

NEUROPATHIES GENE PANEL DG 2.15 (119 genes)

Releasedate: 31-01-2019

<i>Gene</i>	<i>Median Coverage</i>	<i>% covered > 10x</i>	<i>% covered > 20x</i>	<i>Associated Phenotype description and OMIM disease ID</i>
AARS	124.3	100	99.6	Charcot-Marie-Tooth disease, axonal, type 2N, 613287 Epileptic encephalopathy, early infantile, 29, 616339
ABCD1	76	74.7	68	Adrenoleukodystrophy, 300100 Adrenomyeloneuropathy, adult, 300100
ABHD12	107	97.3	88	Polyneuropathy, hearing loss, ataxia, retinitis pigmentosa, and cataract, 612674
AIFM1	106.2	100	99.7	Combined oxidative phosphorylation deficiency 6, 300816 Cowchock syndrome, 310490 Deafness, X-linked 5, 300614
ARHGEF10	132.8	99.8	98	?Slowed nerve conduction velocity, AD, 608236
ARSA	97.8	100	99.7	Metachromatic leukodystrophy, 250100
ATAD3A	87.9	89	86.2	Harel-Yoon syndrome, 617183
ATL1	161	99.7	97.9	Neuropathy, hereditary sensory, type ID, 613708 Spastic paraplegia 3A, autosomal dominant, 182600
ATL3	125	98.1	93.8	Neuropathy, hereditary sensory, type IF, 615632
ATP7A	133.2	99.7	97.8	Menkes disease, 309400 Occipital horn syndrome, 304150 Spinal muscular atrophy, distal, X-linked 3, 300489
BICD2	158.6	100	99.9	Spinal muscular atrophy, lower extremity-predominant, 2, AD, 615290
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
CCT5	164.5	99.9	99.1	Neuropathy, hereditary sensory, with spastic paraplegia, 256840
CNTNAP1	161.2	99.2	97.5	Hypomyelinating neuropathy, congenital, 3, 618186 Lethal congenital contracture syndrome 7, 616286

COX6A1	180.6	100	99.4	Charcot-Marie-Tooth disease, recessive intermediate D, 616039
CTDP1	105	86.6	83.6	Congenital cataracts, facial dysmorphism, and neuropathy, 604168
CYP27A1	175.1	98.3	96.1	Cerebrotendinous xanthomatosis, 213700
DCAF8	124.3	100	99.9	?Giant axonal neuropathy 2, autosomal dominant, 610100
DCTN1	131.6	99.7	98.3	Neuropathy, distal hereditary motor, type VIIB, 607641 Perry syndrome, 168605 {Amyotrophic lateral sclerosis, susceptibility to}, 105400
DHTKD1	141	99.6	98.2	2-aminoadipic 2-oxoadipic aciduria, 204750 ?Charcot-Marie-Tooth disease, axonal, type 2Q, 615025
DNAJB2	102	100	100	Spinal muscular atrophy, distal, autosomal recessive, 5, 614881
DNM2	127.4	97.5	94.4	Centronuclear myopathy 1, 160150 Charcot-Marie-Tooth disease, axonal type 2M, 606482 Charcot-Marie-Tooth disease, dominant intermediate B, 606482 Lethal congenital contracture syndrome 5, 615368
DNMT1	113.4	99.2	98.3	Cerebellar ataxia, deafness, and narcolepsy, autosomal dominant, 604121 Neuropathy, hereditary sensory, type IE, 614116
DST	154.1	99.7	98.2	?Neuropathy, hereditary sensory and autonomic, type VI, 614653 Epidermolysis bullosa simplex, autosomal recessive 2, 615425
DYNC1H1	179.8	100	99.6	Charcot-Marie-Tooth disease, axonal, type 20, 614228 Mental retardation, autosomal dominant 13, 614563 Spinal muscular atrophy, lower extremity-predominant 1, AD, 158600
EGR2	124.4	100	100	Charcot-Marie-Tooth disease, type 1D, 607678 Dejerine-Sottas disease, 145900 Hypomyelinating neuropathy, congenital, 1, 605253
ELP1	142.6	99.7	98.1	Dysautonomia, familial, 223900
EXOSC8	80.1	91.8	76.9	Pontocerebellar hypoplasia, type 1C, 616081
FBLN5	119.6	91.8	91.1	Cutis laxa, autosomal dominant 2, 614434 Cutis laxa, autosomal recessive, type IA, 219100 Macular degeneration, age-related, 3, 608895 Neuropathy, hereditary, with or without age-related macular degeneration, 608895
FGD4	111.9	99.3	97.3	Charcot-Marie-Tooth disease, type 4H, 609311

FIG4	154.9	99.8	98.4	?Polymicrogyria, bilateral temporooccipital, 612691 Amyotrophic lateral sclerosis 11, 612577 Charcot-Marie-Tooth disease, type 4J, 611228 Yunis-Varon syndrome, 216340
FLVCR1	139.5	99.2	95.8	Ataxia, posterior column, with retinitis pigmentosa, 609033
GALC	100.6	98.9	94.6	Krabbe disease, 245200
GAN	190	100	99.9	Giant axonal neuropathy-1, 256850
GARS	125.7	99.9	98.5	Charcot-Marie-Tooth disease, type 2D, 601472 Neuropathy, distal hereditary motor, type VA, 600794
GDAP1	163.1	99.3	96.1	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
GJB1	229.8	100	99.8	Charcot-Marie-Tooth neuropathy, X-linked dominant, 1, 302800
GJB3	308.9	100	100	Deafness, autosomal dominant 2B, 612644 Deafness, autosomal dominant, with peripheral neuropathy, 0 Deafness, autosomal recessive, 0 Deafness, digenic, GJB2/GJB3, 220290 Erythrokeratoderma variabilis et progressiva 1, 133200
GLA	81.3	99.7	97.6	Fabry disease, 301500 Fabry disease, cardiac variant, 301500
GNB4	152.6	100	99.7	Charcot-Marie-Tooth disease, dominant intermediate F, 615185
HARS	159.4	100	100	Charcot-Marie-Tooth disease, axonal, type 2W, 616625 Usher syndrome type 3B, 614504
HINT1	60	98.5	88	Neuromyotonia and axonal neuropathy, autosomal recessive, 137200
HK1	143.4	100	99.9	Hemolytic anemia due to hexokinase deficiency, 235700 Neuropathy, hereditary motor and sensory, Russe type, 605285 Retinitis pigmentosa 79, 617460
HMBS	109	100	99.8	Porphyria, acute intermittent, 176000 Porphyria, acute intermittent, nonerythroid variant, 176000
HOXD10	137.7	100	99.8	Charcot-Marie-Tooth disease, foot deformity of, 192950 Vertical talus, congenital, 192950

HSPB1	39.7	93.7	81.8	Charcot-Marie-Tooth disease, axonal, type 2F, 606595 Neuropathy, distal hereditary motor, type IIB, 608634
HSPB3	290.9	100	100	?Neuronopathy, distal hereditary motor, type IIC, 613376
HSPB8	156.4	100	100	Charcot-Marie-Tooth disease, axonal, type 2L, 608673 Neuropathy, distal hereditary motor, type IIA, 158590
IFRD1	133.9	98.7	96	No OMIM phenotype (Lin et al.J. Hum. Genet. 2018Echeveste et al. Parkinsonism Relat Disord. 2017Brkanac et al.Am J Hum Genet. 2009)
IGHMBP2	107.8	99.3	96	Charcot-Marie-Tooth disease, axonal, type 2S, 616155 Neuronopathy, distal hereditary motor, type VI, 604320
INF2	79.2	84.1	81.1	Charcot-Marie-Tooth disease, dominant intermediate E, 614455 Glomerulosclerosis, focal segmental, 5, 613237
KARS	122.6	100	99.3	?Charcot-Marie-Tooth disease, recessive intermediate, B, 613641 Deafness, autosomal recessive 89, 613916
KIF1A	114	99.2	96.1	Mental retardation, autosomal dominant 9, 614255 Neuropathy, hereditary sensory, type IIC, 614213 Spastic paraplegia 30, autosomal recessive, 610357
KIF1B	154.8	100	99.5	?Charcot-Marie-Tooth disease, type 2A1, 118210 Pheochromocytoma, 171300 {Neuroblastoma, susceptibility to, 1}, 256700
KIF5A	136.1	100	99.9	Myoclonus, intractable, neonatal, 617235 Spastic paraplegia 10, autosomal dominant, 604187 {Amyotrophic lateral sclerosis, susceptibility to, 25}, 617921
LITAF	126.3	94.8	91.5	Charcot-Marie-Tooth disease, type 1C, 601098

LMNA	89.2	97.9	91.3	<p>Cardiomyopathy, dilated, 1A, 115200</p> <p>Charcot-Marie-Tooth disease, type 2B1, 605588</p> <p>Emery-Dreifuss muscular dystrophy 2, autosomal dominant, 181350</p> <p>Emery-Dreifuss muscular dystrophy 3, autosomal recessive, 616516</p> <p>Heart-hand syndrome, Slovenian type, 610140</p> <p>Hutchinson-Gilford progeria, 176670</p> <p>Lipodystrophy, familial partial, type 2, 151660</p> <p>Malouf syndrome, 212112</p> <p>Mandibuloacral dysplasia, 248370</p> <p>Muscular dystrophy, congenital, 613205</p> <p>Restrictive dermopathy, lethal, 275210</p>
LRSAM1	130.4	100	99.7	Charcot-Marie-Tooth disease, axonal, type 2P, 614436
MARS	125.2	99.7	97.3	<p>Charcot-Marie-Tooth disease, axonal, type 2U, 616280</p> <p>Interstitial lung and liver disease, 615486</p>
MED25	103.9	99.1	95.7	<p>?Charcot-Marie-Tooth disease, type 2B2, 605589</p> <p>Basel-Vanagait-Smirin-Yosef syndrome, 616449</p>
MFN2	150.6	100	99.9	<p>Charcot-Marie-Tooth disease, axonal, type 2A2A, 609260</p> <p>Charcot-Marie-Tooth disease, axonal, type 2A2B, 617087</p> <p>Hereditary motor and sensory neuropathy VIA, 601152</p>
MME	101.1	98.2	93.6	<p>?Spinocerebellar ataxia 43, 617018</p> <p>Charcot-Marie-Tooth disease, axonal, type 2T, 617017</p>
MORC2	135.9	100	99.7	Charcot-Marie-Tooth disease, axonal, type 2Z, 616688
MPZ	123.4	100	99.3	<p>Charcot-Marie-Tooth disease, dominant intermediate D, 607791</p> <p>Charcot-Marie-Tooth disease, type 1B, 118200</p> <p>Charcot-Marie-Tooth disease, type 2I, 607677</p> <p>Charcot-Marie-Tooth disease, type 2J, 607736</p> <p>Dejerine-Sottas disease, 145900</p> <p>Hypomyelinating neuropathy, congenital, 2, 618184</p> <p>Roussy-Levy syndrome, 180800</p>
MTMR2	106.6	100	99.2	Charcot-Marie-Tooth disease, type 4B1, 601382
MYH14	102	97.7	91.5	<p>?Peripheral neuropathy, myopathy, hoarseness, and hearing loss, 614369</p> <p>Deafness, autosomal dominant 4A, 600652</p>
NAGLU	108.7	92.4	90.4	<p>?Charcot-Marie-Tooth disease, axonal, type 2V, 616491</p> <p>Mucopolysaccharidosis type IIIB (Sanfilippo B), 252920</p>

NDRG1	128.4	99.9	98.8	Charcot-Marie-Tooth disease, type 4D, 601455
NEFH	111.6	96.6	87.5	?{Amyotrophic lateral sclerosis, susceptibility to}, 105400 Charcot-Marie-Tooth disease, axonal, type 2CC, 616924
NEFL	164.6	99.7	98.1	Charcot-Marie-Tooth disease, dominant intermediate G, 617882 Charcot-Marie-Tooth disease, type 1F, 607734 Charcot-Marie-Tooth disease, type 2E, 607684
NGF	257.6	100	100	Neuropathy, hereditary sensory and autonomic, type V, 608654
NTRK1	130.6	99.7	97.7	Insensitivity to pain, congenital, with anhidrosis, 256800 Medullary thyroid carcinoma, familial, 155240
PDK3	105.1	96.4	94.3	?Charcot-Marie-Tooth disease, X-linked dominant, 6, 300905
PEX1	115.8	97.7	95.4	Heimler syndrome 1, 234580 Peroxisome biogenesis disorder 1A (Zellweger), 214100 Peroxisome biogenesis disorder 1B (NALD/IRD), 601539
PEX7	113.5	89.6	82	Peroxisome biogenesis disorder 9B, 614879 Rhizomelic chondrodysplasia punctata, type 1, 215100
PHYH	74.6	97.5	90.8	Refsum disease, 266500
PLA2G6	117.5	99.9	98.4	Infantile neuroaxonal dystrophy 1, 256600 Neurodegeneration with brain iron accumulation 2B, 610217 Parkinson disease 14, autosomal recessive, 612953
PLEKHG5	86.9	96.2	89	Charcot-Marie-Tooth disease, recessive intermediate C, 615376 Spinal muscular atrophy, distal, autosomal recessive, 4, 611067
PMM2	141.1	99.9	99.4	Congenital disorder of glycosylation, type Ia, 212065
PMP22	111.2	96.7	91.9	?Neuropathy, inflammatory demyelinating, 139393 Charcot-Marie-Tooth disease, type 1A, 118220 Charcot-Marie-Tooth disease, type 1E, 118300 Dejerine-Sottas disease, 145900 Neuropathy, recurrent, with pressure palsies, 162500 Roussy-Levy syndrome, 180800
PRDM12	112.2	91	87.7	Neuropathy, hereditary sensory and autonomic, type VIII, 616488

PRPS1	149.5	100	100	Arts syndrome, 301835 Charcot-Marie-Tooth disease, X-linked recessive, 5, 311070 Deafness, X-linked 1, 304500 Gout, PRPS-related, 300661 Phosphoribosylpyrophosphate synthetase superactivity, 300661
PRX	115.6	99.8	98.3	Charcot-Marie-Tooth disease, type 4F, 614895 Dejerine-Sottas disease, 145900
PSAP	114.4	99.9	99	Combined SAP deficiency, 611721 Gaucher disease, atypical, 610539 Krabbe disease, atypical, 611722 Metachromatic leukodystrophy due to SAP-b deficiency, 249900
RAB7A	157.9	100	100	Charcot-Marie-Tooth disease, type 2B, 600882
REEP1	78.3	76.3	75.7	?Neuronopathy, distal hereditary motor, type VB, 614751 Spastic paraplegia 31, autosomal dominant, 610250
RETREG1	126.1	95.6	90.1	Neuropathy, hereditary sensory and autonomic, type IIB, 613115
SBF1	107.9	98.5	96.5	Charcot-Marie-Tooth disease, type 4B3, 615284
SBF2	117	99.6	96.8	Charcot-Marie-Tooth disease, type 4B2, 604563
SCN10A	165.3	100	99.5	Episodic pain syndrome, familial, 2, 615551
SCN11A	138.1	99.2	97.6	Episodic pain syndrome, familial, 3, 615552 Neuropathy, hereditary sensory and autonomic, type VII, 615548
SCN9A	146.5	98.5	97	Epilepsy, generalized, with febrile seizures plus, type 7, 613863 Erythralgia, primary, 133020 Febrile seizures, familial, 3B, 613863 HSAN2D, autosomal recessive, 243000 Insensitivity to pain, congenital, 243000 Paroxysmal extreme pain disorder,, 167400 Small fiber neuropathy, 133020 {Dravet syndrome, modifier of}, 607208
SCO2	113.1	100	100	Cardioencephalomyopathy, fatal infantile, due to cytochrome c oxidase deficiency 1, 604377 Myopia 6, 608908
SEPT9	118.7	99.7	96.8	Amyotrophy, hereditary neuralgic, 162100 Leukemia, acute myeloid, therapy-related, 0 Ovarian carcinoma, 0

SETX	163.2	99.9	99.1	Amyotrophic lateral sclerosis 4, juvenile, 602433 Spinocerebellar ataxia, autosomal recessive 1, 606002
SH3TC2	121.3	100	99.7	Charcot-Marie-Tooth disease, type 4C, 601596 Mononeuropathy of the median nerve, mild, 613353
SLC12A6	141.8	100	99.9	Agenesis of the corpus callosum with peripheral neuropathy, 218000
SLC25A46	205.7	95.9	87.3	Neuropathy, hereditary motor and sensory, type VIB, 616505
SLC52A2	177.6	100	100	Brown-Vialetto-Van Laere syndrome 2, 614707
SLC52A3	119.6	100	100	?Fazio-Londe disease, 211500 Brown-Vialetto-Van Laere syndrome 1, 211530
SLC5A7	117.1	100	99.9	Myasthenic syndrome, congenital, 20, presynaptic, 617143 Neuronopathy, distal hereditary motor, type VIIA, 158580
SOX10	65.8	98.2	91.3	PCWH syndrome, 609136 Waardenburg syndrome, type 2E, with or without neurologic involvement, 611584 Waardenburg syndrome, type 4C, 613266
SPG11	129.2	99.2	96.9	Amyotrophic lateral sclerosis 5, juvenile, 602099 Charcot-Marie-Tooth disease, axonal, type 2X, 616668 Spastic paraplegia 11, autosomal recessive, 604360
SPTBN4	83.9	96.6	89.1	Neurodevelopmental disorder with hypotonia, neuropathy, and deafness, 617519
SPTLC1	115.5	99	93.9	Neuropathy, hereditary sensory and autonomic, type IA, 162400
SPTLC2	160.2	100	100	Neuropathy, hereditary sensory and autonomic, type IC, 613640
SURF1	96.2	88.3	88.3	Charcot-Marie-Tooth disease, type 4K, 616684 Leigh syndrome, due to COX IV deficiency, 256000
SYT2	101.3	100	99	Myasthenic syndrome, congenital, 7, presynaptic, 616040
TDP1	122.9	98.7	95.3	Spinocerebellar ataxia, autosomal recessive with axonal neuropathy, 607250
TFG	121.5	93.9	90.7	?Spastic paraplegia 57, autosomal recessive, 615658 Hereditary motor and sensory neuropathy, Okinawa type, 604484
TRIM2	157.7	93.6	91.4	Charcot-Marie-Tooth disease, type 2R, 615490

TRPV4	172.4	99.5	98.7	?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 Scapuloperoneal spinal muscular atrophy, 181405 SED, Maroteaux type, 184095 Spinal muscular atrophy, distal, congenital nonprogressive, 600175 Spondylometaphyseal dysplasia, Kozlowski type, 184252 [Sodium serum level QTL 1], 613508
TTR	152.3	94.6	94.6	Amyloidosis, hereditary, transthyretin-related, 105210 Carpal tunnel syndrome, familial, 115430 [Dystransthyretinemic hyperthyroxinemia], 145680
TUBB3	136.1	98.1	96.9	Cortical dysplasia, complex, with other brain malformations 1, 614039 Fibrosis of extraocular muscles, congenital, 3A, 600638
TWINK	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
VCP	144.8	99.9	99.5	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
WARS	123.8	99.7	98	Neuronopathy, distal hereditary motor, type IX, 617721
WNK1	167.7	99.9	99.5	Neuropathy, hereditary sensory and autonomic, type II, 201300 Pseudohypoaldosteronism, type IIC, 614492
YARS	122.4	100	100	Charcot-Marie-Tooth disease, dominant intermediate C, 608323

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 : December 31st, 2018.

This list is accurate for panel version DG 2.15

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