

The Genetic Nomenclature of Recessive Cerebellar Ataxias

Malco Rossi, MD, PhD ¹ Mathieu Anheim, MD, PhD,^{2,3,4} Alexandra Durr, MD, PhD,^{5,6} Christine Klein, MD,^{7,8} Michel Koenig, MD, PhD,⁹ Matthis Synofzik, MD,^{10,11} Connie Marras, MD, PhD,¹² and Bart P. van de Warrenburg, MD, PhD,^{13*} on behalf of the International Parkinson and Movement Disorder Society Task Force on Classification and Nomenclature of Genetic Movement Disorders

¹Movement Disorders Section, Neuroscience Department, Raul Carrea Institute for Neurological Research, Buenos Aires, Argentina

²Département de Neurologie, Hôpitaux Universitaires de Strasbourg, Hôpital de Hautepierre, Strasbourg, France

³Institut de Génétique et de Biologie Moléculaire et Cellulaire, INSERM-U964/CNRS-UMR7104/Université de Strasbourg, Illkirch, France

⁴Fédération de Médecine Translationnelle de Strasbourg, Université de Strasbourg, Strasbourg, France

⁵Brain and Spine Institute, Sorbonne Université, Inserm U1127, CNRS UMR 7225, Pitié-Salpêtrière University Hospital, Paris, France

⁶Department of Genetics, AP-HP, Pitié-Salpêtrière University Hospital, 7501 Paris, France

⁷Institute of Neurogenetics, University of Luebeck, Luebeck, Germany

⁸Department of Neurology, University Hospital Schleswig-Holstein, Campus Lübeck, Germany

⁹Laboratoire de Génétique de Maladies Rares, EA7402, Institut Universitaire de Recherche Clinique, Université de Montpellier, CHU Montpellier, Montpellier, France

¹⁰Department of Neurodegenerative Diseases, Hertie Institute for Clinical Brain Research, University of Tübingen, Tübingen, Germany

¹¹German Center for Neurodegenerative Diseases, Tübingen, Germany

¹²Toronto Western Hospital Morton, Gloria Shulman Movement Disorders Centre, and the Edmond J. Safra Program in Parkinson's Disease, University of Toronto, Toronto, Canada

¹³Department of Neurology, Donders Institute for Brain, Cognition & Behaviour, Radboud University Medical Center, Nijmegen, The Netherlands

ABSTRACT: The recessive cerebellar ataxias are a large group of degenerative and metabolic disorders, the diagnostic management of which is difficult because of the enormous clinical and genetic heterogeneity. Because of several limitations, the current classification systems provide insufficient guidance for clinicians and researchers. Here, we propose a new nomenclature for the genetically confirmed recessive cerebellar ataxias according to the principles and criteria laid down by the International Parkinson and Movement Disorder Society Task Force on Classification and Nomenclature of Genetic Movement Disorders. We apply stringent criteria for considering an association between gene and phenotype to be established. The newly proposed list of recessively inherited cerebellar ataxias includes 62 disorders that were assigned an ATX prefix, followed by the gene name, because these typically present with ataxia as a predominant and/or

consistent feature. An additional 30 disorders that often combine ataxia with a predominant or consistent other movement disorder received a double prefix (e.g., ATX/HSP). We also identified a group of 89 entities that usually present with complex nonataxia phenotypes, but may occasionally present with cerebellar ataxia. These are listed separately without the ATX prefix. This new, transparent and adaptable nomenclature of the recessive cerebellar ataxias will facilitate the clinical recognition of recessive ataxias, guide diagnostic testing in ataxia patients, and help in interpreting genetic findings. © 2018 International Parkinson and Movement Disorder Society

Key Words: genetics; recessive ataxias; movement disorders; nomenclature

*Corresponding author: Dr. Bart van de Warrenburg, Expert Centre for Genetic Movement Disorders, Department of Neurology, Donders Institute for Brain Cognition & Behaviour, Radboud University Medical Center, PO Box 9101, 6500 HB Nijmegen, The Netherlands; bart.vandewarrenburg@radboudumc.nl

Relevant conflicts of interests/financial disclosures: M.S. received speaker honoraria from Actelion Pharmaceuticals. M.A. received speaker honoraria and travel grants from Actelion Pharmaceuticals.

Received: 24 January 2018; **Revised:** 15 March 2018; **Accepted:** 25 March 2018

Published online 00 Month 2018 in Wiley Online Library (wileyonlinelibrary.com). DOI: 10.1002/mds.27415

© 2018 International Parkinson and Movement Disorder Society

The recessive cerebellar ataxias are a large group of clinically and genetically heterogeneous degenerative and metabolic disorders.¹ Although cerebellar ataxia is the shared clinical theme, the phenotype is often complex as other neurological features and sometimes systemic manifestations co-occur. The enormous heterogeneity complicates clinical management, particularly in the diagnostic phase. The currently used classification systems have several limitations and are therefore of limited use in terms of both providing guidance for clinicians in the diagnostic process and making understandable the field of recessive cerebellar ataxias as well as its boundaries. For example, the classification of autosomal recessive spinocerebellar ataxias's (SCAR) (for SCA-recessive) is far from complete, mainly consisting of relatively newer entities (the most common recessive ataxia, Friedreich ataxia, is not included) and also contains unconfirmed entities. Moreover, this SCAR classification is partly paralleled and duplicated, yet with different numbers, by another autosomal recessive cerebellar ataxia (ARCA) classification, the ARCA classification (for a recent criticism of these autosomal recessive ataxia nomenclatures, see the review by Synofzik and Schule²). These problems of existing incomplete, inconsistent, and partly parallel nomenclatures are not unique to recessive ataxias, as the classifications of other genetic movement disorders are similarly flawed.³ For this reason, the International Parkinson and Movement Disorder Society Task Force on Classification and Nomenclature of Genetic Movement Disorders was installed to take on the task of designing a uniform, adaptable, and phenotype-first nomenclature for genetic movement disorders. A certain genetic entity is included only after stringent, predefined criteria are met.⁴ Although the review process and the classification procedure incorporate clinical data, the proposed nomenclature does not constitute a new clinical classification of the recessive ataxias. The first paper of this task force included the proposed nomenclature for genetic parkinsonism, dystonia, autosomal dominant cerebellar ataxia, hereditary spastic paraplegia, paroxysmal movement disorder, neurodegeneration with brain iron accumulation, and primary familial brain calcification.⁴

Here we present the proposed nomenclature for the recessive cerebellar ataxias, based on the same principles, criteria, and recommendations as provided by the task force.

Search Strategy

The search aimed to have a fully comprehensive list of all autosomal and X-linked recessive diseases that have been reported to present with ataxia. The Online Mendelian Inheritance in Man database was searched

using the term “recessive ataxia.” Each disease that was identified was then checked in PubMed, GeneReviews, OrphaNet, and the Washington University in St. Louis neuromuscular section webpage (<http://neuromuscular.wustl.edu/ataxia/recatax.html>). To ensure that no diseases were missed, the same search strategy without time restrictions was applied in PubMed using the terms “recessive ataxia” or “cerebellar ataxia” and updated to August 2017. Only publications written in English were reviewed. Additional publications were identified by using the respective gene name as keyword, for example, SYNE1. A complete list of publications reviewed can be found in the supplementary material.

Review Process

In total, 355 recessive, genetic entities were identified in which ataxia had been reported. A total of 48 disorders for which a causative gene had not been identified were excluded from further analysis. We applied the following criteria that lend support to causality according to the U.S. National Human Genome Research Institute: (1) the presence of the variant in multiple, unrelated affected individuals; (2) evidence for segregation or statistical association of the variant with disease; (3) the variant should be conserved across different species; and (4) the variant should be predicted to alter the normal biochemical effect of a gene product as supported by functional evidence in human tissue, well-established cellular or animal models, or other biochemical or histological abnormalities, if possible. The members of the recessive ataxias working group (MA, AD, MK, MS and BW) each reviewed a subset of entities assigned to them. The review process consisted of selecting those diseases in which cerebellar ataxia was associated with mutations in the gene of interest in at least 2 independent reports. The review process was focused on cerebellar ataxias and not on purely sensory ataxias or those disorders with mainly cerebellar hypoplasia as a cardinal feature without the clinical presence of cerebellar ataxia. Congenital ataxias, Joubert syndrome, and primary metabolic diseases, such as peroxisomal and lysosomal disorders, were included if cerebellar ataxia was documented at any disease stage.

During the early stages of the process, it was decided to use the ataxia (ATX) prefix for both dominantly and recessively inherited ataxias. The first paper that had proposed a classification for the dominant ataxias⁴ used spinocerebellar ataxias (SCA) as the prefix. However, to be consistent with the classification of other movement disorders,⁴ we propose the same prefix independent of the mode of inheritance, in particular, as several ataxias can be inherited in both an autosomal dominant or autosomal recessive fashion (eg, *SPTBN2*, *AFG3L2*, *ITPR1*, *OPA1*; for a review, see ref. 5). As

the term SCA is quite strongly associated with dominant ataxias, we decided to use the ATX prefix instead. In the International Parkinson and Movement Disorder Society Genetic Mutation Online Database⁶ (MDSGene; available at <http://www.mdsgene.org>), which systematically links reported mutations to movement disorder phenotypes, the dominant ataxias will also carry the ATX prefix.

According to the recommendations of the International Parkinson and Movement Disorder Society Task Force on Classification and Nomenclature of Genetic Movement Disorders,⁴ the ATX prefix was assigned to those disorders that included ataxia as a prominent and/or consistent feature. When another movement disorder was also a prominent feature that generally coexisted with ataxia in several patients, a double prefix was assigned (e.g., ataxia (ATX)/Hereditary spastic paraplegia (HSP)). For those genes (eg, TPP1) that gave rise to allelic disorders (autosomal recessive spinocerebellar ataxia type 7 with pure ataxia and without myoclonus as well as neuronal ceroid lipofuscinosis type 2 with myoclonus and ataxia) that both present with prominent and consistent ataxia, a double prefix was established of which the order reflects the phenotype of the more prevalent or common presentation (in this case, ATX/MYC-TPP1). However, when another phenotype (eg, myoclonic epilepsy in POLG mutations) was the most common presentation or if ataxia was not usually predominant, the disease would appear in the proposed new list for the recessive ataxias, but would not receive the ATX prefix. To be considered for an ATX prefix, the association with ataxia must have been reported by more than one group of investigators. Last, a prefix was not assigned for complex or mixed phenotypes that usually showed a different predominant, nonmovement disorder clinical presentation (eg, mental retardation or epilepsy), as these probably would require a nonmovement disorder prefix that has not been established.

An initial draft list was prepared along with any doubts or queries, which were then reviewed by a working group member who had not initially reviewed the entity. At various stages of the process, all authors of this work held telephone conferences to discuss the criteria, discrepancies, and open issues. A cross-check with members of the chorea, myoclonus and hereditary spastic paraplegia working groups was done to ensure that the disorders that received the ATX prefix were referred to in a similar way with respect to the entities listed in the new nomenclature lists of the abovementioned disorders. Discrepancies were resolved by discussion until consensus was reached.

The Proposed New List for the Recessive Cerebellar Ataxias

The newly proposed list of recessively inherited cerebellar ataxias is shown in Table 1. This list includes

62 disorders that usually present with ataxia as a predominant and/or consistent feature, which were assigned the ATX prefix (group A, Table 1). It also lists 30 diseases that usually combine ataxia with other predominant and/or consistent movement disorders (eg, spastic paraparesis) and that received a mixed prefix (eg, ATX/HSP or HSP/ATX, depending on the phenotype that is consistent with the majority of cases; group B, Table 1). A group of 89 entities that usually present with other complex and/or nonataxia phenotypes but that may occasionally manifest cerebellar ataxia were not assigned an ATX prefix (group C, Table 1). Supplementary Table 1 lists the recessive diseases that appeared in our initial list following the abovementioned search strategy, but that upon review were classified as diseases with only occasional ataxia presentation, or diseases for which the presence of ataxia or the genetic finding itself remains unconfirmed. The posterior column ataxia with retinitis pigmentosa, a purely sensory ataxia as a result of mutations in the FLVCR1 gene, was excluded.

Discussion

Here we propose a new nomenclature of the genetically confirmed recessive cerebellar ataxias according to the principles and criteria of the International Parkinson and Movement Disorder Society Task Force on Classification and Nomenclature of Genetic Movement Disorders.

We identified and listed almost 100 different, confirmed genetic entities that present with ataxia as a dominant or relevant phenotype. We also provided a list of a similar number of entities that more commonly have nonataxia presentations but that may present with ataxia. We believe that this resource is a useful guide for the selection and interpretation of genetic testing when confronted with a patient with ataxia. A unique feature of our work is separating out those entities for which the genetic association remains unconfirmed. An important limitation of the previous nomenclature was the potential for perpetuating spurious associations.³ Requiring independent confirmation of associations before assigning prefixes should minimize such occurrences.

There have been earlier attempts to classify recessive cerebellar ataxias, a complex task because of the large clinical and genetic heterogeneity. Previous classifications have used phenotypic precision (ie, a list of the recessive ataxia genes per phenotypic category), for example, by separating Friedreich from non-Friedreich ataxias or by distinguishing those with from those without a peripheral neuropathy.^{1,7} However, this system does not reflect the within-genotype heterogeneity and is also difficult to update as it might not be clear what the consistent phenotype is for a new ataxia

TABLE 1. The proposed new list for the recessive cerebellar ataxias^a

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
ATX-ABCB7 ¹²	Sideroblastic anemia with spino-cerebellar ataxia	A. Disorders that present with ataxia as a predominant or consistent feature Intention tremor, pyramidal signs, hypochromic, microcytic anemia, abnormal pigmentation, skin atrophy	301310	XR	None
ATX-ABHD13	Polyneuropathy, hearing loss, ataxia, retinitis pigmentosa, and cataract (PHARC)	Retinopathy, cataract, hearing loss, intention tremor, pyramidal signs, peripheral neuropathy, pes cavus	612674	AR	None
ATX-ADCK3 ^{14,15}		Developmental delay, muscle weakness, pes cavus, exercise intolerance, myoclonus, dystonia, headache, stroke-like episodes, seizures	612016	AR	SCAF9, ARCA2
ATX-AH1 ^{16,17}		Developmental delay, morphological abnormalities, oculomotor apraxia, nystagmus, retinopathy, spasticity, scoliosis, seizures, renal failure, respiratory dysfunction	608629	AR	JBTS3
ATX-ALDH5A1 ¹⁸	Succinic semialdehyde dehydrogenase deficiency	Developmental delay, mental retardation, hyperkinesia, hyporeflexia, psychiatric symptoms, abnormal eye movements, seizures	271980	AR	None
ATX-ALG6 ¹⁹		Developmental delay, psychiatric symptoms, nystagmus, strabismus, peripheral neuropathy, muscle weakness, seizures, skeletal deformities, coagulation anomalies	603147	AR	None
ATX-ANO10 ²⁰		Cognitive impairment, nystagmus, hypermetric saccades, tortuous conjunctival vessels, pyramidal signs, intention tremor, proximal lower limbs atrophy, fasciculations, seizures, pes cavus	613728	AR	SCAR10, ARCA3
ATX-APT ^{21,22}		Hypometric saccades, oculomotor apraxia, nystagmus, ophthalmoplegia, peripheral neuropathy, scoliosis, pes cavus, choreoathetosis, tremor, dystonia, cognitive decline	208920	AR	AOA1
ATX-ARL13B ²³	Ataxia-telangiectasia (including variant ataxia-telangiectasia)	Developmental delay, oculomotor apraxia, retinopathy, respiratory dysfunction	612291	AR	JBTS8
ATX-ATM ²⁴		Telangiectases and other skin alterations, oculomotor apraxia, dystonia, chorea, myoclonus, tremor, peripheral neuropathy, distal muscular atrophy, short stature, hypogonadism, respiratory dysfunction, immunodeficiency, predisposition to neoplasia, glucose intolerance	208900	AR	None
ATX-BCKDHB ²⁵	Maple syrup urine disease	Maple syrup urine odor, life-threatening metabolic decompensation, lethargy, coma, hypoglycemia, ketosis, lactic acidosis, hallucinations, seizures, mental retardation if untreated, vomiting, pancreatitis	248600	AR	None
ATX-BTD4 ²⁶	Biotinidase deficiency	Developmental delay, optic atrophy, vision and hearing loss, seizures, metabolic ketacidosis, organic aciduria, skin problems, alopecia, hepatosplenomegaly, breathing problems	253260	AR	None
ATX-C10orf2 ²⁷	Hepatocerebral type of Mitochondrial DNA depletion syndrome	Psychomotor retardation, psychiatric symptoms, ophthalmoplegia, nystagmus, optic atrophy, hearing loss, peripheral neuropathy, myopathy, status epilepticus, epileptic encephalopathy, headaches, liver disease, hypergonadotrophic hypogonadism	271245	AR	None
ATX-CA84 ²⁸	Cerebellar ataxia and mental retardation with or without quadrupedal locomotion type 3	Mental retardation, dysarthria, quadrupedal gait, tremor	613227	AR	None
ATX-CEP290 ^{29,30}		Mental retardation, congenital amaurosis, oculomotor apraxia, retinopathy, retinal coloboma, nystagmus, nephronophthisis, neonatal breathing dysregulation	610188	AR	JBTS5
ATX-COX20 ³¹	Mitochondrial complex IV deficiency or cytochrome c oxidase deficiency	Developmental delay, mental retardation, pyramidal signs, peripheral neuropathy, dystonia, lactic acidosis retinopathy, optic atrophy, respiratory insufficiency	220110	AR/Mi	None
ATX-CWF19L1 ³²	Cerebrotendinous xanthomatosis (CTX)	Developmental delay, intellectual disability, tremor, hyperreflexia in lower limbs	616127	AR	SCAR17
ATX-CYP27A1 ³³		Tuberous skin and tendon xanthomas, xanthelemas, cataracts, chronic diarrhea, cognitive decline, psychiatric symptoms, peripheral neuropathy, parkinsonism, dystonia, myoclonus, spastic paraplegia, pseudobulbar palsy, seizures	213700	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
ATX-DNAJC19 ³⁴	3-methylglutaconic aciduria, type V	Developmental delay, mental retardation, growth retardation, optic atrophy, muscle weakness, dilated cardiomyopathy, long Q-T syndrome, genitourinary deformities	610198	AR	None
ATX-FXN ^{35,36}	Friedreich ataxia (FRDA)	Nystagmus, square wave jerks, optic atrophy, hearing loss, peripheral sensory neuropathy, pes cavus, hammertoes, muscle weakness, amyotrophy, extensor plantar responses, spasticity, spastic ataxia, chorea, scoliosis, hypertrophic cardiomyopathy, diabetes	229300	AR	None
ATX-GRID2 ³⁷		Developmental delay, cognitive impairment, esotropia, nystagmus, oculomotor apraxia, tonic upgaze, pale optic discs, retinopathy, pyramidal signs, muscle atrophy, joint contractures, scoliosis	616204	AR	SCAR18
ATX-GRN ³⁸		Dementia, myoclonic retinopathy, optic atrophy, seizures	614706	AR	CLN11
ATX-ITPR1 ³⁹	Gillespie syndrome	Developmental delay, mental retardation, aniridia or iris hypoplasia, scalloped pupillary margins of iris, nystagmus, visual impairments, postural tremor	206700	AR	None
ATX-KCNJ10 ⁴⁰	Seizures, sensorineural deafness, ataxia, mental retardation, and electrolyte imbalance (SESAME syndrome)	Developmental delay, mental retardation, sensorineural deafness, intention tremor, peripheral neuropathy, seizures, short stature, salt craving, enuresis, polydipsia, polyuria, electrolyte imbalance	612780	AR	None
ATX-KIAA0226 ⁴¹	Salih ataxia	Developmental delay, mental retardation, nystagmus, abnormal saccadic eye movements, seizures	615705	AR	SCAR15
ATX-L2HGDH ⁴²	L-2-hydroxyglutaric aciduria or academia	Psychomotor regression, mental retardation, cognitive impairment, hearing loss, strabismus, optic atrophy, nystagmus, spastic tetraparesis, facial dyskinesia, rigidity, dystonia, intention tremor, action-induced negative myoclonus, pyramidal signs, seizures, macrocephaly	236792	AR	None
ATX-MAN2B1 ⁴³	Alpha-mannosidosis	Developmental delay, mental retardation, growth retardation, sensorineural deafness, nystagmus, pyramidal signs, macrocephaly, facial dysmorphism, skeletal deformities, hepatosplenomegaly	248500	AR	None
ATX-MRE11A ⁴⁴	Ataxia-telangiectasia-like disorder type 1	Hypometric saccades, oculomotor apraxia, nystagmus, chorea, dystonia, myoclonus, tremor, hyporeflexia, distal muscle atrophy	604391	AR	None
ATX-MSTO1 ⁴⁵		Myopathy, developmental delay, growth impairment, pigmentary retinopathy with papillary pallor, tremor, skeletal abnormalities, pes cavus, dysmorphism	617619	AR	None
ATX-NPC1 ⁴⁶	Niemann-Pick disease type C1	Developmental regression, cognitive impairment, psychiatric symptoms, loss of speech, vertical supranuclear gaze palsy, dystonia, intention tremor, spasticity, seizures, hepatosplenomegaly, cholestatic jaundice, gelastic cataplexy	257220	AR	None
ATX-NPC2 ⁴⁷	Niemann-Pick disease type C2	Similar to ATX-NPC1 with severe pulmonary involvement and respiratory failure	607625	AR	None
ATX-OFD1 ⁴⁸		Developmental delay, mental retardation, recurrent infections, hirsutism, postaxial polydactyly, cystic renal disease, facial dysmorphism, macrocephaly	300804	XR	JBTS10
ATX-OPHN1 ⁴⁹	X-linked mental retardation with cerebellar hypoplasia and distinctive facial appearance	Developmental delay, mental retardation, spasticity, psychiatric symptoms, seizures, macrocephaly, facial dysmorphism, microphaly, hypoplastic scrotum, cryptorchidism, strabismus	300486	XR	None
ATX-OTC ⁵⁰ (Heterozygous females)	Ornithine transcarbamylase deficiency	Episodic extreme irritability, episodic vomiting and lethargy, protein avoidance, coma, delayed growth, developmental delay, seizures	311250	XR	None
ATX-PEX ⁵¹	Peroxisome biogenesis disorder 9B or Zellweger spectrum disorder	Developmental delay, cognitive impairment, cataracts, retinopathy, anosmia, hearing loss, muscle weakness, pes cavus, peripheral neuropathy	614879	AR	None
ATX-PEX10 ⁵²	Peroxisome biogenesis disorder 6B or Zellweger spectrum disorder	Mental retardation, intention tremor, peripheral neuropathy, pyramidal signs, distal muscle atrophy, pes cavus, dysmetric saccades, impaired smooth pursuit, nystagmus, diabetes	614871	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
ATX-PHYH ⁶³	Refsum disease or hereditary motor and sensory neuropathy type IV	Muscle weakness and atrophy, peripheral neuropathy, sensory impairment, pes cavus, anopia, sensorineural deafness, retinopathy, ichthyosis, shortening of the metacarpals and metatarsals, multiple epiphyseal dysplasia, cardiomyopathy, sudden death	266500	AR	None
ATX-PMIM2 ⁵⁴	Congenital disorder of glycosylation type Ia (CDG1a) or Jaeken syndrome	Developmental delay, psychomotor retardation, cognitive impairment, strabismus, nystagmus, retinopathy, peripheral neuropathy, stroke-like episodes, seizures, microcephaly, morphological abnormalities, abnormal subcutaneous fat tissue distribution, pericardial effusion, hepatomegaly, liver steatosis, diarrhea, renal cysts, nephrotic syndrome, thrombotic events, hypothyroidism, hypergonadotropic hypogonadism, scoliosis, osteoporosis	212065	AR	None
ATX-PMPCA ⁵⁵		Developmental delay, mental retardation, visuospatial defects, nystagmus, dysmetric saccades, pes cavus, hyperreflexia, spasticity, tremor, short stature	213200	AR	SCAR2
ATX-POLR3B ⁵⁶	Hypomyelinating leukodystrophy type 8 with or without oligodontia and/or hypogonadotropic hypogonadism	Developmental delay, cognitive decline, mental retardation, nystagmus, abnormal saccades, vertical gaze palsy, myopia, spasticity, tremor, oligodontia, hypodontia, delayed dentition, short stature, hypogonadism	614381	AR	None
ATX-PRKGG ⁵⁷	Infantile-onset multisystem neurological, endocrine, and pancreatic disease (IMNEPD)	Peripheral neuropathy, pyramidal signs, cognitive impairment, depression, myoclonus, tremor	605361	AR	None
ATX-PTRH2 ⁵⁸	RIDDLE syndrome	Developmental delay, failure to thrive, poor postnatal growth, poor expressive speech, peripheral neuropathy, distal muscle weakness, foot and hand deformities, hypothyroidism, pancreatic endocrine insufficiency, facial dysmorphism, brachycephaly, short stature	616263	AR	None
ATX-RNF168 ⁵⁹	Gordon Holmes syndrome	Learning difficulties, immunodeficiency, dry skin, progressive pulmonary fibrosis and failure, ocular telangiectasia, short stature, microcephaly, dysmorphic features	611943	AR	None
ATX-RNF216 ⁶⁰		Mental retardation, dementia, psychiatric symptoms, chorea, sensorineural deafness, hypogonadism	212840	AR	None
ATX-SETX ⁶¹		Saccadic pursuit, oculomotor apraxia, nystagmus, strabismus, intention tremor, head tremor, dystonia, chorea, pyramidal signs, peripheral neuropathy, distal muscle atrophy and weakness, pes cavus, scoliosis	606002	AR	SCAR1, AOA2
ATX-SIL1 ⁶²	Marinesco-Sjögren syndrome	Developmental delay or regression, mental retardation, growth retardation, microcephaly, facial dysmorphism, short stature, congenital cataracts, nystagmus, strabismus, spasticity, muscle weakness and atrophy, peripheral neuropathy, scoliosis, skeletal deformities, hypogonadism	248800	AR	None
ATX-SLC17A5 ⁶³	Salla disease or Finnish type sialuria and the variant syndrome of infantile form of sialic acid storage disorder	Developmental delay, mental retardation, rigidity, spasticity, seizures, visceromegaly, facial dysmorphism, hypopigmented skin	604369 and 269920	AR	None
ATX-SLC33A1 ⁶⁴	Congenital cataracts, hearing loss, and neurodegeneration (CCHLND)	Psychomotor retardation, nystagmus, congenital cataracts, hearing loss, seizures	614482	AR	None
ATX-SLC52A2 ^{65,66} Brown-Vialetto-Van Laere syndrome type 2)		Optic atrophy, blindness, cochlear degeneration, deafness	271250	AR	SCAR3
ATX-SNX14 ⁶⁷		Developmental delay, mental retardation, autistic behavior, macrocephaly, sensorineural hearing loss, nystagmus, apraxia, spasticity, extensor plantar responses, hyporeflexia, seizures, hypertrichosis, scoliosis, distal skeletal deformities, brachycamptodactyly, facial dysmorphism	616354	AR	SCAR20
ATX-SPTBN2 ⁶⁸		Developmental delay, cognitive impairment, speech delay, intention tremor, spasticity, hyperreflexia, hypometric saccades, nystagmus, abnormal eye movements with convergent squint	615386	AR	SCAR14

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
ATX-SRD5A3 ⁶⁹	Congenital disorder of glycosylation, type Iq	Developmental delay, mental retardation, coloboma, nystagmus, facial dysmorphism, hypertrophicosis, skin abnormalities, coagulation defects, microcytic anemia	612379	AR	None
ATX-STUB1 ⁷⁰		Nystagmus, external ophthalmoplegia, pyramidal signs, tremor, myoclonus, cognitive impairment, peripheral neuropathy, reduced vibration sense in lower limbs, hypogonadism	615768	AR	SCAR16
ATX-SYNE1 ⁷¹		Nystagmus, abnormal saccades and slow or jerky pursuit, hyperreflexia in lower limbs, motor neuron involvement, respiratory dysfunction due to multisystemic neuromuscular compromise, mental retardation	610743	AR	SCAR8, ARCA1
ATX-TTPA ⁷²	Ataxia with vitamin E deficiency	Nystagmus, retinopathy, proprioception loss, areflexia in lower limbs, peripheral neuropathy, extensor plantar response, head titubation or tremor, dystonia, hypoaesthesia, tendon xanthomas, pes cavus, hammer toes, kyphoscoliosis	277460	AR	None
ATX-TMEM216 ⁷³		Developmental delay, failure to thrive, mental retardation, impaired saccades, oculomotor apraxia, nystagmus, optic nerve coloboma, chorioretinal coloboma, retinopathy, esotropia, polydactyly, nephronophthisis, renal cysts, hypoplastic genitalia, episodic hyperpnea or apnea, facial dysmorphism, macrocephaly	608091	AR	JBTS2
ATX-TMEM67 ^{74,75} , ATX-RPGRIP1L ⁷⁶⁻⁷⁸ , ATX-CC2D2A ⁷⁹	COACH syndrome (cerebellar vermis hypoplasia, oligophrenia, ataxia, ocular coloboma, and hepatic fibrosis) and allelic disorders	Developmental delay, mental retardation, oculomotor apraxia, ocular coloboma, retinopathy, nystagmus, facial dysmorphism, polydactyly, pyramidal signs, seizures, splenomegaly, renal failure, liver disease, breathing dysregulation	216360, 611560, 610688	AR	JBTS6, JBTS7
ATX-TMEM231 ⁸⁰		Developmental delay, oculomotor apraxia, psychiatric symptoms, polydactyly, syndactyly, renal cysts, retinopathy	614970	AR	JBTS20
ATX-TTC19 ⁸¹	Mitochondrial complex III deficiency nuclear type 2	Developmental delay, cognitive impairment, apraxia, psychiatric symptoms, dysphonia, nystagmus, bradykinesia, dystonia, muscle atrophy and weakness, pyramidal signs	615157	AR	None
ATX-VLDLR ⁸²	Cerebellar ataxia, mental retardation, and dysequilibrium syndrome type 1	Developmental delay, mental retardation, lack of speech development, strabismus, postnatal cataracts, nystagmus, saccadic visual pursuit, quadrupedal gait, intention tremor, hyperreflexia, seizures, pes planus, short stature	224050	AR	None
ATX-WDR73 ⁸³	Galloway-Mowat syndrome	Delayed psychomotor development, mental retardation, oculomotor apraxia, optic atrophy, retinopathy, seizures, spastic quadriplegia, dystonia, hyperreflexia, skin abnormalities (osmophilic skin vessels), skeletal deformities, genitourinary affection, facial dysmorphias, microcephaly, short stature, intrauterine growth retardation	251300	AR	SCAR5
ATX-WDR81 ⁸⁴	Cerebellar ataxia, mental retardation, and dysequilibrium syndrome type 2	Developmental delay, mental retardation, strabismus, facial dysmorphism, quadrupedal locomotion, poor or absence language development, tremor, hyporeflexia, hirsutism, small hands and feet, thoracic kyphosis, short stature	610185	AR	None
ATX/HSP-AFG3L2 ⁸⁵	B. Combined ataxias	(disorders where ataxia frequently coexists with other predominant or consistent movement disorders)	614487	AR	SPAX5
ATX/HSP-DARS2 ⁸⁶	Leukoencephalopathy with brain stem and spinal cord involvement and lactate elevation	Spastic paraparesis, oculomotor apraxia, dystonia, myoclonus, myoclonic epilepsy, generalized tonic-clonic seizures, distal muscle atrophy, peripheral neuropathy	611105	AR	None
ATX/HSP-FOLR1 ⁸⁷	Neurodegeneration as a result of cerebral folate transport deficiency	Spastic paraparesis, developmental regression, mental retardation, visual disturbances, sensorineural hearing loss, chorea, generalized tonic-clonic, atonic and myoclonic seizures	613068	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
ATX/HSP-GJC2 ⁸⁸	Hypomyelinating leukodystrophy-2 or Pelizaeus-Merzbacher-like disease	Spastic paraparesis, developmental delay, mental retardation, lack of independent ambulation, poor head and trunk control in infancy, optic atrophy, rotary nystagmus, myopia, facial weakness, tremor, head titubation, dystonia, spasticity, seizures, peripheral neuropathy	608804	AR	None
ATX/HSP-HEXA ⁸⁹	Tay-Sachs disease or GM2-gangliosidosis type I	Spastic paraparesis, cognitive decline, psychiatric symptoms, late spasticity, dystonia, peripheral neuropathy, macular pallor with prominence of fovea centralis (cherry red spot), blindness, muscular weakness and atrophy, seizures	272800	AR	None
ATX/HSP-HEXB ⁹⁰	Sandhoff disease or GM2-Gangliosidosis type II	Spastic paraparesis, progressive mental and motor deterioration, macrocephaly, macular pallor with prominence of fovea centralis (cherry red spot), blindness, dysmorphic features, startle reaction, hyperreflexia, muscular atrophy, fasciculations, cardiomegaly, episodic abdominal pain, chronic diarrhea, hepatosplenomegaly, macroglossia, high lumbar gibbus	268800	AR	None
ATX/HSP-PNPLA6 ⁹¹⁻⁹⁵	PNPLA6-related disorders: Boucher-Neuhauser syndrome, Laurence-Moon syndrome and Oliver-McFarlane syndrome	Visual impairment due to chorioretinal dystrophy, nystagmus, distal muscle atrophy, intention tremor, spastic paraparesis, cognitive impairment, peripheral neuropathy, hypogonadism, tri-omegaly, short stature	215470, 245800, 275400, 612020	AR	SPG39
ATX/HSP-POLR3A ⁹⁶	Hypomyelinating leukodystrophy type 7 with or without oligodentia and/or hypogonadotropic hypogonadism or 4H syndrome	Spastic paraparesis, developmental delay, cognitive decline, mental retardation, optic atrophy, nystagmus, abnormal saccades, vertical gaze palsy, pyramidal signs, postural tremor, dystonia, seizures, peripheral neuropathy, oligodontia, hypodontia, delayed dentition, hypogonadotropic hypogonadism, short stature	607694	AR	None
ATX/HSP-SACS ⁹⁷	Autosomal recessive spastic ataxia of Charlevoix-Saguenay (ARSACS) or autosomal recessive spastic ataxia type 6	Spastic paraparesis, delayed walking development, retinal striation, nystagmus, impaired smooth pursuit, pes cavus, hammertoes, finger deformities, hyperreflexia, ankle areflexia, spasticity, extensor plantar responses, scoliosis, distal muscle weakness and atrophy, peripheral neuropathy, dystonia, erectile dysfunction	270550	AR	SPAX6
HSP/ATX-B4GALNT1 ⁹⁸		Spastic paraparesis, distal amyotrophy, nonprogressive cognitive impairment, sensory polyneuropathy, pes cavus, stereotypies, emotional lability, psychiatric symptoms, seizures	609195	AR	SPG26
HSP/ATX-CAPN1 ⁹⁹		Spasticity in lower and upper limbs, other pyramidal signs, peripheral neuropathy, pes cavus, pes valgus, nystagmus	616907	AR	SPG76
HSP/ATX-CLCN2 ¹⁰⁰		Spastic paraparesis, learning disabilities, headache, optic neuropathy, chorioretinopathy, visual field defects	615651	AR	None
HSP/ATX-CYP7B1 ¹⁰¹	Leukoencephalopathy with ataxia	Spastic paraparesis, other pyramidal signs besides spasticity, cognitive impairment, nystagmus, optic atrophy, cataracts, altered saccadic eye movements, pes cavus, sensation deficits	270800	AR	SPG5A
HSP/ATX-GAN ¹⁰²		Spastic paraparesis, distal limb muscle weakness and atrophy as a result of peripheral neuropathy, distal sensory impairment, kinky or curly hair, foot or hand deformities, scoliosis, nystagmus, facial weakness	256850	AR	None
HSP/ATX-GBA2 ¹⁰³		Spastic paraparesis, other pyramidal signs, muscle weakness, pseudobulbar dysarthria, cognitive impairment, mental retardation, congenital cataracts, nystagmus, hearing loss, head tremor, peripheral neuropathy, pes cavus, scoliosis, infertility, small testicles, hypogonadism in males	614409	AR	SPG46
HSP/ATX-KIF1C ^{104,105}		Spastic paraparesis, chorea, myoclonus, dystonia, developmental delay, mild mental retardation, hypodontia, ptosis, short stature, sensorineural deafness, pes planus	611302	AR	SPAX2, SPG58
HSP/ATX-MLC1 ¹⁰⁶	Megalencephalic leukoencephalopathy with subcortical cysts	Spastic paraparesis, developmental delay, mental retardation, seizures, macrocephaly, spasticity	604004	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
HSP/ATX-PLP1 ¹⁰⁷		Spastic paraparesis, lower limb weakness, other pyramidal signs, mental retardation, upper limb spasticity, pes cavus, joint contractures, nystagmus, optic atrophy, tremor	312920	XR	None
HSP/ATX-SPG7 ¹⁰⁸		Spastic paraparesis, optic atrophy, chronic external ophthalmoplegia-like phenotype, nystagmus, decreased vibratory sense in the lower limbs, pes cavus, scoliosis	607259	AR/AD	SPG7
HSP/ATX-UOHL1 ¹⁰⁹	Childhood-onset neurodegeneration with optic atrophy	Spastic paraparesis, other pyramidal signs, myotonia, myokymia, head titubation, intellectual impairment, impaired distal sensation to vibration and position, optic atrophy, nystagmus	615491	AR	None
HSP/ATX/NBIA-FA2H ¹¹⁰	Fatty acid hydroxylase-associated neurodegeneration (FAHN)	Spastic paraparesis, cognitive decline, optic nerve atrophy, seizures, dystonia, parkinsonism	612319	AR	SPG35
ATX/MYC-TPP1 ^{111,112} (allelic disorders: Neuronal ceroid lipofuscinosis type 2 and autosomal recessive spinocerebellar ataxia type 7)		Myoclonus, developmental regression, speech and language difficulties, nystagmus, diplopia, hypermetric saccades, progressive vision loss, retinopathy, postural tremor, pyramidal signs, spastic paraparesis, decreased vibration sense, fasciculations, seizures	609270, 204500	AR	SCAR7, CLN2
MYC/ATX-CSTB ¹¹³	Myoclonic epilepsy of Unverricht and Lundborg	Myoclonic epilepsy, stimulus sensitive segmental or generalized myoclonus, action myoclonus, generalized tonic-clonic or absence seizures, mental and motor deterioration	254800	AR	None
MYC/ATX-EPM2A ¹¹⁴	Lafora disease	Myoclonic or other types of seizures, focal visual seizures, drop attacks cognitive decline, psychosis, myoclonus	607566	AR	None
MYC/ATX-GOSR2 ¹¹⁵		Myoclonic, absence and tonic-clonic seizures, drop attacks, action myoclonus, tremor, areflexia, scoliosis, pes cavus, syndactyly	614018	AR	None
MYC/ATX-KCTD7 ¹¹⁶	Progressive myoclonic epilepsy type 3 with or without intracellular inclusions	Myoclonic epilepsy, secondary generalization seizures, neurologic regression following seizure onset, mental retardation, hyperreflexia, opsoclonus, optic atrophy, visual loss, microcephaly	611726	AR	None
MYC/ATX-NEU1 ¹¹⁷	Neuraminidase deficiency or sialidosis type I and II	Myoclonus, mental retardation, seizures, hyperreflexia, muscle atrophy, skeletal malformations, hepatosplenomegaly, cardiomyopathy, progressive vision loss, cherry-red spots, cataracts, nystagmus, sensorineural hearing loss, short stature, dysmorphic features	256550	AR	None
MYC/ATX-NHLRC1 ¹¹⁸	Lafora disease	Myoclonic or other types of seizures, focal visual seizures, drop attacks cognitive decline, psychosis, myoclonus	608072	AR	None
DYT/ATX-ATP7B ¹¹⁹	Wilson disease	Dystonia, occasionally parkinsonism, chorea, flapping tremor, rest, action, and intention tremor, orofacial dyskinesias, liver disease, Kayser Fleischer rings, psychiatric symptoms	277900	AR	None
PxMD-DYT-ATX-PRRT2 ¹²⁰ (Biallelic mutations)		Paroxysmal non-kinesigenic dyskinesia, prolonged episodes of ataxia, motor tics, seizures, hemiplegic migraine, learning difficulties, psychiatric symptoms	614386	AR	None
HSP-ACP33 ¹²¹	C. Disorders that usually present with other phenotypes but can have occasionally ataxia (no ATX prefix)	Spastic paraparesis, other pyramidal signs, apraxia, bulbar dysfunction, developmental delay, cognitive impairment, akinetic mutism, dyskinesias, peripheral neuropathy	248900	AR	SPG21
HSP-DDHD2 ¹²²		Spastic paraparesis, other pyramidal signs, developmental delay, mental retardation, lower limbs weakness, strabismus, facial dysmorphism, short stature, optic nerve hypoplasia	615033	AR	SPG54
HSP-KIAA0415 ¹²³		Spastic paraparesis, lower limb weakness, dystonia, myoclonus, parkinsonism, peripheral neuropathy	613647	AR	SPG48
HSP-KIAA1840 ¹²⁴		Spastic paraparesis, lower limb atrophy, weakness, peripheral neuropathy, pes cavus, parkinsonism, cognitive impairment, mental retardation, retinopathy, pigmented macular degeneration, nystagmus	604360	AR	SPG11

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
HSP-KIF1A ¹²⁵		Spastic paraparesis, lower limb atrophy and weakness, peripheral neuropathy, saccadic ocular pursuit	610357	AR	SPG30
HSP-SPARTIN ¹²⁴		Spastic paraplegia and upper limb spasticity, distal muscle atrophy, pes cavus, hammer toes, short stature, dysmorphism, developmental delay	275900	AR	SPG20
HSP-ZFWE26 ¹²⁶		Spastic paraparesis, mental retardation, cognitive impairment, psychiatric symptoms, parkinsonism, distal amyotrophy, pes cavus, peripheral neuropathy, retinopathy, hearing loss, pigmentary maculopathy	270700	AR	SPG15
MYC-CLN5 ¹²⁷		Myoclonus, tremor, mental retardation, cognitive decline, visual loss, glaucoma, retinopathy, nystagmus, hyperreflexia, seizures	256731	AR	CLN5
MYC-CLN6 ¹²⁸		Myoclonus, dystonia, bradykinesia, myoclonic epilepsy, dementia, mental retardation, psychiatric symptoms, blindness	204300	AR	CLN6
MYC-SCARB2 ¹²⁹	Progressive myoclonic epilepsy type 4 with or without renal failure	Action and resting myoclonus, intention and postural tremor, horizontal saccades, seizures, nephrotic syndrome, renal failure	254900	AR	None
DYT/PARK-GLB1 ¹³⁰		Dystonia, parkinsonism, pyramidal signs, cognitive deficits, skeletal abnormalities, short stature, corneal clouding, cardiomyopathy	230600	AR	None
DYT/PARK-SPR ¹³¹	Septaplerin reductase deficiency	Dystonia, parkinsonism, motor and speech delay, truncal hypotonia, limb hypertonia and hyperreflexia, oculogyric crises, psychiatric symptoms, autonomic dysfunction, diurnal fluctuation	612716	AR	None
NBIA/DYT/PARK-CP ¹³²	Aceruloplasminemia	Dystonia, parkinsonism, chorea, cognitive impairment, retinopathy, blepharospasm, systemic hemosiderosis, diabetes, anemia	604290	AR	None
NBIA/DYT-PANK2 ¹³³	Pantothenate kinase-associated neurodegeneration (PKAN)	Dystonia, parkinsonism, chorea, tremor, spasticity, cognitive decline, apraxia of eyelid opening, retinopathy, optic atrophy, psychiatric symptoms, muscle atrophy	234200	AR	NBIA1
NBIA/DYT/PARK-PLA2G6 ¹³⁴	PLA2G6-associated neurodegeneration or Infantile neuroaxonal dystrophy type 1	Dystonia, parkinsonism, cognitive decline, pyramidal signs, psychiatric symptoms (adult phenotype), ataxia (childhood phenotype)	256600/610217	AR	NBIA2, PARK14
AAAS ¹³⁵	Achalasia-addisonianism-alacrimia syndrome or Triple-A syndrome or Allgrove syndrome	Developmental delay, mental retardation, pyramidal signs, distal muscle weakness and atrophy, achalasia, autonomic dysfunction, anisocoria, peripheral neuropathy, adrenal insufficiency, alacrima, optic atrophy, short stature, hyperpigmentation, hyperkeratosis of the palms and soles	231550	AR	None
AARS2 ¹³⁶	Progressive leukoencephalopathy with ovarian failure	Dystonia, tremor, developmental delay, cognitive decline, apraxia, psychiatric symptoms, nystagmus, spasticity, premature ovarian failure	615889	AR	None
ABCD1 ¹³⁷	X-linked adrenoleukodystrophy	Visual disturbances, sensation deficits, spastic paraplegia, autonomic failure, adrenal dysfunction	300100	XR	None
ARX ¹³⁸	Partington syndrome	Dystonia, mental retardation, spasticity, seizures, morphological abnormalities	309510	XR	None
ATAD3A ^{139,140}	Harel-Yoon syndrome	Developmental delay, intellectual disability, optic atrophy, nystagmus, spasticity, distal limb muscle atrophy, peripheral neuropathy, scoliosis, dysmorphism, hypertrophic cardiomyopathy	617183	AR/AD	None
AUH ¹⁴¹	3-methylglutaconic aciduria type I	Dystonia, developmental delay, failure to thrive, mental retardation, spastic quadriplegia, cognitive impairment, hyperreflexia, metabolic acidosis, febrile seizures, optic atrophy	250950	AR	None
BCS1L, COX10, COX15, FOXRED1, NDUFAF2, NDUFS3, NDUFS4, NDUFAF6, NDUFS7, NDUFS8, NDUFA10, SDHA, SURF1 ¹⁴²	Leigh syndrome	Dystonia, failure to thrive, psychomotor retardation, mental retardation, pyramidal signs, seizures, psychiatric symptoms, lactic acidosis, hypertrochosis, respiratory failure, pigmentary retinopathy, ptosis, strabismus, nystagmus, optic atrophy, ophthalmoplegia	256000	Mi/AR	None
C5orf42 ¹⁴³		Developmental delay, oculomotor apraxia, polydactyly, syndactyly, abnormal breathing pattern	614615	AR	JBTS17

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
CTC1 ¹⁴⁴	Cerebroretinal microangiopathy with calcifications and cysts or Coats plus syndrome	Dystonia, tremor, intrauterine growth retardation, growth retardation, seizures, spasticity, hemiplegia, cognitive decline, pyramidal signs, bone marrow failure, skin and hair abnormalities, intracranial calcifications, skeletal deformities, intestinal bleeding, retinopathy, optic atrophy, blindness	612199	AR	None
CTDP1 ¹⁴⁵	Congenital cataracts with facial dysmorphism and neuropathy	Chorea, developmental delay, congenital cataracts, nystagmus, cognitive impairment, pyramidal signs, peripheral neuropathy, hypo- or hypergonadotrophic hypogonadism, acute rhabdomyolysis, pes cavus, talipes equinovarus, skeletal deformities, facial dysmorphism	604168	AR	None
CTSA ¹⁴⁶	Galactosialidosis	Myoclonus, mental retardation, seizures, angiokeratoma, facial dysmorphism, conjunctival telangiectases, macular cherry red spot, hearing loss, hemangiomas, hepatosplenomegaly, dysostosis multiplex, cardiac valvular disease	256540	AR	None
CTSF ¹⁴⁷		Myoclonus, tremor, dementia, perioral dyskinesias, pyramidal signs, seizures, psychiatric symptoms	615362	AR	CLN13
CUL4B ¹⁴⁸	Cabezas type of X-linked syndromic mental retardation	Tremor, mental retardation, speech delay, hypogonadism, short stature, facial dysmorphism, skeletal abnormalities, seizures, psychiatric symptoms, central obesity, macrocephaly	300354	XR	None
DKC1 ¹⁴⁹	X-linked dyskeratosis congenita (ataxia reported in the severe variant: Hoyeraal-Hreidarsson syndrome)	Intrauterine growth retardation, developmental delay, mental retardation, microcephaly, multi-system involvement, bone marrow failure resulting in immunodeficiency, enteropathy, strabismus, cataracts, optic atrophy, sparse eyelashes, conjunctival leukoplakia, short stature, pulmonary fibrosis, liver failure, skin atrophy, nail dystrophy, carcinomas, leukemia	305000	XR	None
DLD ¹⁵⁰	Dihydropyrimidine dehydrogenase deficiency or Maple syrup urine disease type II	Dystonia, developmental delay, episodic encephalopathy, seizures, lactic or metabolic acidosis, recurrent vomiting, hepatomegaly, liver dysfunction, hypertrophic cardiomyopathy, microcephaly	246900	AR	None
EIF2B1, EIF2B2, EIF2B3, EIF2B4, EIF2B5 ¹⁵¹	Leukoencephalopathies with vanishing white matter	Mental and motor retardation or regression, cognitive impairment, psychiatric symptoms, optic atrophy, pyramidal signs, seizures, ovarian failure, clinical features worsened by head trauma or fever, macrocephaly, lethargy	603896	AR	None
ERCC4 ^{152,153}	XFE progeroid syndrome and the allelic entity Xeroderma pigmentosum complementation group F/Cockayne syndrome	Dwarfism, cachexia, microcephaly, photosensitivity, wizened appearance, scoliosis, delayed sexual maturity, chorea, tremor, mental retardation, cognitive decline, nystagmus, astigmatism, deep-set eyes, hearing impairment, short stature, seborrheic keratosis-like papules, abnormal pigmentation, skin cancer susceptibility, plantar warts	610965, 278760	AR	None
EXOSC3 ¹⁵⁴	Pontocerebellar hypoplasia type 1B	Tremor, developmental delay, poor growth, axial hypotonia, spasticity, hyperreflexia, lack of speech, seizures, peripheral neuropathy, muscle atrophy and weakness, tongue atrophy and fasciculations, foot deformities, joint contractures, respiratory insufficiency, oculomotor apraxia, nystagmus, strabismus, retinopathy, microcephaly	614678	AR	None
GALC ¹⁵⁵	Krabbe disease or galactocerebrosidase deficiency	Failure to thrive, developmental delay or regression, deafness, blindness, optic atrophy, nystagmus, hypersensitive to stimuli, irritability, spastic tetraparesis, seizures, muscular, decerebrate posturing, peripheral neuropathy, pes cavus, tongue atrophy, episodic fever	245200	AR	None
GBA ¹⁵⁶	Gaucher disease type III or subacute neuronopathic type	Myoclonus, developmental delay, dementia, psychiatric symptoms, spastic paraparesis, horizontal supranuclear gaze palsy, abnormal saccade eye movements, strabismus, seizures, short stature, hepatosplenomegaly, pancytopenia	231000	AR	None
GJC2 ¹⁵⁷		Tremor, slow saccades, sensorineural hearing loss, strabismus, spastic paraparesis, other pyramidal signs, pes cavus, scoliosis, seizures, cognitive impairment	613206	AR	SPG44
GPR56 or ADGRG1 ¹⁵⁸	Bilateral frontoparietal polymicrogyria	Developmental delay, mental retardation, pyramidal signs, esotropia, exotropia, strabismus, nystagmus, seizures	606854	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
HEPACAM ¹⁵⁹	Autosomal recessive megalencephalic leukoencephalopathy with subcortical cysts type 2A	Developmental delay, mental retardation, cognitive decline, spasticity, seizures, macrocephaly	613925	AR	None
HIBCH ¹⁶⁰	3-hydroxyisobutyryl-CoA hydrolase deficiency	Dystonia, myoclonus, developmental delay or regression, seizures, nystagmus, strabismus, facial dysmorphism, head titubation, persistent vomiting	250620	AR	None
HSD17B4 ¹⁶¹	Perrault syndrome type 1 (allelic with Peroxisomal D-bifunctional protein deficiency)	Sensorineural deafness, ovarian dysgenesis, primary amenorrhea, developmental delay, mental retardation, spastic diplegia, pes cavus, pes equinovarus, peripheral neuropathy, nystagmus, short stature, scoliosis	233400	AR	None
LRPPRC ¹⁶²	French Canadian type of Leigh syndrome	Tremor, developmental delay, mental retardation, failure to thrive, language delay, seizures, lactic acidosis, metabolic crises, strabismus, facial dysmorphism, liver dysfunction	220111	AR	None
LYST ¹⁶³	Chediak-Higashi syndrome	Parkinsonism, tremor, mental retardation, cranial nerve palsies, spastic paraparesis, peripheral neuropathy, foot drop, seizures, anemia, recurrent cutaneous and systemic pyogenic infections, severe immune deficiency, hair hypopigmentation, reduced visual acuity, nystagmus, strabismus, reduced iris pigmentation, macular hypoplasia, hepatosplenomegaly, jaundice	214500	AR	None
MAG ¹⁶⁴		Developmental delay, cognitive impairment, impaired distal vibration sense, peripheral neuropathy, distal muscle atrophy in lower limbs, spastic paraplegia, optic atrophy, nystagmus, visual impairment	616680	AR	SPG75
MECP2 ¹⁶⁵	Lubs X-linked mental retardation syndrome	Chorea, psychomotor retardation, macro- or microcephaly, facial dysmorphism, seizures, spasticity, recurrent respiratory infections, cryptorchidism, asymmetric skull, stereotypic hand movements, autistic features, depression, compulsions, psychosis	300260	XR	None
MFS08 ¹⁶⁶		Developmental regression, mental retardation, cognitive impairment, optic atrophy, retinopathy, blindness, seizures, myoclonus	610951	AR	CLN7
MKS1 ¹⁶⁷		Developmental delay, intellectual disability, nystagmus, oculomotor apraxia, retinopathy	617121	AR	JBTS28
MMACHC ¹⁶⁸	cb1C type of combined methylmalonic aciduria and homocystinuria	Tremor, failure to thrive, developmental delay, mental retardation, dementia, retinopathy, visual deficits, nystagmus, facial dysmorphism, seizures, hypergonadotropic hypogonadism, anemia, renal failure, microcephaly	277400	AR	None
MPV17 ¹⁶⁹	Mitochondrial DNA depletion syndrome-6 or Navajo neurohepatopathy	Dystonia, neonatal jaundice, failure to thrive, developmental delay, peripheral neuropathy, hypo-areflexia, pain insensitivity, acral ulceration and osteomyelitis leading to autoamputation, painless fractures due to injury, distal muscle weakness, lactic acidosis, systemic infections, liver dysfunction, Reye syndrome-like episodes, short stature	256810	AR	None
MTFMT ¹⁷⁰	Combined oxidative phosphorylation deficiency type 15	Tremor, developmental delay, cognitive impairment, pyramidal signs, seizures, strabismus, nystagmus, optic atrophy, short stature, obesity, cardiopathy	614947	AR	None
MTTP ¹⁷¹	Abetalipoproteinemia	Peripheral neuropathy, retinopathy, acanthocytosis, steatorrhea (celiac-like syndrome), hepatic steatosis	200100	AR	None
MVK ¹⁷²	Mevalonic aciduria	Developmental delay, psychomotor retardation, failure to thrive, recurrent febrile crises with lymphadenopathy, hepatosplenomegaly, anemia, morbilliform rash, kyphoscoliosis, arthralgias, facial dysmorphism, nystagmus, central cataracts, retinal dystrophy, microcephaly	610377	AR	None
NPHP1 ¹⁷³		Developmental delay, mental retardation, congenital head tilt, abnormal eye movements, nystagmus, oculomotor apraxia, hypometric saccades, tubulointerstitial medullary cystic kidney disease, nephronophthisis, renal failures	609583	AR	JBTS4
NUBPL ¹⁷⁴	Mitochondrial complex I deficiency	Developmental delay, strabismus, nystagmus, contractures, spasticity, cognitive decline	252010	AR/Mi/AD	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
OFA1 ¹⁷⁵	Behr syndrome or infantile hereditary optic atrophy with neurologic abnormalities	Tremor, developmental delay, mental retardation, optic atrophy, progressive visual loss, nystagmus, spasticity, pyramidal signs, myopathy, posterior column sensory loss, peripheral neuropathy, tendon and muscular contractures	210000	AR	None
OFA3 ¹⁷⁶	3-methylglutaconic aciduria type III or Costeff syndrome	Chorea, cognitive impairment, optic atrophy, visual loss, cataracts, nystagmus, pyramidal signs	258501	AR	None
PDHX or PDX1 ¹⁷⁷	Lactic acidemia due to PDX1 deficiency	Dystonia, developmental delay, mental retardation, microcephaly, optic atrophy, hypertelorism, facial dysmorphism, spastic quadriplegia, seizures, lactic or metabolic acidosis	245349	AR	None
PEX2 ¹⁷⁸	Peroxisome biogenesis disorder 5B or Zellweger spectrum disorders	Tremor, developmental delay, peripheral neuropathy, pes cavus, hypoaacusia, slow saccades, oculomotor apraxia, nystagmus, retinopathy, strabismus	614867	AR	None
PEX6 ²⁵	Microcephaly, seizures, and developmental delay and allelic disorders (eg, autosomal recessive axonal Charcot-Marie-Tooth)	Optic atrophy, blindness, cochlear degeneration, deafness	271250	AR	SCAR3
PNKP ¹⁷⁹⁻¹⁸¹	Purine nucleoside phosphorylase deficiency or Nucleoside phosphorylase deficiency	Dystonia, developmental delay, mental retardation, cognitive impairment, oculomotor apraxia, tetraplegia, impaired vibration sense, peripheral neuropathy, pes cavus, hammetoes, distal muscle weakness and atrophy, seizures, microcephaly	616267, 613402, 605610	AR	AOA4
PNP ¹⁸²	PNP	Tremor, developmental delay, failure to thrive, mental retardation, spastic diplegia, tetraparesis, behavioral disorder, autoimmune hemolytic anemia, frequent infections, splenomegaly, pneumonia	613179	AR	None
POLG ¹⁸³⁻¹⁸⁶	Allelic disorders: Mitochondrial recessive ataxia syndrome (MIRAS) or sensory ataxic neuropathy, dysarthria, and ophthalmoparesis (SANDO), Autosomal recessive progressive external ophthalmoplegia with mitochondrial DNA deletions type 1 and Mitochondrial DNA depletion syndrome 4A (Alpers-Huttenlocher syndrome) and 4B (MNGIE type)	Tremor, developmental delay, spastic diplegia, cognitive impairment, psychiatric symptoms, dysarthria, nystagmus, upward gaze paresis, blepharoptosis, ophthalmoparesis, cataracts, progressive external ophthalmoplegia, optic atrophy, dyschromatopsia, cortical blindness, migraine, seizures, peripheral neuropathy, muscle weakness and atrophy, hypogonadism, seizures, stroke-like episodes, gastroparesis, intestinal pseudo-obstruction, cardiomyopathy, hepatic dysfunction	607459, 258450, 203700, 613662	AR	None
POLR1C ¹⁸⁷	Hypomyelinating leukodystrophy type 11 (allelic with Treacher Collins syndrome type 3)	Tremor, developmental delay, intellectual disability, spasticity, myopia, dental abnormalities, head titubation	616494	AR	None
PRF1 ^{188,189}	Familial hemophagocytic lymphohistiocytosis type 2 and the allelic disorder of recurrent immune-mediated neurodegeneration	Developmental delay, failure to thrive, meningitis, encephalitis, hemiplegia, tetraplegia, seizures, coma, pancytopenia, coagulation abnormalities, lymphadenopathy, fever, edema, liver dysfunction Neurodegeneration triggered by infections, recurrent subacute post-viral onset of ataxia, primary immunodeficiency	603553, 170280	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
PRPS1 ¹⁹⁰⁻¹⁹²	Allelic disorders or continuum with Arts syndrome, X-linked recessive Charcot-Marie-Tooth disease-5 or Rosenberg-Chutorian syndrome and Hyperuricemia, mental retardation and sensorineural deafness with PRPS1 superactivity	Developmental delay, mental retardation, poor growth, sensorineural hearing loss, optic atrophy, retinopathy, nystagmus, muscle weakness, hyperreflexia, peripheral neuropathy, distal muscle weakness and atrophy, distal sensory impairment, pes cavus, flaccid tetraplegia, immune deficiency, recurrent respiratory tract infections, uric acid urolithiasis, secondary renal insufficiency, gout, gouty arthritis	301835, 300661, 311070	XR	None
PRX ¹⁹³		Distal and proximal lower limb muscle weakness and atrophy, peripheral neuropathy, pes cavus, scoliosis, delayed motor development	614895	AR	None
RARS ¹⁹⁴		Tremor, developmental delay, mental retardation, pyramidal signs, nystagmus, altered smooth pursuit, microcephaly	616140	AR	None
RELN ¹⁹⁵	Norman-Roberts type of Lissencephaly	Microcephaly, facial dysmorphism, mental retardation, nystagmus, seizures, congenital lymphedema	257320	AR	None
ROGD1 ¹⁹⁶	Kohlschutter-Tonz syndrome	Developmental delay, mental retardation, cognitive impairment, spasticity, seizures, amelogenesis imperfecta, enamel hypoplasia, discolored teeth	226750	AR	None
RRM2B ¹⁹⁷	Mitochondrial DNA depletion syndrome 8B (MNGIE type)	Failure to thrive, lactic acidosis, proximal renal tubulopathy, seizures, external ophthalmoplegia, ptosis, gastrointestinal dysmotility, cachexia, peripheral neuropathy	612075	AR	None
RTN4IP1 ¹⁹⁸	Optic atrophy type 10 with or without ataxia, mental retardation, and seizures	Mental retardation, photophobia, nystagmus, reduced visual acuity, color vision impairment of red/green axis, optic atrophy, central scotoma seizures	616732	AR	None
SLC2A1 ¹⁹⁹	Hartnup disease	Dystonia, developmental delay, seizures, myoclonic epilepsy, spasticity	606777	AR/AD	None
SLC6A19 ²⁰⁰		Delayed cognitive development, psychiatric symptoms, seizures, hypertonia, light-sensitive dermatitis, atrophic glossitis	234500	AR	None
SLC16A2 ²⁰¹	Allan-Herndon-Dudley syndrome or monocarboxylate transporter type 8 deficiency	Dystonia, developmental delay, mental retardation, pyramidal signs, amyotrophy, behavior disorders, scoliosis, nystagmus, facial dysmorphism, microcephaly, pectus excavatum	300523	XR	None
SLC19A3 ²⁰²	Thiamine metabolism dysfunction syndrome type 2 or biotin-thiamine-responsive basal ganglia disease	Dystonia, psychomotor retardation, encephalopathy, coma, psychiatric symptoms, external ophthalmoplegia, nystagmus, ptosis, gaze palsy, seizures, pyramidal signs, paraparesis, rigidity	607483	AR	None
SLC25A15 ²⁰³	Hyperornithinemia-hyperammonemia-homocitrullinuria (HHH) syndrome	Psychomotor retardation, failures to thrive, mental retardation, lethargy, episodic confusion, acute encephalopathy, coma, pyramidal signs, myoclonic epilepsy, decreased vibration sense, coagulopathy as a result of liver dysfunction	238970	AR	None
SLC25A46 ²⁰⁴	Hereditary motor and sensory neuropathy type VIIB or Charcot-Marie-Tooth disease type 6B	Myoclonus, delayed development, optic atrophy, pyramidal signs, peripheral neuropathy, distal sensory impairment, pes cavus, morphological abnormalities	616505	AR	None
SLC52A2 ²⁰⁵	Brown-Vialetto-Van Laere syndrome type 2	Cranial nerve palsies, bulbar palsy, optic atrophy, nystagmus, visual loss, absent pupillary reflex, sensorineural hearing loss, peripheral neuropathy, muscle weakness and atrophy, tongue fasciculations, psychiatric symptoms, claw hands, scoliosis, respiratory insufficiency	614707	AR	None
SNORD118 ²⁰⁶	Leukoencephalopathy, brain calcifications, and cysts	Dystonia, tremor, seizures, spasticity, hemiplegia, cognitive decline, pyramidal signs	614561	AR	None

(Continued)

TABLE 1. Continued

New designation	Alternate name	Main clinical features	OMIM	Inheritance pattern	Locus symbol
SUOX ²⁰⁷	Sulfocysteinuria or sulfite oxidase deficiency	Dystonia, developmental delay, infantile hemiplegia, seizures, behavior disorders, fine hair, eczema, delayed teething, ectopia lentis	272300	AR	None
SURF1 ²⁰⁸	Peripheral neuropathy	Distal muscle weakness and atrophy, kyphoscoliosis, nystagmus, sensorineural hearing loss	616684	AR	None
TCTN1 ²⁰⁹	Cognitive impairment, limb abnormalities	Developmental delay, absent speech, pyramidal signs, nystagmus, hyperopia, polydactyly, talipes equinovarus	614173	AR	JBTS13
TCTN2 ²¹⁰	Developmental delay	Absent speech, pyramidal signs, nystagmus, hyperopia, polydactyly, talipes equinovarus	616654	AR	JBTS24
TCTN3 ²¹¹	Mental retardation, abnormal eye movements, facial dysmorphism, scoliosis, polydactyly, camptodactyly, breathing anomalies, ventricular septal defect, horseshoe kidney	Mental retardation, abnormal eye movements, facial dysmorphism, scoliosis, polydactyly, camptodactyly, breathing anomalies, ventricular septal defect, horseshoe kidney	614815	AR	JBTS18
TRAPPC11 ²¹²	Limb-girdle muscular dystrophy type 2S	Dystonia, chorea, tremor, developmental delay, proximal muscle weakness, muscle cramps, scapular winging, scoliosis, hip dysplasia, cataracts, strabismus, myopia, microcephaly, short stature	615356	AR	None
TRNT1 ²¹³	Congenital sideroblastic anemia with B-cell immunodeficiency, periodic fevers, and developmental delay	Severe sideroblastic anemia, developmental delay, growth retardation, lactic acidosis, recurrent fevers, brittle hair, nephrocalcinosis, cardiomyopathy, retinopathy, sensorineural hearing loss, seizures	616084	AR	None
VAR2 ²¹⁴	Combined oxidative phosphorylation deficiency type 20	Developmental delay, ptosis, progressive external ophthalmoplegia, seizures, facial dysmorphism, microcephaly	615917	AR	None
VRK1 ²¹⁵	Psychomotor retardation	Mental retardation, microcephaly, nystagmus, muscle weakness, distal spinal muscular atrophy, fasciculations, peripheral neuropathy, hyperreflexia, foot deformities, skeletal contractures, arthrogyrosis, scoliosis, respiratory insufficiency	607596	AR	None
WFS1 ²¹⁶	Wolfram syndrome-1 or Diabetes insipidus and mellitus with optic atrophy and deafness	Parkinsonism, tremor, myoclonus, mental retardation or dementia, poor growth, optic atrophy, retinopathy, ptosis, nystagmus, sensorineural hearing loss, hyposmia, seizures, peripheral neuropathy, stroke-like episodes, psychiatric symptoms, diabetes mellitus and diabetes insipidus, hypothyroidism, hydronephrosis, testicular atrophy, cardiomyopathy	222300	AR	None
XRCC4 ²¹⁷	Short stature, microcephaly, and endocrine dysfunction syndrome	Intrauterine growth failure, developmental delay, cognitive impairment, apraxia, pyramidal signs, peripheral neuropathy, dyslipidemia, diabetes mellitus, hypothyroidism, anemia, acanthosis nigricans, cryptorchidism, renal dysgenesis, malpositioned teeth, facial dysmorphism, short stature, microcephaly	616541	AR	None

AR, autosomal recessive; AD, autosomal dominant; Mi, mitochondrial; OMIM, Online Mendelian Inheritance in Man; XR, X-linked recessive.

^aA complete reference list can be found in the Supplementary Material.

gene. Recessive cerebellar ataxias have also been classified according to their presumed molecular pathogenesis, such as metabolic alterations, mitochondrial dysfunction, altered calcium signaling, defective DNA repair, abnormal protein folding, and chaperone dysfunction.^{2,8,9} This is a useful approach for ataxia researchers and for future therapeutic targeting but is of limited value to clinicians. Last, more recently the field has begun to refer to recessive ataxias as SCAR's (for SCA recessive). However, this list is incomplete (only 26 entities so far) and also contains unconfirmed entities, such as SCAR11 (SYT14), SCAR12 (WVOX), SCAR19 (SLC9A1), SCAR21 (SCYL1), SCAR22 (VWA3B), SCAR23 (TDP2) and SCAR24 (UBA5), which did not meet our criteria to be part of the final list (Table 1). Moreover, this SCAR classification partly parallels and duplicates the ARCA classification, yet with different numbers. For example, ATX-SYNE1 is dubbed ARCA1 in 1 nomenclature system, but SCAR8 in the other.

A recently conducted systematic review of autosomal recessive ataxias conducted by Beaudin and colleagues¹⁰ identified 45 genetically confirmed disorders that have ataxia as a core feature. An important difference from their approach is that we included genetic entities that present with ataxia as part of a more complex phenotype. Recently, the comprehensive features of 67 recessively inherited entities that may present with ataxia were gathered within a knowledgebase that was the basis of an automated algorithm for the diagnosis of recessive cerebellar ataxias.¹¹ During the review process of our working group, we have applied strict criteria. In part related to this inclusive principle, we encountered classification difficulties for certain entries, for example, in relation to the criteria of prominent or consistent ataxia. Also, in some cases, it was difficult to distinguish purely sensory ataxias (eg, the posterior column ataxia with retinitis pigmentosa as a result of mutations in the FLVCR1 gene) from disorders that combine sensory and cerebellar ataxia, such as Friedreich ataxia, ataxia with vitamin E deficiency, or ataxia as a result of mutations in the MTTP gene (also known as abetalipoproteinemia) or in the POLG gene (sensory ataxic neuropathy, dysarthria, and ophthalmoparesis). If there was a clear cerebellar contribution to the clinical manifestation of ataxia, such an entity was included in the final list.

Prefix designation is work in progress, as prefixes may change over time when further clinical descriptions or case reports have been published. Also, for the system to be fully comprehensive, prefixes need to be defined for nonmovement disorder phenotypes. This requires other subspecialties within neurology and neurogenetics to adopt this system. We specifically designed our system with adaptability in mind. This also requires periodic

updating, which is facilitated by the availability of an online tool, the International Parkinson and Movement Disorder Society genetic mutation database (<http://www.mdsgene.org>), which is a more suitable medium than static publications of this new genetic nomenclature of the recessive cerebellar ataxias. The updated lists will also be accessible on the Task Forces section of the MDS website (<https://www.movementdisorders.org/MDS/About/Committees-Other-Groups/MDS-Task-Forces/Task-Force-on-Nomenclature-in-Movement-Disorders.htm>).

We encourage all neurologists, particularly movement disorders specialists and pediatric neurologists, and researchers in the field of ataxia to adopt and use this newly transparent and adaptable nomenclature system of the recessive cerebellar ataxias as it will facilitate the clinical recognition of numerous recessive ataxias, guide diagnostic testing in ataxia patients, and help in interpreting genetic findings. ■

Acknowledgments: MA, AD, MK, MS and BvW were supported in the frame of the E-Rare-3 network PREPARE (BMBF 01GM1607) by the European Union's Horizon 2020 research and innovation program under the ERA-NET Cofund action N° 643578.

References

1. Anheim M, Tranchant C, Koenig M. The autosomal recessive cerebellar ataxias. *N Engl J Med* 2012;366:636-646.
2. Synofzik M, Schule R. Overcoming the divide between ataxias and spastic paraplegias: shared phenotypes, genes, and pathways. *Mov Disord* 2017;32:332-345.
3. Marras C, Lohmann K, Lang A, Klein C. Fixing the broken system of genetic locus symbols: Parkinson disease and dystonia as examples. *Neurology* 2012;78:1016-1024.
4. Marras C, Lang A, van de Warrenburg BP, et al. Nomenclature of genetic movement disorders: recommendations of the International Parkinson and Movement Disorder Society Task Force. *Mov Disord* 2016;31:436-457.
5. Synofzik M, Nemeth A. The cerebellum in children and adults. In: Manto M, Huisman T, eds. *Handbook of Clinical Neurology*. Elsevier, New York, US; 2008.
6. Lill CM, Mashychev A, Hartmann C, et al. Launching the Movement Disorders Society Genetic Mutation Database (MDSGene). *Mov Disord* 2016;31:607-609.
7. Harding AE. Clinical features and classification of inherited ataxias. *Adv Neurol* 1993;61:1-14.
8. Vermeer S, van de Warrenburg BP, Willemsen MA, et al. Autosomal recessive cerebellar ataxias: the current state of affairs. *J Med Genet* 2011;48:651-659.
9. De Michele G, Coppola G, Coccozza S, Filla A. A pathogenetic classification of hereditary ataxias: is the time ripe? *J Neurol* 2004;251:913-922.
10. Beaudin M, Klein CJ, Rouleau GA, Dupre N. Systematic review of autosomal recessive ataxias and proposal for a classification. *Cerebellum Ataxias* 2017;4:3.
11. Renaud M, Tranchant C, Martin JVT, et al. A recessive ataxia diagnosis algorithm for the next generation sequencing era. *Ann Neurol* 2017;82:892-899.
12. Allikmets R, Raskind WH, Hutchinson A, Schueck ND, Dean M, Koeller DM. Mutation of a putative mitochondrial iron transporter gene (ABC7) in X-linked sideroblastic anemia and ataxia (XLSA/A). *Hum Mol Genet* 1999;8:743-749.
13. Fiskerstrand T, H'Mida-Ben Brahim D, Johansson S, et al. Mutations in ABHD12 cause the neurodegenerative disease PHARC: an inborn error of endocannabinoid metabolism. *Am J Hum Genet* 2010;87:410-417.

14. Lagier-Tourenne C, Tazir M, Lopez LC, et al. ADCK3, an ancestral kinase, is mutated in a form of recessive ataxia associated with coenzyme Q10 deficiency. *Am J Hum Genet* 2008;82:661-672.
15. Mollet J, Delahodde A, Serre V, et al. CABC1 gene mutations cause ubiquinone deficiency with cerebellar ataxia and seizures. *Am J Hum Genet* 2008;82:623-630.
16. Ferland RJ, Eyaid W, Collura RV, et al. Abnormal cerebellar development and axonal decussation due to mutations in AHI1 in Joubert syndrome. *Nat Genet* 2004;36:1008-1013.
17. Dixon-Salazar T, Silhavy JL, Marsh SE, et al. Mutations in the AHI1 gene, encoding joubertin, cause Joubert syndrome with cortical polymicrogyria. *Am J Hum Genet* 2004;75:979-987.
18. Chambliss KL, Hinson DD, Trettel F, et al. Two exon-skipping mutations as the molecular basis of succinic semialdehyde dehydrogenase deficiency (4-hydroxybutyric aciduria). *Am J Hum Genet* 1998;63:399-408.
19. Imbach T, Burda P, Kuhnert P, et al. A mutation in the human ortholog of the *Saccharomyces cerevisiae* ALG6 gene causes carbohydrate-deficient glycoprotein syndrome type-Ic. *Proc Natl Acad Sci U S A* 1999;96:6982-6987.
20. Vermeer S, Hoischen A, Meijer RP, et al. Targeted next-generation sequencing of a 12.5 Mb homozygous region reveals ANO10 mutations in patients with autosomal-recessive cerebellar ataxia. *Am J Hum Genet* 2010;87:813-819.
21. Date H, Onodera O, Tanaka H, et al. Early-onset ataxia with ocular motor apraxia and hypoalbuminemia is caused by mutations in a new HIT superfamily gene. *Nat Genet* 2001;29:184-188.
22. Moreira MC, Barbot C, Tachi N, et al. The gene mutated in ataxia-ocular apraxia 1 encodes the new HIT/Zn-finger protein aprataxin. *Nat Genet* 2001;29:189-193.
23. Cantagrel V, Silhavy JL, Bielas SL, et al. Mutations in the cilia gene ARL13B lead to the classical form of Joubert syndrome. *Am J Hum Genet* 2008;83:170-179.
24. Savitsky K, Bar-Shira A, Gilad S, et al. A single ataxia telangiectasia gene with a product similar to PI-3 kinase. *Science* 1995;268:1749-1753.
25. Nobukuni Y, Mitsubuchi H, Akaboshi I, et al. Maple syrup urine disease. Complete defect of the E1 beta subunit of the branched chain alpha-ketoacid dehydrogenase complex due to a deletion of an 11-bp repeat sequence which encodes a mitochondrial targeting leader peptide in a family with the disease. *J Clin Invest* 1991;87:1862-1866.
26. Pomponio RJ, Reynolds TR, Cole H, Buck GA, Wolf B. Mutational hotspot in the human biotinidase gene causes profound biotinidase deficiency. *Nat Genet* 1995;11:96-98.
27. Nikali K, Suomalainen A, Saharinen J, et al. Infantile onset spinocerebellar ataxia is caused by recessive mutations in mitochondrial proteins Twinkle and Twinky. *Hum Mol Genet* 2005;14:2981-2990.
28. Turkmen S, Guo G, Garshasbi M, et al. CA8 mutations cause a novel syndrome characterized by ataxia and mild mental retardation with predisposition to quadrupedal gait. *PLoS Genet* 2009;5:e1000487.
29. Sayer JA, Otto EA, O'Toole JF, et al. The centrosomal protein nephrocystin-6 is mutated in Joubert syndrome and activates transcription factor ATF4. *Nat Genet* 2006;38:674-681.
30. Valente EM, Silhavy JL, Brancati F, et al. Mutations in CEP290, which encodes a centrosomal protein, cause pleiotropic forms of Joubert syndrome. *Nat Genet* 2006;38:623-625.
31. Szklarzyk R, Wanschers BF, Nijtmans LG, et al. A mutation in the FAM36A gene, the human ortholog of COX20, impairs cytochrome c oxidase assembly and is associated with ataxia and muscle hypotonia. *Hum Mol Genet* 2013;22:656-667.
32. Burns R, Majczenko K, Xu J, et al. Homozygous splice mutation in CWF19L1 in a Turkish family with recessive ataxia syndrome. *Neurology* 2014;83:2175-2182.
33. Cali JJ, Hsieh CL, Francke U, Russell DW. Mutations in the bile acid biosynthetic enzyme sterol 27-hydroxylase underlie cerebrotendinous xanthomatosis. *J Biol Chem* 1991;266:7779-7783.
34. Davey KM, Parboosingh JS, McLeod DR, et al. Mutation of DNAJC19, a human homologue of yeast inner mitochondrial membrane co-chaperones, causes DCMA syndrome, a novel autosomal recessive Barth syndrome-like condition. *J Med Genet* 2006;43:385-393.
35. Delatycki MB, Knight M, Koenig M, Cossee M, Williamson R, Forrest SM. G130V, a common FRDA point mutation, appears to have arisen from a common founder. *Hum Genet* 1999;105:343-346.
36. Lodi R, Cooper JM, Bradley JL, et al. Deficit of in vivo mitochondrial ATP production in patients with Friedreich ataxia. *Proc Natl Acad Sci U S A* 1999;96:11492-11495.
37. Utine GE, Haliloglu G, Salanci B, et al. A homozygous deletion in GRID2 causes a human phenotype with cerebellar ataxia and atrophy. *J Child Neurol* 2013;28:926-932.
38. Smith KR, Damiano J, Franceschetti S, et al. Strikingly different clinicopathological phenotypes determined by progranulin-mutation dosage. *Am J Hum Genet* 2012;90:1102-1107.
39. Gerber S, Alzayady KJ, Burglen L, et al. Recessive and dominant de novo ITPR1 mutations cause Gillespie syndrome. *Am J Hum Genet* 2016;98:971-980.
40. Scholl UI, Choi M, Liu T, et al. Seizures, sensorineural deafness, ataxia, mental retardation, and electrolyte imbalance (SeSAME syndrome) caused by mutations in KCNJ10. *Proc Natl Acad Sci U S A* 2009;106:5842-5847.
41. Assoum M, Salih MA, Drouot N, et al. Rundataxin, a novel protein with RUN and diacylglycerol binding domains, is mutant in a new recessive ataxia. *Brain* 2010;133:2439-2447.
42. Topcu M, Jobard F, Halliez S, et al. L-2-Hydroxyglutaric aciduria: identification of a mutant gene C14orf160, localized on chromosome 14q22.1. *Hum Mol Genet* 2004;13:2803-2811.
43. Nilssen O, Berg T, Riise HM, et al. alpha-Mannosidosis: functional cloning of the lysosomal alpha-mannosidase cDNA and identification of a mutation in two affected siblings. *Hum Mol Genet* 1997;6:717-726.
44. Stewart GS, Maser RS, Stankovic T, et al. The DNA double-strand break repair gene hMRE11 is mutated in individuals with an ataxia-telangiectasia-like disorder. *Cell* 1999;99:577-587.
45. Nasca A, Scotton C, Zaharieva I, et al. Recessive mutations in MSTO1 cause mitochondrial dynamics impairment, leading to myopathy and ataxia. *Hum Mutat* 2017;38:970-977.
46. Carstea ED, Morris JA, Coleman KG, et al. Niemann-Pick C1 disease gene: homology to mediators of cholesterol homeostasis. *Science* 1997;277:228-231.
47. Naureckiene S, Sleat DE, Lackland H, et al. Identification of HE1 as the second gene of Niemann-Pick C disease. *Science* 2000;290:2298-2301.
48. Coene KL, Roepman R, Doherty D, et al. OFD1 is mutated in X-linked Joubert syndrome and interacts with LCA5-encoded lebercilin. *Am J Hum Genet* 2009;85:465-481.
49. Billuart P, Bienvenu T, Ronce N, et al. Oligophrenin-1 encodes a rhoGAP protein involved in X-linked mental retardation. *Nature* 1998;392:923-926.
50. Tuchman M, Jaleel N, Morizono H, Sheehy L, Lynch MG. Mutations and polymorphisms in the human ornithine transcarbamylase gene. *Hum Mutat* 2002;19:93-107.
51. Braverman N, Chen L, Lin P, et al. Mutation analysis of PEX7 in 60 probands with rhizomelic chondrodysplasia punctata and functional correlations of genotype with phenotype. *Hum Mutat* 2002;20:284-297.
52. Warren DS, Morrell JC, Moser HW, Valle D, Gould SJ. Identification of PEX10, the gene defective in complementation group 7 of the peroxisome-biogenesis disorders. *Am J Hum Genet* 1998;63:347-359.
53. Mihalik SJ, Morrell JC, Kim D, Sacksteder KA, Watkins PA, Gould SJ. Identification of PAHX, a Refsum disease gene. *Nat Genet* 1997;17:185-189.
54. Matthijs G, Schollen E, Pardon E, et al. Mutations in PMM2, a phosphomannomutase gene on chromosome 16p13, in carbohydrate-deficient glycoprotein type I syndrome (Jaeken syndrome). *Nat Genet* 1997;16:88-92.
55. Jobling RK, Assoum M, Gakh O, et al. PMPCA mutations cause abnormal mitochondrial protein processing in patients with non-progressive cerebellar ataxia. *Brain* 2015;138:1505-1517.
56. Saitou H, Osaka H, Sasaki M, et al. Mutations in POLR3A and POLR3B encoding RNA Polymerase III subunits cause an

- autosomal-recessive hypomyelinating leukoencephalopathy. *Am J Hum Genet* 2011;89:644-651.
57. Najmabadi H, Hu H, Garshabi M, et al. Deep sequencing reveals 50 novel genes for recessive cognitive disorders. *Nature* 2011;478:57-63.
 58. Hu H, Matter ML, Issa-Jahns L, et al. Mutations in PTRH2 cause novel infantile-onset multisystem disease with intellectual disability, microcephaly, progressive ataxia, and muscle weakness. *Ann Clin Transl Neurol* 2014;1:1024-1035.
 59. Stewart GS, Panier S, Townsend K, et al. The RIDDLE syndrome protein mediates a ubiquitin-dependent signaling cascade at sites of DNA damage. *Cell* 2009;136:420-434.
 60. Margolin DH, Kousi M, Chan YM, et al. Ataxia, dementia, and hypogonadotropism caused by disordered ubiquitination. *N Engl J Med* 2013;368:1992-2003.
 61. Moreira MC, Klur S, Watanabe M, et al. Senataxin, the ortholog of a yeast RNA helicase, is mutant in ataxia-ocular apraxia 2. *Nat Genet* 2004;36:225-227.
 62. Anttonen AK, Mahjneh I, Hamalainen RH, et al. The gene disrupted in Marinesco-Sjogren syndrome encodes SIL1, an HSPA5 cochaperone. *Nat Genet* 2005;37:1309-1311.
 63. Verheijen FW, Verbeek E, Aula N, et al. A new gene, encoding an anion transporter, is mutated in sialic acid storage diseases. *Nat Genet* 1999;23:462-465.
 64. Huppke P, Brendel C, Kalscheuer V, et al. Mutations in SLC33A1 cause a lethal autosomal-recessive disorder with congenital cataracts, hearing loss, and low serum copper and ceruloplasmin. *Am J Hum Genet* 2012;90:61-68.
 65. Foley AR, Menezes MP, Pandraud A, et al. Treatable childhood neuropathy caused by mutations in riboflavin transporter RFVT2. *Brain* 2014;137:44-56.
 66. Guissart C, Drouot N, Oncel I, et al. Genes for spinocerebellar ataxia with blindness and deafness (SCABD/SCAR3, MIM# 271250 and SCABD2). *Eur J Hum Genet* 2016;24:1154-1159.
 67. Thomas AC, Williams H, Seto-Salvia N, et al. Mutations in SNX14 cause a distinctive autosomal-recessive cerebellar ataxia and intellectual disability syndrome. *Am J Hum Genet* 2014;95:611-621.
 68. Lise S, Clarkson Y, Perkins E, et al. Recessive mutations in SPTBN2 implicate beta-III spectrin in both cognitive and motor development. *PLoS Genet* 2012;8:e1003074.
 69. Cantagrel V, Lefeber DJ, Ng BG, et al. SRD5A3 is required for converting polyprenol to dolichol and is mutated in a congenital glycosylation disorder. *Cell* 2010;142:203-217.
 70. Shi Y, Wang J, Li JD, et al. Identification of CHIP as a novel causative gene for autosomal recessive cerebellar ataxia. *PLoS One* 2013;8:e81884.
 71. Gros-Louis F, Dupre N, Dion P, et al. Mutations in SYNE1 lead to a newly discovered form of autosomal recessive cerebellar ataxia. *Nat Genet* 2007;39:80-85.
 72. Ouahchi K, Arita M, Kayden H, et al. Ataxia with isolated vitamin E deficiency is caused by mutations in the alpha-tocopherol transfer protein. *Nat Genet* 1995;9:141-145.
 73. Edvardson S, Shaag A, Zenvirt S, et al. Joubert syndrome 2 (JBTS2) in Ashkenazi Jews is associated with a TMEM216 mutation. *Am J Hum Genet* 2010;86:93-97.
 74. Brancati F, Iannicelli M, Travaglini L, et al. MKS3/TMEM67 mutations are a major cause of COACH Syndrome, a Joubert Syndrome related disorder with liver involvement. *Hum Mutat* 2009;30:E432-E442.
 75. Baala L, Romano S, Khaddour R, et al. The Meckel-Gruber syndrome gene, MKS3, is mutated in Joubert syndrome. *Am J Hum Genet* 2007;80:186-194.
 76. Doherty D, Parisi MA, Finn LS, et al. Mutations in 3 genes (MKS3, CC2D2A and RPGRIP1L) cause COACH syndrome (Joubert syndrome with congenital hepatic fibrosis). *J Med Genet* 2010;47:8-21.
 77. Arts HH, Doherty D, van Beersum SE, et al. Mutations in the gene encoding the basal body protein RPGRIP1L, a nephrocystin-4 interactor, cause Joubert syndrome. *Nat Genet* 2007;39:882-888.
 78. Delous M, Baala L, Salomon R, et al. The ciliary gene RPGRIP1L is mutated in cerebello-oculo-renal syndrome (Joubert syndrome type B) and Meckel syndrome. *Nat Genet* 2007;39:875-881.
 79. Gorden NT, Arts HH, Parisi MA, et al. CC2D2A is mutated in Joubert syndrome and interacts with the ciliopathy-associated basal body protein CEP290. *Am J Hum Genet* 2008;83:559-571.
 80. Srouf M, Hamdan FF, Schwartzentruber JA, et al. Mutations in TMEM231 cause Joubert syndrome in French Canadians. *J Med Genet* 2012;49:636-641.
 81. Ghezzi D, Arzuffi P, Zordan M, et al. Mutations in TTC19 cause mitochondrial complex III deficiency and neurological impairment in humans and flies. *Nat Genet* 2011;43:259-263.
 82. Boycott KM, Flavelle S, Bureau A, et al. Homozygous deletion of the very low density lipoprotein receptor gene causes autosomal recessive cerebellar hypoplasia with cerebral gyral simplification. *Am J Hum Genet* 2005;77:477-483.
 83. Colin E, Huynh Cong E, Mollet G, et al. Loss-of-function mutations in WDR73 are responsible for microcephaly and steroid-resistant nephrotic syndrome: Galloway-Mowat syndrome. *Am J Hum Genet* 2014;95:637-648.
 84. Gulsuner S, Tekinay AB, Doerschner K, et al. Homozygosity mapping and targeted genomic sequencing reveal the gene responsible for cerebellar hypoplasia and quadrupedal locomotion in a consanguineous kindred. *Genome Res* 2011;21:1995-2003.
 85. Pierson TM, Adams D, Bonn F, et al. Whole-exome sequencing identifies homozygous AFG3L2 mutations in a spastic ataxia-neuropathy syndrome linked to mitochondrial m-AAA proteases. *PLoS Genet* 2011;7:e1002325.
 86. Scheper GC, van der Kloek T, van Anel RJ, et al. Mitochondrial aspartyl-tRNA synthetase deficiency causes leukoencephalopathy with brain stem and spinal cord involvement and lactate elevation. *Nat Genet* 2007;39:534-539.
 87. Steinfeld R, Grapp M, Kraetzner R, et al. Folate receptor alpha defect causes cerebral folate transport deficiency: a treatable neurodegenerative disorder associated with disturbed myelin metabolism. *Am J Hum Genet* 2009;85:354-363.
 88. Uhlenberg B, Schuelke M, Ruschendorf F, et al. Mutations in the gene encoding gap junction protein alpha 12 (connexin 46.6) cause Pelizaeus-Merzbacher-like disease. *Am J Hum Genet* 2004;75:251-260.
 89. Myerowitz R, Costigan FC. The major defect in Ashkenazi Jews with Tay-Sachs disease is an insertion in the gene for the alpha-chain of beta-hexosaminidase. *J Biol Chem* 1988;263:18587-18589.
 90. O'Dowd BF, Klavins MH, Willard HF, Gravel R, Lowden JA, Mahuran DJ. Molecular heterogeneity in the infantile and juvenile forms of Sandhoff disease (O-variant GM2 gangliosidosis). *J Biol Chem* 1986;261:12680-12685.
 91. Synofzik M, Gonzalez MA, Lourenco CM, et al. PNPLA6 mutations cause Boucher-Neuhauser and Gordon Holmes syndromes as part of a broad neurodegenerative spectrum. *Brain* 2014;137:69-77.
 92. Rainier S, Bui M, Mark E, et al. Neuropathy target esterase gene mutations cause motor neuron disease. *Am J Hum Genet* 2008;82:780-785.
 93. Wiethoff S, Bettencourt C, Paudel R, et al. Pure cerebellar ataxia with homozygous mutations in the PNPLA6 gene. *Cerebellum* 2016;16:262-267.
 94. Hufnagel RB, Arno G, Hein ND, et al. Neuropathy target esterase impairments cause Oliver-McFarlane and Laurence-Moon syndromes. *J Med Genet* 2015;52:85-94.
 95. Kmoch S, Majewski J, Ramamurthy V, et al. Mutations in PNPLA6 are linked to photoreceptor degeneration and various forms of childhood blindness. *Nat Commun* 2015;6:5614.
 96. Bernard G, Chouery E, Putorti ML, et al. Mutations of POLR3A encoding a catalytic subunit of RNA polymerase Pol III cause a recessive hypomyelinating leukodystrophy. *Am J Hum Genet* 2011;89:415-423.
 97. Engert JC, Berube P, Mercier J, et al. ARSACS, a spastic ataxia common in northeastern Quebec, is caused by mutations in a new gene encoding an 11.5-kb ORF. *Nat Genet* 2000;24:120-125.
 98. Boukhris A, Schule R, Loureiro JL, et al. Alteration of ganglioside biosynthesis responsible for complex hereditary spastic paraplegia. *Am J Hum Genet* 2013;93:118-123.
 99. Gan-Or Z, Bouslam N, Birouk N, et al. Mutations in CAPN1 cause autosomal-recessive hereditary spastic paraplegia. *Am J Hum Genet* 2016;98:1038-1046.

100. Depienne C, Bugiani M, Dupuits C, et al. Brain white matter oedema due to CIC-2 chloride channel deficiency: an observational analytical study. *Lancet Neurol* 2013;12:659-668.
101. Tsaousidou MK, Ouahchi K, Warner TT, et al. Sequence alterations within CYP7B1 implicate defective cholesterol homeostasis in motor-neuron degeneration. *Am J Hum Genet* 2008;82:510-515.
102. Bomont P, Cavalier L, Blondeau F, et al. The gene encoding giganonin, a new member of the cytoskeletal BTB/kelch repeat family, is mutated in giant axonal neuropathy. *Nat Genet* 2000;26:370-374.
103. Martin E, Schule R, Smets K, et al. Loss of function of glucocerebrosidase GBA2 is responsible for motor neuron defects in hereditary spastic paraplegia. *Am J Hum Genet* 2013;92:238-244.
104. Bouslam N, Bouhouche A, Benomar A, et al. A novel locus for autosomal recessive spastic ataxia on chromosome 17p. *Hum Genet* 2007;121:413-420.
105. Dor T, Cinnamon Y, Raymond L, et al. KIF1C mutations in two families with hereditary spastic paraparesis and cerebellar dysfunction. *J Med Genet* 2014;51:137-142.
106. Leegwater PA, Yuan BQ, van der Steen J, et al. Mutations of MLC1 (KIAA0027), encoding a putative membrane protein, cause megalencephalic leukoencephalopathy with subcortical cysts. *Am J Hum Genet* 2001;68:831-838.
107. Saugier-Verber P, Munnich A, Bonneau D, et al. X-linked spastic paraplegia and Pelizaeus-Merzbacher disease are allelic disorders at the proteolipid protein locus. *Nat Genet* 1994;6:257-262.
108. Casari G, De Fusco M, Ciarmatori S, et al. Spastic paraplegia and OXPHOS impairment caused by mutations in paraplegin, a nuclear-encoded mitochondrial metalloprotease. *Cell* 1998;93:973-983.
109. Bilguvar K, Tyagi NK, Ozkara C, et al. Recessive loss of function of the neuronal ubiquitin hydrolase UCHL1 leads to early-onset progressive neurodegeneration. *Proc Natl Acad Sci U S A* 2013;110:3489-3494.
110. Edvardson S, Hama H, Shaag A, et al. Mutations in the fatty acid 2-hydroxylase gene are associated with leukodystrophy with spastic paraparesis and dystonia. *Am J Hum Genet* 2008;83:643-648.
111. Sun Y, Almomani R, Breedveld GJ, et al. Autosomal recessive spinocerebellar ataxia 7 (SCAR7) is caused by variants in TPP1, the gene involved in classic late-infantile neuronal ceroid lipofuscinosis 2 disease (CLN2 disease). *Hum Mutat* 2013;34:706-713.
112. Sleat DE, Donnelly RJ, Lackland H, et al. Association of mutations in a lysosomal protein with classical late-infantile neuronal ceroid lipofuscinosis. *Science* 1997;277:1802-1805.
113. Pennacchio LA, Lehesjoki AE, Stone NE, et al. Mutations in the gene encoding cystatin B in progressive myoclonus epilepsy (EPM1). *Science* 1996;271:1731-1734.
114. Minassian BA, Lee JR, Herbrick JA, et al. Mutations in a gene encoding a novel protein tyrosine phosphatase cause progressive myoclonus epilepsy. *Nat Genet* 1998;20:171-174.
115. Corbett MA, Schwake M, Bahlo M, et al. A mutation in the Golgi Qb-SNARE gene GOSR2 causes progressive myoclonus epilepsy with early ataxia. *Am J Hum Genet* 2011;88:657-663.
116. Van Bogaert P, Azizieh R, Desir J, et al. Mutation of a potassium channel-related gene in progressive myoclonic epilepsy. *Ann Neurol* 2007;61:579-586.
117. Bonten E, van der Spoel A, Fornerod M, Grosveld G, d'Azzo A. Characterization of human lysosomal neuraminidase defines the molecular basis of the metabolic storage disorder sialidosis. *Genes Dev* 1996;10:3156-3169.
118. Chan EM, Young EJ, Ianzano L, et al. Mutations in NHLRC1 cause progressive myoclonus epilepsy. *Nat Genet* 2003;35:125-127.
119. Bull PC, Thomas GR, Rommens JM, Forbes JR, Cox DW. The Wilson disease gene is a putative copper transporting P-type ATPase similar to the Menkes gene. *Nat Genet* 1993;5:327-337.
120. Delcourt M, Riant F, Mancini J, et al. Severe phenotypic spectrum of biallelic mutations in PRRT2 gene. *J Neurol Neurosurg Psychiatry* 2015;86:782-785.
121. Simpson MA, Cross H, Proukakis C, et al. Maspardin is mutated in mast syndrome, a complicated form of hereditary spastic paraplegia associated with dementia. *Am J Hum Genet* 2003;73:1147-1156.
122. Schuur-Hoeijmakers JH, Geraghty MT, Kamsteeg EJ, et al. Mutations in DDHD2, encoding an intracellular phospholipase A(1), cause a recessive form of complex hereditary spastic paraplegia. *Am J Hum Genet* 2012;91:1073-1081.
123. Slabicki M, Theis M, Krastev DB, et al. A genome-scale DNA repair RNAi screen identifies SPG48 as a novel gene associated with hereditary spastic paraplegia. *PLoS Biol* 2010;8:e1000408.
124. Stevanin G, Santorelli FM, Azzedine H, et al. Mutations in SPG11, encoding spatacsin, are a major cause of spastic paraplegia with thin corpus callosum. *Nat Genet* 2007;39:366-372.
125. Erlich Y, Edvardson S, Hodges E, et al. Exome sequencing and disease-network analysis of a single family implicate a mutation in KIF1A in hereditary spastic paraparesis. *Genome Res* 2011;21:658-664.
126. Hanein S, Martin E, Boukhris A, et al. Identification of the SPG15 gene, encoding spastizin, as a frequent cause of complicated autosomal-recessive spastic paraplegia, including Kjellin syndrome. *Am J Hum Genet* 2008;82:992-1002.
127. Savukoski M, Klockars T, Holmberg V, Santavuori P, Lander ES, Peltonen L. CLN5, a novel gene encoding a putative transmembrane protein mutated in Finnish variant late infantile neuronal ceroid lipofuscinosis. *Nat Genet* 1998;19:286-288.
128. Arsov T, Smith KR, Damiano J, et al. Kufs disease, the major adult form of neuronal ceroid lipofuscinosis, caused by mutations in CLN6. *Am J Hum Genet* 2011;88:566-573.
129. Berkovic SF, Dibbens LM, Oshlack A, et al. Array-based gene discovery with three unrelated subjects shows SCARB2/LIMP-2 deficiency causes myoclonus epilepsy and glomerulosclerosis. *Am J Hum Genet* 2008;82:673-684.
130. Nishimoto J, Nanba E, Inui K, Okada S, Suzuki K. GM1-gangliosidosis (genetic beta-galactosidase deficiency): identification of four mutations in different clinical phenotypes among Japanese patients. *Am J Hum Genet* 1991;49:566-574.
131. Bonafe L, Thony B, Penzien JM, Czarnecki B, Blau N. Mutations in the sepiapterin reductase gene cause a novel tetrahydrobiopterin-dependent monoamine-neurotransmitter deficiency without hyperphenylalaninemia. *Am J Hum Genet* 2001;69:269-277.
132. Harris ZL, Takahashi Y, Miyajima H, Serizawa M, MacGillivray RT, Gitlin JD. Aceruloplasminemia: molecular characterization of this disorder of iron metabolism. *Proc Natl Acad Sci U S A* 1995;92:2539-2543.
133. Zhou B, Westaway SK, Levinson B, Johnson MA, Gitschier J, Hayflick SJ. A novel pantothenate kinase gene (PANK2) is defective in Hallervorden-Spatz syndrome. *Nat Genet* 2001;28:345-349.
134. Morgan NV, Westaway SK, Morton JE, et al. PLA2G6, encoding a phospholipase A2, is mutated in neurodegenerative disorders with high brain iron. *Nat Genet* 2006;38:752-754.
135. Tullio-Pelet A, Salomon R, Hadj-Rabia S, et al. Mutant WD-repeat protein in triple-A syndrome. *Nat Genet* 2000;26:332-335.
136. Dallabona C, Diodato D, Kevelam SH, et al. Novel (ovario) leukodystrophy related to AARS2 mutations. *Neurology* 2014;82:2063-2071.
137. Kemp S, Pujol A, Waterham HR, et al. ABCD1 mutations and the X-linked adrenoleukodystrophy mutation database: role in diagnosis and clinical correlations. *Hum Mutat* 2001;18:499-515.
138. Stromme P, Mangelsdorf ME, Shaw MA, et al. Mutations in the human ortholog of Aristaless cause X-linked mental retardation and epilepsy. *Nat Genet* 2002;30:441-445.
139. Desai R, Frazier AE, Durigon R, et al. ATAD3 gene cluster deletions cause cerebellar dysfunction associated with altered mitochondrial DNA and cholesterol metabolism. *Brain* 2017;140:1595-1610.
140. Harel T, Yoon WH, Garone C, et al. Recurrent de novo and biallelic variation of ATAD3A, encoding a mitochondrial membrane protein, results in distinct neurological syndromes. *Am J Hum Genet* 2016;99:831-845.
141. Ijlst L, Loupatty FJ, Ruiten JP, Duran M, Lehnert W, Wanders RJ. 3-Methylglutaconic aciduria type I is caused by mutations in AUH. *Am J Hum Genet* 2002;71:1463-1466.

142. DiMauro S, De Vivo DC. Genetic heterogeneity in Leigh syndrome. *Ann Neurol* 1996;40:5-7.
143. Srour M, Schwartzentruber J, Hamdan FF, et al. Mutations in CSORF42 cause Joubert syndrome in the French Canadian population. *Am J Hum Genet* 2012;90:693-700.
144. Anderson BH, Kasher PR, Mayer J, et al. Mutations in CTC1, encoding conserved telomere maintenance component 1, cause Coats plus. *Nat Genet* 2012;44:338-342.
145. Varon R, Gooding R, Steglich C, et al. Partial deficiency of the C-terminal-domain phosphatase of RNA polymerase II is associated with congenital cataracts facial dysmorphism neuropathy syndrome. *Nat Genet* 2003;35:185-189.
146. Zhou XY, van der Spoel A, Rottier R, et al. Molecular and biochemical analysis of protective protein/cathepsin A mutations: correlation with clinical severity in galactosialidosis. *Hum Mol Genet* 1996;5:1977-1987.
147. Smith KR, Dahl HH, Canafoglia L, et al. Cathepsin F mutations cause type B Kufs disease, an adult-onset neuronal ceroid lipofuscinosis. *Hum Mol Genet* 2013;22:1417-1423.
148. Tarpey PS, Raymond FL, O'Meara S, et al. Mutations in CUL4B, which encodes a ubiquitin E3 ligase subunit, cause an X-linked mental retardation syndrome associated with aggressive outbursts, seizures, relative macrocephaly, central obesity, hypogonadism, pes cavus, and tremor. *Am J Hum Genet* 2007;80:345-352.
149. Heiss NS, Knight SW, Vulliamy TJ, et al. X-linked dyskeratosis congenita is caused by mutations in a highly conserved gene with putative nucleolar functions. *Nat Genet* 1998;19:32-38.
150. Liu TC, Kim H, Arizmendi C, Kitano A, Patel MS. Identification of two missense mutations in a dihydrolipoamide dehydrogenase-deficient patient. *Proc Natl Acad Sci U S A* 1993;90:5186-5190.
151. Leegwater PA, Vermeulen G, Konst AA, et al. Subunits of the translation initiation factor eIF2B are mutant in leukoencephalopathy with vanishing white matter. *Nat Genet* 2001;29:383-388.
152. Niedernhofer LJ, Garinis GA, Raams A, et al. A new progeroid syndrome reveals that genotoxic stress suppresses the somatotropic axis. *Nature* 2006;444:1038-1043.
153. Sijbers AM, de Laat WL, Ariza RR, et al. Xeroderma pigmentosum group F caused by a defect in a structure-specific DNA repair endonuclease. *Cell* 1996;86:811-822.
154. Wan J, Yourshaw M, Mamsa H, et al. Mutations in the RNA exosome component gene EXOSC3 cause pontocerebellar hypoplasia and spinal motor neuron degeneration. *Nat Genet* 2012;44:704-708.
155. Sakai N, Inui K, Fujii N, et al. Krabbe disease: isolation and characterization of a full-length cDNA for human galactocerebrosidase. *Biochem Biophys Res Commun* 1994;198:485-491.
156. Dahl N, Lagerstrom M, Erikson A, Pettersson U. Gaucher disease type III (Norrbotnian type) is caused by a single mutation in exon 10 of the glucocerebrosidase gene. *Am J Hum Genet* 1990;47:275-278.
157. Orthmann-Murphy JL, Salsano E, Abrams CK, et al. Hereditary spastic paraplegia is a novel phenotype for GJA12/GJC2 mutations. *Brain* 2009;132:426-438.
158. Piao X, Hill RS, Bodell A, et al. G protein-coupled receptor-dependent development of human frontal cortex. *Science* 2004;303:2033-2036.
159. Lopez-Hernandez T, Ridder MC, Montolio M, et al. Mutant GlialCAM causes megalencephalic leukoencephalopathy with subcortical cysts, benign familial macrocephaly, and macrocephaly with retardation and autism. *Am J Hum Genet* 2011;88:422-432.
160. Loupatty FJ, Clayton PT, Ruitter JP, et al. Mutations in the gene encoding 3-hydroxyisobutyryl-CoA hydrolase results in progressive infantile neurodegeneration. *Am J Hum Genet* 2007;80:195-199.
161. Pierce SB, Walsh T, Chisholm KM, et al. Mutations in the DBP-deficiency protein HSD17B4 cause ovarian dysgenesis, hearing loss, and ataxia of Perrault Syndrome. *Am J Hum Genet* 2010;87:282-288.
162. Mootha VK, Lepage P, Miller K, et al. Identification of a gene causing human cytochrome c oxidase deficiency by integrative genomics. *Proc Natl Acad Sci U S A* 2003;100:605-610.
163. Barbosa MD, Barrat FJ, Tchernev VT, et al. Identification of mutations in two major mRNA isoforms of the Chediak-Higashi syndrome gene in human and mouse. *Hum Mol Genet* 1997;6:1091-1098.
164. Novarino G, Fenstermaker AG, Zaki MS, et al. Exome sequencing links corticospinal motor neuron disease to common neurodegenerative disorders. *Science* 2014;343:506-511.
165. Meins M, Lehmann J, Gerresheim F, et al. Submicroscopic duplication in Xq28 causes increased expression of the MECP2 gene in a boy with severe mental retardation and features of Rett syndrome. *J Med Genet* 2005;42:e12.
166. Siintola E, Topcu M, Aula N, et al. The novel neuronal ceroid lipofuscinosis gene MFSD8 encodes a putative lysosomal transporter. *Am J Hum Genet* 2007;81:136-146.
167. Romani M, Micalizzi A, Kraoua I, et al. Mutations in B9D1 and MKS1 cause mild Joubert syndrome: expanding the genetic overlap with the lethal ciliopathy Meckel syndrome. *Orphanet J Rare Dis* 2014;9:72.
168. Lerner-Ellis JP, Tirone JC, Pawelek PD, et al. Identification of the gene responsible for methylmalonic aciduria and homocystinuria, cblC type. *Nat Genet* 2006;38:93-100.
169. Spinazzola A, Santer R, Akman OH, et al. Hepatocerebral form of mitochondrial DNA depletion syndrome: novel MPV17 mutations. *Arch Neurol* 2008;65:1108-1113.
170. Tucker EJ, Hershman SG, Kohrer C, et al. Mutations in MTFMT underlie a human disorder of formylation causing impaired mitochondrial translation. *Cell Metab* 2011;14:428-434.
171. Shoulders CC, Brett DJ, Bayliss JD, et al. Abetalipoproteinemia is caused by defects of the gene encoding the 97 kDa subunit of a microsomal triglyceride transfer protein. *Hum Mol Genet* 1993;2:2109-2116.
172. Prietsch V, Mayatepek E, Krastel H, et al. Mevalonate kinase deficiency: enlarging the clinical and biochemical spectrum. *Pediatrics* 2003;111:258-261.
173. Parisi MA, Bennett CL, Eckert ML, et al. The NPHP1 gene deletion associated with juvenile nephronophthisis is present in a subset of individuals with Joubert syndrome. *Am J Hum Genet* 2004;75:82-91.
174. Calvo SE, Tucker EJ, Compton AG, et al. High-throughput, pooled sequencing identifies mutations in NUBPL and FOXRED1 in human complex I deficiency. *Nat Genet* 2010;42:851-858.
175. Bonneau D, Colin E, Oca F, et al. Early-onset Behr syndrome due to compound heterozygous mutations in OPA1. *Brain* 2014;137:e301.
176. Anikster Y, Kleta R, Shaag A, Gahl WA, Elpeleg O. Type III 3-methylglutaconic aciduria (optic atrophy plus syndrome, or Costeff optic atrophy syndrome): identification of the OPA3 gene and its founder mutation in Iraqi Jews. *Am J Hum Genet* 2001;69:1218-1224.
177. Aral B, Benelli C, Ait-Ghezala G, et al. Mutations in PDX1, the human lipoyl-containing component X of the pyruvate dehydrogenase-complex gene on chromosome 11p1, in congenital lactic acidosis. *Am J Hum Genet* 1997;61:1318-1326.
178. Shimozawa N, Imamura A, Zhang Z, et al. Defective PEX gene products correlate with the protein import, biochemical abnormalities, and phenotypic heterogeneity in peroxisome biogenesis disorders. *J Med Genet* 1999;36:779-781.
179. Bras J, Alonso I, Barbot C, et al. Mutations in PNKP cause recessive ataxia with oculomotor apraxia type 4. *Am J Hum Genet* 2015;96:474-479.
180. Shen J, Gilmore EC, Marshall CA, et al. Mutations in PNKP cause microcephaly, seizures and defects in DNA repair. *Nat Genet* 2010;42:245-249.
181. Pedrosa JL, Rocha CR, Macedo-Souza LI, et al. Mutation in PNKP presenting initially as axonal Charcot-Marie-Tooth disease. *Neurol Genet* 2015;1:e30.
182. Williams SR, Gekeler V, McIvor RS, Martin DW Jr. A human purine nucleoside phosphorylase deficiency caused by a single base change. *J Biol Chem* 1987;262:2332-2338.
183. Van Goethem G, Martin JJ, Dermaut B, et al. Recessive POLG mutations presenting with sensory and ataxic neuropathy in compound heterozygote patients with progressive external ophthalmoplegia. *Neuromuscul Disord* 2003;13:133-142.
184. Van Goethem G, Dermaut B, Lofgren A, Martin JJ, Van Broeckhoven C. Mutation of POLG is associated with progressive

- external ophthalmoplegia characterized by mtDNA deletions. *Nat Genet* 2001;28:211-212.
185. Naviaux RK, Nguyen KV. POLG mutations associated with Alpers' syndrome and mitochondrial DNA depletion. *Ann Neurol* 2004;55:706-712.
 186. Van Goethem G, Schwartz M, Lofgren A, Dermaut B, Van Broeckhoven C, Vissing J. Novel POLG mutations in progressive external ophthalmoplegia mimicking mitochondrial neurogastrointestinal encephalomyopathy. *Eur J Hum Genet* 2003;11:547-549.
 187. Thiffault I, Wolf NI, Forget D, et al. Recessive mutations in POLR1C cause a leukodystrophy by impairing biogenesis of RNA polymerase III. *Nat Commun* 2015;6:7623.
 188. Stepp SE, Dufourcq-Lagelouse R, Le Deist F, et al. Perforin gene defects in familial hemophagocytic lymphohistiocytosis. *Science* 1999;286:1957-1959.
 189. Dias C, McDonald A, Sincan M, et al. Recurrent subacute post-viral onset of ataxia associated with a PRF1 mutation. *Eur J Hum Genet* 2013;21:1232-1239.
 190. de Brouwer AP, Williams KL, Duley JA, et al. Arts syndrome is caused by loss-of-function mutations in PRPS1. *Am J Hum Genet* 2007;81:507-518.
 191. Roessler BJ, Nosal JM, Smith PR, et al. Human X-linked phosphoribosylpyrophosphate synthetase superactivity is associated with distinct point mutations in the PRPS1 gene. *J Biol Chem* 1993;268:26476-26481.
 192. Kim HJ, Sohn KM, Shy ME, et al. Mutations in PRPS1, which encodes the phosphoribosyl pyrophosphate synthetase enzyme critical for nucleotide biosynthesis, cause hereditary peripheral neuropathy with hearing loss and optic neuropathy (cmtx5). *Am J Hum Genet* 2007;81:552-558.
 193. Guilbot A, Williams A, Ravise N, et al. A mutation in periaxin is responsible for CMT4F, an autosomal recessive form of Charcot-Marie-Tooth disease. *Hum Mol Genet* 2001;10:415-421.
 194. Wolf NI, Salomons GS, Rodenburg RJ, et al. Mutations in RARS cause hypomyelination. *Ann Neurol* 2014;76:134-139.
 195. Hong SE, Shugart YY, Huang DT, et al. Autosomal recessive lissencephaly with cerebellar hypoplasia is associated with human RELN mutations. *Nat Genet* 2000;26:93-96.
 196. Schossig A, Wolf NI, Fischer C, et al. Mutations in ROGDI cause Kohlschütter-Tonz syndrome. *Am J Hum Genet* 2012;90:701-707.
 197. Bourdon A, Minai L, Serre V, et al. Mutation of RRM2B, encoding p53-controlled ribonucleotide reductase (p53R2), causes severe mitochondrial DNA depletion. *Nat Genet* 2007;39:776-780.
 198. Angebault C, Guichet PO, Talmar-Amar Y, et al. Recessive mutations in RTN4IP1 cause isolated and syndromic optic neuropathies. *Am J Hum Genet* 2015;97:754-760.
 199. Seidner G, Alvarez MG, Yeh JI, et al. GLUT-1 deficiency syndrome caused by haploinsufficiency of the blood-brain barrier hexose carrier. *Nat Genet* 1998;18:188-191.
 200. Kleta R, Romeo E, Ristic Z, et al. Mutations in SLC6A19, encoding B0AT1, cause Hartnup disorder. *Nat Genet* 2004;36:999-1002.
 201. Dumitrescu AM, Liao XH, Best TB, Brockmann K, Refetoff S. A novel syndrome combining thyroid and neurological abnormalities is associated with mutations in a monocarboxylate transporter gene. *Am J Hum Genet* 2004;74:168-175.
 202. Gerards M, Kamps R, van Oevelen J, et al. Exome sequencing reveals a novel Moroccan founder mutation in SLC19A3 as a new cause of early-childhood fatal Leigh syndrome. *Brain* 2013;136:882-890.
 203. Miyamoto T, Kanazawa N, Kato S, et al. Diagnosis of Japanese patients with HHH syndrome by molecular genetic analysis: a common mutation, R179X. *J Hum Genet* 2001;46:260-262.
 204. Abrams AJ, Hufnagel RB, Rebelo A, et al. Mutations in SLC25A46, encoding a UGO1-like protein, cause an optic atrophy spectrum disorder. *Nat Genet* 2015;47:926-932.
 205. Johnson JO, Gibbs JR, Megarbane A, et al. Exome sequencing reveals riboflavin transporter mutations as a cause of motor neuron disease. *Brain* 2012;135:2875-2882.
 206. Jenkinson EM, Rodero MP, Kasher PR, et al. Mutations in SNORD118 cause the cerebral microangiopathy leukoencephalopathy with calcifications and cysts. *Nat Genet* 2016;48:1185-1192.
 207. Kisker C, Schindelin H, Pacheco A, et al. Molecular basis of sulfite oxidase deficiency from the structure of sulfite oxidase. *Cell* 1997;91:973-983.
 208. Echaniz-Laguna A, Ghezzi D, Chassagne M, et al. SURF1 deficiency causes demyelinating Charcot-Marie-Tooth disease. *Neurology* 2013;81:1523-1530.
 209. Garcia-Gonzalo FR, Corbit KC, Siererol-Piquer MS, et al. A transition zone complex regulates mammalian ciliogenesis and ciliary membrane composition. *Nat Genet* 2011;43:776-784.
 210. Sang L, Miller JJ, Corbit KC, et al. Mapping the NPHP-JBTS-MKS protein network reveals ciliopathy disease genes and pathways. *Cell* 2011;145:513-528.
 211. Thomas S, Legendre M, Saunier S, et al. TCTN3 mutations cause Mohr-Majewski syndrome. *Am J Hum Genet* 2012;91:372-378.
 212. Bogershausen N, Shahrzad N, Chong JX, et al. Recessive TRAPPC11 mutations cause a disease spectrum of limb girdle muscular dystrophy and myopathy with movement disorder and intellectual disability. *Am J Hum Genet* 2013;93:181-190.
 213. Chakraborty PK, Schmitz-Abe K, Kennedy EK, et al. Mutations in TRNT1 cause congenital sideroblastic anemia with immunodeficiency, fevers, and developmental delay (SIFD). *Blood* 2014;124:2867-2871.
 214. Taylor RW, Pyle A, Griffin H, et al. Use of whole-exome sequencing to determine the genetic basis of multiple mitochondrial respiratory chain complex deficiencies. *JAMA* 2014;312:68-77.
 215. Renbaum P, Kellerman E, Jaron R, et al. Spinal muscular atrophy with pontocerebellar hypoplasia is caused by a mutation in the VRK1 gene. *Am J Hum Genet* 2009;85:281-289.
 216. Strom TM, Hortnagel K, Hofmann S, et al. Diabetes insipidus, diabetes mellitus, optic atrophy and deafness (DIDMOAD) caused by mutations in a novel gene (wolframin) coding for a predicted transmembrane protein. *Hum Mol Genet* 1998;7:2021-2028.
 217. Shaheen R, Faqeh E, Ansari S, et al. Genomic analysis of primordial dwarfism reveals novel disease genes. *Genome Res* 2014;24:291-299.

Supporting Data

Additional Supporting Information may be found in the online version of this article at the publisher's website.