RESEARCH REPORT

Development of a Tandem Mass Spectrometry Method for Rapid Measurement of Medium- and Very-Long-Chain Acyl-CoA Dehydrogenase Activity in Fibroblasts

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Abstract Mitochondrial fatty acid oxidation is a vital biochemical process for energy metabolism. Among the known fatty-acid metabolism disorders, very-long-chain acyl-CoA dehydrogenase (VLCAD) deficiency and medium-chain acyl-CoA dehydrogenase (MCAD) deficiency count among the most frequent. Both are potentially very serious diseases as they carry a risk of severe neurological post-crisis sequelae, and even sudden death. Diagnosis relies on plasma acylcarnitine profile analysis and urine organic acid analysis, followed by genetic testing to confirm diagnosis. However, in some cases, it is crucial to run a specific diagnostic assay for enzyme activity, which is generally performed in leukocytes or fibroblasts. The aim of this study was to address this need, first by developing a MCAD and VLCAD enzyme activity-specific diagnostic assay in fibroblasts (by measuring the reaction products, i.e. enoyl-CoA) via a rapid LC–MS/MS-based technique, and

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then by testing MCAD-deficient patients $(n = 6)$, VLCADdeficient patients ($n = 10$), and control patients ($n = 12$). MCAD activity was significantly different in the MCADdeficiency (MCADD) group (mean $= 0.07$ nmol C8:1 formed/min/mg protein) compared to the control group (mean $= 0.36$ nmol C8:1 formed/min/mg protein). All MCADD patients showed less than 35% residual MCAD activity. VLCAD activity was significantly decreased in the VLCADD group (mean $= 0.06$ nmol C16:1 formed/min/ mg protein) compared to the control group (mean $= 0.86$ nmol C16:1 formed/min/mg protein, respectively). All VLCADD patients showed less than 35% residual VLCAD activity. This technique allowed also to confirm that a novel $ACADVL$ gene mutation (c.1400T>C) is responsible for a defective VLCAD activity (residual activity at 10%).

Abbreviations

Introduction

Mitochondrial fatty acid (FA) oxidation is essential for energy production (Houten and Wanders [2010\)](#page-6-0), and a

deficiency in the process leads to inability to utilize FA, often resulting in fasting hypoketotic hypoglycemia. Of the 24 known fatty-acid metabolism disorders, autosomalrecessive deficiencies in very-long-chain acyl-CoA dehydrogenase (VLCAD, EC1.3.8.9) (OMIM 609575) and medium-chain acyl-CoA dehydrogenase (MCAD, EC1.3.8.7) (OMIM 201450) count among the more frequent. Both enzymes catalyze the first step of the mitochondrial FA oxidation cycle leading to dehydrogenation of acyl-CoA to enoyl-CoA (Houten and Wanders [2010](#page-6-0); Wanders et al. [1999\)](#page-7-0). This step requires flavin adenine dinucleotide (FAD) bound to Electron Transfer Flavoprotein (ETF), which is then reduced into FADH2. VLCAD is bound to the inner mitochondrial membrane whereas MCAD is a matrix enzyme (Houten and Wanders [2010](#page-6-0); Wanders et al. [1999](#page-7-0)). MCAD acts on medium-chain FA (4–14 carbon atoms, optimum activity with 6-carbon atoms). VLCAD acts on long-chain FA (12–22 carbon atoms, optimum activity with 16-carbon atoms) (Hashimoto [1992\)](#page-6-0). Clinical presentation of VLCAD deficiency (VLCADD) may include cardiomyopathy, encephalopathy, hypoglycemia, and rhabdomyolysis (Arnold et al. [2009](#page-6-0)). MCAD deficiency (MCADD) is associated with hypoketotic hypoglycemia, hepatic symptoms that are predominantly related to intercurrent illnesses or prolonged fasting. Sudden death related to heartbeat disorders may also occur in adults (Feillet et al. [2012\)](#page-6-0).

For VLCADD and MCADD, diagnosis is first suspected by plasma acylcarnitine and urine organic acid analyses, and has to be confirmed by molecular biology (Feillet et al. [2012;](#page-6-0) Vianey-Saban et al. [1998\)](#page-7-0).

MCADD management specifies that enzyme activity must confirm diagnosis if molecular biology is not contributory. In addition, as a post mortem plasma acylcarnitine profile is very difficult to interpret, an enzyme activity assay can prove important in the exploration of sudden death (Feillet et al. [2012](#page-6-0)).

Strategy is similar for VLCADD where an enzyme activity assay may be even more valuable given the heterogeneity of *ACADVL* gene mutations (Vianey-Saban et al. [1998](#page-7-0)).

A method that measures the production of octenoyl-CoA (C8:1-CoA) (after addition of octanoyl-CoA as substrate) or hexadecenoyl-CoA (C16:1-CoA) (after addition of palmitoyl-CoA as substrate) by tandem mass spectrometry in lymphocytes was published (Tajima et al. [2008](#page-7-0); ter Veld et al. [2009](#page-7-0)). The method uses ferrocenium hexafluorophosphate, an artificial electron acceptor for MCAD or VLCAD, to specifically determine MCAD and VLCAD activity within a day, but has never been further developed for analysis in fibroblasts.

Methods

Patients

Skin fibroblasts (75 cm² flask per strain, i.e. around 6.10^6 cells) grown in Ham's culture medium added with 12% fetal calf serum are harvested at confluence (Centre de Biotechnologie Cellulaire, Hospices Civils de Lyon, Bron). After trypsinization, washing in phosphate-buffered saline (PBS) and aspirating the supernatant, dry-harvested fibroblast pellets (12 controls, 10 known VLCAD-deficient patients, 6 known MCAD-deficient patients) were frozen at -80° C for storage.

The pellets were then resuspended with $300 \mu L$ of ammonium acetate (10 mM, $pH = 8$) and sonicated, in ice, for three \times 15 sec bursts at 30 sec intervals.

Tests were also carried out in control-sample fibroblasts to evaluate analytical performance.

Reagents

Ammonium acetate and 2-(N-morpholino)ethanesulfonic acid (MES) were purchased from Merck (Darmstadt, Germany). Acetonitrile, octanoyl-CoA (C8:0), hexadecenoyl-CoA (C16:0), and ferrocenium hexafluorophosphate were obtained from Sigma-Aldrich (Deisenhofen, Germany).

Protein Assay

The total protein assay in sonicated fibroblast homogenates was performed in a 1:20 dilution on an ABX Pentra400[®] benchtop analyzer (Horiba Medical) using a colorimetric technique based on the bicinchoninic acid method (Thermo Scientific, Rockford, USA).

Procedures

Sonicated fibroblast homogenates, adjusted to a protein concentration of 0.2 g/L with ammonium acetate 10 mM, $pH = 8$, were diluted (1:1 V/V) in ferrocenium hexafluorophosphate solution (400 μ M, pH = 8) to a final volume of 100 µL. The enzymatic reaction was started at 37° C by adding the substrate in MES buffer (20 mM, $pH = 6$): 4 µL of C8:0 (5 mM) to assay MCAD activity; 4 μ L of C16:0 (5 mM) to assay VLCAD activity. After 15 min incubation, the reaction was stopped by adding 100 mL of ice-cold acetonitrile. A T0 timepoint measure was taken for each condition by simultaneously adding substrate and acetonitrile. After centrifugation at 14,000 g for 8 min at 4° C, the supernatant was transferred to autosampler vials with inserts for HPLC–mass spectrometry.

The analyte samples were injected (20 μ L for the MCAD assay, $5 \mu L$ for the VLCAD assay) and chromatographic separation was carried out using a C18 Uptisphere column $(50 \times 2.1$ mm, particle size: 3 um) (Interchim, Montlucon, France). Flowrate was 200 µL/min. Eluent was a 70:30 (V/V) mixture (for MCAD assay) or 45:55 (V/V) mixture (for VLCAD assay) of ammonium acetate (10 mM, $pH = 8$) and acetonitrile. After chromatographic separation, the samples were analyzed by electrospray ionization–tandem mass spectrometry (ESI-MS/MS, Api 3200, Sciex, Les Ulis, France) in positive ion mode. The products derived directly from the MCAD-catalyzed reaction, i.e. octenoyl-CoA (C8:1), and the VLCAD-catalyzed reaction, i.e. hexadecenoyl-CoA (C16:1), were quantified. The products of the second step in the FA oxidation process were also assayed, i.e. 3-hydroxyoctanoyl-CoA (C8OH) and 3 hydroxypalmitoyl-CoA (C16OH), respectively. The product ions were studied in multiple reaction monitoring (MRM) mode, with each transition corresponding to substrate (C8:0 m/z 894.2>387.3, and C16:0 m/z 1006.4>499.4) or to products of the enzymatic reaction (C8:1 m/z 892.2>385.3, and C8OH m/z 910.2>403.3 or C16:1 m/z 1004.4>497.4 and C16OH m/z 1022.4>515.4). Working up from the peak areas obtained, the quantities of reaction products were calculated from substrate area at timepoint T0. As C8:1, C16:1, C8OH, and C16OH are not commercially available, response of the substrate and response of the enzymatic reaction products were considered identical.

Expression of Results

Enzymatic activities are expressed in nmol C8:1 (MCAD) or C16:1 (VLCAD) formed per minute per mg of proteins. For patients with MCADD or VLCADD, residual activity was evaluated as a ratio of the activity level of the control enzyme, i.e. VLCAD for MCAD-deficient patients and MCAD for VLCAD-deficient patients. The MCAD-to-VLCAD or VLCAD-to-MCAD ratios are then normalized by the mean of the ratios of the control group tested in the same experiment, thus giving (MCAD-to-VLCAD)-tocontrol and (VLCAD-to-MCAD)-to-control ratios.

Statistical Analysis

Group-wise results (control, MCADD, VLCADD) for MCAD or VLCAD enzymatic activity are reported as means (SD; $CI_{95%}$). Between-group comparisons (control, MCADD, VLCADD) of MCAD (or VLCAD) activity were performed by one-way ANOVA, followed by a Tukey's post hoc test for pairwise comparisons between significantly different groups. All statistical analyses were performed using GraphPad® Prism 5 software. Statistical significance was set at $p < 0.05$.

Results

Optimization of the Method

The final selected analytical parameters were set as follows: protein concentration $= 0.1$ g/L, incubation time $= 15$ min, ferrocenium hexafluorophosphate concentration $= 200 \mu M$, final substrate (C8 or C16) concentration $= 0.2$ mmol/L.

Intra-assay coefficient of variation (CV) assessed via 10 repeated measures on the same control cell line was 4.3% (mean 0.47 nmol/min/mg protein) for MCAD activity and 10.8% (mean 0.74 nmol/min/mg protein) for VLCAD activity. Inter-assay CV assessed via 5 measurements carried out over 5 days on the same control cell line was 19% (mean 0.41 nmol/min/mg protein) for MCAD activity and 31% (mean 0.92 nmol/min/mg protein) for VLCAD activity.

Figure [1](#page-3-0) illustrates chromatograms obtained in MRM mode for a patient with MCADD (Fig. [1a](#page-3-0)) or VLCADD (Fig. [1](#page-3-0)b), compared to a control patient with a decrease in signal intensity for the products C8:1 or C16:1 and C8OH or C16OH without concomitant change in signal intensity for substrate C8 or C16 (in surplus).

Population Studied

Three patient groups were studied: a group of 12 control patients, a group of 10 previously diagnosed VLCADdeficient patients $(P1-P10)$, and a group of six previously diagnosed MCAD-deficient patients (P11–P16). For each patient, MCAD and VLCAD activity assays were performed in duplicate within a same series and over two independent experimental runs.

Key clinical data on patients P1–P16 are reported in Table [1.](#page-5-0) All enzyme deficiencies were confirmed by molecular biology except for patient P11 who was only heterozygous for the frequent c.985A>G mutation. Diagnosis was confirmed for this patient by an enzymatic activity assay using ETF as electron acceptor (Bertrand et al. [1992](#page-6-0)). All the identified mutations (Table [1\)](#page-5-0) have been described as deleterious in the literature and/or by predictions software (Alamut[®], Interactive biosoftware).

Enzymatic Activities in the Study Population

Mean MCAD activity in the control group was 0.36 nmol C8:1 formed/min/mg protein (SD: 0.07; $CI_{95%}: 0.31-0.40$). Mean MCAD activity in known VLCAD-deficient patients was 0.30 nmol C8:1 formed/min/mg protein (SD: 0.09; $CI_{95%}: 0.25–0.35$. Mean MCAD activity in known MCAD-deficient patients was 0.07 nmol C8:1 formed/ min/mg protein (SD: 0.08; CI_{95%}: 0.00-0.13), which is significantly different to the control and "VLCAD-deficiency" groups $(p < 0.0001)$ (Fig. [2a](#page-6-0)).

Fig. 1 MRM-mode chromatograms obtained after MCAD (a) and VLCAD (b) enzyme reactions. For each molecule, peaks are shown at

identical scale between control patient and known MCAD-deficient (a) or known VLCAD-deficient (b) patient. (a) Substrate octanoyl-

Mean VLCAD activity in the control group was 0.86 nmol C16:1 formed/min/mg protein (SD: 0.028; $CI_{95%}: 0.68–1.04$. Mean VLCAD activity in known MCAD-deficient patients was 0.73 nmol C16:1 formed/ min/mg protein (SD: 0.49; $CI_{95\%}$: 0.35–1.11). Mean VLCAD activity in known VLCAD-deficient patients was 0.06 nmol C16:1 formed/min/mg protein (SD: 0.07; $CI_{95%}$: 0.02–0.10), which is significantly different to the control and "MCAD-deficiency" groups $(p \lt 0.0001)$ (Fig. [2](#page-6-0)b).

All the known VLCADD and MCADD patients showed less than 35% residual VLCAD or MCAD activity (Table [1\)](#page-5-0).

Discussion

MCADD and VLCADD are autosomal-recessive inborn errors of metabolism. Diagnosis is first suggested by plasma acylcarnitine profile analysis and urine organic acid analysis, and generally has to be confirmed by molecular biology. But sometimes a specific MCAD and VLCAD enzyme activity assay may prove necessary in cases involving non-contributive genetics or cases where patients refuse consent for genetic testing. In this context we developed a tandem mass spectrometry assay of MCAD/VLCAD activity in fibroblasts.

We first optimized the technique originally developed by Tajima et al. [\(2008\)](#page-7-0) and ter Veld et al. ([2009](#page-7-0)) in lymphocytes. Wanders et al. [\(2010\)](#page-7-0) also used the ferrocenium hexafluorophosphate in a HPLC coupled to UV-detection assay of MCAD/VLCAD activity in fibroblasts. Although we ultimately retained the same ferrocenium hexafluorophosphate concentration (200 μ M) and the same amount of substrate (20 nmol) as the original method, we opted for a 15 min incubation time at 37° C (instead of 5 min originally) and a 0.1 g/L protein concentration (instead of 0.03 g/L originally). Analytical performances for VLCAD activity are weaker than for MCAD activity, while the CVs remain acceptable for a tandem mass spectrometry technique. This can be explained by the fact that VLCAD is a membrane bound enzyme and therefore offers more random substrate availability than MCAD which is a matrix-soluble enzyme, despite the sonication step.

We applied this new method to analyze fibroblasts from MCAD-deficient or VLCAD-deficient patients. The activity of the deficient enzyme was significantly lower in the corresponding patient groups. Each individual patient with MCADD or VLCADD showed a significant decrease in the corresponding enzyme activity.

However, three particular cases draw our attention.

In patient P1, who died at day 2 of life with only post mortem organic acid profile evocative of long-chain FAO defect, two mutations were identified in ACADVL gene. Mutation c.1837C>T (p.Arg613Trp) has previously been described in the literature (Souri et al. [1996\)](#page-6-0). The second substitution, c.1400T>C (p.Ile467Thr), has never been described (zero Pubmed, Ensembl, Clinvar, or ExAc Browser hits) and was classified as intermediate-severity by the prediction software. Both c.1837C>T and c.1400T>C mutations were properly segregated in the parents. The enzyme activity assay demonstrates its full utility in this case and enables diagnosis to be confirmed with a mean 10% residual VLCAD activity.

For patient P11, presenting with MCADD of good prognosis, only one heterozygous mutation was identified in ACADM gene. No other mutation was found in exons and flanking intronic regions of the second allele. The MCAD residual activity of 8% confirms plasma acylcarnitine and urine organic acid analyses. This method is easier and shorter to achieve than the previous method using ETF (Bertrand et al. [1992](#page-6-0)).

For patient P13, presenting with MCADD of good prognosis, two mutations were identified in the ACADM gene, both of which have been described in the literature (McKinney et al. [2004,](#page-6-0) Waddell et al. [2006](#page-7-0)). Mutation c.127G>A (p.Glu43Lys) has been described in homozygous state in a patient who also presented a mutation in heterozygous state on the gene coding for SCAD (McKinney et al. [2004\)](#page-6-0). Mutation c.617G>A, which leads to substitution of a positively charged side-chain amino acid (arginine) at position 206 by a same-family amino acid (histidine), has been described in a compound heterozygous c.617G>A/ c.985A>G patient (Waddell et al. [2006\)](#page-7-0). Both mutations are classified as intermediate-severity by prediction software, which is an understandable conclusion given how c.617G>A is a conservative mutation. The enzyme activity assay demonstrates its utility and enables diagnosis to be confirmed with a mean 33% residual MCAD activity compared to controls. French consensus guidelines on diagnostic screening have set the confirmatory cut-off as less than 25% residual activity (Feillet et al. [2012\)](#page-6-0). However, the guidelines were put together based on results obtained with the reference technique using ETF in fluorescence analysis, and it is now time to redefine a new threshold (possibly 35%) with this technique. Furthermore, the clinical picture and type of mutations point towards a moderate form of MCAD deficiency and explain why residual activity was found to be around 33%. Analysis of other patients is needed to define this new threshold.

toyl-CoA (C16) and products hexadecenoyl-CoA (C16:1) and 3 hydroxypalmitoyl-CoA (C16OH) were assayed

^{⁄-} Fig. 1 (continued) CoA (C8) and products octenoyl-CoA (C8:1) and 3-hydroxyoctanoyl-CoA (C8OH) were assayed. (b) Substrate palmi-

Table 1 Main clinical and genetic data, and residual activities of the 6 MCAD-deficient and the 10 VLCAD-deficient patients Table 1 Main clinical and genetic data, and residual activities of the 6 MCAD-deficient and the 10 VLCAD-deficient patients

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min/mg protein) of the deficient enzyme (MCAD or VLCAD) was ratioed to the activity of the control enzyme (VLCAD or MCAD), and this ratio was then normalized by the mean control-group

ratio to give a residual activity in %

No mutation found in exons and flanking intronic regions

Fig. 2 Mean MCAD (a) and VLCAD (b) enzyme activities per patient group: control patients $(n = 12)$, known VLCAD-deficient patients $(n = 10)$ (VLCADD) and known MCAD-deficient patients

Conclusion

The tandem mass spectrometry assay of MCAD/VLCAD activity in fibroblasts developed here offers powerful selective diagnostic screening of MCAD-deficient and VLCAD-deficient patients.

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Take-Home Message

The reader will learn about the usefulness of MCAD and VLCAD activity measurement and the performance of the method developed in fibroblasts.

Conflict of Interest

Damien Bouvier, Christine Vianey-Saban, Séverine Ruet, and Cécile Acquaviva declare that they have no conflict of interest.

Compliance with ethics guidelines: Informed consent was obtained from all patients for being included in the study.

Authors Contribution

Damien Bouvier and Severine Ruet have performed technical part of the study.

 $(n = 6)$ (MCADD). For each patient, enzymatic reaction was performed in duplicate over two independent experiments. $\frac{p}{p}$ < 0.0001

Damien Bouvier, Christine Vianey-Saban, Severine Ruet, and Cécile Acquaviva have performed analysis and interpretation of the data.

Damien Bouvier drafted the article.

Christine Vianey-Saban and Cécile Acquaviva revised the article.

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