravalvar aortic stenosis, 185500					
ELOVL4	Ichthyosis, spastic quadriplegia, and mental retardation, 614457	605512	71	100	100	98
	Spinocerebellar ataxia 34, 133190					
	Stargardt disease 3, 600110					

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
ELP1	Dysautonomia, familial, 223900	603722	No coverage data			
EMD	Emery-Dreifuss muscular dystrophy 1, 310300	300384	105	100	100	100
EMG1	Bowen-Conradi syndrome, 211180	611531	78	100	100	100
EMX2	Schizencephaly, 269160	600035	164	100	100	100
ENAM	Amelogenesis imperfecta, type IB, 104500	606585	60	100	100	98
	Amelogenesis imperfecta, type IC, 204650					
ENG	Telangiectasia, hereditary hemorrhagic, type 1, 187300	131195	98	100	100	99
ENO3	?Glycogen storage disease XIII, 612932	131370	148	100	100	100
ENPP1	Arterial calcification, generalized, of infancy, 1, 208000	173335	57	97	93	85
	Cole disease, 615522					
	{Diabetes mellitus, non-insulin-dependent, susceptibility to}, 125853					
	Hypophosphatemic rickets, 2, 613312					
	{Obesity, susceptibility to}, 601665					
ENTPD1	Spastic paraparesis 64, 615683	601752	68	100	100	96
EOGT	Adams-Oliver syndrome 4, 615297	614789	60	100	100	94
EP300	Colorectal cancer, somatic, 114500	602700	109	100	100	97
	Menke-Hennekam syndrome 2, 618333					
	Rubinstein-Taybi syndrome 2, 613684					
EPAS1	Erythrocytosis, familial, 4, 611783	603349	112	100	100	99
EPB41	Elliptocytosis-1, 611804	130500	66	100	100	98
EPB42	Spherocytosis, type 5, 612690	177070	126	100	100	100
EPCAM	Colorectal cancer, hereditary nonpolyposis, type 8, 613244	185535	58	100	100	93
	Diarrhea 5, with tufting enteropathy, congenital, 613217					
EPG5	Vici syndrome, 242840	615068	69	100	100	97
EPHA2	Cataract 6, multiple types, 116600	176946	141	100	100	99
EPHB2	?Bleeding disorder, platelet-type, 22, 618462	600997	147	98	98	98
	{Prostate cancer/brain cancer susceptibility, somatic}, 603688					
EPHX1	?Hypercholanemia, familial, 607748	132810	128	100	100	99
EPM2A	Epilepsy, progressive myoclonic 2A (Lafora), 254780	607566	86	89	87	85
EPX	[Eosinophil peroxidase deficiency], 261500	131399	116	100	100	100
ERBB2	Adenocarcinoma of lung, somatic, 211980	164870	134	100	98	98
	Gastric cancer, somatic, 613659					
	Glioblastoma, somatic, 137800					
	Ovarian cancer, somatic					

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
ERBB3	{?Erythroleukemia, familial, susceptibility to}, 133180 ?Lethal congenital contractual syndrome 2, 607598	190151	104	100	100	100
ERBB4	Amyotrophic lateral sclerosis 19, 615515	600543	65	100	99	93
ERCC1	Cerebrooculofacioskeletal syndrome 4, 610758	126380	71	100	100	94
ERCC2	?Cerebrooculofacioskeletal syndrome 2, 610756 Trichothiodystrophy 1, photosensitive, 601675 Xeroderma pigmentosum, group D, 278730	126340	97	100	99	98
ERCC3	Trichothiodystrophy 2, photosensitive, 616390 Xeroderma pigmentosum, group B, 610651	133510	82	100	100	99
ERCC4	Fanconi anemia, complementation group Q, 615272 XFE progeroid syndrome, 610965 Xeroderma pigmentosum, group F, 278760 Xeroderma pigmentosum, type F/Cockayne syndrome, 278760	133520	83	100	100	97
ERCC5	Cerebrooculofacioskeletal syndrome 3, 616570 Xeroderma pigmentosum, group G, 278780 Xeroderma pigmentosum, group G/Cockayne syndrome, 278780	133530	85	100	100	98
ERCC6	Cerebrooculofacioskeletal syndrome 1, 214150 Cockayne syndrome, type B, 133540 De Sanctis-Cacchione syndrome, 278800 {Lung cancer, susceptibility to}, 211980 {Macular degeneration, age-related, susceptibility to, 5}, 613761 Premature ovarian failure 11, 616946 UV-sensitive syndrome 1, 600630	609413	88	100	100	97
ERCC6L2	Bone marrow failure syndrome 2, 615715	615667	57	100	98	93
ERCC8	Cockayne syndrome, type A, 216400 UV-sensitive syndrome 2, 614621	609412	81	100	99	91
ERF	Chitayat syndrome, 617180 Craniosynostosis 4, 600775	611888	137	100	100	100
ERLIN2	Spastic paraparesis 18, 611225	611605	65	100	100	95
ESCO2	Roberts syndrome, 268300 SC phocomelia syndrome, 269000	609353	58	100	100	95
ESPN	Deafness 36, 609006 Deafness, neurosensory, without vestibular involvement, 609006	606351	101	97	88	81

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
ESR1	{Atherosclerosis, susceptibility to} Breast cancer, somatic, 114480 Estrogen resistance, 615363 {HDL response to hormone replacement, augmented} {Migraine, susceptibility to}, 157300 {Myocardial infarction, susceptibility to}, 608446	133430	115	100	100	99
ESRRB	Deafness 35, 608565	602167	142	100	100	100
ETFA	Glutaric acidemia IIA, 231680	608053	60	100	100	96
ETFB	Glutaric acidemia IIB, 231680	130410	106	100	100	100
ETFDH	Glutaric acidemia IIC, 231680	231675	85	100	100	99
ETHE1	Ethylmalonic encephalopathy, 602473	608451	113	100	100	98
ETV6	Leukemia, acute myeloid, somatic, 601626 Thrombocytopenia 5, 616216	600618	131	100	100	99
EVC	Ellis-van Creveld syndrome, 225500 ?Weyers acrofacial dysostosis, 193530	604831	98	96	94	94
EVC2	Ellis-van Creveld syndrome, 225500 Weyers acrofacial dysostosis, 193530	607261	87	100	99	95
EWSR1	Ewing sarcoma, 612219 Neuroepithelioma, 612219	133450	72	100	100	95
EXOSC3	Pontocerebellar hypoplasia, type 1B, 614678	606489	114	100	100	96
EXPH5	Epidermolysis bullosa, nonspecific, 615028	612878	61	100	100	98
EXT1	Chondrosarcoma, 215300 Exostoses, multiple, type 1, 133700	608177	70	100	100	96
EXT2	Exostoses, multiple, type 2, 133701 Seizures, scoliosis, and macrocephaly syndrome, 616682	608210	98	100	100	97
EXTL3	Immunoskeletal dysplasia with neurodevelopmental abnormalities, 617425	605744	150	100	100	100
EYA1	Anterior segment anomalies with or without cataract, 602588 Branchiootic syndrome 1, 602588 Branchiootorenal syndrome 1, with or without cataracts, 113650 ?Otofaciocervical syndrome, 166780	601653	86	100	100	98
EYA4	?Cardiomyopathy, dilated, 1J, 605362 Deafness 10, 601316	603550	62	100	100	94
EYS	Retinitis pigmentosa 25, 602772	612424	72	100	100	96
EZH2	Weaver syndrome, 277590	601573	79	100	100	97
F10	Factor X deficiency, 227600	613872	172	100	100	97

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
F11	Factor XI deficiency, 612416 Factor XI deficiency, 612416	264900	63	100	100	96
F12	Angioedema, hereditary, type III, 610618 Factor XII deficiency, 234000	610619	135	100	100	100
F13A1	Factor XIII A deficiency, 613225 {Myocardial infarction, protection against}, 608446 {Venous thrombosis, protection against}, 188050	134570	77	100	100	99
F13B	Factor XIII B deficiency, 613235	134580	83	100	100	98
F2	Dysprothrombinemia, 613679 Hypoprothrombinemia, 613679 {Pregnancy loss, recurrent, susceptibility to, 2}, 614390 {Stroke, ischemic, susceptibility to}, 601367 Thrombophilia due to thrombin defect, 188050	176930	102	100	100	99
F5	{Budd-Chiari syndrome}, 600880 Factor V deficiency, 227400 {Pregnancy loss, recurrent, susceptibility to, 1}, 614389 {Stroke, ischemic, susceptibility to}, 601367 Thrombophilia due to activated protein C resistance, 188055 {Thrombophilia, susceptibility to, due to factor V Leiden}, 188055	612309	113	100	100	97
F7	Factor VII deficiency, 227500 {Myocardial infarction, decreased susceptibility to}, 608446	613878	138	100	100	100
F8	Hemophilia A, 306700	300841	48	100	99	88
F9	{Deep venous thrombosis, protection against}, 300807 Hemophilia B, 306900 Thrombophilia, due to factor IX defect, 300807 {Warfarin sensitivity}, 122700	300746	58	100	100	96
FA2H	Spastic paraparesis 35, 612319	611026	83	100	99	91
FADD	Infections, recurrent, with encephalopathy, hepatic dysfunction, and cardiovascular malformations, 613759	602457	143	100	100	100
FAH	Tyrosinemia, type I, 276700	613871	99	100	100	100
FAM111A	Gracile bone dysplasia, 602361 Kenny-Caffey syndrome, type 2, 127000	615292	75	100	100	100
FAM111B	Poikiloderma, hereditary fibrosis, with tendon contractures, myopathy, and pulmonary fibrosis, 615704	615584	65	100	100	99
FAM126A	Leukodystrophy, hypomyelinating, 5, 610532	610531	63	100	100	98
FAM161A	Retinitis pigmentosa 28, 606068	613596	64	100	100	96
FAM20A	Amelogenesis imperfecta, type IG (enamel-renal syndrome), 204690	611062	110	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FAM20C	Raine syndrome, 259775	611061	125	100	100	100
FAM83H	Amelogenesis imperfecta, type IIIA, 130900	611927	114	100	100	98
FAN1	Interstitial nephritis, karyomegalic, 614817	613534	75	100	100	98
FANCA	Fanconi anemia, complementation group A, 227650	607139	102	100	100	97
FANCB	Fanconi anemia, complementation group B, 300514	300515	45	100	97	83
FANCC	Fanconi anemia, complementation group C, 227645	613899	69	100	99	91
FANCD2	Fanconi anemia, complementation group D2, 227646	613984	71	100	100	96
FANCE	Fanconi anemia, complementation group E, 600901	613976	123	100	99	90
FANCF	Fanconi anemia, complementation group F, 603467	613897	170	100	100	100
FANCG	Fanconi anemia, complementation group G, 614082	602956	126	100	100	100
FANCI	Fanconi anemia, complementation group I, 609053	611360	72	100	100	98
FANCL	Fanconi anemia, complementation group L, 614083	608111	57	100	99	88
FANCM	?Premature ovarian failure 15, 618096 Spermatogenic failure 28, 618086	609644	75	100	99	95
FARS2	Combined oxidative phosphorylation deficiency 14, 614946 Spastic paraplegia 77, 617046	611592	111	100	100	100
FAS	Autoimmune lymphoproliferative syndrome, type IA, 601859 {Autoimmune lymphoproliferative syndrome}, 601859 Squamous cell carcinoma, burn scar-related, somatic	134637	264	100	100	96
FASLG	Autoimmune lymphoproliferative syndrome, type IB, 601859 {Lung cancer, susceptibility to}, 211980	134638	65	100	100	100
FAT1	No OMIM phenotype	600976	75	100	100	99
FAT4	Hennekam lymphangiectasia-lymphedema syndrome 2, 616006 Van Maldergem syndrome 2, 615546	612411	90	100	100	99
FBLN1	Synpolydactyly, 3/3'4, associated with metacarpal and metatarsal synostoses, 608180	135820	117	100	99	99
FBLN5	Cutis laxa 2, 614434 Cutis laxa, type IA, 219100 Macular degeneration, age-related, 3, 608895 Neuropathy, hereditary, with or without age-related macular degeneration, 608895	604580	106	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FBN1	Acromicric dysplasia, 102370 Ectopia lentis, familial, 129600 Geleophysic dysplasia 2, 614185 MASS syndrome, 604308 Marfan lipodystrophy syndrome, 616914 Marfan syndrome, 154700 Stiff skin syndrome, 184900 Weill-Marchesani syndrome 2, dominant, 608328	134797	186	100	100	100
FBN2	Contractural arachnodactyly, congenital, 121050 Macular degeneration, early-onset, 616118	612570	74	100	100	97
FBP1	Fructose-1,6-bisphosphatase deficiency, 229700	611570	119	100	100	100
FBXL4	Mitochondrial DNA depletion syndrome 13 (encephalomyopathic type), 615471	605654	67	100	100	98
FBXO38	Neuronopathy, distal hereditary motor, type IID, 615575	608533	83	100	100	96
FBXO7	Parkinson disease 15, 260300	605648	86	100	100	97
FCGR3A	Immunodeficiency 20, 615707	146740	244	100	100	98
FCGR3B	Neutropenia, alloimmune neonatal	610665	210	100	100	100
FCN3	Immunodeficiency due to ficolin 3 deficiency, 613860	604973	103	100	100	97
FECH	Protoporphria, erythropoietic, 1, 177000	612386	64	100	100	96
FERMT1	Kindler syndrome, 173650	607900	70	100	99	92
FERMT3	Leukocyte adhesion deficiency, type III, 612840	607901	114	100	100	97
FGA	Afibrinogenemia, congenital, 202400 Amyloidosis, familial visceral, 105200 Dysfibrinogenemia, congenital, 616004 Hypodysfibrinogenemia, congenital, 616004	134820	110	100	100	98
FGB	Afibrinogenemia, congenital, 202400 Dysfibrinogenemia, congenital, 616004 Hypofibrinogenemia, congenital, 202400	134830	78	100	100	99
FGD1	Aarskog-Scott syndrome, 305400 Mental retardation syndromic 16, 305400	300546	71	100	98	95
FGD4	Charcot-Marie-Tooth disease, type 4H, 609311	611104	67	100	98	92
FGF10	Aplasia of lacrimal and salivary glands, 180920 LADD syndrome, 149730	602115	54	100	98	90
FGF14	Spinocerebellar ataxia 27, 609307	601515	68	100	100	98
FGF16	Metacarpal 4-5 fusion, 309630	300827	98	100	100	95
FGF17	Hypogonadotropic hypogonadism 20 with or without anosmia, 615270	603725	142	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FGF23	Hypophosphatemic rickets, 193100 Osteomalacia, tumor-induced Tumoral calcinosis, hyperphosphatemic, familial, 2, 617993	605380	88	100	100	97
FGF3	Deafness, congenital with inner ear agenesis, microtia, and microdontia, 610706	164950	122	100	100	100
FGF8	Hypogonadotropic hypogonadism 6 with or without anosmia, 612702	600483	134	100	94	94
FGF9	Multiple synostoses syndrome 3, 612961	600921	80	100	100	100
FGFR1	Encephalocraniocutaneous lipomatosis, somatic mosaic, 613001 Hartsfield syndrome, 615465 Hypogonadotropic hypogonadism 2 with or without anosmia, 147950 Jackson-Weiss syndrome, 123150 Osteoglophonic dysplasia, 166250 Pfeiffer syndrome, 101600 Trigonocephaly 1, 190440	136350	102	100	100	98
FGFR2	Antley-Bixler syndrome without genital anomalies or disordered steroidogenesis, 207410 Apert syndrome, 101200 Beare-Stevenson cutis gyrata syndrome, 123790 Bent bone dysplasia syndrome, 614592 Craniofacial-skeletal-dermatologic dysplasia, 101600 Craniosynostosis, nonspecific Crouzon syndrome, 123500 Gastric cancer, somatic, 613659 Jackson-Weiss syndrome, 123150 LADD syndrome, 149730 Pfeiffer syndrome, 101600 Saethre-Chotzen syndrome, 101400 Scaphocephaly and Axenfeld-Rieger anomaly Scaphocephaly, maxillary retrusion, and mental retardation, 609579	176943	74	100	100	95

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FGFR3	Achondroplasia, 100800 Bladder cancer, somatic, 109800 CATSHL syndrome, 610474 Cervical cancer, somatic, 603956 Colorectal cancer, somatic, 114500 Crouzon syndrome with acanthosis nigricans, 612247 Hypochondroplasia, 146000 LADD syndrome, 149730 Muenke syndrome, 602849 Nevus, epidermal, somatic, 162900 SADDAN, 616482 Spermatocytic seminoma, somatic, 273300 Thanatophoric dysplasia, type I, 187600 Thanatophoric dysplasia, type II, 187601	134934	122	100	100	98
FGG	Afibrinogenemia, congenital, 202400 Dysfibrinogenemia, congenital, 616004 Hypodysfibrinogenemia, 616004 Hypofibrinogenemia, congenital, 202400	134850	69	100	100	97
FH	Fumarase deficiency, 606812 Leiomyomatosis and renal cell cancer, 150800	136850	82	99	94	87
FHL1	Emery-Dreifuss muscular dystrophy 6, 300696 Myopathy, with postural muscle atrophy, 300696 Reducing body myopathy 1a, severe, infantile or early childhood onset, 300717 Reducing body myopathy 1b, with late childhood or adult onset, 300718 Scapuloperoneal myopathy dominant, 300695 ?Uruguay faciocardiomusculoskeletal syndrome, 300280	300163	92	100	100	94
FIG4	Amyotrophic lateral sclerosis 11, 612577 Charcot-Marie-Tooth disease, type 4J, 611228 ?Polymicrogyria, bilateral temporooccipital, 612691 Yunis-Varon syndrome, 216340	609390	59	100	99	94
FIGLA	Premature ovarian failure 6, 612310	608697	86	100	100	96
FKBP10	Bruck syndrome 1, 259450 Osteogenesis imperfecta, type XI, 610968	607063	131	100	100	100
FKBP14	Ehlers-Danlos syndrome, kyphoscoliotic type, 2, 614557	614505	96	100	100	99

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FKRP	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 5, 613153 Muscular dystrophy-dystroglycanopathy (congenital with or without mental retardation), type B, 5, 606612 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 5, 607155	606596	143	100	100	100
FKTN	Cardiomyopathy, dilated, 1X, 611615 Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 4, 253800 Muscular dystrophy-dystroglycanopathy (congenital without mental retardation), type B, 4, 613152 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 4, 611588	607440	85	100	100	100
FLCN	Birt-Hogg-Dube syndrome, 135150 Colorectal cancer, somatic, 114500 Pneumothorax, primary spontaneous, 173600 Renal carcinoma, chromophobe, somatic, 144700	607273	127	100	100	100
FLG	{Dermatitis, atopic, susceptibility to, 2}, 605803 Ichthyosis vulgaris, 146700	135940	378	100	100	100
FLNA	Cardiac valvular dysplasia, 314400 Congenital short bowel syndrome, 300048 ?FG syndrome 2, 300321 Frontometaphyseal dysplasia 1, 305620 Heterotopia, periventricular, 1, 300049 Intestinal pseudoobstruction, neuronal, 300048 Melnick-Needles syndrome, 309350 Otopalatodigital syndrome, type I, 311300 Otopalatodigital syndrome, type II, 304120 Terminal osseous dysplasia, 300244	300017	110	100	100	100
FLNB	Atelosteogenesis, type I, 108720 Atelosteogenesis, type III, 108721 Boomerang dysplasia, 112310 Larsen syndrome, 150250 Spondylocarpotarsal synostosis syndrome, 272460	603381	110	100	100	99
FLNC	Cardiomyopathy, familial hypertrophic, 26 Cardiomyopathy, familial restrictive 5, 617047 Myopathy, distal, 4, 614065 Myopathy, myofibrillar, 5, 609524	102565	133	100	100	100
FLRT3	Hypogonadotropic hypogonadism 21 with anosmia, 615271	604808	70	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FLT3	Leukemia, acute lymphoblastic, somatic, 613065 Leukemia, acute myeloid, reduced survival in, somatic, 601626 Leukemia, acute myeloid, somatic, 601626	136351	68	100	100	94
FLT4	Hemangioma, capillary infantile, somatic, 602089 Lymphatic malformation 1, 153100	136352	132	100	100	100
FLVCR1	Ataxia, posterior column, with retinitis pigmentosa, 609033	609144	98	100	100	96
FLVCR2	Proliferative vasculopathy and hydranencephaly-hydrocephaly syndrome, 225790	610865	140	100	100	99
FMO3	Trimethylaminuria, 602079	136132	69	100	100	95
FMR1	Fragile X syndrome, 300624 Fragile X tremor/ataxia syndrome, 300623 Premature ovarian failure 1, 311360	309550	41	100	96	76
FN1	Glomerulopathy with fibronectin deposits 2, 601894 Plasma fibronectin deficiency, 614101 Spondylometaphyseal dysplasia, corner fracture type, 184255	135600	80	100	100	96
FOLR1	Neurodegeneration due to cerebral folate transport deficiency, 613068	136430	111	100	100	100
FOXC1	Anterior segment dysgenesis 3, multiple subtypes, 601631 Axenfeld-Rieger syndrome, type 3, 602482	601090	80	100	95	89
FOXC2	Lymphedema-distichiasis syndrome, 153400 Lymphedema-distichiasis syndrome with renal disease and diabetes mellitus, 153400	602402	137	100	100	100
FOXE1	Bamforth-Lazarus syndrome, 241850 {Thyroid cancer, nonmedullary, 4}, 616534	602617	142	100	100	97
FOXE3	Anterior segment dysgenesis 2, multiple subtypes, 610256 {Aortic aneurysm, familial thoracic 11, susceptibility to}, 617349 Cataract 34, multiple types, 612968	601094	74	87	79	74
FOXF1	Alveolar capillary dysplasia with misalignment of pulmonary veins, 265380	601089	159	100	100	100
FOXG1	Rett syndrome, congenital variant, 613454	164874	127	97	90	84
FOXI1	Enlarged vestibular aqueduct, 600791	601093	161	100	100	100
FOXL2	Blepharophimosis, epicanthus inversus, and ptosis, type 1, 110100 Blepharophimosis, epicanthus inversus, and ptosis, type 2, 110100 Premature ovarian failure 3, 608996	605597	114	100	94	88
FOXN1	T-cell immunodeficiency, congenital alopecia, and nail dystrophy, 601705	600838	136	100	100	100
FOXO1	Rhabdomyosarcoma, alveolar, 268220	136533	93	100	98	96
FOXP1	Mental retardation with language impairment and with or without autistic features, 613670	605515	80	100	100	98
FOXP2	Speech-language disorder-1, 602081	605317	72	100	100	98
FOXP3	Immunodysregulation, polyendocrinopathy, and enteropathy, 304790	300292	69	100	99	88

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
FOXRED1	Mitochondrial complex I deficiency, nuclear type 19, 618241	613622	91	100	100	100
FRAS1	Fraser syndrome 1, 219000	607830	79	100	100	97
FREM1	Bifid nose with or without anorectal and renal anomalies, 608980	608944	83	100	100	98
	Manitoba oculotrichoanal syndrome, 248450					
	Trigonocephaly 2, 614485					
FREM2	Cryptophthalmos, unilateral or bilateral, isolated, 123570	608945	104	100	100	100
	Fraser syndrome 2, 617666					
FRMD7	Nystagmus 1, congenital, 310700	300628	50	100	98	89
	Nystagmus, infantile periodic alternating, 310700					
FSCN2	Retinitis pigmentosa 30, 607921	607643	139	100	100	100
FSHB	Hypogonadotropic hypogonadism 24 without anosmia, 229070	136530	81	100	100	100
FSHR	Ovarian dysgenesis 1, 233300	136435	71	100	100	98
	Ovarian hyperstimulation syndrome, 608115					
	Ovarian response to FSH stimulation, 276400					
FTCD	Glutamate formiminotransferase deficiency, 229100	606806	96	99	96	93
FTL	Hyperferritinemia-cataract syndrome, 600886	134790	144	100	100	100
	L-ferritin deficiency, dominant and recessive, 615604					
	Neurodegeneration with brain iron accumulation 3, 606159					
FTO	Growth retardation, developmental delay, facial dysmorphism, 612938	610966	90	100	100	99
	{Obesity, susceptibility to, BMIQ14}, 612460					
FTSJ1	Mental retardation 9/44, 309549	300499	91	100	100	93
FUCA1	Fucosidosis, 230000	612280	83	100	100	98
FUS	Amyotrophic lateral sclerosis 6, with or without frontotemporal dementia, 608030	137070	77	100	100	97
	Essential tremor, hereditary, 4, 614782					
FUT6	Fucosyltransferase 6 deficiency, 613852	136836	228	100	100	100
FUZ	{Neural tube defects, susceptibility to}, 182940	610622	109	100	100	100
FXN	Friedreich ataxia, 229300	606829	41	100	91	64
	Friedreich ataxia with retained reflexes, 229300					
FXYD2	Hypomagnesemia 2, renal, 154020	601814	101	100	100	100
FYCO1	Cataract 18, 610019	607182	115	100	100	99
FZD4	Exudative vitreoretinopathy 1, 133780	604579	90	100	100	99
	Retinopathy of prematurity, 133780					
FZD6	Nail disorder, nonsyndromic congenital, 10, (claw-shaped nails), 614157	603409	62	100	100	98
G6PC	Glycogen storage disease Ia, 232200	613742	113	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
G6PC3	Dursun syndrome, 612541 Neutropenia, severe congenital 4, 612541	611045	103	100	100	100
G6PD	Hemolytic anemia, G6PD deficient (favism), 300908 {Resistance to malaria due to G6PD deficiency}, 611162	305900	96	100	100	100
GAA	Glycogen storage disease II, 232300	606800	156	100	100	100
GABRA1	{Epilepsy, childhood absence, susceptibility to, 4}, 611136 {Epilepsy, juvenile myoclonic, susceptibility to, 5}, 611136 Epileptic encephalopathy, early infantile, 19, 615744	137160	76	100	100	100
GABRB3	{Epilepsy, childhood absence, susceptibility to, 5}, 612269 Epileptic encephalopathy, early infantile, 43, 617113	137192	95	100	100	98
GABRG2	Epilepsy, generalized, with febrile seizures plus, type 3, 607681 Epileptic encephalopathy, early infantile, 74, 618396 Febrile seizures, familial, 8, 607681	137164	73	94	92	91
GAD1	?Cerebral palsy, spastic quadriplegic, 1, 603513	605363	86	100	100	99
GALC	Krabbe disease, 245200	606890	54	100	99	89
GALE	Galactose epimerase deficiency, 230350	606953	116	100	100	100
GALK1	Galactokinase deficiency with cataracts, 230200	604313	133	100	100	99
GALNS	Mucopolysaccharidosis IVA, 253000	612222	81	100	99	95
GALNT3	Tumoral calcinosis, hyperphosphatemic, familial, 1, 211900	601756	69	100	100	97
GALT	Galactosemia, 230400	606999	151	100	100	100
GAMT	Cerebral creatine deficiency syndrome 2, 612736	601240	85	100	99	95
GAN	Giant axonal neuropathy-1, 256850	605379	86	100	100	99
GARS1	Charcot-Marie-Tooth disease, type 2D, 601472 Neuropathy, distal hereditary motor, type VA, 600794	600287	72	100	100	97
GATA1	Anemia, with/without neutropenia and/or platelet abnormalities, 300835 Leukemia, megakaryoblastic, with or without Down syndrome, somatic, 190685 Thrombocytopenia with beta-thalassemia, 314050 Thrombocytopenia, with or without dyserythropoietic anemia, 300367	305371	75	100	100	96
GATA2	Emberger syndrome, 614038 Immunodeficiency 21, 614172 {Leukemia, acute myeloid, susceptibility to}, 601626 {Myelodysplastic syndrome, susceptibility to}, 614286	137295	100	100	100	100
GATA3	Hypoparathyroidism, sensorineural deafness, and renal dysplasia, 146255	131320	203	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GATA4	Atrial septal defect 2, 607941 Atrioventricular septal defect 4, 614430 ?Testicular anomalies with or without congenital heart disease, 615542 Tetralogy of Fallot, 187500 Ventricular septal defect 1, 614429	600576	81	100	87	80
GATA5	Congenital heart defects, multiple types, 5, 617912	611496	78	100	99	95
GATA6	Atrial septal defect 9, 614475 Atrioventricular septal defect 5, 614474 Pancreatic agenesis and congenital heart defects, 600001 Persistent truncus arteriosus, 217095 Tetralogy of Fallot, 187500	601656	128	100	95	91
GATAD1	?Cardiomyopathy, dilated, 2B, 614672	614518	73	100	100	97
GATAD2B	Mental retardation 18, 615074	614998	67	100	100	99
GATM	Cerebral creatine deficiency syndrome 3, 612718	602360	65	100	100	98
GBA	Gaucher disease, perinatal lethal, 608013 Gaucher disease, type I, 230800 Gaucher disease, type II, 230900 Gaucher disease, type III, 231000 Gaucher disease, type IIIC, 231005 {Lewy body dementia, susceptibility to}, 127750 {Parkinson disease, late-onset, susceptibility to}, 168600	606463	172	100	100	100
GBA2	Spastic paraparesis 46, 614409	609471	137	100	100	100
GBE1	Glycogen storage disease IV, 232500 Polyglucosan body disease, adult form, 263570	607839	67	100	99	92
GCDH	Glutaric aciduria, type I, 231670	608801	112	100	100	100
GCH1	Dystonia, DOPA-responsive, with or without hyperphenylalaninemia, 128230 Hyperphenylalaninemia, BH4-deficient, B, 233910	600225	52	100	100	93
GCK	Diabetes mellitus, noninsulin-dependent, late onset, 125853 Diabetes mellitus, permanent neonatal, 606176 Hyperinsulinemic hypoglycemia, familial, 3, 602485 MODY, type II, 125851	138079	127	100	100	100
GCLC	Hemolytic anemia due to gamma-glutamylcysteine synthetase deficiency, 230450 {Myocardial infarction, susceptibility to}, 608446	606857	88	100	100	97
GCM2	Hyperparathyroidism 4, 617343 Hypoparathyroidism, familial isolated, 146200	603716	84	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GCNT2	Adult i phenotype without cataract, 110800 [Blood group, li], 110800 Cataract 13 with adult i phenotype, 116700	600429	81	100	100	98
GCSH	?Glycine encephalopathy, 605899	238330	93	100	89	62
GDAP1	Charcot-Marie-Tooth disease, axonal, type 2K, 607831 Charcot-Marie-Tooth disease, axonal, with vocal cord paresis, 607706 Charcot-Marie-Tooth disease, recessive intermediate, A, 608340 Charcot-Marie-Tooth disease, type 4A, 214400	606598	77	100	100	100
GDF1	Congenital heart defects, multiple types, 6, 613854 Right atrial isomerism (Ivemark), 208530	602880	69	100	93	81
GDF2	Telangiectasia, hereditary hemorrhagic, type 5, 615506	605120	122	100	100	100
GDF3	Klippel-Feil syndrome 3, 613702 Microphthalmia with coloboma 6, 613703 Microphthalmia, isolated 7, 613704	606522	120	100	100	100
GDF5	?Acromesomelic dysplasia, Hunter-Thompson type, 201250 Brachydactyly, type A1, C, 615072 Brachydactyly, type A2, 112600 Brachydactyly, type C, 113100 Chondrodysplasia, Grebe type, 200700 Du Pan syndrome, 228900 Multiple synostoses syndrome 2, 610017 {Osteoarthritis-5}, 612400 Symphalangism, proximal, 1B, 615298	601146	143	100	100	100
GDF6	Klippel-Feil syndrome 1, 118100 Leber congenital amaurosis 17, 615360 Microphthalmia with coloboma 6, digenic, 613703 Microphthalmia, isolated 4, 613094 Multiple synostoses syndrome 4, 617898	601147	124	100	100	100
GDI1	Mental retardation 41, 300849	300104	97	100	100	100
GDNF	Central hypoventilation syndrome, 209880 {Hirschsprung disease, susceptibility to, 3}, 613711 {Pheochromocytoma, modifier of}, 171300	600837	81	100	100	100
GFAP	Alexander disease, 203450	137780	81	100	100	99
GFER	Myopathy progressive, with congenital cataract, hearing loss, and developmental delay, 613076	600924	86	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GFI1	?Neutropenia, nonimmune chronic idiopathic, of adults, 607847 ?Neutropenia, severe congenital 2, 613107	600871	136	100	100	100
GFI1B	Bleeding disorder, platelet-type, 17, 187900	604383	193	100	100	100
GFM1	Combined oxidative phosphorylation deficiency 1, 609060	606639	72	100	100	96
GFPT1	Myasthenia, congenital, 12, with tubular aggregates, 610542	138292	59	100	100	92
GGCX	Pseudoxanthoma elasticum-like disorder with multiple coagulation factor deficiency, 610842 Vitamin K-dependent clotting factors, combined deficiency of, 1, 277450	137167	82	100	100	100
GH1	Growth hormone deficiency, isolated, type IA, 262400 Growth hormone deficiency, isolated, type IB, 612781 Growth hormone deficiency, isolated, type II, 173100 Kowarski syndrome, 262650	139250	177	100	100	100
GHR	Growth hormone insensitivity, partial, 604271 {Hypercholesterolemia, familial, modifier of}, 143890 Increased responsiveness to growth hormone, 604271 Laron dwarfism, 262500	600946	77	100	100	98
GHRHR	Growth hormone deficiency, isolated, type IV, 618157	139191	87	100	100	98
GHSR	Growth hormone deficiency, isolated partial, 615925	601898	137	100	100	100
GIGYF2	{Parkinson disease 11}, 607688	612003	66	100	99	94
GIPC3	Deafness 15, 601869	608792	115	100	95	93
GJA1	Atrioventricular septal defect 3, 600309 Craniometaphyseal dysplasia, 218400 Erythrokeratodermia variabilis et progressiva 3, 617525 Hypoplastic left heart syndrome 1, 241550 Oculodentodigital dysplasia, 164200 Oculodentodigital dysplasia, 257850 Palmoplantar keratoderma with congenital alopecia, 104100 Syndactyly, type III, 186100	121014	101	100	100	100
GJA3	Cataract 14, multiple types, 601885	121015	138	100	100	100
GJA5	Atrial fibrillation, familial, 11, 614049 Atrial standstill, digenic (GJA5/SCN5A), 108770	121013	211	100	100	100
GJA8	Cataract 1, multiple types, 116200	600897	145	100	100	100
GJB1	Charcot-Marie-Tooth neuropathy dominant, 1, 302800	304040	91	100	100	99

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GJB2	Bart-Pumphrey syndrome, 149200 Deafness 3A, 601544 Deafness 1A, 220290 Hystrix-like ichthyosis with deafness, 602540 Keratitis-ichthyosis-deafness syndrome, 148210 Keratoderma, palmoplantar, with deafness, 148350 Vohwinkel syndrome, 124500	121011	150	100	100	100
GJB3	Deafness 2B, 612644 Deafness, with peripheral neuropathy Deafness Deafness, digenic, GJB2/GJB3, 220290 Erythrokeratoderma variabilis et progressiva 1, 133200	603324	182	100	100	100
GJB4	Erythrokeratoderma variabilis et progressiva 2, 617524	605425	208	100	100	100
GJB6	Deafness 3B, 612643 Deafness 1B, 612645 Deafness, digenic GJB2/GJB6, 220290 Ectodermal dysplasia 2, Clouston type, 129500	604418	75	100	100	99
GJC2	Leukodystrophy, hypomyelinating, 2, 608804 Lymphatic malformation 3, 613480 Spastic paraparesis 44, 613206	608803	85	96	86	76
GK	Glycerol kinase deficiency, 307030	300474	46	99	88	66
GLA	Fabry disease, 301500 Fabry disease, cardiac variant, 301500	300644	101	100	100	100
GLB1	GM1-gangliosidosis, type I, 230500 GM1-gangliosidosis, type II, 230600 GM1-gangliosidosis, type III, 230650 Mucopolysaccharidosis type IVB (Morquio), 253010	611458	105	100	100	98
GLDC	Glycine encephalopathy, 605899	238300	67	100	98	91
GLE1	Congenital arthrogryposis with anterior horn cell disease, 611890 Lethal congenital contracture syndrome 1, 253310	603371	73	100	100	96
GLI2	Culler-Jones syndrome, 615849 Holoprosencephaly 9, 610829	165230	163	100	100	99

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GLI3	Greig cephalopolysyndactyl syndrome, 175700 {Hypothalamic hamartomas, somatic}, 241800 Pallister-Hall syndrome, 146510 Polydactyly, postaxial, types A1 and B, 174200 Polydactyly, preaxial, type IV, 174700	165240	110	100	100	99
GLIS2	Nephronophthisis 7, 611498	608539	117	100	100	100
GLIS3	Diabetes mellitus, neonatal, with congenital hypothyroidism, 610199	610192	113	100	100	100
GLMN	Glomuvenous malformations, 138000	601749	69	100	98	88
GLRA1	Hyperekplexia 1, 149400	138491	85	100	100	96
GLRB	Hyperekplexia 2, 614619	138492	63	100	100	98
GLRX5	Anemia, sideroblastic, 3, pyridoxine-refractory, 616860 Spasticity, childhood-onset, with hyperglycinemia, 616859	609588	68	100	91	81
GLUD1	Hyperinsulinism-hyperammonemia syndrome, 606762	138130	99	100	97	91
GLUL	Glutamine deficiency, congenital, 610015	138290	92	100	100	100
GLYCTK	D-glyceric aciduria, 220120	610516	136	100	100	100
GM2A	GM2-gangliosidosis, AB variant, 272750	613109	94	100	100	100
GMPPA	Alacrima, achalasia, and mental retardation syndrome, 615510	615495	114	100	100	100
GMPPB	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	615320	189	100	100	100
GMPS	No OMIM phenotype	600358	74	100	99	95
GNA11	Hypocalcemia 2, 615361 Hypocalciuric hypercalcemia, type II, 145981	139313	119	100	100	100
GNA12	Pituitary ACTH-secreting adenoma Ventricular tachycardia, idiopathic, 192605	139360	125	100	100	100
GNA3	Auriculocondylar syndrome 1, 602483	139370	64	100	100	97
GNAL	Dystonia 25, 615073	139312	71	100	100	94
GNAO1	Epileptic encephalopathy, early infantile, 17, 615473 Neurodevelopmental disorder with involuntary movements, 617493	139311	98	100	100	99
GNAQ	Capillary malformations, congenital, 1, somatic, mosaic, 163000 Sturge-Weber syndrome, somatic, mosaic, 185300	600998	94	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GNAS	ACTH-independent macronodular adrenal hyperplasia, 219080 McCune-Albright syndrome, somatic, mosaic, 174800 Osseous heteroplasia, progressive, 166350 Pituitary adenoma 3, multiple types, somatic, 617686 Pseudohypoparathyroidism Ia, 103580 Pseudohypoparathyroidism Ib, 603233 Pseudohypoparathyroidism Ic, 612462 Pseudopseudohypoparathyroidism, 612463	139320	180	100	100	97
GNAT1	Night blindness, congenital stationary 3, 610444 Night blindness, congenital stationary, type 1G, 616389	139330	166	100	100	100
GNAT2	Achromatopsia 4, 613856	139340	72	100	100	98
GNB4	Charcot-Marie-Tooth disease, dominant intermediate F, 615185	610863	76	100	100	96
GNE	Nonaka myopathy, 605820 Sialuria, 269921	603824	93	100	100	99
GNMT	Glycine N-methyltransferase deficiency, 606664	606628	105	100	100	100
GNPAT	Rhizomelic chondrodysplasia punctata, type 2, 222765	602744	78	100	100	96
GNPTAB	Mucolipidosis II alpha/beta, 252500 Mucolipidosis III alpha/beta, 252600	607840	63	100	99	95
GNPTG	Mucolipidosis III gamma, 252605	607838	153	100	99	94
GNRH1	?Hypogonadotropic hypogonadism 12 with or without anosmia, 614841	152760	55	100	100	100
GNRHR	Hypogonadotropic hypogonadism 7 without anosmia, 146110	138850	98	100	100	99
GNS	Mucopolysaccharidosis type IID, 252940	607664	68	100	100	98
GOLGA5	No OMIM phenotype	606918	54	100	99	95
GORAB	Geroderma osteodysplasticum, 231070	607983	66	100	100	98
GOSR2	Epilepsy, progressive myoclonic 6, 614018	604027	66	100	95	90
GOT1	Aspartate aminotransferase, serum level of, QTL1, 614419	138180	88	100	100	100
GP1BA	Bernard-Soulier syndrome, type A1 (recessive), 231200 Bernard-Soulier syndrome, type A2 (dominant), 153670 {Nonarteritic anterior ischemic optic neuropathy, susceptibility to}, 258660 von Willebrand disease, platelet-type, 177820	606672	124	100	100	98
GP1BB	Bernard-Soulier syndrome, type B, 231200 Giant platelet disorder, isolated, 231200	138720	65	100	99	84
GP6	Bleeding disorder, platelet-type, 11, 614201	605546	93	100	100	99
GP9	Bernard-Soulier syndrome, type C, 231200	173515	187	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GPC3	Simpson-Golabi-Behmel syndrome, type 1, 312870 Wilms tumor, somatic, 194070	300037	53	100	99	91
GPC6	Omodyplasia 1, 258315	604404	76	100	100	97
GPD1	Hypertriglyceridemia, transient infantile, 614480	138420	80	100	100	100
GPD1L	Brugada syndrome 2, 611777	611778	113	100	100	100
GPHN	Molybdenum cofactor deficiency C, 615501	603930	68	100	100	98
GPI	Hemolytic anemia, nonspherocytic, due to glucose phosphate isomerase deficiency, 613470	172400	130	100	100	100
GPR143	Nystagmus 6, congenital, 300814 Ocular albinism, type I, Nettleship-Falls type, 300500	300808	43	100	89	76
GPR179	Night blindness, congenital stationary (complete), 1E, 614565	614515	135	100	100	100
GPSM2	Chudley-McCullough syndrome, 604213	609245	93	100	100	96
GRHL2	Corneal dystrophy, posterior polymorphous, 4, 618031 Deafness 28, 608641 Ectodermal dysplasia/short stature syndrome, 616029	608576	81	100	100	99
GRHL3	Van der Woude syndrome 2, 606713	608317	121	100	100	100
GRHPR	Hyperoxaluria, primary, type II, 260000	604296	96	100	100	100
GRIA3	Mental retardation 94, 300699	305915	51	100	98	88
GRIK2	Mental retardation, 6, 611092	138244	94	100	100	98
GRIN1	Neurodevelopmental disorder with or without hyperkinetic movements and seizures, 614254 Neurodevelopmental disorder with or without hyperkinetic movements and seizures, 617820	138249	132	100	100	99
GRIN2A	Epilepsy, focal, with speech disorder and with or without mental retardation, 245570	138253	103	100	100	100
GRIN2B	Epileptic encephalopathy, early infantile, 27, 616139 Mental retardation 6, 613970	138252	122	100	100	99
GRIP1	Fraser syndrome 3, 617667	604597	84	100	100	99
GRK1	Oguchi disease-2, 613411	180381	114	100	100	99
GRM1	Spinocerebellar ataxia 44, 617691 Spinocerebellar ataxia 13, 614831	604473	136	100	100	99
GRM6	Night blindness, congenital stationary (complete), 1B, 257270	604096	127	100	95	90
GRN	Aphasia, primary progressive, 607485 Ceroid lipofuscinosi, neuronal, 11, 614706 Frontotemporal lobar degeneration with ubiquitin-positive inclusions, 607485	138945	155	100	100	100
GRXCR1	Deafness 25, 613285	613283	100	100	100	99
GSC	Short stature, auditory canal atresia, mandibular hypoplasia, skeletal abnormalities, 602471	138890	131	100	100	100
GSDME	Deafness 5, 600994	608798	94	100	100	100
GSN	Amyloidosis, Finnish type, 105120	137350	95	100	100	99

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
GSS	Glutathione synthetase deficiency, 266130 Hemolytic anemia due to glutathione synthetase deficiency, 231900	601002	89	100	100	99
GTF2H5	Trichothiodystrophy 3, photosensitive, 616395	608780	53	100	100	100
GTPBP3	Combined oxidative phosphorylation deficiency 23, 616198	608536	148	100	100	100
GUCA1A	Cone dystrophy-3, 602093 Cone-rod dystrophy 14, 602093	600364	108	100	100	100
GUCA1B	Retinitis pigmentosa 48, 613827	602275	156	100	100	100
GUCY1A1	Moyamoya 6 with achalasia, 615750	139396	76	100	99	96
GUCY2C	Diarrhea 6, 614616 Meconium ileus, 614665	601330	80	100	100	97
GUCY2D	?Choroidal dystrophy, central areolar 1, 215500 Cone-rod dystrophy 6, 601777 Leber congenital amaurosis 1, 204000 Night blindness, congenital stationary, type 1I, 618555	600179	112	100	100	99
GUCY2F	No OMIM phenotype	300041	44	100	96	84
GUSB	Mucopolysaccharidosis VII, 253220	611499	106	100	100	97
GYG1	?Glycogen storage disease XV, 613507 Polyglucosan body myopathy 2, 616199	603942	60	100	100	96
GYS1	Glycogen storage disease 0, muscle, 611556	138570	116	100	100	100
GYS2	Glycogen storage disease 0, liver, 240600	138571	67	100	100	98
H19	Beckwith-Wiedemann syndrome, 130650 Silver-Russell syndrome, 180860 Wilms tumor 2, 194071	103280	No coverage data			
H6PD	Cortisone reductase deficiency 1, 604931	138090	170	100	100	100
HADH	3-hydroxyacyl-CoA dehydrogenase deficiency, 231530 Hyperinsulinemic hypoglycemia, familial, 4, 609975	601609	77	100	100	100
HADHA	Fatty liver, acute, of pregnancy, 609016 HELLP syndrome, maternal, of pregnancy, 609016 LCHAD deficiency, 609016 Trifunctional protein deficiency, 609015	600890	99	100	100	97
HADHB	Trifunctional protein deficiency, 609015	143450	66	100	100	97
HAMP	Hemochromatosis, type 2B, 613313	606464	161	100	100	100
HAND2	No OMIM phenotype	602407	99	100	100	100
HARS1	Charcot-Marie-Tooth disease, axonal, type 2W, 616625 Usher syndrome type 3B, 614504	142810	104	100	100	99

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HARS2	?Perrault syndrome 2, 614926	600783	76	100	100	99
HAX1	Neutropenia, severe congenital 3, 610738	605998	99	100	100	99
HBA1	Erythrocytosis, 7, 617981 Heinz body anemias, alpha-, 140700 Hemoglobin H disease, nondeletional, 613978 Methemoglobinemia, alpha type, 617973 Thalassemias, alpha-, 604131	141800	215	100	100	100
HBA2	Erythrocytosis 7, 617981 Heinz body anemia, 140700 Hemoglobin H disease, deletional and nondeletional, 613978 Thalassemia, alpha-, 604131	141850	113	76	76	75
HBB	Delta-beta thalassemia, 141749 Erythrocytosis 6, 617980 Heinz body anemia, 140700 Hereditary persistence of fetal hemoglobin, 141749 {Malaria, resistance to}, 611162 Methmoglobinemia, beta type, 617971 Sickle cell anemia, 603903 Thalassemia, beta, 613985 Thalassemia-beta, dominant inclusion-body, 603902	141900	125	100	100	100
HBD	Thalassemia due to Hb Lepore Thalassemia, delta-	142000	144	100	100	100
HBG1	Fetal hemoglobin quantitative trait locus 1, 141749	142200	49	55	43	30
HBG2	Cyanosis, transient neonatal, 613977 Fetal hemoglobin quantitative trait locus 1, 141749	142250	191	100	100	100
HCCS	Linear skin defects with multiple congenital anomalies 1, 309801	300056	45	100	99	84
HCFC1	Mental retardation 3 (methylmalonic aciduria and homocysteinemia, cblX type), 309541	300019	79	100	97	92
HCN4	Brugada syndrome 8, 613123 Sick sinus syndrome 2, 163800	605206	103	100	100	99
HCRT	?Narcolepsy 1, 161400	602358	139	100	100	100
HDAC4	No OMIM phenotype	605314	112	100	100	100
HDAC6	?Chondrodysplasia with platyspondyly, distinctive brachydactyly, hydrocephaly, and microphthalmia, 300863	300272	85	100	100	97
HDAC8	Cornelia de Lange syndrome 5, 300882	300269	47	100	100	91

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
HEPACAM	Megalencephalic leukoencephalopathy with subcortical cysts 2A, 613925 Megalencephalic leukoencephalopathy with subcortical cysts 2B, remitting, with or without mental retardation, 613926	611642	97	100	100	99
HERC2	Mental retardation 38, 615516 [Skin/hair/eye pigmentation 1, blond/brown hair], 227220 [Skin/hair/eye pigmentation 1, blue/nonblue eyes], 227220	605837	102	100	99	96
HES7	Spondylocostal dysostosis 4, 613686	608059	61	95	86	76
HESX1	Growth hormone deficiency with pituitary anomalies, 182230 Pituitary hormone deficiency, combined, 5, 182230 Septooptic dysplasia, 182230	601802	71	100	100	95
HEXA	GM2-gangliosidosis, several forms, 272800 [Hex A pseudodeficiency], 272800 Tay-Sachs disease, 272800	606869	86	100	100	98
HEXB	Sandhoff disease, infantile, juvenile, and adult forms, 268800	606873	119	100	100	97
HEY2	No OMIM phenotype	604674	160	100	99	96
HFE	{Alzheimer disease, susceptibility to}, 104300 Hemochromatosis, 235200 {Microvascular complications of diabetes 7}, 612635 {Porphyria cutanea tarda, susceptibility to}, 176100 {Porphyria variegata, susceptibility to}, 176200 [Transferrin serum level QTL2], 614193	613609	97	100	100	100
HFM1	Premature ovarian failure 9, 615724	615684	63	100	97	81
HGD	Alkaptonuria, 203500	607474	63	100	100	96
HGF	Deafness 39, 608265	142409	73	100	100	96
HGSNAT	Mucopolysaccharidosis type IIIC (Sanfilippo C), 252930 Retinitis pigmentosa 73, 616544	610453	70	94	94	92
HIBCH	3-hydroxyisobutyryl-CoA hydrolase deficiency, 250620	610690	51	100	94	75
HINT1	Neuromyotonia and axonal neuropathy, 137200	601314	86	100	100	100
HJV	Hemochromatosis, type 2A, 602390	608374	117	100	100	100
HK1	Hemolytic anemia due to hexokinase deficiency, 235700 Neurodevelopmental disorder with visual defects and brain anomalies, 618547 Neuropathy, hereditary motor and sensory, Russe type, 605285 Retinitis pigmentosa 79, 617460	142600	102	100	100	99
HLCS	Holocarboxylase synthetase deficiency, 253270	609018	98	100	100	99

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HMBS	Porphyria, acute intermittent, 176000 Porphyria, acute intermittent, nonerythroid variant, 176000	609806	88	100	100	100
HMGCL	HMG-CoA lyase deficiency, 246450	613898	94	100	100	100
HMGCS2	HMG-CoA synthase-2 deficiency, 605911	600234	78	100	100	99
HMOX1	Heme oxygenase-1 deficiency, 614034 {Pulmonary disease, chronic obstructive, susceptibility to}, 606963	141250	112	100	100	99
HMX1	Oculoauricular syndrome, 612109	142992	38	94	72	49
HNF1A	Diabetes mellitus, insulin-dependent, 20, 612520 {Diabetes mellitus, insulin-dependent}, 222100 {Diabetes mellitus, noninsulin-dependent, 2}, 125853 Hepatic adenoma, somatic, 142330 MODY, type III, 600496 Renal cell carcinoma, 144700	142410	157	100	100	100
HNF1B	Diabetes mellitus, noninsulin-dependent, 125853 {Renal cell carcinoma}, 144700 Renal cysts and diabetes syndrome, 137920	189907	112	100	100	100
HNF4A	{Diabetes mellitus, noninsulin-dependent}, 125853 Fanconi renotubular syndrome 4, with maturity-onset diabetes of the young, 616026 MODY, type I, 125850	600281	116	100	100	98
HNRNPA1	Amyotrophic lateral sclerosis 20, 615426 ?Inclusion body myopathy with early-onset Paget disease without frontotemporal dementia 3, 615424	164017	61	100	99	95
HOGA1	Hyperoxaluria, primary, type III, 613616	613597	122	100	100	100
HOXA1	Athabaskan brainstem dysgenesis syndrome, 601536 Bosley-Salih-Alorainy syndrome, 601536	142955	126	100	100	100
HOXA11	Radio-ulnar synostosis with amegakaryocytic thrombocytopenia 1, 605432	142958	72	100	100	100
HOXA13	?Guttmacher syndrome, 176305 Hand-foot-uterus syndrome, 140000	142959	80	86	78	74
HOXB1	Facial paresis, hereditary congenital, 3, 614744	142968	115	100	100	100
HOXC13	Ectodermal dysplasia 9, hair/nail type, 614931	142976	180	100	100	100
HOXD10	Charcot-Marie-Tooth disease, foot deformity of, 192950 Vertical talus, congenital, 192950	142984	139	100	100	100

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HOXD13	Brachydactyly, type D, 113200 Brachydactyly, type E, 113300 ?Brachydactyly-syndactyly syndrome, 610713 Syndactyly, type V, 186300 Synpolydactyly 1, 186000	142989	137	100	100	100
HPD	Hawkinsinuria, 140350 Tyrosinemia, type III, 276710	609695	107	100	100	99
HPGD	Cranoosteopathia, 259100 Digital clubbing, isolated congenital, 119900 Hypertrophic osteoarthropathy, primary 1, 259100	601688	73	100	100	98
HPRT1	HPRT-related gout, 300323 Lesch-Nyhan syndrome, 300322	308000	43	100	99	81
HPS1	Hermansky-Pudlak syndrome 1, 203300	604982	103	100	100	97
HPS3	Hermansky-Pudlak syndrome 3, 614072	606118	66	100	100	95
HPS4	Hermansky-Pudlak syndrome 4, 614073	606682	103	100	100	98
HPS5	Hermansky-Pudlak syndrome 5, 614074	607521	68	100	100	97
HPS6	Hermansky-Pudlak syndrome 6, 614075	607522	130	100	100	93
HPSE2	Urofacial syndrome 1, 236730	613469	73	100	98	92
HR	Alopecia universalis, 203655 Atrichia with papular lesions, 209500 Hypotrichosis 4, 146550	602302	111	100	99	97
HRAS	Bladder cancer, somatic, 109800 Congenital myopathy with excess of muscle spindles, 218040 Costello syndrome, 218040 Nevus sebaceous or woolly hair nevus, somatic, 162900 Schimmelpenning-Feuerstein-Mims syndrome, somatic mosaic, 163200 Spitz nevus or nevus spilus, somatic, 137550 Thyroid carcinoma, follicular, somatic, 188470	190020	179	100	100	100
HRG	Thrombophilia due to HRG deficiency, 613116 Thrombophilia due to elevated HRG, 613116	142640	94	100	100	99
HSD11B1	Cortisone reductase deficiency 2, 614662	600713	68	100	100	99
HSD11B2	Apparent mineralocorticoid excess, 218030	614232	147	92	86	83
HSD17B10	HSD10 mitochondrial disease, 300438	300256	76	100	100	100
HSD17B3	Pseudohermaphroditism, male, with gynecomastia, 264300	605573	67	100	99	94

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HSD17B4	D-bifunctional protein deficiency, 261515 Perrault syndrome 1, 233400	601860	62	100	98	93
HSD3B2	Adrenal hyperplasia, congenital, due to 3-beta-hydroxysteroid dehydrogenase 2 deficiency, 201810	613890	143	100	100	100
HSD3B7	Bile acid synthesis defect, congenital, 1, 607765	607764	115	100	100	100
HSF4	Cataract 5, multiple types, 116800	602438	114	100	100	97
HSPB1	Charcot-Marie-Tooth disease, axonal, type 2F, 606595 Neuropathy, distal hereditary motor, type IIB, 608634	602195	116	100	100	98
HSPB3	?Neuronopathy, distal hereditary motor, type IIC, 613376	604624	125	100	100	100
HSPB8	Charcot-Marie-Tooth disease, axonal, type 2L, 608673 Neuropathy, distal hereditary motor, type IIA, 158590	608014	220	100	100	98
HSPD1	Leukodystrophy, hypomyelinating, 4, 612233 Spastic paraplegia 13, 605280	118190	65	100	99	85
HSPG2	Dyssegmental dysplasia, Silverman-Handmaker type, 224410 Schwartz-Jampel syndrome, type 1, 255800	142461	105	99	99	99
HTR1A	Periodic fever, menstrual cycle dependent, 614674	109760	147	100	100	100
HTRA1	CARASIL syndrome, 600142	602194	93	96	87	83
	Cerebral arteriopathy, with subcortical infarcts and leukoencephalopathy, type 2, 616779					
	{Macular degeneration, age-related, 7}, 610149					
	{Macular degeneration, age-related, neovascular type}, 610149					
HTRA2	3-methylglutaconic aciduria, type VIII, 617248 {Parkinson disease 13}, 610297	606441	143	100	100	98
HTT	Huntington disease, 143100 Lopes-Macié-Rodan syndrome, 617435	613004	85	99	99	97
HUWE1	Mental retardation syndromic, Turner type, 309590	300697	53	100	96	83
HYAL1	?Mucopolysaccharidosis type IX, 601492	607071	103	100	100	100
HYDIN	Ciliary dyskinesia, primary, 5, 608647	610812	46	82	69	57
HYLS1	Hydrocephalus syndrome, 236680	610693	74	100	100	100
ICK	Endocrine-cerebroosteodysplasia, 612651 {Epilepsy, juvenile myoclonic, susceptibility to, 10}, 617924	612325	59	100	99	91
ICOS	Immunodeficiency, common variable, 1, 607594	604558	57	100	100	95
IDH2	D-2-hydroxyglutaric aciduria 2, 613657	147650	105	100	100	100
IDH3B	Retinitis pigmentosa 46, 612572	604526	117	100	100	100
IDS	Mucopolysaccharidosis II, 309900	300823	74	100	99	95

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IDUA	Mucopolysaccharidosis Ih, 607014 Mucopolysaccharidosis Ih/s, 607015 Mucopolysaccharidosis Is, 607016	252800	138	100	97	91
IER3IP1	Microcephaly, epilepsy, and diabetes syndrome, 614231	609382	83	100	100	77
IFITM5	Osteogenesis imperfecta, type V, 610967	614757	131	100	100	100
IFNGR1	{H. pylori infection, susceptibility to}, 600263 {Hepatitis B virus infection, susceptibility to}, 610424 Immunodeficiency 27A, mycobacteriosis, AR, 209950 Immunodeficiency 27B, mycobacteriosis, AD, 615978 {Tuberculosis infection, protection against}, 607948 {Tuberculosis, susceptibility to}, 607948	107470	68	100	100	95
IFT122	Cranioectodermal dysplasia 1, 218330	606045	111	100	100	99
IFT140	Retinitis pigmentosa 80, 617781 Short-rib thoracic dysplasia 9 with or without polydactyly, 266920	614620	113	100	100	98
IFT172	Retinitis pigmentosa 71, 616394 Short-rib thoracic dysplasia 10 with or without polydactyly, 615630	607386	75	100	100	96
IFT43	?Cranioectodermal dysplasia 3, 614099 ?Retinitis pigmentosa 81, 617871 Short-rib thoracic dysplasia 18 with polydactyly, 617866	614068	75	100	100	100
IFT80	Short-rib thoracic dysplasia 2 with or without polydactyly, 611263	611177	62	100	98	87
IGBP1	Corpus callosum, agenesis of, with mental retardation, ocular coloboma and micrognathia, 300472	300139	69	100	100	90
IGF1	Growth retardation with deafness and mental retardation due to IGF1 deficiency, 608747	147440	71	100	100	100
IGF1R	Insulin-like growth factor I, resistance to, 270450	147370	120	100	100	100
IGF2R	Hepatocellular carcinoma, somatic, 114550	147280	101	100	99	97
IGFALS	Acid-labile subunit, deficiency of, 615961	601489	103	100	100	100
IGFBP7	Retinal arterial macroaneurysm with supravalvular pulmonic stenosis, 614224	602867	60	100	95	87
IGHMBP2	Charcot-Marie-Tooth disease, axonal, type 2S, 616155 Neuronopathy, distal hereditary motor, type VI, 604320	600502	101	100	100	99
IGLL1	Agammaglobulinemia 2, 613500	146770	132	100	100	100
IGSF1	Hypothyroidism, central, and testicular enlargement, 300888	300137	62	100	99	93
IHH	Acrocapitofemoral dysplasia, 607778 Brachydactyly, type A1, 112500	600726	133	100	100	100
IKBKB	Immunodeficiency 15A, 618204 Immunodeficiency 15B, 615592	603258	86	100	100	97

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IKBKG	Ectodermal dysplasia and immunodeficiency 1, 300291 Ectodermal, dysplasia, anhidrotic, lymphedema and immunodeficiency, 300301 Immunodeficiency 33, 300636 Immunodeficiency, isolated, 300584 Incontinentia pigmenti, 308300 Invasive pneumococcal disease, recurrent isolated, 2, 300640	300248	24	38	33	31
IKZF1	Immunodeficiency, common variable, 13, 616873	603023	166	100	100	100
IKZF5	No OMIM phenotype	606238	85	100	100	99
IL10RA	Inflammatory bowel disease 28, early onset, 613148	146933	125	100	100	99
IL10RB	{Hepatitis B virus, susceptibility to}, 610424 Inflammatory bowel disease 25, early onset, 612567	123889	69	100	100	97
IL11RA	Craniosynostosis and dental anomalies, 614188	600939	103	100	100	100
IL17F	?Candidiasis, familial, 6, 613956	606496	79	100	100	100
IL17RA	Immunodeficiency 51, 613953	605461	125	100	100	100
IL17RD	Hypogonadotropic hypogonadism 18 with or without anosmia, 615267	606807	102	100	100	99
IL1RAPL1	Mental retardation 21/34, 300143	300206	52	100	98	88
IL1RN	{Gastric cancer risk after H. pylori infection}, 137215 Interleukin 1 receptor antagonist deficiency, 612852 {Microvascular complications of diabetes 4}, 612628	147679	76	100	100	90
IL21R	[IgE, elevated level of], 147050 Immunodeficiency 56, 615207	605383	113	100	100	100
IL2RA	{Diabetes, mellitus, insulin-dependent, susceptibility to, 10}, 601942 Immunodeficiency 41 with lymphoproliferation and autoimmunity, 606367	147730	96	100	100	100
IL2RG	Combined immunodeficiency, moderate, 312863 Severe combined immunodeficiency, 300400	308380	54	100	99	91
IL31RA	?Amyloidosis, primary localized cutaneous, 2, 613955	609510	74	100	100	96
IL36RN	Psoriasis 14, pustular, 614204	605507	82	100	100	100
IL7R	Severe combined immunodeficiency, T-cell negative, B-cell/natural killer cell-positive type, 608971	146661	60	100	100	99
ILDR1	Deafness 42, 609646	609739	95	100	100	97
IMPAD1	Chondrodysplasia with joint dislocations, GPAPP type, 614078	614010	146	100	100	97
IMPDH1	Leber congenital amaurosis 11, 613837 Retinitis pigmentosa 10, 180105	146690	99	100	96	91
IMPG2	Macular dystrophy, vitelliform, 5, 616152 Retinitis pigmentosa 56, 613581	607056	68	100	100	97

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INF2	Charcot-Marie-Tooth disease, dominant intermediate E, 614455 Glomerulosclerosis, focal segmental, 5, 613237	610982	117	96	94	92
ING1	Squamous cell carcinoma, head and neck, somatic, 275355	601566	156	100	100	100
INPP5E	Joubert syndrome 1, 213300 Mental retardation, truncal obesity, retinal dystrophy, and micropenis, 610156	613037	113	100	100	97
INPPL1	Opsismodysplasia, 258480	600829	111	100	100	99
INS	Diabetes mellitus, insulin-dependent, 2, 125852	176730	106	100	100	100
	Diabetes mellitus, permanent neonatal, 606176					
	Hyperproinsulinemia, 616214					
	Maturity-onset diabetes of the young, type 10, 613370					
INSL3	Cryptorchidism, 219050	146738	69	93	78	78
INSR	Diabetes mellitus, insulin-resistant, with acanthosis nigricans, 610549	147670	115	100	98	95
	Hyperinsulinemic hypoglycemia, familial, 5, 609968					
	Leprechaunism, 246200					
	Rabson-Mendenhall syndrome, 262190					
INVS	Nephronophthisis 2, infantile, 602088	243305	81	100	100	97
IQCB1	Senior-Loken syndrome 5, 609254	609237	54	100	94	78
IQSEC2	Mental retardation 1/78, 309530	300522	66	98	92	85
IRAK4	IRAK4 deficiency, 607676	606883	72	100	99	91
	Invasive pneumococcal disease, recurrent isolated, 1, 610799					
IRF1	Gastric cancer, somatic, 613659	147575	107	100	100	98
	Myelodysplastic syndrome, preleukemic					
	Myelogenous leukemia, acute					
	Nonsmall cell lung cancer, somatic, 211980					
IRF4	[Skin/hair/eye pigmentation, variation in, 8], 611724	601900	147	100	100	100
IRF6	{Orofacial cleft 6}, 608864	607199	121	100	100	100
	Popliteal pterygium syndrome 1, 119500					
	van der Woude syndrome, 119300					
IRF8	Immunodeficiency 32A, mycobacteriosis, 614893	601565	136	100	100	99
	Immunodeficiency 32B, monocyte and dendritic cell deficiency, 226990					
IRGM	{Inflammatory bowel disease (Crohn disease) 19}, 612278	608212	54	78	78	78
	{Mycobacterium tuberculosis, protection against}, 607948					
IRX5	Hamamy syndrome, 611174	606195	99	100	100	100
ISCU	Myopathy with lactic acidosis, hereditary, 255125	611911	75	100	100	100
ITCH	Autoimmune disease, multisystem, with facial dysmorphism, 613385	606409	61	96	95	90

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ITGA2	?Glycoprotein Ia deficiency, 614200	192974	74	100	100	97
ITGA2B	Bleeding disorder, platelet-type, 16, 187800 Glanzmann thrombasthenia, 273800 Thrombocytopenia, neonatal alloimmune, BAK antigen related	607759	125	100	100	99
ITGA3	Interstitial lung disease, nephrotic syndrome, and epidermolysis bullosa, congenital, 614748	605025	137	100	100	100
ITGA6	Epidermolysis bullosa, junctional, with pyloric stenosis, 226730	147556	84	100	100	95
ITGA7	Muscular dystrophy, congenital, due to ITGA7 deficiency, 613204	600536	117	100	97	96
ITGA8	Renal hypodysplasia/aplasia 1, 191830	604063	60	100	99	94
ITGB2	Leukocyte adhesion deficiency, 116920	600065	140	100	100	100
ITGB3	Bleeding disorder, platelet-type, 16, 187800 Glanzmann thrombasthenia, 273800 {Myocardial infarction, susceptibility to}, 608446 Purpura, posttransfusion Thrombocytopenia, neonatal alloimmune	173470	87	100	100	98
ITGB4	Epidermolysis bullosa of hands and feet, 131800 Epidermolysis bullosa, junctional, non-Herlitz type, 226650 Epidermolysis bullosa, junctional, with pyloric atresia, 226730	147557	135	99	98	97
ITK	Lymphoproliferative syndrome 1, 613011	186973	76	100	100	95
ITM2B	Dementia, familial British, 176500 Dementia, familial Danish, 117300 ?Retinal dystrophy with inner retinal dysfunction and ganglion cell abnormalities, 616079	603904	79	100	100	97
ITPR1	Gillespie syndrome, 206700 Spinocerebellar ataxia 15, 606658 Spinocerebellar ataxia 29, congenital nonprogressive, 117360	147265	90	100	100	98
IVD	Isovaleric acidemia, 243500	607036	107	100	100	98
IYD	Thyroid dyshormonogenesis 4, 274800	612025	104	100	100	100
JAG1	Alagille syndrome 1, 118450 ?Deafness, congenital heart defects, and posterior embryotoxon, 617992 Tetralogy of Fallot, 187500	601920	96	100	100	96
JAK2	{Budd-Chiari syndrome, somatic}, 600880 Erythrocytosis, somatic, 133100 Leukemia, acute myeloid, somatic, 601626 Myelofibrosis, somatic, 254450 Polycythemia vera, somatic, 263300 Thrombocythemia 3, 614521	147796	63	100	99	93

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JAK3	SCID, T-negative/B-positive type, 600802	600173	104	100	100	99
JAM3	Hemorrhagic destruction of the brain, subependymal calcification, and cataracts, 613730	606871	70	100	100	97
JPH2	Cardiomyopathy, hypertrophic, 17, 613873	605267	109	100	100	98
JPH3	Huntington disease-like 2, 606438	605268	180	100	100	99
JUP	Arrhythmogenic right ventricular dysplasia 12, 611528 Naxos disease, 601214	173325	106	100	100	98
KANK1	Cerebral palsy, spastic quadriplegic, 2, 612900	607704	105	100	100	99
KANSL1	Koolen-De Vries syndrome, 610443	612452	85	100	100	97
KARS1	?Charcot-Marie-Tooth disease, recessive intermediate, B, 613641 Deafness 89, 613916	601421	126	100	100	97
KAT6A	Mental retardation 32, 616268	601408	95	100	100	99
KAT6B	Genitopatellar syndrome, 606170 SBBYSS syndrome, 603736	605880	107	100	100	98
KBTBD13	Nemaline myopathy 6, 609273	613727	177	100	100	100
KCNA1	Episodic ataxia/myokymia syndrome, 160120	176260	126	100	100	100
KCNA5	Atrial fibrillation, familial, 7, 612240	176267	147	100	100	99
KCNC3	Spinocerebellar ataxia 13, 605259	176264	91	94	78	64
KCND3	Brugada syndrome 9, 616399 Spinocerebellar ataxia 19, 607346	605411	167	100	100	99
KCNE1	Jervell and Lange-Nielsen syndrome 2, 612347 Long QT syndrome 5, 613695	176261	252	100	100	100
KCNE2	Atrial fibrillation, familial, 4, 611493 Long QT syndrome 6, 613693	603796	81	100	100	100
KCNE3	?Brugada syndrome 6, 613119	604433	103	100	100	100
KCNH2	Long QT syndrome 2, 613688 {Long QT syndrome 2, acquired, susceptibility to}, 613688 Short QT syndrome 1, 609620	152427	121	100	100	99
KCNJ1	Bartter syndrome, type 2, 241200	600359	60	100	100	100
KCNJ10	Enlarged vestibular aqueduct, digenic, 600791 SESAME syndrome, 612780	602208	171	100	100	100
KCNJ11	Diabetes mellitus, transient neonatal, 3, 610582 {Diabetes mellitus, type 2, susceptibility to}, 125853 Diabetes, permanent neonatal, with or without neurologic features, 606176 Hyperinsulinemic hypoglycemia, familial, 2, 601820 Maturity-onset diabetes of the young, type 13, 616329	600937	128	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
KCNJ13	Leber congenital amaurosis 16, 614186 Snowflake vitreoretinal degeneration, 193230	603208	73	100	100	99
KCNJ2	Andersen syndrome, 170390 Atrial fibrillation, familial, 9, 613980 Short QT syndrome 3, 609622	600681	96	100	100	100
KCNJ5	Hyperaldosteronism, familial, type III, 613677 Long QT syndrome 13, 613485	600734	148	100	100	100
KCNK3	Pulmonary hypertension, primary, 4, 615344	603220	185	100	100	99
KCNK9	Birk-Barel mental retardation dysmorphism syndrome, 612292	605874	129	100	100	100
KCNMA1	?Cerebellar atrophy, developmental delay, and seizures, 617643 Paroxysmal nonkinesigenic dyskinesia, 3, with or without generalized epilepsy, 609446	600150	88	100	100	96
KCNQ1	Atrial fibrillation, familial, 3, 607554 Jervell and Lange-Nielsen syndrome, 220400 Long QT syndrome 1, 192500 {Long QT syndrome 1, acquired, susceptibility to}, 192500 Short QT syndrome 2, 609621	607542	160	97	94	93
KCNQ1OT1	Beckwith-Wiedemann syndrome, 130650	604115	No coverage data			
KCNQ2	Epileptic encephalopathy, early infantile, 7, 613720 Myokymia, 121200 Seizures, benign neonatal, 1, 121200	602235	131	100	100	100
KCNQ3	Seizures, benign neonatal, 2, 121201	602232	108	100	100	96
KCNQ4	Deafness 2A, 600101	603537	147	98	95	91
KCNQ5	Mental retardation 46, 617601	607357	85	100	98	95
KCNT1	Epilepsy, nocturnal frontal lobe, 5, 615005 Epileptic encephalopathy, early infantile, 14, 614959	608167	120	100	99	99
KCNV2	Retinal cone dystrophy 3B, 610356	607604	130	100	100	100
KCTD1	Scalp-ear-nipple syndrome, 181270	613420	90	99	98	93
KCTD7	Epilepsy, progressive myoclonic 3, with or without intracellular inclusions, 611726	611725	159	100	100	100
KDM5C	Mental retardation, syndromic, Claes-Jensen type, 300534	314690	88	100	99	96
KDM6A	Kabuki syndrome 2, 300867	300128	52	100	95	80
KDM6B	Neurodevelopmental disorder with coarse facies and mild distal skeletal abnormalities, 618505	611577	132	100	98	95
KDR	Hemangioma, capillary infantile, somatic, 602089 {Hemangioma, capillary infantile, susceptibility to}, 602089	191306	68	100	99	96
KERA	Cornea plana 2, 217300	603288	61	100	100	98
KHDC3L	Hydatidiform mole, recurrent, 2, 614293	611687	144	100	100	100

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KIF11	Microcephaly with or without chorioretinopathy, lymphedema, or mental retardation, 152950	148760	73	100	99	93
KIF1A	Mental retardation 9, 614255 Neuropathy, hereditary sensory, type IIC, 614213 Spastic paraparesis 30, 610357	601255	101	100	100	98
KIF1B	?Charcot-Marie-Tooth disease, type 2A1, 118210 {Neuroblastoma, susceptibility to, 1}, 256700 Pheochromocytoma, 171300	605995	71	100	100	97
KIF1C	Spastic ataxia 2, 611302	603060	141	100	100	100
KIF20A	No OMIM phenotype	605664	74	100	99	95
KIF21A	Fibrosis of extraocular muscles, congenital, 1, 135700 Fibrosis of extraocular muscles, congenital, 3B, 135700	608283	68	100	99	93
KIF22	Spondyloepiphyseal dysplasia with joint laxity, type 2, 603546	603213	158	100	100	100
KIF2A	Cortical dysplasia, complex, with other brain malformations 3, 615411	602591	84	100	99	90
KIF5A	{Amyotrophic lateral sclerosis, susceptibility to, 25}, 617921 Myoclonus, intractable, neonatal, 617235 Spastic paraparesis 10, 604187	602821	90	100	100	96
KIF7	Acrocallosal syndrome, 200990 ?Al-Gazali-Bakalinova syndrome, 607131 ?Hydrocephalus syndrome 2, 614120 Joubert syndrome 12, 200990	611254	107	98	96	93
KIFBP	Goldberg-Shprintzen megacolon syndrome, 609460	609367	76	100	100	98
KIRREL3	No OMIM phenotype	607761	114	100	100	100
KISS1	?Hypogonadotropic hypogonadism 13 with or without anosmia, 614842	603286	127	100	100	100
KISS1R	Hypogonadotropic hypogonadism 8 with or without anosmia, 614837 ?Precocious puberty, central, 1, 176400	604161	165	100	100	100
KIT	Gastrointestinal stromal tumor, familial, 606764 Germ cell tumors, somatic, 273300 Leukemia, acute myeloid, somatic, 601626 Mastocytosis, cutaneous, 154800 Mastocytosis, systemic, somatic, 154800 Piebaldism, 172800	164920	79	100	100	99
KITLG	Deafness 69, unilateral or asymmetric, 616697 Hyperpigmentation with or without hypopigmentation, 145250 [Skin/hair/eye pigmentation 7, blond/brown hair], 611664	184745	78	100	100	97
KL	?Tumoral calcinosis, hyperphosphatemic, familial, 3, 617994	604824	120	99	98	97

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
KLF1	Blood group--Lutheran inhibitor, 111150 Dyserythropoietic anemia, congenital, type IV, 613673 [Hereditary persistence of fetal hemoglobin], 613566	600599	115	100	100	100
KLF11	Maturity-onset diabetes of the young, type VII, 610508	603301	137	100	100	99
KLF6	Gastric cancer, somatic, 613659 Prostate cancer, somatic, 176807	602053	146	100	100	100
KLHDC8B	{Hodgkin lymphoma, susceptibility to}, 236000	613169	90	100	100	100
KLHL10	Spermatogenic failure 11, 615081	608778	85	100	100	100
KLHL3	Pseudohypoaldosteronism, type IID, 614495	605775	86	100	100	97
KLHL40	Nemaline myopathy 8, 615348	615340	123	100	100	99
KLHL41	Nemaline myopathy 9, 615731	607701	77	100	100	100
KLHL7	Cold-induced sweating syndrome 3, 617055 Retinitis pigmentosa 42, 612943	611119	70	100	100	98
KLK4	Amelogenesis imperfecta, type IIA1, 204700	603767	140	100	100	100
KLKB1	Fletcher factor (prekallikrein) deficiency, 612423	229000	74	100	100	96
KLLN	Cowden syndrome 4, 615107	612105	98	100	100	100
KMT2A	Leukemia, myeloid/lymphoid or mixed-lineage, 159555 Wiedemann-Steiner syndrome, 605130	159555	71	100	100	98
KMT2D	Kabuki syndrome 1, 147920	602113	114	100	100	99
KNL1	Microcephaly 4, primary, 604321	609173	60	100	99	95
KPTN	Mental retardation 41, 615637	615620	150	100	100	100
KRAS	Arteriovenous malformation of the brain, somatic, 108010 Bladder cancer, somatic, 109800 Breast cancer, somatic, 114480 Cardiofaciocutaneous syndrome 2, 615278 Gastric cancer, somatic, 137215 Leukemia, acute myeloid, 601626 Lung cancer, somatic, 211980 Noonan syndrome 3, 609942 Oculoectodermal syndrome, somatic, 600268 Pancreatic carcinoma, somatic, 260350 RAS-associated autoimmune leukoproliferative disorder, 614470 Schimmelpenning-Feuerstein-Mims syndrome, somatic mosaic, 163200	190070	83	100	100	87

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KRIT1	Cavernous malformations of CNS and retina, 116860 Cerebral cavernous malformations-1, 116860 Hyperkeratotic cutaneous capillary-venous malformations associated with cerebral capillary malformations, 116860	604214	68	100	100	94
KRT1	Epidermolytic hyperkeratosis, 113800 Ichthyosis histrix, Curth-Macklin type, 146590 Ichthyosis, cyclic, with epidermolytic hyperkeratosis, 607602 Keratosis palmoplantaris striata III, 607654 Palmoplantar keratoderma, epidermolytic, 144200 Palmoplantar keratoderma, nonepidermolytic, 600962	139350	108	100	100	100
KRT10	Epidermolytic hyperkeratosis, 113800 Ichthyosis with confetti, 609165 Ichthyosis, cyclic, with epidermolytic hyperkeratosis, 607602	148080	121	100	100	100
KRT12	Meesmann corneal dystrophy, 122100	601687	94	100	100	100
KRT13	White sponge nevus 2, 615785	148065	126	100	100	100
KRT14	Dermatopathia pigmentosa reticularis, 125595 Epidermolysis bullosa simplex, Dowling-Meara type, 131760 Epidermolysis bullosa simplex, Koebner type, 131900 Epidermolysis bullosa simplex, Weber-Cockayne type, 131800 Epidermolysis bullosa simplex, recessive 1, 601001 Naegeli-Franceschetti-Jadassohn syndrome, 161000	148066	119	100	100	99
KRT16	Pachyonychia congenita 1, 167200 Palmoplantar keratoderma, nonepidermolytic, focal, 613000	148067	149	100	100	100
KRT17	Pachyonychia congenita 2, 167210 Steatocystoma multiplex, 184500	148069	85	100	98	94
KRT18	Cirrhosis, cryptogenic, 215600 {Cirrhosis, noncryptogenic, susceptibility to}, 215600	148070	91	100	100	100
KRT2	Ichthyosis bullosa of Siemens, 146800	600194	129	100	100	100
KRT3	Meesmann corneal dystrophy, 122100	148043	134	100	100	99
KRT4	White sponge nevus 1, 193900	123940	132	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
KRT5	Dowling-Degos disease 1, 179850 Epidermolysis bullosa simplex, Dowling-Meara type, 131760 Epidermolysis bullosa simplex, Koebner type, 131900 Epidermolysis bullosa simplex, Weber-Cockayne type, 131800 Epidermolysis bullosa simplex, recessive 1, 601001 Epidermolysis bullosa simplex-MCR, 609352 Epidermolysis bullosa simplex-MP, 131960	148040	151	100	100	100
KRT6A	Pachyonychia congenita 3, 615726	148041	213	100	100	100
KRT6B	Pachyonychia congenita 4, 615728	148042	214	100	100	100
KRT6C	Palmoplantar keratoderma, nonepidermolytic, focal or diffuse, 615735	612315	164	100	100	100
KRT74	?Ectodermal dysplasia 7, hair/nail type, 614929 ?Hypotrichosis 3, 613981 Woolly hair, 194300	608248	151	100	100	100
KRT8	Cirrhosis, cryptogenic, 215600 {Cirrhosis, noncryptogenic, susceptibility to}, 215600	148060	91	100	100	97
KRT81	Monilethrix, 158000	602153	113	76	72	70
KRT83	Erythrokeratoderma variabilis et progressiva 5, 617756 Monilethrix, 158000	602765	131	100	100	100
KRT85	Ectodermal dysplasia 4, hair/nail type, 602032	602767	124	100	100	99
KRT86	Monilethrix, 158000	601928	126	79	75	73
KRT9	Palmoplantar keratoderma, epidermolytic, 144200	607606	119	100	100	100
L1CAM	CRASH syndrome, 303350 Corpus callosum, partial agenesis of, 304100 Hydrocephalus due to aqueductal stenosis, 307000 Hydrocephalus with Hirschsprung disease, 307000 Hydrocephalus with congenital idiopathic intestinal pseudoobstruction, 307000 MASA syndrome, 303350	308840	101	100	100	100
L2HGDH	L-2-hydroxyglutaric aciduria, 236792	609584	74	100	100	97
LAMA2	Muscular dystrophy, congenital, merosin deficient or partially deficient, 607855 Muscular dystrophy, limb-girdle 23, 618138	156225	71	100	99	96
LAMA3	Epidermolysis bullosa, generalized atrophic benign, 226650 Epidermolysis bullosa, junctional, Herlitz type, 226700 Laryngoonychocutaneous syndrome, 245660	600805	73	100	99	96
LAMA4	Cardiomyopathy, dilated, 1JJ, 615235	600133	71	100	100	97
LAMB1	Lissencephaly 5, 615191	150240	94	100	100	97

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
LAMB2	Nephrotic syndrome, type 5, with or without ocular abnormalities, 614199 Pierson syndrome, 609049	150325	153	100	100	100
LAMB3	Amelogenesis imperfecta, type IA, 104530 Epidermolysis bullosa, junctional, Herlitz type, 226700 Epidermolysis bullosa, junctional, non-Herlitz type, 226650	150310	120	100	100	100
LAMC2	Epidermolysis bullosa, junctional, Herlitz type, 226700 Epidermolysis bullosa, junctional, non-Herlitz type, 226650	150292	106	100	100	99
LAMC3	Cortical malformations, occipital, 614115	604349	130	100	100	99
LAMP2	Danon disease, 300257	309060	44	100	95	77
LAMTOR2	Immunodeficiency due to defect in MAPBP-interacting protein, 610798	610389	108	100	100	100
LARGE1	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 6, 613154 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 6, 608840	603590	97	100	100	99
LARP7	Alazami syndrome, 615071	612026	67	100	99	92
LARS2	?Hydrops, lactic acidosis, and sideroblastic anemia, 617021 Perrault syndrome 4, 615300	604544	79	100	100	99
LBR	Greenberg skeletal dysplasia, 215140 Pelger-Huet anomaly, 169400 Pelger-Huet anomaly with mild skeletal anomalies, 618019 ?Reynolds syndrome, 613471	600024	68	100	100	96
LCA5	Leber congenital amaurosis 5, 604537	611408	74	100	100	99
LCAT	Fish-eye disease, 136120 Norum disease, 245900	606967	130	100	100	97
LCT	Lactase deficiency, congenital, 223000	603202	114	100	100	100
LDB3	Cardiomyopathy, dilated, 1C, with or without LVNC, 601493 Cardiomyopathy, hypertrophic, 24, 601493 Left ventricular noncompaction 3, 601493 Myopathy, myofibrillar, 4, 609452	605906	123	100	100	100
LDHA	Glycogen storage disease XI, 612933	150000	62	100	99	88
LDHB	[Lactate dehydrogenase-B deficiency], 614128	150100	58	100	100	98
LDLR	Hypercholesterolemia, familial, 1, 143890 LDL cholesterol level QTL2, 143890	606945	209	100	100	100
LDLRAP1	Hypercholesterolemia, familial, 4, 603813	605747	121	100	100	100
LEF1	Sebaceous tumors, somatic	153245	74	100	100	100
LEFTY2	No OMIM phenotype	601877	144	100	100	99

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
LEMD3	Buschke-Ollendorff syndrome, 166700 Osteopoikilosis with or without melorheostosis, 166700	607844	84	100	100	96
LEP	Obesity, morbid, due to leptin deficiency, 614962	164160	129	100	100	100
LEPR	Obesity, morbid, due to leptin receptor deficiency, 614963	601007	80	100	100	97
LFNG	Spondylocostal dysostosis 3, 609813	602576	129	86	84	83
LGI1	Epilepsy, familial temporal lobe, 1, 600512	604619	83	100	100	98
LHB	Hypogonadotropic hypogonadism 23 with or without anosmia, 228300	152780	99	100	100	100
LHCGR	Leydig cell adenoma, somatic, with precocious puberty, 176410	152790	71	100	100	94
	Leydig cell hypoplasia with hypergonadotropic hypogonadism, 238320					
	Leydig cell hypoplasia with pseudohermaphroditism, 238320					
	Luteinizing hormone resistance, female, 238320					
	Precocious puberty, male, 176410					
LHFPL5	Deafness 67, 610265	609427	175	100	100	100
LHX3	Pituitary hormone deficiency, combined, 3, 221750	600577	98	100	100	98
LHX4	Pituitary hormone deficiency, combined, 4, 262700	602146	93	100	100	100
LIAS	Hyperglycinemia, lactic acidosis, and seizures, 614462	607031	78	100	100	95
LIFR	Stuve-Wiedemann syndrome/Schwartz-Jampel type 2 syndrome, 601559	151443	61	100	98	89
LIG1	No OMIM phenotype	126391	94	100	100	99
LIG4	LIG4 syndrome, 606593 {Multiple myeloma, resistance to}, 254500	601837	71	100	100	100
LIM2	Cataract 19, multiple types, 615277	154045	120	100	100	100
LINS1	Mental retardation 27, 614340	610350	57	100	100	94
LIPA	Cholesteryl ester storage disease, 278000 Wolman disease, 278000	613497	68	100	100	96
LIPC	{Diabetes mellitus, noninsulin-dependent}, 125853 Hepatic lipase deficiency, 614025 [High density lipoprotein cholesterol level QTL 12], 612797	151670	82	100	100	98
LIPH	Hypotrichosis 7, 604379 Woolly hair 2 with or without hypotrichosis, 604379	607365	55	100	100	95
LIPN	Ichthyosis, congenital 8, 613943	613924	78	100	99	92
LITAF	Charcot-Marie-Tooth disease, type 1C, 601098	603795	89	100	100	100
LMAN1	Combined factor V and VIII deficiency, 227300	601567	72	100	97	86

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
LMBR1	Acheiropody, 200500 Hypoplastic or aplastic tibia with polydactyly, 188740 Laurin-Sandrow syndrome, 135750 Polydactyly, preaxial type II, 174500 Syndactyly, type IV, 186200 Triphalangeal thumb, type I, 174500 Triphalangeal thumb-polysyndactyly syndrome, 174500	605522	54	100	98	86
LMBRD1	Methylmalonic aciduria and homocystinuria, cblF type, 277380	612625	65	100	98	85
LMF1	Lipase deficiency, combined, 246650	611761	129	100	100	98
LMNA	Cardiomyopathy, dilated, 1A, 115200 Charcot-Marie-Tooth disease, type 2B1, 605588 Emery-Dreifuss muscular dystrophy 2, 181350 Emery-Dreifuss muscular dystrophy 3, 616516 Heart-hand syndrome, Slovenian type, 610140 Hutchinson-Gilford progeria, 176670 Lipodystrophy, familial partial, type 2, 151660 Malouf syndrome, 212112 Mandibuloacral dysplasia, 248370 Muscular dystrophy, congenital, 613205 Restrictive dermopathy, lethal, 275210	150330	143	100	100	100
LMNB1	Leukodystrophy, adult-onset, 169500	150340	66	100	100	97
LMOD3	Nemaline myopathy 10, 616165	616112	89	100	100	100
LMX1B	Nail-patella syndrome, 161200	602575	118	100	100	99
LOR	Vohwinkel syndrome with ichthyosis, 604117	152445	83	100	100	100
LOX	Aortic aneurysm, familial thoracic 10, 617168	153455	119	100	100	100
LOXHD1	Deafness 77, 613079	613072	118	100	100	99
LPAR6	Hypotrichosis 8, 278150 Woolly hair 1, with or without hypotrichosis, 278150	609239	87	100	100	100
LPIN1	Myoglobinuria, acute recurrent, 268200	605518	78	100	100	97
LPIN2	Majeed syndrome, 609628	605519	77	100	100	98
LPL	Combined hyperlipidemia, familial, 144250 [High density lipoprotein cholesterol level QTL 11], 238600 Lipoprotein lipase deficiency, 238600	609708	110	100	100	99
	Leukemia, acute myeloid, 601626	600700	97	100	100	100
	Lipoma					

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
LRAT	Leber congenital amaurosis 14, 613341 Retinal dystrophy, early-onset severe, 613341 Retinitis pigmentosa, juvenile, 613341	604863	159	100	100	100
LRBA	Immunodeficiency, common variable, 8, with autoimmunity, 614700	606453	65	100	99	94
LRIG2	Urofacial syndrome 2, 615112	608869	82	100	100	97
LRIT3	Night blindness, congenital stationary (complete), 1F, 615058	615004	78	100	100	100
LRMDA	Albinism, oculocutaneous, type VII, 615179	614537	91	100	95	93
LRP2	Donnai-Barrow syndrome, 222448	600073	69	100	100	96
LRP4	Cenani-Lenz syndactyly syndrome, 212780 ?Myasthenic syndrome, congenital, 17, 616304 Sclerosteosis 2, 614305	604270	104	100	99	98
LRP5	[Bone mineral density variability 1], 601884 Exudative vitreoretinopathy 4, 601813 Hyperostosis, endosteal, 144750 Osteopetrosis 1, 607634 Osteoporosis-pseudoglioma syndrome, 259770 {Osteoporosis}, 166710 Osteosclerosis, 144750 Polycystic liver disease 4 with or without kidney cysts, 617875 van Buchem disease, type 2, 607636	603506	146	100	99	98
LRPAP1	Myopia 23, 615431	104225	135	100	100	100
LRPPRC	Leigh syndrome, French-Canadian type, 220111	607544	60	100	99	91
LRRC6	Ciliary dyskinesia, primary, 19, 614935	614930	79	100	99	93
LRRC8A	?Agammaglobulinemia 5, 613506	608360	174	100	100	100
LRRK2	{Parkinson disease 8}, 607060	609007	78	100	100	96
LRSAM1	Charcot-Marie-Tooth disease, axonal, type 2P, 614436	610933	99	100	100	97
LRTOMT	Deafness 63, 611451	612414	111	100	100	100
LTBP2	Glaucoma 3, primary congenital, D, 613086 Microspherophakia and/or megalocornea, with ectopia lentis and with or without secondary glaucoma, 251750 ?Weill-Marchesani syndrome 3, recessive, 614819	602091	106	100	100	99
LTBP3	Dental anomalies and short stature, 601216 Geleophysic dysplasia 3, 617809	602090	130	100	99	96
LTBP4	Cutis laxa, type IC, 613177	604710	139	100	100	99
LYST	Chediak-Higashi syndrome, 214500	606897	72	100	99	95

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LYZ	Amyloidosis, renal, 105200	153450	62	100	100	100
LZTFL1	Bardet-Biedl syndrome 17, 615994	606568	79	100	100	93
LZTS1	Esophageal squamous cell carcinoma, somatic, 133239	606551	132	100	100	100
MACF1	Lissencephaly 9 with complex brainstem malformation, 618325	608271	68	100	99	94
MAD1L1	Lymphoma, somatic Prostate cancer, somatic, 176807	602686	125	100	100	100
MAF	Ayme-Gripp syndrome, 601088 Cataract 21, multiple types, 610202	177075	80	84	80	76
MAFB	Duane retraction syndrome 3, 617041 Multicentric carpotarsal osteolysis syndrome, 166300	608968	136	100	100	100
MAGEL2	Schaaf-Yang syndrome, 615547	605283	88	100	98	93
MAGT1	Immunodeficiency, with magnesium defect, Epstein-Barr virus infection and neoplasia, 300853	300715	47	100	99	85
MAK	Retinitis pigmentosa 62, 614181	154235	82	100	100	97
MAML2	Mucoepidermoid salivary gland carcinoma	607537	90	100	100	99
MAMLD1	Hypospadias 2, 300758	300120	96	100	99	95
MAN1B1	Mental retardation 15, 614202	604346	122	100	100	100
MAN2B1	Mannosidosis, alpha-, types I and II, 248500	609458	115	100	100	100
MANBA	Mannosidosis, beta, 248510	609489	85	100	100	95
MAOA	{Antisocial behavior}, 300615 Brunner syndrome, 300615	309850	49	100	99	92
MAP2K1	Cardiofaciocutaneous syndrome 3, 615279	176872	79	100	100	96
MAP2K2	Cardiofaciocutaneous syndrome 4, 615280	601263	117	100	100	95
MAP3K1	46XY sex reversal 6, 613762	600982	69	100	98	94
MAP3K8	Lung cancer, somatic, 211980	191195	68	100	100	100
MAPT	Dementia, frontotemporal, with or without parkinsonism, 600274 {Parkinson disease, susceptibility to}, 168600 Pick disease, 172700 Supranuclear palsy, progressive, 601104 Supranuclear palsy, progressive atypical, 260540	157140	120	100	100	100
MARS2	?Combined oxidative phosphorylation deficiency 25, 616430 Spastic ataxia 3, 611390	609728	157	100	100	100
MARVELD2	Deafness 49, 610153	610572	92	100	100	97
MASP1	3MC syndrome 1, 257920	600521	112	100	100	99
MASP2	MASP2 deficiency, 613791	605102	85	100	100	97
MAST1	Mega-corpus-callosum syndrome with cerebellar hypoplasia and cortical malformations, 618273	612256	143	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
MASTL	No OMIM phenotype	608221	65	99	99	96
MAT1A	Hypermethioninemia, persistent, due to methionine adenosyltransferase I/III deficiency, 250850 Methionine adenosyltransferase deficiency, 250850	610550	107	100	100	100
MAT2A	No OMIM phenotype	601468	83	100	100	96
MATN3	Epiphyseal dysplasia, multiple, 5, 607078 {Osteoarthritis susceptibility 2}, 140600 ?Spondyloepimetaphyseal dysplasia, 608728	602109	73	98	90	84
MATR3	Amyotrophic lateral sclerosis 21, 606070	164015	63	100	100	95
MBD5	Mental retardation 1, 156200	611472	81	100	100	99
MBTPS2	IFAP syndrome with or without BRESHECK syndrome, 308205 Keratosis follicularis spinulosa decalvans, 308800 ?Olmsted syndrome, 300918 Osteogenesis imperfecta, type XIX, 301014	300294	46	100	99	90
MC2R	Glucocorticoid deficiency, due to ACTH unresponsiveness, 202200	607397	102	100	100	100
MC4R	Obesity (BMIQ20), 618406 {Obesity, resistance to (BMIQ20)}, 618306	155541	71	100	100	100
MCC	Colorectal cancer, somatic, 114500	159350	94	100	100	98
MCCC1	3-Methylcrotonyl-CoA carboxylase 1 deficiency, 210200	609010	74	100	100	94
MCCC2	3-Methylcrotonyl-CoA carboxylase 2 deficiency, 210210	609014	64	100	100	96
MCEE	Methylmalonyl-CoA epimerase deficiency, 251120	608419	77	100	100	95
MCFD2	Factor V and factor VIII, combined deficiency of, 613625	607788	87	100	100	100
MCM4	Immunodeficiency 54, 609981	602638	88	100	100	97
MCM6	Lactase persistence/nonpersistence, 223100	601806	70	100	100	97
MCOLN1	Mucolipidosis IV, 252650	605248	129	100	100	100
MCPH1	Microcephaly 1, primary, 251200	607117	85	94	94	92
MECP2	{Autism susceptibility 3}, 300496 Encephalopathy, neonatal severe, 300673 Mental retardation syndromic, Lubs type, 300260 Mental retardation, syndromic 13, 300055 Rett syndrome, 312750 Rett syndrome, atypical, 312750 Rett syndrome, preserved speech variant, 312750	300005	113	100	100	96
MED12	Lujan-Fryns syndrome, 309520 Ohdo syndrome, 300895 Opitz-Kaveggia syndrome, 305450	300188	68	100	100	97

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MED13L	Mental retardation and distinctive facial features with or without cardiac defects, 616789 Transposition of the great arteries, dextro-looped 1, 608808	608771	73	100	100	97
MED17	Microcephaly, postnatal progressive, with seizures and brain atrophy, 613668	603810	94	100	100	99
MED23	Mental retardation 18, 614249	605042	63	100	100	96
MED25	Basel-Vanagait-Smirin-Yosef syndrome, 616449 ?Charcot-Marie-Tooth disease, type 2B2, 605589	610197	137	100	100	99
MEF2C	Chromosome 5q14.3 deletion syndrome, 613443 Mental retardation, stereotypic movements, epilepsy, and/or cerebral malformations, 613443	600662	93	100	99	96
MEFV	Familial Mediterranean fever, AD, 134610 Familial Mediterranean fever, AR, 249100	608107	105	100	100	100
MEGF10	Myopathy, areflexia, respiratory distress, and dysphagia, early-onset, 614399 Myopathy, areflexia, respiratory distress, and dysphagia, early-onset, mild variant, 614399	612453	94	100	100	98
MEGF8	Carpenter syndrome 2, 614976	604267	134	100	99	98
MEN1	Adrenal adenoma, somatic Angiofibroma, somatic Carcinoid tumor of lung Lipoma, somatic Multiple endocrine neoplasia 1, 131100 Parathyroid adenoma, somatic	613733	182	100	100	100
MEOX1	Klippel-Feil syndrome 2, 214300	600147	62	100	99	90
MERTK	Retinitis pigmentosa 38, 613862	604705	97	100	100	96
MESP2	Spondylocostal dysostosis 2, 608681	605195	154	100	100	100
MET	?Deafness 97, 616705 Hepatocellular carcinoma, childhood type, somatic, 114550 {Osteofibrous dysplasia, susceptibility to}, 607278 Renal cell carcinoma, papillary, 1, familial and somatic, 605074	164860	64	100	100	96
MFAP5	Aortic aneurysm, familial thoracic 9, 616166	601103	50	100	100	92
MFN2	Charcot-Marie-Tooth disease, axonal, type 2A2A, 609260 Charcot-Marie-Tooth disease, axonal, type 2A2B, 617087 Hereditary motor and sensory neuropathy VIA, 601152	608507	114	100	100	98
MFRP	Microphthalmia, isolated 5, 611040 Nanophthalmos 2, 609549	606227	100	100	100	100
MFSD8	Ceroid lipofuscinosi, neuronal, 7, 610951 Macular dystrophy with central cone involvement, 616170	611124	68	100	100	95
MGAT2	Congenital disorder of glycosylation, type IIa, 212066	602616	104	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
MGME1	Mitochondrial DNA depletion syndrome 11, 615084	615076	58	94	94	92
MGP	Keutel syndrome, 245150	154870	64	100	98	89
MIB1	Left ventricular noncompaction 7, 615092	608677	69	100	100	97
MICU1	Myopathy with extrapyramidal signs, 615673	605084	52	100	98	82
MID1	Opitz GBBB syndrome, type I, 300000	300552	89	100	100	93
MINPP1	{Thyroid carcinoma, follicular}, 188470	605391	135	100	100	100
MIP	Cataract 15, multiple types, 615274	154050	123	100	100	100
MIR17HG	Feingold syndrome 2, 614326	609415	No coverage data			
MIR184	EDICT syndrome, 614303	613146	No coverage data			
MIR96	Deafness 50, 613074	611606	No coverage data			
MITF	COMMAD syndrome, 617306 {Melanoma, cutaneous malignant, susceptibility to, 8}, 614456 Tietz albinism-deafness syndrome, 103500 Waardenburg syndrome, type 2A, 193510 Waardenburg syndrome/ocular albinism, digenic, 103470	156845	87	100	100	99
MKKS	Bardet-Biedl syndrome 6, 605231 McKusick-Kaufman syndrome, 236700	604896	80	100	100	100
MKRN3	Precocious puberty, central, 2, 615346	603856	107	100	100	100
MKS1	Bardet-Biedl syndrome 13, 615990 Joubert syndrome 28, 617121 Meckel syndrome 1, 249000	609883	117	100	100	99
MLC1	Megalencephalic leukoencephalopathy with subcortical cysts, 604004	605908	79	100	100	94
MLH1	Colorectal cancer, hereditary nonpolyposis, type 2, 609310 Mismatch repair cancer syndrome, 276300 Muir-Torre syndrome, 158320	120436	203	100	100	100
MLH3	Colorectal cancer, hereditary nonpolyposis, type 7, 614385 Colorectal cancer, somatic, 114500 {Endometrial cancer, susceptibility to}, 608089	604395	62	100	100	99
MLLT10	Leukemia, acute myeloid, 601626	602409	79	100	100	97
MLLT11	No OMIM phenotype	604684	76	100	100	100
MLPH	Griselli syndrome, type 3, 609227	606526	107	100	100	100
MLYCD	Malonyl-CoA decarboxylase deficiency, 248360	606761	65	100	96	92
MMAA	Methylmalonic aciduria, vitamin B12-responsive, 251100	607481	78	100	100	98
MMAB	Methylmalonic aciduria, vitamin B12-responsive, due to defect in synthesis of adenosylcobalamin, cblB complementation type, 251110	607568	82	100	100	98

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MMACHC	Methylmalonic aciduria and homocystinuria, cblC type, 277400	609831	133	100	100	100
MMADHC	Homocystinuria, cblD type, variant 1, 277410	611935	63	100	100	90
	Methylmalonic aciduria and homocystinuria, cblD type, 277410					
	Methylmalonic aciduria, cblD type, variant 2, 277410					
MMP1	COPD, rate of decline of lung function in, 606963 {Epidermolysis bullosa dystrophica, modifier of}, 226600	120353	63	100	100	98
MMP13	Metaphyseal anadysplasia 1, 602111	600108	78	100	100	95
	Metaphyseal dysplasia, Spahr type, 250400					
	Spondyloepimetaphyseal dysplasia, Missouri type, 602111					
MMP2	Multicentric osteolysis, nodulosis, and arthropathy, 259600	120360	125	100	100	100
MMP20	Amelogenesis imperfecta, type IIA2, 612529	604629	68	100	100	95
MMP21	Heterotaxy, visceral, 7, autosomal, 616749	608416	81	100	96	88
MMP9	Metaphyseal anadysplasia 2, 613073	120361	118	100	100	97
MMUT	Methylmalonic aciduria, mut(0) type, 251000	609058	75	100	100	95
MN1	Meningioma, 607174	156100	151	100	100	100
MNX1	Curarino syndrome, 176450	142994	44	77	69	57
MOCS1	Molybdenum cofactor deficiency A, 252150	603707	105	100	100	99
MOCS2	Molybdenum cofactor deficiency B, 252160	603708	67	100	100	96
MOG	?Narcolepsy 7, 614250	159465	91	100	100	100
MOGS	Congenital disorder of glycosylation, type IIb, 606056	601336	130	100	100	100
MPC1	Mitochondrial pyruvate carrier deficiency, 614741	614738	74	100	100	92
MPDU1	Congenital disorder of glycosylation, type If, 609180	604041	87	100	100	99
MPDZ	Hydrocephalus, congenital, 2, with or without brain or eye anomalies, 615219	603785	69	100	99	95
MPI	Congenital disorder of glycosylation, type Ib, 602579	154550	154	100	100	99
MPL	Myelofibrosis with myeloid metaplasia, somatic, 254450	159530	109	100	100	100
	Thrombocythemia 2, 601977					
	Thrombocytopenia, congenital amegakaryocytic, 604498					
MPLKIP	Trichothiodystrophy 4, nonphotosensitive, 234050	609188	56	100	100	100
MPO	{Alzheimer disease, susceptibility to}, 104300	606989	137	100	100	100
	{Lung cancer, protection against, in smokers}					
	Myeloperoxidase deficiency, 254600					
MPV17	Charcot-Marie-Tooth disease, axonal, type 2EE, 618400	137960	108	100	100	100
	Mitochondrial DNA depletion syndrome 6 (hepatocerebral type), 256810					

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MPZ	Charcot-Marie-Tooth disease, dominant intermediate D, 607791 Charcot-Marie-Tooth disease, type 1B, 118200 Charcot-Marie-Tooth disease, type 2I, 607677 Charcot-Marie-Tooth disease, type 2J, 607736 Dejerine-Sottas disease, 145900 Hypomyelinating neuropathy, congenital, 2, 618184 Roussy-Levy syndrome, 180800	159440	125	100	100	96
MRAP	Glucocorticoid deficiency 2, 607398	609196	172	100	100	100
MRE11	Ataxia-telangiectasia-like disorder 1, 604391	600814	55	100	97	83
MRPL3	Combined oxidative phosphorylation deficiency 9, 614582	607118	64	100	100	90
MRPS16	Combined oxidative phosphorylation deficiency 2, 610498	609204	120	100	100	100
MRPS22	Combined oxidative phosphorylation deficiency 5, 611719 Ovarian dysgenesis 7, 618117	605810	75	100	100	96
MRTFA	Megakaryoblastic leukemia, acute	606078	94	98	97	92
MS4A1	Immunodeficiency, common variable, 5, 613495	112210	80	100	100	98
MSH2	Colorectal cancer, hereditary nonpolyposis, type 1, 120435 Mismatch repair cancer syndrome, 276300 Muir-Torre syndrome, 158320	609309	191	100	100	100
MSH3	Endometrial carcinoma, somatic, 608089 Familial adenomatous polyposis 4, 617100	600887	81	100	100	97
MSH6	Colorectal cancer, hereditary nonpolyposis, type 5, 614350 {Endometrial cancer, familial}, 608089 Mismatch repair cancer syndrome, 276300	600678	213	100	100	100
MSR1	Barrett esophagus/esophageal adenocarcinoma, 614266	153622	94	100	99	92
MSRB3	Deafness 74, 613718	613719	72	100	100	93
MSTN	Muscle hypertrophy, 614160	601788	80	100	99	94
MSX1	Ectodermal dysplasia 3, Witkop type, 189500 Orofacial cleft 5, 608874 Tooth agenesis, selective, 1, with or without orofacial cleft, 106600	142983	144	100	100	100
MSX2	Craniosynostosis 2, 604757 Parietal foramina 1, 168500 Parietal foramina with cleidocranial dysplasia, 168550	123101	82	100	100	98
MTAP	Diaphyseal medullary stenosis with malignant fibrous histiocytoma, 112250	156540	72	100	99	90
MTFMT	Combined oxidative phosphorylation deficiency 15, 614947 Mitochondrial complex I deficiency, nuclear type 27, 618248	611766	78	100	100	90

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
MTHFR	Homocystinuria due to MTHFR deficiency, 236250 {Neural tube defects, susceptibility to}, 601634 {Schizophrenia, susceptibility to}, 181500 {Thromboembolism, susceptibility to}, 188050 {Vascular disease, susceptibility to}	607093	99	100	100	99
MTM1	Myotubular myopathy, 310400	300415	46	100	95	75
MTMR2	Charcot-Marie-Tooth disease, type 4B1, 601382	603557	72	100	100	95
MTO1	Combined oxidative phosphorylation deficiency 10, 614702	614667	116	97	94	93
MTPAP	?Spastic ataxia 4, 613672	613669	97	100	100	97
MTR	Homocystinuria-megaloblastic anemia, cblG complementation type, 250940 {Neural tube defects, folate-sensitive, susceptibility to}, 601634	156570	80	100	100	97
MTRR	Homocystinuria-megaloblastic anemia, cbl E type, 236270 {Neural tube defects, folate-sensitive, susceptibility to}, 601634	602568	87	100	100	98
MTTP	Abetalipoproteinemia, 200100 {Metabolic syndrome, protection against}, 605552	157147	67	100	100	97
MUC1	Medullary cystic kidney disease 1, 174000	158340	78	98	92	87
MUSK	Fetal akinesia deformation sequence 1, 208150 Myasthenic syndrome, congenital, 9, associated with acetylcholine receptor deficiency, 616325	601296	85	100	100	99
MUTYH	Adenomas, multiple colorectal, 608456 Colorectal adenomatous polyposis, with pilomatrixomas, 132600 Gastric cancer, somatic, 613659	604933	175	100	100	100
MVK	Hyper-IgD syndrome, 260920 Mevalonic aciduria, 610377 Porokeratosis 3, multiple types, 175900	251170	100	100	100	100
MXI1	Neurofibrosarcoma, somatic Prostate cancer, somatic, 176807	600020	66	97	93	88
MYBPC1	Arthrogryposis, distal, type 1B, 614335 Lethal congenital contracture syndrome 4, 614915 Myopathy, congenital, with tremor, 618524	160794	62	100	100	94
MYBPC3	Cardiomyopathy, dilated, 1MM, 615396 Cardiomyopathy, hypertrophic, 4, 115197 Left ventricular noncompaction 10, 615396	600958	169	100	100	98
MYCN	Feingold syndrome 1, 164280	164840	169	100	100	100
MYD88	Macroglobulinemia, Waldenstrom, somatic, 153600 Pyogenic bacterial infections, recurrent, due to MYD88 deficiency, 612260	602170	157	100	100	100

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MYF6	No OMIM phenotype	159991	132	100	100	100
MYH11	Aortic aneurysm, familial thoracic 4, 132900	160745	136	100	100	99
MYH14	Deafness 4A, 600652 ?Peripheral neuropathy, myopathy, hoarseness, and hearing loss, 614369	608568	114	100	99	97
MYH2	Proximal myopathy and ophthalmoplegia, 605637	160740	85	100	100	98
MYH3	Arthrogryposis, distal, type 2A (Freeman-Sheldon), 193700 Arthrogryposis, distal, type 2B3 (Sheldon-Hall), 618436 Contractures, pterygia, and variable skeletal fusions syndrome 1A, 178110 Contractures, pterygia, and variable skeletal fusions syndrome 1B, 618469	160720	95	100	100	98
MYH6	Atrial septal defect 3, 614089 Cardiomyopathy, dilated, 1EE, 613252 Cardiomyopathy, hypertrophic, 14, 613251 {Sick sinus syndrome 3}, 614090	160710	145	100	99	98
MYH7	Cardiomyopathy, dilated, 1S, 613426 Cardiomyopathy, hypertrophic, 1, 192600 Laing distal myopathy, 160500 Left ventricular noncompaction 5, 613426 Myopathy, myosin storage, 608358 Myopathy, myosin storage, 255160 Scapuloperoneal syndrome, myopathic type, 181430	160760	176	100	100	100
MYH7B	No OMIM phenotype	609928	115	100	100	99
MYH8	Carney complex variant, 608837 Trismus-pseudocampodactyly syndrome, 158300	160741	79	100	100	96
MYH9	Deafness 17, 603622 Macrothrombocytopenia and granulocyte inclusions with or without nephritis or sensorineural hearing loss, 155100	160775	125	100	100	99
MYL2	Cardiomyopathy, hypertrophic, 10, 608758	160781	170	100	100	100
MYL3	Cardiomyopathy, hypertrophic, 8, 608751	160790	118	100	100	100
MYLK	Aortic aneurysm, familial thoracic 7, 613780	600922	121	100	100	99
MYLK2	Cardiomyopathy, hypertrophic, 1, digenic, 192600	606566	111	100	100	100
MYO15A	Deafness 3, 600316	602666	134	100	99	98
MYO1A	No OMIM phenotype	601478	97	100	100	100
MYO1E	Glomerulosclerosis, focal segmental, 6, 614131	601479	76	100	100	98
MYO3A	Deafness 30, 607101	606808	73	100	99	95
MYO5A	Griselli syndrome, type 1, 214450	160777	71	100	99	94

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MYO5B	Microvillus inclusion disease, 251850	606540	103	100	100	98
MYO6	Deafness 22, 606346 Deafness 22, with hypertrophic cardiomyopathy, 606346 Deafness 37, 607821	600970	82	100	99	94
MYO7A	Deafness 11, 601317 Deafness 2, 600060 Usher syndrome, type 1B, 276900	276903	109	100	100	99
MYOC	Glaucoma 1A, primary open angle, 137750	601652	140	100	100	100
MYOT	Myopathy, myofibrillar, 3, 609200 Myopathy, spheroid body, 182920	604103	81	100	100	96
MYOZ2	Cardiomyopathy, hypertrophic, 16, 613838	605602	78	100	100	100
MYPN	Cardiomyopathy, dilated, 1KK, 615248 Cardiomyopathy, familial restrictive, 4, 615248 Cardiomyopathy, hypertrophic, 22, 615248 Nemaline myopathy 11, 617336	608517	80	100	100	97
MYRF	Cardiac-urogenital syndrome, 618280 Encephalitis/encephalopathy, mild, with reversible myelin vacuolization, 618113	608329	118	98	97	96
NAA10	?Microphthalmia, syndromic 1, 309800 Ogden syndrome, 300855 dominant	300013	89	100	100	94
NAA15	Mental retardation 50, 617787	608000	79	100	100	95
NAGA	Kanzaki disease, 609242 Schindler disease, type I, 609241 Schindler disease, type III, 609241	104170	118	100	100	100
NAGLU	?Charcot-Marie-Tooth disease, axonal, type 2V, 616491 Mucopolysaccharidosis type IIIB (Sanfilippo B), 252920	609701	112	100	97	93
NAGS	N-acetylglutamate synthase deficiency, 237310	608300	110	100	100	100
NALCN	Congenital contractures of the limbs and face, hypotonia, and developmental delay, 616266 Hypotonia, infantile, with psychomotor retardation and characteristic facies 1, 615419	611549	69	100	100	97
NANOS1	Spermatogenic failure 12, 615413	608226	96	99	96	92
NBAS	Infantile liver failure syndrome 2, 616483 Short stature, optic nerve atrophy, and Pelger-Huet anomaly, 614800	608025	66	100	99	94
NBEAL2	Gray platelet syndrome, 139090	614169	139	100	99	99
NBN	Aplastic anemia, 609135 Leukemia, acute lymphoblastic, 613065 Nijmegen breakage syndrome, 251260	602667	68	100	99	88

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NCF1	Chronic granulomatous disease due to deficiency of NCF-1, 233700	608512	108	68	65	61
NCF2	Chronic granulomatous disease due to deficiency of NCF-2, 233710	608515	94	100	100	98
NCF4	?Granulomatous disease, chronic, cytochrome b-positive, type III, 613960	601488	121	100	100	100
NCOA4	No OMIM phenotype	601984	72	100	100	99
NCSTN	Acne inversa, familial, 1, 142690	605254	91	100	100	98
NDE1	Lissencephaly 4 (with microcephaly), 614019 ?Microhydranencephaly, 605013	609449	109	100	100	100
NDN	Prader-Willi syndrome, 176270	602117	99	100	100	97
NDP	Exudative vitreoretinopathy 2, 305390 Norrie disease, 310600	300658	77	100	100	100
NDRG1	Charcot-Marie-Tooth disease, type 4D, 601455	605262	97	100	100	100
NDUFA1	Mitochondrial complex I deficiency, nuclear type 12, 301020	300078	110	100	100	100
NDUFA10	Mitochondrial complex I deficiency, nuclear type 22, 618243	603835	80	100	95	85
NDUFA11	Mitochondrial complex I deficiency, nuclear type 14, 618236	612638	117	100	100	100
NDUFA12	?Mitochondrial complex I deficiency, nuclear type 23, 618244	614530	72	100	100	100
NDUFA2	?Mitochondrial complex I deficiency, nuclear type 13, 618235	602137	104	100	100	100
NDUFA9	Mitochondrial complex I deficiency, nuclear type 26, 618247	603834	82	100	100	100
NDUFAF1	Mitochondrial complex I deficiency, nuclear type 11, 618234	606934	63	100	100	95
NDUFAF2	Mitochondrial complex I deficiency, nuclear type 10, 618233	609653	83	100	92	78
NDUFAF3	Mitochondrial complex I deficiency, nuclear type 18, 618240	612911	130	100	100	100
NDUFAF4	Mitochondrial complex I deficiency, nuclear type 15, 618237	611776	95	100	100	100
NDUFAF5	Mitochondrial complex I deficiency, nuclear type 16, 618238	612360	70	100	100	96
NDUFAF6	Mitochondrial complex I deficiency, nuclear type 17, 618239	612392	46	100	100	90
NDUFB3	Mitochondrial complex I deficiency, nuclear type 25, 618246	603839	31	100	87	55
NDUFS1	Mitochondrial complex I deficiency, nuclear type 5, 618226	157655	69	100	100	94
NDUFS2	Mitochondrial complex I deficiency, nuclear type 6, 618228	602985	79	100	100	99
NDUFS3	Mitochondrial complex I deficiency, nuclear type 8, 618230	603846	112	100	100	100
NDUFS4	Mitochondrial complex I deficiency, nuclear type 1, 252010	602694	83	100	100	99
NDUFS6	Mitochondrial complex I deficiency, nuclear type 9, 618232	603848	70	100	100	100
NDUFS7	Mitochondrial complex I deficiency, nuclear type 3, 618224	601825	130	100	100	98
NDUFS8	Mitochondrial complex I deficiency, nuclear type 2, 618222	602141	140	100	100	100
NDUFV1	Mitochondrial complex I deficiency, nuclear type 4, 618225	161015	129	100	100	100
NDUFV2	Mitochondrial complex I deficiency, nuclear type 7, 618229	600532	48	100	95	80
NEB	Nemaline myopathy 2, 256030	161650	69	88	87	84

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NECTIN1	Cleft lip/palate-ectodermal dysplasia syndrome, 225060 Orofacial cleft 7, 225060	600644	112	100	100	100
NECTIN4	Ectodermal dysplasia-syndactyly syndrome 1, 613573	609607	121	100	100	100
NEFL	Charcot-Marie-Tooth disease, dominant intermediate G, 617882 Charcot-Marie-Tooth disease, type 1F, 607734 Charcot-Marie-Tooth disease, type 2E, 607684	162280	157	100	100	100
NEK1	{Amyotrophic lateral sclerosis, susceptibility to, 24}, 617892 Short-rib thoracic dysplasia 6 with or without polydactyly, 263520	604588	56	100	98	89
NEU1	Sialidosis, type I, 256550 Sialidosis, type II, 256550	608272	155	100	100	100
NEUROD1	{Diabetes mellitus, noninsulin-dependent}, 125853 Maturity-onset diabetes of the young 6, 606394	601724	115	100	100	100
NEUROG3	Diarrhea 4, malabsorptive, congenital, 610370	604882	161	100	100	100
NEXMIF	Mental retardation 98, 300912	300524	48	100	100	95
NEXN	Cardiomyopathy, dilated, 1CC, 613122 Cardiomyopathy, hypertrophic, 20, 613876	613121	85	100	100	99
NF1	Leukemia, juvenile myelomonocytic, 607785 Neurofibromatosis, familial spinal, 162210 Neurofibromatosis, type 1, 162200 Neurofibromatosis-Noonan syndrome, 601321 Watson syndrome, 193520	613113	55	97	90	79
NF2	Meningioma, NF2-related, somatic, 607174 Neurofibromatosis, type 2, 101000 Schwannomatosis, somatic, 162091	607379	141	100	100	100
NFIX	Marshall-Smith syndrome, 602535 Sotos syndrome 2, 614753	164005	154	100	100	100
NFKB2	Immunodeficiency, common variable, 10, 615577	164012	143	100	100	96
NFKBIA	Ectodermal dysplasia and immunodeficiency 2, 612132	164008	131	100	100	100
NFU1	Multiple mitochondrial dysfunctions syndrome 1, 605711	608100	66	100	100	95
NGF	Neuropathy, hereditary sensory and autonomic, type V, 608654	162030	131	100	100	100
NGLY1	Congenital disorder of deglycosylation, 615273	610661	78	100	100	97
NHEJ1	Severe combined immunodeficiency with microcephaly, growth retardation, and sensitivity to ionizing radiation, 611291	611290	55	100	100	96
NHLRC1	Epilepsy, progressive myoclonic 2B (Lafora), 254780	608072	144	100	100	100
NHP2	Dyskeratosis congenita 2, 613987	606470	122	100	100	100

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NHS	Cataract 40, 302200 Nance-Horan syndrome, 302350	300457	57	100	96	90
NIN	?Seckel syndrome 7, 614851	608684	79	100	99	95
NIPA1	Spastic paraplegia 6, 600363	608145	107	100	100	97
NIPAL4	Ichthyosis, congenital 6, 612281	609383	97	100	100	99
NIPBL	Cornelia de Lange syndrome 1, 122470	608667	65	100	98	92
NKX2-1	Chorea, hereditary benign, 118700 Choreoathetosis, hypothyroidism, and neonatal respiratory distress, 610978 {Thyroid cancer, nonmedullary, 1}, 188550	600635	95	100	100	100
NKX2-5	Atrial septal defect 7, with or without AV conduction defects, 108900 Conotruncal heart malformations, variable, 217095 Hypoplastic left heart syndrome 2, 614435 Hypothyroidism, congenital nongoitrous, 5, 225250 Tetralogy of Fallot, 187500 Ventricular septal defect 3, 614432	600584	101	100	100	97
NKX2-6	Conotruncal heart malformations, 217095 Persistent truncus arteriosus, 217095	611770	145	100	100	100
NKX3-2	Spondylo-megaepiphyseal-metaphyseal dysplasia, 613330	602183	108	100	100	100
NLGN4X	{Asperger syndrome susceptibility 2}, 300497 {Autism susceptibility 2}, 300495 Mental retardation, 300495	300427	179	100	100	100
NLRP12	Familial cold autoinflammatory syndrome 2, 611762	609648	135	100	100	100
NLRP3	CINCA syndrome, 607115 Deafness 34, with or without inflammation, 617772 Familial cold inflammatory syndrome 1, 120100 Keratoendothelitis fugax hereditaria, 148200 Muckle-Wells syndrome, 191900	606416	125	100	100	100
NLRP7	Hydatidiform mole, recurrent, 1, 231090	609661	167	100	100	100
NME8	Ciliary dyskinesia, primary, 6, 610852	607421	65	100	100	93
NMNAT1	Leber congenital amaurosis 9, 608553	608700	90	100	100	98
NNT	Glucocorticoid deficiency 4, with or without mineralocorticoid deficiency, 614736	607878	67	100	100	96
NOBOX	Premature ovarian failure 5, 611548	610934	86	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
NOD2	Blau syndrome, 186580 {Inflammatory bowel disease 1, Crohn disease}, 266600 {Psoriatic arthritis, susceptibility to}, 607507 {Yao syndrome}, 617321	605956	113	100	100	99
NODAL	Heterotaxy, visceral, 5, 270100	601265	119	100	100	100
NOG	Brachydactyly, type B2, 611377 Multiple synostoses syndrome 1, 186500 Stapes ankylosis with broad thumbs and toes, 184460 Symphalangism, proximal, 1A, 185800 Tarsal-carpal coalition syndrome, 186570	602991	198	100	100	100
NOL3	?Myoclonus, familial, 1, 614937	605235	132	100	100	100
NOP10	Dyskeratosis congenita 1, 224230	606471	147	100	100	100
NOP56	Spinocerebellar ataxia 36, 614153	614154	93	100	100	99
NOTCH1	Adams-Oliver syndrome 5, 616028 Aortic valve disease 1, 109730	190198	117	100	99	98
NOTCH2	Alagille syndrome 2, 610205 Hajdu-Cheney syndrome, 102500	600275	98	100	99	98
NOTCH3	Cerebral arteriopathy with subcortical infarcts and leukoencephalopathy 1, 125310 Lateral meningocele syndrome, 130720 ?Myofibromatosis, infantile 2, 615293	600276	111	99	95	91
NPC1	Niemann-Pick disease, type C1, 257220 Niemann-Pick disease, type D, 257220	607623	87	100	100	99
NPC2	Niemann-pick disease, type C2, 607625	601015	95	100	100	100
NPHP1	Joubert syndrome 4, 609583 Nephronophthisis 1, juvenile, 256100 Senior-Loken syndrome-1, 266900	607100	58	100	98	88
NPHP3	Meckel syndrome 7, 267010 Nephronophthisis 3, 604387 Renal-hepatic-pancreatic dysplasia 1, 208540	608002	66	100	99	94
NPHP4	Nephronophthisis 4, 606966 Senior-Loken syndrome 4, 606996	607215	110	100	100	100
NPHS1	Nephrotic syndrome, type 1, 256300	602716	102	100	100	100
NPHS2	Nephrotic syndrome, type 2, 600995	604766	84	100	100	100
NPM1	Leukemia, acute myeloid, somatic, 601626	164040	72	100	96	92

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NPPA	Atrial fibrillation, familial, 6, 612201 Atrial standstill 2, 615745	108780	120	100	100	100
NPR2	Acromesomelic dysplasia, Maroteaux type, 602875 Epiphyseal chondrodysplasia, Miura type, 615923 Short stature with nonspecific skeletal abnormalities, 616255	108961	116	100	100	100
NR0B1	Adrenal hypoplasia, congenital, 300200 46XY sex reversal 2, dosage-sensitive, 300018	300473	120	100	100	99
NR0B2	Obesity, mild, early-onset, 601665	604630	72	100	100	96
NR2E3	Enhanced S-cone syndrome, 268100 Retinitis pigmentosa 37, 611131	604485	123	100	100	99
NR2F1	Bosch-Boonstra-Schaaf optic atrophy syndrome, 615722	132890	197	100	100	100
NR3C1	Glucocorticoid resistance, 615962	138040	80	100	100	98
NR3C2	Hypertension, early-onset, with exacerbation in pregnancy, 605115 Pseudohypoaldosteronism type I, 177735	600983	89	100	100	98
NR4A3	Chondrosarcoma, extraskeletal myxoid, 612237	600542	114	100	100	100
NR5A1	Adrenocortical insufficiency, 612964 Premature ovarian failure 7, 612964 Spermatogenic failure 8, 613957 46, XX sex reversal 4, 617480 46XY sex reversal 3, 612965	184757	107	100	100	100
NRAS	Colorectal cancer, somatic, 114500 Epidermal nevus, somatic, 162900 Melanocytic nevus syndrome, congenital, somatic, 137550 Neurocutaneous melanosis, somatic, 249400 Noonan syndrome 6, 613224 ?RAS-associated autoimmune lymphoproliferative syndrome type IV, somatic, 614470 Schimmelpenning-Feuerstein-Mims syndrome, somatic mosaic, 163200 Thyroid carcinoma, follicular, somatic, 188470	164790	63	100	100	98
NRL	Retinal degeneration, clumped pigment type Retinitis pigmentosa 27, 613750	162080	108	100	100	100
NRXN1	Pitt-Hopkins-like syndrome 2, 614325 {Schizophrenia, susceptibility to, 17}, 614332	600565	104	100	100	97
NSD1	Leukemia, acute myeloid, 601626 Sotos syndrome 1, 117550	606681	77	100	100	98
NSD3	No OMIM phenotype	607083	55	100	98	90

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NSDHL	CHILD syndrome, 308050 CK syndrome, 300831	300275	68	100	100	93
NSMF	Hypogonadotropic hypogonadism 9 with or without anosmia, 614838	608137	96	100	95	95
NSUN2	Mental retardation 5, 611091	610916	85	100	99	92
NT5C2	Spastic paraplegia 45, 613162	600417	54	100	98	89
NT5C3A	Anemia, hemolytic, due to UMPH1 deficiency, 266120	606224	61	100	99	89
NT5E	Calcification of joints and arteries, 211800	129190	90	100	100	98
NTF4	Glaucoma 1, open angle, 1O, 613100	162662	176	100	100	100
NTRK1	Insensitivity to pain, congenital, with anhidrosis, 256800 Medullary thyroid carcinoma, familial, 155240	191315	119	100	100	99
NTRK2	Epileptic encephalopathy, early infantile, 58, 617830 Obesity, hyperphagia, and developmental delay, 613886	600456	92	100	100	96
NUBPL	Mitochondrial complex I deficiency, nuclear type 21, 618242	613621	60	100	100	97
NUMA1	Leukemia, acute promyelocytic, somatic, 612376	164009	129	100	100	98
NUP214	{Encephalopathy, acute, infection-induced, susceptibility to, 9}, 618426 Leukemia, T-cell acute lymphoblastic, somatic, 613065 Leukemia, acute myeloid, somatic, 601626	114350	111	100	100	98
NUP62	Striatonigral degeneration, infantile, 271930	605815	107	100	100	100
NYX	Night blindness, congenital stationary (complete), 1A, 310500	300278	132	100	100	100
OAT	Gyrate atrophy of choroid and retina with or without ornithinemia, 258870	613349	64	100	96	85
OBSL1	3-M syndrome 2, 612921	610991	135	100	100	100
OCA2	Albinism, brown oculocutaneous, 203200 Albinism, oculocutaneous, type II, 203200 [Skin/hair/eye pigmentation 1, blond/brown hair], 227220 [Skin/hair/eye pigmentation 1, blue/nonblue eyes], 227220	611409	93	100	100	97
OCLN	Pseudo-TORCH syndrome 1, 251290	602876	61	96	84	79
OCRL	Dent disease 2, 300555 Lowe syndrome, 309000	300535	43	100	95	80
ODAPH	Amelogenesis imperfecta, type IIA4, 614832	614829	88	100	100	96
OFD1	Joubert syndrome 10, 300804 Orofaciodigital syndrome I, 311200 ?Retinitis pigmentosa 23, 300424 Simpson-Golabi-Behmel syndrome, type 2, 300209	300170	43	100	95	75
OGG1	Renal cell carcinoma, clear cell, somatic, 144700	601982	101	100	100	100

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OPA1	Behr syndrome, 210000 {Glaucoma, normal tension, susceptibility to}, 606657 ?Mitochondrial DNA depletion syndrome 14 (encephalocardiomyopathic type), 616896 Optic atrophy 1, 165500 Optic atrophy plus syndrome, 125250	605290	59	100	98	89
OPA3	3-methylglutaconic aciduria, type III, 258501 Optic atrophy 3 with cataract, 165300	606580	130	100	100	100
OPHN1	Mental retardation, with cerebellar hypoplasia and distinctive facial appearance, 300486	300127	52	100	96	84
OPLAH	5-oxoprolinase deficiency, 260005	614243	132	100	100	99
OPN1LW	Blue cone monochromacy, 303700 Colorblindness, protan, 303900	300822	121	89	89	88
OPN1MW	Blue cone monochromacy, 303700 Colorblindness, deutan, 303800	300821	47	54	45	42
OPN1SW	Colorblindness, tritan, 190900	613522	78	100	100	100
OPTN	Amyotrophic lateral sclerosis 12, 613435 Glaucoma 1, open angle, E, 137760 {Glaucoma, normal tension, susceptibility to}, 606657	602432	81	100	100	96
ORAI1	Immunodeficiency 9, 612782 Myopathy, tubular aggregate, 2, 615883	610277	175	99	99	97
ORC1	Meier-Gorlin syndrome 1, 224690	601902	86	100	100	97
ORC4	Meier-Gorlin syndrome 2, 613800	603056	67	100	100	91
ORC6	Meier-Gorlin syndrome 3, 613803	607213	70	100	100	98
OSMR	Amyloidosis, primary localized cutaneous, 1, 105250	601743	74	100	100	97
OSTM1	Osteopetrosis 5, 259720	607649	79	100	100	97
OTC	Ornithine transcarbamylase deficiency, 311250	300461	58	100	98	85
OTOA	Deafness 22, 607039	607038	78	77	74	72
OTOF	Auditory neuropathy, 1, 601071 Deafness 9, 601071	603681	113	100	100	100
OTOG	Deafness 18B, 614945	604487	121	100	100	99
OTOGL	Deafness 84B, 614944	614925	76	100	100	95
OTX2	Microphthalmia, syndromic 5, 610125 Pituitary hormone deficiency, combined, 6, 613986 Retinal dystrophy, early-onset, with or without pituitary dysfunction, 610125	600037	100	100	100	100
OXCT1	Succinyl CoA:3-oxoacid CoA transferase deficiency, 245050	601424	74	100	100	96
P2RX1	No OMIM phenotype	600845	103	100	100	100

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P2RX2	Deafness 41, 608224	600844	136	100	100	99
P2RY12	Bleeding disorder, platelet-type, 8, 609821	600515	67	100	100	100
P3H1	Osteogenesis imperfecta, type VIII, 610915	610339	123	100	100	100
P3H2	Myopia, high, with cataract and vitreoretinal degeneration, 614292	610341	65	100	100	95
PABPN1	Oculopharyngeal muscular dystrophy, 164300	602279	89	89	76	71
PACS1	Schuurs-Hoeijmakers syndrome, 615009	607492	92	100	100	97
PAFAH1B1	Lissencephaly 1, 607432	601545	82	100	96	90
	Subcortical laminar heterotopia, 607432					
PAH	[Hyperphenylalaninemia, non-PKU mild], 261600 Phenylketonuria, 261600	612349	68	100	99	95
PAK3	Mental retardation 30/47, 300558	300142	48	100	96	83
PALB2	{Breast cancer, susceptibility to}, 114480	610355	69	100	100	97
	Fanconi anemia, complementation group N, 610832					
	{Pancreatic cancer, susceptibility to, 3}, 613348					
PANK2	HARP syndrome, 607236	606157	81	100	100	99
	Neurodegeneration with brain iron accumulation 1, 234200					
PAPSS2	Brachyolmia 4 with mild epiphyseal and metaphyseal changes, 612847	603005	94	100	100	95
PARK7	Parkinson disease 7 early-onset, 606324	602533	68	100	99	87
PAX2	Glomerulosclerosis, focal segmental, 7, 616002	167409	144	100	100	98
	Papillorenal syndrome, 120330					
PAX3	Craniofacial-deafness-hand syndrome, 122880	606597	104	100	100	100
	Rhabdomyosarcoma 2, alveolar, 268220					
	Waardenburg syndrome, type 1, 193500					
	Waardenburg syndrome, type 3, 148820					
PAX4	{Diabetes mellitus, ketosis-prone, susceptibility to}, 612227	167413	74	100	100	99
	Diabetes mellitus, type 2, 125853					
	Maturity-onset diabetes of the young, type IX, 612225					

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PAX6	Aniridia, 106210 Anterior segment dysgenesis 5, multiple subtypes, 604229 Cataract with late-onset corneal dystrophy, 106210 ?Coloboma of optic nerve, 120430 ?Coloboma, ocular, 120200 Foveal hypoplasia 1, 136520 Keratitis, 148190 ?Morning glory disc anomaly, 120430 Optic nerve hypoplasia, 165550	607108	73	100	100	96
PAX8	Hypothyroidism, congenital, due to thyroid dysgenesis or hypoplasia, 218700	167415	111	100	100	97
PAX9	Tooth agenesis, selective, 3, 604625	167416	215	100	98	98
PC	Pyruvate carboxylase deficiency, 266150	608786	141	100	100	100
PCARE	Retinitis pigmentosa 54, 613428	613425	105	100	100	98
PCBD1	Hyperphenylalaninemia, BH4-deficient, D, 264070	126090	87	100	100	99
PCCA	Propionicacidemia, 606054	232000	68	100	100	95
PCCB	Propionicacidemia, 606054	232050	72	100	98	95
PCDH15	Deafness 23, 609533 Usher syndrome, type 1D/F digenic, 601067 Usher syndrome, type 1F, 602083	605514	72	100	100	95
PCDH19	Epileptic encephalopathy, early infantile, 9, 300088	300460	113	100	100	97
PCM1	No OMIM phenotype	600299	79	100	99	94
PCNT	Microcephalic osteodysplastic primordial dwarfism, type II, 210720	605925	121	100	100	99
PCSK1	Obesity with impaired prohormone processing, 600955 {Obesity, susceptibility to, BMIQ12}, 612362	162150	68	100	100	97
PCSK9	Hypercholesterolemia, familial, 3, 603776 {Low density lipoprotein cholesterol level QTL 1}, 603776	607786	116	100	98	95
PCYT1A	Spondylometaphyseal dysplasia with cone-rod dystrophy, 608940	123695	74	100	99	85
PDCD10	Cerebral cavernous malformations 3, 603285	609118	48	100	99	82
PDE11A	Pigmented nodular adrenocortical disease, primary, 2, 610475	604961	77	100	99	93
PDE4D	Acrodyostosis 2, with or without hormone resistance, 614613	600129	70	100	97	94
PDE6A	Retinitis pigmentosa 43, 613810	180071	74	100	100	96
PDE6B	Night blindness, congenital stationary 2, 163500 Retinitis pigmentosa-40, 613801	180072	125	100	100	100
PDE6C	Cone dystrophy 4, 613093	600827	67	100	100	96
PDE6G	Retinitis pigmentosa 57, 613582	180073	147	100	100	100

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PDE6H	Achromatopsia 6, 610024 Retinal cone dystrophy 3, 610024	601190	47	100	100	96
PDE8B	Pigmented nodular adrenocortical disease, primary, 3, 614190 Striatal degeneration, 609161	603390	72	100	99	96
PDGFB	Basal ganglia calcification, idiopathic, 5, 615483 Dermatofibrosarcoma protuberans, 607907 Meningioma, SIS-related, 607174	190040	101	100	100	100
PDGFRA	Gastrointestinal stromal tumor/GIST-plus syndrome, somatic or familial, 175510 Hypereosinophilic syndrome, idiopathic, resistant to imatinib, 607685	173490	79	100	98	95
PDGFRB	Basal ganglia calcification, idiopathic, 4, 615007 Kosaki overgrowth syndrome, 616592 Myeloproliferative disorder with eosinophilia, 131440 Myofibromatosis, infantile, 1, 228550 Premature aging syndrome, Penttinen type, 601812	173410	116	100	100	100
PDGFRL	Colorectal cancer, somatic, 114500 Hepatocellular cancer, somatic, 114550	604584	129	100	100	100
PDHA1	Pyruvate dehydrogenase E1-alpha deficiency, 312170	300502	49	100	94	84
PDHB	Pyruvate dehydrogenase E1-beta deficiency, 614111	179060	65	100	100	95
PDP1	Pyruvate dehydrogenase phosphatase deficiency, 608782	605993	87	100	100	100
PDS5B	No OMIM phenotype	605333	52	100	98	89
PDSS1	Coenzyme Q10 deficiency, primary, 2, 614651	607429	60	100	93	85
PDSS2	Coenzyme Q10 deficiency, primary, 3, 614652	610564	75	100	97	87
PDX1	{Diabetes mellitus, type II, susceptibility to}, 125853 MODY, type IV, 606392 Pancreatic agenesis 1, 260370	600733	79	100	100	89
PDYN	Spinocerebellar ataxia 23, 610245	131340	134	100	100	100
PDZD7	Deafness 57, 618003 {Retinal disease in Usher syndrome type IIA, modifier of}, 276901 Usher syndrome, type IIC, GPR98/PDZD7 digenic, 605472	612971	84	100	100	99
PEPD	Prolidase deficiency, 170100	613230	100	100	100	98
PER2	?Advanced sleep phase syndrome, familial, 1, 604348	603426	84	100	100	97
PET100	Mitochondrial complex IV deficiency, 220110	614770	85	92	66	66
PEX1	Heimler syndrome 1, 234580 Peroxisome biogenesis disorder 1A (Zellweger), 214100 Peroxisome biogenesis disorder 1B (NALD/IRD), 601539	602136	62	100	99	94

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PEX10	Peroxisome biogenesis disorder 6A (Zellweger), 614870 Peroxisome biogenesis disorder 6B, 614871	602859	100	100	100	97
PEX11B	?Peroxisome biogenesis disorder 14B, 614920	603867	137	100	100	100
PEX12	Peroxisome biogenesis disorder 3A (Zellweger), 614859 Peroxisome biogenesis disorder 3B, 266510	601758	67	100	100	94
PEX13	Peroxisome biogenesis disorder 11A (Zellweger), 614883 Peroxisome biogenesis disorder 11B, 614885	601789	69	100	100	99
PEX14	Peroxisome biogenesis disorder 13A (Zellweger), 614887	601791	142	100	100	100
PEX16	Peroxisome biogenesis disorder 8A (Zellweger), 614876 Peroxisome biogenesis disorder 8B, 614877	603360	134	100	96	94
PEX19	Peroxisome biogenesis disorder 12A (Zellweger), 614886	600279	67	100	100	98
PEX2	Peroxisome biogenesis disorder 5A (Zellweger), 614866 Peroxisome biogenesis disorder 5B, 614867	170993	63	100	100	100
PEX26	Peroxisome biogenesis disorder 7A (Zellweger), 614872 Peroxisome biogenesis disorder 7B, 614873	608666	118	100	100	100
PEX3	Peroxisome biogenesis disorder 10A (Zellweger), 614882 ?Peroxisome biogenesis disorder 10B, 617370	603164	57	100	100	95
PEX5	Peroxisome biogenesis disorder 2A (Zellweger), 214110 Peroxisome biogenesis disorder 2B, 202370 Rhizomelic chondrodysplasia punctata, type 5, 616716	600414	111	100	100	100
PEX6	Heimler syndrome 2, 616617 Peroxisome biogenesis disorder 4A (Zellweger), 614862 Peroxisome biogenesis disorder 4B, 614863	601498	107	100	99	95
PEX7	Peroxisome biogenesis disorder 9B, 614879 Rhizomelic chondrodysplasia punctata, type 1, 215100	601757	57	100	100	95
PFKM	Glycogen storage disease VII, 232800	610681	92	100	100	99
PFN1	Amyotrophic lateral sclerosis 18, 614808	176610	159	100	100	100
PGAM2	Glycogen storage disease X, 261670	612931	156	100	100	100
PGAP2	Hyperphosphatasia with mental retardation syndrome 3, 614207	615187	147	100	100	100
PGAP3	Hyperphosphatasia with mental retardation syndrome 4, 615716	611801	115	100	100	100
PGK1	Phosphoglycerate kinase 1 deficiency, 300653	311800	47	100	100	89
PGM1	Congenital disorder of glycosylation, type I α , 614921	171900	72	100	100	99
PHEX	Hypophosphatemic rickets dominant, 307800	300550	48	100	98	89
PHF6	Borjeson-Forssman-Lehmann syndrome, 301900	300414	51	100	96	82
PHF8	Mental retardation syndrome, Siderius type, 300263	300560	61	100	98	87

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PHGDH	Neu-Laxova syndrome 1, 256520 Phosphoglycerate dehydrogenase deficiency, 601815	606879	138	100	100	100
PHKA1	Muscle glycogenosis, 300559	311870	47	100	97	82
PHKA2	Glycogen storage disease, type IXa1, 306000 Glycogen storage disease, type IXa2, 306000	300798	56	100	99	90
PHKB	Phosphorylase kinase deficiency of liver and muscle, 261750	172490	76	100	100	98
PHKG2	Cirrhosis due to liver phosphorylase kinase deficiency Glycogen storage disease IXc, 613027	172471	123	100	100	100
PHOX2A	Fibrosis of extraocular muscles, congenital, 2, 602078	602753	61	100	100	86
PHOX2B	Central hypoventilation syndrome, congenital, with or without Hirschsprung disease, 209880 Neuroblastoma with Hirschsprung disease, 613013 {Neuroblastoma, susceptibility to, 2}, 613013	603851	149	100	100	100
PHRF1	No OMIM phenotype	611780	143	100	100	98
PHYH	Refsum disease, 266500	602026	121	100	100	94
PICALM	Leukemia, acute myeloid, somatic, 601626	603025	60	100	98	85
PIEZ01	Dehydrated hereditary stomatocytosis with or without pseudohyperkalemia and/or perinatal edema, 194380 Lymphatic malformation 6, 616843	611184	136	100	100	99
PIEZ02	Arthrogryposis, distal, type 3, 114300 Arthrogryposis, distal, type 5, 108145 Arthrogryposis, distal, with impaired proprioception and touch, 617146 ?Marden-Walker syndrome, 248700	613629	70	100	99	94
PIGA	Multiple congenital anomalies-hypotonia-seizures syndrome 2, 300868 Paroxysmal nocturnal hemoglobinuria, somatic, 300818	311770	67	100	100	97
PIGL	CHIME syndrome, 280000	605947	98	100	100	100
PIGM	Glycosylphosphatidylinositol deficiency, 610293	610273	85	100	100	100
PIGN	Multiple congenital anomalies-hypotonia-seizures syndrome 1, 614080	606097	63	100	99	89
PIGO	Hyperphosphatasia with mental retardation syndrome 2, 614749	614730	126	100	100	100
PIGV	Hyperphosphatasia with mental retardation syndrome 1, 239300	610274	96	100	100	100

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PIK3CA	Breast cancer, somatic, 114480 CLAPO syndrome, somatic, 613089 CLOVE syndrome, somatic, 612918 Colorectal cancer, somatic, 114500 Cowden syndrome 5, 615108 Gastric cancer, somatic, 613659 Hepatocellular carcinoma, somatic, 114550 Keratosis, seborrheic, somatic, 182000 Macrodactyly, somatic, 155500 Megalencephaly-capillary malformation-polymicrogyria syndrome, somatic, 602501 Nevus, epidermal, somatic, 162900 Nonsmall cell lung cancer, somatic, 211980 Ovarian cancer, somatic, 167000	171834	85	100	99	96
PIK3CD	Immunodeficiency 14, 615513	602839	137	100	99	98
PIK3R1	?Agammaglobulinemia 7, 615214 Immunodeficiency 36, 616005 SHORT syndrome, 269880	171833	80	100	100	97
PIK3R2	Megalencephaly-polymicrogyria-polydactyly-hydrocephalus syndrome 1, 603387	603157	89	95	93	90
PIK3R5	Ataxia-oculomotor apraxia 3, 615217	611317	107	100	100	100
PIKFYVE	Corneal fleck dystrophy, 121850	609414	67	100	99	95
PINK1	Parkinson disease 6, early onset, 605909	608309	100	98	93	88
PIP5K1C	Lethal congenital contractual syndrome 3, 611369	606102	130	100	95	95
PITPNM3	Cone-rod dystrophy 5, 600977	608921	99	100	99	97
PITX1	Clubfoot, congenital, with or without deficiency of long bones and/or mirror-image polydactyly, 119800 Liebenberg syndrome, 186550	602149	109	100	100	97
PITX2	Anterior segment dysgenesis 4, 137600 Axenfeld-Rieger syndrome, type 1, 180500 Ring dermoid of cornea, 180550	601542	149	100	100	100
PITX3	Anterior segment dysgenesis 1, multiple subtypes, 107250 Cataract 11, multiple types, 610623 Cataract 11, syndromic, 610623	602669	96	100	100	100
PJVK	Deafness 59, 610220	610219	93	100	100	99
PKD1	Polycystic kidney disease 1, 173900	601313	93	97	95	90
PKD1L1	Heterotaxy, visceral, 8, autosomal, 617205	609721	69	100	98	91
PKD2	Polycystic kidney disease 2, 613095	173910	62	100	97	89

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PKHD1	Polycystic kidney disease 4, with or without hepatic disease, 263200	606702	82	100	100	98
PKLR	Adenosine triphosphate, elevated, of erythrocytes, 102900 Pyruvate kinase deficiency, 266200	609712	156	100	100	100
PKP1	Ectodermal dysplasia/skin fragility syndrome, 604536	601975	116	100	100	99
PKP2	Arrhythmogenic right ventricular dysplasia 9, 609040	602861	147	100	98	96
PLA2G4A	Gastrointestinal ulceration, recurrent, with dysfunctional platelets, 618372	600522	68	100	100	96
PLA2G5	[Fleck retina, familial benign], 228980	601192	89	100	100	100
PLA2G6	Infantile neuroaxonal dystrophy 1, 256600 Neurodegeneration with brain iron accumulation 2B, 610217 Parkinson disease 14, 612953	603604	118	100	100	99
PLA2G7	{Asthma, susceptibility to}, 600807 {Atopy, susceptibility to}, 147050 Platelet-activating factor acetylhydrolase deficiency, 614278	601690	99	100	100	97
PLAG1	Adenomas, salivary gland pleomorphic, somatic, 181030	603026	74	100	100	100
PLAU	{Alzheimer disease, late-onset, susceptibility to}, 104300 Quebec platelet disorder, 601709	191840	149	100	100	100
PLCB1	Epileptic encephalopathy, early infantile, 12, 613722	607120	60	100	99	94
PLCB4	Auriculocondylar syndrome 2, 614669	600810	61	100	100	94
PLCD1	Nail disorder, nonsyndromic congenital, 3, (leukonychia), 151600	602142	115	100	100	100
PLCE1	Nephrotic syndrome, type 3, 610725	608414	79	100	100	97
PLCG2	Autoinflammation, antibody deficiency, and immune dysregulation syndrome, 614878 Familial cold autoinflammatory syndrome 3, 614468	600220	107	100	100	99
PLEC	Epidermolysis bullosa simplex with muscular dystrophy, 226670 ?Epidermolysis bullosa simplex with nail dystrophy, 616487 Epidermolysis bullosa simplex with pyloric atresia, 612138 Epidermolysis bullosa simplex, Ogna type, 131950 Muscular dystrophy, limb-girdle 17, 613723	601282	146	100	100	100
PLEKHG5	Charcot-Marie-Tooth disease, recessive intermediate C, 615376 Spinal muscular atrophy, distal, 4, 611067	611101	85	100	100	96
PLEKHM1	Osteopetrosis 3, 618107 ?Osteopetrosis 6, 611497	611466	174	100	100	99
PLG	Dysplasminogenemia, 217090 Plasminogen deficiency, type I, 217090	173350	81	100	100	98
PLIN1	Lipodystrophy, familial partial, type 4, 613877	170290	96	100	100	99

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PLN	Cardiomyopathy, dilated, 1P, 609909 Cardiomyopathy, hypertrophic, 18, 613874	172405	112	100	100	100
PLOD1	Ehlers-Danlos syndrome, kyphoscoliotic type, 1, 225400	153454	100	100	100	99
PLOD2	Bruck syndrome 2, 609220	601865	58	100	99	90
PLOD3	Lysyl hydroxylase 3 deficiency, 612394	603066	100	100	100	99
PLP1	Pelizaeus-Merzbacher disease, 312080 Spastic paraparesis 2, 312920	300401	93	100	99	97
PLS3	Bone mineral density QTL18, osteoporosis, 300910	300131	48	100	96	83
PML	Leukemia, acute promyelocytic, PML/RARA type	102578	136	100	100	100
PMM2	Congenital disorder of glycosylation, type Ia, 212065	601785	72	100	100	97
PMP22	Charcot-Marie-Tooth disease, type 1A, 118220 Charcot-Marie-Tooth disease, type 1E, 118300 Dejerine-Sottas disease, 145900 ?Neuropathy, inflammatory demyelinating, 139393 Neuropathy, recurrent, with pressure palsies, 162500 Roussy-Levy syndrome, 180800	601097	105	100	100	100
PMPCA	Spinocerebellar atrophy 2, 213200	613036	133	100	100	100
PMS2	Colorectal cancer, hereditary nonpolyposis, type 4, 614337 Mismatch repair cancer syndrome, 276300	600259	148	100	96	93
PNKP	Ataxia-oculomotor apraxia 4, 616267 Microcephaly, seizures, and developmental delay, 613402	605610	104	100	100	98
PNP	Immunodeficiency due to purine nucleoside phosphorylase deficiency, 613179	164050	70	100	100	97
PNPLA1	Ichthyosis, congenital 10, 615024	612121	102	100	100	99
PNPLA2	Neutral lipid storage disease with myopathy, 610717	609059	114	100	100	100
PNPLA6	Boucher-Neuhauser syndrome, 215470 ?Laurence-Moon syndrome, 245800 Oliver-McFarlane syndrome, 275400 Spastic paraparesis 39, 612020	603197	123	100	100	99
PNPO	Pyridoxamine 5'-phosphate oxidase deficiency, 610090	603287	69	100	100	94
PNPT1	Combined oxidative phosphorylation deficiency 13, 614932 Deafness 70, 614934	610316	57	100	97	83
POC1A	Short stature, onychodysplasia, facial dysmorphism, and hypotrichosis, 614813	614783	90	100	100	100
POF1B	?Premature ovarian failure 2B, 300604	300603	45	100	93	79
POFUT1	Dowling-Degos disease 2, 615327	607491	114	100	100	100

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POGLUT1	Dowling-Degos disease 4, 615696 ?Muscular dystrophy, limb-girdle 21, 617232	615618	54	100	98	91
POLD1	{Colorectal cancer, susceptibility to, 10}, 612591 Mandibular hypoplasia, deafness, progeroid features, and lipodystrophy syndrome, 615381	174761	114	100	99	96
POLE	{Colorectal cancer, susceptibility to, 12}, 615083 FILS syndrome, 615139 IMAGE-I syndrome, 618336	174762	117	100	100	100
POLG	Mitochondrial DNA depletion syndrome 4A (Alpers type), 203700 Mitochondrial DNA depletion syndrome 4B (MNGIE type), 613662 Mitochondrial recessive ataxia syndrome (includes SANDO and SCAE), 607459 Progressive external ophthalmoplegia 1, 157640 Progressive external ophthalmoplegia 1, 258450	174763	114	100	100	99
POLG2	Mitochondrial DNA depletion syndrome 16 (hepatic type), 618528 Progressive external ophthalmoplegia with mitochondrial DNA deletions 4, 610131	604983	140	100	100	95
POLH	Xeroderma pigmentosum, variant type, 278750	603968	74	100	100	98
POLR1C	Leukodystrophy, hypomyelinating, 11, 616494 Treacher Collins syndrome 3, 248390	610060	92	100	100	98
POLR1D	Treacher Collins syndrome 2, 613717	613715	81	100	100	100
POLR3A	Leukodystrophy, hypomyelinating, 7, with or without oligodontia and/or hypogonadotropic hypogonadism, 607694 Wiedemann-Rautenstrauch syndrome, 264090	614258	84	100	100	98
POLR3B	Leukodystrophy, hypomyelinating, 8, with or without oligodontia and/or hypogonadotropic hypogonadism, 614381	614366	76	100	99	93
POMC	Obesity, adrenal insufficiency, and red hair due to POMC deficiency, 609734 {Obesity, early-onset, susceptibility to}, 601665	176830	132	100	100	100
POMGNT1	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 3, 253280 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 3, 613151 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 3, 613157 Retinitis pigmentosa 76, 617123	606822	93	100	100	99
POMGNT2	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies, type A, 8, 614830 Muscular dystrophy-dystroglycanopathy (limb-girdle) type C, 8, 618135	614828	154	100	100	100
POMP	Keratosis linearis with ichthyosis congenita and sclerosing keratoderma, 601952 Proteasome-associated autoinflammatory syndrome 2, 618048	613386	45	100	100	87

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POMT1	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 1, 236670 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 1, 613155 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 1, 609308	607423	103	100	100	100
POMT2	Muscular dystrophy-dystroglycanopathy (congenital with brain and eye anomalies), type A, 2, 613150 Muscular dystrophy-dystroglycanopathy (congenital with mental retardation), type B, 2, 613156 Muscular dystrophy-dystroglycanopathy (limb-girdle), type C, 2, 613158	607439	87	100	100	99
POR	Antley-Bixler syndrome with genital anomalies and disordered steroidogenesis, 201750 Disordered steroidogenesis due to cytochrome P450 oxidoreductase, 613571	124015	170	100	100	100
PORCN	Focal dermal hypoplasia, 305600	300651	102	100	100	98
POU1F1	Pituitary hormone deficiency, combined, 1, 613038	173110	83	100	100	100
POU3F4	Deafness 2, 304400	300039	120	100	100	100
POU4F3	Deafness 15, 602459	602460	247	100	100	100
PPA2	?Sudden cardiac failure, alcohol-induced, 617223 Sudden cardiac failure, infantile, 617222	609988	70	100	94	76
PPARG	Carotid intimal medial thickness 1, 609338 {Diabetes, type 2}, 125853 Insulin resistance, severe, digenic, 604367 Lipodystrophy, familial partial, type 3, 604367 [Obesity, resistance to] Obesity, severe, 601665	601487	71	100	100	100
PPIB	Osteogenesis imperfecta, type IX, 259440	123841	96	100	100	100
PPM1D	Breast cancer, somatic, 114480 Jansen de Vries syndrome, 617450	605100	88	100	100	99
PPM1K	?Maple syrup urine disease, mild variant, 615135	611065	77	100	100	98
PPOX	Porphyria variegata, 176200	600923	106	100	100	100
PPP1CB	Noonan syndrome-like disorder with loose anagen hair 2, 617506	600590	54	100	100	95
PPP1R3A	Insulin resistance, severe, digenic, 125853	600917	66	100	100	98
PPP2R1B	Lung cancer, somatic, 211980	603113	77	100	100	96
PPP2R2B	Spinocerebellar ataxia 12, 604326	604325	77	100	99	95
PPT1	Ceroid lipofuscinosi, neuronal, 1, 256730	600722	84	100	100	100
PQBP1	Renpenning syndrome, 309500	300463	100	100	100	100
PRCC	Renal cell carcinoma, papillary, 605074	179755	114	100	100	96
PRCD	Retinitis pigmentosa 36, 610599	610598	83	100	100	100
PRDM16	Cardiomyopathy, dilated, 1LL, 615373 Left ventricular noncompaction 8, 615373	605557	191	100	100	99

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PRDM5	Brittle cornea syndrome 2, 614170	614161	76	100	100	96
PRDM6	Patent ductus arteriosus 3, 617039	616982	85	100	100	98
PRELID2	No OMIM phenotype	No ID	56	100	100	92
PRF1	Aplastic anemia, 609135 Hemophagocytic lymphohistiocytosis, familial, 2, 603553 Lymphoma, non-Hodgkin, 605027	170280	120	100	100	100
PRG4	Campodontaly-arthropathy-coxa vara-pericarditis syndrome, 208250	604283	157	100	100	100
PRICKLE1	Epilepsy, progressive myoclonic 1B, 612437	608500	76	100	100	98
PRICKLE2	No OMIM phenotype	608501	113	100	100	100
PRIMPOL	Myopia 22, 615420	615421	49	95	89	78
PRKAG2	Cardiomyopathy, hypertrophic 6, 600858 Glycogen storage disease of heart, lethal congenital, 261740 Wolff-Parkinson-White syndrome, 194200	602743	123	100	100	97
PRKAR1A	Acrody sostosis 1, with or without hormone resistance, 101800 Adrenocortical tumor, somatic Carney complex, type 1, 160980 Myxoma, intracardiac, 255960 Pigmented nodular adrenocortical disease, primary, 1, 610489	188830	86	100	100	99
PRKCA	Pituitary tumor, invasive	176960	126	100	100	100
PRKCG	Spinocerebellar ataxia 14, 605361	176980	122	100	100	99
PRKCSH	Polycystic liver disease 1, 174050	177060	142	100	100	98
PRKD1	Congenital heart defects and ectodermal dysplasia, 617364	605435	93	100	100	95
PRKG1	Aortic aneurysm, familial thoracic 8, 615436	176894	69	100	100	98
PRKN	Adenocarcinoma of lung, somatic, 211980 Ovarian cancer, somatic, 167000 Parkinson disease, juvenile, type 2, 600116	602544	96	100	100	99
PRKRA	Dystonia 16, 612067	603424	81	100	100	96
PRLR	Hyperprolactinemia, 615555 Multiple fibroadenomas of the breast, 615554	176761	67	100	100	99

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PRNP	Cerebral amyloid angiopathy, PRNP-related, 137440 Creutzfeldt-Jakob disease, 123400 Gerstmann-Straussler disease, 137440 Huntington disease-like 1, 603218 Insomnia, fatal familial, 600072 {Kuru, susceptibility to}, 245300 Prion disease with protracted course, 606688	176640	138	100	100	100
PROC	Thrombophilia due to protein C deficiency, 176860 Thrombophilia due to protein C deficiency, 612304	612283	149	100	100	100
PRODH	Hyperprolinemia, type I, 239500 {Schizophrenia, susceptibility to, 4}, 600850	606810	98	100	95	86
PROK2	Hypogonadotropic hypogonadism 4 with or without anosmia, 610628	607002	61	100	100	99
PROKR2	Hypogonadotropic hypogonadism 3 with or without anosmia, 244200	607123	210	100	100	100
PROM1	Cone-rod dystrophy 12, 612657 Macular dystrophy, retinal, 2, 608051 Retinitis pigmentosa 41, 612095 Stargardt disease 4, 603786	604365	84	100	99	93
PROP1	Pituitary hormone deficiency, combined, 2, 262600	601538	103	100	100	92
PROS1	Thrombophilia due to protein S deficiency, 612336 Thrombophilia due to protein S deficiency, 614514	176880	57	100	93	78
PRPF3	Retinitis pigmentosa 18, 601414	607301	77	100	98	94
PRPF31	Retinitis pigmentosa 11, 600138	606419	101	100	100	99
PRPF6	Retinitis pigmentosa 60, 613983	613979	108	100	100	99
PRPF8	Retinitis pigmentosa 13, 600059	607300	102	100	100	98
PRPH2	Choroidal dystrophy, central areolar 2, 613105 Leber congenital amaurosis 18, 608133 Macular dystrophy, patterned, 1, 169150 Macular dystrophy, vitelliform, 3, 608161 Retinitis pigmentosa 7 and digenic form, 608133 Retinitis punctata albescens, 136880	179605	162	100	100	100
PRPS1	Arts syndrome, 301835 Charcot-Marie-Tooth disease recessive, 5, 311070 Deafness 1, 304500 Gout, PRPS-related, 300661 Phosphoribosylpyrophosphate synthetase superactivity, 300661	311850	50	100	99	90

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PRRT2	Convulsions, familial infantile, with paroxysmal choreoathetosis, 602066 Episodic kinesigenic dyskinesia 1, 128200 Seizures, benign familial infantile, 2, 605751	614386	123	100	100	100
PRRX1	Agnathia-otocephaly complex, 202650	167420	94	100	100	100
PRSS1	Pancreatitis, hereditary, 167800 Trypsinogen deficiency, 614044	276000	239	100	100	100
PRSS12	Mental retardation 1, 249500	606709	94	100	100	99
PRSS56	Microphthalmia, isolated 6, 613517	613858	95	100	100	100
PRX	Charcot-Marie-Tooth disease, type 4F, 614895 Dejerine-Sottas disease, 145900	605725	198	100	100	100
PSAP	Combined SAP deficiency, 611721 Gaucher disease, atypical, 610539 Krabbe disease, atypical, 611722 Metachromatic leukodystrophy due to SAP-b deficiency, 249900	176801	121	100	100	97
PSAT1	Neu-Laxova syndrome 2, 616038 ?Phosphoserine aminotransferase deficiency, 610992	610936	64	100	100	97
PSENEN	Acne inversa, familial, 2, with or without Dowling-Degos disease, 613736	607632	88	100	100	100
PSMB8	Proteasome-associated autoinflammatory syndrome 1 and digenic forms, 256040	177046	124	100	100	100
PSMC3IP	Ovarian dysgenesis 3, 614324	608665	108	100	100	100
PSPH	Phosphoserine phosphatase deficiency, 614023	172480	51	100	96	84
PSTPIP1	Pyogenic sterile arthritis, pyoderma gangrenosum, and acne, 604416	606347	109	100	100	99
PTCH1	Basal cell carcinoma, somatic, 605462 Basal cell nevus syndrome, 109400 Holoprosencephaly 7, 610828	601309	91	100	98	96
PTCH2	Basal cell carcinoma, somatic, 605462 Basal cell nevus syndrome, 109400 Medulloblastoma, somatic, 155255	603673	108	100	100	100
PTDSS1	Lenz-Majewski hyperostotic dwarfism, 151050	612792	66	100	100	98
PTEN	Cowden syndrome 1, 158350 {Glioma susceptibility 2}, 613028 Lhermitte-Duclos syndrome, 158350 Macrocephaly/autism syndrome, 605309 {Meningioma}, 607174 Prostate cancer, somatic, 176807	601728	115	85	78	76

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PTF1A	Pancreatic agenesis 2, 615935 Pancreatic and cerebellar agenesis, 609069	607194	136	100	100	94
PTGIS	Hypertension, essential, 145500	601699	101	100	100	95
PTH	Hypoparathyroidism, 146200 Hypoparathyroidism, 146200	168450	85	100	94	88
PTH1R	Chondrodysplasia, Blomstrand type, 215045 Eiken syndrome, 600002 Failure of tooth eruption, primary, 125350 Metaphyseal chondrodysplasia, Murk Jansen type, 156400	168468	123	100	100	100
PTHLH	Brachydactyly, type E2, 613382	168470	137	100	100	100
PTPN11	LEOPARD syndrome 1, 151100 Leukemia, juvenile myelomonocytic, somatic, 607785 Metachondromatosis, 156250 Noonan syndrome 1, 163950	176876	77	100	99	90
PTPN12	Colon cancer, somatic, 114500	600079	56	100	99	94
PTPN14	Choanal atresia and lymphedema, 613611	603155	110	100	100	98
PTPRC	{Hepatitis C virus, susceptibility to}, 609532 Severe combined immunodeficiency, T cell-negative, B-cell/natural killer-cell positive, 608971	151460	73	100	99	95
PTPRJ	Colon cancer, somatic, 114500	600925	83	97	97	94
PTPRO	Nephrotic syndrome, type 6, 614196	600579	72	100	100	98
PTPRQ	Deafness 73, 617663 Deafness 84A, 613391	603317	80	100	99	95
PTS	Hyperphenylalaninemia, BH4-deficient, A, 261640	612719	86	100	100	95
PUF60	Verheij syndrome, 615583	604819	130	100	100	100
PUS1	Myopathy, lactic acidosis, and sideroblastic anemia 1, 600462	608109	88	100	99	95
PYCR1	Cutis laxa, type IIB, 612940 Cutis laxa, type IIIB, 614438	179035	93	100	100	100
PYGL	Glycogen storage disease VI, 232700	613741	84	100	100	96
PYGM	McArdle disease, 232600	608455	112	100	100	100
QARS1	Microcephaly, progressive, seizures, and cerebral and cerebellar atrophy, 615760	603727	108	100	100	100
QDPR	Hyperphenylalaninemia, BH4-deficient, C, 261630	612676	94	100	100	97
RAB18	Warburg micro syndrome 3, 614222	602207	85	100	100	97
RAB23	Carpenter syndrome, 201000	606144	104	100	100	100
RAB27A	Griscelli syndrome, type 2, 607624	603868	54	100	100	91
RAB28	Cone-rod dystrophy 18, 615374	612994	55	100	100	92

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RAB33B	Smith-McCort dysplasia 2, 615222	605950	74	100	100	100
RAB39B	Mental retardation 72, 300271 Waisman syndrome, 311510	300774	59	100	100	100
RAB3GAP1	Warburg micro syndrome 1, 600118	602536	67	100	100	97
RAB3GAP2	Martsolf syndrome, 212720 Warburg micro syndrome 2, 614225	609275	66	100	99	92
RAB40AL	No OMIM phenotype	300405	197	100	100	100
RAB7A	Charcot-Marie-Tooth disease, type 2B, 600882	602298	91	100	100	100
RAC2	Neutrophil immunodeficiency syndrome, 608203	602049	99	100	100	100
RAD21	Cornelia de Lange syndrome 4, 614701 ?Mungan syndrome, 611376	606462	69	100	99	92
RAD50	Nijmegen breakage syndrome-like disorder, 613078	604040	98	100	100	98
RAD51	{Breast cancer, susceptibility to}, 114480 ?Fanconi anemia, complementation group R, 617244 Mirror movements 2, 614508	179617	52	88	88	85
RAD51C	{Breast-ovarian cancer, familial, susceptibility to, 3}, 613399 Fanconi anemia, complementation group O, 613390	602774	64	100	100	93
RAD54B	Colon cancer, somatic, 114500 Lymphoma, non-Hodgkin, somatic, 605027	604289	89	100	99	95
RAD54L	Adenocarcinoma, colonic, somatic {Breast cancer, invasive ductal}, 114480 Lymphoma, non-Hodgkin, somatic, 605027	603615	95	100	100	99
RAF1	Cardiomyopathy, dilated, 1NN, 615916 LEOPARD syndrome 2, 611554 Noonan syndrome 5, 611553	164760	78	100	99	95
RAG1	Alpha/beta T-cell lymphopenia with gamma/delta T-cell expansion, severe cytomegalovirus infection, and autoimmunity, 609889 Combined cellular and humoral immune defects with granulomas, 233650 Omenn syndrome, 603554 Severe combined immunodeficiency, B cell-negative, 601457	179615	101	100	100	100
RAG2	Combined cellular and humoral immune defects with granulomas, 233650 Omenn syndrome, 603554 Severe combined immunodeficiency, B cell-negative, 601457	179616	71	100	100	100
RAI1	Smith-Magenis syndrome, 182290	607642	187	100	100	100
RAP1GDS1	Lymphocytic leukemia, acute T-cell	179502	75	100	100	98

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RAPSN	Fetal akinesia deformation sequence 2, 618388 Myasthenic syndrome, congenital, 11, associated with acetylcholine receptor deficiency, 616326	601592	137	100	100	100
RARB	Microphthalmia, syndromic 12, 615524	180220	85	100	100	99
RARS2	Pontocerebellar hypoplasia, type 6, 611523	611524	66	100	99	93
RASA1	Basal cell carcinoma, somatic, 605462 Capillary malformation-arteriovenous malformation 1, 608354	139150	61	100	98	90
RAX2	Cone-rod dystrophy 11, 610381 ?Macular degeneration, age-related, 6, 613757	610362	78	100	100	100
RB1	Bladder cancer, somatic, 109800 Osteosarcoma, somatic, 259500 Retinoblastoma, 180200 Retinoblastoma, trilateral, 180200 Small cell cancer of the lung, somatic, 182280	614041	163	100	100	98
RB1CC1	Breast cancer, somatic, 114480	606837	63	100	99	92
RBBP8	Jawad syndrome, 251255 Pancreatic carcinoma, somatic Seckel syndrome 2, 606744	604124	57	100	99	92
RBM10	TARP syndrome, 311900	300080	81	100	96	88
RBM20	Cardiomyopathy, dilated, 1DD, 613172	613171	136	100	100	100
RBM28	?Alopecia, neurologic defects, and endocrinopathy syndrome, 612079	612074	70	100	100	96
RBM8A	Thrombocytopenia-absent radius syndrome, 274000	605313	99	100	100	100
RBP4	Microphthalmia, isolated, with coloboma 10, 616428 Retinal dystrophy, iris coloboma, and comedogenic acne syndrome, 615147	180250	128	100	100	100
RBPJ	Adams-Oliver syndrome 3, 614814	147183	77	100	99	89
RD3	Leber congenital amaurosis 12, 610612	180040	153	100	100	100
RDH12	Leber congenital amaurosis 13, 612712	608830	104	100	100	100
RDH5	Fundus albipunctatus, 136880	601617	149	100	100	100
RDX	Deafness 24, 611022	179410	54	100	93	76
RECQL4	Baller-Gerold syndrome, 218600 RAPADILINO syndrome, 266280 Rothmund-Thomson syndrome, 268400	603780	148	100	100	99
REEP1	?Neuronopathy, distal hereditary motor, type VB, 614751 Spastic paraparesis 31, 610250	609139	72	100	100	99
RELN	{Epilepsy, familial temporal lobe, 7}, 616436 Lissencephaly 2 (Norman-Roberts type), 257320	600514	74	100	100	97

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REN	[Hyperproreninemia] Hyperuricemic nephropathy, familial juvenile 2, 613092 Renal tubular dysgenesis, 267430	179820	93	100	100	100
RET	Central hypoventilation syndrome, congenital, 209880 {Hirschsprung disease, protection against}, 142623 {Hirschsprung disease, susceptibility to, 1}, 142623 Medullary thyroid carcinoma, 155240 Multiple endocrine neoplasia IIA, 171400 Multiple endocrine neoplasia IIB, 162300 Pheochromocytoma, 171300	164761	173	100	100	100
RETREG1	Neuropathy, hereditary sensory and autonomic, type IIB, 613115	613114	73	100	98	86
RFT1	Congenital disorder of glycosylation, type IIn, 612015	611908	67	100	99	93
RFTN2	No OMIM phenotype	618215	73	100	98	92
RFX5	Bare lymphocyte syndrome, type II, complementation group C, 209920 Bare lymphocyte syndrome, type II, complementation group E, 209920	601863	100	100	100	98
RFX6	Mitchell-Riley syndrome, 615710	612659	91	100	99	95
RFXANK	MHC class II deficiency, complementation group B, 209920	603200	118	100	100	100
RFXAP	Bare lymphocyte syndrome, type II, complementation group D, 209920	601861	148	100	100	100
RGR	Retinitis pigmentosa 44, 613769	600342	101	100	98	91
RGS9	Bradyopsia, 608415	604067	108	100	100	98
RGS9BP	Bradyopsia, 608415	607814	153	100	100	100
RHAG	Anemia, hemolytic, Rh-null, regulator type, 268150 Overhydrated hereditary stomatocytosis, 185000	180297	62	100	100	94
RHBDF2	Tylosis with esophageal cancer, 148500	614404	100	100	99	96
RHCE	[Blood group, Rhesus] Rh-null disease, amorph type, 617970	111700	207	100	100	99
RHO	Night blindness, congenital stationary 1, 610445 Retinitis pigmentosa 4 or recessive, 613731 Retinitis punctata albescens, 136880	180380	131	100	100	100
RIMS1	Cone-rod dystrophy 7, 603649	606629	94	100	100	98
RIN2	Macrocephaly, alopecia, cutis laxa, and scoliosis, 613075	610222	102	100	100	99
RIPK4	CHAND syndrome, 214350 Popliteal pterygium syndrome, Bartsocas-Papas type, 263650	605706	171	100	100	100
RIT1	Noonan syndrome 8, 615355	609591	72	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
RLBP1	Bothnia retinal dystrophy, 607475 Fundus albipunctatus, 136880 Newfoundland rod-cone dystrophy, 607476 Retinitis punctata albescens, 136880	180090	111	100	100	100
RMND1	Combined oxidative phosphorylation deficiency 11, 614922	614917	67	100	100	93
RMRP	Anauxetic dysplasia 1, 607095 Cartilage-hair hypoplasia, 250250 Metaphyseal dysplasia without hypotrichosis, 250460	157660	No coverage data			
RNASEH2A	Aicardi-Goutieres syndrome 4, 610333	606034	109	100	100	100
RNASEH2B	Aicardi-Goutieres syndrome 2, 610181	610326	62	100	98	88
RNASEH2C	Aicardi-Goutieres syndrome 3, 610329	610330	323	100	100	100
RNASEL	Prostate cancer 1, 601518	180435	85	100	100	99
RNASET2	Leukoencephalopathy, cystic, without megalencephaly, 612951	612944	106	100	100	99
RNF123	No OMIM phenotype	614472	114	100	100	100
RNF135	No OMIM phenotype	611358	83	100	100	99
RNF139	Renal cell carcinoma, 144700	603046	91	100	100	100
RNF145	No OMIM phenotype	No ID	50	99	90	68
RNF168	RIDDLE syndrome, 611943	612688	93	100	100	99
RNF170	Ataxia, sensory, 1, 608984	614649	69	100	97	85
RNF212	Recombination rate QTL 1, 612042	612041	87	100	100	97
RNF216	Cerebellar ataxia and hypogonadotropic hypogonadism, 212840	609948	66	100	97	86
RNF6	Esophageal carcinoma, somatic, 133239	604242	73	100	100	99
ROBO2	Vesicoureteral reflux 2, 610878	602431	76	100	100	98
ROBO3	Gaze palsy, familial horizontal, with progressive scoliosis, 1, 607313	608630	106	100	100	100
ROBO4	Aortic valve disease 8, 618496	607528	91	100	100	98
ROGDI	Kohlschutter-Tonz syndrome, 226750	614574	90	100	98	93
ROM1	Retinitis pigmentosa 7, digenic form, 608133	180721	110	100	100	100
ROR2	Brachydactyly, type B1, 113000 Robinow syndrome, 268310	602337	134	100	100	99
RP1	Retinitis pigmentosa 1, 180100	603937	73	100	100	97
RP1L1	Occult macular dystrophy, 613587	608581	165	100	100	100
RP2	Retinitis pigmentosa 2, 312600	300757	67	100	100	98
RPE65	Leber congenital amaurosis 2, 204100 Retinitis pigmentosa 20, 613794	180069	72	100	100	98

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
RPGR	Cone-rod dystrophy, 1, 304020 Macular degeneration atrophic, 300834 Retinitis pigmentosa 3, 300029 Retinitis pigmentosa, and sinorespiratory infections, with or without deafness, 300455	312610	39	81	72	60
RPGRIP1	Cone-rod dystrophy 13, 608194 Leber congenital amaurosis 6, 613826	605446	88	100	100	96
RPGRIP1L	COACH syndrome, 216360 Joubert syndrome 7, 611560 Meckel syndrome 5, 611561	610937	60	98	96	89
RPIA	Ribose 5-phosphate isomerase deficiency, 608611	180430	77	100	100	98
RPL11	Diamond-Blackfan anemia 7, 612562	604175	62	100	100	100
RPL35A	Diamond-Blackfan anemia 5, 612528	180468	86	100	100	100
RPL5	Diamond-Blackfan anemia 6, 612561	603634	54	99	93	85
RPS10	Diamond-Blackfan anemia 9, 613308	603632	61	100	100	97
RPS14	Macrocytic anemia, refractory, due to 5q deletion, somatic, 153550	130620	134	100	100	100
RPS17	Diamond-Blackfan anemia 4, 612527	180472	2	5	5	5
RPS19	Diamond-Blackfan anemia 1, 105650	603474	97	100	100	92
RPS24	Diamond-blackfan anemia 3, 610629	602412	105	100	100	100
RPS26	Diamond-Blackfan anemia 10, 613309	603701	81	100	100	100
RPS6KA3	Coffin-Lowry syndrome, 303600 Mental retardation 19, 300844	300075	46	100	92	70
RPS7	Diamond-Blackfan anemia 8, 612563	603658	54	100	100	89
RPSA	Asplenia, isolated congenital, 271400	150370	84	100	100	100
RRAS2	Ovarian carcinoma	600098	68	100	99	86
RRM2B	Mitochondrial DNA depletion syndrome 8A (encephalomyopathic type with renal tubulopathy), 612075 Mitochondrial DNA depletion syndrome 8B (MNGIE type), 612075 Progressive external ophthalmoplegia with mitochondrial DNA deletions 5, 613077	604712	90	100	100	100
RS1	Retinoschisis, 312700	300839	44	100	99	74
RSPH1	Ciliary dyskinesia, primary, 24, 615481	609314	80	100	100	95
RSPH4A	Ciliary dyskinesia, primary, 11, 612649	612647	100	100	100	100
RSPH9	Ciliary dyskinesia, primary, 12, 612650	612648	125	100	100	100
RSPO1	Palmoplantar hyperkeratosis and true hermaphroditism, 610644 Palmoplantar hyperkeratosis with squamous cell carcinoma of skin and sex reversal, 610644	609595	93	100	100	100
RSPO4	Anonychia congenita, 206800	610573	116	100	100	100

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RTEL1	Dyskeratosis congenita 4, 615190 Dyskeratosis congenita 5, 615190 Pulmonary fibrosis and/or bone marrow failure, telomere-related, 3, 616373	608833	131	100	100	99
RTL4	No OMIM phenotype	No ID	No coverage data			
RTN2	Spastic paraplegia 12, 604805	603183	135	100	100	100
RTTN	Microcephaly, short stature, and polymicrogyria with seizures, 614833	610436	71	100	99	94
RUNX1	Leukemia, acute myeloid, 601626 Platelet disorder, familial, with associated myeloid malignancy, 601399	151385	76	100	100	98
RUNX2	Cleidocranial dysplasia, 119600 Cleidocranial dysplasia, forme fruste, dental anomalies only, 119600 Cleidocranial dysplasia, forme fruste, with brachydactyly, 119600 Metaphyseal dysplasia with maxillary hypoplasia with or without brachydactyly, 156510	600211	107	100	100	100
RXFP2	No OMIM phenotype	606655	54	100	99	90
RYR1	Central core disease, 117000 King-Denborough syndrome, 145600 {Malignant hyperthermia susceptibility 1}, 145600 Minicore myopathy with external ophthalmoplegia, 255320 Neuromuscular disease, congenital, with uniform type 1 fiber, 117000	180901	152	99	99	99
RYR2	Arrhythmogenic right ventricular dysplasia 2, 600996 Ventricular tachycardia, catecholaminergic polymorphic, 1, 604772	180902	128	100	100	97
SACS	Spastic ataxia, Charlevoix-Saguenay type, 270550	604490	70	100	100	99
SAG	Oguchi disease-1, 258100 Retinitis pigmentosa 47, 613758	181031	97	100	100	98
SALL1	Townes-Brocks branchiootorenal-like syndrome, 107480 Townes-Brocks syndrome 1, 107480	602218	120	100	100	99
SALL4	Duane-radial ray syndrome, 607323 IVIC syndrome, 147750	607343	155	100	100	97
SAMD9	MIRAGE syndrome, 617053 Tumoral calcinosis, familial, normophosphatemic, 610455	610456	68	100	100	100
SAMHD1	Aicardi-Goutieres syndrome 5, 612952 ?Chilblain lupus 2, 614415	606754	64	100	99	88
SAR1B	Chylomicron retention disease, 246700	607690	79	100	100	97
SARS2	Hyperuricemia, pulmonary hypertension, renal failure, and alkalosis, 613845	612804	86	100	100	99
SART3	No OMIM phenotype	611684	94	100	100	97
SAT1	No OMIM phenotype	313020	66	100	100	97

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
SATB2	Glass syndrome, 612313	608148	95	100	100	97
SBDS	{Aplastic anemia, susceptibility to}, 609135 Shwachman-Diamond syndrome, 260400	607444	79	100	100	99
SBF2	Charcot-Marie-Tooth disease, type 4B2, 604563	607697	72	100	100	95
SC5D	Lathosterolemia, 607330	602286	88	100	100	100
SCARB2	Epilepsy, progressive myoclonic 4, with or without renal failure, 254900	602257	89	100	100	98
SCARF2	Van den Ende-Gupta syndrome, 600920	613619	101	100	99	97
SCN10A	Episodic pain syndrome, familial, 2, 615551	604427	110	100	100	99
SCN11A	Episodic pain syndrome, familial, 3, 615552 Neuropathy, hereditary sensory and autonomic, type VII, 615548	604385	88	100	100	97
SCN1A	Epilepsy, generalized, with febrile seizures plus, type 2, 604403 Epileptic encephalopathy, early infantile, 6 (Dravet syndrome), 607208 Febrile seizures, familial, 3A, 604403 Migraine, familial hemiplegic, 3, 609634	182389	92	100	100	99
SCN1B	Atrial fibrillation, familial, 13, 615377 Brugada syndrome 5, 612838 Cardiac conduction defect, nonspecific, 612838 Epilepsy, generalized, with febrile seizures plus, type 1, 604233 Epileptic encephalopathy, early infantile, 52, 617350	600235	169	100	97	93
SCN2A	Epileptic encephalopathy, early infantile, 11, 613721 Seizures, benign familial infantile, 3, 607745	182390	95	100	100	97
SCN2B	Atrial fibrillation, familial, 14, 615378	601327	117	100	100	100
SCN3A	Epilepsy, familial focal, with variable foci 4, 617935 Epileptic encephalopathy, early infantile, 62, 617938	182391	84	100	100	97
SCN3B	Atrial fibrillation, familial, 16, 613120 Brugada syndrome 7, 613120	608214	92	100	100	100
SCN4A	Hyperkalemic periodic paralysis, type 2, 170500 Hypokalemic periodic paralysis, type 2, 613345 Myasthenic syndrome, congenital, 16, 614198 Myotonia congenita, atypical, acetazolamide-responsive, 608390 Paramyotonia congenita, 168300	603967	143	100	100	100
SCN4B	Atrial fibrillation, familial, 17, 611819 Long QT syndrome-10, 611819	608256	111	100	100	100

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SCN5A	Atrial fibrillation, familial, 10, 614022 Brugada syndrome 1, 601144 Cardiomyopathy, dilated, 1E, 601154 Heart block, nonprogressive, 113900 Heart block, progressive, type IA, 113900 Long QT syndrome-3, 603830 Sick sinus syndrome 1, 608567 {Sudden infant death syndrome, susceptibility to}, 272120 Ventricular fibrillation, familial, 1, 603829	600163	192	100	100	100
SCN8A	Cognitive impairment with or without cerebellar ataxia, 614306 Epileptic encephalopathy, early infantile, 13, 614558 ?Myoclonus, familial, 2, 618364 Seizures, benign familial infantile, 5, 617080	600702	110	100	100	98
SCN9A	{Dravet syndrome, modifier of}, 607208 Epilepsy, generalized, with febrile seizures plus, type 7, 613863 Erythermalgia, primary, 133020 Febrile seizures, familial, 3B, 613863 HSAN2D, 243000 Insensitivity to pain, congenital, 243000 Paroxysmal extreme pain disorder, 167400 Small fiber neuropathy, 133020	603415	84	100	100	98
SCNN1A	Bronchiectasis with or without elevated sweat chloride 2, 613021 ?Liddle syndrome 3, 618126 Pseudohypoaldosteronism, type I, 264350	600228	108	100	100	98
SCNN1B	Bronchiectasis with or without elevated sweat chloride 1, 211400 Liddle syndrome 1, 177200 Pseudohypoaldosteronism, type I, 264350	600760	109	100	100	99
SCNN1G	Bronchiectasis with or without elevated sweat chloride 3, 613071 Liddle syndrome 2, 618114 Pseudohypoaldosteronism, type I, 264350	600761	122	100	100	98
SCO1	Mitochondrial complex IV deficiency, 220110	603644	99	100	100	99
SCO2	Cardioencephalomyopathy, fatal infantile, due to cytochrome c oxidase deficiency 1, 604377 Myopia 6, 608908	604272	125	100	100	100
SCP2	?Leukoencephalopathy with dystonia and motor neuropathy, 613724	184755	54	100	100	92

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SDCCAG8	Bardet-Biedl syndrome 16, 615993 Senior-Loken syndrome 7, 613615	613524	92	100	100	96
SDHA	Cardiomyopathy, dilated, 1GG, 613642 Leigh syndrome, 256000 Mitochondrial respiratory chain complex II deficiency, 252011 Paragangliomas 5, 614165	600857	134	100	97	91
SDHAF1	Mitochondrial complex II deficiency, 252011	612848	82	100	100	100
SDHAF2	Paragangliomas 2, 601650	613019	140	100	100	100
SDHB	Gastrointestinal stromal tumor, 606764 Paraganglioma and gastric stromal sarcoma, 606864 Paragangliomas 4, 115310 Pheochromocytoma, 171300	185470	149	100	100	100
SDHC	Gastrointestinal stromal tumor, 606764 Paraganglioma and gastric stromal sarcoma, 606864 Paragangliomas 3, 605373	602413	170	100	100	100
SDHD	Mitochondrial complex II deficiency, 252011 Paraganglioma and gastric stromal sarcoma, 606864 Paragangliomas 1, with or without deafness, 168000 Pheochromocytoma, 171300	602690	164	100	100	100
SEC23A	Craniolenticulosutural dysplasia, 607812	610511	61	100	97	90
SEC23B	?Cowden syndrome 7, 616858 Dyserythropoietic anemia, congenital, type II, 224100	610512	74	100	100	97
SEC63	Polycystic liver disease 2, 617004	608648	54	100	99	87
SECISBP2	Thyroid hormone metabolism, abnormal, 609698	607693	82	100	100	95
SELENON	Muscular dystrophy, rigid spine, 1, 602771 Myopathy, congenital, with fiber-type disproportion, 255310	606210	132	90	90	90
SEMA3E	?CHARGE syndrome, 214800	608166	64	100	100	94
SEMA4A	Cone-rod dystrophy 10, 610283 Retinitis pigmentosa 35, 610282	607292	108	100	100	100
SEPSECS	Pontocerebellar hypoplasia type 2D, 613811	613009	78	100	100	96
SEPTIN12	Spermatogenic failure 10, 614822	611562	85	100	100	99
SEPTIN9	Amyotrophy, hereditary neuralgic, 162100 Leukemia, acute myeloid, therapy-related Ovarian carcinoma	604061	125	100	100	100
SERAC1	3-methylglutaconic aciduria with deafness, encephalopathy, and Leigh-like syndrome, 614739	614725	65	100	99	86

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SERPINA1	Emphysema due to AAT deficiency, 613490 Emphysema-cirrhosis, due to AAT deficiency, 613490 Hemorrhagic diathesis due to antithrombin Pittsburgh, 613490 {Pulmonary disease, chronic obstructive, susceptibility to}, 606963	107400	90	100	100	100
SERPINA3	Alpha-1-antichymotrypsin deficiency Cerebrovascular disease, occlusive	107280	106	100	100	100
SERPINA6	Corticosteroid-binding globulin deficiency, 611489	122500	106	100	100	100
SERPINA7	[Thyroxine-binding globulin QTL], 300932	314200	50	100	98	85
SERPINB6	?Deafness 91, 613453	173321	85	100	100	98
SERPINB7	Palmoplantar keratoderma, Nagashima type, 615598	603357	70	100	100	97
SERPINCl	Thrombophilia due to antithrombin III deficiency, 613118	107300	96	100	100	99
SERPIND1	Thrombophilia due to heparin cofactor II deficiency, 612356	142360	77	100	100	99
SERPINE1	Plasminogen activator inhibitor-1 deficiency, 613329 {Transcription of plasminogen activator inhibitor, modulator of}	173360	132	100	100	100
SERPINF1	Osteogenesis imperfecta, type VI, 613982	172860	111	100	99	95
SERPINF2	Alpha-2-plasmin inhibitor deficiency, 262850	613168	150	100	100	98
SERPING1	Angioedema, hereditary, types I and II, 106100 Complement component 4, partial deficiency of, 120790	606860	112	100	100	99
SERPINH1	Osteogenesis imperfecta, type X, 613848 {Preterm premature rupture of the membranes, susceptibility to}, 610504	600943	159	100	100	100
SERPINI1	Encephalopathy, familial, with neuroserpin inclusion bodies, 604218	602445	73	100	100	98
SETBP1	Mental retardation 29, 616078 Schinzel-Giedion midface retraction syndrome, 269150	611060	102	99	98	97
SETD1A	No OMIM phenotype	611052	135	100	99	98
SETD5	Mental retardation 23, 615761	615743	85	100	100	98
SETX	Amyotrophic lateral sclerosis 4, juvenile, 602433 Spinocerebellar ataxia, with axonal neuropathy 2, 606002	608465	70	100	100	97
SF3B1	Myelodysplastic syndrome, somatic, 614286	605590	69	100	100	94
SF3B4	Acrofacial dysostosis 1, Nager type, 154400	605593	84	100	100	99
SFTPA2	Pulmonary fibrosis, idiopathic, 178500	178642	191	100	100	100
SFTPB	Surfactant metabolism dysfunction, pulmonary, 1, 265120	178640	86	100	100	99
SFTPC	Surfactant metabolism dysfunction, pulmonary, 2, 610913	178620	100	100	100	100
SFXN4	Combined oxidative phosphorylation deficiency 18, 615578	615564	74	100	100	99
SGCA	Muscular dystrophy, limb-girdle 3, 608099	600119	149	100	98	96
SGCB	Muscular dystrophy, limb-girdle 4, 604286	600900	54	100	95	89

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SGCD	Cardiomyopathy, dilated, 1L, 606685 Muscular dystrophy, limb-girdle 6, 601287	601411	60	100	100	97
SGCE	Dystonia-11, myoclonic, 159900	604149	84	100	100	94
SGCG	Muscular dystrophy, limb-girdle 5, 253700	608896	78	100	100	98
SGSH	Mucopolysaccharidosis type IIIA (Sanfilippo A), 252900	605270	103	100	94	90
SH2B3	Erythrocytosis, somatic, 133100 Myelofibrosis, somatic, 254450 Thrombocythemia, somatic, 187950	605093	115	100	100	100
SH2D1A	Lymphoproliferative syndrome, 1, 308240	300490	56	100	98	83
SH3BP2	Cherubism, 118400	602104	127	91	91	91
SH3PXD2B	Frank-ter Haar syndrome, 249420	613293	122	100	100	99
SH3TC2	Charcot-Marie-Tooth disease, type 4C, 601596 Mononeuropathy of the median nerve, mild, 613353	608206	89	100	100	99
SHANK3	Phelan-McDermid syndrome, 606232 {Schizophrenia 15}, 613950	606230	117	99	93	86
SHH	Holoprosencephaly 3, 142945 Microphthalmia with coloboma 5, 611638 Schizencephaly, 269160 Single median maxillary central incisor, 147250	600725	119	100	100	99
SHOC2	Noonan-like syndrome with loose anagen hair, 607721	602775	61	100	99	95
SHOX	Langer mesomelic dysplasia, 249700 Leri-Weill dyschondrosteosis, 127300 Short stature, idiopathic familial, 300582	312865	0	0	0	0
SHROOM4	Stocco dos Santos X-linked mental retardation syndrome, 300434	300579	70	100	100	97
SI	Sucrase-isomaltase deficiency, congenital, 222900	609845	60	100	99	92
SIGMAR1	?Amyotrophic lateral sclerosis 16, juvenile, 614373 ?Spinal muscular atrophy, distal, 2, 605726	601978	135	100	100	100
SIL1	Marinesco-Sjögren syndrome, 248800	608005	100	100	100	100
SIM1	No OMIM phenotype	603128	99	100	100	99
SIX1	Branchioototic syndrome 3, 608389 Deafness 23, 605192	601205	119	100	100	100
SIX3	Holoprosencephaly 2, 157170 Schizencephaly, 269160	603714	158	100	100	96
SIX5	Branchiootorenal syndrome 2, 610896	600963	75	100	95	85
SIX6	Optic disc anomalies with retinal and/or macular dystrophy, 212550	606326	233	100	100	100

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SKI	Shprintzen-Goldberg syndrome, 182212	164780	123	100	100	99
SKIV2L	Trichohepatoenteric syndrome 2, 614602	600478	124	100	100	100
SLC10A2	Bile acid malabsorption, primary, 613291	601295	78	100	100	96
SLC11A2	Anemia, hypochromic microcytic, with iron overload 1, 206100	600523	57	100	99	90
SLC12A1	Bartter syndrome, type 1, 601678	600839	72	100	100	98
SLC12A3	Gitelman syndrome, 263800	600968	113	100	100	100
SLC12A6	Agenesis of the corpus callosum with peripheral neuropathy, 218000	604878	67	100	99	94
SLC16A1	Erythrocyte lactate transporter defect, 245340	600682	83	100	100	99
	Hyperinsulinemic hypoglycemia, familial, 7, 610021					
	Monocarboxylate transporter 1 deficiency, 616095					
SLC16A12	Cataract 47, juvenile, with microcornea, 612018	611910	89	100	100	100
SLC16A2	Allan-Herndon-Dudley syndrome, 300523	300095	75	100	100	95
SLC17A5	Salla disease, 604369	604322	86	100	100	97
	Sialic acid storage disorder, infantile, 269920					
SLC17A8	Deafness 25, 605583	607557	66	100	100	99
SLC19A2	Thiamine-responsive megaloblastic anemia syndrome, 249270	603941	68	100	100	99
SLC19A3	Thiamine metabolism dysfunction syndrome 2 (biotin- or thiamine-responsive encephalopathy type 2), 607483	606152	73	100	100	98
SLC1A3	Episodic ataxia, type 6, 612656	600111	99	100	100	100
SLC20A2	Basal ganglia calcification, idiopathic, 1, 213600	158378	99	100	100	95
SLC22A12	Hypouricemia, renal, 220150	607096	117	100	100	97
SLC22A18	Breast cancer, somatic, 114480	602631	107	100	100	98
	Lung cancer, somatic, 211980					
	Rhabdomyosarcoma, somatic, 268210					
SLC22A5	Carnitine deficiency, systemic primary, 212140	603377	118	100	100	100
SLC24A1	Night blindness, congenital stationary (complete), 1D, 613830	603617	105	100	100	99
SLC24A5	Albinism, oculocutaneous, type VI, 113750	609802	77	100	100	100
	[Skin/hair/eye pigmentation 4, fair/dark skin], 113750					
SLC25A1	Combined D-2- and L-2-hydroxyglutaric aciduria, 615182	190315	102	100	100	100
	?Myasthenic syndrome, congenital, 23, presynaptic, 618197					
SLC25A12	Epileptic encephalopathy, early infantile, 39, 612949	603667	88	100	100	97
SLC25A13	Citrullinemia, adult-onset type II, 603471	603859	98	100	99	93
	Citrullinemia, type II, neonatal-onset, 605814					
SLC25A15	Hyperornithinemia-hyperammonemia-homocitrullinemia syndrome, 238970	603861	132	100	100	98

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SLC25A19	Microcephaly, Amish type, 607196 Thiamine metabolism dysfunction syndrome 4 (progressive polyneuropathy type), 613710	606521	105	100	100	100
SLC25A20	Carnitine-acylcarnitine translocase deficiency, 212138	613698	66	100	100	97
SLC25A22	Epileptic encephalopathy, early infantile, 3, 609304	609302	122	100	100	100
SLC25A3	Mitochondrial phosphate carrier deficiency, 610773	600370	105	100	100	100
SLC25A38	Anemia, sideroblastic, 2, pyridoxine-refractory, 205950	610819	109	100	100	100
SLC25A4	Mitochondrial DNA depletion syndrome 12A (cardiomyopathic type) AD, 617184	103220	112	100	100	100
	Mitochondrial DNA depletion syndrome 12B (cardiomyopathic type) AR, 615418					
	Progressive external ophthalmoplegia with mitochondrial DNA deletions 2, 609283					
SLC26A2	Achondrogenesis Ib, 600972	606718	68	100	100	100
	Atelosteogenesis, type II, 256050					
	De la Chapelle dysplasia, 256050					
	Diastrophic dysplasia, 222600					
	Diastrophic dysplasia, broad bone-platyspondylic variant, 222600					
SLC26A3	Epiphyseal dysplasia, multiple, 4, 226900	126650	72	100	100	95
	Diarrhea 1, secretory chloride, congenital, 214700					
	Deafness 4, with enlarged vestibular aqueduct, 600791					
SLC26A4	Pendred syndrome, 274600	605646	68	100	100	94
	?Deafness 61, 613865					
SLC26A8	Spermatogenic failure 3, 606766	608480	64	100	99	93
SLC27A4	Ichthyosis prematurity syndrome, 608649	604194	133	100	100	100
SLC29A3	Histiocytosis-lymphadenopathy plus syndrome, 602782	612373	152	100	99	99
SLC2A1	Dystonia 9, 601042	138140	127	100	100	100
	{Epilepsy, idiopathic generalized, susceptibility to, 12}, 614847					
	GLUT1 deficiency syndrome 1, infantile onset, severe, 606777					
	GLUT1 deficiency syndrome 2, childhood onset, 612126					
	Stomatin-deficient cryohydrocytosis with neurologic defects, 608885					
SLC2A10	Arterial tortuosity syndrome, 208050	606145	125	100	100	100
SLC2A2	{Diabetes mellitus, noninsulin-dependent}, 125853	138160	63	100	100	96
	Fanconi-Bickel syndrome, 227810					
SLC2A9	Hypouricemia, renal, 2, 612076	606142	107	100	100	97
	{Uric acid concentration, serum, QTL 2}, 612076					
SLC30A10	Hypermanganesemia with dystonia 1, 613280	611146	148	100	100	100
SLC30A2	Zinc deficiency, transient neonatal, 608118	609617	104	100	100	100

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SLC33A1	Congenital cataracts, hearing loss, and neurodegeneration, 614482 Spastic paraparesis 42, 612539	603690	68	100	98	86
SLC34A1	?Fanconi renotubular syndrome 2, 613388 Hypercalcemia, infantile, 2, 616963 Nephrolithiasis/osteoporosis, hypophosphatemic, 1, 612286	182309	136	100	100	100
SLC34A2	Pulmonary alveolar microlithiasis, 265100	604217	120	100	100	99
SLC34A3	Hypophosphatemic rickets with hypercalciuria, 241530	609826	121	100	98	93
SLC35A1	Congenital disorder of glycosylation, type IIf, 603585	605634	66	100	100	97
SLC35A2	Congenital disorder of glycosylation, type IIm, 300896, Somatic mosaicism	314375	75	100	100	99
SLC35C1	Congenital disorder of glycosylation, type IIC, 266265	605881	146	100	100	100
SLC35D1	Schneckenbecken dysplasia, 269250	610804	57	100	99	85
SLC36A2	Hyperglycinuria, 138500 Iminoglycinuria, digenic, 242600	608331	102	100	100	100
SLC37A4	Glycogen storage disease Ib, 232220 Glycogen storage disease Ic, 232240	602671	96	100	100	95
SLC38A8	Foveal hypoplasia 2, with or without optic nerve misrouting and/or anterior segment dysgenesis, 609218	615585	83	100	99	94
SLC39A13	Ehlers-Danlos syndrome, spondylodysplastic type, 3, 612350	608735	129	100	100	100
SLC39A4	Acrodermatitis enteropathica, 201100	607059	125	100	100	100
SLC3A1	Cystinuria, 220100	104614	93	100	100	97
SLC40A1	Hemochromatosis, type 4, 606069	604653	63	100	100	97
SLC45A2	Albinism, oculocutaneous, type IV, 606574 [Skin/hair/eye pigmentation 5, black/nonblack hair], 227240 [Skin/hair/eye pigmentation 5, dark/fair skin], 227240 [Skin/hair/eye pigmentation 5, dark/light eyes], 227240	606202	93	100	100	97
SLC46A1	Folate malabsorption, hereditary, 229050	611672	121	100	100	98

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SLC4A1	[Blood group, Diego], 110500 [Blood group, Froese], 601551 [Blood group, Swann], 601550 [Blood group, Waldner], 112010 [Blood group, Wright], 112050 Cryohydrocytosis, 185020 [Malaria, resistance to], 611162 Ovalocytosis, SA type, 166900 Renal tubular acidosis, distal, AD, 179800 Renal tubular acidosis, distal, AR, 611590 Spherocytosis, type 4, 612653	109270	122	100	100	100
SLC4A11	Corneal dystrophy, Fuchs endothelial, 4, 613268 Corneal endothelial dystrophy and perceptive deafness, 217400 Corneal endothelial dystrophy, 217700	610206	125	100	100	99
SLC4A4	Renal tubular acidosis, proximal, with ocular abnormalities, 604278	603345	61	100	100	97
SLC52A3	Brown-Vialetto-Van Laere syndrome 1, 211530 ?Fazio-Londe disease, 211500	613350	109	100	100	100
SLC5A1	Glucose/galactose malabsorption, 606824	182380	104	100	100	96
SLC5A2	Renal glucosuria, 233100	182381	113	100	100	99
SLC5A5	Thyroid dyshormonogenesis 1, 274400	601843	94	100	99	95
SLC5A7	Myasthenic syndrome, congenital, 20, presynaptic, 617143 Neuronopathy, distal hereditary motor, type VIIA, 158580	608761	82	100	100	99
SLC6A19	Hartnup disorder, 234500 Hyperglycinuria, 138500 Iminoglycinuria, digenic, 242600	608893	115	100	100	100
SLC6A2	?Orthostatic intolerance, 604715	163970	101	100	100	100
SLC6A20	Hyperglycinuria, 138500 Iminoglycinuria, digenic, 242600	605616	139	100	100	100
SLC6A3	{Nicotine dependence, protection against}, 188890 Parkinsonism-dystonia, infantile, 1, 613135	126455	102	100	100	100
SLC6A5	Hyperekplexia 3, 614618	604159	85	100	100	96
SLC6A8	Cerebral creatine deficiency syndrome 1, 300352	300036	103	100	98	95
SLC7A14	Retinitis pigmentosa 68, 615725	615720	112	100	100	100
SLC7A7	Lysinuric protein intolerance, 222700	603593	79	100	100	97
SLC7A9	Cystinuria, 220100	604144	100	100	100	99

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SLC9A3R1	Nephrolithiasis/osteoporosis, hypophosphatemic, 2, 612287	604990	116	100	100	100
SLC9A6	Mental retardation syndromic, Christianson type, 300243	300231	64	100	96	84
SLCO1B1	Hyperbilirubinemia, Rotor type, digenic, 237450	604843	61	100	99	90
SLCO1B3	Hyperbilirubinemia, Rotor type, digenic, 237450	605495	66	100	99	92
SLCO2A1	Hypertrophic osteoarthropathy, primary 2, 614441	601460	97	100	100	99
SLTRK1	Tourette syndrome, 137580 ?Trichotillomania, 613229	609678	89	100	100	100
SLTRK6	Deafness and myopia, 221200	609681	75	100	100	99
SLURP1	Meleda disease, 248300	606119	98	100	100	100
SLX4	Fanconi anemia, complementation group P, 613951	613278	118	100	100	99
SMAD2	No OMIM phenotype	601366	69	100	100	97
SMAD3	Loeys-Dietz syndrome 3, 613795	603109	188	100	100	100
SMAD4	Juvenile polyposis/hereditary hemorrhagic telangiectasia syndrome, 175050 Myhre syndrome, 139210 Pancreatic cancer, somatic, 260350 Polyposis, juvenile intestinal, 174900	600993	79	100	100	98
SMAD6	Aortic valve disease 2, 614823 {Craniosynostosis 7, susceptibility to}, 617439	602931	165	100	97	87
SMAD9	Pulmonary hypertension, primary, 2, 615342	603295	85	100	100	99
SMARCA2	Nicolaides-Baraitser syndrome, 601358	600014	94	98	98	96
SMARCA4	Coffin-Siris syndrome 4, 614609 {Rhabdoid tumor predisposition syndrome 2}, 613325	603254	127	100	100	100
SMARCAD1	Adermatoglyphia, 136000 Basan syndrome, 129200 Huriez syndrome, 181600	612761	61	100	98	89
SMARCAL1	Schimke immunoosseous dysplasia, 242900	606622	88	100	100	98
SMARCB1	Coffin-Siris syndrome 3, 614608 {Rhabdoid tumor predisposition syndrome 1}, 609322 Rhabdoid tumors, somatic, 609322 {Schwannomatosis-1, susceptibility to}, 162091	601607	130	100	100	99
SMARCC2	Coffin-Siris syndrome 8, 618362	601734	74	100	100	94
SMC1A	Cornelia de Lange syndrome 2, 300590	300040	71	100	100	97
SMC3	Cornelia de Lange syndrome 3, 610759	606062	69	100	97	88
SMCHD1	Bosma arhinia microphthalmia syndrome, 603457 Fascioscapulohumeral muscular dystrophy 2, digenic, 158901	614982	63	100	98	89

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SMN1	Spinal muscular atrophy-1, 253300 Spinal muscular atrophy-2, 253550 Spinal muscular atrophy-3, 253400 Spinal muscular atrophy-4, 271150	600354	4	7	7	7
SMO	Basal cell carcinoma, somatic, 605462 Curry-Jones syndrome, somatic mosaic, 601707	601500	124	100	100	100
SMOC1	Microphthalmia with limb anomalies, 206920	608488	106	100	100	97
SMOC2	Dentin dysplasia, type I, with microdontia and misshapen teeth, 125400	607223	88	100	99	94
SMPD1	Niemann-Pick disease, type A, 257200 Niemann-Pick disease, type B, 607616	607608	131	100	100	99
SMPX	Deafness 4, 300066	300226	51	100	100	83
SMS	Mental retardation, Snyder-Robinson type, 309583	300105	42	95	89	68
SNAI2	Piebaldism, 172800 Waardenburg syndrome, type 2D, 608890	602150	66	100	100	100
SNAP29	Cerebral dysgenesis, neuropathy, ichthyosis, and palmoplantar keratoderma syndrome, 609528	604202	148	100	100	100
SNCA	Dementia, Lewy body, 127750 Parkinson disease 1, 168601 Parkinson disease 4, 605543	163890	52	100	100	98
SNCB	Dementia, Lewy body, 127750	602569	130	100	100	100
SNIP1	Psychomotor retardation, epilepsy, and craniofacial dysmorphism, 614501	608241	100	100	100	100
SNRNP200	Retinitis pigmentosa 33, 610359	601664	102	100	100	98
SNRPE	Hypotrichosis 11, 615059	128260	37	100	97	69
SNRPN	Prader-Willi syndrome, 176270	182279	96	100	100	100
SNTA1	Long QT syndrome 12, 612955	601017	111	100	94	82
SNX10	Osteopetrosis 8, 615085	614780	77	100	100	98
SOBP	Mental retardation, anterior maxillary protrusion, and strabismus, 613671	613667	165	100	98	95
SOD1	Amyotrophic lateral sclerosis 1, 105400	147450	74	100	100	100
SOS1	?Fibromatosis, gingival, 1, 135300 Noonan syndrome 4, 610733	182530	71	100	100	95
SOST	Craniodiaphyseal dysplasia, 122860 Sclerosteosis 1, 269500 Van Buchem disease, 239100	605740	163	100	100	100
SOX10	PCWH syndrome, 609136 Waardenburg syndrome, type 2E, with or without neurologic involvement, 611584 Waardenburg syndrome, type 4C, 613266	602229	72	100	97	89

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SOX17	Vesicoureteral reflux 3, 613674	610928	157	100	100	100
SOX18	Hypotrichosis-lymphedema-telangiectasia syndrome, 607823 Hypotrichosis-lymphedema-telangiectasia-renal defect syndrome, 137940	601618	73	95	90	81
SOX2	Microphthalmia, syndromic 3, 206900 Optic nerve hypoplasia and abnormalities of the central nervous system, 206900	184429	183	100	100	100
SOX3	Mental retardation, with isolated growth hormone deficiency, 300123 Panhypopituitarism, 312000	313430	64	100	95	89
SOX9	Acampomelic campomelic dysplasia, 114290 Campomelic dysplasia, 114290 Campomelic dysplasia with autosomal sex reversal, 114290	608160	157	100	100	100
SP110	Hepatic venoocclusive disease with immunodeficiency, 235550 {Mycobacterium tuberculosis, susceptibility to}, 607948	604457	61	100	100	96
SP7	Osteogenesis imperfecta, type XII, 613849	606633	129	100	100	100
SPAG1	Ciliary dyskinesia, primary, 28, 615505	603395	64	100	99	91
SPART	Troyer syndrome, 275900	607111	71	100	100	98
SPAST	Spastic paraplegia 4, 182601	604277	64	100	100	94
SPATA16	?Spermatogenic failure 6, 102530	609856	64	100	100	98
SPATA7	Leber congenital amaurosis 3, 604232 Retinitis pigmentosa, juvenile, 604232	609868	62	100	99	94
SPECC1L	?Facial clefting, oblique, 1, 600251 Hypertelorism, Teebi type, 145420 Opitz GBBB syndrome, type II, 145410	614140	64	100	99	92
SPG11	Amyotrophic lateral sclerosis 5, juvenile, 602099 Charcot-Marie-Tooth disease, axonal, type 2X, 616668 Spastic paraplegia 11, 604360	610844	84	100	100	97
SPG21	Mast syndrome, 248900	608181	66	100	95	88
SPG7	Spastic paraplegia 7, 607259	602783	125	100	100	97
SPINK1	{Fibrocalculus pancreatic diabetes, susceptibility to}, 608189 Pancreatitis, hereditary, 167800 Tropical calcific pancreatitis, 608189	167790	79	100	100	97
SPINK5	Netherton syndrome, 256500	605010	66	100	99	95
SPINT2	Diarrhea 3, secretory sodium, congenital, syndromic, 270420	605124	94	100	100	100
SPR	Dystonia, dopa-responsive, due to sepiapterin reductase deficiency, 612716	182125	102	100	100	100
SPRED1	Legius syndrome, 611431	609291	57	100	98	92
SPRY4	Hypogonadotropic hypogonadism 17 with or without anosmia, 615266	607984	104	100	100	96

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SPTA1	Elliptocytosis-2, 130600 Pyropoikilocytosis, 266140 Spherocytosis, type 3, 270970	182860	68	100	100	96
SPTAN1	Epileptic encephalopathy, early infantile, 5, 613477	182810	95	100	100	98
SPTB	Anemia, neonatal hemolytic, fatal or near-fatal, 617948 Elliptocytosis-3, 617948 Spherocytosis, type 2, 616649	182870	124	100	100	100
SPTBN2	Spinocerebellar ataxia 5, 600224 Spinocerebellar ataxia 14, 615386	604985	121	100	100	100
SPTLC1	Neuropathy, hereditary sensory and autonomic, type IA, 162400	605712	62	100	99	94
SPTLC2	Neuropathy, hereditary sensory and autonomic, type IC, 613640	605713	81	100	100	96
SQSTM1	Frontotemporal dementia and/or amyotrophic lateral sclerosis 3, 616437 Myopathy, distal, with rimmed vacuoles, 617158 Neurodegeneration with ataxia, dystonia, and gaze palsy, childhood-onset, 617145 Paget disease of bone 3, 167250	601530	129	100	100	96
SRC	Colon cancer, advanced, somatic, 114500 ?Thrombocytopenia 6, 616937	190090	119	100	100	100
SRCAP	Floating-Harbor syndrome, 136140	611421	139	100	100	100
SRD5A2	Pseudovaginal perineoscrotal hypospadias, 264600	607306	56	100	99	86
SRD5A3	Congenital disorder of glycosylation, type Ig, 612379 Kahrizi syndrome, 612713	611715	111	100	100	96
SRP72	Bone marrow failure syndrome 1, 614675	602122	56	100	99	89
SRPX2	?Rolandic epilepsy, mental retardation, and speech dyspraxia, 300643	300642	59	100	99	95
SRY	46XX sex reversal 1, 400045 dominant 46XY sex reversal 1, 400044	480000	159	100	100	100
SSTR5	Somatostatin analog, resistance to	182455	168	100	100	100
ST14	Ichthyosis, congenital 11, 602400	606797	111	100	100	99
ST3GAL3	?Epileptic encephalopathy, early infantile, 15, 615006 Mental retardation 12, 611090	606494	79	100	100	96
ST3GAL5	Salt and pepper developmental regression syndrome, 609056	604402	55	99	92	84
STAC3	Myopathy, congenital, Baily-Bloch, 255995	615521	82	100	100	98
STAMBP	Microcephaly-capillary malformation syndrome, 614261	606247	72	100	100	97
STAR	Lipoid adrenal hyperplasia, 201710	600617	118	100	100	100

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STAT1	Immunodeficiency 31A, mycobacteriosis, 614892 Immunodeficiency 31B, mycobacterial and viral infections, 613796 Immunodeficiency 31C, 614162	600555	64	100	99	94
STAT3	Autoimmune disease, multisystem, infantile-onset, 1, 615952 Hyper-IgE recurrent infection syndrome, 147060	102582	90	100	100	99
STAT5B	Growth hormone insensitivity with immunodeficiency, 245590 Leukemia, acute promyelocytic, somatic, 102578	604260	107	100	100	97
STIL	Microcephaly 7, primary, 612703	181590	65	100	100	97
STIM1	Immunodeficiency 10, 612783 Myopathy, tubular aggregate, 1, 160565 Stormorken syndrome, 185070	605921	95	100	100	99
STK11	Melanoma, malignant, somatic Pancreatic cancer, somatic, 260350 Peutz-Jeghers syndrome, 175200 Testicular tumor, somatic, 273300	602216	161	100	100	100
STK4	T-cell immunodeficiency, recurrent infections, autoimmunity, and cardiac malformations, 614868	604965	82	100	100	99
STOX1	Preeclampsia/eclampsia 4, 609404	609397	62	89	89	89
STRA6	Microphthalmia, isolated, with coloboma 8, 601186 Microphthalmia, syndromic 9, 601186	610745	98	100	100	100
STRADA	Polyhydramnios, megalencephaly, and symptomatic epilepsy, 611087	608626	96	100	100	98
STRC	Deafness 16, 603720	606440	66	53	48	46
STS	Ichthyosis, 308100	300747	90	97	97	95
STXBP1	Epileptic encephalopathy, early infantile, 4, 612164	602926	76	100	100	99
STXBP2	Hemophagocytic lymphohistiocytosis, familial, 5, 613101	601717	121	100	100	100
SUCLA2	Mitochondrial DNA depletion syndrome 5 (encephalomyopathic with or without methylmalonic aciduria), 612073	603921	56	100	97	86
SUCLG1	Mitochondrial DNA depletion syndrome 9 (encephalomyopathic type with methylmalonic aciduria), 245400	611224	88	100	100	100
SUFU	Basal cell nevus syndrome, 109400 Joubert syndrome 32, 617757 Medulloblastoma, desmoplastic, 155255 {Meningioma, familial, susceptibility to}, 607174	607035	104	100	100	100
SUGCT	Glutaric aciduria III, 231690	609187	73	100	100	93
SUMF1	Multiple sulfatase deficiency, 272200	607939	88	100	100	99
SUMO1	?Orofacial cleft 10, 613705	601912	72	100	97	85
SUOX	Sulfite oxidase deficiency, 272300	606887	136	100	100	100

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SURF1	Charcot-Marie-Tooth disease, type 4K, 616684 Leigh syndrome, due to COX IV deficiency, 256000	185620	88	94	89	87
SYCP3	Pregnancy loss, recurrent, 4, 270960 Spermatogenic failure 4, 270960	604759	45	100	100	89
SYN1	Epilepsy, with variable learning disabilities and behavior disorders, 300491 dominant	313440	60	100	99	85
SYNE1	Arthrogryposis multiplex congenita, myogenic type, 618484 Emery-Dreifuss muscular dystrophy 4, 612998 Spinocerebellar ataxia 8, 610743	608441	77	100	100	97
SYNE2	Emery-Dreifuss muscular dystrophy 5, 612999	608442	77	100	100	96
SYNE4	Deafness 76, 615540	615535	107	100	100	100
SYNGAP1	Mental retardation 5, 612621	603384	158	98	98	98
SYNJ1	Epileptic encephalopathy, early infantile, 53, 617389 Parkinson disease 20, early-onset, 615530	604297	62	100	99	92
SYP	Mental retardation 96, 300802	313475	69	100	100	98
SYT14	?Spinocerebellar ataxia 11, 614229	610949	61	100	95	86
SZT2	Epileptic encephalopathy, early infantile, 18, 615476	615463	117	100	100	100
TAB2	Congenital heart defects, nonsyndromic, 2, 614980	605101	76	100	100	99
TAC3	Hypogonadotropic hypogonadism 10 with or without anosmia, 614839	162330	93	100	100	95
TACR3	Hypogonadotropic hypogonadism 11 with or without anosmia, 614840	162332	136	100	100	100
TACSTD2	Corneal dystrophy, gelatinous drop-like, 204870	137290	228	100	100	100
TAF1	Dystonia-Parkinsonism, 314250 Mental retardation, syndromic 33, 300966	313650	54	100	97	87
TAF2	Mental retardation 40, 615599	604912	64	100	99	92
TAL1	Leukemia, T-cell acute lymphocytic, somatic, 613065	187040	64	97	89	82
TAL2	Leukemia, T-cell acute lymphocytic, somatic, 613065	186855	87	100	100	100
TALDO1	Transaldolase deficiency, 606003	602063	136	100	100	100
TAP1	Bare lymphocyte syndrome, type I, 604571	170260	125	100	100	100
TAP2	Bare lymphocyte syndrome, type I, due to TAP2 deficiency, 604571	170261	106	100	100	100
TAPBP	Bare lymphocyte syndrome, type I, 604571	601962	106	100	100	100
TARDBP	Amyotrophic lateral sclerosis 10, with or without FTD, 612069 Frontotemporal lobar degeneration, TARDBP-related, 612069	605078	124	100	100	95
TAT	Tyrosinemia, type II, 276600	613018	65	100	100	98
TAZ	Barth syndrome, 302060	300394	97	100	98	90
TBC1D20	Warburg micro syndrome 4, 615663	611663	74	100	93	93

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TBC1D24	DOORS syndrome, 220500 Deafness 86, 614617 Deafness 65, 616044 Epilepsy, rolandic, with proxysmal exercise-induce dystonia and writer's cramp, 608105 Epileptic encephalopathy, early infantile, 16, 615338 Myoclonic epilepsy, infantile, familial, 605021	613577	158	100	100	100
TBCE	Encephalopathy, progressive, with amyotrophy and optic atrophy, 617207 Hypoparathyroidism-retardation-dysmorphism syndrome, 241410 Kenny-Caffey syndrome, type 1, 244460	604934	59	100	97	88
TBP	{Parkinson disease, susceptibility to}, 168600 Spinocerebellar ataxia 17, 607136	600075	80	100	100	96
TBX1	Conotruncal anomaly face syndrome, 217095 DiGeorge syndrome, 188400 Tetralogy of Fallot, 187500 Velocardiofacial syndrome, 192430	602054	94	94	85	80
TBX15	Cousin syndrome, 260660	604127	77	100	100	99
TBX19	Adrenocorticotrophic hormone deficiency, 201400	604614	126	100	100	99
TBX20	Atrial septal defect 4, 611363	606061	80	100	100	97
TBX21	Asthma and nasal polyps, 208550 {Asthma, aspirin-induced, susceptibility to}, 208550	604895	139	100	100	99
TBX22	?Abruzzo-Erickson syndrome, 302905 Cleft palate with ankyloglossia, 303400	300307	60	100	99	90
TBX3	Ulnar-mammary syndrome, 181450	601621	100	100	100	99
TBX4	Ischiocoxopodopatellar syndrome with or without pulmonary arterial hypertension, 147891	601719	145	100	100	100
TBX5	Holt-Oram syndrome, 142900	601620	77	100	100	99
TBXAS1	Ghosal hematodiaphyseal syndrome, 231095 ?Thromboxane synthase deficiency, 614158	274180	74	100	99	93
TCAP	Cardiomyopathy, hypertrophic, 25, 607487 Muscular dystrophy, limb-girdle 7, 601954	604488	102	100	100	100
TCF12	Craniosynostosis 3, 615314	600480	72	100	100	98
TCF4	Corneal dystrophy, Fuchs endothelial, 3, 613267 Pitt-Hopkins syndrome, 610954	602272	73	100	100	96
TCIRG1	Osteopetrosis 1, 259700	604592	117	100	100	97
TCN2	Transcobalamin II deficiency, 275350	613441	118	100	100	100
TCOF1	Treacher Collins syndrome 1, 154500	606847	115	100	100	99

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TCTN1	Joubert syndrome 13, 614173	609863	94	100	100	98
TCTN2	Joubert syndrome 24, 616654 ?Meckel syndrome 8, 613885	613846	84	100	100	97
TCTN3	Joubert syndrome 18, 614815 Orofaciodigital syndrome IV, 258860	613847	66	100	100	97
TDGF1	Forebrain defects	187395	127	100	100	100
TDP1	?Spinocerebellar ataxia, with axonal neuropathy 1, 607250	607198	53	100	100	92
TDRD7	Cataract 36, 613887	611258	70	100	100	98
TEAD1	Sveinsson chorioretinal atrophy, 108985	189967	82	100	100	98
TECPR2	Spastic paraplegia 49, 615031	615000	102	100	100	99
TECR	Mental retardation 14, 614020	610057	140	100	100	100
TECTA	Deafness 8/12, 601543 Deafness 21, 603629	602574	123	100	100	98
TEK	Glaucoma 3, primary congenital, E, 617272 Venous malformations, multiple cutaneous and mucosal, 600195	600221	80	100	100	99
TENM3	?Microphthalmia, isolated, with coloboma 9, 615145 Microphthalmia, syndromic 15, 615145	610083	86	100	100	99
TERC	{Aplastic anemia}, 614743 Dyskeratosis congenita 1, 127550 {Pulmonary fibrosis, idiopathic, susceptibility to}, 614743	602322	No coverage data			
TET2	Myelodysplastic syndrome, somatic, 614286	612839	71	100	100	99
TEX28	No OMIM phenotype	300092	No coverage data			
TF	Atransferrinemia, 209300	190000	79	100	100	99
TFAP2A	Branchiooculofacial syndrome, 113620	107580	95	100	100	100
TFAP2B	Char syndrome, 169100 Patent ductus arteriosus 2, 617035	601601	142	100	100	100
TFE3	Renal cell carcinoma, papillary, 1, 300854	314310	71	100	99	90
TFG	Hereditary motor and sensory neuropathy, Okinawa type, 604484 ?Spastic paraplegia 57, 615658	602498	87	100	100	94
TFR2	Hemochromatosis, type 3, 604250	604720	101	100	100	97
TG	{Autoimmune thyroid disease, susceptibility to, 3}, 608175 Thyroid dyshormonogenesis 3, 274700	188450	97	100	100	99
TGFB1	Camurati-Engelmann disease, 131300 {Cystic fibrosis lung disease, modifier of}, 219700 Inflammatory bowel disease, immunodeficiency, and encephalopathy, 618213	190180	111	100	99	95

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
TGFB2	Loeys-Dietz syndrome 4, 614816	190220	88	100	100	98
TGFB3	Arrhythmogenic right ventricular dysplasia 1, 107970 Loeys-Dietz syndrome 5, 615582	190230	104	100	100	100
TGFBI	Corneal dystrophy, Avellino type, 607541 Corneal dystrophy, Groenouw type I, 121900 Corneal dystrophy, Reis-Bucklers type, 608470 Corneal dystrophy, Thiel-Behnke type, 602082 Corneal dystrophy, epithelial basement membrane, 121820 Corneal dystrophy, lattice type I, 122200 Corneal dystrophy, lattice type IIIA, 608471	601692	82	100	100	98
TGFBR1	Loeys-Dietz syndrome 1, 609192 {Multiple self-healing squamous epithelioma, susceptibility to}, 132800	190181	171	94	93	93
TGFBR2	Colorectal cancer, hereditary nonpolyposis, type 6, 614331 Esophageal cancer, somatic, 133239 Loeys-Dietz syndrome 2, 610168	190182	210	100	100	100
TGIF1	Holoprosencephaly 4, 142946	602630	152	100	100	100
TGM1	Ichthyosis, congenital 1, 242300	190195	125	100	100	100
TGM5	Peeling skin syndrome 2, 609796	603805	113	100	100	100
TGM6	Spinocerebellar ataxia 35, 613908	613900	117	100	100	99
TH	Segawa syndrome, recessive, 605407	191290	93	100	99	94
THAP1	Dystonia 6, torsion, 602629	609520	87	100	100	98
THBD	{Hemolytic uremic syndrome, atypical, susceptibility to}, 612926 Thrombophilia due to thrombomodulin defect, 614486	188040	195	100	100	100
THOC6	Beaulieu-Boycott-Innes syndrome, 613680	615403	210	100	100	100
THPO	Thrombocythemia 1, 187950	600044	118	100	100	100
THRA	Hypothyroidism, congenital, nongoitrous, 6, 614450	190120	121	100	100	100
THRΒ	Thyroid hormone resistance, 188570 Thyroid hormone resistance, 274300 Thyroid hormone resistance, selective pituitary, 145650	190160	88	100	100	99
TIA1	Welander distal myopathy, 604454	603518	73	100	100	96
TIMM8A	Mohr-Tranebjærg syndrome, 304700	300356	138	100	100	100
TIMP3	Sorsby fundus dystrophy, 136900	188826	104	100	100	100
TINF2	Dyskeratosis congenita 3, 613990 Revesz syndrome, 268130	604319	155	100	100	100

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TJP2	Cholestasis, progressive familial intrahepatic 4, 615878 Hypercholanemia, familial, 607748	607709	83	100	100	99
TK2	Mitochondrial DNA depletion syndrome 2 (myopathic type), 609560 ?Progressive external ophthalmoplegia with mitochondrial DNA deletions 3, 617069	188250	66	100	100	95
TLL1	Atrial septal defect 6, 613087	606742	67	100	100	96
TLR4	No OMIM phenotype	603030	61	100	100	99
TMC1	Deafness 36, 606705 Deafness 7, 600974	606706	61	100	100	94
TMC6	Epidermolytic hyperkeratosis, 226400	605828	86	100	100	97
TMC8	Epidermolytic hyperkeratosis 2, 618231	605829	118	100	100	100
TMCO1	Craniofacial dysmorphism, skeletal anomalies, and mental retardation syndrome, 213980	614123	108	100	100	96
TMEM126A	Optic atrophy 7, 612989	612988	68	100	100	93
TMEM138	Joubert syndrome 16, 614465	614459	54	100	100	100
TMEM165	Congenital disorder of glycosylation, type IIk, 614727	614726	108	100	100	100
TMEM181	No OMIM phenotype	613209	83	100	100	97
TMEM216	Joubert syndrome 2, 608091 Meckel syndrome 2, 603194	613277	87	100	100	95
TMEM231	Joubert syndrome 20, 614970 Meckel syndrome 11, 615397	614949	96	100	95	90
TMEM237	Joubert syndrome 14, 614424	614423	57	100	98	87
TMEM38B	Osteogenesis imperfecta, type XIV, 615066	611236	68	100	100	98
TMEM67	{Bardet-Biedl syndrome 14, modifier of}, 615991 COACH syndrome, 216360 Joubert syndrome 6, 610688 Meckel syndrome 3, 607361 Nephronophthisis 11, 613550 ?RHYSN syndrome, 602152	609884	77	100	100	94
TMEM70	Mitochondrial complex V (ATP synthase) deficiency, nuclear type 2, 614052	612418	93	100	100	95
TMIE	Deafness 6, 600971	607237	85	100	100	99
TMLHE	{Autism, susceptibility to 6}, 300872	300777	32	89	81	67
TMPRSS15	Enterokinase deficiency, 226200	606635	68	100	98	91
TMPRSS3	Deafness 8/10, 601072	605511	74	100	100	99
TMPRSS6	Iron-refractory iron deficiency anemia, 206200	609862	118	100	100	100
TNC	Deafness 56, 615629	187380	110	100	100	98
TNFRSF10B	Squamous cell carcinoma, head and neck, 275355	603612	95	100	100	99

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TNFRSF11A	Osteolysis, familial expansile, 174810 Osteopetrosis 7, 612301 {Paget disease of bone 2, early-onset}, 602080	603499	97	95	95	95
TNFRSF11B	Paget disease of bone 5, juvenile-onset, 239000	602643	91	100	100	99
TNFRSF13B	Immunodeficiency, common variable, 2, 240500 Immunoglobulin A deficiency 2, 609529	604907	106	100	100	100
TNFRSF13C	Immunodeficiency, common variable, 4, 613494	606269	78	100	91	73
TNFRSF1A	{Multiple sclerosis, susceptibility to, 5}, 614810 Periodic fever, familial, 142680	191190	97	100	98	94
TNFSF11	Osteopetrosis 2, 259710	602642	56	100	100	94
TNNC1	Cardiomyopathy, dilated, 1Z, 611879 Cardiomyopathy, hypertrophic, 13, 613243	191040	127	100	100	100
TNNI2	Arthrogryposis, distal, type 2B1, 601680	191043	151	100	100	100
TNNI3	Cardiomyopathy, dilated, 1FF, 613286 ?Cardiomyopathy, dilated, 2A, 611880 Cardiomyopathy, familial restrictive, 1, 115210 Cardiomyopathy, hypertrophic, 7, 613690	191044	165	100	100	97
TNNT1	Nemaline myopathy 5, Amish type, 605355	191041	88	100	99	94
TNNT2	Cardiomyopathy, dilated, 1D, 601494 Cardiomyopathy, familial restrictive, 3, 612422 Cardiomyopathy, hypertrophic, 2, 115195 Left ventricular noncompaction 6, 601494	191045	149	100	100	100
TNNT3	Arthrogryposis, distal, type 2B2, 618435	600692	114	100	100	100
TNXB	Ehlers-Danlos syndrome, classic-like, 1, 606408 Vesicoureteral reflux 8, 615963	600985	151	94	92	91
TOP1	DNA topoisomerase I, camptothecin-resistant	126420	76	100	100	93
TOP2A	DNA topoisomerase II, resistance to inhibition of, by amsacrine	126430	71	100	100	97
TOPORS	Retinitis pigmentosa 31, 609923	609507	73	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
TP53	{Adrenocortical carcinoma, pediatric}, 202300 {Basal cell carcinoma 7}, 614740 Bone marrow failure syndrome 5, 618165 Breast cancer, somatic, 114480 {Choroid plexus papilloma}, 260500 {Colorectal cancer}, 114500 {Glioma susceptibility 1}, 137800 Hepatocellular carcinoma, somatic, 114550 Li-Fraumeni syndrome, 151623 Nasopharyngeal carcinoma, somatic, 607107 {Osteosarcoma}, 259500 Pancreatic cancer, somatic, 260350	191170	167	100	100	100
TP63	ADULT syndrome, 103285 Ectrodactyly, ectodermal dysplasia, and cleft lip/palate syndrome 3, 604292 Hay-Wells syndrome, 106260 Limb-mammary syndrome, 603543 Orofacial cleft 8, 618149 Rapp-Hodgkin syndrome, 129400 Split-hand/foot malformation 4, 605289	603273	115	100	100	98
TPI1	Hemolytic anemia due to triosephosphate isomerase deficiency, 615512	190450	123	100	98	96
TPK1	Thiamine metabolism dysfunction syndrome 5 (episodic encephalopathy type), 614458	606370	59	100	100	89
TPM1	Cardiomyopathy, dilated, 1Y, 611878 Cardiomyopathy, hypertrophic, 3, 115196 Left ventricular noncompaction 9, 611878	191010	145	100	100	100
TPM2	Arthrogryposis, distal, type 1A, 108120 Arthrogryposis, distal, type 2B4, 108120 CAP myopathy 2, 609285 Nemaline myopathy 4, 609285	190990	98	100	100	100
TPM3	CAP myopathy 1, 609284 Myopathy, congenital, with fiber-type disproportion, 255310 Nemaline myopathy 1 or recessive, 609284	191030	69	100	100	99
TPMT	{Thiopurines, poor metabolism of, 1}, 610460	187680	75	100	100	89
TPO	Thyroid dyshormonogenesis 2A, 274500	606765	116	100	100	100
TPP1	Ceroid lipofuscinoses, neuronal, 2, 204500 Spinocerebellar ataxia 7, 609270	607998	97	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
TPRN	Deafness 79, 613307	613354	94	92	86	79
TRAPPC11	Muscular dystrophy, limb-girdle 18, 615356	614138	60	100	97	90
TRAPPC2	Spondyloepiphyseal dysplasia tarda, 313400	300202	52	100	91	70
TRAPPC9	Mental retardation 13, 613192	611966	100	100	100	98
TRDN	Ventricular tachycardia, catecholaminergic polymorphic, 5, with or without muscle weakness, 615441	603283	53	100	94	79
TREM2	Polycystic lipomembranous osteodysplasia with sclerosing leukoencephalopathy 2, 618193	605086	107	100	100	100
TREX1	Aicardi-Goutieres syndrome 1, dominant and recessive, 225750 Chilblain lupus, 610448 {Systemic lupus erythematosus, susceptibility to}, 152700 Vasculopathy, retinal, with cerebral leukodystrophy, 192315	606609	234	100	100	100
TRHR	Thyrotropin-releasing hormone resistance, generalized	188545	94	100	100	100
TRIM24	No OMIM phenotype	603406	72	100	100	97
TRIM32	?Bardet-Biedl syndrome 11, 615988 Muscular dystrophy, limb-girdle 8, 254110	602290	107	100	100	100
TRIM33	No OMIM phenotype	605769	66	100	99	92
TRIM37	Mulibrey nanism, 253250	605073	61	100	100	94
TRIOBP	Deafness 28, 609823	609761	182	100	99	97
TRIP11	Achondrogenesis, type IA, 200600 Osteochondrodysplasia, 184260	604505	66	100	98	90
TRMU	{Deafness, modifier of}, 580000 Liver failure, transient infantile, 613070	610230	97	100	100	98
TRPA1	?Episodic pain syndrome, familial, 1, 615040	604775	61	100	98	88
TRPC6	Glomerulosclerosis, focal segmental, 2, 603965	603652	86	100	100	96
TRPM1	Night blindness, congenital stationary (complete), 1C, 613216	603576	85	100	100	99
TRPM4	Erythrokeratoderma variabilis et progressiva 6, 618531 Progressive familial heart block, type IB, 604559	606936	124	100	100	99
TRPM6	Hypomagnesemia 1, intestinal, 602014	607009	75	100	100	96
TRPS1	Trichorhinophalangeal syndrome, type I, 190350 Trichorhinophalangeal syndrome, type III, 190351	604386	85	100	100	100
TRPV3	Olmsted syndrome, 614594 ?Palmoplantar keratoderma, nonepidermolytic, focal 2, 616400	607066	106	100	100	99

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
TRPV4	?Avascular necrosis of femoral head, primary, 2, 617383 Brachyolmia type 3, 113500 Digital arthropathy-brachydactyly, familial, 606835 Hereditary motor and sensory neuropathy, type IIc, 606071 Metatropic dysplasia, 156530 Parastremmatic dwarfism, 168400 SED, Maroteaux type, 184095 Scapuloperoneal spinal muscular atrophy, 181405 [Sodium serum level QTL 1], 613508 Spinal muscular atrophy, distal, congenital nonprogressive, 600175 Spondylometaphyseal dysplasia, Kozlowski type, 184252	605427	109	100	100	100
TSC1	Focal cortical dysplasia, type II, somatic, 607341 Lymphangiomyomatosis, 606690 Tuberous sclerosis-1, 191100	605284	167	100	100	100
TSC2	?Focal cortical dysplasia, type II, somatic, 607341 Lymphangiomyomatosis, somatic, 606690 Tuberous sclerosis-2, 613254	191092	184	100	100	100
TSEN2	Pontocerebellar hypoplasia type 2B, 612389	608753	63	87	86	78
TSEN34	?Pontocerebellar hypoplasia type 2C, 612390	608754	87	100	100	100
TSEN54	Pontocerebellar hypoplasia type 2A, 277470 Pontocerebellar hypoplasia type 4, 225753 ?Pontocerebellar hypoplasia type 5, 610204	608755	105	100	96	96
TSFM	Combined oxidative phosphorylation deficiency 3, 610505	604723	77	100	100	100
TSG101	No OMIM phenotype	601387	78	100	100	93
TSHB	Hypothyroidism, congenital, nongoitrous 4, 275100	188540	95	100	100	100
TSHR	Hyperthyroidism, familial gestational, 603373 Hyperthyroidism, nonautoimmune, 609152 Hypothyroidism, congenital, nongoitrous, 1, 275200 Thyroid adenoma, hyperfunctioning, somatic Thyroid carcinoma with thyrotoxicosis	603372	100	100	100	99
TSHZ1	Aural atresia, congenital, 607842	614427	142	98	98	98
TSPAN12	Exudative vitreoretinopathy 5, 613310	613138	67	100	100	92
TSPAN7	Mental retardation 58, 300210	300096	55	100	98	83
TSPEAR	?Deafness 98, 614861 Ectodermal dysplasia 14, hair/tooth type with or without hypohidrosis, 618180	612920	105	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
TSPYL1	Sudden infant death with dysgenesis of the testes syndrome, 608800	604714	133	100	100	100
TTBK2	Spinocerebellar ataxia 11, 604432	611695	74	100	100	99
TTC19	Mitochondrial complex III deficiency, nuclear type 2, 615157	613814	53	100	92	77
TTC21B	Nephronophthisis 12, 613820	612014	75	100	99	94
	Short-rib thoracic dysplasia 4 with or without polydactyly, 613819					
TTC37	Trichohepatoenteric syndrome 1, 222470	614589	58	100	99	93
TTC7A	Gastrointestinal defects and immunodeficiency syndrome, 243150	609332	110	100	100	99
TTC8	Bardet-Biedl syndrome 8, 615985	608132	66	100	100	93
	?Retinitis pigmentosa 51, 613464					
TTI2	Mental retardation 39, 615541	614426	66	100	100	98
TTN	Cardiomyopathy, dilated, 1G, 604145	188840	68	100	99	97
	Cardiomyopathy, familial hypertrophic, 9, 613765					
	Muscular dystrophy, limb-girdle 10, 608807					
	Myopathy, myofibrillar, 9, with early respiratory failure, 603689					
	Salih myopathy, 611705					
	Tibial muscular dystrophy, tardive, 600334					
TTPA	Ataxia with isolated vitamin E deficiency, 277460	600415	61	100	98	91
TTR	Amyloidosis, hereditary, transthyretin-related, 105210	176300	93	100	100	100
	Carpal tunnel syndrome, familial, 115430 [Dystransthyretinemic hyperthyroxinemia], 145680					
TUBA1A	Lissencephaly 3, 611603	602529	110	100	100	100
TUBA8	Cortical dysplasia, complex, with other brain malformations 8, 613180	605742	118	100	100	100
TUBB1	Macrothrombocytopenia, TUBB1-related, 613112	612901	153	100	100	99
TUBB2A	Cortical dysplasia, complex, with other brain malformations 5, 615763	615101	99	99	82	74
TUBB2B	Cortical dysplasia, complex, with other brain malformations 7, 610031	612850	119	100	87	78
TUBB3	Cortical dysplasia, complex, with other brain malformations 1, 614039	602661	269	100	99	95
	Fibrosis of extraocular muscles, congenital, 3A, 600638					
TUBB4A	Dystonia 4, torsion, 128101	602662	242	100	100	99
	Leukodystrophy, hypomyelinating, 6, 612438					
TUBG1	Cortical dysplasia, complex, with other brain malformations 4, 615412	191135	190	100	100	100
TUBGCP6	Microcephaly and chorioretinopathy, 1, 251270	610053	159	100	100	99
TUFM	Combined oxidative phosphorylation deficiency 4, 610678	602389	158	100	100	100
TULP1	Leber congenital amaurosis 15, 613843	602280	108	100	100	100
	Retinitis pigmentosa 14, 600132					
TUSC3	Mental retardation 7, 611093	601385	72	100	100	95

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TWIST1	Craniosynostosis 1, 123100 Robinow-Sorauf syndrome, 180750 Saethre-Chotzen syndrome with or without eyelid anomalies, 101400 Sweeney-Cox syndrome, 617746	601622	132	100	100	95
TWIST2	Ablepharon-macrostomia syndrome, 200110 Barber-Say syndrome, 209885 Focal facial dermal dysplasia 3, Setleis type, 227260	607556	150	100	100	100
TWNK	Mitochondrial DNA depletion syndrome 7 (hepatocerebral type), 271245 Perrault syndrome 5, 616138 Progressive external ophthalmoplegia with mitochondrial DNA deletions 3, 609286	606075	145	100	100	100
TYK2	Immunodeficiency 35, 611521	176941	128	100	100	99
TYMP	Mitochondrial DNA depletion syndrome 1 (MNGIE type), 603041	131222	102	100	100	100
TYR	Albinism, oculocutaneous, type IA, 203100 Albinism, oculocutaneous, type IB, 606952 {Melanoma, cutaneous malignant, susceptibility to, 8}, 601800 [Skin/hair/eye pigmentation 3, blue/green eyes], 601800 [Skin/hair/eye pigmentation 3, light/dark/freckling skin], 601800 Waardenburg syndrome/albinism, digenic, 103470	606933	83	100	100	98
TYROBP	Polycystic lipomembranous osteodysplasia with sclerosing leukoencephalopathy 1, 221770	604142	108	100	100	100
TYRP1	Albinism, oculocutaneous, type III, 203290 [Skin/hair/eye pigmentation, variation in, 11 (Melanesian blond hair)], 612271	115501	73	100	100	98
UBA1	Spinal muscular atrophy 2, infantile, 301830	314370	94	100	100	100
UBE2A	Mental retardation syndromic, Nascimento-type, 300860	312180	57	100	99	81
UBE3A	Angelman syndrome, 105830	601623	61	100	100	96
UBE3B	Kaufman oculocerebrofacial syndrome, 244450	608047	116	100	100	99
UBIAD1	Corneal dystrophy, Schnyder type, 121800	611632	173	100	100	100
UBQLN2	Amyotrophic lateral sclerosis 15, with or without frontotemporal dementia, 300857	300264	79	100	100	100
UBR1	Johanson-Blizzard syndrome, 243800	605981	61	100	99	93
UGT1A1	[Bilirubin, serum level of, QTL1], 601816 Crigler-Najjar syndrome, type I, 218800 Crigler-Najjar syndrome, type II, 606785 [Gilbert syndrome], 143500 Hyperbilirubinemia, familial transient neonatal, 237900	191740	130	100	100	100

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
UMOD	Glomerulocystic kidney disease with hyperuricemia and isosthenuria, 609886 Hyperuricemic nephropathy, familial juvenile 1, 162000 Medullary cystic kidney disease 2, 603860	191845	95	100	100	97
UMPS	Orotic aciduria, 258900	613891	74	100	100	98
UNG	Immunodeficiency with hyper IgM, type 5, 608106	191525	104	100	100	100
UPB1	Beta-ureidopropionase deficiency, 613161	606673	96	100	100	100
UPF3B	Mental retardation, syndromic 14, 300676	300298	74	100	98	87
UQCRB	Mitochondrial complex III deficiency, nuclear type 3, 615158	191330	75	100	100	100
UQCRC2	Mitochondrial complex III deficiency, nuclear type 5, 615160	191329	95	100	100	96
UQCRCQ	Mitochondrial complex III deficiency, nuclear type 4, 615159	612080	123	100	100	100
UROC1	?Urocanase deficiency, 276880	613012	112	100	100	99
UROD	Porphyria cutanea tarda, 176100 Porphyria, hepatoerythropoietic, 176100	613521	87	100	100	97
UROS	Porphyria, congenital erythropoietic, 263700	606938	60	100	100	97
USB1	Poikiloderma with neutropenia, 604173	613276	93	100	100	99
USH1C	Deafness 18A, 602092 Usher syndrome, type 1C, 276904	605242	80	100	98	92
USH1G	Usher syndrome, type 1G, 606943	607696	180	100	100	100
USH2A	Retinitis pigmentosa 39, 613809 Usher syndrome, type 2A, 276901	608400	82	100	100	98
USP21	No OMIM phenotype	604729	150	100	100	99
USP9Y	Spermatogenic failure, 2, 415000	400005	76	100	98	93
UTP4	No OMIM phenotype	607456	85	100	100	97
UVSSA	UV-sensitive syndrome 3, 614640	614632	123	100	100	99
VANGL1	Caudal regression syndrome, 600145 {Neural tube defects, susceptibility to}, 182940	610132	99	100	100	100
VANGL2	Neural tube defects, 182940	600533	131	100	100	100
VAPB	Amyotrophic lateral sclerosis 8, 608627 Spinal muscular atrophy, late-onset, Finkel type, 182980	605704	71	100	100	98
VAX1	?Microphthalmia, syndromic 11, 614402	604294	104	97	91	87
VCAN	Wagner syndrome 1, 143200	118661	67	100	100	99
VCL	Cardiomyopathy, dilated, 1W, 611407 Cardiomyopathy, hypertrophic, 15, 613255	193065	89	100	99	93

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
VCP	Amyotrophic lateral sclerosis 14, with or without frontotemporal dementia, 613954 Charcot-Marie-Tooth disease, type 2Y, 616687 Inclusion body myopathy with early-onset Paget disease and frontotemporal dementia 1, 167320	601023	90	100	100	98
VDR	?Osteoporosis, involutional, 166710 Rickets, vitamin D-resistant, type IIA, 277440	601769	87	100	100	100
VHL	Erythrocytosis, familial, 2, 263400 Hemangioblastoma, cerebellar, somatic Pheochromocytoma, 171300 Renal cell carcinoma, somatic, 144700 von Hippel-Lindau syndrome, 193300	608537	183	100	100	100
VIM	Cataract 30, pulverulent, 116300	193060	106	100	100	96
VIPAS39	Arthrogryposis, renal dysfunction, and cholestasis 2, 613404	613401	64	100	100	98
VKORC1	Vitamin K-dependent clotting factors, combined deficiency of, 2, 607473 Warfarin resistance, 122700	608547	113	100	100	99
VLDLR	Cerebellar hypoplasia and mental retardation with or without quadrupedal locomotion 1, 224050	192977	70	100	100	97
VPS13A	Choreoacanthocytosis, 200150	605978	72	100	99	91
VPS13B	Cohen syndrome, 216550	607817	75	100	99	96
VPS33B	Arthrogryposis, renal dysfunction, and cholestasis 1, 208085	608552	80	100	100	98
VPS35	{Parkinson disease 17}, 614203	601501	68	100	99	94
VPS37A	Spastic paraparesis 53, 614898	609927	46	100	99	86
VPS45	Neutropenia, severe congenital, 5, 615285	610035	74	100	100	98
VRK1	Pontocerebellar hypoplasia type 1A, 607596	602168	53	100	99	92
VSX1	?Craniofacial anomalies and anterior segment dysgenesis syndrome, 614195 Keratoconus 1, 148300	605020	87	100	100	99
VSX2	Microphthalmia with coloboma 3, 610092 Microphthalmia, isolated 2, 610093	142993	96	100	100	100
VWF	von Willebrand disease, type 1, 193400 von Willebrand disease, types 2A, 2B, 2M, and 2N, 613554 von Willibrand disease, type 3, 277480	613160	90	98	98	96
WAS	Neutropenia, severe congenital, 300299 Thrombocytopenia, 313900 Thrombocytopenia, intermittent, 313900 Wiskott-Aldrich syndrome, 301000	300392	66	97	84	76
WASHC5	Ritscher-Schinzel syndrome 1, 220210 Spastic paraparesis 8, 603563	610657	62	100	100	94

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
WDPCP	?Bardet-Biedl syndrome 15, 615992 ?Congenital heart defects, hamartomas of tongue, and polysyndactyly, 217085	613580	72	100	99	93
WDR11	Hypogonadotropic hypogonadism 14 with or without anosmia, 614858	606417	83	100	100	96
WDR19	?Cranoectodermal dysplasia 4, 614378 Nephronophthisis 13, 614377 Senior-Loken syndrome 8, 616307 ?Short-rib thoracic dysplasia 5 with or without polydactyly, 614376	608151	71	100	100	97
WDR34	Short-rib thoracic dysplasia 11 with or without polydactyly, 615633	613363	116	100	100	100
WDR35	Cranoectodermal dysplasia 2, 613610 Short-rib thoracic dysplasia 7 with or without polydactyly, 614091	613602	62	100	100	94
WDR36	Glaucoma 1, open angle, G, 609887	609669	74	100	100	95
WDR45	Neurodegeneration with brain iron accumulation 5, 300894	300526	92	100	100	100
WDR60	Short-rib thoracic dysplasia 8 with or without polydactyly, 615503	615462	80	100	99	94
WDR62	Microcephaly 2, primary, with or without cortical malformations, 604317	613583	139	100	100	100
WDR72	Amelogenesis imperfecta, type IIA3, 613211	613214	70	100	100	96
WDR81	Cerebellar ataxia, mental retardation, and dysequilibrium syndrome 2, 610185 Hydrocephalus, congenital, 3, with brain anomalies, 617967	614218	155	100	100	100
WFS1	?Cataract 41, 116400 Deafness 6/14/38, 600965 {Diabetes mellitus, noninsulin-dependent, association with}, 125853 Wolfram syndrome 1, 222300 Wolfram-like syndrome, 614296	606201	196	100	100	100
WHRN	Deafness 31, 607084 Usher syndrome, type 2D, 611383	607928	119	100	100	100
WIPF1	?Wiskott-Aldrich syndrome 2, 614493	602357	73	100	100	96
WNK1	Neuropathy, hereditary sensory and autonomic, type II, 201300 Pseudohypoaldosteronism, type IIC, 614492	605232	90	100	100	99
WNK4	Pseudohypoaldosteronism, type IIB, 614491	601844	130	100	100	100
WNT1	Osteogenesis imperfecta, type XV, 615220 {Osteoporosis, early-onset, susceptibility to}, 615221	164820	199	100	100	100
WNT10A	Odontoonychodermal dysplasia, 257980 Schopf-Schulz-Passarge syndrome, 224750 Tooth agenesis, selective, 4, 150400	606268	126	100	100	100
WNT10B	Split-hand/foot malformation 6, 225300 Tooth agenesis, selective, 8, 617073	601906	129	100	100	100

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WNT3	?Tetra-amelia syndrome 1, 273395	165330	151	100	100	99
WNT4	Mullerian aplasia and hyperandrogenism, 158330 ?SERKAL syndrome, 611812	603490	231	100	92	92
WNT5A	Robinow syndrome 1, 180700	164975	127	100	100	99
WNT7A	Fuhrmann syndrome, 228930 Ulna and fibula, absence of, with severe limb deficiency, 276820	601570	150	100	100	100
WRAP53	Dyskeratosis congenita 3, 613988	612661	148	100	100	100
WRN	Werner syndrome, 277700	604611	63	100	100	95
WT1	Denys-Drash syndrome, 194080 Frasier syndrome, 136680 Meacham syndrome, 608978 Mesothelioma, somatic, 156240 Nephrotic syndrome, type 4, 256370 Wilms tumor, type 1, 194070	607102	117	100	100	98
WWOX	Epileptic encephalopathy, early infantile, 28, 616211 Esophageal squamous cell carcinoma, somatic, 133239 Spinocerebellar ataxia 12, 614322	605131	92	100	100	100
XDH	Xanthinuria, type I, 278300	607633	81	100	100	96
XIAP	Lymphoproliferative syndrome, 2, 300635	300079	47	100	95	80
XK	McLeod syndrome with or without chronic granulomatous disease, 300842	314850	62	100	100	96
XPA	Xeroderma pigmentosum, group A, 278700	611153	71	100	100	93
XPC	Xeroderma pigmentosum, group C, 278720	613208	108	100	100	98
XPNPEP3	Nephronophthisis-like nephropathy 1, 613159	613553	63	100	100	96
YAP1	Coloboma, ocular, with or without hearing impairment, cleft lip/palate, and/or mental retardation, 120433	606608	73	100	99	94
YARS1	Charcot-Marie-Tooth disease, dominant intermediate C, 608323	603623	96	100	100	99
YARS2	Myopathy, lactic acidosis, and sideroblastic anemia 2, 613561	610957	127	100	100	99
ZAP70	Autoimmune disease, multisystem, infantile-onset, 2, 617006 Immunodeficiency 48, 269840	176947	151	100	100	100
ZBTB16	Leukemia, acute promyelocytic, PL2F/RARA type Skeletal defects, genital hypoplasia, and mental retardation, 612447	176797	128	100	100	100
ZBTB24	Immunodeficiency-centromeric instability-facial anomalies syndrome 2, 614069	614064	73	100	100	100
ZC4H2	Wieacker-Wolff syndrome, 314580	300897	78	100	99	93
ZDHC9	Mental retardation syndromic, Raymond type, 300799	300646	44	100	97	83
ZEB1	Corneal dystrophy, Fuchs endothelial, 6, 613270 Corneal dystrophy, posterior polymorphous, 3, 609141	189909	84	100	100	98

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x
ZEB2	Mowat-Wilson syndrome, 235730	605802	85	100	100	100
ZFP57	Diabetes mellitus, transient neonatal, 1, 601410	612192	101	100	100	100
ZFPM2	Diaphragmatic hernia 3, 610187 Tetralogy of Fallot, 187500 46XY sex reversal 9, 616067	603693	82	100	100	100
ZFYVE26	Spastic paraplegia 15, 270700	612012	90	100	100	98
ZFYVE27	Spastic paraplegia 33, 610244	610243	99	100	100	100
ZIC1	Craniosynostosis 6, 616602	600470	255	100	100	100
ZIC2	Holoprosencephaly 5, 609637	603073	161	96	95	93
ZIC3	Congenital heart defects, nonsyndromic, 1, 306955 Heterotaxy, visceral, 1, 306955 VACTERL association, 314390	300265	101	100	100	97
ZMPSTE24	Mandibuloacral dysplasia with type B lipodystrophy, 608612 Restrictive dermopathy, lethal, 275210	606480	59	100	100	96
ZMYND10	Ciliary dyskinesia, primary, 22, 615444	607070	130	100	100	100
ZNF335	Microcephaly 10, primary, 615095	610827	110	100	100	100
ZNF423	Joubert syndrome 19, 614844 Nephronophthisis 14, 614844	604557	166	100	100	100
ZNF469	Brittle cornea syndrome 1, 229200	612078	169	100	100	100
ZNF513	?Retinitis pigmentosa 58, 613617	613598	121	100	100	100
ZNF592	No OMIM phenotype	613624	141	100	100	100
ZNF644	Myopia 21, 614167	614159	63	100	100	99
ZNF711	Mental retardation 97, 300803	314990	54	100	96	85
ZNF750	Seborrhea-like dermatitis with psoriasiform elements, 610227	610226	146	100	100	100
ZNF81	No OMIM phenotype	314998	49	100	100	90

- Gene symbols according HGNC

- OMIM release used: 8-9-2019

- "No OMIM phenotypes" indicates a gene without a current OMIM association

- OMIM phenotypes between "[]", indicate "nondiseases," mainly genetic variations that lead to apparently abnormal laboratory test values

- OMIM phenotypes between "{}", indicate risk factors

- OMIM phenotypes with a question mark, "?", before the disease name indicates an unconfirmed or possibly spurious mapping

- The statistics above are based on a set of 100 samples

- Median depth is the median of the mean sequence depth over the protein coding exons ($\pm 10\text{bp}$ flanking introns) of the longest transcript

- % Covered 10x, 20x and 30 x describes the percentage of a gene's coding sequence ($\pm 10\text{bp}$ flanking introns) that is covered at least 10x, 20x or 30x

HGNC approved gene symbol	Phenotype description including OMIM phenotype ID(s)	OMIM gene ID	median depth	% covered >10x	% covered >20x	% covered >30x