

MOVEMENT DISORDERS GENE PANEL DG 2.13 (284 genes)

<i>Gene</i>	<i>Median coverage</i>	<i>% covered > 10x</i>	<i>% covered > 20x</i>	<i>Associated Phenotype description and OMIM disease ID</i>
AARS2	126.2	100	99	Combined oxidative phosphorylation deficiency 8, 614096 Leukoencephalopathy, progressive, with ovarian failure, 615889
ABCB7	131.5	99	98	Anemia, sideroblastic, with ataxia, 301310
ABCD1	76	74	68	Adrenoleukodystrophy, 300100 Adrenomyeloneuropathy, adult, 300100
ABHD12	107	97	88	Polyneuropathy, hearing loss, ataxia, retinitis pigmentosa, and cataract, 612674
ACTB	129	99	94	Baraitser-Winter syndrome 1, 243310 ?Dystonia, juvenile-onset, 607371
ADAR	125	100	99	Aicardi-Goutieres syndrome 6, 615010 Dyschromatosis symmetrica hereditaria, 127400
ADCY5	129.2	92	89	Dyskinesia, familial, with facial myokymia, 606703
ADGRG1	149.7	100	100	Polymicrogyria, bilateral frontoparietal, 606854 Polymicrogyria, bilateral perisylvian, 615752
AFG3L2	121	91	84	Spastic ataxia 5, autosomal recessive, 614487 Spinocerebellar ataxia 28, 610246
AIMP1	84.8	97	89	Leukodystrophy, hypomyelinating, 3, 260600
ALDH18A1	131.1	100	99	Cutis laxa, autosomal dominant 3, 616603 Cutis laxa, autosomal recessive, type IIIA, 219150 Spastic paraplegia 9A, autosomal dominant, 601162 Spastic paraplegia 9B, autosomal recessive, 616586
ALDH3A2	125.7	95	94	Sjogren-Larsson syndrome, 270200
ALS2	170.2	99	99	Amyotrophic lateral sclerosis 2, juvenile, 205100 Primary lateral sclerosis, juvenile, 606353 Spastic paralysis, infantile onset ascending, 607225
ANO10	116.7	98	96	Spinocerebellar ataxia, autosomal recessive 10, 613728
ANO3	138.8	99	98	Dystonia 24, 615034
AP4B1	147.4	100	99	Spastic paraplegia 47, autosomal recessive, 614066
AP4E1	98.7	99	97	Spastic paraplegia 51, autosomal recessive, 613744 Stuttering, familial persistent, 1, 184450

AP4M1	127.2	99	96	Spastic paraplegia 50, autosomal recessive, 612936
AP4S1	65.8	71	69	Spastic paraplegia 52, autosomal recessive, 614067
APTX	118.9	94	91	Ataxia, early-onset, with oculomotor apraxia and hypoalbuminemia, 208920
ARG1	167.8	100	100	Argininemia, 207800
ARSA	97.8	100	99	Metachromatic leukodystrophy, 250100
ARX	29.1	75	59	Epileptic encephalopathy, early infantile, 1, 308350 Hydranencephaly with abnormal genitalia, 300215 Lissencephaly, X-linked 2, 300215 Mental retardation, X-linked 29 and others, 300419 Partington syndrome, 309510 Proud syndrome, 300004
ASPA	127.6	99	95	Canavan disease, 271900
ATCAY	146.1	100	99	Ataxia, cerebellar, Cayman type, 601238
ATL1	161	99	97	Neuropathy, hereditary sensory, type ID, 613708 Spastic paraplegia 3A, autosomal dominant, 182600
ATM	109.7	99	94	Ataxia-telangiectasia, 208900 Lymphoma, B-cell non-Hodgkin, somatic Lymphoma, mantle cell, somatic T-cell prolymphocytic leukemia, somatic {Breast cancer, susceptibility to}, 114480
ATP13A2	117.4	100	98	Kufor-Rakeb syndrome, 606693 ?Ceroid lipofuscinosis, neuronal, 12, 606693
ATP1A2	190.8	100	99	Alternating hemiplegia of childhood, 104290 Migraine, familial basilar, 602481 Migraine, familial hemiplegic, 2, 602481
ATP1A3	177.3	100	100	Alternating hemiplegia of childhood 2, 614820 CAPOS syndrome, 601338 Dystonia-12, 128235
ATP2B3	135.2	99	97	?Spinocerebellar ataxia, X-linked 1, 302500
ATP7B	168.9	100	99	Wilson disease, 277900
B4GALNT1	151	95	90	Spastic paraplegia 26, autosomal recessive, 609195
BCAP31	70.6	93	82	Deafness, dystonia, and cerebral hypomyelination, 300475
BCKDHA	171.5	100	99	Maple syrup urine disease, type Ia, 248600
BCKDHB	112.6	88	81	Maple syrup urine disease, type Ib, 248600

BSCL2	113.5	100	100	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
C12orf65	88.2	97	91	Combined oxidative phosphorylation deficiency 7, 613559 Spastic paraplegia 55, autosomal recessive, 615035
C19orf12	93.9	100	99	Neurodegeneration with brain iron accumulation 4, 614298 ?Spastic paraplegia 43, autosomal recessive, 615043
CA8	114.6	96	93	Cerebellar ataxia and mental retardation with or without quadrupedal locomotion 3, 613227
CACNA1A	87.8	92	89	Episodic ataxia, type 2, 108500 Migraine, familial hemiplegic, 1, 141500 Migraine, familial hemiplegic, 1, with progressive cerebellar ataxia, 141500 Spinocerebellar ataxia 6, 183086
CACNA1G	132.8	99	97	Spinocerebellar ataxia 42, 616795
CACNB4	106.1	96	94	Episodic ataxia, type 5, 613855 {Epilepsy, idiopathic generalized, susceptibility to, 9}, 607682 {Epilepsy, juvenile myoclonic, susceptibility to, 6}, 607682
CAMTA1	185.9	99	98	Cerebellar ataxia, nonprogressive, with mental retardation, 614756
CAPN1	146	100	100	Spastic paraplegia 76, autosomal recessive, 616907
CCT5	164.5	99	99	Neuropathy, hereditary sensory, with spastic paraplegia, 256840
CIZ1	152.8	98	95	No OMIM phenotype Cervical dystonia, primary (Xiao (2012) Ann Neurol 71, 458)
CLCN2	108.6	100	99	Leukoencephalopathy with ataxia, 615651 {Epilepsy, idiopathic generalized, susceptibility to, 11}, 607628 {Epilepsy, juvenile absence, susceptibility to, 2}, 607628 {Epilepsy, juvenile myoclonic, susceptibility to, 8}, 607628
COASY	168.5	100	100	Neurodegeneration with brain iron accumulation 6, 615643
COL4A1	92.8	97	94	Angiopathy, hereditary, with nephropathy, aneurysms, and muscle cramps, 611773 Brain small vessel disease with or without ocular anomalies, 607595 Porencephaly 1, 175780 ?Retinal arteries, tortuosity of, 180000 {Hemorrhage, intracerebral, susceptibility to}, 614519
COQ2	89.3	96	93	Coenzyme Q10 deficiency, primary, 1, 607426 {Multiple system atrophy, susceptibility to}, 146500
COQ4	89.8	88	84	Coenzyme Q10 deficiency, primary, 7, 616276

COQ8A	134.3	100	99	Coenzyme Q10 deficiency, primary, 4, 612016
COQ9	91.4	99	96	Coenzyme Q10 deficiency, primary, 5, 614654
COX20	58.1	83	65	Mitochondrial complex IV deficiency, 220110
CP	120	93	89	Cerebellar ataxia, 604290 Hemosiderosis, systemic, due to aceruloplasminemia, 604290 [Hypoceruloplasminemia, hereditary], 604290
CSF1R	139.5	99	98	Leukoencephalopathy, diffuse hereditary, with spheroids, 221820
CSTB	82.5	97	82	Epilepsy, progressive myoclonic 1A (Unverricht and Lundborg), 254800
CYP27A1	175.1	98	96	Cerebrotendinous xanthomatosis, 213700
CYP2U1	119.2	93	91	Spastic paraplegia 56, autosomal recessive, 615030
CYP7B1	93.2	94	87	Bile acid synthesis defect, congenital, 3, 613812 Spastic paraplegia 5A, autosomal recessive, 270800
DBT	102.1	97	93	Maple syrup urine disease, type II, 248600
DCAF17	91.9	95	89	Woodhouse-Sakati syndrome, 241080
DCC	138.5	100	99	Colorectal cancer, somatic, 114500 Esophageal carcinoma, somatic 133239 Mirror movements 1, 157600
DCTN1	131.6	99	98	Neuropathy, distal hereditary motor, type VIIB, 607641 Perry syndrome, 168605 {Amyotrophic lateral sclerosis, susceptibility to}, 105400
DDC	101	99	95	Aromatic L-amino acid decarboxylase deficiency, 608643
DDHD1	141.8	97	94	Spastic paraplegia 28, autosomal recessive, 609340
DDHD2	149.7	99	98	Spastic paraplegia 54, autosomal recessive, 615033
DLAT	91.6	99	96	Pyruvate dehydrogenase E2 deficiency, 245348
DLD	123.5	99	98	Dihydrolipoamide dehydrogenase deficiency, 246900
DNAL4	72.6	99	93	?Mirror movements 3, 616059
DNMT1	113.4	99	98	Cerebellar ataxia, deafness, and narcolepsy, autosomal dominant, 604121 Neuropathy, hereditary sensory, type IE, 614116
ECHS1	112.8	99	97	Mitochondrial short-chain enoyl-CoA hydratase 1 deficiency, 616277
EIF2B1	149.9	100	100	Leukoencephalopathy with vanishing white matter, 603896
EIF2B2	131.9	100	99	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896
EIF2B3	163.8	100	100	Leukoencephalopathy with vanishing white matter, 603896

EIF2B4	146	100	99	Leukoencephaly with vanishing white matter, 603896 Ovarioleukodystrophy, 603896
EIF2B5	128	99	97	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896
ELOVL4	91.9	99	98	Ichthyosis, spastic quadriplegia, and mental retardation, 614457 Stargardt disease 3, 600110 ?Spinocerebellar ataxia 34, 133190
ELOVL5	120.9	100	99	Spinocerebellar ataxia 38, 615957
ERLIN2	156	100	99	Spastic paraplegia 18, autosomal recessive, 611225
FA2H	94.1	87	79	Spastic paraplegia 35, autosomal recessive, 612319
FAM126A	125.2	97	95	Leukodystrophy, hypomyelinating, 5, 610532
FAR1	80.4	96	92	Peroxisomal fatty acyl-CoA reductase 1 disorder, 616154
FBXO7	189.4	98	93	Parkinson disease 15, autosomal recessive, 260300
FGF14	190.1	100	99	Spinocerebellar ataxia 27, 609307
FLVCR1	139.5	99	95	Ataxia, posterior column, with retinitis pigmentosa, 609033
FOLR1	150.4	100	100	Neurodegeneration due to cerebral folate transport deficiency, 613068
FRMD7	114.2	99	98	Nystagmus 1, congenital, X-linked, 310700 Nystagmus, infantile periodic alternating, X-linked, 310700
FTL	147.7	99	93	Hyperferritinemia-cataract syndrome, 600886 L-ferritin deficiency, dominant and recessive, 615604 Neurodegeneration with brain iron accumulation 3, 606159
GALC	100.6	98	94	Krabbe disease, 245200
GAN	190	100	99	Giant axonal neuropathy-1, 256850
GBA	240.3	100	100	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
GBA2	176.2	99	99	Spastic paraplegia 46, autosomal recessive, 614409
GCDH	147.6	99	99	Glutaricaciduria, type I, 231670
GCH1	74.4	97	86	Dystonia, DOPA-responsive, with or without hyperphenylalaninemia, 128230 Hyperphenylalaninemia, BH4-deficient, B, 233910

GDAP2	131.1	99	97	No OMIM phenotype
GFAP	102.6	91	90	Alexander disease, 203450
GJC2	41.9	68	58	Leukodystrophy, hypomyelinating, 2, 608804 Lymphedema, hereditary, IC, 613480 Spastic paraplegia 44, autosomal recessive, 613206
GLB1	94.3	99	97	GM1-gangliosidosis, type I, 230500 GM1-gangliosidosis, type II, 230600 GM1-gangliosidosis, type III, 230650 Mucopolysaccharidosis type IVB (Morquio), 253010
GNAL	134.9	94	91	Dystonia 25, 615073
GOSR2	127.2	95	95	Epilepsy, progressive myoclonic 6, 614018
GPR143	61.5	85	75	Nystagmus 6, congenital, X-linked, 300814 Ocular albinism, type I, Nettleship-Falls type, 300500
GRID2	175.4	100	99	Spinocerebellar ataxia, autosomal recessive 18, 616204
GRIN1	150.7	100	99	Mental retardation, autosomal dominant 8, 614254
GRIN2B	189.4	99	99	Epileptic encephalopathy, early infantile, 27, 616139 Mental retardation, autosomal dominant 6, 613970
GRM1	185.8	100	99	Spinocerebellar ataxia, autosomal recessive 13, 614831
HEXB	129.7	99	94	Sandhoff disease, infantile, juvenile, and adult forms, 268800
HPRT1	58.2	96	84	HPRT-related gout, 300323 Lesch-Nyhan syndrome, 300322
HSD17B4	95.1	93	90	D-bifunctional protein deficiency, 261515 Perrault syndrome 1, 233400
HSPD1	96.5	98	93	Leukodystrophy, hypomyelinating, 4, 612233 Spastic paraplegia 13, autosomal dominant, 605280
IBA57	113.3	93	89	?Multiple mitochondrial dysfunctions syndrome 3, 615330 ?Spastic paraplegia 74, autosomal recessive, 616451
ITPR1	161.4	100	99	Gillespie syndrome, 206700 Spinocerebellar ataxia 15, 606658 Spinocerebellar ataxia 29, congenital nonprogressive, 117360
KATNB1	141.7	100	100	Lissencephaly 6, with microcephaly, 616212
KCNA1	167	100	99	Episodic ataxia/myokymia syndrome, 160120
KCNA2	157.7	100	99	Epileptic encephalopathy, early infantile, 32, 616366
KCNC1	199.2	100	100	Epilepsy, progressive myoclonic 7, 616187

KCNC3	144	68	59	Spinocerebellar ataxia 13, 605259
KCND3	182.5	99	99	Brugada syndrome 9, 616399 Spinocerebellar ataxia 19, 607346
KCNJ10	213.4	89	89	Enlarged vestibular aqueduct, digenic, 600791 SESAME syndrome, 612780
KCNJ6	156.6	100	99	Keppen-Lubinsky syndrome, 614098
KCNMA1	120.4	94	93	Generalized epilepsy and paroxysmal dyskinesia, 609446
KCTD7	166.7	95	95	Epilepsy, progressive myoclonic 3, with or without intracellular inclusions, 611726
KIDINS220	155.4	99	99	Spastic paraplegia, intellectual disability, nystagmus and obesity, 617296
KIF1A	114	99	96	Mental retardation, autosomal dominant 9, 614255 Neuropathy, hereditary sensory, type IIC, 614213 Spastic paraplegia 30, autosomal recessive, 610357
KIF1C	121.3	99	99	Spastic ataxia 2, autosomal recessive, 611302
KIF5A	136.1	100	99	Spastic paraplegia 10, autosomal dominant, 604187
KMT2B	120.3	94	91	Dystonia 28, childhood-onset, 617284
L1CAM	133.3	99	98	Corpus callosum, partial agenesis of, 304100 CRASH syndrome, 303350 Hydrocephalus due to aqueductal stenosis, 307000 Hydrocephalus with congenital idiopathic intestinal pseudoobstruction, 307000 Hydrocephalus with Hirschsprung disease, 307000 MASA syndrome, 303350
LAMA1	137.5	100	99	Poretti-Boltshauser syndrome, 615960
LAMB1	169.8	100	99	Lissencephaly 5, 615191
LMNB1	123.3	99	99	Leukodystrophy, adult-onset, autosomal dominant, 169500
MARS2	173.2	100	100	Spastic ataxia 3, autosomal recessive, 611390 ?Combined oxidative phosphorylation deficiency 25, 616430
MECP2	87.3	99	93	Encephalopathy, neonatal severe, 300673 Mental retardation, X-linked syndromic, Lubs type, 300260 Mental retardation, X-linked, syndromic 13, 300055 Rett syndrome, 312750 Rett syndrome, atypical, 312750 Rett syndrome, preserved speech variant, 312750 {Autism susceptibility, X-linked 3}, 300496
MECR	108.1	98	96	Dystonia, childhood-onset, with optic atrophy and basal ganglia abnormalities, 617282

MICU1	134.2	96	88	Myopathy with extrapyramidal signs, 615673
MLC1	103.4	100	99	Megalencephalic leukoencephalopathy with subcortical cysts, 604004
MMADHC	77	89	75	Homocystinuria, cblD type, variant 1, 277410 Methylmalonic aciduria and homocystinuria, cblD type, 277410 Methylmalonic aciduria, cblD type, variant 2, 277410
MRE11	51.2	95	82	Ataxia-telangiectasia-like disorder, 604391
MTHFR	126.1	98	97	Homocystinuria due to MTHFR deficiency, 236250 {Neural tube defects, susceptibility to}, 601634 {Schizophrenia, susceptibility to}, 181500 {Thromboembolism, susceptibility to}, 188050 {Vascular disease, susceptibility to}
MTPAP	109.6	98	93	Ataxia, spastic, 4, 613672
MTPP	132.4	99	98	Abetalipoproteinemia, 200100 {Metabolic syndrome, protection against}, 605552
NANS	106.1	100	99	Sponyloepimetaphyseal dysplasia, Genevieve type, 610442
NEFL	164.6	99	98	Charcot-Marie-Tooth disease, type 1F, 607734 Charcot-Marie-Tooth disease, type 2E, 607684
NEXMIF	139.2	99	99	Mental retardation, X-linked 98, 300912
NF2	100.2	100	99	Meningioma, NF2-related, somatic, 607174 Neurofibromatosis, type 2, 101000 Schwannomatosis, 162091
NIPA1	174.3	99	99	Spastic paraplegia 6, autosomal dominant, 600363
NKX2-1	52	96	83	Chorea, hereditary benign, 118700 Choreoathetosis, hypothyroidism, and neonatal respiratory distress, 610978 {Thyroid cancer, nonmedullary, 1}, 188550
NOL3	76.3	93	83	Myoclonus, familial cortical, 614937
NPC1	147.9	99	97	Niemann-Pick disease, type C1, 257220 Niemann-Pick disease, type D, 257220 {Nasopharyngeal carcinoma 1}
NPC2	140.7	100	99	Niemann-pick disease, type C2, 607625
NUP62	111.6	100	99	Striatonigral degeneration, infantile, 271930
OCLN	220.7	100	100	Band-like calcification with simplified gyration and polymicrogyria, 251290

OPA1	122.5	99	94	Behr syndrome,210000 Optic atrophy 1, 165500 Optic atrophy plus syndrome, 125250 ?Mitochondrial DNA depletion syndrome 14 (encephalocardiomyopathic type),616896 {Glaucoma, normal tension, susceptibility to}, 606657
PANK2	146.6	99	93	HARP syndrome, 607236 Neurodegeneration with brain iron accumulation 1, 234200
PAX6	119.9	100	99	Aniridia, 106210 Cataract with late-onset corneal dystrophy, 106210 Coloboma of optic nerve, 120430 Coloboma, ocular, 120200 Foveal hypoplasia 1, 136520 Keratitis, 148190 Optic nerve hypoplasia, 165550 Peters anomaly, 604229 ?Morning glory disc anomaly, 120430
PDE10A	119.8	81	80	Dyskinesia,limb and orofacial,infantile-onset,616921 Striatal degeneration,autosomal dominant,616922
PDE8B	111	99	98	Pigmented nodular adrenocortical disease, primary, 3, 614190 Striatal degeneration, autosomal dominant, 609161
PDGFB	95.1	100	100	Basal ganglia calcification, idiopathic, 5, 615483 Dermatofibrosarcoma protuberans, 607907 Meningioma, SIS-related, 607174
PDGFRB	147.1	99	96	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
PDHA1	109.8	98	92	Pyruvate dehydrogenase E1-alpha deficiency, 312170
PDHX	132.5	98	94	Lacticacidemia due to PDX1 deficiency,245349
PDSS1	116.7	88	78	Coenzyme Q10 deficiency, primary, 2, 614651
PDSS2	126.8	96	93	Coenzyme Q10 deficiency, primary, 3, 614652
PDYN	107.1	100	99	Spinocerebellar ataxia 23, 610245
PEX10	111.8	96	90	Peroxisome biogenesis disorder 6A (Zellweger), 614870 Peroxisome biogenesis disorder 6B, 614871

PEX2	147.1	100	100	Peroxisome biogenesis disorder 5A (Zellweger), 614866 Peroxisome biogenesis disorder 5B, 614867
PEX7	113.5	89	82	Peroxisome biogenesis disorder 9B, 614879 Rhizomelic chondrodysplasia punctata, type 1, 215100
PHYH	74.6	97	90	Refsum disease, 266500
PIK3R5	110.1	100	99	Ataxia-oculomotor apraxia 3, 615217
PLA2G6	117.5	99	98	Infantile neuroaxonal dystrophy 1, 256600 Neurodegeneration with brain iron accumulation 2B, 610217 Parkinson disease 14, autosomal recessive, 612953
PLP1	129.2	100	99	Pelizaeus-Merzbacher disease, 312080 Spastic paraplegia 2, X-linked, 312920
PMM2	141.1	99	99	Congenital disorder of glycosylation, type Ia, 212065
PMPCA	120.8	99	96	Spinocerebellar ataxia, autosomal recessive 2, 213200
PNKD	99.8	100	99	Paroxysmal nonkinesigenic dyskinesia, 118800
PNKP	93	99	97	Ataxia-oculomotor apraxia 4, 616267 Microcephaly, seizures, and developmental delay, 613402
PNPLA6	122.1	99	98	Boucher-Neuhauser syndrome, 215470 Oliver-McFarlane syndrome, 275400 Spastic paraplegia 39, autosomal recessive, 612020 ?Laurence-Moon syndrome, 245800
POLG	114.4	100	99	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, autosomal dominant 1, 157640 Progressive external ophthalmoplegia, autosomal recessive 1, 258450
POLR1C	117	99	96	Leukodystrophy, hypomyelinating, 11, 616494 Treacher Collins syndrome 3, 248390
POLR3A	137.4	100	99	Leukodystrophy, hypomyelinating, 7, with or without oligodontia and/or hypogonadotropic hypogonadism, 607694
POLR3B	146.4	99	98	Leukodystrophy, hypomyelinating, 8, with or without oligodontia and/or hypogonadotropic hypogonadism, 614381
PRF1	122.5	91	90	Aplastic anemia, 609135 Hemophagocytic lymphohistiocytosis, familial, 2, 603553 Lymphoma, non-Hodgkin, 605027

PRKCG	116.1	99	94	Spinocerebellar ataxia 14, 605361
PRKRA	179.6	99	98	Dystonia 16, 612067
PRRT2	78.9	99	98	Convulsions, familial infantile, with paroxysmal choreoathetosis, 602066 Episodic kinesigenic dyskinesia 1, 128200 Seizures, benign familial infantile, 2, 605751
PSAP	114.4	99	99	Combined SAP deficiency, 611721 Gaucher disease, atypical, 610539 Krabbe disease, atypical, 611722 Metachromatic leukodystrophy due to SAP-b deficiency, 249900
PUM1	158	100	99	No OMIM phenotype
PYCR2	127.6	100	97	Leukodystrophy, hypomyelinating, 10, 616420
RAB18	82.7	97	86	Warburg micro syndrome 3, 614222
RAB3GAP1	124.2	99	98	Warburg micro syndrome 1, 600118
RAB3GAP2	94.1	98	93	Martsolf syndrome, 212720 Warburg micro syndrome 2, 614225
RAD51	123.2	89	89	Mirror movements 2, 614508 {Breast cancer, susceptibility to}, 114480 ?Fanconi anemia, complementation group R, 617244
RARS	86.4	92	85	Leukodystrophy, hypomyelinating, 9, 616140
REEP1	78.3	76	75	Spastic paraplegia 31, autosomal dominant, 610250 ?Neuronopathy, distal hereditary motor, type VB, 614751
RNASEH2A	142.1	100	99	Aicardi-Goutieres syndrome 4, 610333
RNASEH2B	103.8	93	87	Aicardi-Goutieres syndrome 2, 610181
RNASEH2C	209.2	100	99	Aicardi-Goutieres syndrome 3, 610329
RNF170	147.1	98	91	Ataxia, sensory, 1, autosomal dominant, 608984
RNF216	137.1	99	98	Cerebellar ataxia and hypogonadotropic hypogonadism, 212840
RTN2	104.8	99	96	Spastic paraplegia 12, autosomal dominant, 604805
RUBCN	104.1	98	97	?Spinocerebellar ataxia, autosomal recessive 15, 615705
SACS	154.5	100	99	Spastic ataxia, Charlevoix-Saguenay type, 270550
SAMD9L	165.7	100	99	Ataxia-pancytopenia syndrome, 159550
SAMHD1	127.9	99	96	Aicardi-Goutieres syndrome 5, 612952 ?Chilblain lupus 2, 614415
SCN11A	138.1	99	97	Episodic pain syndrome, familial, 3, 615552 Neuropathy, hereditary sensory and autonomic, type VII, 615548

SCN8A	198.3	100	99	Epileptic encephalopathy, early infantile, 13, 614558 ?Cognitive impairment with or without cerebellar ataxia, 614306
SERAC1	112.5	98	94	3-methylglutaconic aciduria with deafness, encephalopathy, and Leigh-like syndrome, 614739
SETX	163.2	99	99	Amyotrophic lateral sclerosis 4, juvenile, 602433 Spinocerebellar ataxia, autosomal recessive 1, 606002
SGCE	88.9	93	90	Dystonia-11, myoclonic, 159900
SIL1	154.4	99	98	Marinesco-Sjogren syndrome, 248800
SLC12A6	141.8	100	99	Agenesis of the corpus callosum with peripheral neuropathy, 218000
SLC16A2	60.3	92	82	Allan-Herndon-Dudley syndrome, 300523
SLC19A3	186.4	100	99	Thiamine metabolism dysfunction syndrome 2 (biotin- or thiamine-responsive encephalopathy type 2), 607483
SLC1A3	121.9	100	99	Episodic ataxia, type 6, 612656
SLC20A2	119	99	97	Basal ganglia calcification, idiopathic, 1, 213600
SLC25A15	192.5	98	95	Hyperornithinemia-hyperammonemia-homocitrullinemia syndrome, 238970
SLC2A1	190.1	92	92	Dystonia 9, 601042 GLUT1 deficiency syndrome 1, infantile onset, severe, 606777 GLUT1 deficiency syndrome 2, childhood onset, 612126 Stomatin-deficient cryohydrocytosis with neurologic defects, 608885 {Epilepsy, idiopathic generalized, susceptibility to, 12}, 614847
SLC30A10	164.4	100	100	Hypermanganesemia with dystonia 1, 613280
SLC33A1	140.9	96	90	Congenital cataracts, hearing loss, and neurodegeneration, 614482 Spastic paraplegia 42, autosomal dominant, 612539
SLC39A14	107.7	99	97	Hypermanganesemia with dystonia 2, 617013
SLC52A2	177.6	100	100	Brown-Vialetto-Van Laere syndrome 2, 614707
SLC52A3	119.6	100	100	Brown-Vialetto-Van Laere syndrome 1, 211530 Fazio-Londe disease, 211500
SLC6A3	145.7	100	99	Parkinsonism-dystonia, infantile, 613135 {Nicotine dependence, protection against}, 188890
SLC9A1	160.9	100	100	?Lichtenstein-Knorr syndrome, 616291
SMPD1	123.5	99	97	Niemann-Pick disease, type A, 257200 Niemann-Pick disease, type B, 607616
SNCA	129.8	100	100	Dementia, Lewy body, 127750 Parkinson disease 1, 168601 Parkinson disease 4, 605543

SNORD118				Leukoencephalopathy, brain calcifications, and cysts,614561
SNX14	70.1	95	82	Spinocerebellar ataxia, autosomal recessive 20, 616354
SOX10	65.8	98	91	PCWH syndrome, 609136 Waardenburg syndrome, type 2E, with or without neurologic involvement, 611584 Waardenburg syndrome, type 4C, 613266
SPART	132.4	99	98	Troyer syndrome, 275900
SPAST	63.8	93	81	Spastic paraplegia 4, autosomal dominant, 182601
SPG11	129.2	99	96	Amyotrophic lateral sclerosis 5, juvenile, 602099 Charcot-Marie-Tooth disease, axonal, type 2X, 616668 Spastic paraplegia 11, autosomal recessive, 604360
SPG21	121.4	98	94	Mast syndrome, 248900
SPG7	119.2	93	92	Spastic paraplegia 7, autosomal recessive, 607259
SPR	166.5	98	90	Dystonia, dopa-responsive, due to sepiapterin reductase deficiency, 612716
SPTBN2	118	99	99	Spinocerebellar ataxia 5, 600224 Spinocerebellar ataxia, autosomal recessive 14, 615386
STUB1	176.1	100	98	Spinocerebellar ataxia, autosomal recessive 16, 615768
SUOX	212.6	100	100	Sulfite oxidase deficiency, 272300
SYNE1	136.6	98	97	Emery-Dreifuss muscular dystrophy 4, autosomal dominant, 612998 Spinocerebellar ataxia, autosomal recessive 8, 610743
TAF1	112.4	99	96	Dystonia-Parkinsonism, X-linked, 314250 Mental retardation, X-linked, syndromic 33, 300966
TANGO2	145.3	100	100	Metabolic encephalomyopathic crises,recurrent,with rhabdomyolysis,cardiac arrhythmias and neurodegeneration,616878
TBC1D20	145.7	94	94	Warburg micro syndrome 4, 615663
TDP1	122.9	98	95	Spinocerebellar ataxia, autosomal recessive with axonal neuropathy, 607250
TDP2	165.1	99	98	Spinocerebellar ataxia, autosomal recessive, 616949
TECPR2	161.1	100	99	Spastic paraplegia 49, autosomal recessive, 615031
TENM4	160.8	99	99	Tremor, hereditary essential, 5, 616736
TGM6	149.7	99	98	Spinocerebellar ataxia 35, 613908
TH	68.2	97	88	Segawa syndrome, recessive, 605407
THAP1	122	100	100	Dystonia 6, torsion, 602629
TIMM8A	46	94	78	Jensen syndrome, 311150 Mohr-Tranebjaerg syndrome, 304700

TMEM106B	120.2	99	96	No OMIM phenotype Hypomyelinating leukodystrophy (Simons (2017) Brain 140, 3105)
TMEM240	112.2	99	97	Spinocerebellar ataxia 21, 607454
TMEM67	72.9	93	83	COACH syndrome, 216360 Joubert syndrome 6, 610688 Meckel syndrome 3, 607361 Nephronophthisis 11, 613550 {Bardet-Biedl syndrome 14, modifier of}, 615991
TOR1A	185	100	99	Dystonia-1, torsion, 128100 {Dystonia-1, modifier of}
TPP1	146.3	100	100	Ceroid lipofuscinosis, neuronal, 2, 204500 Spinocerebellar ataxia, autosomal recessive 7, 609270
TREM2	149	99	99	Nasu-Hakola disease, 221770
TREX1	242.4	100	100	Aicardi-Goutieres syndrome 1, dominant and recessive, 225750 Chilblain lupus, 610448 Vasculopathy, retinal, with cerebral leukodystrophy, 192315 {Systemic lupus erythematosus, susceptibility to}, 152700
TSEN54	82.9	95	92	Pontocerebellar hypoplasia type 2A, 277470 Pontocerebellar hypoplasia type 4, 225753 ?Pontocerebellar hypoplasia type 5, 610204
TTBK2	123.2	100	98	Spinocerebellar ataxia 11, 604432
TTC19	92.1	80	72	Mitochondrial complex III deficiency, nuclear type 2, 615157
TTPA	101.5	83	76	Ataxia with isolated vitamin E deficiency, 277460
TUBA1A	113.2	99	97	Lissencephaly 3, 611603
TUBB4A	121.2	96	95	Dystonia 4, torsion, autosomal dominant, 128101 Leukodystrophy, hypomyelinating, 6, 612438
TUBG1	164.2	100	100	Cortical dysplasia, complex, with other brain malformations 4, 615412
TWNK	178.8	100	100	Mitochondrial DNA depletion syndrome 7 (hepatocerebral type), 271245 Perrault syndrome 5, 616138 Progressive external ophthalmoplegia with mitochondrial DNA deletions, autosomal dominant 3, 609286
TYROBP	95.2	100	99	Nasu-Hakola disease, 221770
VAMP1	131.5	100	100	Spastic ataxia 1, autosomal dominant, 108600

VCP	144.8	99	99	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
VLDLR	200.9	99	99	Cerebellar hypoplasia and mental retardation with or without quadrupedal locomotion 1, 224050
VPS13A	69.5	95	85	Choreoacanthocytosis, 200150
VPS13D	158.6	99	99	No OMIM phenotype Schizophrenia (McCarthy (2014) Mol Psychiatry 19, 652)
VPS37A	73.6	86	66	Spastic paraplegia 53, autosomal recessive, 614898
VRK1	124.8	97	94	Pontocerebellar hypoplasia type 1A, 607596
WASHC5	146.6	99	98	Ritscher-Schinzel syndrome 1, 220210 Spastic paraplegia 8, autosomal dominant, 603563
WDR45	75	97	90	Neurodegeneration with brain iron accululation 5, 300894
WDR81	163.3	99	99	Cerebellar ataxia, mental retardation, and dysequilibrium syndrome 2, 610185
WWOX	130.9	100	99	Epileptic encephalopathy, early infantile, 28, 616211 Esophageal squamous cell carcinoma, somatic, 133239 Spinocerebellar ataxia, autosomal recessive 12, 614322
XK	96.8	99	99	McLeod syndrome with or without chronic granulomatous disease, 300842
XPR1	131.7	100	99	Basal ganglia calcification, idiopathic, 6, 616413
XRCC1	106.6	99	97	?Spinocerebellar ataxia, autosomal recessive 26,617633
ZC4H2	78.6	99	98	Wieacker-Wolff syndrome, 314580
ZFYVE26	120.3	99	99	Spastic paraplegia 15, autosomal recessive, 270700
ZFYVE27	118.3	100	100	Spastic paraplegia 33, autosomal dominant, 610244
ZNF592	150.1	100	99	Spinocerebellar ataxia, autosomal recessive 5, 251300

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

Median Coverage describes the average number of reads seen across 50 exomes.

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

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

Genes with Median Coverage and % Covered 10x/20x denoting NC are non-coding genes for which coverage statistics could not be generated.

OMIM release used for OMIM disease identifiers and descriptions : April 18th, 2018.

This list is accurate for panel version DG 2.13

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