CASE REPORT

NDUFS8-related Complex I Deficiency Extends Phenotype from "PEO Plus" to Leigh Syndrome

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Abstract With over 1,000 nuclear genes that could potentially cause a mitochondrial disorder, the current diagnostic approach requires targeted molecular analysis, guided by a combination of clinical and biochemical features. However, the expanding molecular and clinical spectrum means that this approach does not always yield a result. Here we report the unusual clinical presentation of "Progressive External Ophthalmoplegia (PEO) plus" Leigh syndrome in three children from a consanguineous family where exome sequencing identified mutations in NDUFS8. NDUFS8 is a nuclearencoded structural core protein of complex I, and mutations are expected to cause infantile onset and severe disease. Our patients had a later onset, milder and a clinically distinct phenotype, and this gene would not normally be considered in this context. Being untargeted to specific genes, whole exome analysis has the potential to re-write the phenotype and reveal an unexpected molecular aetiology, as illustrated by this family.

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Introduction

Mitochondrial complex I (NADH ubiquinone oxidoreductase) is the largest enzyme of the mitochondrial respiratory chain and catalyzes the transfer of electrons from NADH through the respiratory chain (Distelmaier et al. 2009; Pagniez-Mammeri et al. 2012). It is composed of 37 nuclear-encoded and 7 mitochondrial-encoded subunits, based on a recent study, which showed that the NDUFA4 subunit is actually a subunit of complex IV (Balsa et al. 2012), and requires several additional proteins for its assembly. Mutations in many of these genes were reported in patients with different clinical presentations associated with isolated complex I deficiency, which is the most frequently encountered defect, causing ~25–35 % of all disorders of the mitochondrial energy metabolism (Distelmaier et al. 2009; Pagniez-Mammeri et al. 2012; Nouws et al. 2012).

Mutations in the mitochondrial subunit genes (MTND1-MTND6) of complex I cause maternally inherited disease with variable clinical presentations such as Leber's hereditary optic neuropathy, mitochondrial myopathy, encephalopathy, lactic acidosis and stroke-like episodes or Leigh syndrome. More frequent clinical presentations of nuclear gene defects are Leigh syndrome (25 %), neonatal cardiomyopathy with lactic acidosis, leukoencephalopathies or other undefined encephalomyopathies, which follow Mendelian inheritance. In general, mutations in assembly factor genes result in later onset and milder phenotypes compared to deficiencies of nuclear-encoded structural complex I subunit genes (Nouws et al. 2012). The number of nuclear mutations responsible for isolated complex I deficiency is increasing rapidly, which was recently facilitated by the new high-throughput technologies. However, thorough clinical reports on patients are still lacking (Haack et al. 2012; Calvo et al. 2010).

Here we report three children with a mild and variable clinical presentation due to a homozygous mutation in a nuclear complex I structural subunit gene, a totally unexpected finding detected through non-targeted whole exome sequencing.

Case Reports

Patient 1

The now 13-year-old index patient (Fig. 1a) is the first of three children of consanguineous Afghan parents (first cousins). Our index patient was born at term after uneventful pregnancy. He exhibited generalized muscular hypotonia from infancy with delayed motor milestones (sitting at 16 months, walking at 24 months, frequent falls). His speech development was delayed (first words at age 24 months). Facial hypotonia and mild ptosis were also noted.

From 5 years of age he suffered from a slowly progressive muscle weakness, and developed dysarthria and an ataxic gait. There was a marked worsening of his symptoms, especially of muscle strength and respiratory function, frequently caused by infections. Currently, at the age of 13 years he is able to walk only a few steps independently. He complains of pain after physical exercise and for longer distances he uses a wheelchair.

On clinical examination his height was on the 3rd percentile (144 cm) and weight 2 kg below the 3rd percentile (31.2 kg). He showed facial hypotonia, mild ptosis, external ophthalmoplegia (Fig. 1a) and had intermittent hypersalivation and difficulties in swallowing. He had atrophic muscles with thoracolumbal scoliosis and contractures; deep tendon reflexes were reduced. Further symptoms were dystonic posturing, decelerated movements and an ataxic gait. He had a reduced intelligence quotient (IQ) 61 (Snijders Oomen nonverbal Test for Intelligence), was not able to read or write apart from his own name and attends special schooling.

Laboratory measurements including creatine kinase (CK) were normal, but serum lactate was mildly elevated (23.5 mg/dl, normal<19.8 mg/dl). Echocardiography detected a borderline left ventricular dysfunction (fractional shortening 25.6 %); electrocardiogram was normal.

Cranial magnetic resonance imaging (MRI) at 5 years of age showed bilateral increased signals in the putamen. MRI at age 9 years showed increased signals in the bilateral putamen, nucleus caudatus and frontal subcortical region (Fig. 1b), consistent with mitochondrial encephalopathy (Leigh syndrome). Substitution with creatine and riboflavin was initiated at 10 years of age, with no clearly demonstrable effect.

Patient 2

The younger brother of the index patient had a normal motor and mental development until 5 years of age, when he developed dysarthria. Severe myopia was diagnosed at 6 years of age, followed by slowly progressive muscle weakness leading to frequent falls. Currently, at 12 years of age (Fig. 1a), his weight is on 25th percentile (35 kg) and his height on 75th percentile (154 cm). He showed mild proximal muscular hypotonia with difficulties in uprising from a squatting position. His exercise tolerance was reduced, and he climbed stairs by holding on to the hand rail. He also complained of pain after exercise. His symptoms worsen during febrile infections.

Clinical examination revealed mild ptosis and ophthalmoplegia with intermittent horizontal nystagmus and dysarthria, but normal deep tendon reflexes. He had mild cognitive impairment (Kaufman-Assessment Battery for Children: IQ 73); he attends a school for handicapped children. Cranial MRI at 8 years of age showed abnormal signal in the bilateral putamen (Fig. 1b). Echocardiography detected borderline left ventricular dysfunction (fractional shortening 25.6 %); electrocardiogram was normal, as well as his laboratory findings including CK and lactate.

Patient 3

The third child of the family is a girl, currently 9 years old. She reached her motor and mental milestones age appropriately. At 7 years of age she had mild generalized muscular hypotonia, scapular winging and mild knee contractures. Deep tendon reflexes were normal. She had difficulties in rising from a squatting position, but her walking distance was unlimited. She had myopia, intermittent nystagmus and dysmetria, but no gait ataxia. She had normal cognitive development until 9 years of age (Kaufman-Assessment Battery for Children: IQ 84), and she had been visiting a mainstream primary school from age 7. Because of mild symptoms, no laboratory findings or MRI were performed. Echocardiography and electrocardiogram were normal.

Results

Muscle biopsy of patient 1 at 5 years of age detected nonspecific variation of fibre size, no ragged-fed fibres in the Gomori-Trichrome stain and enzyme histochemistry for NADH-CoQ-Oxidoreductase (NADH), cytochrome c



						Number of	homozygous po	ossibly pathoge	nic mutations
Patient	Variant Type	Total predicted (*)	Homozygous on Target (a)	1000 Genomes & dbSNP135 (b)	Homozygous rare/novel Variants (c)	Homozy- gous	Homozygous protein altering (d)	Homozygous predicted disease causing (e)	Homozygous mtochondrial
Patient 1	SBV	362,804	23,077	22,598	165				
Fallent	Indel	29,731	1,351	986	43	60	20	15	1
Patient 2	SBV	333,390	22,324	21,926	140	02	30	15	
	Indel	28,893	1,221	902	26				



Fig. 1 (a) The picture shows patient 1 (13 years) and patient 2 (12 years). Note external ophthalmoplegia on upgaze in patient 1. (b) Brain MRI of patient 1 at 9 years of age showed hyperintense lesions in the bilateral putamen and nucleus caudatus as well as in

the bilateral frontal subcortical region on T2w sequences without enhancement after gadolinium. The lesions in the bilateral putamen and nucleus caudatus were hypointense on T1 images. There was a slight ventricular asymmetry, but the configuration of the oxidase (COX) and succinate dehydrogenase (SDH) revealed no abnormalities (data not shown).

Biochemical analysis of the respiratory chain enzymes in *muscle* was done by standard methods and showed a reduction of complex I activity, NADH-CoQ-Oxidoreductase (0.05U/U Citrate Synthase, normal range: 0.17–0.56). The activities of complex II, III and IV were normal.

Genetic analyses: mitochondrial DNA deletions, mtDNA depletion and point mutations were excluded by standard methods. Genetic analysis of *POLG* was negative.

Whole exome sequencing was performed as described previously in patient 1 and 2 (Horvath et al. 2012). We detected 62 novel homozygous changes shared in both patients, 30 were protein altering and 15 of these were predicted to be disease causing. Only one of these homozygous changes was in a mitochondrial complex I related gene, in *NDUFS8* (Fig. 1c). The mutation c.160C>T, p.Arg54Trp (Fig. 1d) is predicted to be disease causing (Mutation Taster) (De Sury et al. 1998) and was absent in 222 control chromosomes. This mutation cosegregated within the family as determined by Sanger sequencing, it was found heterozygous in the healthy parents and homozygous in all three affected siblings.

Discussion

Although mutations in several nuclear and mitochondrial genes have been reported in association with complex I deficiencies, there are few reports describing clear genotype-phenotype correlations to direct diagnostic genetic testing (Tuppen et al. 2010). The patients reported here have – unlike deficiencies of structural nuclear complex I subunit genes - a subtle onset, slowly progressive neuromuscular condition, with ptosis, eye movement abnormalities, exercise intolerance, muscle weakness and pain, dysarthria and ataxia, resembling "progressive external ophthalmoplegia (PEO) plus". Cognitive development was variable, although differences in the younger siblings could be explained by an earlier phase of the disease. Brain MRI showed Leigh syndrome and isolated complex I deficiency was detected in skeletal muscle of patient 1, pointing to a mitochondrial disease. The high number of nuclear genes involved in the structure and assembly of complex I and the lack of genotype/phenotype data

prompted us to perform whole exome sequencing. We detected 15 possible disease causing homozygous changes in the two affected patients, but only c.160C>T, p.Arg54Trp altered a complex I related gene, *NDUFS8*. This mutation is most likely causative for the disease in our patients, since it co-segregates with the symptoms, and alters a highly conserved amino acid, but does not disrupt the [4Fe-4S] cluster, and thus may result in some residual protein function.

By searching the literature we identified eight patients carrying NDUFS8 mutations to date (Table 1); however, the published clinical information was very brief in five cases (Haack et al. 2012; Calvo et al. 2010; Tuppen et al. 2010; Procaccio et al. 2004; Loeffen et al. 1998). Some patients had a severe disease with feeding difficulties, respiratory problems, epilepsy and hypertrophic cardiomyopathy and died within the first months of life (Tuppen et al. 2010; Loeffen et al. 1998). Others, including our patients show normal or mildly impaired motor development in the first years of life and develop a slowly progressive neurological disease at the end of the first decade (Procaccio and Wallace 2004, this paper). All reported mutations are missense mutations (Table 1, Fig. 1d), and no patients are reported to date with nonsense mutations in NDUFS8, suggesting that a complete absence of this protein may have deleterious effect leading to intrauterine lethality. Relatively mild phenotypes associated with Leigh syndrome are not so uncommon in mitochondrial disorders due to mutations in some nuclear assembly genes, as reported in patients with mutations in SURF1 (Piekutowska-Abramczuk et al. 2009).

Defining the molecular basis of human diseases is important for genetic counselling and for prenatal genetic testing. However, limited clinical data on new and existing disease genes makes it difficult to know which genes to target. More detailed clinical descriptions of patients carrying mutations in different genes would be highly valuable for clinicians, who need to diagnose and counsel patients with mitochondrial disease. This will be facilitated by next generation sequencing of either large panels of relevant genes (Calvo et al. 2010), whole exome (Haack et al. 2012) and ultimately whole genome analysis. Our chapter shows the success of this approach.

altering and 15 of these were predicted to be disease causing. Only one of these homozygous changes was in a gene encoding a known mitochondrial protein, NDUFS8, a complex I related gene. (d) Localization of the c.160C>T, pArg54Trp mutation in the *NDUFS8* gene. 4Fe-4S refers to the iron-sulphur binding domains

Fig. 1 (continued) cerebellum and brainstem were normal. Brain MRI of patient 2 at 8 years of age showed an abnormal symmetrical signal in the bilateral putamen (hypointense on T1 images, hyperintense on T2 images) but no other malformations of the brain structures. (c) Exome sequencing detected 62 novel homozygous changes shared in both patients; 30 were protein

Reference	Onset	Age/death	Clinical presentation	Brain MRI	Muscle biopsy	RC in cell lines	Mutation
Previously descri	bed patient:	S					
Loeffen et al. 1998	5 weeks	Died at 11 weeks	Poor feeding, apnea, cyanosis, spasticity, seizures, hypertrophic cardiomyopathy	TS	Histology n.d. complex I↓ (39% of normal)	Fibroblasts complex I↓ (69% of normal)	c.236C>T/p.Pro79Leu c.305G>A/p.Arg102His compound heterozygous
Procaccio and Wallace 2004	7 years	Alive at 9 years	Toe walking, nystagmus, dysarthria, dystonia, ataxic gait	LS	Histology normcomplex I↓ (31% of normal)	Lymphoblast complex I↓ (43% of normal)	c.254C>T/p.Pro85Leu c.413G>A/p.Arg138His compound heterozygous
Haack et al. 2012	n.d.	n.d.	Mitochondrial encephalopathy, cardiomyopathy, muscular hypotonia, respiratory insufficiency	n.d.	Histology n.d. complex I↓ (38% of normal)	Fibroblastscomplex I (52% of normal)	c.279C>T/p.Arg79Trp c.476C>A/p.Ala159Asp compound heterozygous
	n.d.	n.d.	Muscular hypotonia, dyskinesia, epilepsy, lactic acidosis	TS	Histology n.d. complex 1↓ (8% of normal)	Fibroblastscomplex I (54% of normal)	c.187G>C/p.Glu63Gln homozygous
	n.d.	n.d.	Muscular hypotonia, hypertrophic cardiomyopathy. lactic acidosis	LS	n.d.	n.d.	
Tuppen et al. 2010	4 weeks	Died at 12 weeks	Poor feedine, apriea, lactic acidosis, seizures, failure to thrive, respiratory insufficiency	n.d.	Histology n.d. complex I↓ (30% of normal)	n.d.	c.236C>T, p.Pro79Leu homozygous
Calvo et al. 2010sibs	n.d.	n.d.	Mitochondrial encephalomyopathy	Leuko- encepha- lopathy	Complex I	n.d.	c.460G>A, p.Gly154Ser homozygous
Patients describe	d in this pa	per					
P1	Birth	Alive at 13 years	Developmental delay, PEO, muscular hypotonia, ataxic gait, dysarthria, dystonia, resniratory problems	TS	Histology normcomplex I↓ (25% of normal)	n.d.	c.160C>T, p.Arg54Trp homozygous
P2	5 years	Alive at 11 vears	Dysarthria, muscular hypotonia, myopia, ntosis, PFO, swallowing difficulties	LS	n.d.	n.d.	
P3	9 years	Alive at 9 years	Muscular hypotonia, scapular winging, dysmetria, nystagmus, myopia	n.d.	n.d.	n.d.	

Table 1 Summary of all previously reported patients compared to our patients with NDUFS8 mutations

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Abbreviations: + present, n.d. not determined; PEO progressive external ophthalmoparesis, LS Leigh syndrome, RC respiratory chain

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Synopsis

Here we report three children with a mild and unusual clinical presentation of Leigh syndrome with progressive external ophthalmoparesis due to a homozygous mutation in a nuclear complex I structural subunit gene detected through non-targeted whole exome sequencing.

Author Contribution

Adela Della Marina and Ulrike Schara participated in the clinical follow-up of the patients and in the drafting the manuscript.

Angela Pyle performed whole exome sequencing and participated in the drafting of the manuscript.

Claudia Möller-Hartmann analyzed and interpreted the MRI images.

Elke Holinski-Feder, Angela Abicht and Birgit Czermin did the diagnostic work-up.

Hanns Lochmüller was involved in the biochemical and histological analyses.

Helen Griffin and Mauro Santibanez-Koref did the bioinformatic analysis of the exome data.

Patrick F. Chinnery and Rita Horvath are responsible for the clinical and diagnostic work-up, the study designs and for drafting of the manuscript.

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Ethical Issues

The study was approved by the County Durham & Tees Valley 1 Research Ethics Committee (08/H0905/106 "Nuclear genes in mitochondrial disease").

The parents gave written consent for the study and for presenting the data and photographs of the patients.

Competing Interest Statement

The authors have no competing interests.

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