

MOVEMENT DISORDERS GENE PANEL DG 2.11 (275 genes)

<i>Gene</i>	<i>Median</i>	<i>% covered > 10x</i>	<i>% covered > 20x</i>	<i>Associated Phenotype description and OMIM disease ID</i>
AARS2	126.4	99	99	Combined oxidative phosphorylation deficiency 8, 614096 Leukoencephalopathy, progressive, with ovarian failure, 615889
ABCB7	132.4	99	98	Anemia, sideroblastic, with ataxia, 301310
ABCD1	76.6	74	68	Adrenoleukodystrophy, 300100 Adrenomyeloneuropathy, adult, 300100
ABHD12	107	97	87	Polyneuropathy, hearing loss, ataxia, retinitis pigmentosa, and cataract, 612674
ACTB	129.3	99	94	Baraitser-Winter syndrome 1, 243310 ?Dystonia, juvenile-onset, 607371
ADAR	122.6	100	99	Aicardi-Goutieres syndrome 6, 615010 Dyschromatosis symmetrica hereditaria, 127400
ADCK3	134.7	100	99	Coenzyme Q10 deficiency, primary, 4, 612016
ADCY5	129.5	92	89	Dyskinesia, familial, with facial myokymia, 606703
AFG3L2	121.1	91	84	Spastic ataxia 5, autosomal recessive, 614487 Spinocerebellar ataxia 28, 610246
AIMP1	84.7	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	131.7	100	99	Sjogren-Larsson syndrome, 270200
ALS2	167.7	99	98	Amyotrophic lateral sclerosis 2, juvenile, 205100 Primary lateral sclerosis, juvenile, 606353 Spastic paraparesis, infantile onset ascending, 607225
ANO10	116.6	98	96	Spinocerebellar ataxia, autosomal recessive 10, 613728
ANO3	138.7	99	98	Dystonia 24, 615034
AP4B1	146.6	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	127.3	99	96	Spastic paraplegia 50, autosomal recessive, 612936
AP4S1	65	71	68	Spastic paraplegia 52, autosomal recessive, 614067
AP5Z1	97	99	96	Spastic paraplegia 48, autosomal recessive, 613647
APTX	119.1	94	91	Ataxia, early-onset, with oculomotor apraxia and hypoalbuminemia, 208920
ARG1	167.9	100	100	Argininemia, 207800
ARSA	97.9	100	99	Metachromatic leukodystrophy, 250100
ARX	29.3	76	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	127.5	99	95	Canavan disease, 271900
ATCAY	149.5	100	99	Ataxia, cerebellar, Cayman type, 601238
ATL1	160.3	99	97	Neuropathy, hereditary sensory, type ID, 613708 Spastic paraplegia 3A, autosomal dominant, 182600
ATM	109.8	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	117.5	100	98	Kufor-Rakeb syndrome, 606693 ?Ceroid lipofuscinosis, neuronal, 12, 606693
ATP1A2	191.1	100	99	Alternating hemiplegia of childhood, 104290 Migraine, familial basilar, 602481 Migraine, familial hemiplegic, 2, 602481
ATP1A3	177.6	100	100	Alternating hemiplegia of childhood 2, 614820 CAPOS syndrome, 601338 Dystonia-12, 128235
ATP2B3	136.3	99	97	?Spinocerebellar ataxia, X-linked 1, 302500
ATP7B	169.1	100	99	Wilson disease, 277900
B4GALNT1	151.3	95	90	Spastic paraplegia 26, autosomal recessive, 609195

BCAP31	71.2	93	82	Deafness, dystonia, and cerebral hypomyelination, 300475
BCKDHA	172.8	100	99	Maple syrup urine disease, type Ia, 248600
BCKDHB	112.6	88	81	Maple syrup urine disease, type Ib, 248600
BSCL2	113.7	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
C10orf2	179	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
C12orf65	88.1	97	91	Combined oxidative phosphorylation deficiency 7, 613559 Spastic paraplegia 55, autosomal recessive, 615035
C19orf12	94.3	100	99	Neurodegeneration with brain iron accumulation 4, 614298 ?Spastic paraplegia 43, autosomal recessive, 615043
CA8	114.5	96	92	Cerebellar ataxia and mental retardation with or without quadrupedal locomotion 3, 613227
CACNA1A	87.9	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	133	99	97	Spinocerebellar ataxia 42, 616795
CACNB4	107.6	97	95	Episodic ataxia, type 5, 613855 {Epilepsy, idiopathic generalized, susceptibility to, 9}, 607682 {Epilepsy, juvenile myoclonic, susceptibility to, 6}, 607682
CAMTA1	186.2	99	98	Cerebellar ataxia, nonprogressive, with mental retardation, 614756
CAPN1	146.3	100	100	Spastic paraplegia 76, autosomal recessive, 616907
CCT5	164.6	99	99	Neuropathy, hereditary sensory, with spastic paraplegia, 256840
CIZ1	155.8	98	95	No OMIM phenotype Cervical dystonia, primary (Xiao (2012) Ann Neurol 71, 458)

CLCN2	108.7	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	89.2	96	93	Coenzyme Q10 deficiency, primary, 1, 607426 {Multiple system atrophy, susceptibility to}, 146500
COQ9	91.6	99	96	Coenzyme Q10 deficiency, primary, 5, 614654
COX20	57.7	82	65	Mitochondrial complex IV deficiency, 220110
CP	120	93	89	Cerebellar ataxia, 604290 Hemosiderosis, systemic, due to aceruloplasminemia, 604290 [Hypoceruloplasminemia, hereditary], 604290
CSF1R	140	99	98	Leukoencephalopathy, diffuse hereditary, with spheroids, 221820
CSTB	100	98	87	Epilepsy, progressive myoclonic 1A (Unverricht and Lundborg), 254800
CYP27A1	175.2	98	96	Cerebrotendinous xanthomatosis, 213700
CYP2U1	119.2	93	91	Spastic paraplegia 56, autosomal recessive, 615030
CYP7B1	92.8	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	139.3	100	99	Colorectal cancer, somatic, 114500 Esophageal carcinoma, somatic 133239 Mirror movements 1, 157600
DCTN1	131.9	99	98	Neuropathy, distal hereditary motor, type VIIIB, 607641 Perry syndrome, 168605 {Amyotrophic lateral sclerosis, susceptibility to}, 105400

DDC	101.1	99	95	Aromatic L-amino acid decarboxylase deficiency, 608643
DDHD1	141.7	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	72.7	99	93	?Mirror movements 3, 616059
DNMT1	113.5	99	98	Cerebellar ataxia, deafness, and narcolepsy, autosomal dominant, 604121 Neuropathy, hereditary sensory, type IE, 614116
ECHS1	112.7	99	97	Mitochondrial short-chain enoyl-CoA hydratase 1 deficiency, 616277
EIF2B1	149.9	100	100	Leukoencephalopathy with vanishing white matter, 603896
EIF2B2	132	100	99	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896
EIF2B3	164.1	100	100	Leukoencephalopathy with vanishing white matter, 603896
EIF2B4	146.1	100	99	Leukoencephaly with vanishing white matter, 603896 Ovarioleukodystrophy, 603896
EIF2B5	128	99	98	Leukoencephalopathy with vanishing white matter, 603896 Ovarioleukodystrophy, 603896
EIF4G1	135.1	100	99	{Parkinson disease 18}, 614251
ELOVL4	91.9	99	98	Ichthyosis, spastic quadriplegia, and mental retardation, 614457 Stargardt disease 3, 600110 ?Spinocerebellar ataxia 34, 133190
ELOVL5	121.9	100	99	Spinocerebellar ataxia 38, 615957
ERLIN2	156.2	100	99	Spastic paraplegia 18, autosomal recessive, 611225
FA2H	94.2	87	79	Spastic paraplegia 35, autosomal recessive, 612319
FAM126A	124.9	97	95	Leukodystrophy, hypomyelinating, 5, 610532
FAR1	80.3	96	92	Peroxisomal fatty acyl-CoA reductase 1 disorder, 616154
FBXO7	189.7	98	93	Parkinson disease 15, autosomal recessive, 260300
FGF14	190.3	100	99	Spinocerebellar ataxia 27, 609307
FLVCR1	139.4	99	95	Ataxia, posterior column, with retinitis pigmentosa, 609033
FOLR1	150.5	100	100	Neurodegeneration due to cerebral folate transport deficiency, 613068
FRMD7	114.9	99	98	Nystagmus 1, congenital, X-linked, 310700 Nystagmus, infantile periodic alternating, X-linked, 310700

FTL	148.1	98	93	Hyperferritinemia-cataract syndrome, 600886 L-ferritin deficiency, dominant and recessive, 615604 Neurodegeneration with brain iron accumulation 3, 606159
GALC	100.5	98	94	Krabbe disease, 245200
GAN	190.2	100	99	Giant axonal neuropathy-1, 256850
GBA	240.7	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	176.3	99	99	Spastic paraparesis 46, autosomal recessive, 614409
GCDH	133.7	94	91	Glutaric aciduria, type I, 231670
GCH1	74.4	97	86	Dystonia, DOPA-responsive, with or without hyperphenylalaninemia, 128230 Hyperphenylalaninemia, BH4-deficient, B, 233910
GDAP2	130.9	99	97	No OMIM phenotype
GFAP	99.1	99	98	Alexander disease, 203450
GJC2	42.1	68	58	Leukodystrophy, hypomyelinating, 2, 608804 Lymphedema, hereditary, IC, 613480 Spastic paraparesis 44, autosomal recessive, 613206
GLB1	91.6	99	97	GM1-gangliosidosis, type I, 230500 GM1-gangliosidosis, type II, 230600 GM1-gangliosidosis, type III, 230650 Mucopolysaccharidosis type IVB (Morquio), 253010
GNAL	134.8	94	91	Dystonia 25, 615073
GOSR2	123.2	97	96	Epilepsy, progressive myoclonic 6, 614018
GPR143	60.5	85	74	Nystagmus 6, congenital, X-linked, 300814 Ocular albinism, type I, Nettleship-Falls type, 300500
GPR56	150.9	100	100	Polymicrogyria, bilateral frontoparietal, 606854 Polymicrogyria, bilateral perisylvian, 615752
GRID2	175.4	100	99	Spinocerebellar ataxia, autosomal recessive 18, 616204
GRM1	185.7	100	99	Spinocerebellar ataxia, autosomal recessive 13, 614831
HEXB	129.7	99	93	Sandhoff disease, infantile, juvenile, and adult forms, 268800

HPRT1	58.6	96	84	HPRT-related gout, 300323 Lesch-Nyhan syndrome, 300322
HSPD1	99.8	98	93	Leukodystrophy, hypomyelinating, 4, 612233 Spastic paraplegia 13, autosomal dominant, 605280
IBA57	113.6	93	89	?Multiple mitochondrial dysfunctions syndrome 3, 615330 ?Spastic paraplegia 74, autosomal recessive, 616451
ITPR1	161.5	100	99	Gillespie syndrome, 206700 Spinocerebellar ataxia 15, 606658 Spinocerebellar ataxia 29, congenital nonprogressive, 117360
KCNA1	167	100	99	Episodic ataxia/myokymia syndrome, 160120
KCNA2	162.2	100	100	Epileptic encephalopathy, early infantile, 32, 616366
KCNC1	199.4	100	100	Epilepsy, progressive myoclonic 7, 616187
KCNC3	129.7	63	54	Spinocerebellar ataxia 13, 605259
KCND3	182.9	99	99	Brugada syndrome 9, 616399 Spinocerebellar ataxia 19, 607346
KCNJ10	238.8	100	99	Enlarged vestibular aqueduct, digenic, 600791 SESAME syndrome, 612780
KCNJ6	156	100	99	Keppen-Lubinsky syndrome, 614098
KCNMA1	131.4	100	99	Generalized epilepsy and paroxysmal dyskinesia, 609446
KCTD7	148.9	94	92	Epilepsy, progressive myoclonic 3, with or without intracellular inclusions, 611726
KIAA0196	146.6	99	98	Ritscher-Schinzel syndrome 1, 220210 Spastic paraplegia 8, autosomal dominant, 603563
KIAA0226	104.4	98	97	?Spinocerebellar ataxia, autosomal recessive 15, 615705
KIAA2022	140.1	99	99	Mental retardation, X-linked 98, 300912
KIDINS220	155.4	99	99	Spastic paraplegia, intellectual disability, nystagmus and obesity, 617296
KIF1A	114.2	99	96	Mental retardation, autosomal dominant 9, 614255 Neuropathy, hereditary sensory, type IIC, 614213 Spastic paraplegia 30, autosomal recessive, 610357
KIF1C	121.5	100	99	Spastic ataxia 2, autosomal recessive, 611302
KIF5A	136.3	100	99	Spastic paraplegia 10, autosomal dominant, 604187
KMT2B	120.5	94	91	Dystonia 28, childhood-onset, 617284

L1CAM	133.8	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	137.3	100	99	Poretti-Boltshauser syndrome, 615960
LMNB1	123.3	99	99	Leukodystrophy, adult-onset, autosomal dominant, 169500
MARS2	173.3	100	100	Spastic ataxia 3, autosomal recessive, 611390 ?Combined oxidative phosphorylation deficiency 25, 616430
MECP2	87.9	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	134.1	96	89	Myopathy with extrapyramidal signs, 615673
MLC1	103.6	100	99	Megalencephalic leukoencephalopathy with subcortical cysts, 604004
MMADHC	77	89	74	Homocystinuria, cbID type, variant 1, 277410 Methylmalonic aciduria and homocystinuria, cbID type, 277410 Methylmalonic aciduria, cbID type, variant 2, 277410
MRE11A	51.2	95	82	Ataxia-telangiectasia-like disorder, 604391
MTHFR	131.3	100	100	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	107.7	99	93	Ataxia, spastic, 4, 613672
MTTP	132.3	99	98	Abetalipoproteinemia, 200100 {Metabolic syndrome, protection against}, 605552

NEFL	165	99	98	Charcot-Marie-Tooth disease, type 1F, 607734 Charcot-Marie-Tooth disease, type 2E, 607684
NIPA1	174.6	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, monmedullary, 1}, 188550
NOL3	76.2	93	83	Myoclonus, familial cortical, 614937
NPC1	148.1	99	97	Niemann-Pick disease, type C1, 257220 Niemann-Pick disease, type D, 257220 {Nasopharyngeal carcinoma 1}
NPC2	140.9	100	99	Niemann-pick disease, type C2, 607625
NUP62	111.6	100	99	Striatonigral degeneration, infantile, 271930
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	121.9	100	100	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	147	99	99	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	95.2	100	100	Basal ganglia calcification, idiopathic, 5, 615483 Dermatofibrosarcoma protuberans, 607907 Meningioma, SIS-related, 607174
PDGFRB	147.2	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	110.5	98	92	Pyruvate dehydrogenase E1-alpha deficiency, 312170
PDHX	132.5	98	94	Lacticacidemia due to PDX1 deficiency, 245349
PDSS1	117.2	88	78	Coenzyme Q10 deficiency, primary, 2, 614651
PDSS2	126.9	96	93	Coenzyme Q10 deficiency, primary, 3, 614652
PDYN	107.1	100	99	Spinocerebellar ataxia 23, 610245
PEX10	112.1	96	90	Peroxisome biogenesis disorder 6A (Zellweger), 614870 Peroxisome biogenesis disorder 6B, 614871
PEX2	147	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	74.7	97	90	Refsum disease, 266500
PIK3R5	110.3	100	99	Ataxia-oculomotor apraxia 3, 615217
PLA2G6	117.6	99	98	Infantile neuroaxonal dystrophy 1, 256600 Neurodegeneration with brain iron accumulation 2B, 610217 Parkinson disease 14, autosomal recessive, 612953
PLP1	130.4	100	99	Pelizaeus-Merzbacher disease, 312080 Spastic paraplegia 2, X-linked, 312920
PMM2	141.3	99	99	Congenital disorder of glycosylation, type Ia, 212065
PMPCA	120.7	99	96	Spinocerebellar ataxia, autosomal recessive 2, 213200
PNKD	99.9	100	99	Paroxysmal nonkinesigenic dyskinesia, 118800
PNKP	93.1	99	97	Ataxia-oculomotor apraxia 4, 616267 Microcephaly, seizures, and developmental delay, 613402
PNPLA6	122.4	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	117.2	99	96		Leukodystrophy, hypomyelinating, 11, 616494 Treacher Collins syndrome 3, 248390
POLR3A	137.8	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	134.2	100	99		Aplastic anemia, 609135 Hemophagocytic lymphohistiocytosis, familial, 2, 603553 Lymphoma, non-Hodgkin, 605027
PRKCG	114.5	98	93		Spinocerebellar ataxia 14, 605361
PRKRA	178.7	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	114.3	99	99		Combined SAP deficiency, 611721 Gaucher disease, atypical, 610539 Krabbe disease, atypical, 611722 Metachromatic leukodystrophy due to SAP-b deficiency, 249900
PYCR2	127.8	100	97		Leukodystrophy, hypomyelinating, 10, 616420
RAB18	80.4	96	84		Warburg micro syndrome 3, 614222
RAB3GAP1	124.2	99	98		Warburg micro syndrome 1, 600118

RAB3GAP2	94.2	98	93	Martolf syndrome, 212720 Warburg micro syndrome 2, 614225
RAD51	123.3	89	89	Mirror movements 2, 614508 {Breast cancer, susceptibility to}, 114480 ?Fanconi anemia, complementation group R, 617244
RARS	91.5	98	90	Leukodystrophy, hypomyelinating, 9, 616140
REEP1	99.6	96	95	Spastic paraparesis 31, autosomal dominant, 610250 ?Neuronopathy, distal hereditary motor, type VB, 614751
RNASEH2A	142.2	100	99	Aicardi-Goutieres syndrome 4, 610333
RNASEH2B	103.8	93	87	Aicardi-Goutieres syndrome 2, 610181
RNASEH2C	209.6	100	99	Aicardi-Goutieres syndrome 3, 610329
RNF170	147.3	98	91	Ataxia, sensory, 1, autosomal dominant, 608984
RNF216	137	99	98	Cerebellar ataxia and hypogonadotropic hypogonadism, 212840
RTN2	104.8	99	96	Spastic paraparesis 12, autosomal dominant, 604805
SACS	154.4	100	99	Spastic ataxia, Charlevoix-Saguenay type, 270550
SAMD9L	165.3	100	99	Ataxia-pancytopenia syndrome, 159550
SAMHD1	127.8	99	96	Aicardi-Goutieres syndrome 5, 612952 ?Chilblain lupus 2, 614415
SCN11A	138	99	97	Episodic pain syndrome, familial, 3, 615552 Neuropathy, hereditary sensory and autonomic, type VII, 615548
SCN8A	198.5	100	99	Epileptic encephalopathy, early infantile, 13, 614558 ?Cognitive impairment with or without cerebellar ataxia, 614306
SERAC1	112.4	98	94	3-methylglutaconic aciduria with deafness, encephalopathy, and Leigh-like syndrome, 614739
SETX	162.7	99	99	Amyotrophic lateral sclerosis 4, juvenile, 602433 Spinocerebellar atrophy, autosomal recessive 1, 606002
SGCE	88.9	93	90	Dystonia-11, myoclonic, 159900
SIL1	154.8	99	98	Marinesco-Sjogren syndrome, 248800
SLC12A6	141.8	100	99	Agenesis of the corpus callosum with peripheral neuropathy, 218000
SLC16A2	57.7	93	81	Allan-Herndon-Dudley syndrome, 300523
SLC19A3	178	100	99	Thiamine metabolism dysfunction syndrome 2 (biotin- or thiamine-responsive encephalopathy type 2), 607483

SLC1A3	122.2	100	99	Episodic ataxia, type 6, 612656
SLC20A2	119.1	99	97	Basal ganglia calcification, idiopathic, 1, 213600
SLC25A15	192.8	98	95	Hyperornithinemia-hyperammonemia-homocitrullinemia syndrome, 238970
SLC2A1	179.6	100	100	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	161.7	100	99	Hypermanganesemia with dystonia 1, 613280
SLC33A1	140.7	96	90	Congenital cataracts, hearing loss, and neurodegeneration, 614482 Spastic paraplegia 42, autosomal dominant, 612539
SLC39A14	107.9	99	98	Hypermanganesemia with dystonia 2, 617013
SLC52A2	177.8	100	100	Brown-Vialetto-Van Laere syndrome 2, 614707
SLC52A3	119.7	100	100	Brown-Vialetto-Van Laere syndrome 1, 211530 Fazio-Londe disease, 211500
SLC6A3	146	100	99	Parkinsonism-dystonia, infantile, 613135 {Nicotine dependence, protection against}, 188890
SLC9A1	161.2	100	100	?Lichtenstein-Knorr syndrome, 616291
SMPD1	127.5	99	98	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	NC	NC	NC	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
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
SPG20	132.3	99	98	Troyer syndrome, 275900
SPG21	121.5	98	94	Mast syndrome, 248900
SPG7	119.4	93	92	Spastic paraplegia 7, autosomal recessive, 607259
SPR	166.9	98	89	Dystonia, dopa-responsive, due to sepiapterin reductase deficiency, 612716
SPTBN2	118.4	99	99	Spinocerebellar ataxia 5, 600224 Spinocerebellar ataxia, autosomal recessive 14, 615386
STUB1	176.5	100	98	Spinocerebellar ataxia, autosomal recessive 16, 615768
SUOX	213.1	100	100	Sulfite oxidase deficiency, 272300
SYNE1	139.1	99	99	Emery-Dreifuss muscular dystrophy 4, autosomal dominant, 612998 Spinocerebellar ataxia, autosomal recessive 8, 610743
TAF1	113.3	99	96	Dystonia-Parkinsonism, X-linked, 314250 Mental retardation, X-linked, syndromic 33, 300966
TANGO2	145.6	100	100	Metabolic encephalomyopathic crises, recurrent, with rhabdomyolysis, cardiac arrhythmias and neurodegeneration, 616878
TBC1D20	146.1	94	94	Warburg micro syndrome 4, 615663
TDP1	123	98	95	Spinocerebellar ataxia, autosomal recessive with axonal neuropathy, 607250
TDP2	156.8	99	98	Spinocerebellar ataxia, autosomal recessive, 616949
TECPR2	161.3	100	99	Spastic paraplegia 49, autosomal recessive, 615031
TENM4	161	99	99	Tremor, hereditary essential, 5, 616736
TGM6	149.9	99	98	Spinocerebellar ataxia 35, 613908
TH	68.4	97	88	Segawa syndrome, recessive, 605407
THAP1	121.7	100	100	Dystonia 6, torsion, 602629
TIMM8A	46.2	94	78	Jensen syndrome, 311150 Mohr-Tranebjærg syndrome, 304700
TMEM240	112.7	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	185.2	100	99	Dystonia-1, torsion, 128100 {Dystonia-1, modifier of}
TPP1	146.4	100	100	Ceroid lipofuscinosis, neuronal, 2, 204500 Spinocerebellar ataxia, autosomal recessive 7, 609270
TREM2	148.9	99	99	Nasu-Hakola disease, 221770
TREX1	242.7	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	83.1	95	92	Pontocerebellar hypoplasia type 2A, 277470 Pontocerebellar hypoplasia type 4, 225753 ?Pontocerebellar hypoplasia type 5, 610204
TTBK2	148.5	100	99	Spinocerebellar ataxia 11, 604432
TTC19	93.2	85	79	Mitochondrial complex III deficiency, nuclear type 2, 615157
TTPA	101.2	83	76	Ataxia with isolated vitamin E deficiency, 277460
TUBA1A	113.3	99	97	Lissencephaly 3, 611603
TUBB4A	121.5	96	95	Dystonia 4, torsion, autosomal dominant, 128101 Leukodystrophy, hypomyelinating, 6, 612438
TUBG1	164.6	100	100	Cortical dysplasia, complex, with other brain malformations 4, 615412
TYROBP	95.2	100	99	Nasu-Hakola disease, 221770
VAMP1	131.6	100	100	Spastic ataxia 1, autosomal dominant, 108600
VCP	144.9	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	201.2	99	99	Cerebellar hypoplasia and mental retardation with or without quadrupedal locomotion 1, 224050
VPS13A	69.4	95	85	Choreoacanthocytosis, 200150

VPS13D	158	99	99	No OMIM phenotype
VPS37A	72.6	87	68	Spastic paraplegia 53, autosomal recessive, 614898
VRK1	124.8	97	94	Pontocerebellar hypoplasia type 1A, 607596
WDR45	75.6	97	90	Neurodegeneration with brain iron accumulation 5, 300894
WDR81	163.7	99	99	Cerebellar ataxia, mental retardation, and dysequilibrium syndrome 2, 610185
WWOX	131.1	100	99	Epileptic encephalopathy, early infantile, 28, 616211 Esophageal squamous cell carcinoma, somatic, 133239 Spinocerebellar ataxia, autosomal recessive 12, 614322
XK	97.4	99	99	McLeod syndrome with or without chronic granulomatous disease, 300842
XPR1	131.6	100	99	Basal ganglia calcification, idiopathic, 6, 616413
XRCC1	106.8	99	97	?Spinocerebellar ataxia, autosomal recessive 26, 617633
ZC4H2	76.5	99	97	Wieacker-Wolff syndrome, 314580
ZFYVE26	120.3	100	99	Spastic paraplegia 15, autosomal recessive, 270700
ZFYVE27	118.7	100	100	Spastic paraplegia 33, autosomal dominant, 610244
ZNF592	150.3	100	99	Spinocerebellar ataxia, autosomal recessive 5, 251300

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

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 14th, 2017.

This list is accurate for panel version DG 2.11

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