ORIGINAL ARTICLE



High Cytochrome c Oxidase Expression Links to Severe Skeletal Energy Failure by ³¹P-MRS Spectroscopy in Mitochondrial Encephalomyopathy, Lactic Acidosis, and Stroke-Like Episodes

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Keywords

³¹P-MRS; Cytochrome c oxidase; Energy metabolism; Mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes; Muscle biopsy.

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SUMMARY

Aims: The purpose of this study was to evaluate the energy metabolism and mitochondrial function in skeletal muscle from patients with Mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes (MELAS) or chronic progressive external ophthalmoplegia (CPEO) using phosphorus magnetic resonance spectroscopy (³¹P-MRS), to determine whether abnormally increasing cytochrome c oxidase (COX), as detected in muscle biopsy, could be a cause for MELAS. **Methods:** ³¹P-MRS was performed on the quadriceps femoris muscle of 12 healthy volunteers and 11 patients diagnosed as MELAS or CPEO by muscle biopsy and genetic analysis. All subjects experienced a state of rest, 5-min exercise, and 5-min recovery protocol in a supine position. **Results:** Compared to CPEO, MELAS patients typically exhibited COX-positive ragged-red fibers (RRFs) as well as strongly SDH-positive blood vessels (SSVs). However, based on ³¹P-MRS results, MELAS showed a higher inorganic phosphate (Pi)/phosphocreatine (PCr) ratio and lower ATP/PCr ratio during exercise and delayed Pi/PCr and ATP/PCr recovery to normal. **Conclusions:** This study suggests that high COX expression contributes to severe skeletal energy failure by ³¹P-MRS spectroscopy in MELAS.

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Introduction

Mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes (MELAS) is the most common subtype of mitochondrial disorders associated with severe mitochondrial dysfunction and energy abnormalities. The typical pathological characteristics of MELAS are massive mitochondrial proliferation and defects of respiratory chain enzymes. Until now, the pathogenesis of MELAS has been unclear. Previous studies proposed mtDNA mutationinduced mitochondrial neuropathy as the underlying cause, but this theory cannot fully explain the clinical phenotype and the molecular defect relationship.

An alternative explanation suggested cytochrome c oxidase (COX), the fourth complex of respiratory chain, as a specific pathology seen in MELAS [1]. This theory is based on two aspects:

(1) an increased COX activity compared to other mitochondrial encephalopathy such as chronic progressive external ophthalmoplegia (CPEO), in which COX activity was significantly absent, and (2) an observed hypocitrullinemia in MELAS patients. Citrulline is a substrate donor for nitric oxide (NO) generation. It has been proposed that excessive COX activity in RRFs could lead to dysfunction of NO catabolism and subsequent low plasma citrulline levels [2]. However, this interesting hypothesis lacks direct and efficient evidence to assess muscle weakness and exercise intolerance, that is to say, the skeletal muscle energy metabolism itself.

³¹P-MRS is a sensitive and specific method to assess skeletal energy metabolism *in vivo*. ³¹P-MRS allows for noninvasive quantification of high-energy phosphate compounds. A typical ³¹P-MRS from muscles displays seven major peaks representing inorganic phosphate (Pi), phosphocreatine (PCr), phosphomonoester (PME), phosphodiester (PDE), and three peaks of adenosine triphosphate (α -ATP, β -ATP, γ -ATP). Several ³¹P-MRS studies have attempted to investigate abnormal skeletal muscle energy metabolism in patients with mitochondrial encephalomyopathy [3,4]. However, few studies have compared the energy metabolism between MELAS and CPEO patients.

This study attempts to assess energy metabolism and mitochondrial function by 3.0T ³¹P-MRS during a rest-exercise-recovery protocol in MELAS and CPEO patients. These results allow this study to: (1) address the relationship among energy metabolism, respiratory defect (revealed by COX), and mitochondrial proliferation (revealed by RRFs and SSVs), and (2) investigate whether the abnormal COX activity could result in impaired energy metabolism in MELAS patients.

Materials and methods

Patient Selection

From January 2010 to December 2012, eleven patients with the mitochondrial encephalomyopathy were recruited from Drum Tower Hospital of Nanjing University Medical School, Jiangsu Province Hospital of Nanjing Medical University and Nanjing General Hospital of Nanjing Military Command. Twelve healthy control volunteers were recruited and matched in age, but not in gender. The studies were approved by the local ethics committee, and all subjects gave written informed consent.

Mitochondrial Disease Criteria Score

Mitochondrial disease criteria (MDC) score is an improved consensus scoring system as established in Europe in 2006 [5]. The MDC score is based on an evaluation that includes clinical signs and symptoms (I) (muscle presentation IA, central nervous system (CNS) abnormalities IB, multisystem involvement IC), metabolic/ imaging studies (II), and muscle histology (III) (Table S1). Patients in the current study had multiple laboratory examinations including electrocardiography (EKG), electromyography (EMG), electroencephalography (EEG), serum lactate/alanine or cerebrospinal fluid (CSF), lactate/pyruvate ratio and magnetic resonance imaging (MRI) scans, and additional magnetic resonance spectroscopy (MRS). The data from patients' clinical, biochemical, and histological examinations were reviewed and converted into MDC scores, then were compared between MELAS and CPEO groups.

Pathological Study (Muscle Biopsy)

The quadriceps femoris muscle (vastus lateralis) was chosen as the most suitable muscle for biopsy because: (1) a relatively large sample can be acquired without damaging major nerves or blood vessels, and (2) most mitochondrial encephalomyopathy patients have proximal muscle fatigue. The surgical muscle biopsy was performed at the Pathology Department in Drum Tower Hospital. Serial cryostat sections of fresh frozen tissue were stained with routine histochemical reactions. These were generally modified stains including hematoxylin–eosin (HE), adenosine triphosphatase (ATPase pH 9.4, pH 4.6), modified Gomori trichrome (MGT),

periodic acid-Schiff reaction (PAS), oil red O, nicotinamide adenine dinucleotide-tetrazolium reductase (NADH-TR), COX, and succinate dehydrogenase (SDH). For MELAS, COX oxidative enzymatic stains were essential.

Molecular Analysis

All patients received gene confirmation. Total mitochondrial DNA was extracted (eight skeletal muscle samples, three urinary sediment samples) using commercial DNA purification kits (Promega Corporation., Madison, WI, USA). Genotype was detected by polymerase chain reaction (PCR) amplification and sequencing to determine the presence or absence of mtDNA mutation as previously described [6]. The gene mutation in MELAS syndrome is mtDNA 3243 A>G, and in CPEO syndrome, it is a 4977 bp large-scale mtDNA deletion.

Maximum Voluntary Isometric Contraction

To apply an individually adjusted exercise load, the maximum voluntary isometric contraction (MVC) of the quadriceps femoris was measured prior to MRS as previously described [7]. A sandbag load was fastened to the right ankle. Each subject was instructed to lie in a supine position and encouraged to raise the legs to maximize the effect of the ankle load. They were asked not to move the knee and maintain it straight in order to perform maximal quadriceps femoris contraction. Each contraction was maintained for 3 seconds, and the process was repeated after a resting period of 30 seconds. MVC was quantified in three repeated exercises and defined as the average of the three peak measurements [8].

³¹P-MRS Acquisition

Measurements were performed on a 3.0T MRI scanner (Achieva 3.0T TX; Philips Medical Systems, Eindhoven, the Netherlands) [9]. The subjects were placed in a supine position in the magnet bore. They were in a state of rest, exercise, and recovery protocol. For exercise, a weight of 25% MVC was loaded to the right ankle and subjects were asked to kick the ankle to contract the quadriceps muscle as mentioned above. Spectra measurements of Pi, PCr, and α -, β -, γ -ATP were acquired from a 50-mm-diameter surface coil on the surface of the quadriceps femoris. ³¹P-MRS parameters included 31 P-SV, TR = 5000 mseconds, TE = 0.1 mseconds, volume of interest (VOI) = $30 \times 30 \times 120$ mm, NSA 8, Dyn. Scan time for each spectrum was 50 seconds. Spectra were analyzed using a commercially available postprocessing workstation (Extended Workspace [EWS], Philips Medical Systems). Each MR investigation lasted nearly 14 min and comprised three phases: Phase 1: 100 seconds of resting measurements (baseline) of the unloaded resting muscle; Phase 2: 5 min of dynamic loaded with a 25% MVC in resistance; and Phase 3: 5 min of recovery measurements. PME and PDE data were not acquired as they were regarded to play no role in skeletal muscle energy metabolism [10]. The ratios of Pi/PCr and ATP/PCr were calculated at rest, 5-min exercise, and 5-min recovery time phases. Pi/PCr ratio was considered a sensitive indicator of efficiently generating and maintaining high-energy phosphate

COX Links to Energy Failure in MELAS by ³¹P-MRS

compounds [11]. This ratio up to the control level indicated a defect in production and utilization of energy. ATP/PCr ratio was considered a direct evidence of energy generation and maintenance. The ATP value was equal to the sum of the α -ATP, β -ATP, and γ -ATP, three different subtypes of ATP.

Statistical Analyses

The data were expressed as the mean \pm SEM, and the statistical analyses were performed using SPSS 13.0 software (SPSS, Chicago, IL, USA). Comparison of quantitative parameters between the two groups was evaluated using Student's *t*-test, and multiple group comparisons were analyzed by one-way analysis of variance (ANOVA), followed by least significant difference (LSD) or Student–Newman–Keuls (SNK) comparisons. *P*-value below 0.05 was considered significant.

Results

Clinical Information and MDC Scores

According to muscle biopsy and gene detection, six patients were diagnosed as MELAS and five were CPEO. The clinical information and MDC score evaluations of these patients are shown in Table 1. Common symptoms were proximal muscle weakness, exercise intolerance, and muscular atrophy, as well as other secondary manifestations including numbness, bulbar palsy, intention tremor, depression, ataxia, and cardiac conduction defects cardiomyopathy. The average age was 24.91 ± 6.05 years. One patient died during this study. The control group consisted of 12 matched volunteers (25.00 ± 2.45 years). There were no significant differences (P > 0.05) between the total scores of the MELAS group (9.17 ± 0.61) and the CPEO group (8.80 ± 0.86), indicating the two groups had similar disease severity.

Muscle Histopathology in MELAS and CPEO

The morphological anomalies of muscle biopsy associated with mtDNA alteration brought a unique opportunity to investigate the relationship between mitochondrial anomalies and respiratory activity. On the one hand, the abnormal mitochondrial changes included the appearance of ragged-red fibers (RRFs) on muscle biopsies in both MELAS and CPEO. The reason for pathological RRFs was an abundance of proliferated mitochondrial. As illustrated in Figure 1A, HE staining showed variations in myofiber size and several fibers with basophilic sarcoplasmic masses indicative of RRFs. Gomori staining clearly demonstrated RRFs which were irregular in shape and intensely red in color, whereas normal myofibrils are green. Similarly, reflective of compensatory mitochondrial proliferation, specifically in MELAS, was that the same fibers appear "ragged-blue" with the SDH stain and several SSVs, which were not observed in CPEO.

On the other hand, oxidative enzymes were specific markers of respiratory chain complex. COX, the complex IV of the respiratory chain, was only present in mitochondria [12]. Cytochrome

Table 1 Clinical characteristics of MELAS and CPEO patient groups

Patients $(N = 11)$	Age of onset/sex	Clinical manifestation	MDC score	Diagnostic phenotype	mtDNA mutation
1	29/F	Short stature, mildly mentally retarded, headache, cognitive regression, developmental delay, dysphagia	8	MELAS	mt.3243 A>G
2	26/M	Moderate mental retardation, developmental delay, migraine, generalized seizure, blurred vision, family history (diabetes mellitus)	10	MELAS	mt.3243 A>G
3	20/M	Shortness of breath after mild exercise, mild proximal limb weakness, predominantly in the legs, migraine, family history (diabetes mellitus)	7	MELAS	mt.3243 A>G
4	19/F	Muscle weakness, abnormal EMG, seizure, myoclonus, blurred vision, family history (diabetes mellitus)	11	MELAS	mt.3243 A>G
5	30/M	Facies myopathy, migraine, blurred vision, exercise intolerance, and myalgia during walking, myoclonus, blurred speech	10	MELAS	mt.3243 A>G
6	28/M	Short stature, slight headache, abnormal behavior, mentally retarded, developmental delay	9	MELAS	mt.3243 A>G
7	25/F	Progressive heart complaint, cardiomyopathy, ophthalmoplegia, ventricular conduction abnormalities	6	CPEO	mtDNA depletion
8	15/M	Progressive ptosis, blurred vision for 2 years, 1 year ago slurred speech, 2 months ago involuntary movement in hands, gait difficulties, slight intension tremor, limited eye movement, abnormal EMG, died suddenly at 17 years old	9	CPEO/KSS	mtDNA depletion
9	30/F	Progressive ptosis for 10 years, diplopia for 5 years, limited eye motion for 4 years, proximal weakness, abnormal EMG, family history (ophthalmoplegia)	11	CPEO	mtDNA depletion
10	18/M	Progressive ptosis for 3 years, ophthalmoplegia, limited eye motion, vision problem, muscle weakness, exercise intolerance, abnormal EMG	8	CPEO	mtDNA depletion
11	34/F	Progressive ptosis for 8 years, ophthalmoplegia, limited eye movement, abnormal EMG, hearing problem, blurred vision	10	CPEO	mtDNA depletion

F, female; M, male; MELAS, mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes; CPEO, chronic progressive external ophthalmoplegia; KSS, Kearns–Sayre syndrome.



c oxidase activity, therefore, was regarded as an important indicator to demonstrate respiratory chain function. In CPEO with large mtDNA depletion, the damaged RRFs were often deficient for COX histochemical staining, indicative of severely impaired respiratory chain. However, in MELAS associated with mtDNA 3243 A>G mutation, the RRFs and SSVs were typically COX positive and the percentage of COX⁺ fibers was significantly higher than that in CPEO (P < 0.05) as shown in Figure 1B. These observations indicate that MELAS patients may have more residual respiratory chain activity relative to CPEO patients, although both MELAS and CPEO had similar mitochondrial proliferation status.

³¹P-MRS of Quadriceps Femoris Muscle

Before MRS studies, maximal voluntary contraction (MVC) load was estimated as previously described in the healthy volunteers and patients groups. To maintain consistent exercise intensity, 25% MVC was applied to each person. As illustrated in Figure 2A, the MVC of the plantar flexion exercise in patients with MELAS and CPEO was significantly lower than that in healthy volunteers, while there was no significant difference between the MELAS and CPEO groups.

The location of quadriceps on 3.0T MRI is shown in Figure 2B. The quadriceps muscle was chosen because it can produce prominent high-energy phosphate and has high PCr and ATP peaks. A serial ³¹P-MRS procedure at rest, exercise, and recovery phases from a healthy volunteer is shown in Figure 2C. Just after exercise, the Pi peak increased and the PCr peak decreased compared to rest state. During the recovery phase, the Pi gradually decreased and the PCr increased, both of them returned to the normal value of resting phase.

Figure 1 (A). Histopathological features of mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes (MELAS) (patient #1) and chronic progressive external ophthalmoplegia (CPEO) (patient #7) syndrome. (a) HE staining showed scattered vacuolated muscle fibers containing many small basophilic inclusions (RRFs) in both MELAS and CPEO. (b) Gomori staining demonstrated clearly RRFs as well. (c) In MELAS, the SDH staining exhibited intense activity in the subsarcolemmal zones corresponding to RRFs and blood vessel walls (SSVs). In CPEO, there was a complete absence of SSVs and less "ragged-blue" fibers with the SDH stain. (d) Cytochrome c oxidase (COX) staining exhibited increased staining in MELAS, but significantly decreased staining in CPEO. Frozen sections (10 μ m thickness). Scale bar = 50 μ m. (**B**). Percentage of COX+ and COXfibers in RRF and SDH in biopsy of patients with MELAS and CPEO. In MELAS, RRFs and SSVs were typically COX positive, while in CPEO, RRFs were strikingly COX negative. $^{\#}P < 0.05$ compared to CPEO group.

Similar ³¹P-MRS phase information from MELAS and CPEO patients is shown in Figure 2D, including:

- 1. At rest, compared to healthy controls (0.144 \pm 0.021), both MELAS (0.185 \pm 0.023) and CPEO (0.176 \pm 0.027) patients showed an elevated Pi/PCr ratio (*P* < 0.05) due to lower oxidative capacity, while there was no significant difference (*P* > 0.05) in Pi/PCr ratio between the MELAS and CPEO groups.
- 2. Just after exercise (25% MVC), all subjects displayed an elevated Pi/PCr ratio. MELAS patients (0.435 \pm 0.033) acquired an even higher Pi/PCr ratio, significantly different from controls (0.307 \pm 0.023) and CPEO patients (0.388 \pm 0.039) (*P* < 0.05), including MELAS may have more increased nonoxidative glycolytic activity and more impairment in mitochondrial ATP production than CPEO.
- 3. During recovery, ³¹P-MRS is regarded as the most sensitive and specific method to assess skeletal muscle mitochondrial rate of ATP production [13]. In this study, Pi/PCr ratio began to decrease in all subjects from the end of exercise to recovery. This ratio dropped to nearly the resting phase value in healthy controls (0.144 \pm 0.026), while it returned slowly in both MELAS (0.283 \pm 0.029) and CPEO (0.224 \pm 0.016) patients. Mitochondrial encephalomyopathy, lactic acidosis, and stroke-like episodes patients showed the slowest decline (*P* < 0.05), suggestive of a more prolonged rate of Pi/PCr recovery than CPEO. All these observations indicate a more impaired high-energy phosphate transfer in proximal muscles and an even worse mitochondrial function in MELAS.

In healthy controls, ATP/PCr ratio had no significant changes during the rest-exercise-recovery protocol (P > 0.05), because the ATP could be a continuous resynthesis by mitochondrial aerobic





metabolism. However, for both MELAS and CPEO patients, the ATP/PCr ratio declined rapidly after 25% MVC exercise and recovered slowly compared to the control group (P < 0.05). In MELAS, this ratio decreased more dramatically than CPEO during exercise (P < 0.05) and did not return to mean value of resting phase at 5-min recovery, indicative of a worse energy metabolism and less supplement of ATP.

Discussion

A characteristic of MELAS point mutation (mtDNA 3243 A>G) is the occurrence of mitochondrial proliferation in fibers with increased COX activity. The presence of massive mitochondrial proliferation (revealed by RRFs and SSVs) may allow the total COX activity to surpass that seen in unaffected individual fibers. It differs from the CPEO with large deletion mutation (4977 bp mtDNA deletion) where mitochondrial proliferation is commonly seen in COX-negative fibers [14]. Therefore, analysis of the two different genetic subtypes of mitochondrial encephalopathy allowed investigation of the influence of abnormal COX activity in MELAS pathogenesis.

In general, COX histochemistry has proven extremely useful for the diagnosis of mitochondrial diseases [15]. Previous studies showed that COX staining may be decreased, normal, or increased in ragged-red fibers from MELAS patients; however, in CPEO patients, COX activity was significantly absent. This study indicated high expression of COX in six MELAS patients. So compared to CPEO, COX activity was relatively increased, which indicated preserved respiratory chain function in MELAS.

There is a lot of difference between MELAS and CPEO about energy metabolism, because they had different mutation types (generally 3243A>G point mutation in MELAS, mtDNA deletion in CPEO). In vivo ³¹P-MRS has been proposed as a potential tool to evaluate high-energy phosphate compounds, and it has been successfully applied to study mitochondrial dysfunction. It is more sensitive for assessing mitochondrial function than clinical manifestations [16]. Until now, there has been very little research utilizing ³¹P-MRS to explore the energy status between the MELAS and CPEO. This study found that MELAS patients had a higher Pi/PCr ratio and a lower ATP/PCr ratio compared to CPEO, suggesting less direct energy source, delayed energy recovery, and worse energy metabolism [16,17]. These observations were opposed to more resident COX expression in MELAS that represented less respiratory defect. In other words, when COX activity increases beyond normal levels, it may become harmful to the MELAS patient. Therefore, at least in patients evaluated in the current study, abnormally increasing COX expression appears to be an important cause in the pathogenesis of MELAS.

The inherent contradiction in MELAS between the presence of residual enzyme activity (relative to CPEO) and its more severe clinical phenotype, which also called "MELAS paradox," has been reported previously [2]. ³¹P-MRS in the current study demonstrates that MELAS patients with resident COX activity display a

worse energy status compared with CPEO, which further proves the "MELAS paradox" phenomenon. The mechanisms, however, are not clear. An explanation for this is that abnormally increasing COX may lead to dysfunction of NO catabolism. NO plays an integral role in controlling smooth muscle tone and mediating vasodilation and cerebral perfusion. In MELAS, excessive COX binding to NO might cause a relative shortage of circulating NO, which induces endothelial dysfunction and vasodilation retardation, while in CPEO with deficient COX activity, vasodilation will occur normally. Therefore, increased residual COX expression may link to severe skeletal energy failure in MELAS disorders.

Currently, because the clinical manifestations are variable and nonspecific, definite diagnosis of mitochondrial encephalomyopathy still depends on muscle biopsy or gene detection [18]. However, genetic classifications are inevitably incomplete because the responsible mutation is identified in only 20–25% of pediatric patients using routine diagnostic tests [19]. ³¹P-MRS data can be more helpful to elucidate the complex relationship between genotype and phenotype [20]. In addition, ³¹P-MRS is capable of providing quantitative markers to assess the metabolic status of tissues and monitor progression of the disease or treatment response *in vivo* [21]. For example, ³¹P-MRS has proved particularly useful in the therapeutic follow-up of coenzyme Q treatment of mitochondrial disorders [22]. As ³¹P-MRS is entirely noninvasive, it is apparently superior to muscle biopsy or gene detection [18,23]. Taken together, findings of this strongly suggest that: (1) MELAS exhibited increasing COX expression that seemed to have a preservative respiratory function, but had higher Pi/PCr during exercise and delayed ATP/PCr recovery during recovery compared to CPEO, indicating even worse energy metabolism and mito-chondrial dysfunction, (2) this conflict may be partially explained by excessive COX, suggesting that COX might lose the normal function and involved in MELAS pathology, and (3) noninvasive ³¹P-MRS is a promising tool for detecting energy metabolism inefficiency and mitochondrial function impairment in skeletal muscle of mitochondrial encephalomyopathy.

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Disclosures

None.

Conflict of Interest

We declare no conflict of interest.

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Supporting Information

The following supplementary material is available for this article:

Table S1. Mitochondrial disease criteria (MDC, simplified version for beside use)*, 2006.