

Supplementary table 1. CMT genes: inheritance, phenotype and cell pathophysiology.

Gene	OMIM (*)	MoI	Phenotype				OMIM (#)	Localization		Protein function / Pathway	Disrupted process
			Neuropathy type					Schwann cell	Body / axon		
			A	D	I						
Other genes											
<i>AARS1</i>	601065	AD	✓			CMT2N	613287		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing
<i>ABHD12</i>	613599	AR		✓		Polyneuropathy	612674	NA	NA	Lipid hydrolysis	ER and Golgi
<i>AIFM1</i>	300169	XL	✓			CMTX4	310490		✓	Oxidative phosphorylation - apoptosis	Mitochondria
<i>ARHGEF10</i>	608136	AD		✓		Slowed nerve conduction velocities	608236	✓		Guanine nucleotide exchange factor	Endosomal sorting and cell signaling
<i>ATPIA1</i>	182310	AD	✓			CMT2DD	618036		✓	Cation transport ATPase	Channel
<i>ATP7A</i>	300011	XL	✓			Spinal muscular atrophy/distal motor neuropathy Menkes disease	300489 309400		✓	Copper transport ATPase	Channel
<i>BAG3</i>	603883	AD	✓			Myopathy, myofibrillar, 6	612954	NA	NA	Co-chaperone	Apoptosis
<i>BSCL2</i>	606158	AD	✓			dHMN type VC	619112		✓	Formation of lipid droplets	ER and Golgi
<i>CADM3</i>	609743	AD	✓			CMT2FF	619519	NA	NA	Calcium-independent cell-cell adhesion	Cell-cell interaction
<i>CNTNAP1</i>	602346	AR	✓	✓		Hypomyelinating neuropathy, congenital, 3	618186	NA	NA	Signaling between axons and myelinating glial cells	Axon guidance
<i>COA7</i>	615623	AR	✓			Spinocerebellar ataxia, AR, with axonal neuropathy 3	618387	NA	NA	Assembly of mitochondrial oxidative phosphorylation complexes	NA
<i>DCTN1</i>	601143	AD				dHMN type VIIB	607641		✓	Dynein-mediated retrograde transport of vesicles and organelles along microtubules	Cytoskeleton, axonal transport
<i>DCTN2</i>	607376	AD	✓			NA	NA	NA	NA	Modulation of dynein-organelles binding	Cytoskeleton
<i>DGAT2</i>	606983	AD	✓			NA	NA		✓	Synthesis of triglycerides	ER and Golgi
<i>DHTKD1</i>	614984	AD	✓			CMT2Q	615025		✓	Degradation of amino acids	Mitochondria
<i>DNAJB2</i>	604139	AR	✓			Spinal muscular atrophy, distal, AR, 5	614881		✓	Co-chaperone	Proteasome and protein aggregation

<i>DNMT1</i>	126375	AD	✓			dHSN type IE	614116		✓	Maintenance of methylation patterns of genomic DNA	Nuclear envelope, mRNA processing
<i>DNM2</i>	602378	AD			✓	CMT2B CMT2M Centronuclear myopathy 1	606482 606482 160150	✓		GTPase, fission-fusion apparatus	Endosomal sorting and cell signaling
<i>DRP2</i>	300052	XL			✓	NA	NA	✓		Maintenance of membrane-associated complexes	Cytoskeleton
<i>DYNC1H1</i>	600112	AD	✓			CMT20	614228		✓	Retrograde axonal transport	Cytoskeleton, axonal transport
<i>EGR2</i>	129010	AD AR		✓		CMT1D	607678	✓		Transcription regulator	Transcription, mRNA processing
<i>FGD4</i>	611104	AR		✓		CMT4H	609311	✓		Regulates cell signaling involved in myelin production and actin cytoskeleton	Cytoskeleton
<i>FIG4</i>	609390	AR		✓		CMT4J	611228	✓		Abnormal transport of intracellular organelles	Endosomal sorting and cell signaling
<i>GARS1</i>	600287	AD	✓			CMT2D dHMN type VA	601472 600794		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing
<i>GNB4</i>	610863	AD			✓	CMTDIF	615185		✓	Signal transduction	Endosomal sorting and cell signaling
<i>HARS1</i>	142810	AD	✓	✓		CMT2W	616625		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing
<i>HSPB1</i>	602195	AD	✓			CMT2F dHMN type IIB	606595 608634		✓	Microtubule regulator	Proteasome and protein aggregation
<i>HSPB3</i>	604624	AD	✓			dHMN type IIC	613376		✓	Microtubule regulator	Proteasome and protein aggregation
<i>HSPB8</i>	608014	AD	✓			CMT2L dHMN type IIA	608673 158590		✓	Microtubule regulator	Proteasome and protein aggregation
<i>IGHMBP2</i>	600502	AR	✓			CMT2S dHMN type VI	616155 604320		✓	Helicase	Nuclear envelope, mRNA processing
<i>INF2</i>	610982	AD			✓	CMTDIE	614455	✓		Actin polymerization and filament severing	Cytoskeleton
<i>KARS</i>	609574	NA				NA	NA		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing
<i>KIF1B</i>	605995	AD	✓			CMT2A1	118210		✓	Intracellular organelle transport	Cytoskeleton, axonal transport

<i>KIF5A</i>	602821	AD	✓			Spastic paraplegia 10	604187		✓	Intracellular organelle transport	Cytoskeleton, axonal transport
<i>LITAF</i>	603795	AD			✓	CMT1C	603795	✓		Regulation of endosome to lysosome trafficking and cell signaling	Endosomal sorting and cell signaling
<i>LMNA</i>	150330	AR	✓			CMT2B1	605588		✓	Intermediate filament protein of nuclear envelope	Nuclear envelope, mRNA processing
<i>LRSAMI</i>	610933	AD AR	✓			CMT2P	614436		✓	E3 ubiquitin ligase, regulates cell adhesion molecules	Proteasome and protein aggregation
<i>MARS1</i>	156560	AD	✓			CMT2U	616280		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing
<i>MCM3AP</i>	603294	AR	✓	✓		Peripheral neuropathy, autosomal recessive, with or without impaired intellectual development	618124		✓	DNA replication	
<i>MME</i>	120520	AR AD	✓			CMT2T	120520		✓	Destruction of opioid peptides	Endosomal sorting and cell signaling
<i>MORC2</i>	616661	AD	✓			CMT2Z	616688		✓	Regulation of heterochromatin condensation	Nuclear envelope, mRNA processing
<i>MPV17</i>	137960	AR	✓			CMT2EE	137960		✓	Metabolism of reactive oxygen species	Mitochondria
<i>MTMR2</i>	603557	AR			✓	CMT4B1	603557	✓		Modification of chemical messengers involved in signal transduction	Endosomal sorting and cell signaling
<i>NAGLU</i>	609701	AD	✓			CMT2V	616491		✓	Degradation of heparan sulfate	Endosomal sorting and cell signaling
<i>NDRG1</i>	605262	AR			✓	CMT4D	605262	✓		Signaling protein shuttling between cytoplasm and nucleus	Endosomal sorting and cell signaling
<i>NEFH</i>	162230	AD	✓			CMT2CC	616924		✓	Growth arrest and cell differentiation	Cytoskeleton
<i>NEFL</i>	162280	AD AR	✓	✓		CMTDIG CMT1F CMT2E	617882 607734 607684		✓	Intermediate filament in neurons	Cytoskeleton, axonal transport
<i>PDK3</i>	300906	XL	✓			CMTX6	300905		✓	Regulates pyruvate dehydrogenase complex	Mitochondria
<i>PLEKHG5</i>	611101	AR			✓	CMTRIC	615376		✓	Nuclear factor kB-activator	Nuclear envelope, mRNA processing
<i>PMP2</i>	170715	AD			✓	CMT1G	618279	✓		Myelin assembly	Myelin assembly

<i>PNKP</i>	605610	AR	✓			CMT2B2	605589		✓	DNA repair	Nuclear envelope, mRNA processing
<i>PRPS1</i>	311850	XL				CMTX5	311070		✓	Purine and pyrimidine biosynthesis	Nuclear envelope, mRNA processing
<i>PRX</i>	605725	AR	✓			CMT4F	614895	✓		Membrane-protein interaction stabilizing myelin sheath	Cytoskeleton
<i>PTRH2</i>	608625	AR				Infantile-onset multisystem neurologic, endocrine, and pancreatic disease	616263	NA	NA	Peptidyl-tRNA hydrolase	Proteasome and protein metabolism
<i>RAB7A</i>	602298	AD	✓			CMT2B	600882		✓	Vesicular transport and membrane trafficking	Endosomal sorting and cell signaling
<i>SBF1</i>	603560	AR		✓		CMT4B3	615284	✓		Endosomal trafficking	Endosomal sorting and cell signaling
<i>SBF2</i>	607697	AR		✓		CMT4B2	604563	✓		Development of Schwann cells	Endosomal sorting and cell signaling
<i>SCO2</i>	604272	AR	✓			Motor neuropathy ^[7]	NA	NA	NA	Cytochrome c oxidase synthesis	Gene expression
<i>SETX</i>	608465	AD	✓			Amyotrophic lateral sclerosis 4, juvenile Spinocerebellar ataxia, AR, with axonal neuropathy 2	602433 606002		✓	Probable RNA/DNA helicase	Nuclear envelope, mRNA processing
<i>SIGMAR1</i>	601978	AR	✓			Amyotrophic lateral sclerosis 16, juvenile Spinal muscular atrophy, distal, AR, 2	614373 605726		✓	Chaperone	ER and Golgi
<i>SGPL1</i>	603729	AR	✓			Motoneuropathy	NA		✓	Sphingolipid catabolism	ER and Golgi
<i>SPG11</i>	610844	AR	✓			CMT2X	616668		✓	Cytoskeleton stability and synaptic vesicle transport	Cytoskeleton, axonal transport
<i>SPTLC1</i>	605712	AD	✓			Neuropathy, hereditary sensory and autonomic, type IA	162400		✓	Serine palmitoyltransferase	Toxin
<i>TRIM2</i>	614141	AR	✓			CMT2R	614141		✓	E3 ubiquitin ligase	Proteasome and protein aggregation
<i>TRPV4</i>	605427	AD	✓			CMT2C dHMN type VIII	606071 600175		✓	Calcium homeostasis, cytoskeleton remodeling	Channel
<i>VCP</i>	611745	AD	✓			NA	NA		✓	AAA ATPase family	ER and Golgi
<i>VWAI</i>	611901	AR	✓			dHMN	619216	NA	NA	Matrix assembly	

WARS	191050	AD	✓			dHMN type IX	617721		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing
YARSI	603623	AD			✓	CMTDIC	608323		✓	Aminoacyl tRNA synthetase	Nuclear envelope, mRNA processing

MoI: Mode of Inheritance; A: axonal; D: demyelinating; I: intermediate; AR: autosomal recessive; AD: autosomal dominant; XL: X-linked; CMT: Charcot-Marie-Tooth; dHMN: distal hereditary motor neuropathy; dHSN: distal hereditary sensory neuropathy DI: dominant intermediate; NA: data not available. Information about genes and protein functions was extracted from databases such as OMIM (<https://www.omim.org/>), GeneCards (<https://www.genecards.org/>), Reactome (<https://reactome.org/>) and UniProt (<https://www.uniprot.org/>).