

SECONDARY MITOCHONDRIAL DYSFUNCTION IN PROPIONIC ACIDURIA, A PATHOGENIC ROLE FOR ENDOGENOUS MITOCHONDRIAL TOXINS

Marina A. Schwab*, Sven W. Sauer*, Jürgen G. Okun*, Leo G. J. Nijtmans†, Richard J. T. Rodenburg†, Lambert P. van den Heuvel†, Stefan Dröse‡, Ulrich Brandt‡, Georg F. Hoffmann*, Henk ter Laak†, Stefan Kölker*§¹, and Jan A. M. Smeitink†§.

From the *Department of General Pediatrics, Division of Inherited Metabolic Diseases, University Children's Hospital Heidelberg, Im Neuenheimer Feld 150, D-69120 Heidelberg, Germany; the †Department of Pediatrics, Radboud University Nijmegen Medical Center - Nijmegen Center for Mitochondrial Disorders (NCMD), Geert Grooteplein 10, NL-6500 HB Nijmegen, The Netherlands; the ‡Molecular Bioenergetics Group, Gustav-Emden-Zentrum der Biologischen Chemie, Johann Wolfgang Goethe University, Theodor-Stern-Kai 7, D-60590 Frankfurt/Main, Germany.

Short title: Mitochondrial dysfunction in propionic aciduria

Mitochondrial dysfunction during acute metabolic crises is considered an important pathomechanism in inherited disorders of propionate metabolism, i.e. propionic and methylmalonic acidurias. Biochemically, these disorders are characterized by accumulation of propionyl-Coenzyme A (CoA) and metabolites of alternative propionate oxidation. Here we demonstrate uncompetitive inhibition of pyruvate dehydrogenase complex (PDHc) by propionyl-CoA in purified porcine enzyme and in submitochondrial particles from bovine heart being in the same range as the inhibition induced by acetyl-CoA, the physiological product and known inhibitor of PDHc. Evaluation of homologous monocarboxylic CoA esters showed a chain-length specificity for PDHc inhibition. In contrast to CoA esters, free fatty acids did not inhibit PDHc activity. In addition to PDHc inhibition, analysis of respiratory chain and tricarboxylic acid enzymes also revealed an inhibition of propionyl-CoA on respiratory chain complex III and α -ketoglutarate dehydrogenase complex.

To test whether impairment of mitochondrial energy metabolism is involved in the pathogenesis of propionic aciduria, we performed a thorough bioenergetic analysis in muscle biopsy specimens of two patients. In line with the *in vitro* results, oxidative phosphorylation was severely compromised in both patients. Furthermore, expression of respiratory chain complexes I-IV, and the amount of mitochondrial DNA were strongly decreased, and ultrastructural mitochondrial abnormalities were found, highlighting severe mitochondrial dysfunction. In conclusion, our results favour the hypothesis that toxic metabolites, in particular propionyl-CoA, are involved in the pathogenesis of inherited disorders of propionate metabolism, sharing mechanistic similarities with propionate toxicity in microorganisms.

Key words: propionic aciduria, propionyl-coenzyme A, mitochondrial dysfunction, mitochondrial DNA, oxidative phosphorylation, pyruvate dehydrogenase complex

INTRODUCTION

Propionyl-CoA is an intermediate metabolite in the final common catabolic pathways of the amino acids L-isoleucine, L-methionine, L-threonine and L-valine as well as odd-chain fatty acids and cholesterol. It is carboxylated by propionyl-CoA carboxylase (EC 6.4.1.3) to methylmalonyl-CoA and then is converted by methylmalonyl-CoA mutase (EC 5.4.99.2) to succinyl-CoA subsequently being fed as anaplerotic precursor into the tricarboxylic acid (TCA) cycle. Inherited deficiency of propionyl-CoA carboxylase, i.e. propionic aciduria (PA), is biochemically characterized by an accumulation of propionate, 3-hydroxypropionate, 2-methylcitrate, and propionylglycine, demonstrating alternative propionate oxidation in these patients. Furthermore, metabolic acidosis, ketosis, increased lactate concentrations, hypoglycemia, and hyperammonemia are found during metabolic derangement, highlighting secondarily compromised energy metabolism and ammonia detoxification [1]. Clinically, PA is complicated by acute life-threatening metabolic crises which are precipitated by catabolic state and result in multiple organ failure or even death if untreated [1]. Like other organic acidurias (e.g. methylmalonic aciduria), age at clinical onset and disease course may vary between individual patients presenting with neonatal metabolic encephalopathy (neonatal onset), recurrent episodes of ketoacidotic coma or Reye-like syndromes (chronic intermittent form), or psychomotor retardation and failure to thrive in the absence of acute crises (chronic progressive form). Similarly to neonatal onset of PA patients, *PCCA*^{-/-} mice, a transgenic mouse model for PA, develop lethal ketoacidosis shortly after birth [2].

Despite improvements in the diagnostic work-up and management of affected patients during the last decades, the neurologic outcome remains disappointing in PA patients [1,3]. In particular mental retardation and movement disorders secondary to cortical atrophy, leukoencephalopathy and basal ganglia injury are still frequently found even in early diagnosed children. The biochemical abnormalities found during acute metabolic crises have led to the suggestion that impairment of oxidative phosphorylation (OXPHOS) is crucial for the pathogenesis of this disease. Notably, accumulating metabolites of alternative propionate oxidation have been suggested to act as endogenous inhibitors of energy metabolism in different *in vitro* models [4-7], however, the pathophysiological impact of these findings on PA still remains unclear.

Here we report severe disturbance of mitochondrial energy metabolism in muscle tissues from two PA patients and demonstrate *in vitro* that propionyl-CoA-induced mitochondrial dysfunction plays a central role in this scenario.

EXPERIMENTAL

Patient 1

This girl was born at term as the second child of non-consanguineous Caucasian parents. The third day of life she was admitted because of progressive feeding refusal, lethargy and abnormal breathing. Laboratory investigations revealed a severe metabolic acidosis (pH 7.01, pCO₂ 18 mm Hg, bicarbonate 5 mM, base excess -25 mM) and hyperammonemia (830 μ M; normal <80 μ M). The amino acids analysis in plasma revealed no abnormalities. Analysis of urine organic acids by gas chromatography/mass spectrometry showed biochemical abnormalities characteristic for PA with increased concentrations of beta-hydroxybutyric acid (6373 mmol/mol creatinine; control range: 0-45 mmol/mol creatinine), beta-hydroxypropionic acid (21461 mmol/mol creatinine; control range: 0-160 mmol/mol creatinine), beta-hydroxyisovaleric acid (176 mmol/mol creatinine; control range: 0-10 mmol/mol creatinine), lactic acid (776 mmol/mol creatinine; control range: 0-270 mmol/mol creatinine), propionylglycine, methylcitrate and tiglylglycine. The last three metabolites were not quantified. The concentration of free carnitine in plasma was decreased (10 μ M; control range 20-200 μ M), and the concentration of total carnitine was low-normal. A carnitine profiling was not done at the time of admission, but later on it repeatedly showed increased concentrations of propionylcarnitine varying between 40-80 μ M (normal < 4 μ M).

Extracorporeal detoxification was performed. Besides a high caloric intake and transient stop of protein intake, metronidazol, lactulose, L-carnitine and biotin were added in the acute treatment phase. At age 10 days she was dismissed from the neonatal intensive care unit and then was nursed at the metabolic ward. The only abnormal clinical sign at that time was a slight axial hypotonia. Maintenance treatment consisted of a natural protein intake (1.0 g/kg per day) supplemented with an amino acid mixture lacking precursor amino acids of propionyl-CoA (i.e., L-isoleucine, L-methionine, L-threonine, and L-valine), and application of L-carnitine and biotin. Analysis of propionyl-CoA carboxylase activity in leukocytes showed deficient activity of this enzyme (0.1 nmol/h/mg protein; control: 2.3 nmol/h/mg protein) and thus confirmed the diagnosis of PA. She was admitted to hospital numerous times due to metabolic derangements; the last one, at age five years, being fatal. The child had a severe myopathic appearance during life, which was, in combination with repeatedly increased blood lactate concentrations (2.0-7.2 mM; normal <2.1 mM), a reason to perform a muscle biopsy for mitochondrial studies.

Patient 2

This boy was born at term following an uncomplicated pregnancy and delivery. At the third day of life tachypnoea and metabolic acidosis led to a detailed metabolic investigation. The investigation of amino acids in plasma revealed an increased concentration of glycine (722 μ M; 80-440 μ M). As for patient 1 urine organic acid analysis showed an increased concentration of beta-hydroxybutyric acid (17818 mmol/mol creatinine), beta-hydroxypropionic acid (25096 mmol/mol creatinine), beta-hydroxyisovaleric

acid (34 mmol/mol creatinine), lactic acid (1322 mmol/mol creatinine), propionylglycine, methylcitrate and tiglylglycine. Propionyl-CoA carboxylase activity in cultured skin fibroblasts was strongly decreased (0.3 nmol/h/mg protein), confirming the diagnosis of PA that was treated according to standard protocols (see also patient 1).

The child always had a myopathic appearance and exercise intolerance. Standard exercise testing at age five years showed an increase of blood lactate from normal concentrations at rest to 20 mM lactate following a short run of about 50 meters. This was the reason for further bioenergetic analyses in muscle tissue (Table 1).

Muscle and Skin Biopsies

Muscle biopsies (Musculus vastus lateralis dextra) and forearm skin punch biopsies were performed in both patients. Furthermore, fibroblast cultures from patients with confirmed diagnosis of PDHc deficiency ($n=3$) and healthy volunteers ($n=12$) were used. Biopsies and subsequent experiments were performed after receipt of informed consent. The study was approved by the Institutional Review Board. Further investigations were performed in muscle tissue (Musculus quadriceps femoris) of wild-type C57Bl/6 mice. Animal care followed the official governmental guidelines and was approved by the government ethics committee.

Human Skin Fibroblast Cultures

Human skin fibroblasts from forearm skin biopsies were cultivated at standard conditions in DMEM at 37 °C supplemented with 10 % fetal calf serum (v/v), 100 μ g/ml of penicillin, 100 μ g/ml of streptomycin, 2.5 μ g/ml of fungizone and 200 μ M of uridine until confluency [8]. Medium was changed twice a week. Each cell culture was tested for contamination with *Mycoplasma spp.* before enzyme analysis.

Preparation of Tissue Extracts

Fibroblast and muscle homogenates as well as submitochondrial particles from bovine heart (SMP) were prepared as previously described [8-10].

PDHc Activity

Spectrophotometric analysis of PDHc activity (E1, EC 1.2.4.1; E2, EC 2.3.1.12; E3, EC 1.8.1.4) was performed in purified porcine PDHc (Sigma-Aldrich, Schnelldorf, Germany), in SMP, and in homogenates from human skin fibroblasts and quadriceps muscle biopsy specimens using a *p*-iodonitrotetrazolium violet coupled system as previously described [8]. In brief, steady state activities of PDHc were recorded using a computer tunable spectrophotometer (SPECTRAMax Plus 384 Microplate Reader, Molecular Devices, Sunnyvale, CA) operating in the dual wavelength mode. Samples were

analyzed in temperature-controlled 96-well plates in a final volume of 300 μ l. PDHc (purified enzyme, 160 milliunits/ml; SMP, 1.2 mg of protein/ml; fibroblast homogenate, 4 mg of protein/ml) was assayed in a buffer containing 0.05 M of potassium phosphate, 2.5 mM of NAD, 5 mM of L-carnitine, 0.2 mM of thiamine pyrophosphate, 0.1 mM of CoA, 0.1% Triton X-100 (v/v), 1 mM of MgCl₂, 1 mg/ml of bovine serum albumin, 0.6 mM of *p*-iodonitrotetrazolium violet, and 6.5 μ M of phenazine methosulfate, which was adjusted to pH 7.5 at 25 °C. PDHc activity was determined as *p*-iodonitrotetrazolium violet reduction at a wavelength of λ =500-750 nm. The specificity of this assay was confirmed by complete inhibition of PDHc activity by the specific E1 inhibitor 3-fluoropyruvate (5 mM).

To investigate the effect of acetyl-CoA and propionyl-CoA, we varied the concentrations of these CoA esters (0-1 mM) and pyruvate (0-0.1 mM). Additional experiments were performed using short-chain (acetyl-CoA [C₂], propionyl-CoA [C₃], butyryl-CoA [C₄]), medium-chain (hexanoyl-CoA [C₆], octanoyl-CoA [C₈], decanoyl-CoA [C₁₀]) and long-chain (myristoyl-CoA [C₁₄], palmitoyl-CoA [C₁₆]) acyl-CoA esters as well as corresponding short- and medium-chain fatty acids (up to 1 mM; all adjusted to pH 7.5).

Spectrophotometric Analysis of α -Ketoglutarate Dehydrogenase Complex (KGDHc) Activity

Spectrophotometric analyses of KGDHc (E1k, EC 1.2.4.2; E2k, EC 2.3.1.61; E3, EC 1.8.1.4) was performed in SMP and muscle homogenates as previously described [10]. In brief, KGDHc (650 milliunits/ml) was assayed in a buffer containing 35 mM of potassium phosphate, 5 mM of MgCl₂, 0.5 mM of EDTA, 0.5 mM of NAD, 0.2 mM of thiamine pyrophosphate, 0.04 mM of CoA-SH, and 2 mM of α -ketoglutarate, which was adjusted to pH 7.4 at 30 °C. KGDHc activity was determined as NAD reduction at a wavelength of λ =340-400 nm. The effect of propionyl-CoA on KGDHc activity was tested in analogy to PDHc (n=8).

Spectrophotometric Analysis of OXPHOS Complexes I-V

The catalytic activities of respiratory chain complexes I-V in SMP and muscle homogenates were investigated as previously described [9-14]. The addition of standard respiratory chain inhibitors (complex I, 2-*n*-decylquinazolin-4-yl-amine [1 μ M]; complex II, thenoyltrifluoroacetone [8 mM]; complex III, antimycin A [1 μ M]; complex IV, NaCN [2 mM]; complex V, oligomycin [80 μ M]) revealed good inhibitory responses (93-100% of control activity, $p < 0.001$ versus controls), confirming a high specific enzyme activity in our assay system. Acyl-CoA esters (C₂, C₃, C₄, C₆) and corresponding fatty acids (all adjusted to pH 7.4) were added and effects on single respiratory chain complexes I-V were determined at concentrations up to 1 mM each (complex I, 0.5 mg of protein/ml; complex II, 0.3 mg/ml of protein; complex III, 0.1 mg/ml of protein, complex IV, 0.01 mg/ml of protein; complex V, 1.1 mg/ml of protein). In analogy, complex III activity was also investigated in isolated complex III from bovine heart (0.1 mg/ml of protein).

Radiometric Analysis of Mitochondrial OXPHOS

Radiometric analysis was performed in freshly prepared quadriceps muscle biopsy specimens from PA patients. 600×g supernatants were prepared and oxidation rates of [1-¹⁴C]pyruvate, [U-¹⁴C]malate and [1,4-¹⁴C]succinate were determined as previously described [14-16].

ATP Production

Spectrophotometric analysis of ATP production was determined in muscle biopsies of PA patients using unlabeled pyruvate, malate, and succinate as described previously [14]. In brief, spectrophotometric analysis of ATP production is coupled to the formation of NADPH ($\lambda=320-400$ nm, 25 °C). The test principle consists of two enzyme reactions. In the first step, glucose (30.3 mM) and ATP are catalysed to glucose 6-phosphate and ADP by hexokinase (EC 2.7.1.1). Subsequently, glucose 6-phosphate dehydrogenase (EC 1.1.1.49) catalyses glucose 6-phosphate and NADP⁺ to 6-phosphogluconolacton and NADPH.

Protein Electrophoresis of Muscle Tissue

One and two dimensional blue native polyacrylamide-gelelectrophoresis (BN-PAGE) and subsequent in-gel activity assays and Western blotting were performed in muscle biopsies of PA patients as described previously [17]. Antibodies against OXPHOS complex I (subunits 39 kDa and ND6), complex II (subunit 70 kDa), complex III (subunit Core 2) were obtained from Molecular Probes (Eugene, Oregon, USA), and against cyclophilin b from Affinity BioReagents (Golden, Colorado, USA).

Electron Microscopy of Muscle Tissue

Electron microscopy in muscle biopsies of PA patients was done according to standard protocols.

Data Analysis

Enzyme activities were normalized to citrate synthase (EC 2.3.3.1) activity and protein concentration of the same sample [18,19]. Data were expressed as mean±SD of at least three independent experiments. One-way analysis of variance (ANOVA) followed by *posthoc* Bonferroni's multiple comparison test (for three or more groups), or Student's *t* test (for two groups) were used to calculate statistical differences between groups. Results are presented as the mean±SD if not indicated differently. Statistics were calculated using SPSS for Windows 12.0 software. $p<0.05$ was considered significant.

RESULTS

Propionyl-CoA Inhibits PDHc

In purified porcine PDHc, propionyl-CoA showed an inhibition of PDHc activity being uncompetitive with respect to pyruvate (Figure 1). Notably, this inhibitory effect was in the same range as for acetyl-CoA, the physiologic product and known inhibitor of PDHc (Figure 2). For propionyl-CoA (0.125-1 mM) the inhibition constant (K_{is}) ranged from 101-422 μ M (3-12 times K_m), and for acetyl-CoA (0.125-1 mM) from 90-256 μ M (2-7 times K_m). The K_m was $35 \pm 0.5 \mu$ M for pyruvate. Similar results were found in SMP from bovine heart (data not shown).

➤ **Figure 1 and 2**

Next, we determined PDHc activity in fibroblast homogenates from two PA patients. Assay conditions allowed enzyme analysis in the absence of intracellularly formed propionyl-CoA. Notably, fibroblasts from both patients revealed PDHc activities within the control range confirming our hypothesis of secondary PDHc inhibition induced by propionyl-CoA (Figure 3). As expected, patients with confirmed PDHc deficiency showed a reduced PDHc activity in fibroblasts (Figure 3).

➤ **Figure 3**

Acyl-CoA Esters Inhibit PDHc in a Chain-length Specific Manner

To investigate whether PDHc inhibition was dependent on the chain length of acyl-CoA esters, we investigated the inhibitory effect of homologous monocarboxylic CoA esters ranging from C₄ to C₁₆ in purified porcine PDHc. In fact, PDHc was also effectively inhibited by C₄, C₁₀, C₁₄, and C₁₆ but less effectively by C₆ and C₈ (Figure 4A). In contrast, short-chain fatty acids (up to 1 mM) did not exert an inhibition of PDHc (Figure 4B). These results were confirmed in SMP from bovine heart (data not shown).

➤ **Figure 4**

Alternative Effects of Propionyl-CoA on Energy Metabolism?

Bioenergetic investigations in muscle tissue of both PA patients showed strongly reduced oxidation of pyruvate and PDHc activity as well as a diminished ATP plus phosphocreatine production (Table 1). In addition, decreased oxidation of succinate and malate as well as of almost all measured enzymes and enzyme complexes of mitochondrial respiratory chain and TCA cycle were found in these patients (Table 1).

➤ **Table 1**

Consistent with the enzymatic measurements is the decrease of the OXPHOS complexes I, II, III and IV in patient 2 observed with one and two dimensional BN-PAGE electrophoresis in combination with western blotting and a complex I in-gel activity assay (Figure 5).

➤ **Figure 5**

The general decrease in the amount of these OXPHOS complexes suggests a destabilisation and increased breakdown of mitochondria. The observed reduction in mitochondrial DNA (mtDNA) content of about 50% in patient 1 and 2 is in line with this suggestion. In analogy with these biochemical results, histopathology of muscle biopsies showed numerous signs of mitochondrial alteration indicating disturbance of mitochondrial energy metabolism, such as inclusion of many lipid droplets in muscle fibers of patient 1 (Figure 6A), cristalline inclusions (Figure 6B) and large swollen mitochondria with dark globular inclusions (Figure 6C).

➤ **Figure 6**

Since bioenergetic analyses in both PA patients suggested additional mechanisms acting in concert with propionyl-CoA-induced PDHc inhibition, we investigated whether propionyl-CoA and other short-chain acyl-CoA esters revealed inhibitory effects on respiratory chain and TCA cycle enzymes. We found no effect of short-chain acyl-CoA esters and fatty acids on respiratory complexes I, II, IV, and V in SMP (data not shown). However, in analogy to muscle tissue of PA patients, propionyl-CoA (1 mM) induced a mild inhibition of complex III activity in SMP ($72\pm 5\%$ of controls) and in isolated complex III from bovine heart ($81\pm 8\%$ of controls). In addition, the activity of KGDHc, a rate-limiting TCA cycle enzyme complex, was also decreased by propionyl-CoA (1 mM propionyl-CoA: $38\pm 3\%$ of controls).

DISCUSSION

The accumulation of metabolites from the alternative oxidative pathways of propionyl-CoA is the biochemical hallmark of PA. During metabolic crisis additional biochemical abnormalities such as lactic acidosis, hyperketosis and hypoglycemia indicate the development of severe mitochondrial dysfunction resulting in impairment of energy metabolism [1]. The present study significantly adds to the understanding of these mechanisms demonstrating 1) a severe bioenergetic disturbance and ultrastructural changes of mitochondria in muscle tissue of PA patients and 2) propionyl-CoA-induced synergistic inhibition of PDHc, KGDHc, and complex III *in vitro*. These results support the hypothesis that *secondary* mitochondrial dysfunction induced by accumulating toxic metabolites plays a major role in the pathomechanisms of propionic aciduria [7]. We clearly demonstrated that propionyl-CoA but not propionate is the major toxic metabolite involved in this scenario.

PDHc is a bioenergetic bottle neck coupling cytosolic anaerobic glycolysis with the mitochondrial TCA cycle and OXPHOS. Not surprisingly, inherited PDHc deficiency results in a severe disturbance of energy metabolism and frequently in a fatal disease course [20]. In analogy, *secondary* inhibition of PDHc by propionyl-CoA can be suggested as an important mechanism inducing energy failure in PA which is confirmed by severely reduced ATP and phosphocreatine concentrations in muscle tissue of PA patients and increased serum lactate concentrations. In line with this, multiple organ failure in PA patients predominantly manifests in tissues with a high energy demand, such as central nervous system (encephalopathy, movement disorders, developmental retardation), skeletal muscle (myopathy), heart muscle (cardiomyopathy), and bone marrow (pancytopenia) [1,3]. Apart from mammalian species, propionyl-CoA-induced inhibition of PDHc has recently been suggested as major mechanism underlying the antibacterial and antifungal properties of propionate [4]. Thus propionyl-CoA-induced mitochondrial dysfunction can be regarded as common final pathway involved in different conditions, i.e. inherited disorders of propionate metabolism and propionate toxicity. Future studies should focus on concentration-dependent effects of propionyl-CoA in tissues of affected children with PA. However, since the amount of a muscle biopsy specimen in children is always very limited because of ethical and functional considerations, this important aspect could not be investigated additionally in the present study.

In addition to PDHc inhibition, alternative mechanisms for propionyl-CoA and metabolites of alternative propionate oxidation should be considered as relevant. In particular inhibition of respiratory chain complex III (this study) and the TCA cycle enzymes KGDHc (this study), citrate synthase [5] and succinyl-CoA synthetase [4] may also contribute to impaired energy metabolism in this disease. Interestingly, inherited deficiency of succinyl-CoA synthetase caused by deleterious mutations in the *SUCLA2* gene has recently been associated with encephalomyopathy and mtDNA depletion [21], linking the TCA cycle with mtDNA homeostasis [21,22]. In addition, increased oxidative stress, which has been demonstrated in an *in vitro* model for disorders of propionate metabolism, induces mtDNA damage

[23,24]. Interestingly, the amount of mtDNA and the activities of OXPHOS complexes I, III, and IV, which are partially encoded by mtDNA, were significantly decreased in muscle tissue of both PA patients, however, it remains unclear whether this result reflects a causal link. Besides mtDNA homeostasis, other *secondary* or *tertiary* targets might be involved but have not yet been identified.

PA shares a variety of biochemical and clinical similarities with methylmalonic aciduria which is caused by inherited deficiency of methylmalonyl-CoA mutase or the synthesis or transport of its cofactor, 5'-adenosylcobalamin [1]. We have recently hypothesized that propionyl-CoA and metabolites deriving from propionyl-CoA, such as 2-methylcitrate, might act as endogenous neurotoxins also in this disease, whereas methylmalonate most likely plays a minor role [13,14,25]. Since the manifestation of secondary metabolic blocks are pathophysiologically relevant in PA and methylmalonic aciduria, it is of interest to investigate whether alternative energy substrates such as succinate and citrate might be beneficial for metabolic maintenance treatment and intensified emergency treatment of these patients helping to restore mitochondrial energy metabolism and to prevent multiple organ failure.

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FOOTNOTES

Abbreviations used: Coenzyme A, CoA; PDHc, pyruvate dehydrogenase complex; TCA, tricarboxylic acid; PA, propionic aciduria; OXPHOS, oxidative phosphorylation; SMP, submitochondrial particles from bovine heart; E1, pyruvate decarboxylase; E2, dihydrolipoyl transacetylase; E3, dihydrolipoyl dehydrogenase; C₂, acetyl-CoA; C₃, propionyl-CoA; C₄, butyryl-CoA; C₆, hexanoyl-CoA; C₈, octanoyl-CoA; C₁₀, decanoyl-CoA; C₁₄, myristoyl-CoA; C₁₆, palmitoyl-CoA; KGDHc, α -ketoglutarate dehydrogenase complex; E1k, α -ketoglutarate dehydrogenase; E2k, dihydrolipoyl succinyltransferase; BN-PAGE, blue native polyacrylamide-gel electrophoresis; ANOVA, one-way analysis of variance; K_{is} , inhibition constant; mtDNA, mitochondrial DNA; K_m , Michaelis-Menten constant; Pyr, pyruvate; n.m., not measured

¹To whom correspondence should be addressed: Stefan Kölker, Department of General Pediatrics, Division of Inborn Metabolic Diseases, Im Neuenheimer Feld 150, D-69120 Heidelberg, Germany. Tel.: +49-6221-561714; Fax: +49-6221-565565; E-mail:Stefan_Koelker@med.uni-heidelberg.de

[§]Both authors contributed equally to the study.

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FIGURE LEGEND

FIGURE 1. Inhibition of PDHc by Propionyl-CoA.

Spectrophotometric analysis in purified PDHc from porcine heart demonstrates uncompetitive inhibition of PDHc activity by propionyl-CoA with respect to pyruvate (Pyr) as demonstrated by a Lineweaver Burk plot. Values are given as the mean \pm SD. Experiments were performed at least in triplicates.

FIGURE 2. Inhibition of PDHc by Acetyl-CoA.

Spectrophotometric analysis in purified PDHc from porcine heart demonstrates uncompetitive inhibition of PDHc activity by acetyl-CoA with respect to pyruvate (Pyr) as demonstrated by a Lineweaver Burk plot. Values are given as the mean \pm SD. Experiments were performed at least in triplicates.

FIGURE 3. Spectrophotometric Analysis of PDHc Activity in Human Skin Fibroblasts.

Spectrophotometric analysis of PDHc activity was performed in human skin fibroblasts from healthy volunteers ($n=12$), patients with propionic aciduria ($n=2$) and primary PDHc deficiency ($n=3$). The horizontal bar represents the mean of the control group and the dotted line indicates the range of the control group.

FIGURE 4. Chain-length specific effects of Acyl-CoA Esters and fatty acids on PDHc activity. (A)

Acyl-CoA Esters: Spectrophotometric analysis of PDHc activity was performed in purified enzyme from porcine heart using short-, medium- and long-chain monocarboxylic acyl-CoA esters. The figure summarizes the I_{50} values of acyl-CoA esters. Values are given as mean \pm SD. Experiments were performed at least in triplicates. * $p<0.01$; ** $p<0.001$ versus control (ANOVA followed by *posthoc* Bonferroni's multiple comparison test, $n=8$). (B) Fatty Acids: In contrast to acyl-CoA esters, corresponding short- and medium-chain fatty acids (up to 1 mM) did not inhibit PDHc activity in purified enzyme from porcine heart. PDHc activity was determined at standard conditions. Values were normalized to 100% (dotted line) and values were given as the mean \pm SD. Experiments were performed at least in triplicates.

FIGURE 5. BN-PAGE of OXPHOS Complexes in Muscle Homogenates.

Protein (40 μ g) of solubilised muscle homogenates from patient 2 (P) and control (C) were analyzed on a 5-15% BN-polyacrylamide gel for the separation of multi-subunit complexes. (A) In-gel activity assay of mitochondrial complex I confirming a decrease in activities of patient 2 compared to control samples. (B) A second gel was run in duplicate and western blot analysis was performed using antibodies against complex I subunit 39 kDa, complex II subunit 70 kDa and complex III subunit Core 2 (C). A second

dimension was run and western blot analysis was performed using antibodies against OXPHOS complexes I (ND6), III (Core 2) and IV (COX2) and cyclophilin B as a loading control. Arrows indicate the first and second dimension.

FIGURE 6. Ultrastructural Changes in Muscle Tissue of PA Patients.

Electromicroscopy was performed in muscle biopsy specimens (Musculus vastus lateralis dextra) of both PA patients. (A) Patient 1: Muscle fiber with many lipid droplets (the large clear globules). (B) Patient 1: Mitochondrion with a crystalline inclusion. (C) Patient 2: Large swollen mitochondrion with disrupted cristae and two mitochondria with a dark globular inclusion. Bars represent 5 μm (A), 100 nm (B), and 1 μm (C).

TABLE 1

Bioenergetical Analysis in Muscle Tissue of PA patients.

Radiochemical analysis of coupled mitochondria (A) using [1-¹⁴C]pyruvate, [U-¹⁴C]malate and [1,4-¹⁴C]succinate. Spectrophotometric analysis of single enzymes of the respiratory chain, TCA cycle, PDHc (B), ATP and phosphocreatine production rate (C), and citrate synthase (D). Analysis were performed in muscle biopsy specimens of both patients. Activities are expressed as nmol/h×mg protein (A), milliunits/unit citrate synthase (B), nmol/h×milliunits citrate synthase (C) and milliunits/mg protein (D). The production of ATP and phosphocreatine was determined to establish the total mitochondrial energy generating capacity of the muscle mitochondria of both patients. n. m., not measured.

Experiment	Patient 1	Patient 2	Control Range
A. Oxidation of pyruvate, malate and succinate			
[1- ¹⁴ C]pyruvate+malate	0.13	0.56	3.61–7.48
[1- ¹⁴ C]pyruvate+carnitine	0.21	0.59	2.84–8.24
[U- ¹⁴ C]malate+pyruvate+malonate	0.18	0.32	4.68–9.62
[U- ¹⁴ C]malate+acetylcarnitine+malonate	0.46	1.32	3.43–7.30
[U- ¹⁴ C]malate+acetylcarnitine+arsenite	0.43	1.45	2.05–6.39
[1,4- ¹⁴ C]succinate+acetylcarnitine	0.27	0.82	2.54–6.39
B. Single enzyme activity			
Complex I	16	34	70-250
Complex II	59	38	67-177
Complex III	446	989	2,200-6,610
Complex IV	507	593	810-3,120
Complex II+coenzymeQ+complex III	87	118	300-970
PDHc	6	10	34-122
PDHc-E1	1.08	1.52	1.23-4.13
PDHc-E3	353	533	760-2,630
KGDHc	n. m.	3	23-49
C. ATP and phosphocreatine			
ATP+phosphocreatine	10	0.0	42-81
D. Mitochondrial reference enzyme			
Citrate synthase	81	147	37-162

Figure 1

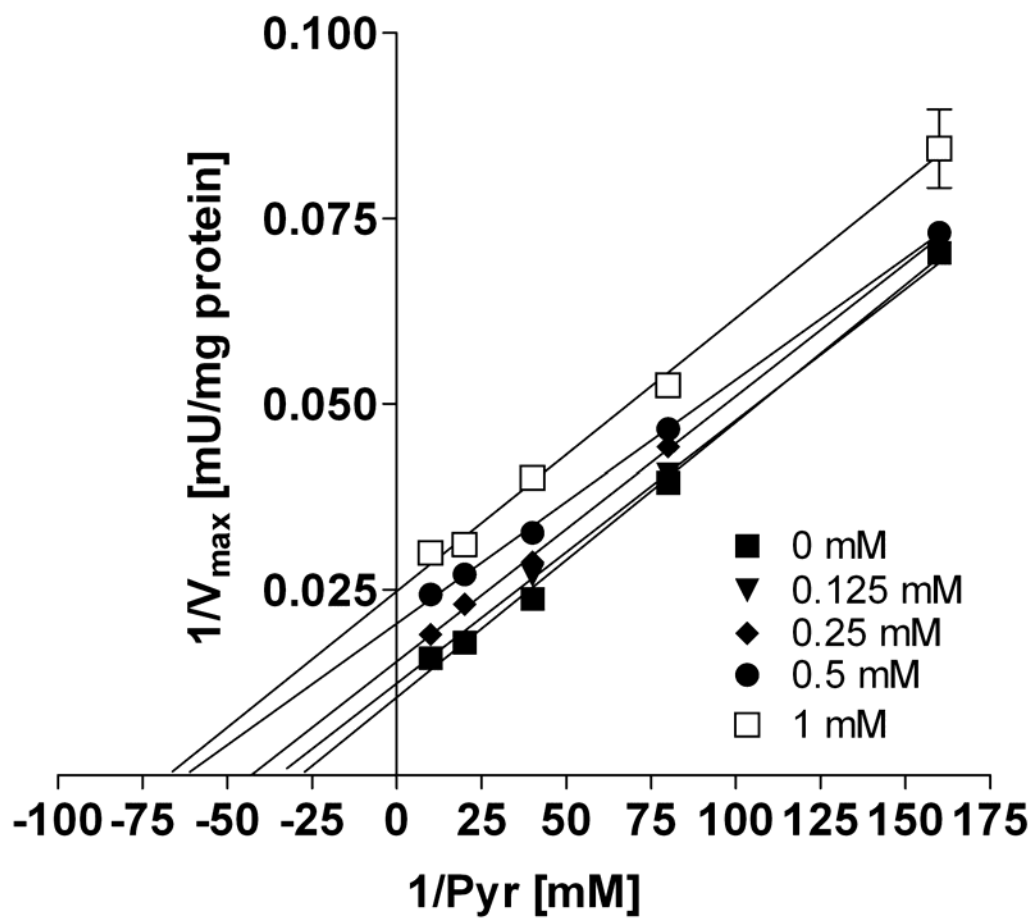


Figure 2

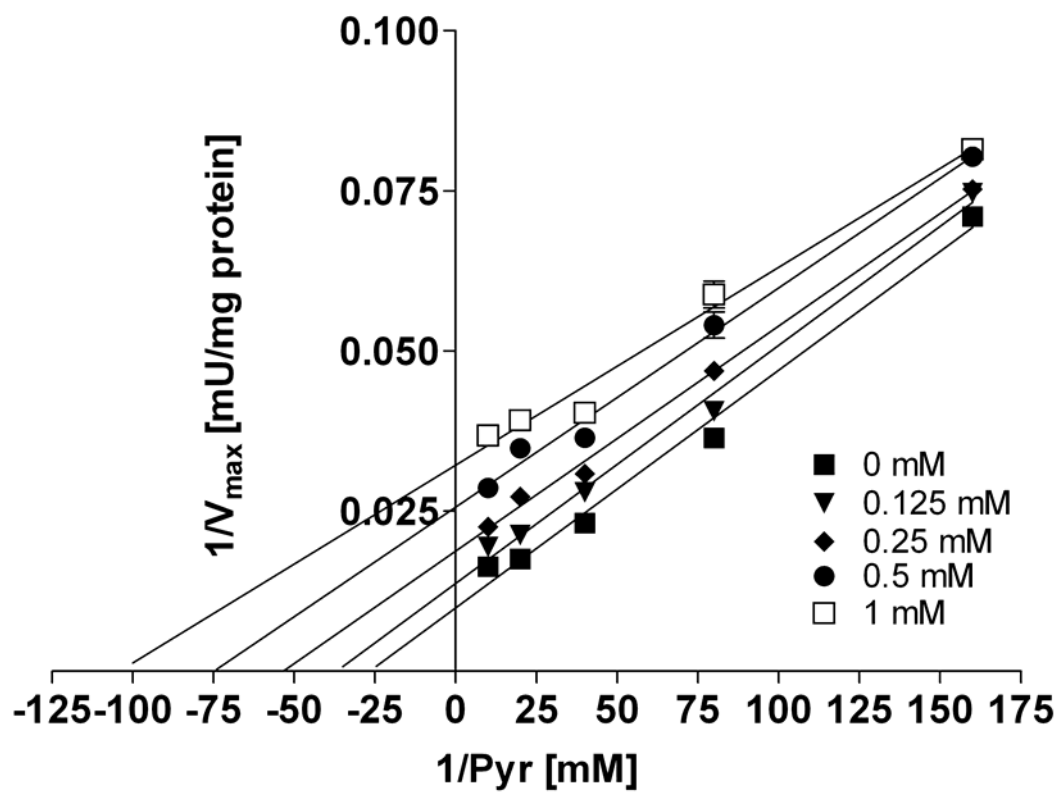


Figure 3

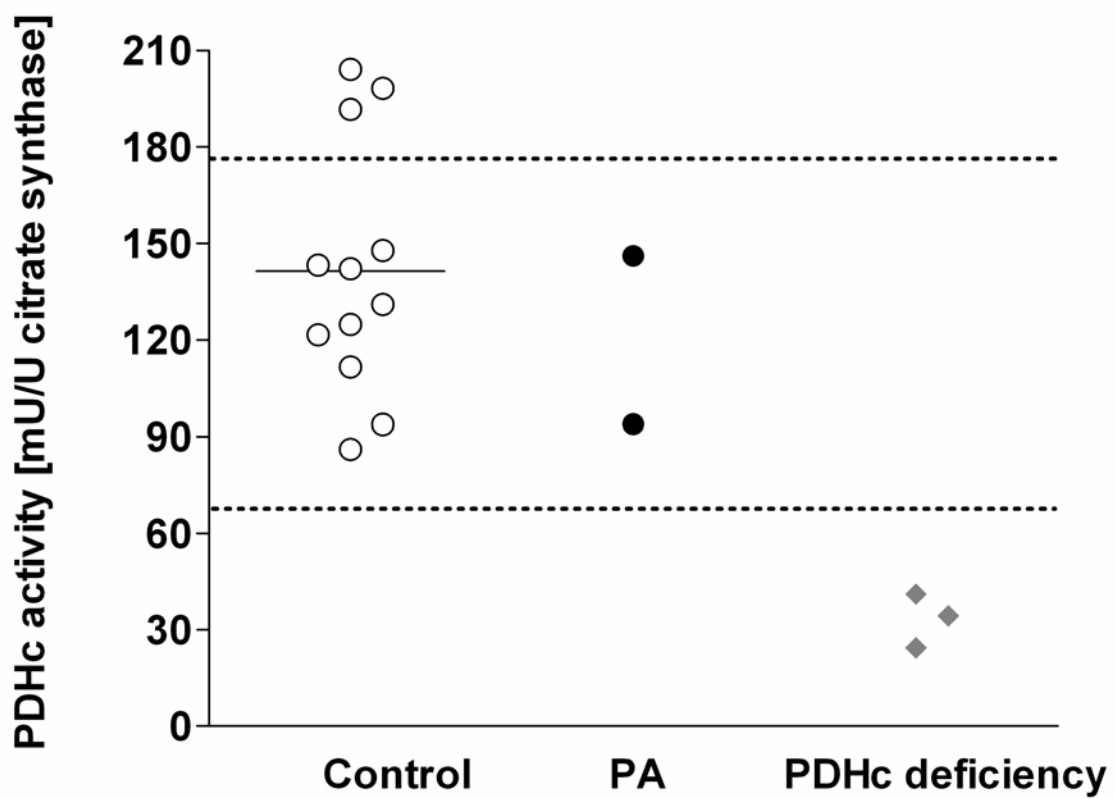


Figure 4

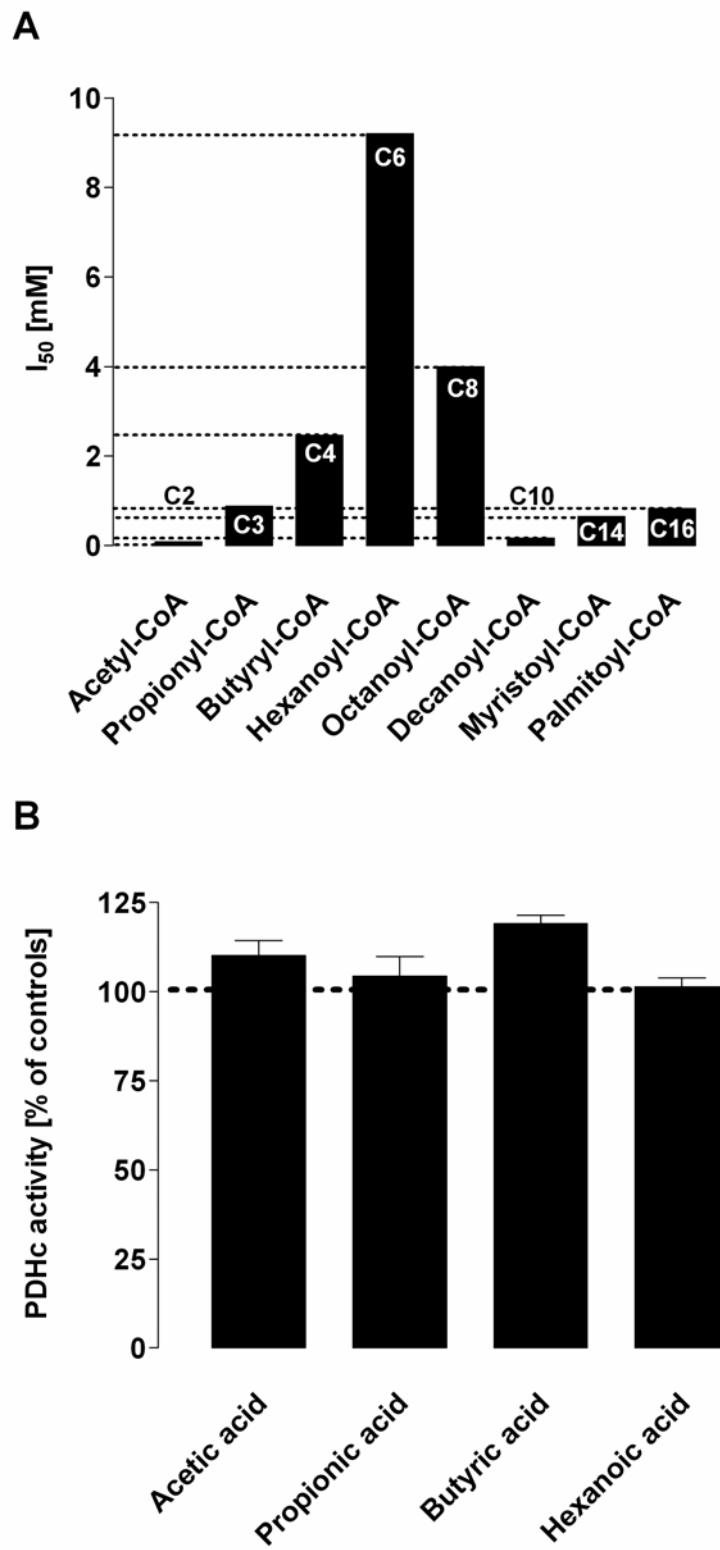


Figure 5

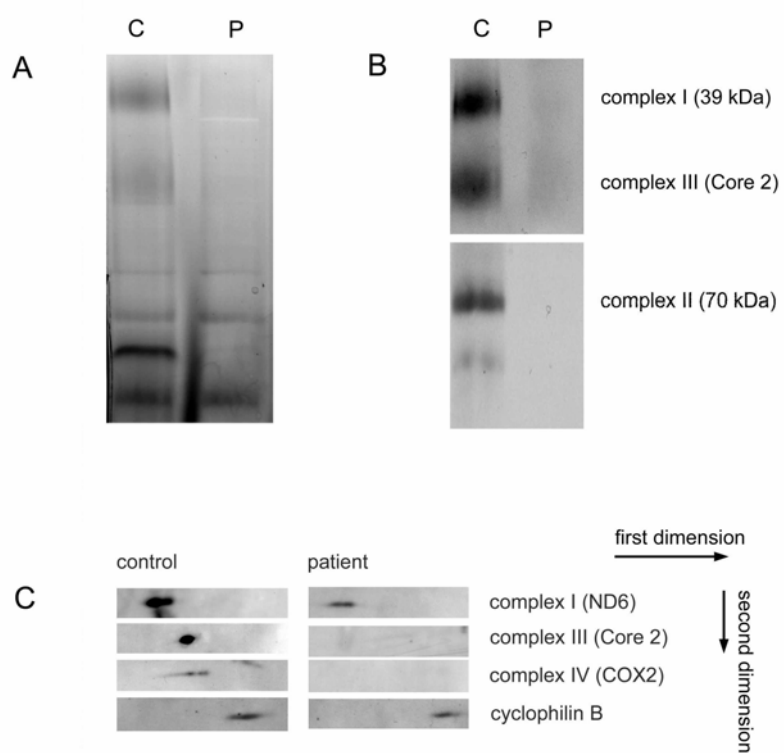


Figure 6

