Rosiglitazone and fenofibrate improve insulin sensitivity of pre-diabetic OLETF rats by reducing malonyl-CoA levels in the liver and skeletal muscle

Zhengshan Zhao\textsuperscript{a,b,c}, Yong-Jik Lee\textsuperscript{a,b}, Soo-Kyung Kim\textsuperscript{b,d}, Hae-Jin Kim\textsuperscript{a}, Wan-Sub Shim\textsuperscript{a}, Chul-Woo Ahn\textsuperscript{a,b}, Hyun-Chul Lee\textsuperscript{a,b}, Bong-Soo Cha\textsuperscript{a,b,⁎}, Zhongmin Alex Mac\textsuperscript{c,⁎}

\textsuperscript{a} Department of Internal Medicine, Yonsei University College of Medicine, Seoul, Republic of Korea
\textsuperscript{b} Brain Korea 21 Project for Medical Science, Yonsei University College of Medicine, Seoul, Korea
\textsuperscript{c} Division of Experimental Diabetes and Aging, Mount Sinai School of Medicine, New York, USA
\textsuperscript{d} Department of Internal Medicine, College of Medicine, Pochon CHA University, Sungnam, Kyonggi-Do, Republic of Korea

\textsuperscript{⁎} Corresponding authors. Cha is to be contacted at Department of Internal Medicine, College of Medicine, Yonsei University, 134 Shinchon-Dong, Seodaemoon-Ku, PO. Box 120-749, Seoul, Republic of Korea. Ma, Division of Experimental Diabetes and Aging, Mount Sinai School of Medicine, New York, NY 10029, USA.

\texttt{E-mail addresses: bsha@yuhs.ac (B.-S. Cha), zhongmin.ma@mssm.edu (Z.A. Ma).}

Abstract

Aims: Rosiglitazone and fenofibrate, specific agonists of the peroxisome proliferator activated receptors-γ (PPAR\textsubscript{γ}) and -α (PPAR\textsubscript{α}), respectively, improve insulin sensitivity in diabetic animals and in patients with type 2 diabetes. Here we investigated how pre-diabetic Otsuka Long–Evans Tokushima Fatty (OLETF) rats fed with normal and high-fat diets respond to these PPAR agonists.

Main methods: Pre-diabetic OLETF rats were subjected to high-fat or standard diets with or without rosiglitazone and fenofibrate for 2 weeks. The metabolism of the rats and the levels of malonyl-CoA and activities of malonyl-CoA decarboxylase (MCD), acetyl-CoA carboxylase (ACC), and AMP-activated protein kinase (AMPK) in metabolic tissues were assessed.

Key findings: Rosiglitazone and fenofibrate significantly improved insulin sensitivity and reduced the levels of plasma triglycerides and free fatty acids in OLETF rats fed with a high-fat diet. Fenofibrate particularly reduced the body weight, fat, and total cholesterol in high fat diet OLETF rats. The highly elevated malonyl-CoA levels in the skeletal muscle and liver of OLETF rat were significantly reduced by rosiglitazone or fenofibrate due to, in part, the increased MCD activities and expression. On the other hand, ACC activities were unchanged in skeletal muscle and decreased in liver in high fat diet group. AMPK activities were dramatically decreased in OLETF rats and not affected by these agonists.

Significance: These results demonstrate that treatment of pre-diabetic OLETF rats—particularly those fed a high-fat diet—with rosiglitazone and fenofibrate significantly improves insulin sensitivity and fatty acid metabolism by increasing the activity of MCD and reducing malonyl-CoA levels in the liver and skeletal muscle.

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Introduction

Rosiglitazone and fenofibrate, specific agonists of peroxisome proliferator activated receptors-γ (PPAR\textsubscript{γ}) and -α (PPAR\textsubscript{α}), respectively, improve insulin sensitivity in diabetic animals (Guerre-Millo et al. 2000; Kramer et al. 2001; Wang et al. 2001), and in people with type 2 diabetes and insulin resistance syndrome (Mudalier and Henry et al. 2000; Kramer et al. 2001; Wang et al. 2001), and in people with type 2 diabetes. Here we investigated how pre-diabetic Otsuka Long–Evans Tokushima Fatty (OLETF) rats fed with normal and high-fat diets respond to these PPAR agonists.

PPAR\textsubscript{γ} is the key transcriptional regulator of adipogenesis and it plays a critical role in glucose homeostasis (Lehmann et al. 1995; Rosen et al. 1999). PPAR\textsubscript{α} is critical for peroxisome proliferation (Lee et al. 1995) and it functions as fatty acid sensor important for the regulation of fatty acid metabolism and energy homeostasis (Blaschke et al. 2006). Although PPAR\textsubscript{γ} and PPAR\textsubscript{α} play specific and separate roles under physiological conditions, their agonists have similar effects on triglyceride levels in the blood of diabetic animals (Chaput et al. 2000). Malonyl-CoA is a potent inhibitor of mitochondrial carnitine palmitoyltransferase (CPT1), the key regulatory enzyme involved in the first committed step of fatty-acid oxidation in mitochondria (Rasmussen et al. 2002), and its levels are precisely regulated in various nutritional states (Chien et al. 2000). Inhibition of CPT1 by malonyl-CoA decreases the uptake of fatty acids into the mitochondria and slows mitochondrial fatty acid oxidation (Mcgarry...
Malonyl-CoA is synthesized by acetyl-CoA carboxylase (ACC) (Thampy 1989) and degraded to acetyl-CoA and carbon dioxide by malonyl-CoA decarboxylase (MCD) (Saha et al. 2000). It has been reported that exercise-induced activation of AMPK controls malonyl-CoA levels by coordinating the functions of MCD and ACC (Park et al. 2002).

Although PPAR agonists markedly reduce blood triglyceride levels in diabetic animals (Chaput et al. 2000), to our knowledge their effect on the regulation of malonyl-CoA has not been investigated in the tissues of non-exercising pre-diabetic rats. In this study we investigated the effect of rosiglitazone and fenofibrate on insulin resistance and malonyl-CoA levels in the liver and skeletal muscles of Otsuka Long–Evans Tokushima Fatty (OLETF) rats. These rats spontaneously develop mild diabetes at 18 weeks of age and are pre-diabetic at 14 weeks of age with normal fasting glucose levels and high insulin levels (Kawano et al. 1992, 1994). We treated 14-week-old pre-diabetic OLETF rats fed with a normal or high-fat diet with rosiglitazone or fenofibrate for 2 weeks and found that these agonists stimulate MCD activity, decrease malonyl-CoA levels in the liver and skeletal muscles, and improve insulin sensitivity.

Materials and methods

Materials

[1, 3-14C]malonyl-CoA was purchased from American Radiolabeled Chemicals, Inc. (St. Louis, MO, USA). [1-314C]sodium bicarbonate from Moravek Biochemicals (Brea, CA, USA), [γ-32P]ATP from Amersham Biosciences (Piscataway, NJ, USA), and rat insulin RIA kit from Linco Research, Inc. (St. Charles, MO, USA). TaqMan EZ RT-PCR Core Reagents and the primers and probes of MCD and cyclophilin were purchased from Applied Biosystems (Foster City, CA, USA). The AMARAASAAALLRRR (AMARA) peptide was synthesized by and purchased from Peptron Inc. (Taejon, Korea). High-fat diet was purchased from BioGenomics, Inc. (Harlan, CA, USA). Whole blood glucose analyzer was obtained from Johnson & Johnson (Miltonias, CA, USA). Protease inhibitors, phosphatase inhibitors, and all other reagents were purchased from Sigma Chemical (St. Louis, MO, USA).

Animals and diet composition

Four-week-old male Long–Evans Tokushima Otsuka (LETO) rats (n = 20) and Otsuka Long–Evans Tokushima Fatty (OLETF) rats (n = 60) were donated by Otsuka Pharmaceuticals (Japan) and were group housed until the experiment was completed. At 14 weeks of age, the rats were randomly assigned to one of seven dietary regimens: 1) LETO (n = 20) standard rat diet, 2) OLETF (n = 10) standard rat diet, 3) OLETF (n = 10) standard rat diet with rosiglitazone (4 mg/kg/d), 4) OLETF (n = 10) standard rat diet with fenofibrate (100 mg/kg/d), 5) OLETF (n = 10) high-fat diet, 6) OLETF (n = 10) high-fat diet with rosiglitazone (4 mg/kg/d), 7) OLETF (n = 10) high-fat diet with fenofibrate (100 mg/kg/d). The standard diet comprises approximately 3.6 kcal/g: 21% protein, 12.5% fat, and 66.5% carbohydrates and the high-fat diet comprises approximately 5.0 kcal/g: 21% protein, 66.5% fat, 12.5% carbohydrates as described (Bi et al. 2007). All rats were kept in the Department of Animal Experiment of Clinical Medical Research Center, Yonsei University Medical College, under controlled conditions of 23 ± 1 °C and 12 h light:12 h dark cycle, and given free access to food and water ad libitum. All rats were cared for as outlined in the Guidelines of Animal Experiments recommended by Korean Academy Sciences.

Body weight and oral glucose tolerance test

Food consumed was checked every day, and body weight was checked every 3 days during the experiments. Oral glucose tolerance tests (OGTT) were conducted at the beginning and end of each experiment. Rats were given glucose orally (2 g/kg) and blood samples were collected from their tails 0, 30, 60, 90, 120 min later. Glucose levels were measured with a glucose analyzer (Milpitas, CA, USA).

Tissue sampling and measurement of fat mass

After 2 weeks of feeding with the normal or high-fat diet in the presence or absence of rosiglitazone or fenofibrate, animals were anesthetized with isoflurane and sacrificed for tissue sampling. Blood was collected by cardiac puncture for triglyceride (TG), total cholesterol (TC), free fatty acid (FFA), and insulin assays. Abdominal subcutaneous fat (SQ fat) and epididymal fat (epididymal pads) were surgically removed through a mid-abdominal incision and the weight of each dissected fat mass was recorded immediately. Skeletal muscles (gastrocnemius), livers, and pancreas were immediately isolated, freeze-clamped in liquid nitrogen, and stored at −80 °C until assays were performed.

Serum lipids and insulin, and pancreatic insulin content measurement

Triglyceride, total cholesterol, and free fatty acid levels were measured with infinity triglycerides reagent, infinity cholesterol reagent, and the ACS-ACOD enzyme method (NEFA ZYME-S, Aiken, Japan). To measure pancreatic insulin contents, pancreases were homogenized with an electric homogenizer in 500 μl of acid ethanol buffer (1.5 ml of 12 mol/L HCl in 100 ml of 70% ethanol) and extracted overnight at 4 °C. Insulin levels in the serum and pancreatic extract were measured with a rat RIA kit according to the manufacturer’s instructions.

Determination of malonyl-CoA levels

Malonyl-CoA was extracted from frozen (−80 °C) skeletal muscles and livers as previously described (King et al. 1988). The 6% perchloric acid extract was maintained at a pH of 2–3. Malonyl-CoA was measured with a modified high-performance liquid chromatography (HPLC) procedure described previously (King et al. 1988; Saddik et al. 1993) on a Beckman System Gold with a UV detector 167. Each sample (100 μl) was run through a pre-column cartridge (C18, size 3 cm, 7 μm) and a Microsorb short-one column (type C18, particle size 3 μm, size 4.6 x 100 mm). Absorbance was set at 254 nm and flow rate at 1 ml/min. A gradient was initiated using two buffers: buffer A consisted of 0.2 M NaH2PO4 (pH 5.0) and buffer B was a mixture of 0.25 M NaH2PO4 and acetonitrile (pH 5.0) in a ratio of 80:20 (v/v). Buffers were filtered using filter pure, Nylon-66 filter membrane (Pierce Chemical Co.). Initial conditions (97% buffer A, 3% buffer B) were maintained for 2.5 min and were changed thereafter to 18% buffer B over 5 min using Beckman’s curve 3. At 15 min the gradient was changed linearly to 37% buffer B over 3 min and subsequently 90% buffer B over 17 min. At 42 min the composition was returned linearly back to 3% buffer B over 0.5 min, and at 50 min column equilibration was complete. Peaks were integrated by a Beckman System Gold software package.

Determination of ACC activities

ACC activities were measured as described previously (Sakamoto et al. 2000). Briefly, approximately 200 mg of frozen tissues were homogenized using an electric homogenizer with a buffer containing Tris–HCl (10 mM), mannitol (200 mM), NaF (50 mM), EDTA (1 mM), 2-mercaptoethanol (10 mM), pH 7.5, and three proteolytic enzyme inhibitors (aprotinin, leupeptin, and antipyrin), at a concentration of 5 mg/L. The homogenate was immediately centrifuged at 48,000 g for 30 min at 4 °C. The fraction containing ACC and AMPK was precipitated from the supernatant by addition of 144 mg ammonium bicarbonate (1 M). The supernatant was centrifuged at 10,000 g for 10 min and ACC activity was measured using a modified assay (Sakamoto et al. 2000).
sulfate/ml and by stirring for 60 min on ice. The precipitate was collected by centrifugation at 48,000 g for 30 min. The pellet was dissolved in 10% of the original volume of the homogenate buffer and centrifuged again to remove insoluble protein. The supernatant was assayed for ACC and AMPK activity.

ACC activity was measured using the CO2 fixation method, as described previously (Sakamoto et al. 2000). Briefly, 5 µl of the above supernatant containing 20 µg of total protein was added to a reaction mixture (final volume, 165 µl) containing tris acetate (60.6 mM), BSA (1 mg/ml), 2-mercaptoethanol (1.32 µmol/l), ATP (2.21 mM), acetyl-CoA (1.06 mM), magnesium acetate (5.0 mM), and NaHCO3 (18.08 mM). Samples were incubated at 37 °C for 10 min, and the reaction was stopped by adding 25 µl of 10% perchloric acid. Samples were centrifuged for 20 min at 3500 rpm, and 160 µl of supernatant was placed in minivials and dried in a fume hood overnight. H2O (100 µl) followed by scintillant, was added to the vials, and the vials were counted. ACC activity was expressed as the amount of malonyl-CoA produced/min/mg protein.

**Determination of AMPK activities**

AMPK activity was measured as previously described (Sakamoto et al. 2000). Briefly, 2 µl of the supernatant prepared as described above for the determination of ACC activity was added to a reaction mixture composed of HEPES-NaOH (40 mM), NaCl (80 mM), glycerol (8% wt/vol), EDTA (0.8 mM), AMARAASAAALARRR (AMARA) peptide (200 µmol/l), dithiothreitol (DTT) (0.8 mM), [γ−32P]ATP (200 µmol/1), MgCl2 (5 mM), and 0.18% Triton X-100. Samples were incubated in the presence or absence of 200 µM AMP for 3 min at 30 °C. 15 µl of this mixture was spotted on 1 cm2 phosphocellulose paper and the paper was washed four times for 10 min each with 150 mM phosphoric acid, followed by a 5 min acetic acid wash. The papers were then dried and counted for radioactivity. AMPK activity was expressed as nmol of 32P incorporated into the AMARA peptide/min/mg protein.

**Determination of MCD activities**

Approximately 300 mg of frozen tissues was homogenized in a buffer of 0.1 M Tris–HCl (pH 8.0), 2 mM PMSE, 5 µM aprotinin, 5 µM leupeptin, and 5 µM pepstatin A. 40 mM β-glycerophosphate, 40 mM NaF, 4 mM NaPi, and 1 mM Na3VO4 were added to inhibit phosphatase activity. The homogenates were then centrifuged at 500 g for 10 min. Powdered (NH4)2SO4 was slowly added to the supernatant with stirring until 40% (243 g/L of ammonium sulfate) saturation was achieved. The mixture was recentrifuged at 14,000 g for 10 min. The supernatant from this spin was dissolved in 0.1 M Tris–HCl (pH 8.0) and MCD activity was determined as described below.

The enzyme activity was assayed by measuring the amount of 14CO2 generated from [3-14C]malonyl-CoA, as previously described (Goodwin and Taegtmeyer 1999) with the following modification. Briefly, a reaction mixture of 10 µmol of Tris–HCl buffer (pH 8.0), 0.01 µmol of dithioerythritol (DTE), 0.02 µmol of [3-14C]malonyl-CoA was incubated with enzyme in a total volume of 0.1 ml for 10 min at 37 °C. The 14CO2 generated by [3-14C] malonyl-CoA was trapped in filter papers soaked with 2 N KOH and assayed by liquid scintillation spectrometry. Enzyme activity is expressed as nmol/min/mg protein.

**RNA extraction and real-time RT-PCR**

Total RNA was extracted from frozen (−80 °C) tissues with TRIzol Reagent (GIBCO BRL) according to the manufacturer’s instructions. Real-time RT-PCR was performed using an ABI PRISM 7700 Sequence Detection System instrument and software (PE Applied Biosystems). Primers and probes for MCD and cyclophilin were used as described previously (Young et al. 2001). MCD primer/probe: forward 5′-CGGCACCTTCTCTAAATACG-3′, reverse 5′-GGGTATAGTGACAGGCTGGA-3′, probe 5′-FAM-AGTGCTGAAAGA GCTGCAAGAATGTT-3′; Cyclophilin primer/probe: forward 5′-CTGTCAGCCAGCCTG-3′, reverse 5′-CTCTTGCTTCTTGGAACTTTGTC-3′, probe 5′-FAM-CCGGCTGGTCCTCCGACTCTTCCGGA-3′. The level of transcripts for the constitutive housekeeping gene cyclophilin was measured in triplicate in each sample to adjust for sample-to-sample differences in RNA concentration. MCD mRNA expression was calculated with the comparative CT (threshold cycle) method and reported as the ratio of MCD transcripts per cyclophilin transcripts.

**Statistical analysis**

Data are presented as means ± SEM. Statistically significant differences between groups were calculated by Tukey’s Multiple Comparison Test of One-Way ANOVA. A value of p < 0.05 is considered significant.

**Results**

**Effect on body weight, and fat mass**

To examine the effects of PPARγ and α agonists on pre-diabetic OLETF rats, we fed 14-week-old OLETF rats with normal or high-fat diets and measured their body weight and fat mass after 2 weeks of treatment with rosiglitazone or fenofibrate (Table 1). We found that the body weights of all groups of OLETF rats were significantly higher than those of LETO controls. Rosiglitazone and fenofibrate treatment did not significantly affect the phenotype of rats fed a normal diet. However, fenofibrate, but not rosiglitazone, significantly reduced body weight and abdominal subcutaneous and epididymal fat mass, in rats fed a high-fat diet.

**Effect on blood lipids, cholesterol, insulin levels, and pancreatic insulin content**

We then measured the levels of blood lipids, total cholesterol, and insulin in serum and in the pancreas in pre-diabetic rats after treatment with rosiglitazone and fenofibrate (Table 2). Serum triglycerides, free fatty acids, total cholesterol, insulin and pancreatic
insulin content were elevated in all high fat diet groups of OLETF rats, indicating insulin resistance and abnormalities in lipid metabolism. In normal-diet groups, the levels of triglycerides and free fatty acids were significantly reduced by treatment with either rosiglitazone or fenofibrate. However, serum triglyceride, free fatty acid, and total cholesterol levels, and insulin were significantly reduced in the presence of either rosiglitazone or fenofibrate in high-fat diet groups, suggesting a significant improvement in lipid metabolism and insulin sensitivity.

**Effects on oral glucose tolerance test**

We next performed the oral glucose tolerance test (OGTT) (2 g/kg body weight) on 16-week-old pre-diabetic OLETF rats treated with rosiglitazone and fenofibrate. As shown in Fig. 1, both groups developed impaired glucose tolerance. This impairment was more severe in the group fed the high-fat diet, despite the fact that these rats still exhibited normal fasting glucose levels. However, treatment of pre-diabetic OLETF rats with rosiglitazone or fenofibrate significantly improved insulin sensitivity in both the normal (Fig. 1A) and the high-fat (Fig. 1B) diet groups. Our results clearly show that PPAR-γ and -α agonists can improve insulin sensitivity during the pre-diabetic period even in the presence of a high-fat diet.

**Effect on malonyl-CoA levels in skeletal muscle and liver**

Since PPAR-γ and -α agonists significantly reduced serum levels of triglycerides and free fatty acids in pre-diabetic rats fed either a normal or a high-fat diet (Table 2), we investigated whether these agonists affect the levels of malonyl-CoA in skeletal muscle and liver. As shown in Fig. 2, malonyl-CoA levels were highly elevated in both the skeletal muscle and livers of pre-diabetic OLETF rats compared to control rats. This elevation was more significant in rats fed with a high-fat diet, suggesting an abnormal regulation of malonyl-CoA in the pre-diabetic OLETF rats. Malonyl-CoA levels in skeletal muscles of

![Fig. 1](image1.png) Results of the oral glucose tolerance test (OGTT). OGTT (2 g/kg body weight) was performed after pre-diabetic OLETF rats were treated for 2 weeks with rosiglitazone and fenofibrate. A. OGTT in 16-week-old OLETF rats fed a normal diet. *p < 0.05, compared with normal control (LETO); †p < 0.05, compared with OLETF control fed a normal diet; ‡p < 0.01 vs. normal diet fed OLETF control; and, *p < 0.05 vs. high fat diet fed OLETF control. Values are means ± SEM of 10 rats in each group.

![Fig. 2](image2.png) Malonyl-CoA levels in the skeletal muscle and livers of OLETF rats fed either normal or high-fat diet. A. Malonyl-CoA levels in skeletal muscle. B. Malonyl-CoA levels in liver. Values are expressed as means ± SEM (n = 10). †p < 0.05, vs. LETO; ‡p < 0.01, vs. OLETF controls in the normal-diet group; *p < 0.01 vs. corresponding OLETF control. Ctrl, control; RSG, rosiglitazone; FNF, fenofibrate.
OLETF rats fed a normal diet were 25% lower in the rosiglitazone-treated animals and 33% lower in the fenofibrate-treated animals. In OLETF rats fed a high-fat diet, rosiglitazone treatment decreased malonyl-CoA levels in skeletal muscles by 35% and fenofibrate treatment by 43% (Fig. 2A). In OLETF rats fed a normal diet, rosiglitazone treatment decreased malonyl-CoA levels in the liver by 24% and fenofibrate treatment by 27% (Fig. 2B). In the high-fat diet group, rosiglitazone treatment reduced malonyl-CoA levels in the liver by 30% and fenofibrate treatment by 41% (Fig. 2B). Taken together, our results clearly indicate that both rosiglitazone and fenofibrate can reduce the levels of malonyl-CoA in skeletal muscle and liver in this pre-diabetic rat model.

Effect on MCD and ACC activities

Since malonyl-CoA levels in tissues are regulated by MCD and ACC, we measured the activities of these two enzymes in skeletal muscle. As shown in Fig. 3A, rosiglitazone and fenofibrate significantly increased MCD activity in rats fed either a normal or a high-fat diet (Fig. 3A). Interestingly, MCD activity was more dramatically enhanced by fenofibrate than by rosiglitazone in rats fed a high-fat diet. However, ACC activity in skeletal muscle was not significantly changed after treatment (Fig. 3B). These results suggest that rosiglitazone and fenofibrate decrease malonyl-CoA levels at least in part by increasing MCD activity in skeletal muscle in OLETF rats fed either a normal or a high-fat diet.

We next examined the effect of rosiglitazone and fenofibrate on MCD and ACC activities in the liver. We found that fenofibrate, but not rosiglitazone, significantly increased the activity of MCD in OLETF rats fed a normal diet. Both rosiglitazone and fenofibrate increased the activity of MCD in rats fed a high-fat diet (Fig. 4A). In contrast to ACC activity in rats fed a normal diet, ACC activity was elevated in rats fed a normal diet (Fig. 4B).
high-fat diet. Rosiglitazone suppressed this elevated ACC activity by 38% and fenofibrate by 87% (Fig. 4B). These results suggested that the decrease we observed in malonyl-CoA levels in the livers of pre-diabetic rats fed a high-fat diet is due to both an increase in degradation by MCD and a decrease in synthesis by ACC.

Effect on AMPK activities in skeletal muscle and liver

It has been previously reported that AMPK regulates MCD and ACC activities under specific conditions, such as during exercise or the electric stimulation of skeletal muscles (Park et al. 2002). To examine whether AMPK regulates MCD and ACC activities in resting state in pre-diabetic OLETF rats, we examined its activity in skeletal muscle and liver. Surprisingly, the activity of AMPK in both tissues was dramatically lower in this pre-diabetic rat model than in control rats (Fig. 5). In addition, rosiglitazone and fenofibrate did not affect AMPK activity in the tissues of OLETF rats regardless of diet. These results suggest that the AMPK signaling network may be defective in the skeletal muscles and livers of OLETF rats.

MCD mRNA expression in skeletal muscle and liver

PPARs are transcriptional regulators. Since PPAR agonists increase MCD activity in both skeletal muscle and liver, we used real-time RT-PCR to examine whether they affect the transcription of MCD in OLETF rats fed a normal or high-fat diet. As shown in Fig. 6A, the levels of MCD mRNA in the skeletal muscles of OLETF rats were significantly lower than that LETO rats and restored by fenofibrate, but not rosiglitazone. Interestingly, we observed that MCD mRNA levels were significantly higher in rats fed a high-fat diet than in rats fed a normal diet. This finding suggests that, consistent with previous reports, increased availability of fatty acids may promote their utilization by increasing the expression of MCD in skeletal muscles (Young et al. 2001). As shown in Fig. 6B, the levels of MCD mRNA in the livers of OLETF rats are similar regardless of whether the rats are fed a normal or a high-fat diet. As in skeletal muscle, fenofibrate, but not rosiglitazone, significantly increased the expression of MCD in the livers of OLETF rats. Our data suggest that PPAR-α, but not PPAR-γ, agonists stimulate the transcription of MCD in skeletal muscle and liver, and that a high-fat diet alters this transcription in skeletal muscle.

Discussion

Clinically, rosiglitazone is used as an insulin-sensitizing drug in type 2 diabetes (Lehmann et al. 1995) due to its ability to regulate genes that control cellular energy homeostasis and insulin action in muscle and liver, whereas fenofibrate is used for the treatment of dyslipidemia mainly due to its ability to lower triglyceride levels (Berger and Moller 2002). The role of these agonists of PPARs in insulin resistance has been investigated extensively in diabetic animals (Guerre-Millo et al. 2000; Kramer et al. 2001; Wang et al. 2001) and in patients with type 2 diabetes and insulin resistance syndrome (Mudaliar and Henry 2001; Saltiel and Olefsky 1996). However, a few studies have emphasized the pre-diabetic phase, during which insulin resistance is already in place despite normal blood glucose levels due to the hyper-secretion of insulin by pancreatic β-cells.

In this study, we examined the effect of the specific agonists of PPAR-γ and PPAR-α, rosiglitazone and fenofibrate, respectively, on 14-week-old OLETF rats fed for 2 weeks with either a normal or a high-fat diet. OLETF rats spontaneously develop mild diabetes at 18 weeks of age and 14-week-old OLETF rats are pre-diabetic, but have normal fasting glucose levels (Kawano et al. 1992, 1994). Our results indicate that, in addition to their different effects on body weight and fat mass (Table 1), rosiglitazone and fenofibrate significantly improve the insulin resistance and reduce the levels of triglycerides, free fatty acids, and total cholesterol in the serum of these pre-diabetic animals (Table 2). Importantly, these agonists of PPAR-γ and PPAR-α also significantly reduce the levels of malonyl-CoA, a key regulator for fatty acid oxidation, in skeletal muscle and liver.

In the liver, malonyl-CoA is important both as an intermediate in the de novo synthesis of fatty acids and as an allosteric inhibitor of CPT1, the enzyme that controls the transfer of long-chain fatty acyl-CoAs into the mitochondria for oxidation. In contrast, malonyl-CoA functions mainly as a regulator of CPT1 in skeletal muscle, where de novo synthesis of fatty acids is minimal (Megarry 2002). Similar to what has been observed in obese humans or those who have type 2 diabetes (Bandyopadhyay et al. 2006), we found that the concentrations of malonyl-CoA in skeletal muscle and liver are highly elevated in pre-diabetic rats fed a normal diet, and that this effect is exacerbated by a high-fat diet. Increased malonyl-CoA levels may promote the de novo synthesis of fatty acids in liver and inhibit the oxidation of fatty acids in both liver and skeletal muscle, causing an accumulation of triglycerides and fatty acids that contribute to insulin resistance. Our data suggest that using PPAR-γ and PPAR-α agonists to lower malonyl-CoA levels may be one way to improve insulin resistance in pre-diabetic animals or humans.

Malonyl-CoA is synthesized by acetyl-CoA carboxylase (ACC) (Thampy 1989) and degraded to acetyl-CoA and carbon dioxide by malonyl-CoA decarboxylase (MCD) (Saha et al. 2000). In this study, we characterized the activities of these two enzymes in non-exercising, pre-diabetic OLETF rats. We found that MCD activities in both skeletal muscle and liver were significantly increased after 2 weeks of rosiglitazone and fenofibrate treatment even in rats fed a high-fat diet. In addition, we determined that the profound activation of MCD by fenofibrate that we observed in this study
was due to an increase in MCD mRNA levels. This finding is consistent with our previous report that PPAR-α enhances MCD transcription (Lee et al. 2004). Our data suggest that PPAR-γ and PPAR-α agonists reduce malonyl-CoA levels in pre-diabetic OLEFTY by enhancing MCD transcription.

In contrast to MCD, the effects of PPAR-γ and PPAR-α agonists on ACC activity in liver differs from their effects in skeletal muscle. Although ACC activity was not affected by either rosiglitazone or fenofibrate in skeletal muscle, it was significantly elevated in the livers of rats fed a high-fat diet. This increased ACC activity was reduced to normal levels by rosiglitazone treatment and suppressed even further by fenofibrate.

ACC is a cytosolic enzyme that catalyzes the carboxylation of cytosolic acetyl-CoA to form malonyl-CoA (Saha and Ruderman 2003). Two principal isoforms have been identified: a 265 kDa protein referred to as ACCα or ACC1, which is predominantly expressed in lipogenic tissues such as liver and adipose tissue; and a 275–280 kDa protein, ACCβ or ACC2, which, although it is present to some extent in liver and other tissues (Bianchi et al. 1990; Kim 1997), is the major isoform expressed in skeletal muscle. ACCα and ACCβ are the products of distinct genes and they have different affinities for their substrate, cytosolic acetyl-CoA (Kim 1997). ACCα plays a major role in fatty acid synthesis and ACCβ is important for fatty acid oxidation (Kim 1997; Ruderman et al. 1999).

The fact that a high-fat diet increases ACC activity in the liver but not the skeletal muscles of pre-diabetic OLEFTY rats, and that this increase is abrogated by treatment with PPAR-γ and PPAR-α agonists, suggests that these conditions specifically modulate ACCα activity and its ability to promote fatty acid synthesis. In particular, the dramatic decrease in ACCα activity we observed in response to fenofibrate may explain the significant reduction in body weight and fat mass in OLEFTY rats fed with a high-fat diet.

Expression of ACCα is regulated by multiple hormones and nutrient status mediated by three promoters and multiple transcriptional factors, such as sterol regulatory element binding proteins (SREBP1a and SREBP1c) and carbohydrate response element binding protein (ChREBP), as well as by mRNA splicing (Barber et al. 2005; Brownsey et al. 2006). In addition, its function is regulated through posttranslational modification, including protein maturation, allosteric regulation, multiple sites of phosphorylation, and protein–protein interaction (Brownsey et al. 2006; Tong 2005). Therefore, further studies are required to determine how this PPAR-α agonist inhibits ACCα activity in the liver under these conditions.

AMPK is a heterotrimeric protein kinase complex that acts as an energy sensor, responding to a rise in AMP levels by increasing ATP-generating pathways and reducing ATP-consuming pathways (Hardie et al. 2003; Kahn et al. 2005). AMPK activation inhibits fatty acid and cholesterol synthesis and gluconeogenesis in the liver and stimulates fatty acid uptake and oxidation, glucose uptake, and mitochondrial biogenesis in skeletal muscle (Kahn et al. 2005). It regulates malonyl-CoA levels by inactivating ACC and activating MCD during exercise or the electric stimulation of skeletal muscles (Hardie and Carling 1997; Saha et al. 2000). It has been hypothesized that dysregulation of the AMPK/malonyl-CoA fuel-sensing and signaling network is a key factor in the development of insulin resistance (Ruderman and Prentki 2004). Thiazolidinedione treatment reportedly stimulates AMPK activity in cultured cells, rat liver, and adipose tissues by increasing the ratio of AMP to ATP (Fryer et al. 2002; Saha et al. 2004).

In contrast, we found that AMPK activity levels are barely detectable in pre-diabetic OLEFTY rats, regardless of whether they were fed with a normal or a high-fat diet. Furthermore, these low basal levels were not affected by treatment with either rosiglitazone or fenofibrate. Consistent with this observation, the activity of ACC—a key target of AMPK activation—in skeletal muscle remained essentially unchanged in rats fed either a normal or a high-fat diet. Our results suggest that AMPK/malonyl-CoA signaling network is defective in this pre-diabetic rat model, which may contribute to the dysregulation of fatty acid metabolism in skeletal muscle and liver.

Conclusion

In this study, we demonstrate that rosiglitazone and fenofibrate improve insulin sensitivity and reduce plasma triglyceride and free fatty acid levels in pre-diabetic rats. This effect is likely due to a reduction of malonyl-CoA levels caused by an increase in MCD activity. In particular, fenofibrate stimulates the transcription of MCD by PPAR-α. The fact that ACC and AMPK activities in skeletal muscle were not affected by these agonists suggests a defect in the AMPK signaling network in this diabetic rat model. Further studies will be required to determine how MCD activity is stimulated by rosiglitazone and why fenofibrate inhibits ACCα activity in the livers of rats fed a high-fat diet. Our data strongly suggest that early intervention with PPAR agonists during the pre-diabetic period will significantly improve fatty acid metabolism and insulin sensitivity and, perhaps, prevent further development of diabetes.

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