

# Mitochondrial Epilepsies

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## The Causal Disease

### Definitions and Epidemiology of Mitochondrial Disease

Mitochondrial diseases are inherited conditions caused by gene defects that directly or indirectly affect the function of the oxidative phosphorylation (OXPHOS) system in the mitochondria [1]. The OXPHOS system comprises five multi-subunit enzyme complexes embedded in the inner mitochondrial membrane. Together, these enzyme complexes are responsible for generating the majority of cellular ATP (Figure 30.1). As a consequence, mitochondrial disorders predominantly affect organs with high energy demands, especially the brain, and epilepsy is a frequent manifestation. Mitochondrial disorders are estimated to affect at least 1 in 5000 births and may be caused by defects in mitochondrial DNA (37 maternally inherited genes located within the mitochondria themselves) or nuclear genes (>300 genes are currently linked to mitochondrial dysfunction).

### Pathology, Physiology and Clinical Features of Mitochondrial Disease

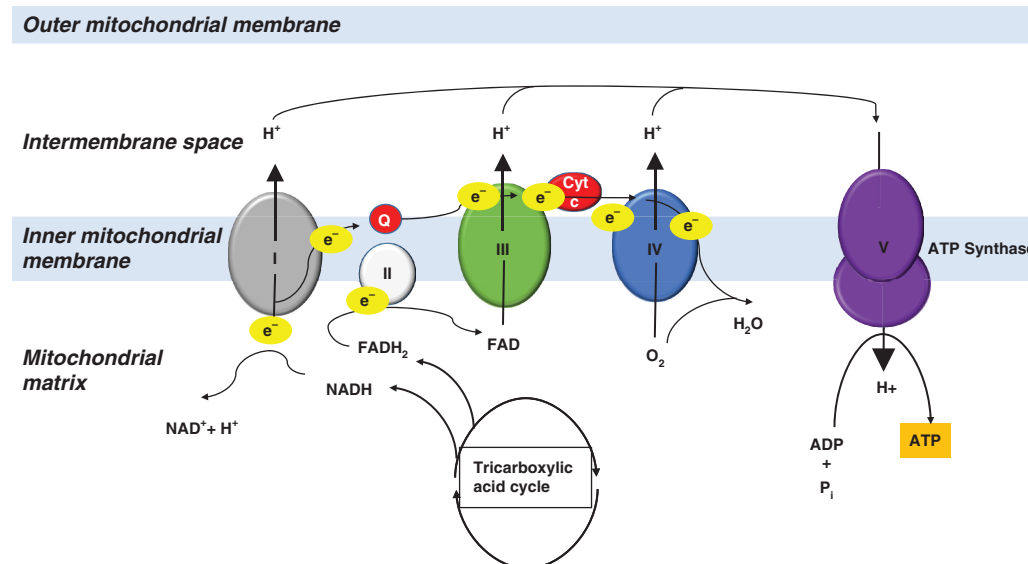
Mitochondrial disease may be classified according to clinical syndrome, biochemical deficiency or class of gene defect. Mitochondrial diseases are frequently multisystem disorders (theoretically any organ may be affected in any combination)

and epilepsy is rarely the sole (or even the first) clinical manifestation [2]. Clinical features frequently observed in mitochondrial disease are shown in Figure 30.2. Some patients have stereotyped manifestations fitting within a classical mitochondrial syndrome, others have nonspecific clinical features. Underlying disease mechanisms include defects of OXPHOS subunits or assembly factors (which may be associated with isolated deficiency of a single enzyme complex), and disorders of cofactor biosynthesis, mtDNA maintenance, and mitochondrial translation, import, membrane lipids and dynamics (all of which may be associated with multiple enzyme deficiencies). Downstream pathophysiological mechanisms are not completely understood and are not purely related to energetic failure. Impairment of other essential mitochondrial functions is also believed to contribute to pathogenesis of mitochondrial epilepsy, including alterations in calcium homeostasis, excitotoxicity, oxidative stress secondary to increased production of reactive oxygen species and depletion of cerebral folate, arginine, citrulline or  $\text{NAD}^+$  [3].

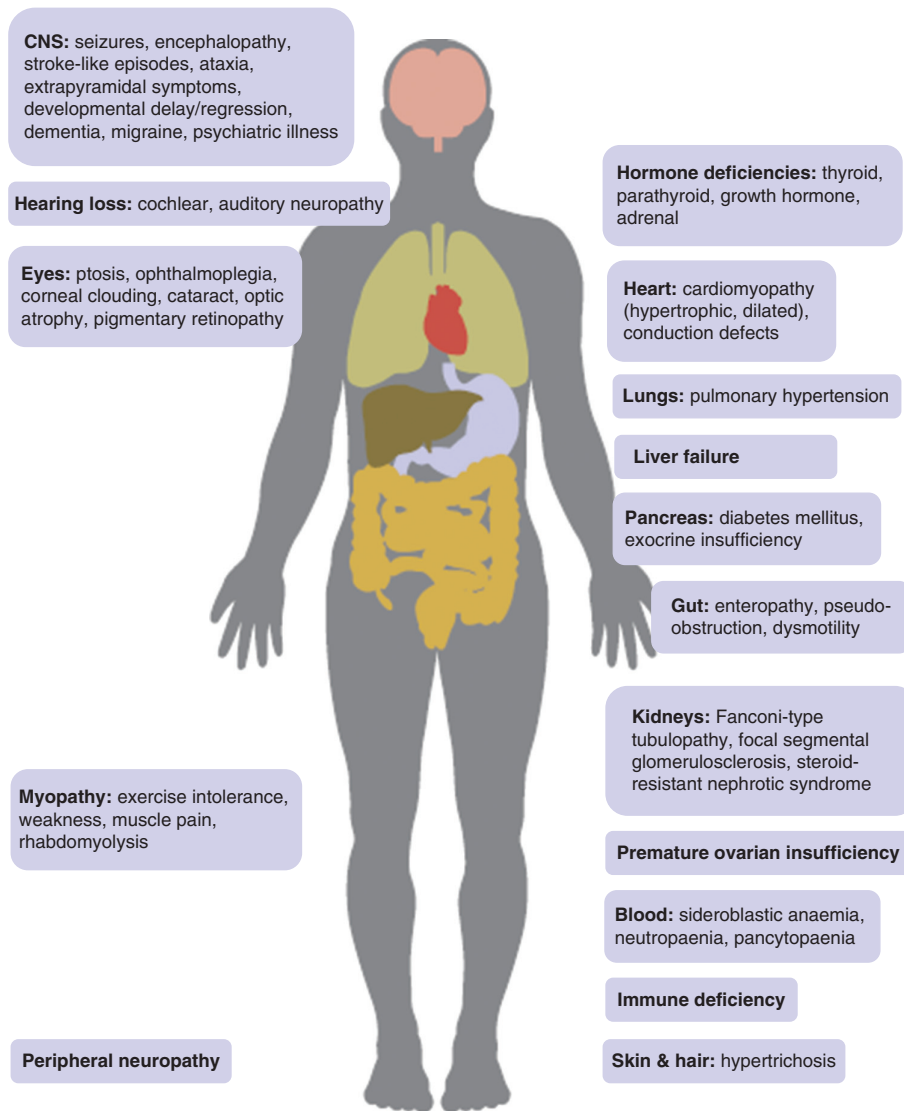
## Epilepsy in Mitochondrial Disease

### Frequency of Epilepsy in Mitochondrial Disease

Seizures occur frequently in both paediatric and adult mitochondrial disease. Reported seizure frequency ranged from 23% in an adult cohort [4] to 35–61% in several paediatric cohorts [5–7].



**Figure 30.1** The mitochondrial oxidative phosphorylation system

**Figure 30.2** Clinical manifestations of mitochondrial disease

## Epilepsy Phenotypes in Mitochondrial Disease

Seizure semiology has been reported in relatively few cohorts of mitochondrial epilepsy (Table 30.1). Epilepsy phenotypes frequently reported in mitochondrial diseases include neonatal refractory status and multiorgan failure, neonatal myoclonic epilepsy, infantile spasms, refractory/recurrent status epilepticus, epilepsy partialis continua and myoclonic epilepsy. Between 20 and 60% of patients have more than one seizure type [8, 9].

## Mitochondrial Syndromic Epilepsies

Some of the classical mitochondrial syndromes most often associated with epilepsy are listed in Table 30.2 and described in the following section.

### Epilepsy Associated with Mitochondrial DNA Mutations

#### MELAS

The syndrome of mitochondrial encephalomyopathy, lactic acidosis and stroke-like episodes (MELAS) usually

presents in the first decade of life with a stroke-like episode, often heralded by a migraine headache and/or focal seizure. Other clinical features include ataxia, sensorineural hearing loss, optic atrophy, cognitive decline, short stature, diabetes mellitus and cardiomyopathy. Eighty percent of cases with MELAS have the same maternally inherited mtDNA mutation, m.3243A>G in the *MT-TL1* gene encoding a transfer RNA (tRNA) for leucine. A review of 110 cases of MELAS revealed seizures in 96%, and myoclonus in 38% [12]. Seizures were the first symptom in 28% of cases. Sixty-two percent had generalised seizures, whilst focal seizures were present in 24%. It should be noted that the m.3243A>G mutation is associated with considerable clinical heterogeneity, and only 10% have the complete MELAS syndrome [13]. Thirty-four percent of patients with m.3243A>G had epilepsy in a large cohort [4]. Since m.3243A>G is present in 1 in 400 individuals (although most are oligosymptomatic or asymptomatic) this mutation makes a sizeable contribution to mitochondrial epilepsy.

**Table 30.1** Seizure semiology reported in mitochondrial disease cohorts

Study	Canafoglia <i>et al.</i> 2001 <i>Neurology</i> [10]	Lee <i>et al.</i> 2008 <i>Epilepsia</i> [11]	El Sabbagh <i>et al.</i> 2010 <i>Epilepsia</i> [9]	Chevallier <i>et al.</i> 2014 <i>Epilepsia</i> [8]
Size of cohort	<i>n</i> = 31	<i>n</i> = 48	<i>n</i> = 56	<i>n</i> = 165
Age range	Mixed (birth–44 years)	Paediatric (1–12 years)	Paediatric (birth–17 years)	Mixed (1–79 years)
Seizure semiology (% of cases)	<i>Juvenile/adult onset (n = 14)</i> Myoclonic epilepsy 43% Tonic–clonic seizures 21% Motor seizures 36% Somatosensory seizures 7% <i>Infantile onset (n = 17)</i> Partial epilepsy 65% Generalised epilepsy 24% West syndrome 12%	Partial seizures 17% Generalised epilepsy 29% Lennox–Gastaut 25% West syndrome 21% Ohtahara 4% Landau–Kleffner 4%	Myoclonic seizures 52% Infantile spasms 18% Focal seizures 23% Tonic seizures 9% Tonic–clonic seizures 9% Multiple seizure types ~60%	Generalised tonic–clonic 37% Complex partial 37% Myoclonus 22% Focal seizures 10% Infantile spasms 7% Atonic seizures 7% Generalised absence 2% Undefined 5% Status epilepticus 2% Epilepsia partialis continua 2% Nonconvulsive status epilepticus 3%

**Table 30.2** Mitochondrial syndromes associated with epilepsy

Syndrome	Clinical features	Seizure types	Gene defects	Inheritance
MELAS	Mitochondrial encephalomyopathy, lactic acidosis, stroke-like episodes, migraine, sensorineural hearing loss, cognitive decline, diabetes, cardiomyopathy	Focal seizures (motor, visual), myoclonic seizures, status epilepticus	<i>MT-TL1</i> (80%), other mtDNA point mutations	Maternal
MERRF	Myoclonic epilepsy with ragged-red fibres, ataxia, sensorineural hearing loss, cognitive decline, multiple lipomata	Progressive myoclonus, GTCS	<i>MT-TK</i> (80%), other mtDNA point mutations	Maternal
NARP	Neurogenic muscle weakness, ataxia, retinitis pigmentosa, sensorineural hearing loss, cognitive decline	GTCS	<i>MT-ATP6</i>	Maternal
Leigh	Subacute necrotising encephalomyelopathy: stepwise neurodevelopmental regression, hypotonia, dystonia, ataxia, ophthalmoparesis, nystagmus, optic atrophy	Multiple seizure types including infantile spasms and other generalised seizures	>89 genes (mtDNA and nuclear)	Maternal, AR, X-linked
Alpers	Neurodevelopmental delay/regression, intractable epilepsy, +/- liver failure	Refractory seizures, often focal initially (may be explosive onset) with subsequent generalisation, EPC, frequent status epilepticus	<i>POLG</i> , <i>FARS2</i> , <i>NARS2</i> , <i>PARS2</i>	AR
MEMSA	Myoclonic epilepsy, myopathy, sensory ataxia	Focal clonic or myoclonic seizures, EPC, GTCS, status epilepticus	<i>POLG</i>	AR

EPC epilepsy partialis continua; GTCS generalised tonic–clonic seizures

**MERRF**

Myoclonic epilepsy with ragged-red fibres (MERRF) is also caused by a maternally inherited mtDNA point mutation, m.8344A>G in the *MT-TK* gene encoding the tRNA for lysine. In an Italian cohort of 34 adults with MERRF, 35% had generalised epilepsy, 24% had myoclonus and 24% had cerebellar ataxia [14]. Other clinical features include myopathy, sensorineural hearing loss, pigmentary retinopathy, cognitive decline and multiple symmetrical lipomatosis. Rarely, patients with MERRF may have stroke-like episodes, and these may be preceded by status epilepticus.

**NARP**

The maternally inherited disorder neuropathy, ataxia and retinitis pigmentosa (NARP) is caused by point mutations in the *MT-ATP6* gene encoding a subunit of complex V. Individuals with the same mutations at high load (>95%) have maternally inherited Leigh syndrome (see below) and frequently present with seizures, often infantile spasms. Seizures are less frequently observed in adults with NARP but generalised tonic-clonic seizures have been reported, associated with generalized spike and wave (GSW) discharges on EEG [15].

**Single Mitochondrial Deletion Disorders**

Large-scale deletions of mtDNA are a relatively frequent cause of mitochondrial disease and are associated with a broad spectrum of phenotypes, ranging in severity from the infantile onset Pearson marrow pancreas syndrome to Kearns-Sayre syndrome to isolated progressive external ophthalmoplegia starting in adult life [16]. Although seizures may occur in Kearns-Sayre syndrome, they were uncommon in a large paediatric cohort [17].

**Epilepsy Associated with Nuclear Gene Defects****Alpers–Huttenlocher syndrome**

The Alpers–Huttenlocher syndrome was originally described as a clinical triad of intractable epilepsy, neurodevelopmental regression and liver failure with characteristic neuropathological features. Focal lesions typically affect the occipital and parietal cortex (especially the striate cortex) and are characterised by neuronal loss, vacuolation of the neuropil and astrocytosis [18]. However since the genetic basis of this disorder was identified as recessive mutations in *POLG*, encoding the catalytic subunit of DNA polymerase gamma (the only DNA polymerase in the mitochondrion), it is increasingly recognised that many affected individuals have seizures and developmental regression without hepatic involvement [19]. The liver failure may be precipitated by sodium valproate treatment (for this reason sodium valproate therapy is typically avoided in mitochondrial epilepsies) but may also occur in the absence of valproate exposure. Alpers–Huttenlocher syndrome is characterised by relentless disease progression and often leads to death in early childhood [19].

**MEGDEL**

The relatively recently reported disorder MEGDEL (3-methylglutaconic aciduria with deafness, encephalopathy, Leigh-like)

is caused by a recessive defect of lipid remodelling at the mitochondrial associated membranes and is frequently associated with seizures. Typically there is neonatal onset of feeding problems, failure to thrive, hypotonia and hypoglycaemia, often associated with hepatic dysfunction. By age 2 years progressive deafness, dystonia, spasticity and developmental regression are apparent. Seizures may start in the neonatal period or later in childhood [20].

**Epilepsy Associated with either Mitochondrial or Nuclear Gene Defects****Leigh Syndrome**

Leigh syndrome is the most frequent presentation of mitochondrial disease and is also traditionally a neuropathological diagnosis, namely focal bilaterally symmetrical subacute necrotic lesions characterised by demyelination, vacuolation, gliosis and capillary proliferation with relative preservation of neurons [21, 22]. However, Leigh syndrome is increasingly diagnosed on the basis of MRI features (bilateral symmetrical signal abnormality involving the basal ganglia and/or brainstem) in association with elevated lactate levels in blood and/or CSF and compatible clinical features [22]. The syndrome is characterised by stepwise neurological deterioration leading to early death, often in childhood. There is extreme genetic heterogeneity, with defects of >89 mitochondrial and nuclear genes linked to Leigh syndrome to date [23]. Thirty-nine percent of a multinational cohort of 130 children with Leigh syndrome had seizures, including 22% with generalised seizures, 14% with focal seizures and 6% with epileptic spasms [24].

**Nonsyndromic Mitochondrial Epilepsies**

Many patients with mitochondrial disease do not present with classical syndromic phenotypes and may have variable multisystem features. Furthermore, an unknown proportion of patients may have pure epilepsy presentations, especially infantile epileptic encephalopathies, since mitochondrial disease may not be suspected in the absence of multisystem manifestations.

**OXPHOS subunit and assembly factor deficiencies**

Epilepsy has been reported in association with mutations of subunits and assembly factors of all five OXPHOS complexes (Table 30.3). Isolated complex I deficiency accounts for ~25–30% of childhood mitochondrial disease and typical clinical presentations include Leigh syndrome and leukoencephalopathies, frequently with seizures [25]. Complex II deficiency is extremely rare but also presents with Leigh syndrome or a leukoencephalopathy with seizures, for example secondary to mutations in the SDHA or SDHB subunits or the SDHAF1 assembly factor. Isolated complex III deficiency is also relatively infrequent, but seizures have been reported in patients with mutations in the mitochondrial subunit cytochrome *b*, one nuclear subunit (UQCRC2) and four assembly factors (BCS1L, TTC19, HCCS and UQCRC2). Isolated complex IV (cytochrome oxidase, COX) deficiency is observed in ~25% cases of childhood onset mitochondrial disease and is genetically extremely heterogeneous, with reported causes including mutations in COX subunits and assembly factors and defects of

**Table 30.3** Genetic defects underlying the mitochondrial epilepsies

Disease mechanism	Biochemical defect	Gene defects and modes of inheritance
<i>OXPHOS subunit defects</i>	Complex I	<b>Mat:</b> MT-ND1 MT-ND3 MT-ND4 MT-ND5 MT-ND6  <b>AR:</b> NDUFV1 NDUFS2 NDUFS3 NDUFS4 NDUFS6 NDUFS7 NDUFS8 NDUF2A NDUF2B NDUF2C NDUF2D NDUF2E NDUF2F NDUF2G NDUF2H NDUF2I NDUF2J NDUF2K NDUF2L NDUF2M NDUF2N NDUF2O NDUF2P NDUF2Q NDUF2R NDUF2S NDUF2T NDUF2U NDUF2V NDUF2W NDUF2X NDUF2Y NDUF2Z NDUF2AA NDUF2AB NDUF2AC NDUF2AD NDUF2AE NDUF2AF NDUF2AG NDUF2AH NDUF2AI NDUF2AJ NDUF2AK NDUF2AL NDUF2AM NDUF2AN NDUF2AO NDUF2AP NDUF2AQ NDUF2AR NDUF2AS NDUF2AT NDUF2AU NDUF2AV NDUF2AW NDUF2AX NDUF2AY NDUF2AZ NDUF2BA NDUF2BB NDUF2BC NDUF2BD NDUF2BE NDUF2BF NDUF2BG NDUF2BH NDUF2BI NDUF2BJ NDUF2BK NDUF2BL NDUF2BM NDUF2BN NDUF2BO NDUF2BP NDUF2BQ NDUF2BR NDUF2BS NDUF2BT NDUF2BU NDUF2BV NDUF2BW NDUF2BX NDUF2BY NDUF2BZ NDUF2CA NDUF2CB NDUF2CC NDUF2CD NDUF2CE NDUF2CF NDUF2CG NDUF2CH NDUF2CI NDUF2CJ NDUF2CK NDUF2CL NDUF2CM NDUF2CN NDUF2CO NDUF2CP NDUF2CQ NDUF2CR NDUF2CS NDUF2CT NDUF2CU NDUF2CV NDUF2CW NDUF2CX NDUF2CY NDUF2CZ NDUF2DA NDUF2DB NDUF2DC NDUF2DD NDUF2DE NDUF2DF NDUF2DG NDUF2DH NDUF2DI NDUF2DJ NDUF2DK NDUF2DL NDUF2DM NDUF2DN NDUF2DO NDUF2DP NDUF2DQ NDUF2DR NDUF2DS NDUF2DT NDUF2DU NDUF2DV NDUF2DW NDUF2DX NDUF2DY NDUF2DZ NDUF2EA NDUF2EB NDUF2EC NDUF2ED NDUF2EE NDUF2EF NDUF2EG NDUF2EH NDUF2EI NDUF2EJ NDUF2EK NDUF2EL NDUF2EM NDUF2EN NDUF2EO NDUF2EP NDUF2EQ NDUF2ER NDUF2ES NDUF2ET NDUF2EU NDUF2EV NDUF2EW NDUF2EX NDUF2EY NDUF2EZ NDUF2FA NDUF2FB NDUF2FC NDUF2FD NDUF2FE NDUF2FF NDUF2FG NDUF2FH NDUF2FI NDUF2FJ NDUF2FK NDUF2FL NDUF2FM NDUF2FN NDUF2FO NDUF2FP NDUF2FQ NDUF2FR NDUF2FS NDUF2FT NDUF2FU NDUF2FV NDUF2FW NDUF2FX NDUF2FY NDUF2FZ NDUF2GA NDUF2GB NDUF2GC NDUF2GD NDUF2GE NDUF2GF NDUF2GG NDUF2GH NDUF2GI NDUF2GJ NDUF2GK NDUF2GL NDUF2GM NDUF2GN NDUF2GO NDUF2GP NDUF2GQ NDUF2GR NDUF2GS NDUF2GT NDUF2GU NDUF2GV NDUF2GW NDUF2GX NDUF2GY NDUF2GZ NDUF2HA NDUF2HB NDUF2HC NDUF2HD NDUF2HE NDUF2HF NDUF2HG NDUF2HH NDUF2HI NDUF2HJ NDUF2HK NDUF2HL NDUF2HM NDUF2HN NDUF2HO NDUF2HP NDUF2HQ NDUF2HR NDUF2HS NDUF2HT NDUF2HU NDUF2HV NDUF2HW NDUF2HX NDUF2HY NDUF2HZ NDUF2IA NDUF2IB NDUF2IC NDUF2ID NDUF2IE NDUF2IF NDUF2IG NDUF2IH NDUF2II NDUF2IJ NDUF2IK NDUF2IL NDUF2IM NDUF2IN NDUF2IO NDUF2IP NDUF2IQ NDUF2IR NDUF2IS NDUF2IT NDUF2IU NDUF2IV NDUF2IW NDUF2IX NDUF2IY NDUF2IZ NDUF2JA NDUF2JB NDUF2JC NDUF2JD NDUF2JE NDUF2JF NDUF2JG NDUF2JH NDUF2JI NDUF2JJ NDUF2JK NDUF2JL NDUF2JM NDUF2JN NDUF2JO NDUF2JP NDUF2JQ NDUF2JR NDUF2JS NDUF2JT NDUF2JU NDUF2JV NDUF2JW NDUF2JX NDUF2JY NDUF2JZ NDUF2KA NDUF2KB NDUF2KC NDUF2KD NDUF2KE NDUF2KF NDUF2KG NDUF2KH NDUF2KI NDUF2KJ NDUF2KK NDUF2KL NDUF2KM NDUF2KN NDUF2KO NDUF2KP NDUF2KQ NDUF2KR NDUF2KS NDUF2KT NDUF2KU NDUF2KV NDUF2KW NDUF2KX NDUF2KY NDUF2KZ NDUF2LA NDUF2LB NDUF2LC NDUF2LD NDUF2LE NDUF2LF NDUF2LG NDUF2LH NDUF2LI NDUF2LJ NDUF2LK NDUF2LL NDUF2LM NDUF2LN NDUF2LO NDUF2LP NDUF2LQ NDUF2LR NDUF2LS NDUF2LT NDUF2LU NDUF2LV NDUF2LW NDUF2LX NDUF2LY NDUF2LZ NDUF2MA NDUF2MB NDUF2MC NDUF2MD NDUF2ME NDUF2MF NDUF2MG NDUF2MH NDUF2MI NDUF2MJ NDUF2MK NDUF2ML NDUF2MM NDUF2MN NDUF2MO NDUF2MP NDUF2MQ NDUF2MR NDUF2MS NDUF2MT NDUF2MU NDUF2MV NDUF2MW NDUF2MX NDUF2MY NDUF2MZ NDUF2NA NDUF2NB NDUF2NC NDUF2ND NDUF2NE NDUF2NF NDUF2NG NDUF2NH NDUF2NI NDUF2NJ NDUF2NK NDUF2NL NDUF2NM NDUF2NN NDUF2NO NDUF2NP NDUF2NQ NDUF2NR NDUF2NS NDUF2NT NDUF2NU NDUF2NV NDUF2NW NDUF2NX NDUF2NY NDUF2NZ NDUF2OA NDUF2OB NDUF2OC NDUF2OD NDUF2OE NDUF2OF NDUF2OG NDUF2OH NDUF2OI NDUF2OJ NDUF2OK NDUF2OL NDUF2OM NDUF2ON NDUF2OO NDUF2OP NDUF2OQ NDUF2OR NDUF2OS NDUF2OT NDUF2OU NDUF2OV NDUF2OW NDUF2OX NDUF2OY NDUF2OZ NDUF2PA NDUF2PB NDUF2PC NDUF2PD NDUF2PE NDUF2PF NDUF2PG NDUF2PH NDUF2PI NDUF2PJ NDUF2PK NDUF2PL NDUF2PM NDUF2PN NDUF2PO NDUF2PP NDUF2PQ NDUF2PR NDUF2PS NDUF2PT NDUF2PU NDUF2PV NDUF2PW NDUF2PX NDUF2PY NDUF2PZ NDUF2QA NDUF2QB NDUF2QC NDUF2QD NDUF2QE NDUF2QF NDUF2QG NDUF2QH NDUF2QI NDUF2QJ NDUF2QK NDUF2QL NDUF2QM NDUF2QN NDUF2QO NDUF2QP NDUF2QQ NDUF2QR NDUF2QS NDUF2QT NDUF2QU NDUF2QV NDUF2QW NDUF2QX NDUF2QY NDUF2QZ NDUF2RA NDUF2RB NDUF2RC NDUF2RD NDUF2RE NDUF2RF NDUF2RG NDUF2RH NDUF2RI NDUF2RJ NDUF2RK NDUF2RL NDUF2RM NDUF2RN NDUF2RO NDUF2RP NDUF2RQ NDUF2RR NDUF2RS NDUF2RT NDUF2RU NDUF2RV NDUF2RW NDUF2RX NDUF2RY NDUF2RZ NDUF2SA NDUF2SB NDUF2SC NDUF2SD NDUF2SE NDUF2SF NDUF2SG NDUF2SH NDUF2SI NDUF2SJ NDUF2SK NDUF2SL NDUF2SM NDUF2SN NDUF2SO NDUF2SP NDUF2SQ NDUF2SR NDUF2SS NDUF2ST NDUF2SU NDUF2SV NDUF2SW NDUF2SX NDUF2SY NDUF2SZ NDUF2TA NDUF2TB NDUF2TC NDUF2TD NDUF2TE NDUF2TF NDUF2TG NDUF2TH NDUF2TI NDUF2TJ NDUF2TK NDUF2TL NDUF2TM NDUF2TN NDUF2TO NDUF2TP NDUF2TQ NDUF2TR NDUF2TS NDUF2TT NDUF2TU NDUF2TV NDUF2TW NDUF2TX NDUF2TY NDUF2TZ NDUF2UA NDUF2UB NDUF2UC NDUF2UD NDUF2UE NDUF2UF NDUF2UG NDUF2UH NDUF2UI NDUF2UJ NDUF2UK NDUF2UL NDUF2UM NDUF2UN NDUF2UO NDUF2UP NDUF2UQ NDUF2UR NDUF2US NDUF2UT NDUF2UU NDUF2UV NDUF2UW NDUF2UX NDUF2UY NDUF2UZ NDUF2VA NDUF2VB NDUF2VC NDUF2VD NDUF2VE NDUF2VF NDUF2VG NDUF2VH NDUF2VI NDUF2VJ NDUF2VK NDUF2VL NDUF2VM NDUF2VN NDUF2VO NDUF2VP NDUF2VQ NDUF2VR NDUF2VS NDUF2VT NDUF2VU NDUF2VV NDUF2VW NDUF2VX NDUF2VY NDUF2VZ NDUF2WA NDUF2WB NDUF2WC NDUF2WD NDUF2WE NDUF2WF NDUF2WG NDUF2WH NDUF2WI NDUF2WJ NDUF2WK NDUF2WL NDUF2WM NDUF2WN NDUF2WO NDUF2WP NDUF2WQ NDUF2WR NDUF2WS NDUF2WT NDUF2WU NDUF2WV NDUF2WW NDUF2WX NDUF2WY NDUF2WZ NDUF2XA NDUF2XB NDUF2XC NDUF2XD NDUF2XE NDUF2XF NDUF2XG NDUF2XH NDUF2XI NDUF2XJ NDUF2XK NDUF2XL NDUF2XM NDUF2XN NDUF2XO NDUF2XP NDUF2XQ NDUF2XR NDUF2XS NDUF2XT NDUF2XU NDUF2XV NDUF2XW NDUF2XX NDUF2XY NDUF2XZ NDUF2YA NDUF2YB NDUF2YC NDUF2YD NDUF2YE NDUF2YF NDUF2YG NDUF2YH NDUF2YI NDUF2YJ NDUF2YK NDUF2YL NDUF2YM NDUF2YN NDUF2YO NDUF2YP NDUF2YQ NDUF2YR NDUF2YS NDUF2YT NDUF2YU NDUF2YV NDUF2YW NDUF2YX NDUF2YY NDUF2YZ NDUF2ZA NDUF2ZB NDUF2ZC NDUF2ZD NDUF2ZE NDUF2ZF NDUF2ZG NDUF2ZH NDUF2ZI NDUF2ZJ NDUF2ZK NDUF2ZL NDUF2ZM NDUF2ZN NDUF2ZO NDUF2ZP NDUF2ZQ NDUF2ZR NDUF2ZS NDUF2ZT NDUF2ZU NDUF2ZV NDUF2ZW NDUF2ZX NDUF2ZY NDUF2ZZ

**Table 30.3** (cont.)

Disease mechanism	Biochemical defect	Gene defects and modes of inheritance
<i>Pyruvate dehydrogenase deficiency</i>	Pyruvate dehydrogenase	<b>AR:</b> PDHB DLD PDHX <b>XL:</b> PDHA1
<i>Defects of cofactor biosynthesis and transport</i>	Coenzyme Q <sub>10</sub>	<b>AR:</b> PDSS2 COQ2 COQ4 COQ6 COQ9 COQ8A COQ8B
	Lipoic acid	<b>AR:</b> LIAS
	Iron sulphur clusters	<b>AR:</b> NFU1 BOLA3 NFS1
	Thiamine	<b>AR:</b> SLC19A3 SLC25A19 TPK1
	NADP	<b>AR:</b> NADK
	Manganese	<b>AR:</b> SLC39A8
<i>Mitochondrial import</i>	Multiple RC enzymes or normal enzyme activities	<b>AR:</b> SLC25A1 SLC25A12 SLC25A22 DNAJC19 TIMM50 AIFM1
<i>Defects of mitochondrial lipid membranes and dynamics</i>	Multiple RC enzymes or normal enzyme activities	<b>AR:</b> SERAC1 AGK DNM1L MFF STAT2 SLC25A46
<i>Defects of mitochondrial quality control</i>	Multiple RC enzymes or normal enzyme activities	<b>AR:</b> CLPB AFG3L2 HSPD1 HTRA2 MIPEP SACS
<i>Respiratory chain toxicity</i>	Multiple RC enzymes, isolated complex IV deficiency or normal enzyme activities	<b>AR:</b> ETHE1 HIBCH ECHS1 NAXE
<i>Other mechanisms</i>	Multiple RC enzymes, isolated complex IV deficiency or normal enzyme activities	<b>AR:</b> APOPT1 TXN2 PPA2

AR autosomal recessive; Mat maternal inheritance; NADP nicotinamide adenine dinucleotide phosphate; XL X-linked.

mtDNA maintenance or mitochondrial translation. Furthermore, COX deficiency may sometimes be a secondary phenomenon with a non-mitochondrial underlying cause. Mutations of the maternally inherited complex V subunit ATP6 have been associated with maternally inherited Leigh syndrome, and affected individuals may present with infantile spasms. Few mutations have been described in nuclear encoded subunits of complex V, but patients with mutations in the ATP5A1 subunit were reported to have seizures, as were patients with mutations in the assembly factors ATPAF2 and TMEM70.



### Mitochondrial DNA Depletion Syndromes

Patients with the mtDNA depletion syndrome (MDDS) have a severe quantitative defect of mtDNA. In the most severely affected cases this may be <5–10% of levels observed in tissues from age-matched healthy control subjects [26]. The underlying problem is an inability to maintain the mtDNA, either because of a defect in the replication machinery, or because of impaired nucleoside supply for mtDNA synthesis. Phenotypically, MDDS can be divided into myopathic, encephalomyopathic and hepatocerebral subgroups. By far the most frequent cause of MDDS is mutation of *POLG*. The Alpers–Huttenlocher syndrome (see above) is one form of hepatocerebral MDDS caused by recessive *POLG* mutations, but recessive *POLG* mutations can also cause other epilepsy syndromes in older children and adolescents, which are collectively known as myoclonic epilepsy, myopathy, sensory ataxia (MEMSA). Other genetic causes of MDDS associated with seizures include mutations of *TWNK* encoding the twinkie helicase, *RRM2B* encoding a subunit of the ribonucleotide reductase, *DGUOK*, *TK2*, *SUCLA2*, *SUCLG1*, *TYMP* and *MPV17* encoding proteins implicated in mitochondrial deoxynucleotide homeostasis, *ABAT* encoding GABA transaminase, and *FBXL4* encoding a protein whose precise function remains unknown [27].

### Disorders of Mitochondrial Translation

In recent years there has been an explosion of reports of defects of mitochondrial translation, particularly mutations of the tRNA synthetases required for aminoacylation of the 22 mitochondrial tRNA molecules prior to their attachment to the mitochondrial ribosome for protein synthesis. Mutations in *RARS2* are associated with intractable seizures and developmental stasis from birth, typically with initial severe lactic acidosis (which later resolves) and pontocerebellar hypoplasia on neuro-imaging, whilst mutations in *FARS2*, *NARS2* and *PARS2* have been reported to cause an Alpers-like phenotype [28]. Other disorders of mitochondrial translation associated with seizures include several defects of mitochondrial tRNA modification and mutations in the mitochondrial translation factors GFM1 and TSFM (Table 30.3).

### Disorders of Cofactor Biosynthesis

Another rapidly growing group of mitochondrial epilepsies are disorders of cofactor biosynthesis, particularly coenzyme Q<sub>10</sub> (CoQ<sub>10</sub>) and iron–sulphur (Fe–S) clusters. CoQ<sub>10</sub> is a lipophilic electron carrier transferring electrons from fatty acid  $\beta$ -oxidation and OXPHOS complexes I and II to complex III, and also acts as a potent intramitochondrial antioxidant. Eight defects of CoQ<sub>10</sub> biosynthesis have now been reported and seven of these are characterised by epilepsy associated with the following clinical features: infantile multisystem encephalomyopathies (*PDSS2*, *COQ2*, *COQ4* and *COQ9*), ataxia (*COQ8A*) or steroid-resistant nephrotic syndrome (*COQ6*, *COQ8B*) [29]. Fe–S clusters are required for the function of several mitochondrial enzymes, including respiratory chain enzyme complexes I–III, the Krebs cycle enzyme aconitase and the electron transfer flavoprotein dehydrogenase. Several

disorders of Fe–S cluster biosynthesis have been reported [30], associated with epilepsy in some cases (*NFU1*, *BOLA3*, *LIAS*, *NFS1*). Defects of supply of other cofactors (including thiamine, NADP, copper and manganese) to the respiratory chain have also been linked to epilepsy (see Table 30.3 for a comprehensive list).

### Disorders of Mitochondrial Import

Mitochondria have complex mechanisms for importing hundreds of solutes and proteins. Mutations in several solute transporters have been linked to epilepsy, including the citrate transporter (*SLC25A1*) and two mitochondrial glutamate carriers (*SLC25A12* and *SLC25A22*). Mutations in *SLC25A22* in particular have been reported in several families associated with severe epilepsy syndromes, including early myoclonic epilepsy with burst suppression and migrating partial seizures of infancy. Defects of mitochondrial protein importers associated with epilepsy include mutations in *TIMM50* and *DNAJC19*.

### Disorders of Mitochondrial Membrane Lipids and Dynamics

Mitochondria are dynamic organelles, constantly undergoing fission and fusion events. Several defects of mitochondrial dynamics have been reported, associated with epilepsy in some cases. For example, several infants with mutations in *DNM1L*, encoding the major mitochondrial fission protein, had intractable seizures. More recently mutations in the mitochondrial fission factor (MFF) and the JAK-STAT cytokine STAT2 have also been reported to cause epilepsy [31].

### Other Disease Mechanisms

Relatively new disease mechanisms for mitochondrial disease recognised more recently include disorders of mitochondrial quality control (e.g. *CLPB*, *AFG3L2*, *HSPD1*, *HTRA2*, *MIPEP* and *SACS* mutations) and toxic damage to the respiratory chain, for example in ethylmalonic encephalopathy and HIBCH and *ECHS1* deficiencies (two disorders of valine degradation) [32, 33].

## Secondary Mitochondrial Epilepsies

Various disorders with prominent epilepsy have been reported to cause secondary respiratory chain dysfunction, for example GM3 synthase deficiency, infantile epileptic encephalopathy caused by *STXBPI* mutations and Dravet syndrome associated with *SCN1A* mutations [34–38].

## Diagnostic Tests for Mitochondrial Disease

Mitochondrial diagnostics is currently in a state of flux, as the field moves from muscle biopsy as a first stage diagnostic test to NGS as the frontline test. However, even when NGS approaches are being utilised, it is important to take a multidisciplinary approach to mitochondrial diagnostics, encompassing multisystem clinical, imaging, neurophysiological, and biochemical (and, where available, histological) information as well as genetic data.

## EEG

The most frequent EEG abnormality reported in mitochondrial disease is nonspecific slowing. EEGs were systematically performed in 109 subjects in one cohort and were abnormal in 61% of these [8]. The most frequent abnormalities were combined epileptiform discharges and slowing (33%), epileptiform discharges (22%), generalised slowing (7%) and focal slowing (3%). In this study 39% of epileptic discharges were focal, and multifocal, generalised or both in the remaining 61%. EEG findings are particularly useful in suggesting the possibility of *POLG* mutations. In Alpers syndrome the initial EEG may reveal unilateral occipital rhythmic high-amplitude delta with superimposed (poly)spikes (RHADS), although later in the disease EEG changes may be generalised and nonspecific [39]. Patients with other *POLG* epileptic syndromes such as MEMSA also have predominantly occipital changes, typically occipital slow wave and epileptic activity early in the disease course [40].

## Imaging

Characteristic neuro-imaging changes are observed in some mitochondrial syndromes. For example, patients with Leigh syndrome typically have bilateral symmetrical T2 hyperintense lesions in the basal ganglia and/or brainstem, and those with MELAS have stroke-like lesions in parieto-occipital regions. Other patients may have a leukoencephalopathy (e.g. those with mutations of subunits of complexes I and II) or pontocerebellar hypoplasia (*RARS2*). However, in many cases with mitochondrial epilepsy neuro-imaging appearances may be nonspecific, such as cerebral or cerebellar atrophy, or even normal.

## Metabolic Investigations

Various metabolites may be elevated in mitochondrial disease, including lactate, the amino acid alanine, fibroblast growth factor 21 (FGF21) and growth and differentiation factor 15 (GDF15) [41, 42]. However none has universally increased levels in mitochondrial disease, making these metabolites unreliable as screening tests.

## Muscle Biopsy

Traditionally, diagnosis of mitochondrial disease rested on biopsy of skeletal muscle (or occasionally another affected tissue such as liver or heart). An open or large bore needle biopsy is required to provide sufficient tissue for the necessary analyses: histochemical, ultrastructural, biochemical and genetic. The ragged-red fibre (characterised by subsarcolemmal proliferation of mitochondria visualised by the modified Gomori trichrome stain), is traditionally viewed as the morphological hallmark of mitochondrial disease, but is mainly a feature of adult mitochondrial disease associated with defective mtDNA (including point mutations and single or multiple deletions). Ragged-red fibres are rarely seen in the paediatric population, but may be a feature of MDDS or disease caused by mtDNA point mutations, including MELAS and MERRF.

Electron microscopy may reveal ultrastructural abnormalities of the mitochondria, such as whorled cristae or crystalline inclusions, but features may be nonspecific. Biochemical investigation of biopsied muscle may include spectrophotometric assays of individual OXPHOS complexes, interrogation of global mitochondrial function using polarography, or assessment of OXPHOS complex assembly using blue native gel electrophoresis. Genetic investigation of mtDNA in muscle may include screening for rearrangements, quantitation of mtDNA copy number, and full sequencing of the mitochondrial genome.

## Genetic Testing

Single gene testing is appropriate when clinical, imaging or EEG data point to a specific syndromic diagnosis, such as MELAS or Alpers syndrome (test for m.3243A>G or *POLG* mutations respectively). However, since >300 genes have now been linked to mitochondrial disease, and >140 of these may be associated with seizures (Table 30.3), single gene testing is frequently an inefficient diagnostic strategy, and NGS is increasingly being employed as a first-line genetic diagnostic test [43]. NGS may be of a candidate gene panel, such as genes implicated in epileptic encephalopathies or the mitochondrial proteome, or of the whole exome or even whole genome sequencing.

## Assessment of Multisystem Involvement

Screening for multisystem involvement, which is a typical finding in infantile and childhood onset mitochondrial disease, is important since it may help to identify a specific syndromic diagnosis. Furthermore, knowledge of multisystem disease manifestations allows appropriate supportive therapy to be provided.

## Principles of Management

### Management of Mitochondrial Disease

Curative therapies for mitochondrial disease are lacking, but many supportive therapies are available for the multisystem manifestations of mitochondrial disorders, for example brow suspension surgery for severe ptosis, hearing aids, replacement of deficient hormones, gastrostomy feeding and pacing for cardiac conduction defects [44]. Other treatments include correction of electrolyte imbalances in patients with renal tubulopathies, CoQ<sub>10</sub> replacement for disorders of CoQ<sub>10</sub> biosynthesis, folinic acid supplementation in cases with cerebral folate deficiency, and L-arginine or L-citrulline supplementation in MELAS syndrome or if plasma levels are low [45]. The role of cofactor and antioxidant supplementation is controversial and currently there is no supportive evidence for these treatments [46].

### Antiepileptic Drugs

Mitochondrial epilepsy is frequently refractory to drug therapy, and the overwhelming majority of affected individuals

receive two or more AEDs [9, 47]. Randomised controlled trials have not been performed to determine which AEDs are most efficacious in mitochondrial disease. However, the clinical and genetic heterogeneity of mitochondrial epilepsy, with many underlying disease mechanisms, would likely hamper the design and interpretation of such trials. Combinations of a sodium channel blocker such as lamotrigine and a benzodiazepine (e.g. clobazam), together with levetiracetam or topiramate as needed, appear to be relatively efficacious, particularly in *POLG*-related epilepsy, the most common mitochondrial epilepsy [47, 48]. Other therapies which have been reported in *POLG* disease include magnesium infusion, L-carnitine supplementation and folinic acid [47], but the anecdotal nature of these reports and the unpredictable natural history of mitochondrial disease make it difficult to ascertain whether these agents are truly efficacious. Sodium valproate is absolutely contraindicated in patients with *POLG* mutations, because of the risk of inducing liver failure [39], but may be well tolerated and effective in controlling seizures in patients with other types of mitochondrial epilepsy. However, most specialists would prefer to use another AED in patients with mitochondrial disease, and only use valproate if other AEDs have failed.

## Experimental Approaches

The ketogenic diet (a high fat, low carbohydrate diet) has been suggested as a treatment for mitochondrial epilepsy, but so far experience is limited to case reports and small

case series [2]. Although a formal clinical trial would help to determine whether the ketogenic diet is efficacious, a modified Atkins diet was poorly tolerated in a small series of patients with mitochondrial myopathy, causing muscle pain and elevation of creatine kinase [49]. This may therefore not be an acceptable treatment for mitochondrial disease. Another option could be supplementation with the fatty acid decanoic acid, which is thought to be the active component of the ketogenic diet and appears to stimulate mitochondrial biogenesis in cell models [50, 51], but clinical trials have yet to be performed. Other experimental approaches currently being explored on a research basis include the development of novel antioxidants and agents to stimulate mitochondrial biogenesis, nucleoside supplementation for some forms of MDDS, and gene therapy for both mtDNA and nuclear gene encoded mitochondrial disorders [45, 52].

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