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Autosomal dominant optic atrophy and cataract “plus” phenotype including axonal neuropathy

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Abstract

Objective

To characterize the phenotype in individuals with OPA3-related autosomal dominant optic atrophy and cataract (ADOAC) and peripheral neuropathy (PN).

Methods

Two probands with multiple affected relatives and one sporadic case were referred for evaluation of a PN. Their phenotype was determined by clinical ± neurophysiological assessment. Neuropathologic examination of sural nerve and skeletal muscle, and ultrastructural analysis of mitochondria in fibroblasts were performed in one case. Exome sequencing was performed in the probands.

Results

The main clinical features in one family (n = 7 affected individuals) and one sporadic case were early-onset cataracts (n = 7), symptoms of gastrointestinal dysmotility (n = 8), and possible/confirmed PN (n = 7). Impaired vision was an early-onset feature in another family (n = 4 affected individuals), in which 3 members had symptoms of gastrointestinal dysmotility and 2 developed PN and cataracts. The less common features among all individuals included symptoms/signs of autonomic dysfunction (n = 3), hearing loss (n = 3), and recurrent pancreatitis (n = 1). In 5 individuals, the neuropathy was axonal and clinically asymptomatic (n = 1), sensory-predominant (n = 2), or motor and sensory (n = 2). In one patient, nerve biopsy revealed a loss of large and small myelinated fibers. In fibroblasts, mitochondria were frequently enlarged with slightly fragmented cristae. The exome sequencing identified OPA3 variants in all probands: a novel variant (c.23T>C) and the known mutation (c.313C>G) in OPA3.

Conclusions

A syndromic form of ADOAC (ADOAC+), in which axonal neuropathy may be a major feature, is described. OPA3 mutations should be included in the differential diagnosis of complex inherited PN, even in the absence of clinically apparent optic atrophy.

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Mutations in OPA3 (figure 1), encoding a mitochondrial protein likely involved in the regulation of mitochondrial fission,\(^1\)–\(^5\) are associated with 2 distinct disorders that share the common feature of bilateral optic atrophy (OA; table e-1, links.lww.com/NXG/A146). Recessive, loss-of-function mutations cause 3-methylglutaconic aciduria type III (MGA3 [MIM 258501]), also known as Costeff syndrome, a rare disorder that is clinically characterized by OA and extrapyramidal dysfunction with onset in the first decade of life, subsequent development of spastic paraparesis and cerebellar signs, and increased urinary excretion of 3-methylglutaconic and 3-methylglutaric acids.\(^6\)–\(^13\) Dominant mutations, in contrast, lead to OA that typically presents within the first 2 decades of life and is often associated with cataracts with variable age at onset (autosomal dominant OA and cataract; ADOAC [MIM 165300]).\(^14\)–\(^18\) Additional features such as hearing loss, cerebellar and extrapyramidal signs, and dysautonomic or gastrointestinal symptoms have been reported less consistently in patients with ADOAC (table e-2). Symptoms or signs suggestive of peripheral neuropathy (PN) have been described to date in 4 individuals with dominant OPA3 mutations,\(^14\),15,18 although it was confirmed by nerve conduction studies (NCSs) in only one of them.\(^18\)

Here, we describe 2 families and one sporadic case with a syndromic form of ADOAC in which PN was a major clinical feature. This report broadens the clinical and genetic spectrum of ADOAC and indicates that OPA3 should be included in the differential diagnosis of the complex inherited PN\(^19\) even in the absence of clinically apparent OA.

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**Figure 1 Schematic of the OPA3 gene and OPA3 protein isoform b**

The OPA3 gene (NCBI RefSeq NG_013332.1; top figure), located in chromosome 19q13.32, contains 3 exons (1, 2, and 3; boxes containing exon numbers) and spans 57.4 kb. The coding regions of exons 2 and 3 and their exon-intron boundaries are highly similar and may have originated by segmental duplication.\(^2\) OPA3 exons are alternatively spliced to generate 2 mRNA transcripts: transcript variant 2 (exon 1 plus exon 2 [NM_025136.3]) and transcript variant 1 (exon 1 plus exon 3 [NM_001017989.2]). Transcript variant 2 seems to be the predominant transcript in most tissues and encodes a 179 amino acid (AA) protein (OPA3 isoform b [NP_079412.1]; bottom figure), Transcript variant 1 encodes a 180 AA protein (OPA3 isoform a [NP_001017989]; not shown). OPA3 isoform b amino acids 1–48 are encoded by exon 1 (dark green) and amino acids 48–179 are encoded by exon 2 (light green). Its N-terminal region contains a putative mitochondrial targeting sequence in AAs 1–18 (as predicted by in silico analysis with MitoProt and TargetP) or 1–30 (as indicated by functional studies with deletion mutants). In silico analysis with MitoFates predicts a mitochondrial processing peptidase cleavage site (red arrow) and 2 TOM20 recognition motifs (AAs 8–12 and 48–52; red boxes). An additional mitochondrial sorting/cleavage signal at position 25–29 (purple box) has been proposed by some authors.\(^3\),\(^8\) The localizations of reported dominant and recessive mutations are shown in the figure (one-letter AA abbreviations are used for simplicity; blue boxes indicate mutations reported in the present study, p.Leu11Gln [p.L11Q] was homozygous in 2 siblings with optic atrophy, extrapyramidal signs including dystonia, and pyramidal signs or ataxia, and it was heterozygous in their mother with later-onset dystonia.\(^2\) No pathogenic mutations have been described in AAs 48–180 of OPA3 isoform a encoded by exon 3 (not shown).
Methods

Patients
The 3 probands were identified at the outpatient clinics of the National Hospital for Neurology and Neurosurgery, London, UK, and the Charcot-Marie-Tooth disease (CMT) Center of Excellence at the University of Pennsylvania, Philadelphia, PA. These probands were recruited into local research protocols to determine the genetic etiology in patients with inherited neuropathies using next-generation sequencing. Diagnostic laboratory tests, neurophysiological studies, MRI scans, and tissue biopsies were performed and analyzed using standard protocols.

Sequencing
DNA samples were extracted from peripheral blood leukocytes and skeletal muscle biopsy specimens using commercial kits. Exome sequencing was performed after target capture using the Illumina TruSeq Exome, Agilent SureSelect Focused Exome, or Agilent SureSelect Human All Exon kit. The Illumina HiSeq2000 or HiSeq2500 instruments were used to produce 100 bp paired-end sequence reads. The following software tools were used to align sequence reads to the human genome assembly 19 (GRCh37) and to call and annotate variants: Novocraft NovoAlign, Burrows-Wheeler Aligner, Picard, Genome Analysis Toolkit, SAMtools, and ANNOVAR. After filtering, candidate variants were evaluated in silico to predict their effects, and validation and cosegregation analysis of OPA3 variants in families A, B, and C were performed by Sanger sequencing (see supplementary material for details, links.lww.com/NXG/A146).

Mitochondrial DNA (mtDNA) was assessed for large-scale rearrangements using long-range PCR and Southern blot of total genomic DNA extracted from skeletal muscle. The entire mtDNA sequence was analyzed with Affymetrix GeneChip Human Mitochondrial Resequencing Array 2.0 as described elsewhere.

Cell imaging
Fibroblast cell lines were established from skin biopsies of proband AII-2 and a healthy age- and sex-matched control using standard methods. Cells were grown in high glucose Dulbecco’s modified Eagle’s medium, supplemented with 10% fetal bovine serum, penicillin/streptomycin, and 0.05 mg/mL uridine. For electron microscopy (EM), fibroblasts were fixed in 3% glutaraldehyde in 0.1 M sodium cacodylate buffer and 5 mM CaCl₂ overnight, then treated with 1% osmium tetroxide for 3 hours at 4°C and embedded in Araldite CY-212 resin. Ultrathin sections (70 μm) were stained with lead citrate and uranyl acetate. Images were taken on a Philips CM10 transmission electron microscope fitted with a MegaView G3 camera and RADIUS Software (Olympus). Mitochondria morphology in fibroblasts was measured blind to disease status.

Standard protocol approvals, registrations, and patient consents
Research protocols were approved by local institutional review board and/or ethics committee. Patients gave written informed consent to participate.

Data availability
Anonymized data not published within this article will be made available by request from any qualified investigator.

Results

Families and overall phenotype
Three unrelated probands were referred to our PN clinics for evaluation. Probands AII-2 and BIII-2 had affected relatives of 3 generations based on history (AI-1, AII-1, AIII-2, AIII-3, AIII-4, BI-1, and BII-3) or examination (AIII-1 and BII-1), while proband CII-1 was a sporadic case (figure 2). The clinical information of all affected individuals is summarized in table 1, and their clinical description is provided in the supplementary material (links.lww.com/NXG/A146).

Family A
The phenotype in the affected members of family A (n = 7) was mostly characterized by early-onset cataracts, gastrointestinal dysmotility symptoms, and PN (based on symptoms/signs or neurophysiological studies), although the data on

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Image of figure 2: Pedigrees of families and segregation analysis of variants c.23T>C (p.Met8Thr) and c.313C>G (p.Gln105Glu) in OPA3 (NM_025136.3)

![Pedigrees of families and segregation analysis of variants c.23T>C (p.Met8Thr) and c.313C>G (p.Gln105Glu) in OPA3 (NM_025136.3)](image-url)

OPA3 variants found in each family and the genotype of tested individuals are indicated in red. Arrow = probands.
these features were incomplete for 3 individuals. Reduced visual acuity was first detected in proband AII-2 during the evaluation of her PN at age 49, and OA was subsequently confirmed (figure 3, A and B). A reduced visual acuity was also observed in AIII-1, but we had no confirmation of an underlying OA. Impaired vision was not reported for other relatives. Intrafamilial phenotypic variability was evident, with patient AIII-1 having more severe symptoms than his siblings, and an earlier and more severe presentation than AII-2.

Family B
Early-onset, bilateral visual impairment leading to severely reduced visual acuities by the age of 30–40 was the initial clinical feature in all affected members of family B (n = 4; originally reported in 199622). Three of them had a diagnosis of OA and reported symptoms of gastrointestinal dysmotility. PN and cataracts were detected in proband BIII-2 and her father BII-1 on follow-up examinations in their adulthood.

Family C
Proband CII-1 was the only affected individual in this family. Her phenotype was similar to that of family A: early-onset cataracts, gastrointestinal dysmotility symptoms, and PN. In addition, she had recurrent episodes of pancreatitis from age 11. Bilateral OA was detected by reverse phenotyping: after a genetic diagnosis of her PN was achieved, by optical coherence tomography.

Peripheral neuropathy
Nine of the 12 affected individuals from the 3 families had a possible or confirmed PN. It was considered possible in 4 individuals (AII-4, AIII-2, AIII-3, and AIII-4) based on the history provided by relatives. In 5 patients (AII-2, AIII-1, BII-1, BIII-2, and CII-1), we confirmed the diagnosis based on clinical and neurophysiological assessment (see supplementary material for an extended description, links.lww.com/NXG/A146). Detailed results of NCS/EMG were available in 4 cases (table e-3).

Table 1 Summary of clinical features in affected individuals from families A, B, and C

| Individual | Bilateral cataracts | Bilateral optic atrophy | Gastrointestinal dysmotility symptoms | Peripheral neuropathy symptoms/signs | PN on NCS/EMG | Hearing loss | Other features |
|------------|---------------------|-------------------------|--------------------------------------|-------------------------------------|---------------|-------------|---------------|
| AI-1       | +                   | n/a                     | +                                    | ±                                   | n/a           | ± 50s       |               |
| AII-1      | +                   | n/a                     | +                                    | ±                                   | n/a           | n/a         |               |
| AII-2      | +                   | n/a                     | + on examination 49 y                | ±                                   | axonal S>M    | + 50s       | Autonomic features, mild ptosis |
| AIII-1     | +                   | ≤ 10 s                  | +                                    | +                                   | axonal S&M    | +           | Parenteral nutrition from his 20s |
| AIII-2     | n/a                 | n/a                     | ±                                    | n/a                                 | n/a           | n/a         |               |
| AIII-3     | + early-onset       | n/a                     | ±                                    | n/a                                 | +             | n/a         |               |
| AIII-4     | + surgery ≥15 y     | n/a                     | ±                                    | +                                   | n/a           | n/a         |               |
| BI-1       | n/a                 | ±                       | n/a                                 | n/a                                 | n/a           | n/a         |               |
| BII-1      | + on examination 65 y | +                       | ±                                    | + on examination 65 y                | axonal S>M    | —           |               |
| BII-3      | n/a                 | + 7 y                   | ±                                    | n/a                                 | n/a           | n/a         |               |
| BIII-2     | + surgery 44 y      | + 5 y                   | + 5 y                               | + 20 s                              | axonal S>M    | + 50s       | Orthostatic hypotension |
| CII-1      | + surgery 14 y      | + on examination 32 y   | + 10 s                              | + ≤ 16 y                            | axonal S&M    | —           | Recurrent pancreatitis, hypertension, PRES |

Abbreviations: n/a = not available or unknown; M = motor; NCS/EMG = nerve conduction studies/electromyography; PN = peripheral neuropathy; PRES = posterior reversible encephalopathy syndrome; S = sensory; ± mild or possible symptoms reported by relatives.

a Bilateral visual loss but no confirmed diagnosis of optic atrophy.
b Abnormal neurophysiological studies.
Asymptomatic neuropathy
In patient BII-1, the PN was asymptomatic and detected on examination at age 65. NCS were consistent with an axonal motor and sensory neuropathy.

Sensory-predominant neuropathy
Probands AII-2 and BIII-2 had a slowly progressive, predominantly sensory neuropathy. AII-2 developed sensory symptoms in her feet in her late 40s and was referred to us for the evaluation of suspected PN at age 49. BIII-2 developed sensory symptoms in her feet in his 20s; her PN was confirmed in her early 30s, but she was referred again to us at age 52 because of symptom worsening. Both patients described reduced sensation restricted to the lower limbs. On serial examinations, we observed distal sensory loss in the upper and lower limbs and signs of mild motor involvement. In both cases, NCS revealed a length-dependent axonal sensory neuropathy. NCS/EMG signs of distal motor involvement were detected only in proband BIII-2.

Motor and sensory neuropathy
Proband CII-1 and patient AIII-1 had a moderate-to-severe progressive motor and sensory neuropathy with onset in their childhood or teens. Proband CII-1 had frequent ankle sprains as a child and was referred to us at age 16 when a PN was suspected during physical therapy for a foot fracture. Patient AIII-1 had pes cavus and limb weakness since his teens and had been diagnosed with inherited neuropathy in his early 20s; he was examined by us at age 31 as part of the family evaluation. Both patients exhibited motor and sensory deficits in upper and lower limbs on examination. Proband CII-1 had 4 neurophysiological studies performed between ages 16 and 32, which revealed a severe motor and sensory axonal neuropathy. In patient AIII-3, a previous neurophysiological study at age 29 confirmed the same diagnosis.

Other clinical features
Symptoms of gastrointestinal dysmotility of variable severity were reported for most individuals. In the best documented cases, symptoms included intermittent constipation (AII-2); episodes of nausea, vomiting, and abdominal pain (CII-1); gastroparesis and episodes of intestinal pseudo-obstruction (BIII-2); and intestinal pseudo-obstruction requiring parenteral nutrition (AIII-1). Patient BIII-2 required emergency surgery for intestinal intussusception in her 50s, and patient AII-1 was also reported to require surgery for her intestinal motility problems.

All probands had features suggestive of autonomic nervous system dysfunction, including orthostatic tachycardia (AII-2), postural hypotension (BIII-2), and episodes of hypertension in the context of gastrointestinal symptoms that, in one...
occurrence, lead to a posterior reversible encephalopathy syndrome (CII-1).

Patient AIII-1 developed hearing loss concurrently to other symptoms. Proband AII-2 complained of mild hearing loss in her 50s, and a bilateral auditory neuropathy was confirmed in her 60s. Patient AI-1 and proband BIII-2 developed hearing loss in their 50s. The less common clinical features are shown in table 1.

**Neuropathology**

Patient AII-2 underwent sural nerve and quadriceps muscle biopsy at age 57. Nerve biopsy showed a severe loss of myelinated fibers that was not selective for large or small fibers, with no apparent active axonal degeneration and minimal regeneration, and no evidence of demyelination (figure 3, C and D). Muscle biopsy revealed occasional angular atrophic fibers, 2 cytochrome oxidase-negative fibers, and one ragged-red fiber. The activity of mitochondrial electron transport chain (ETC) complexes I, II + III, and IV was within control ranges, and screening for single/multiple deletions and point mutations of mtDNA in muscle were negative.

**EM of mitochondria**

Ultrastructural examination of mitochondria with EM was performed in the cultured fibroblasts from proband AII-2 and a healthy control. On EM images, we frequently observed abnormally enlarged mitochondria coupled with slightly fragmented mitochondrial cristae and a concomitant reduction in the area of mitochondrial cristae (figure 4).

**Genetic studies**

In probands AII-2, BIII-2, and CII-1, analysis of common genes associated with inherited neuropathy and/or OA yielded negative results. Exome sequencing was then performed on these patients as 3 independent studies. The analysis focused on nonsynonymous, splice-site, and coding indel variants with minor allele frequency <0.1% in the Exome Aggregation Consortium data set, located in genes known to cause inherited neuropathy and/or OA (table e-4, links.lww.com/NXG/A146).

In proband AII-2, we identified the heterozygous missense variant c.23T>C in exon 1 of OPA3 (table e-5, links.lww.com/NXG/A146). Sanger sequencing detected the mutation in the proband and her affected daughter (AIII-4) but not in the unaffected son (AIII-5), who was homozygous for the wild-type allele (figure 2). c.23T>C is a novel variant absent from public databases and from an in-house control exome database. It affects an evolutionarily conserved nucleotide and is predicted to be deleterious by several bioinformatics tools (table e-6). c.23T>C leads to the substitution of nonpolar, hydrophobic methionine for polar threonine at amino acid (AA) position 8 of OPA3 (p.Met8Thr). This position lies within the N-terminal mitochondrial targeting sequence (MTS) predicted by MitoProt and TargetP software tools (AA 1-18) or by functional studies with deletion mutants (AA 1-30), in which another dominant mutation has been identified (figure 1). In silico analysis with MitoFates suggests that methionine 8 would be the first of 5 AA of a putative consensus motif recognized by the mitochondrial import receptor subunit TOM20 (figure 1) and that p.Met8Thr would eliminate this recognition motif. The functional significance of hydrophobic AA at position 8 of OPA3 is supported by multiple sequence alignment of OPA3 homologs, which shows conservation of methionine and isoleucine at that position from human to zebra fish (figure e-1). Based on these data, and the previous association of dominant OPA3 mutations with OA, cataracts, and PN, we considered c.23T>C as the most plausible genetic etiology in family A.

In probands BIII-2 and CII-1, we identified the heterozygous missense mutation c.313C>G (p.Gln105Glu) in exon 2 of OPA3. The mutation was validated by Sanger sequencing in both cases. In family B, DNA samples from relatives were not available for cosegregation analysis. In family C, the mutation was not found in either parent, confirming that it had occurred de novo in the proband (figure 2). c.313C>G is not reported in public databases but is the most frequent mutation reported in families with dominant OPA3-related OA, sometimes in association with additional features (tables e-1 and e-2, links.lww.com/NXG/A146). This variant was therefore classified as likely pathogenic.

**Figure 4** Ultrastructural examination of mitochondria in cultured skin fibroblasts

(A) Representative electron micrographs showing the ultrastructure of mitochondria in control and patient AII-2 fibroblasts (scale bar = 1 μm). (B) Patient AII-2 mitochondria (n = 94) display a significant increase in area as compared to control mitochondria (n = 123). (C) Data are presented as box plots illustrating 80% of the data distribution; 10th, 25th, median, 75th, and 90th percentiles are shown for these box plots. *p < 0.0005 (Mann-Whitney U test).
Discussion

Here, we describe the members from 3 families in which a novel missense variant (c.23T>C, p.Met8Thr) and a known mutation (c.313C>G, p.Gln105Glu) in OPA3 were the most likely underlying cause of a complex phenotype consisting of OA, cataracts, gastrointestinal dystotility, axonal neuropathy, and possibly autonomic dysfunction and hearing loss. This observation, together with 2 former reports of similar phenotypes associated with the missense substitutions p.Leu79Val and p.Gln105Glu, demonstrates that the dominant OPA3 mutations can cause syndromic forms of ADOAC (ADOAC “plus”).

We also confirm that an axonal neuropathy may be a major clinical feature of ADOAC/+.

Based on the pathogenicity of the same pathologic process rather than distinct PN subtypes. A sensory ganglionopathy seemed unlikely in patients with sensory-predominant forms, in whom the PN was length-dependent and symmetrical, with almost normal joint position sense and no significant limb ataxia.

Despite the limited sample size, our report also suggests a higher prevalence of PN among patients with ADOAC/+ than previously recognized. Possible reasons for this are that the PN may be asymptomatic and/or develop several decades after the onset of the ophthalmologic problems. Therefore, long-term follow-up with neurologic evaluations may be necessary to confirm or exclude a PN in these patients. On the other hand, data from this and previous studies indicate that the dominant OPA3 mutations may exhibit considerable phenotypic variability, so it is also possible that the families or individuals with the same mutation may or may not develop additional features such as PN. This point should be clarified in future studies encompassing larger sample sizes.

Symptoms of gastrointestinal dystotility were reported for most affected individuals from the present series, which so far has only been described in 3 patients with ADOAC/+.

The precise function of the OPA3 protein and the molecular mechanisms by which heterozygous missense mutations in OPA3 cause ADOAC/+ are unknown. OPA3 predominantly localizes to the mitochondria but its intramitochondrial topology is still discussed: the mouse homolog of OPA3 copurifies with the inner mitochondrial membrane, but the studies on subcellular fractions of HeLa cells suggest that OPA3 may be anchored to the mitochondrial outer membrane with the C-terminus exposed to the cytoplasm. Overexpression of OPA3 induces mitochondrial fragmentation, whereas downregulation leads to more elongated and tubular mitochondria. These data support that OPA3 is a mitochondrial membrane protein implicated in the regulation of mitochondrial fission and morphology.

Heterozygous carriers of the recessive OPA3 mutation c.143-1G>C, which abolishes mRNA expression, are asymptomatic. This suggests that dominant missense mutations have a dominant-negative effect or result in a gain-of-function rather than haploinsufficiency. Fibroblasts from ADOAC patients with the p.Val3_Gly4insAlaPro mutation, located in the N-terminal MTS of OPA3, showed increased fragmentation of the mitochondrial network, which mimics OPA3 overexpression and gives support to the gain-of-function hypothesis. Similar findings were observed in HeLa cells transfected with an OPA3 mutant carrying the p.Gly93Ser mutation, located in the hydrophobic region of OPA3 (AA 83-120) that is required for mitochondrial fragmentation. However, no mitochondrial network abnormalities were detected in fibroblasts from one patient with the same mutation. Therefore, it is possible that dominant mutations affecting different regions of OPA3 may exert different deleterious effects.

Studies in a zebra fish model of MGA3 indicate that OPA3 does not have a direct role in mitochondrial ETC function, and consistent with this, we found the normal activities of ETC complexes in skeletal muscle from patient All-2. We did not detect large-scale mtDNA rearrangements in muscle, which argues against a role of OPA3 in mtDNA maintenance. However, EM analysis of mitochondria in fibroblasts from patient All-2 revealed frequently enlarged mitochondria with abnormal cristae. Subtle alterations to the morphology of mitochondrial cristae have also been identified in the retinal tissues of a mouse model of MGA3 caused by the homozygous missense mutation p.Leu122Pro. Therefore, OPA3 function might be important for the structural integrity of mitochondrial cristae, and altered mitochondrial cristae architecture could be an additional mechanism involved in the pathogenesis OPA3-related disease.

Based on the findings from the present study, some parallels can be drawn between OPA3- and OPA1-related diseases. Mutations affecting OPA1, a dynamin-related guanosine triphosphatase involved in mitochondrial dynamics, represent
a major cause of dominant OA (DOA), and ~20% of patients may present a DOA “plus” phenotype, including axonal neuropathy.25 Biallelic mutations in OPA1 can also cause Behr syndrome,26–28 a severe, early-onset neuro-ophtalmologic syndrome that is clinically similar to MGA3. Of interest, the loss of OPA1 causes disruption of mitochondrial cristae structures.29 Overall, these data may indicate shared pathogenic mechanisms between OPA3 and OPA1 defects that warrant further research.

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Appendix Author contributions

| Name                  | Location                      | Role           | Contribution                                      |
|-----------------------|-------------------------------|----------------|--------------------------------------------------|
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Appendix (continued)

| Name                  | Location                                           | Role                                      | Contribution                                                                 |
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| Michael G. Hanna, MD, FRCPath | University College London, London, UK              | Author                                   | Drafting/revising the manuscript for content; Study concept or design; Study supervision; Obtaining funding |
| Steven S. Scherer, MD, PhD | Perelman School of Medicine, Philadelphia, USA | Author                                   | Drafting/revising the manuscript for content; Major role in acquisition of data; Study supervision |
| Henry Houlden, MD, PhD | University College London, London, UK              | Author                                   | Drafting/revising the manuscript for content; Major role in acquisition of data; Study concept and design; Study supervision; Obtaining funding |
| Mary M. Reilly, MD, FRCP | University College London, London, UK              | Author                                   | Drafting/revising the manuscript for content; Major role in acquisition of data; Study concept and design; Study supervision; Obtaining funding |

References

1. Da Cruz S, Xenarios I, Langridge J, Vilboux F, Parone PA, Martinou JC. Proteomic analysis of the mouse liver mitochondrial inner membrane. J Biol Chem 2003;278:41566–41571.
2. Ryu SW, Jeong HJ, Choi M, Karbowska M, Choi C. Optic atrophy 3 as a protein of the mitochondrial outer membrane induces mitochondrial fragmentation. Cell Mol Life Sci 2010;67:2839–2851.
3. Huizing M, Dorward H, Ly J, et al. OPA3, mutated in 3-methylglutaconic aciduria type III, encodes two transcripts targeted primarily to mitochondria. Mol Genet Metab 2010;100:149–154.
4. Powell KA, Davies JR, Taylor E, Wride MA, Votruba M. Mitochondrial localization and ocular expression of mutant Opa1 in a mouse model of 3-methylglutaconic aciduria type III. Invest Ophthalmol Vis Sci 2011;52:4359–4380.
5. Ryu SW, Yoon J, Yim N, Choi K, Choi C. Downregulation of OPA1 is responsible for transforming growth factor-beta-induced mitochondrial elongation and F-actin rearrangement in retinal pigment epithelial ARPE-19 cells. PLoS One 2013;8:e53495.
6. Costel H, Gadoth N, Aptor A, Prusiaik M, Savin H. A familial syndrome of infantile optic atrophy, movement disorder, and spastic paraplegia. Neurology 1989;39:595–597.
7. Elpegd ON, Costeff H, Joseph A, Shental Y, Weitz R, Gibson KM. 3-Methylglutaconic aciduria in the Iraqi-Jewish ‘optic atrophy plus’ (Costeff) syndrome. Dev Med Child Neurol 1994;36:167–172.
8. 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.
9. Kleta R, Skovby F, Chintisens E, Rosenberg T, Gahl WA, Anikster Y. 3-Methylglutaconic aciduria type III in a non-Iraqi-Jewish kindred: clinical and molecular findings. Mol Genet Metab 2002;76:201–206.
10. Neas K, Bennetts B, Carpenter K, et al. OPA3 mutation screening in patients with unexplained 3-methylglutaconic aciduria. J Inherit Metab Dis 2005;28:525–532.
11. Ho G, Walter JH, Chistodoulou J. Costeff optic atrophy syndrome: new clinical case and novel molecular findings. J Inherit Metab Dis 2008;31(suppl 2):419–423.
12. Lam C, Gallo LK, Dineen R, et al. Two novel compound heterozygous mutations in OPA3 in two siblings with OPA3-related 3-methylglutaconic aciduria. Mol Genet Metab Rep 2014;1:114–123.
13. Yalahom G, Anikster Y, Huna-Baron R, et al. Costeff syndrome: clinical features and natural history. J Neurol 2014;261:2275–2282.
14. Reynier P, Amati-Bonneau P, Verny C, et al. OPA3 gene mutations responsible for autosomal dominant optic atrophy and cataract. J Med Genet 2004;41:e110.
15. Ayirignac I, Liauzun C, Leners G, et al. OPA3—related autosomal dominant optic atrophy and cataract with ataxia and areflexia. Eur Neurol 2012;68:108–110.
16. Grau T, Burbulla LL, Engl G, et al. A novel heterozygous OPA3 mutation located in the mitochondrial target sequence results in altered steady-state levels and fragmented mitochondrial network. J Med Genet 2013;50:848–858.
17. Sergouniotis PI, Pevern G, Thioelen DL, et al. Clinical and molecular genetic findings in autosomal dominant OPA3-related optic neuropathy. Neurogenetics 2015;16:69–75.
18. Bourne SC, Townsends KN, Shyr C, et al. Optic atrophy, cataracts, lipodystrophy/lipostrophy, and peripheral neuropathy caused by a de novo OPA3 mutation. Cold Spring Harb Mol Case Stud 2017;3:e001156.
19. Rosser AM, Carr AS, Devine H, et al. Peripheral neuropathy in complex inherited diseases: an approach to diagnosis. J Neurol Neurosurg Psychiatry 2017;88:846–863.
20. Wang K, Li M, Hakonarson H. ANNOVAR: functional annotation of genetic variants from high-throughput sequencing data. Nucleic Acids Res 2010;38:e164.
21. Rahman S, Ecob R, Costello H, et al. Hearing in 44-45 year olds with m.1555A>G, a genetic mutation predisposing to aminoglycoside-induced deafness: a population based cohort study. BMJ Open 2012;2:e000411.
22. Hom XB, Lavine JE. Gastrointestinal complications of mitochondrial disease. Mitochondrion 2006;4:601–607.
23. Gunay-Aygun M, Huizing M, Anikster Y. OPA3-related 3-methylglutaconic aciduria. In: Adam MP, Ardinger HH, Pagon RA, et al, editors. GeneReviews® [Internet]. Seattle: University of Washington, Seattle; 2013.
24. Pet W, Kratz LE, Bernardini I, et al. A model of Costeff Syndrome reveals metabolic and protective functions of mitochondrial OPA3. Development 2010;137:2587–2596.
25. Chao de la Barca JM, Prunier-Mirebeau D, Amati-Bonneau P, et al. OPA3-related disorders: diverse clinical expression, modes of inheritance and pathophysiology. Neurobiol Dis 2016;90:20–26.
26. Bonneau D, Colin E, Oca F, et al. Early-onset Behr syndrome due to compound heterozygous mutations in OPA1. Brain 2014;137:e301.
27. Carelli V, Sabatelli M, Carrozzo R, et al. “Behr syndrome” with OPA1 compound heterozygous mutations. Brain 2015;138:e321.
28. Carelli V, Amati-Bonneau P, Reynier P, et al. Heterozygous OPA1 mutations in Behr syndrome. Brain 2011;134:e169–e170.
29. Olichon A, Baricault L, Gas N, et al. Loss of OPA1 perturbs the mitochondrial inner membrane structure and integrity, leading to cytochrome c release and apoptosis. J Biol Chem 2003;278:7743–7746.
30. Arif B, Kumar KR, Sehler P, et al. A novel OPA3 mutation revealed by exome sequencing: an example of reverse phenotyping. JAMA Neurol 2013;70:783–787.