



# RESEARCH PAPER

# Activation of peroxisome proliferator-activated receptor-α in mice induces expression of the hepatic low-density lipoprotein receptor

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**Background and purpose:** Mutations in the low-density lipoprotein receptor (*LDLR*) gene cause familial hypercholesterolaemia in humans and deletion of the LDLR induces lesion development in mice fed a high-fat diet. LDLR expression is predominantly regulated by sterol regulatory element-binding protein 2 (SREBP2). Fenofibrate, a peroxisome proliferator-activated receptor  $\alpha$  (PPAR $\alpha$ ) ligand, belongs to a drug class used to treat dyslipidaemic patients. We have investigated the effects of fenofibrate on hepatic LDLR expression.

**Experimental approach:** The effects of fenofibrate on hepatic LDLR expression (mRNA and protein) and function were evaluated by both *in vitro* (with AML12 cells) and *in vivo* experiments in mice.

Key results: Fenofibrate increased LDLR expression and LDL binding in a mouse hepatoma cell line, AML12 cells. Fenofibrate restored sterol-inhibited hepatocyte LDLR expression. Mechanistic studies demonstrated that induction of LDLR expression by fenofibrate was dependent on PPAR $\alpha$  and sterol regulatory elements (SRE). Specifically, fenofibrate induced LDLR expression by increasing maturation of SREBP2 and phosphorylation of protein kinase B (Akt) but had no effect on SREBP cleavage-activating protein. *In vivo*, a high-fat diet suppressed LDLR expression in mouse liver while elevating total and LDL cholesterol levels in plasma. However, fenofibrate restored LDLR expression inhibited by high-fat diets in the liver and reduced LDL cholesterol levels in plasma.

**Conclusions and implications:** Our data suggest that fenofibrate increased hepatic LDLR expression in mice by a mechanism involving Akt phosphorylation and LDLR gene transcription mediated by SREBP2.

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Abbreviations: Akt, protein kinase B; LDLR, low-density lipoprotein receptor; PPAR $\alpha$ , peroxisome proliferator-activated receptor  $\alpha$ ; SCAP, SREBP cleavage-activating protein; SRE, sterol regulatory element; SREBP2, sterol regulatory element-binding protein 2

### Introduction

Expression of the low-density lipoprotein receptor (LDLR) is regulated at the transcriptional and/or translational levels. Cholesterol and its derivatives inhibit LDLR expression that results in a reduction of cellular cholesterol uptake. In this feedback regulatory pathway, the sterol regulatory element-binding protein 2 (SREBP2) has a crucial function (Goldstein *et al.*, 2006). SREBPs are transcription factors belonging to the basic helix–loop–helix–leucine zipper (bHLH-Zip) family. They

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consist of three isoforms, namely SREBP-1a, SREBP-1c and SREBP2 (Goldstein *et al.*, 2006). SREBP1 is responsible for fatty acid synthesis and metabolism, whereas SREBP2 controls expression of LDLR and 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase, the rate-limiting enzyme for cholesterol synthesis (Brown and Goldstein, 1997; Shimano, 2001). SREBP 1 and 2 are synthesized in the rough endoplasmic reticulum (ER) as precursor proteins and bind to the SREBP cleavage-activating protein (SCAP). In sterol-depleted cells, SCAP chaperones SREBP from the ER to the Golgi where the SREBP is cleaved by two Golgi residing proteases, site-1 protease (S1P) and site-2 protease (S2P). The released bHLH-Zip domain of SREBP or nuclear form of SREBP (nSREBP) translocates into the nucleus, and then binds to the sterol regulatory element (SRE) in the promoter region of target genes, including the

*LDLR*, and activates transcription. In contrast, cholesterol binds to the sterol-sensing domain of SCAP in lipid-loaded cells and causes a change in SCAP conformation, allowing SCAP to bind to a pair of ER membrane proteins, Insig1 and Insig2. The formation of the SREBP/SCAP/Insig complex prevents the processing of SREBP and inhibits the expression of SREBP target genes (Nohturfft *et al.*, 1999; Yang *et al.*, 2002; Adams *et al.*, 2003; Radhakrishnan *et al.*, 2004).

Peroxisome proliferator-activated receptors (PPARs) including PPARα, PPARβ/δ and PPARγ are ligand-activated transcription factors that regulate lipid metabolism, glucose homoeostasis, inflammation and the development of adipocyte differentiation (Gilde et al., 2006). Both PPARa and PPARγ ligands are anti-atherogenic in mouse models of atherosclerosis (Li et al., 2000; Duez et al., 2002). They have been shown to decrease cardiovascular end points in clinical trials (Helsinki Heart Study, Veterans Affairs High-Density Lipoprotein Cholesterol Intervention Trial (VA-HIT), Fenofibrate Intervention and Event Lowering in Diabetes (FIELD) and Prospective Pioglitazone Clinical Trial in Macrovascular Events (PROactive)) (Dormandy et al., 2005; Keech et al., 2005; Backes et al., 2007). Fibric acid derivatives (such as, fenofibrate) are PPARα ligands used in the treatment of dyslipidaemia in patients with diabetes. They reduce plasma triglyceride levels, increase high-density lipoprotein (HDL) cholesterol levels and shift the distribution of LDL subfractions towards larger, less atherogenic particles (Vu-Dac et al., 1995; Berthou et al., 1996).

Increased hepatic LDLR expression is one of the most effective means to lower plasma cholesterol levels (Bays and Stein, 2003). In some clinical studies, PPAR $\alpha$  agonists have been shown to reduce LDL cholesterol levels, particularly in the presence of hypertriglyceridaemia with unrevealed mechanisms. In patients with hypertriglyceridaemia, LDL cholesterol may increase, but this is due to an increase in particle size but not in particle number (Davidson *et al.*, 2006).

In this study, we hypothesized that fenofibrate, a PPAR $\alpha$  ligand, might reduce plasma LDL cholesterol levels through increased hepatic LDLR expression. We provide studies that show for the first time that fenofibrate increased LDLR expression and uptake of LDL by hepatocytes in a manner dependent on the activation of PPAR $\alpha$ . Our *in vivo* studies further show that administration of fenofibrate in mouse induced liver LDLR expression and reduced plasma LDL cholesterol levels, and that this process occurred though the activation of SREBP2 and phosphorylation of Akt.

## Methods

Cell culture

The AML12 murine hepatocyte cell line (ATCC CRL-2254) (Wu *et al.*, 1994) was purchased from ATCC (Manassas, VA, USA) and cultured in complete growth medium prepared as follows: 1:1 of Dulbecco's modified Eagle's medium (DMEM) and Ham's F12 medium with  $5\,\mu\mathrm{g}\,\mathrm{mL}^{-1}$  insulin,  $5\,\mu\mathrm{g}\,\mathrm{mL}^{-1}$  transferrin,  $5\,\mathrm{ng}\,\mathrm{mL}^{-1}$  selenium,  $40\,\mathrm{ng}\,\mathrm{mL}^{-1}$  dexamethasone, 10% foetal bovine serum (FBS) and  $50\,\mu\mathrm{g}\,\mathrm{mL}^{-1}$  penicillin/ streptomycin. Cells were switched to serum-free medium and received treatments when confluent.

Human kidney 293 cell line was also purchased from ATCC and cultured in complete DMEM medium (10% FBS and  $50 \,\mu\mathrm{g}\,\mathrm{mL}^{-1}$  penicillin/streptomycin).

### Animals

All animal procedures were in strict compliance with NIH guidelines. C57BL wild-type mice (at the age of 10 weeks and about 25 g) were purchased from The Jackson Laboratory (Bar Harbor, ME, USA) and maintained in fully accredited facilities (AAALAC) at Weill Medical College of Cornell University. Mice were divided into three groups at the time of the experiments. Group 1 was left on normal chow, whereas the other two groups were initially fed with a high-fat diet for 2 weeks. They were then further divided as follows: a high-fat diet alone or a high-fat diet plus fenofibrate (50 mg per 100 g food). We assumed that the intake of fenofibrate by mice was about  $50\,\mathrm{mg}\,\mathrm{day}^{-1}$  per kg body weight, based on the following calculation: a 25 g mouse eats 2.5 g food a day, thus the intake of fenofibrate is  $2.5 \times 50/100/0.025 = 50$ . After 10 days of treatment with fenofibrate, mice were killed and blood or liver was individually collected. Plasma was prepared by centrifuging blood samples for 5 min at  $400 \times g$  and was kept at -20 °C for assay of total, LDL and HDL cholesterol levels. The liver was used to extract cellular total RNA.

Isolation of total RNA and northern blot analysis of LDLR, HMG-CoA reductase and SCAP mRNA expression

Total cellular RNA was extracted from cells or liver and used to determine the levels of mouse *LDLR*, *HMGCR* (HMG-CoA reductase), *SCAP* and *GAPDH* (glyceraldehyde 3-phosphate dehydrogenase) mRNA by northern blot analysis as described (Han *et al.*, 2004). Probes were generated by reverse transcription-PCR based on the published sequence. The sequences of primers are *LDLR*-f: GACTGCAAGGACATGA GCGA (781-801); *LDLR*-r: CGGTTGGTGAAGAGCAGATA (1221-1201); *HMGCR*-f: CGTTGGCTGGAGACGGCA (121-138), *HMGCR*-r, CACTGACATGCAGCCGAA (798-781); *SCAP*-f, TTCTGGCAGTCGATGTGTTC (573–592); and *SCAP*-r, GACCA CCAGGTATGGGAAGA (1309-1280).

Western blot analysis of LDLR, SREBP2, PPARa, total Akt and phospho-Akt protein expression

After treatment, cells were washed twice with cold phosphate-buffered saline, then scraped and lysed in ice-cold lysis buffer (50 mm Tris, pH 7.5, 150 mm NaCl, 1% Triton X-100, 1% sodium deoxycholate, 1 mm phenylmethanesulphonyl fluoride, 50 mm sodium fluoride, 1 mm sodium orthovanadate, 50  $\mu g\,mL^{-1}$  aprotinin and 50  $\mu g\,mL^{-1}$  leupeptin). If the lysis buffer was used to extract cellular proteins for total Akt and phospho-Akt analysis, 1 mm  $\beta$ -glycerol phosphate and 2.5 mm sodium pyrophosphate were added. Cellular lysate was centrifuged at  $400\times g$  for 15 min at 4 °C, and supernatants were transferred into a new test tube. After measuring the protein content, lysates were loaded and separated on a 12% SDS-polyacrylamide gel electrophoresis and transferred onto nylon-enhanced nitrocellulose membrane. The

membrane was blocked with a solution of TBS/0.1% Tween 20 (TBS-T) containing 5% fat-free dry milk for 1 h, and then incubated with primary antibody overnight at 4  $^{\circ}\text{C}$  followed by washing for  $3\times10\,\text{min}$  with TBS-T buffer. The blot was re-blocked with TBS/T containing 5% milk, followed by incubation with horseradish peroxidase-conjugated secondary antibody for 1 h at room temperature. After washing  $3\times10\,\text{min}$  with TBS-T, the membrane was incubated for 1 min in a mixture of equal volumes of western blot chemiluminescence reagents 1 and 2, and then exposed to film before development.

Isolation of LDL and analysis of LDL binding to hepatocytes LDL (1.019–1.063 g mL $^{-1}$ ) was isolated from normal human plasma by sequential ultracentrifugation. To conduct the binding of LDL to hepatocytes, LDL was fluorescein conjugated with a reactive succinimidyl-ester of carboxyl-fluorescein by using a labelling kit purchased from Princeton Separations (Adelphia, NJ, USA). Hepatocytes were cultured in a four-well slide chamber. After treatment, cells were washed twice with phosphate-buffered saline and then incubated with 30  $\mu g$  mL $^{-1}$  of labelled LDL in serum-free medium for 2h at 37 °C. After washing twice with phosphate-buffered saline and covering with Vectashield mounting medium (Vector Laboratories Inc., Burlingame, CA, USA), cells were observed with a fluorescent microscope and photographed.

Construction and determination of LDLR promoter activity

A mouse LDLR promoter including the SRE (pLDLR, from -331 to +49, SRE in LDLR (ATCACCCCAT) locates from -213 to -204) was generated by PCR with primers, pLDLR-f, 5'-CGGCCAAGCTTGCTGTGGGAGGAATTTGAGGA-3' and pLDLR-r, 5'-GGCCACCATGGAAGGAGCAGGGCGATGAC-3', and genomic DNA isolated from mouse liver. The primers contained restriction sites for HindIII (5' primer) and NcoI (3' primer), respectively. After the sequence was confirmed, the PCR product was digested by enzymes and then subcloned into pGL4.10 luciferase reporter vector (Promega, Madison, WI, USA) to generate the LDLR promoter (pLDLR-Luc) and amplified. The LDLR promoter with a mutation in SRE (pLDLR-SREmu-Luc) was generated by using a Quik-Change II Site-Directed Mutagenesis Kit (Stratagene, La Jolla, CA, USA), DNA of pLDLR-Luc, and the following primers: pLDLR SREmu-f, 5'-gaagatttttgaaaATCACGGCATtgcagactcct ccccg-3' (CC in SRE motif (in caps) was replaced by GG (underlined)); pLDLR SREmu-r, 5'-ggggaggagtctgcaATGCCG TGATtttcaaaaatcttc-3'.

About 90% confluent 293 cells in 24-well plates were transfected with  $1\,\mu g$  per well of pLDLR-Luc or pLDLR-SREmu-Luc plus Renilla luciferase reporter (RLuc, for internal control) by using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA). After transfection for 8 h, the cells were treated overnight. After washing with phosphate-buffered saline, the cells were lysed and the activity of pLDLR-Luc or pLDLR-SREmu-Luc and RLuc in cellular lysate was determined by using a Dual-Luciferase Reporter Assay System from Promega.

Data analysis

All experiments were repeated three times except as indicated, and representative results are presented. Data generated from the cholesterol assays and the activity assays of LDLR promoters were expressed as mean  $\pm$  s.e.mean and analysed by paired t-test.

### **Materials**

Fenofibrate, GW7647, GW9662, LY294002 and LY303511 were purchased from CalBiochemistry (San Diego, CA, USA) and prepared in dimethyl sulphoxide. Control and PPARa siRNAs, siRNA transfection medium and reagents, and polyclonal anti-PPARα antibody were purchased from Santa Cruz Biotechnology Inc. (Santa Cruz, CA, USA). High-fat diet (21% fat and 0.2% cholesterol) was purchased from Harlan Teklad (Madison, WI, USA). Rabbit anti-mouse LDLR antibody was purchased from R&D Systems (Minneapolis, MN, USA). Rabbit anti-mouse total Akt and phospho-Akt (Ser473) were obtained from Cell Signaling Technology (Beverly, MA, USA). Rabbit anti-aP2 serum was generated by immunization with murine aP2 peptide of DGKSTTIKRKRDGDKLV (Sun et al., 2003). Rabbit-anti mouse-SREBP2-specific polyclonal antibody was a gift of Dr Joseph Goldstein from The University of Southwestern Medical Center. Total, LDL and HDL cholesterol assay kits were purchased from Wako Chemicals (Richmond, VA, USA). All other reagents were obtained from Sigma-Aldrich Chemicals (St Louis, MO, USA) except as indicated.

### Results

Fenofibrate induces LDLR expression in hepatocytes in a PPAR\u00e3-dependent manner

To study the effects of fenofibrate on hepatic LDLR expression, a mouse hepatocyte cell line, AML12 cells, was treated overnight with different concentrations of fenofibrate in serum-free medium. Changes in LDLR mRNA were assessed by northern blot analysis (Figure 1a). Fenofibrate induced hepatocyte LDLR mRNA expression in a concentration-dependent manner. Maximal induction was observed with 25 µM of fenofibrate.

To determine whether the induction of LDLR mRNA by fenofibrate was associated with the induction of LDLR protein, total cellular protein was extracted from cells after treatment with fenofibrate and used to determine LDLR protein expression by western blot analysis. Consistent with the changes in LDLR mRNA expression, LDLR protein was increased by fenofibrate in a concentration-dependent manner, with maximal induction of LDLR protein expression at a concentration of  $10\,\mu\rm M$  (Figure 1a).

The temporal effects of fenofibrate on LDLR expression were assessed by time course studies. Cells were treated with fenofibrate ( $10\,\mu\text{M}$ ) for up to 24 h and changes in LDLR protein were analysed by western blot. Induction of LDLR protein expression by fenofibrate was first observed at 6 h, and maximal expression was observed at 12 h (Figure 1b).

To correlate the induction of LDLR expression in response to fenofibrate with the LDLR function, LDL binding to

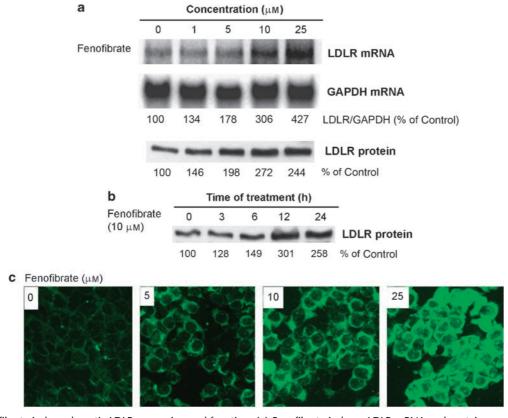


Figure 1 Fenofibrate induces hepatic LDLR expression and function. (a) Fenofibrate induces LDLR mRNA and protein expression: confluent mouse hepatocytes (AML12 cells) in serum-free medium were treated with fenofibrate at the indicated concentrations overnight. Total cellular RNA or proteins were extracted and used to determine the levels of hepatic LDLR mRNA or protein by northern or western blot analysis as described in the Methods section. (b) Fenofibrate induces LDLR protein expression and time course: hepatocytes in serum-free medium were treated with fenofibrate (10 μM) for indicated times. Cellular proteins were extracted and used to determine the levels of LDLR protein by western blot analysis. (c) Fenofibrate induces the binding of LDL to hepatocytes: hepatocytes were treated overnight with fenofibrate at the indicated concentrations. LDL binding assays were performed as described in the Methods section. LDLR, low-density lipoprotein receptor.

AML12 hepatocytes was assessed. Uptake of LDL by hepatocytes was significantly enhanced by fenofibrate in a concentration-dependent manner (Figure 1c).

To study if the induction of hepatic LDLR expression by fenofibrate was dependent on the activation of PPARα, we initially treated hepatocytes with another PPARa agonist, GW7647 (Muoio et al., 2002), and determined its effects on LDLR expression. Similar to fenofibrate, GW7647 induced LDLR expression in a dose-dependent manner (Figure 2a). In contrast, a PPARα antagonist, GW9662 alone (Brown et al., 2001; Leesnitzer et al., 2002), moderately inhibited hepatic LDLR expression. It also blocked fenofibrate-induced hepatic LDLR expression (Figure 2b). Induction of adipose fatty acid-binding protein (A-FABP or aP2) in the liver is also dependent on PPARa activity (Motojima, 2000). Thus, increased aP2 expression in hepatocytes in response to fenofibrate will indicate the activation of PPARa. Indeed, we observed a strong induction of aP2 expression in hepatocytes by fenofibrate (Figure 2c). To further confirm if the induction of hepatic LDLR expression by fenofibrate is PPARα-dependent, we used siRNA to reduce PPARα expression. Results in Figure 2d indicate that PPARα siRNA significantly reduced PPARα levels, whereas a control siRNA had no effect on PPARa. PPARa siRNA alone decreased the expression of hepatocyte LDLR. More importantly, as PPARα siRNA did not completely inhibit PPAR $\alpha$  expression, as compared with PPAR $\alpha$  siRNA alone, the co-treatment of fenofibrate and PPAR $\alpha$  siRNA induced LDLR expression (Figure 2d).

To examine the physiological relevance of fenofibrateinduced hepatic LDLR expression, wild-type mice were fed with normal chow or a high-fat diet or a high-fat diet containing fenofibrate. The effects of fenofibrate on LDLR mRNA expression in the liver, and the levels of total, LDL and HDL cholesterol in plasma were determined. Compared with normal chow, a high-fat diet increased total and LDL cholesterol levels (163 and 139% of control group, respectively, Table 1). However, fenofibrate abolished the elevation of circulating LDL cholesterol. The increased total cholesterol in mice given fenofibrate was partially due to increased HDL cholesterol (Table 1). To study if the reduction of LDL cholesterol in response to fenofibrate was due to the increased LDLR-mediated LDL clearance, the expression of liver LDLR mRNA in mice was assessed by northern blot analysis. As expected, a highfat diet inhibited LDLR mRNA expression in mouse liver (43% of group on normal chow, Figure 3). In contrast, the addition of fenofibrate in the high-fat diet abolished the decline of LDLR mRNA levels (96% of group on normal chow, Figure 3).

Z Huang et al

Fenofibrate restores LDLR expression inhibited by lipid loading LDLR expression is feedback inhibited by cellular sterol levels. Thus, cholesterol or lipid 'loading' will inhibit LDLR

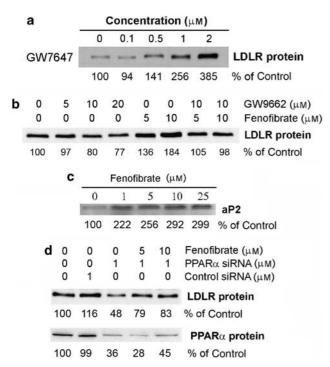


Figure 2 Induction of hepatocyte LDLR expression by fenofibrate is PPARα-dependent. (a) Induction of hepatocyte LDLR expression by GW7647, another PPARα ligand: confluent hepatocytes were treated with another PPARa agonist, GW7647 at the indicated concentrations overnight. The expression of LDLR protein was determined by western blot analysis. (b) PPAR $\alpha$  antagonist, GW9662, blocks fenofibrate-induced LDLR expression: confluent hepatocytes were treated with GW9662 or fenofibrate alone or co-treated with GW9662 and fenofibrate at the indicated concentrations overnight. Changes in LDLR protein was assessed by western blot analysis. (c) Induction of hepatocyte aP2 expression by fenofibrate: confluent hepatocytes were treated with fenofibrate at the indicated concentrations overnight. Expression of aP2 was determined by western blot analysis. (d) PPARa siRNA inhibits LDLR expression: confluent hepatic cells in six-well plates were transfected with control siRNA or PPARα siRNA for 6 h followed by treatment with fenofibrate at the indicated concentrations overnight. Expression of LDLR and PPARα proteins were determined by western blot analysis. LDLR, low-density lipoprotein receptor; PPARa, peroxisome proliferatoractivated receptor  $\alpha$ .

**Table 1** Effect of fenofibrate on mouse plasma cholesterol levels  $(mq L^{-1})$ 

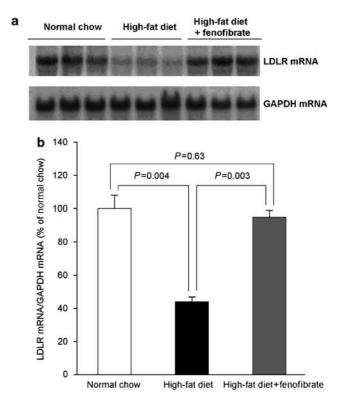
Treatment	Total C	LDL C	HDL C
Control (normal chow)	782 ± 90	196 ± 17	465 ± 50
High-fat diet	1277 ± 38*	271 ± 24*	749 ± 46*
High-fat diet + fenofibrate	1508 ± 48*,#	211 ± 13 <sup>#</sup>	997 ± 45*,#

Abbreviations: C, cholesterol; HDL, high-density lipoprotein; LDL, low-density lipoprotein.

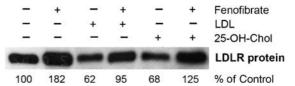
Mice at the age of 10 weeks were divided into three groups. Control group was fed with normal chow. Other two groups were fed a high-fat diet for 2 weeks followed by drug treatment as described in the Methods section. At the end of experiments, plasma and liver were collected from each mouse. Total, LDL and HDL cholesterol levels were determined.

\*Significantly different from control group at P < 0.05; #significantly different from group of high-fat diet alone at P < 0.05, by Student's t-test (n = 5).

expression. This mechanism not only prevents cells from uncontrolled accumulation of lipids but also reduces the clearance of excess LDL cholesterol from the circulation by the liver during hyperlipidaemia. To determine whether fenofibrate would restore LDLR expression suppressed by lipids, cells were treated with LDL or 25-hydroxycholesterol or lipid plus fenofibrate. LDL and 25-hydroxycholesterol inhibited LDLR protein expression in AML12 hepatocytes (Figure 4). This inhibitory effect was abolished in the presence of fenofibrate, suggesting that fenofibrate restores the LDLR expression inhibited by sterols (Figure 4).



**Figure 3** Fenofibrate induces LDLR expression in mouse liver. (a) Mice were treated as described in Table 1. Total RNA was extracted from the mouse liver, and the expression of LDLR mRNA was determined by northern blot analysis. The same blot was re-hybridized with  $^{32}$ P-labeled GAPDH probe for internal control. (b) Summary of northern blot results indicating the significant difference (P<0.05, n=3) between groups of normal chow and high-fat diet, or groups of high-fat diet and high-fat diet+fenofibrate, but not between groups of normal chow and high-fat diet+fenofibrate. GAPDH, glyceraldehyde 3-phosphate dehydrogenase; LDLR, low-density lipoprotein receptor.

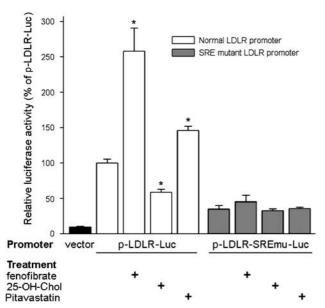


**Figure 4** Fenofibrate restores sterol-inhibited LDLR expression. Hepatocytes were treated with fenofibrate ( $10\,\mu\text{M}$ ) in the presence or absence of LDL ( $100\,\mu\text{g mL}^{-1}$ ) or 25-hydroxycholesterol (25-OH-Chol,  $10\,\mu\text{M}$ ) overnight. Expression of LDLR protein was determined by western blot analysis. LDLR, low-density lipoprotein receptor.

Induction of LDLR expression by fenofibrate occurs through the activation of SREBP2 and Akt phosphorylation

LDLR expression is primarily regulated by the activity of SREBP2, which is dependent on cellular sterol levels. Fenofibrate restored sterol-suppressed LDLR expression, suggesting that fenofibrate may modulate the SREBP2 pathway to activate hepatic LDLR expression. Initially, we determined whether the induction of LDLR expression by fenofibrate was SRE-dependent. The kidney cell line, 293 cells, was transfected with LDLR promoter-luciferase constructs, including normal SRE or mutant SRE, followed by treatment with fenofibrate or 25-hydroxycholesterol or pitavastatin, a HMG-CoA reductase inhibitor. The activity of the promoter containing the normal SRE sequence was increased by pitavastatin (47%) and decreased by 25-hydroxycholesterol (42%), as expected. Fenofibrate treatment significantly increased the activity of the LDLR promoter containing normal SRE (>2-fold) (Figure 5). Mutation in the SRE reduced promoter activity and disabled the regulation of promoter activity by pitavastatin and 25hydroxycholesterol. Fenofibrate did not influence the activity of the SRE mutant LDLR promoter, also suggesting an essential role of SRE in the regulation of LDLR transcription in response to fenofibrate.

SRE-dependent induction of LDLR promoter activity by fenofibrate indicated that fenofibrate may activate SREBP2. Thus, expression and maturation of SREBP2 in response to

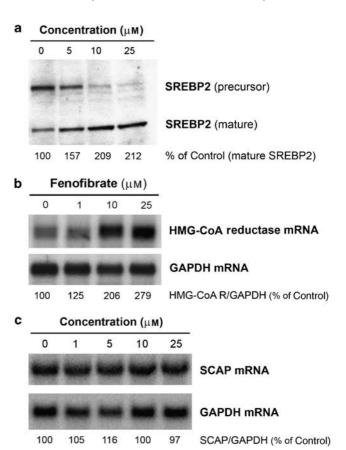


**Figure 5** Induction of hepatic LDLR expression by fenofibrate is sterol regulatory elements (SRE)-dependent. Confluent 293 kidney cells were transfected with plasmid DNA of normal LDLR promoter (pLDLR-luc) or SRE mutant LDLR promoter (pLDLR-SREmu-Luc) plus *Renilla* luciferase reporter (*R*Luc) followed by treatment with fenofibrate (10  $\mu$ M) or 25-hydroxycholesterol (25-OH-Chol, 10  $\mu$ M) or pitavastatin (10  $\mu$ M) overnight in serum-free medium. Promoter activity was analysed as described in the Methods section. The relative luciferase activity of the LDLR promoter was calculated based on the activity of co-transfected *R*luc. The activity of normal LDLR promoter without treatment was defined as 100%. \*Significantly different from normal LDLR promoter without treatment at *P* < 0.05 by Student's *t*-test (*n* = 4). LDLR, low-density lipoprotein receptor.

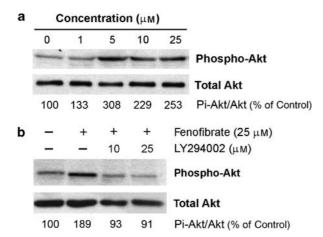
fenofibrate was determined by western blot analysis. Results shown in Figure 6a demonstrate that fenofibrate did not induce SREBP2 expression. However, fenofibrate increased the amount of mature SREBP2 and decreased SREBP2 precursor, suggesting that fenofibrate enhanced the proteolysis of SREBP2. To further confirm the activation of SREBP2 by fenofibrate, we evaluated the expression of HMG-CoA reductase mRNA, another target gene regulated by SREBP2 activity. HMG-CoA reductase mRNA was also increased in response to fenofibrate (Figure 6b).

Levels of SCAP or translocation of the SCAP–SREBP2 complex from the ER to the Golgi affect the maturation of SREBP2. To test if fenofibrate was able to induce SCAP expression, its effect on SCAP mRNA expression was assessed by northern blot analysis. Results shown in Figure 6c indicate that SCAP expression was unaffected by fenofibrate treatment.

The activity of the protein kinase Akt influences the SREBP2 processing and expression of the LDLR (Du *et al.*, 2006). To study if the activation of SREBP2 by fenofibrate



**Figure 6** Fenofibrate induces maturation of sterol regulatory element-binding protein 2 (SREBP2) and expression of 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase, but has no effect on the expression of SREBP cleavage-activating protein (SCAP). (a) Hepatocytes were treated with fenofibrate at the indicated concentrations overnight. Total cellular proteins were extracted and used to determine the levels of precursor of SREBP2 (P) and mature SREBP2 (N) by western blot analysis. Hepatocytes in serum-free medium were treated with fenofibrate at the indicated concentrations overnight. Total cellular RNA was collected and used to determine the expression of HMG-CoA reductase mRNA (b) and SCAP mRNA (c), respectively, by northern blot analysis.



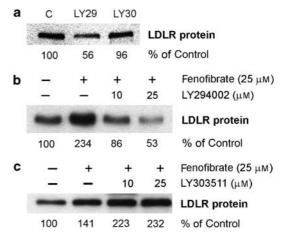
**Figure 7** Fenofibrate increases Akt phosphorylation. Hepatocytes were treated with fenofibrate at the indicated concentrations overnight (a), or treated with fenofibrate ( $25\,\mu\text{M}$ ) in the presence or absence of LY294002 at the indicated concentrations overnight (b). Cellular protein was extracted and used to determine total Akt and phospho-Akt expressions by western blot analysis as described in the Methods section.

was initially related to Akt phosphorylation, total Akt and phosphorylated Akt in response to fenofibrate treatment was assessed by western blot analysis. Figure 7a demonstrates that fenofibrate increased phosphorylated Akt but had no effect on total Akt expression. Furthermore, LY294002, a phosphoinositide 3-kinase (PI3K, the primary activator of Akt phosphorylation) inhibitor (Song *et al.*, 2005), blocked the induction of Akt phosphorylation by fenofibrate (Figure 7b).

To correlate changes in Akt activity with expression of the LDLR in response to fenofibrate, we treated hepatacytes with a PI3K inhibitor, LY294002, or a negative control for the PI3K inhibitor, LY303511. Results in Figure 8a demonstrate that the PI3K inhibitor reduced LDLR expression. The negative control for the PI3K inhibitor had little effect on LDLR expression. We then evaluated whether the PI3K inhibitor was able to block fenofibrate-induced LDLR expression. Consistent with changes in phospho-Akt, the induction of LDLR expression by fenofibrate was blocked by a PI3K inhibitor, LY294002 (Figures 8b), whereas the negative control for PI3K inhibitor (LY303511) slightly enhanced fenofibrate-induced LDLR expression (Figure 8c).

### Discussion

Clearance of excess plasma LDL cholesterol occurs through binding of LDL to the hepatic LDLR. Thus, increased LDLR expression in the liver can reduce plasma LDL cholesterol. Statins, the most commonly used therapeutic agents to lower plasma LDL cholesterol, act to increase liver LDLR expression by inhibiting cholesterol synthesis in hepatocytes. Herein, we demonstrate for the first time that fenofibrate induces the expression of LDLR mRNA and protein in a mouse hepatocyte cell line (AML12 cells) and that the induction of LDLR expression enhances LDL uptake. The induction of hepatic LDLR expression is not a



**Figure 8** Induction of hepatic low-density lipoprotein receptor (LDLR) expression by fenofibrate is blocked by phosphoinositide 3-kinase (PI3K) inhibition. (a) Confluent hepatocytes were treated with the PI3K inhibitor (LY294002, LY29,  $10\,\mu\text{M}$ ) or the negative control for the PI3K inhibitor (LY303511, LY30,  $10\,\mu\text{M}$ ) overnight. (b) Cells were treated with fenofibrate (25  $\mu\text{M}$ ) in the absence or presence of PI3K inhibitor (LY294002) at the indicated concentrations overnight. (c) Cells were treated with fenofibrate (25  $\mu\text{M}$ ) in the absence or presence of the negative control for the PI3K inhibitor (LY303511) at the indicated concentrations overnight. Changes in LDLR protein were assessed by western blot analysis.

fenofibrate-specific effect but rather a PPAR $\alpha$  ligand-dependent effect (that is, dependent on the activation of PPAR $\alpha$ ). Administration of fenofibrate induced expression of the LDLR in mouse liver reducing the plasma levels of LDL cholesterol. The increase in hepatic LDLR expression by fenofibrate is sustainable and dependent on the SRE and the activity of SREBP2, the primary transcriptional regulator of LDLR expression. Moreover, our data show that increased phosphorylation of Akt results in enhanced SREBP2 processing.

Induction of hepatic LDLR expression by fenofibrate was SRE-dependent (Figure 5). Suppression of hepatic LDLR expression by lipid treatment in vitro or by feeding mice a high-fat diet is also dependent on the action of lipids on SRE in the LDLR promoter (Brown and Goldstein, 1997). Therefore, fenofibrate was able to restore sterol-inhibited hepatic LDLR expression both *in vitro* and *in vivo* (Figures 3 and 4). The SRE-dependent induction of hepatic LDLR expression by fenofibrate suggests that the activity of SREBP2 is required. Indeed, we observed that fenofibrate reduced the precursor of SREBP2 but increased mature SREBP2, indicating that the proteolysis of SREBP2 is enhanced. In addition, we observed that fenofibrate increased the expression of HMG-CoA reductase, another target gene of SREBP2. SCAP is the molecule chaperoning SREBP2 from the ER to the Golgi for SREBP2 processing. Interestingly, we observed that the expression of SCAP was not altered in fenofibrate-treated AML12 hepatocytes. Thus, the enhancement of ER-to-Golgi transport of SCAP-SREBP2 most likely occurs by a different mechanism.

Protein kinase B (Akt) is a critical regulator of cell growth, proliferation and survival, and is itself activated by PI3K (Song *et al.*, 2005). Akt activity is also involved in

SREBP2-mediated gene transcription. Inhibition of PI3K by a specific inhibitor, LY294002, reduced SREBP2 processing from its precursor to a mature form, thereby inhibiting the expression of SREBP2 target genes, including the LDLR (Du et al., 2006). Induction of SREBP maturation by sterol depletion (for example, statin treatment) or by growth factors (for example, epidermal growth factor, vascular endothelial growth factor and insulin/insulin-like growth factor-1) in different cell types can involve the activation of Akt phosphorylation. This process can also be blocked by PI3K-specific inhibitors (Borradaile et al., 2003; Zhou et al., 2004; Porstmann et al., 2005). Inhibition of Akt phosphorylation does not affect the binding of SCAP-Insig1, but disrupts the transport of SCAP-SREBP2 from the ER to the Golgi (Du et al., 2006). We interpret our studies to suggest that Akt activity can play a critical role in fenofibrate-induced hepatic LDLR expression, as fenofibrate induced phoshorylation of Akt (Figure 7). Fenofibrateinduced Akt activity, as well as hepatic LDLR expression, was blocked by a PI3K inhibitor (Figures 7 and 8). Taken together, our data demonstrate that fenofibrate can induce hepatic LDLR expression by the activation of Akt and consequently the maturation of SREBP2.

The effects of fenofibrate on serum total and LDL cholesterol levels have been well examined in patients. Although it did not greatly lower serum total and LDL cholesterol levels (~10%) in patients with isolated hypercholesterolaemia (LDL cholesterol>1300 mg L $^{-1}$  but triglyceride levels are normal; Wierzbicki, 2006), fenofibrate did significantly reduce serum total and LDL cholesterol levels in patients with combined hypercholesterolaemia (LDL cholesterol >  $1300 \,\mathrm{mg}\,\mathrm{L}^{-1}$ , triglycerides >  $2000 \,\mathrm{mg}\,\mathrm{L}^{-1}$ and  $< 10000 \,\mathrm{mg} \,\mathrm{L}^{-1}$ ), either as a monotherapy or a combined therapy with a statin (Frost et al., 2001; Malik et al., 2001; Athyros et al., 2002; Melenovsky et al., 2002; Sebestjen et al., 2002; Vega et al., 2003; Wang et al., 2003; Derosa et al., 2004; Winkler et al., 2004; Saklamaz et al., 2005; Grundy et al., 2005; Iovine et al., 2006; Muhlestein et al., 2006; Arca et al., 2007; Chatley et al., 2007). In these studies, fenofibrate alone at a dose about 200 mg day<sup>-1</sup> decreased total cholesterol by 10-20% in combined hypercholesterolaemic patients after a few months (1–6) treatment. A similar decrease was observed in serum LDL cholesterol. In a similar period of treatment, statin therapy alone reduced total and LDL cholesterol by a greater degree than fenofibrate (that is, 20-30% by cerivastatin and simvastatin, and 30-40% by atorvastatin). Statins have little effect on HDL and only a moderate effect on triglycerides. In contrast, fenofibrate can increase HDL cholesterol and apolipoprotein AI/II levels, and significantly decrease triglycerides. LDL consists of a heterogeneous group of particles. Among these particles, the small or dense subfractions of LDL are believed the most atherogenic. Statins decrease all subfractions of LDL to a similar degree. In contrast, fenofibrate can decrease the small or dense LDL more than the large or light LDL. Thus, statins and fenofibrate have different impacts on the composition of LDL in serum (Frost et al., 2001; Vega et al., 2003; Winkler et al., 2004; Arca et al., 2007). For example, atorvastatin decreased the dense LDL (dLDL) apoB in total LDL apoB by 5%, whereas fenofibrate reduced it by 36% (Winkler et al., 2004). Simvastatin alone had little effect on LDL large/small ratio (1.12 vs 0.94 by placebo). In combination with fenofibrate, simvastatin increased this ratio to 2.37 (Vega et al., 2003). More importantly, a combination therapy of fenofibrate and a statin improved lipid profiles in patients with combined hypercholesterolaemia in a synergistic manner and lowered the dose of statin, potentially reducing side effects (Athyros et al., 2002; Vega et al., 2003; Derosa et al., 2004; Grundy et al., 2005; Muhlestein et al., 2006). Fenofibrate also demonstrated several beneficial effects on non-lipid biochemical risk factors, such as C-reactive protein, Lp(a) and oxidation of LDL (Athyros et al., 2002; Melenovsky et al., 2002; Wang et al., 2003). Combined hypercholesterolaemia is very often observed in type II diabetic patients. These studies suggest the application of fenofibrate alone or in combination with a statin in the treatment of lipid disorders in these patients.

Effects of fibrates on cholesterol homoeostatsis were also investigated in different animal models. Administration of fenofibrate to rats reduced plasma cholesterol levels associated with increased LDLR expression in the liver (Staels et al., 1992). In contrast, fenofibrate induced expression of cholesterol synthesis genes in mice (Le Jossic-Corcos et al., 2004). Administration of fenofibrate to mice increased the expression of HMG-CoA reductase and farnesyl pyrophosphate synthases, although the effect of fenofibrate on plasma cholesterol levels and expression of LDLR in the liver was not examined in this report. Induction of HMG-CoA reductase and farnesyl pyrophosphate synthase expression by fibrates was PPARα-dependent. Interestingly, fenofibrate induced expression of SREBP1 but not SREBP2 at mRNA levels in mouse liver (Le Jossic-Corcos et al., 2004). Similarly, we observed that fenofibrate enhanced the proteolysis, but not the expression of SREBP2 protein (Figure 6a). Effect of fibrates on the expression of SREBP target genes including LDLR may depend on the ligand and the duration of treatment. WY14643 is a more selective PPARα ligand but had less inductive effect than fenofibrate on HMG-CoA reductase in mouse liver (Le Jossic-Corcos et al., 2004). Fenofibrate induced LDLR expression in rat liver only after 3 days of administration, whereas WY14643 had no effect on LDLR expression in mouse liver after the same duration of treatment (Staels et al., 1992; Anderson et al., 2004). Gemfibrozil induced LDLR expression, whereas clofibrate had no effect on LDLR expression in rat liver. In contrast to the results in clinical studies, administration of fibrates to rats reduced apoA-I levels in plasma and suppressed expression of apoA genes in the liver (Staels et al., 1992). In a recent report, fenofibrate decreased plasma LDL cholesterol levels in human apolipoprotein (apo) E2 knockin mice (Hennuyer et al., 2005). Proprotein convertase subtilisin/kexin type 9 (PCSK9) is a natural inhibitor of LDLR expression. Recently, fenofibrate has been reported to suppress the expression of PCSK9 in human hepatocytes. This may be another potential mechanism by which fenofibrate induces hepatic LDLR expression (Kourimate et al., 2008).

In summary, our studies demonstrate for the first time that fenofibrate increased hepatic LDLR expression in a PPAR $\alpha$ -dependent manner and reduced plasma LDL cholesterol

levels in mice. Activation of SREBP2 and Akt phosphorylation was required for fenofibrate-induced hepatic LDLR expression. Whether a similar PPAR $\alpha$ -dependent mechanism operates in humans remains to be determined.

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### Conflict of interest

The authors state no conflict of interest.

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