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Modulation of ammonium perfluorooctanoate-induced hepatic damage by genetically different PPAR α in mice

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Abstract

Perfluorooctanoic acid is a ligand for peroxisome proliferator-activated receptor (PPAR α). Ammonium perfluorooctanoate (APFO) at 0.1 and 0.3 mg/kg doses activated mouse PPAR α , but not human PPAR α . This study aimed to clarify whether milligram-order APFO can activate human PPAR α , and the receptor is involved in APFO-induced chronic hepatic damage. Male Sv/129 wild-type (mPPAR α), *Ppara*-null, and humanized PPAR α (hPPAR α) mice (8 weeks old) were divided into three groups. The first was treated with water and the other two with 1.0 and 5.0 mg/kg APFO for 6 weeks, orally, respectively. Both doses activated mouse and human PPAR α to a similar or lower degree in the latter. APFO dose dependently increased hepatic triglyceride levels in *Ppara*-null and hPPAR α mice, but conversely decreased those in mPPAR α ones. APFO-induced hepatic damage differed markedly among the three genotyped groups: single-cell necrosis was observed in all genotyped mice; inflammatory cells and macrovesicular steatosis only in *Ppara*-null mice; and microvesicular steatosis and hydropic degenerations in hPPAR α and *Ppara*-null mice. The molecular mechanism underlying these differences may be attributable to those of gene expressions involved in lipid homeostasis (PPAR α , β - and ω -oxidation enzymes, and diacylglycerol acyl-transferases) and uncoupling protein 2. Thus, milligram-order APFO activated both mouse and human PPAR α in a different manner, which may reflect histopathologically different types of hepatic damage.

Keywords

Hepatic damage; Human; Mouse; Perfluorooctanoic acid; Peroxisome proliferator-activated receptor

Introduction

Perfluorooctanoic acid (PFOA) (CAS, 335-67-1), an organofluoro compound, is used in industrial surfactants, emulsifiers, and numerous consumer products (Butenhoff et al. 2006). Because the biological half-life in humans is reported as 3.5–4.4 years (Olsen et al. 2007),

PFOA will probably be added to the list of Persistent Organic Pollutants in the near future (World Wildlife Fund 2005).

A variety of toxicities associated with PFOA exposure have been investigated and revealed by many studies using ammonium perfluorooctanoate (APFO). Since PFOA is hardly excreted from the body (Kennedy et al. 2004) and is accumulated mostly in the liver (Lau et al. 2007), many studies have focused on the risk of hepatic damage, such as peroxisome proliferation (Nakamura et al. 2009), hepato-cyte necrosis (Butenhoff et al. 2002), hepatocellular adenoma (Biegel et al. 2001), and hepatobiliary injury (Minata et al. 2010). Recently, though Minata et al. (2010) showed that APFO caused cholestasis, this finding was seen mainly in peroxisome proliferator-activated receptor α (*Ppara*)-null mice, not *mPPAR α* mice. Wolf et al. (2008a) reported that lipid droplets were characteristically observed in the livers of *Ppara*-null mice, but not in *mPPAR α* mice. Thus, these results suggest that *PPAR α* may play an important role in the pathogenesis of APFO-induced hepatosteatosis and cholestasis. However, its precise mechanism has not yet been fully understood.

Epidemiologically, PFOA may influence lipid metabolism: Olsen et al. (2003a) reported a positive association between PFOA and serum total cholesterol (TC) and triglycerides (TG). In contrast, no such association was reported in another factory (Olsen et al. 2003b). Therefore, it is very important to clarify whether PFOA influences the lipid metabolism using experimental animals.

Since PFOA is an agonist for *PPAR α* (Ikeda et al. 1985), its activation enhances the activities of peroxisomal and mitochondrial β -oxidation enzymes for fatty acids and inhibits the secretion of very low-density lipoproteins and cholesterol from the liver, as well as reducing total cholesterol and TG in serum and the accumulation of lipids in the liver (Berthiaume and Wallace 2002). However, the functional activation is thought to differ among species. Additionally, constitutive expression of *PPAR α* is quite different between mice or rats and humans, whose expression is thought to be 1/10 of the experimental animals (Palmer et al. 1998). *PPAR α* -humanized (*PPAR α ^{Tet-OFF}*) mice that expressed human *PPAR α* only in the liver of *Ppara*-null background mice have been established (Cheung et al. 2004). This mouse model has been recognized as a useful tool in determining the human *PPAR α* function. As for the effects of Wy-14,643 on hepatic peroxisomal and mitochondrial β -oxidation enzymes, there were few differences in the inductions between *mPPAR α* and *hPPAR α* mice. Ramdhan et al. (2010) reported that one of the trichloroethylene metabolites, trichloroacetic acid, activated not only mouse *PPAR α* but also human *PPAR α* , though the exposure concentration of trichloroethylene was 1,000 and 2,000 ppm, respectively. However, the fact that expressions of human *PPAR α* mRNA and protein are higher in *hPPAR α* mice than in those of *mPPAR α* (Nakamura et al. 2009) may suggest a weaker function of human *PPAR α* compared with mouse *PPAR α* . Indeed, although microgram-order APFO was unable to activate human *PPAR α* , it did activate mouse *PPAR α* (Nakamura et al. 2009). Therefore, it is very important to clarify whether APFO that is higher than that in a previous study (Nakamura et al. 2009) can activate human *PPAR α* and also to determine how the species difference in the function is involved in PFOA-induced hepatic damage when we extrapolate from animal to human data.

Additionally, PFOA is also found to be an agonist for PPAR γ (Vanden Heuvel et al. 2006), which has anti-inflammatory power (Jiang et al. 1998), even though contrary opinions have been reported (Chawla et al. 2001). This receptor is also accepted as a master transcriptional regulator of lipid and glucose metabolism (Spiegelman 1998).

In this study, we compared the effects of relatively high dosages of APFO (0, 1.0, and 5.0 mg/kg) on the PPAR α and the target gene expressions as well as the involvement of this receptor in hepatic damage using wild-type, *Ppara*-null, and *hPPAR α* mice. The molecular mechanisms were also clarified by analyzing the mRNA and protein expressions of related genes. Although relatively low-dose APFO could not activate human PPAR α (Nakamura et al. 2009), higher doses clearly activated the receptor. Our results also suggest that the species difference in the function may determine the characteristic features of hepatic damage caused by PFOA.

Materials and methods

Experimental animals

This study was conducted according to the Guidelines for Animal Experiments of the Nagoya University Animal Center. Three genotyped male and female mice, i.e., wild-type (*mPPAR α*), *Ppara*-null (Lee et al. 1995), and *hPPAR α ^{Tet-OFF}* (*hPPAR α*) (Cheung et al. 2004) mice with an Sv/129 genetic background were bred and reared as described elsewhere (Nakamura et al. 2009). All mice were housed in a temperature- and light-controlled environment (25°C, 12 h light/dark cycle) and maintained on stock rodent chow (Nippon Clea, Tokyo, Japan) and tap water ad libitum. At 8 weeks old, the offspring male mice ($n = 8-10$) of each strain were assigned to the following treatment groups: treated with distilled water daily for 6 weeks by gavage (control group); treated with 1.0 or 5.0 mg/kg APFO (Tokyo Kasei Kagaku, Tokyo, Japan), respectively, for 6 weeks by gavages (Table 1). Since 0.1 and 0.3 mg/kg APFO activated mouse PPAR α , but not human PPAR α (Nakamura et al. 2009), about tenfold doses were selected in this experiment. Since we planned to investigate not only PPAR α -directed hepatic damage but also reproductive toxicity of PFOA, we selected six-week exposure to the chemicals. The results of reproductive toxicity will be reported elsewhere. Macroscopically, there was no abnormal sign in all animals throughout the treatments. On the day following the last dose (18–20 h later), all mice were killed by decapitation, and the blood and livers were removed. A part of each liver was fixed by 10% buffered formalin. The remaining liver samples were snap frozen in liquid nitrogen and stored at -80°C until used. Plasma was collected after centrifuging blood at 3,500g for 10 min and stored at -80°C until used. The numbers of samples except for the histopathological analyses used were indicated in the Tables and Figure legends, and all measurements were performed in duplicate or triplicate.

Analysis of protein concentrations

Each tissue was homogenized with a threefold volume of 10 mM phosphate buffer (pH 7.4) containing 0.25 M sucrose. Protein concentrations of the homogenate and nuclear fraction samples were measured using a Protein Assay Kit (Bio-Rad, Tokyo, Japan).

Western blotting

A nuclear fraction was extracted from a part of the frozen liver using a CellLytic™ NuCLEAR™ Extraction Kit (SIGMA, Tokyo, Japan). The nuclear fractions (NF κ B p65, p50, p52, and PPAR α) and liver homogenates (other proteins) for electrophoresis were subjected to 10 or 12.5% polyacrylamide gel electrophoresis and transferred to nitrocellulose membranes. Immunostaining was conducted as described elsewhere (Nakajima et al. 2000).

Real-time quantitative PCR

Total RNA was isolated from the liver using the RNeasy Mini Kit (QIAGEN, Tokyo, Japan). Real-time quantitative PCR analysis was performed as described elsewhere (Nakamura et al. 2009; Ramdhan et al. 2010).

Lipid concentrations in plasma and liver

Lipid from livers was extracted using the method of Folch et al. (1957). TG and TC in the liver and plasma were measured using TG-IE and T-Cho E kits (Wako, Osaka, Japan), respectively.

Histopathological analysis

Small blocks of liver tissues from each mouse (five animals randomly selected from each group) fixed in 10% neutral buffered formalin were embedded in paraffin and sliced into 4- μ m sections. Tissue sections of the livers were stained with hematoxylin and eosin (H & E) and examined under a BZ-8000 (Keyence, Osaka, Japan) light microscope. Severities of steatosis, lobular inflammation, and hepatocyte degeneration were scored by a pathologist in a blinded fashion referring to the methods of Brunt et al. (1999) and Ramdhan et al. (2010) with the following minor modifications: (1) grade of steatosis: 0, none (0–5% of parenchymal involvement by steatosis); 1, mild (5–33% of parenchymal involvement by steatosis); 2, moderate (33–66%); 3, severe (>66%); (2) grade of lobular inflammation: 0, none; 1, mild; 2, moderate; 3, severe; (3) single-cell necrosis and hepatocyte hydropic degeneration: 0, absent; 1, present; 2, frequent.

Alanine aminotransferase measurements

Plasma ALT activities were measured using a Transaminase C II Test kit purchased from Wako (Osaka, Japan).

Statistical analysis

The Steel-Dwass method in case of pathological scoring and Tukey-Kramer HSD post hoc test in the other cases were conducted to compare the effects of treatment among each treated group of each genotyped mouse, and also among the control groups of mPPAR α , Ppar α -null, and hPPAR α mice. Values of $P < 0.05$ were considered to indicate statistical significance.

Results

Body and liver weight

No significant differences were observed in body weight before and after APFO treatments among the control groups of *mPPAR α* , *Ppara*-null, and *hPPAR α* mice (Table 1). APFO treatments did not induce an increase in body weight in any genotyped mice, whereas they did increase the liver weight and the ratio of liver per body in all genotyped mice in a dose-dependent manner. In the 1.0 mg/kg treatment group, the ratio increases were most prominent in *mPPAR α* mice (1.9-fold), while in the 5.0 mg/kg dose group, they were most prominent in *Ppara*-null mice (2.9-fold).

Histopathological evaluation

Apparent macrovesicular and microvesicular steatosis were not observed in the liver of control *Ppara*-null mice (Fig. 1b, Table 2). APFO induced macrovesicular steatosis only in the liver of *Ppara*-null mice in dose-dependent fashion, while it induced microvesicular steatosis in both *Ppara*-null as well as *hPPAR α* mice; the degree of these fat accumulations was not influenced by APFO dosages (Fig. 1, Table 2). High-dose APFO (5.0 mg/kg) induced lobular inflammation only in *Ppara*-null mice. Although high-dose APFO induced single-cell necrosis in all genotyped mice, the severity appeared to be greater in *mPPAR α* mice than in *Ppara*-null and *hPPAR α* mouse lines. Interestingly, high-dose APFO characteristically induced hydropic degeneration of hepatocytes in *Ppara*-null (arrows in Fig. 1k) and *hPPAR α* mice (Fig. 1i), but not in *mPPAR α* mice. Hypertrophied hepatocytes with prominent eosinophilic cytoplasm, which sometimes appear in the livers of mice treated with potent *PPAR α* activators, were detected in APFO-treated *mPPAR α* (Fig. 1g, arrowheads in Fig. 1j) and *hPPAR α* mice (Fig. 1i). We might note that hepatocholangiolar proliferation was only fleetingly observed in *Ppara*-null mice exposed to 5.0 mg/kg PFOA, but obvious cholestasis, evidenced by intracellular bile droplets and a bile plug, was not observed in the livers of any genotyped mice.

Plasma ALT levels

High-dose APFO increased plasma ALT activities in all genotyped mouse lines, though the elevations were very small, and no differences were noted in these increases among three genotyped mice (*mPPAR α* mice, 2.3-fold vs control; *Ppara*-null, 1.3-fold; *hPPAR α* mice, 1.4-fold) (Table 1).

Plasma and hepatic TG and TC levels

APFO treatment dose dependently decreased plasma TG levels in *mPPAR α* mice (0.70-fold at 1.0 mg/kg and 0.50-fold at 5.0 mg/kg), while not influencing the levels in *Ppara*-null and *hPPAR α* mice. In contrast, APFO did not influence hepatic TG levels in *mPPAR α* mice, though 1.0 mg APFO slightly increased those levels. APFO treatments significantly increased hepatic TG levels in *Ppara*-null mice (3.2-fold and 5.3-fold at 1.0 and 5.0 mg/kg, respectively), but only slightly increased them in *hPPAR α* mice at 1.0 and 5.0 mg/kg (2.0-fold and 3.0-fold, respectively). Consistent with histopathological findings, hepatic TG

contents in the control *Ppara*-null mice were similar to those in the control *mPpara* mice and *hPPAR α* mice.

High-dose APFO (5.0 mg/kg) treatment dose dependently decreased plasma TC levels in *mPPAR α* and *Ppara*-null mice, but not in *hPPAR α* mice. In contrast, PFOA increased hepatic TC levels in *Ppara*-null mice, but not in *mPPAR α* and *hPPAR α* mice.

Analysis of mRNA levels

Because APFO influenced plasma and hepatic TG levels differently in three genotyped mice, we investigated hepatic β -oxidation enzymes that are PPAR α -target genes and are involved in fatty acid metabolism. PPAR α mRNA expression was significantly greater in *hPPAR α* mice than in *mPPAR α* mice, results similar to those of Nakamura et al. (2009) (Fig. 2). Constitutive expressions of CYP4A10, PT, and VLCAD mRNA were significantly lower in *Ppara*-null mice than in *mPPAR α* and *hPPAR α* mice, while those of PH were lower only in *hPPAR α* mice.

Although APFO treatment did not influence the PPAR α -mRNA levels in either *mPPAR α* or *hPPAR α* mice, increased expressions of hepatic PPAR α -target genes, CYP4A10, PT, PH, MCAD, VLCAD, and proliferation cell nuclear antigen (PCNA) were observed in both mouse lines. In general, their increases were the same (VLCAD, MCAD, and PCNA) or roughly twofold greater (PT and PH) in the former than in the latter, while CYP4A10-mRNA was around fivefold higher in *mPPAR α* mice compared with that in *hPPAR α* mice. Next, we measured the effects of APFO treatment on triglyceride-synthesizing enzymes (Yen et al. 2008). APFO was found to increase DGAT1-mRNA levels in *Ppara*-null and *hPPAR α* mice (1.7- to 2.2-fold), but at a 5.0 mg/kg dose alone it decreased the mRNA in *mPPAR α* (0.7-fold). In contrast, APFO slightly reduced DGAT2-mRNA levels in *mPPAR α* and *hPPAR α* mice (~ 0.6-fold). Because DGAT1 and 2 are PPAR γ target genes (Ranganathan et al. 2006; Festuccia et al. 2009), the expression was also investigated: both APFO dosages increased PPAR γ mRNA levels in *Ppara*-null mice in a dose-dependent fashion (7- to 13-fold), and only slightly in *hPPAR α* mice (2- to 3-fold); in *mPPAR α* mice, they increased only at a 1.0 mg/kg dose (3.5-fold).

Since obvious inflammatory cell infiltration was seen in *Ppara*-null mice, but not in *mPPAR α* and *hPPAR α* mice, we measured NF κ B subunits: APFO treatments slightly induced p65-mRNA levels only in *Ppara*-null mice (2.2-fold), but even decreased them in *mPPAR α* (0.6-fold); in p50-mRNA, PFOA significantly decreased at 1.0 and 5.0 mg/kg dosages in *mPPAR α* mice (0.7- to 0.8-fold). As for p52-mRNA, the low- and high-dose APFO significantly raised the levels in *Ppara*-null (1.5-fold and 1.9-fold) and the high dose in *mPPAR α* mice (1.4-fold), although none of them increased the mRNA levels in *hPPAR α* mice. We also measured the pro-inflammatory cytokine TNF α -mRNA (Moriya et al. 2009) and a mitochondrial antioxidant UCP2-mRNA (Nègre-Salvayre et al. 1997). High-dose PFOA significantly increased TNF α -mRNA in *mPPAR α* , *Ppara*-null, and *hPPAR α* mice (2.9-fold, 1.9-fold, and 1.9-fold, respectively). An obvious difference was observed in the increase in UCP2-mRNA levels; the increase was dose dependent, with the high dose increasing by 28-fold in *mPPAR α* mice, while it raised the levels fivefold in *hPPAR α* mice. No increase in mRNA was observed in *Ppara*-null mice.

Western blot analysis

In the control group, PPAR α expression was greater in *hPPAR α* mice than in *mPPAR α* mice and surprisingly decreased APFO treatments only in the latter (Fig. 3). The expressions of PT and PH were significantly lower in *Ppara*-null mice than in *mPPAR α* and *hPPAR α* mice, while that of VLCAD was significantly higher in *hPPAR α* mice than in *mPPAR α* and *Ppara*-null mice. The levels of PT, PH, and VLCAD in *mPPAR α* mice appeared to reach the maximum at low-dose APFO and did not increase further at the high dose, while in *hPPAR α* mice, the increase tended to elevate in a dose-dependent fashion. APFO elevated MCAD protein only in *mPPAR α* mice at the high dose. The protein expression of the cell proliferation marker, PCNA, was higher in all APFO-treated *mPPAR α* and *hPPAR α* mice compared with respective controls.

The levels of p65 and p52 protein in the control group were lower in *Ppara*-null and *hPPAR α* mice than in *mPPAR α* mice, while TNF α levels were higher in *hPPAR α* mice than in *mPPAR α* and *Ppara*-null mice. APFO treatments increased the protein of p65 only in *Ppara*-null (1.8- to 1.9-fold) and *hPPAR α* mice (1.4- to 1.5-fold), but did not increase p50 levels. In contrast, APFO increased p52 levels in all genotyped mice at both dosages (1.5- to 2.0-fold). High-dose APFO significantly increased TNF α levels in all genotyped mice, while increases were also observed when compared with the low-dose treatment.

Discussion

We reported that microgram-order APFO could not activate human PPAR α but did activate mouse PPAR α (Nakamura et al. 2009). The current experiment has clearly shown that milligram-order APFO (1.0–5.0 mg/kg) activated both PPAR α to a similar or even smaller extent in human PPAR α . Therefore, the PPAR α function of humans might be weak compared with that of mice, due to the higher expression of human PPAR α compared with that of mouse PPAR α in the mouse livers used in this experiment. In that connection, Takacs and Abbott (2007) measured the lowest effective concentrations of PFOA at which mouse and human PPAR α were activated in an in vitro system, showing them to be 10 and 30 μ M, respectively. Wolf et al. (2008b) also reported that the activation of mouse PPAR α by APFO was generally higher compared with that of human PPAR α . Thus, there were clear-cut species differences in PPAR α activation between mice and humans, similar to those by trichloroethylene exposure (Ramdhan et al. 2010).

The first point for discussion in the present study is that APFO-induced hepatic damage was quite different histopathologically among *mPPAR α* , *Ppara*-null, and *hPPAR α* mice, unlike such differences in plasma ALT activities. Hypertrophic hepatocytes with eosinophilic cytoplasm were observed in *mPPAR α* and *hPPAR α* mice, suggesting the presence of PPAR α activation. This finding was consistent with the results that typical PPAR α target genes (e.g., CYP4A10, PH, PCNA, and UCP-2) were induced in both mouse groups. In contrast, inflammatory cell infiltrations were frequently observed in *Ppara*-null mice, which were also supported by the inflammation signaling analysis referring to p65 mRNA and protein expressions. It is of interest to note that APFO decreased p65 and p50 expressions in *mPPAR α* mice, which may have resulted from the strong activation of PPAR α in the mouse line. Mouse PPAR α induced by APFO treatment may completely inhibit the import of p50

and p65 directly into the nucleus or via I κ B α and thereby the inhibiting inflammation, since this receptor was shown to possess such a function (Moriya et al. 2009). However, the function of human PPAR α may be weak compared with that of mice and thus could not fully inhibit the inflammatory signaling in hPPAR α mice as it does in mPPAR α mice. The increase in p52 mRNA and protein were not correlated with inflammatory cell infiltrations and may be related instead to the increase in plasma ALT activity in all genotyped mice.

Our second point involves the histopathological differences in steatosis among three genotyped mice after APFO treatments: microvesicular steatosis was seen in Ppara-null and hPPAR α mice, whereas macrovesicular steatosis was observed only in Ppara-null mice. These findings were not replicated in mPPAR α mice. In line with these results, APFO treatments also increased hepatic TG and TC levels in Ppara-null and hPPAR α mice, though the increase in TC was not significant in the latter mouse line. In mPPAR α mice, 5.0 mg/kg dose decreased the TG level, which may be related to the elevation of β -oxidation enzymes. APFO treatments induced fatty acid β -oxidation enzymes more in mPPAR α mice than in hPPAR α mice, but very few in Ppara-null mice. In addition, APFO increased the expressions of DGAT1 in Ppara-null and hPPAR α mice more prominently compared with those in mPPAR α mice. As for DGAT2, the exposure decreased the expressions in mPPAR α and hPPAR α mice, but did not influence those in Ppara-null mice. Taken together, less or lower induction of fatty acid β -oxidation enzymes and higher induction of DGAT1 by APFO may, in part, reflect the increase of macrovesicular and/or microvesicular steatosis in Ppara-null and hPPAR α mice. However, we were unable to explain why macrovesicular steatosis was found only in Ppara-null mice. Further study is warranted to determine whether or not a PPAR γ or increase in cholesterol is involved in this regard in the liver of Ppara-null mice.

A question may arise as to why DGAT1 was induced in the liver of Ppara-null and hPPAR α mice, but not in mPPAR α mice, though increased in the 1.0 mg/kg dose group. DGAT1 is expressed in organs that produce large amounts of TG, such as liver, small intestine, and adipose tissue (Cases et al. 1998), which may be regulated by PPAR γ (Ranganathan et al. 2006; Festuccia et al. 2009). The mRNA levels of PPAR γ were greatly increased in Ppara-null and only slightly in hPPAR α and mPPAR α mice, but not in mPPAR α mice at 5.0 mg/kg dose. Thus, the activation of PPAR γ may be related to the increased DGAT1 in all genotyped mice. PFOA treatment also induces mitochondrial biogenesis at the transcriptional level with a preferential stimulation of mitochondrial DNA transcription, which occurs by way of the activation of a PPAR γ coactivator-1 α pathway (Walters et al. 2009). Indeed, PFOA is also reported to act as an agonist for PPAR γ (Vanden Heuvel et al. 2006). However, the question still remains why PPAR γ could not be activated by APFO treatment in mPPAR α mice. Since the activation of PPAR α by APFO was weak in hPPAR α mice and showed no activation at all in Ppara-null, this may result in increasing the return PPAR γ .

Our third concern in this study is why APFO treatments increased pro-inflammatory cytokine TNF α in all geno-typed mice. TNF α is thought to be an index of Kupffer cell activation (Yoshida et al. 2001), which may be related to the increase in necrotic (Morgan et al. 2008) or inflammatory cells (Dasarathy 2008). APFO treatment increased necrotic cells in the livers of mPPAR α and hPPAR α mice, and inflammatory cell infiltrations in the livers

of *Ppara*-null mice, both of which may be related to the increase in TNF α levels; these phenomena are also related to the rise in ALT activity.

Finally, in addition to macro/microvesicular steatosis, hydropic degeneration was characteristically observed in hPPAR α and *Ppara*-null mice. Minata et al. (2010) recently reported that PFOA in the liver was easily excreted into the bile duct in mPPAR α mice, but that the excretion in *Ppara*-null mice was less than half that in mPPAR α mice. Therefore, PFOA is thought to be much more accumulated in *Ppara*-null, and perhaps less so in hPPAR α mice, compared with that in mPPAR α mice. PFOA accumulated in hepatocytes poses a potential risk for mitochondrial dysfunction: APFO treatment results in an increase in the production of oxidative stress (Panaretakis et al. 2001) and in an enhancement of mitochondrial inner membrane permeability, which may disturb the mitochondrial inter-membranous electrochemical gradient, thus reducing the ATP production rate (Starkov and Wallace 2002). A powerful induction of the UCP2 expression following APFO treatment found only in mPPAR α mice may serve as protection against mitochondrial damage, because UCP2 is known to be a target gene of PPAR α and can inhibit oxidative stress production in mitochondria (Nègre-Salvayre et al. 1997; Kizaki et al. 2002). Very low levels of UCP2 induction in hPPAR α mice and non-induction in *Ppara*-null mice may not be enough to protect mitochondria from PFOA-induced oxidative stress. Furthermore, cellular stress induced by an excessive PFOA accumulation may also damage the endoplasmic reticulum, cytoskeleton, and microtubules, thus impairing excretion of proteins, lipids, and bile acids from hepatocytes. We consider that these abnormalities may eventually lead to the appearance of hydropic degeneration of hepatocytes, preferentially in *Ppara*-null and hPPAR α mice.

In this study, steatosis was not observed in the control *Ppara*-null mice. This is consistent with the previous report that 24-week-old control Sv/129 *Ppara*-null mice demonstrated neither apparent steatosis nor increases in hepatic TG contents compared with control wild-type mice (Tanaka et al. 2010). Okiyama et al. (2009) showed the presence of macrovesicular steatosis in the control group of *Ppara*-null mice, but they have used fat-rich liquid diet as a control. Furthermore, Rosen et al. (2010) reported the presence of significant macrovesicular steatosis in the control *Ppara*-null mice; such a discrepancy may be derived from the marked difference in mouse age (6- to 9-month-old mice in Rosen's study vs. 14-week-old mice in this study). It is plausible that steatosis may become obvious with age especially in *Ppara*-null mice, since mitochondrial β -oxidation ability is constitutively lower in *Ppara*-null mice than in wild-type mice (Aoyama et al. 1998).

In conclusion, PPAR α function may be very important in protecting against hepatic damage caused by PFOA, a result similar to the findings reported by many laboratories (Minata et al. 2010; Wolf et al. 2008b). In this regard, the human PPAR α function may be weak compared with that of mice. In addition, the expression of the receptor in human liver was 1/10 lower than that in mice or rats (Palmer et al. 1998). Therefore, hepatic damage may be induced if humans are exposed to high doses of PFOA. In such cases, histopathological findings may resemble those of *Ppara*-null mice more than mPPAR α mice.

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Abbreviations

ALT	Alanine aminotransferase
APFO	Ammonium perfluorooctanoate
CV	Central vein
CYP4A10	Cytochrome P450 4A10
DGAT1	Diacylglycerol acyltransferase 1
DGAT2	Diacylglycerol acyltransferase 2
H & E	Hematoxylin and eosin
hPPARα mice	Humanized PPAR α mice
MCAD	Medium chain acyl-CoA dehydrogenase
NFκB	Nuclear factor kappa B
PCNA	Proliferation cell nuclear antigen
PH	Peroxisomal bifunctional protein
PPARα	Peroxisome proliferator-activated receptor α
PPARγ	Peroxisome proliferator-activated receptor γ
PT	Peroxisomal thiolase
TC	Total cholesterol
TG	Triglycerides
TNFα	Tumor necrosis factor α
UCP2	Uncoupling protein 2
VLCAD	Very long chain acyl-CoA dehydrogenase

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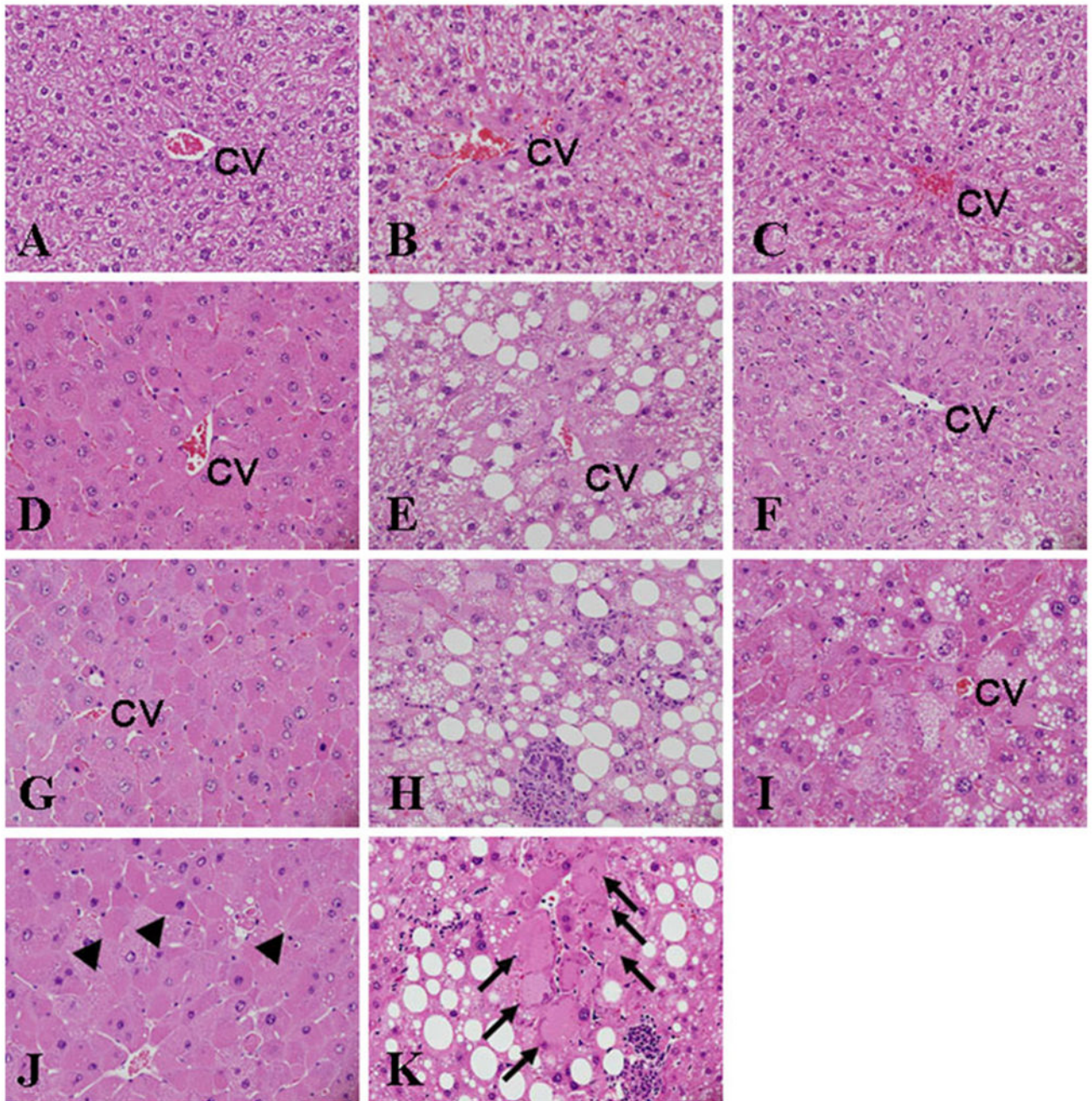


Fig. 1. Liver histopathological findings after APFO treatment. Liver section photomicrograph taken from *mPPARα* (a), *Ppara*-null (b) and *hPPARα* (c) mice treated with 0 mg/kg of APFO. Liver from *mPPARα* (d), *Ppara*-null (e) and *hPPARα* (f) mice treated with 1.0 mg/kg of APFO. Liver from *mPPARα* (g), *Ppara*-null (h), *hPPARα* (i) mice treated with 5.0 mg/kg of APFO. PV, portal vein; CV, central vein; original magnification $\times 400$. *Arrowheads* (j) and *arrows* (k) indicate hypertrophied and hydropic hepatocytes with eosinophilic cytoplasm in *mPPARα* and *Ppara*-null mice treated with 5.0 mg/kg APFO, respectively

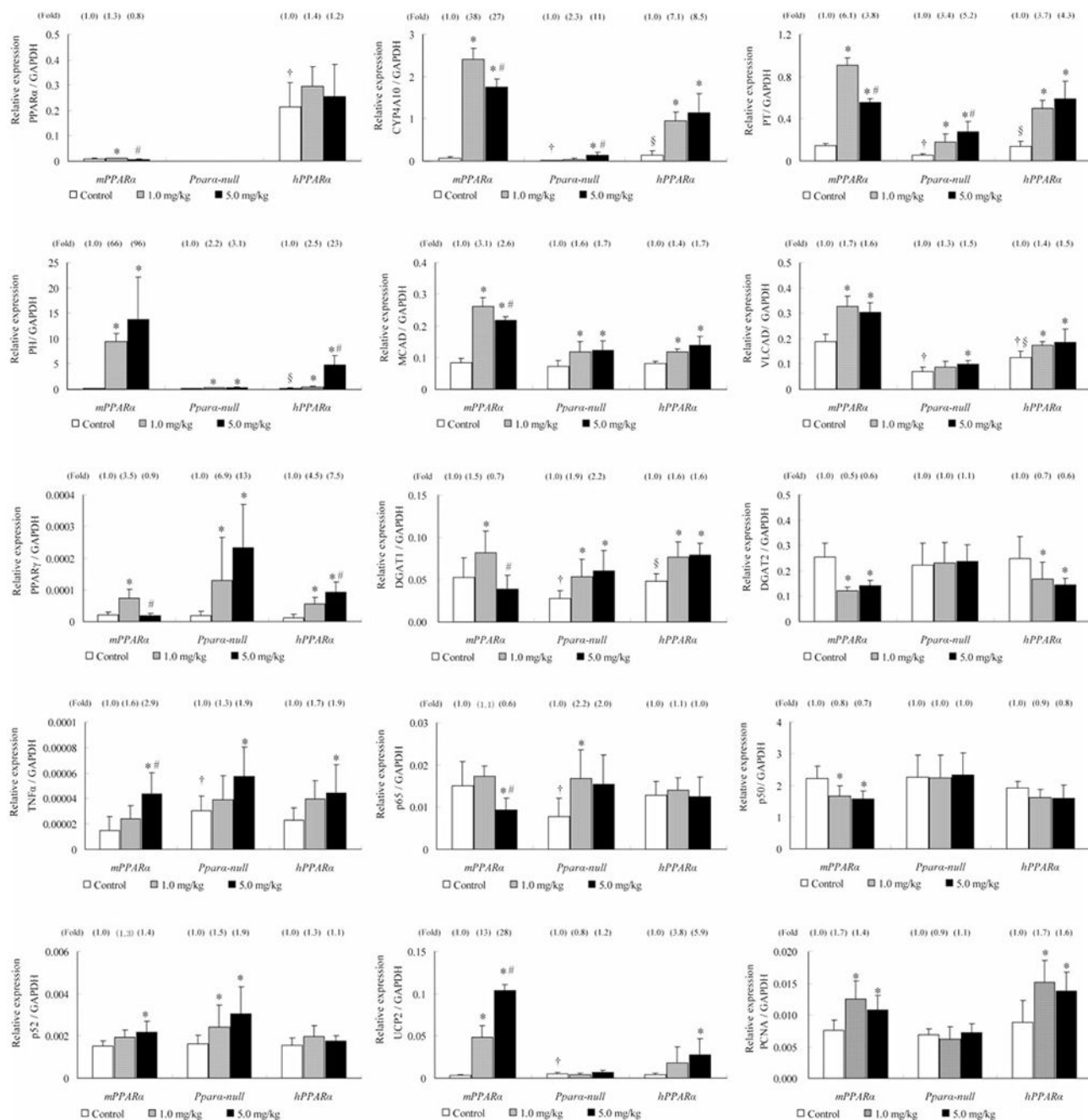
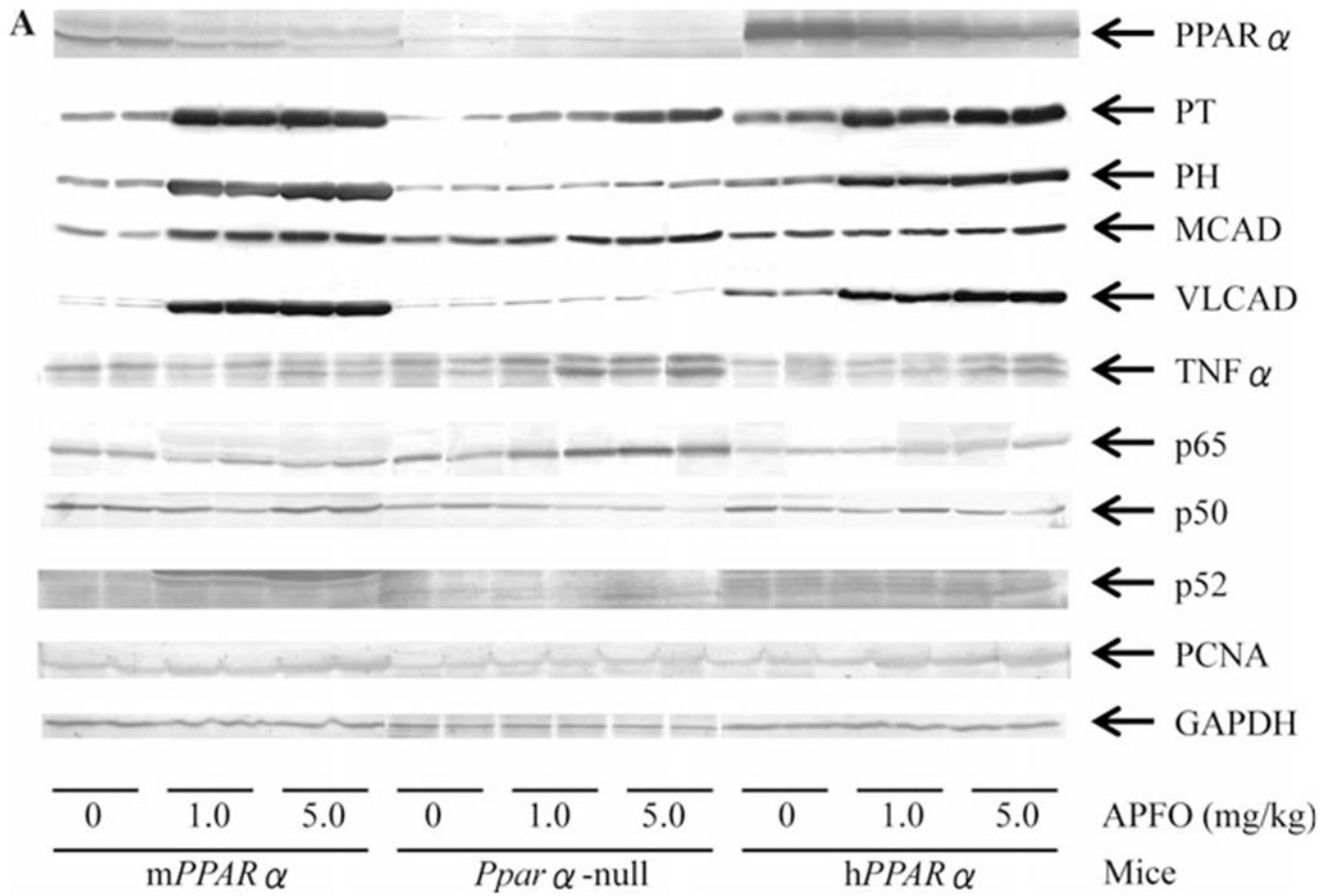
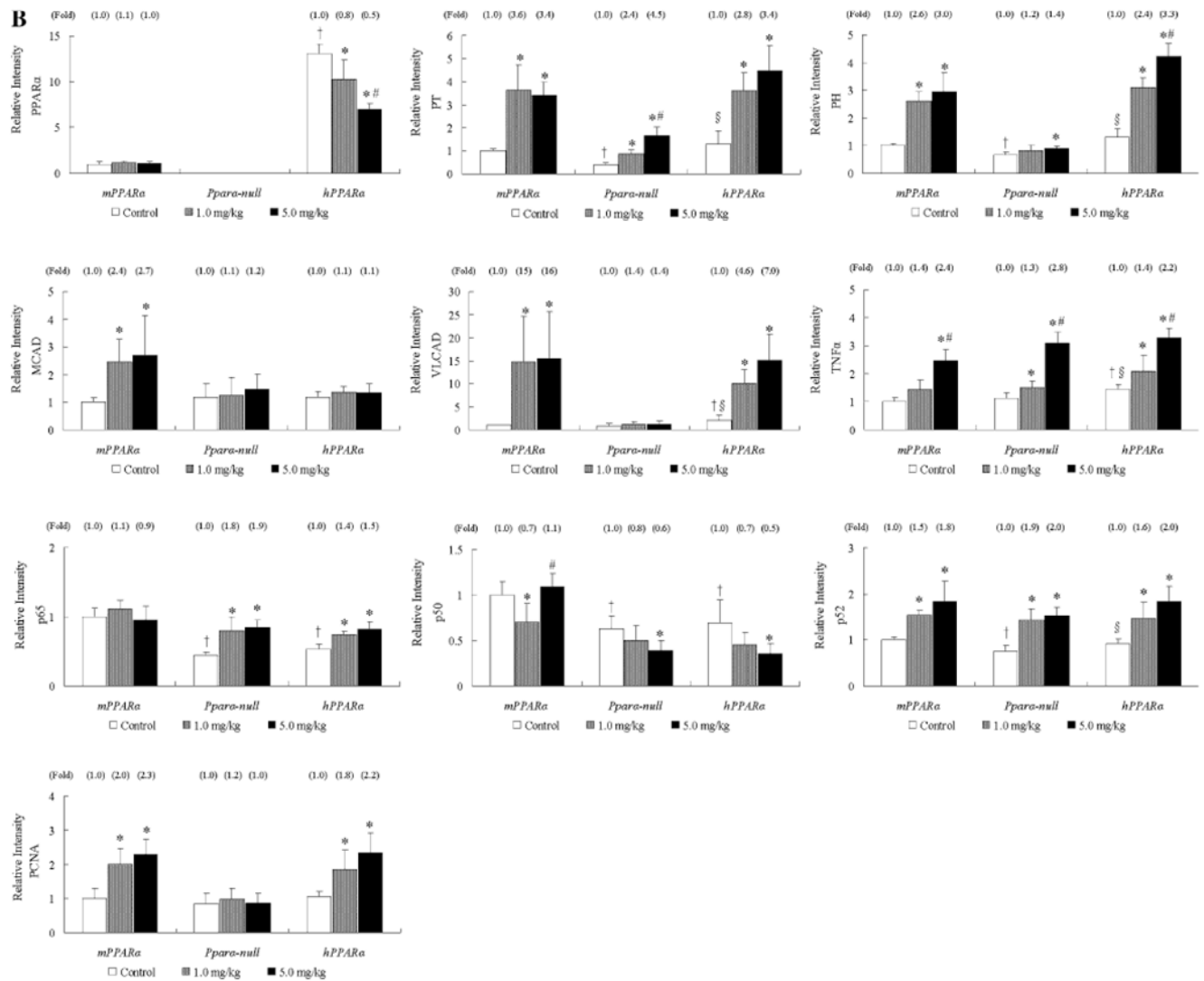


Fig. 2. mRNA expressions of PPARα and the related genes in livers from *mPPARα*, *Ppara*-null and *hPPARα* mice treated with APFO. Values represent means ± SD for 8–10 mice per group. *Significantly different from respective control (0 mg/kg) group ($P < 0.05$). #Significantly different from respective low-dose (1.0 mg/kg) group ($P < 0.05$). †Significantly different from control of *mPPARα* ($P < 0.05$). ‡Significantly different from control of *Ppara*-null mice ($P < 0.05$).



**Fig. 3.**

Protein expressions of PPAR α and the related genes in livers from *mPPAR α* , *Ppara*-null and *hPPAR α* mice treated with APFO. **a** Representative Western blot analyses. As an internal standard, GAPDH was stained. **b** Each band was quantified by densitometric analysis. Histogram presents means \pm SD for 8–10 mice per group, and the mean from each control group in *mPPAR α* mice was assigned a value of 1.0. *Significantly different from respective control (0 mg/kg) group ($P < 0.05$). #Significantly different from respective low-dose (1.0 mg/kg) group ($P < 0.05$). †Significantly different from control of *mPPAR α* ($P < 0.05$). §Significantly different from control of *Ppara*-null mice ($P < 0.05$)

Table 1

Body and liver weight, plasma ALT activity, plasma and hepatic TG and TC levels in *mPPAR α* , *Ppara*-null, and *hPPAR α* mice treated with APFO

	APFO		
	Control	1.0 mg/kg	5.0 mg/kg
Body weight (g) at the start of APFO treatments			
<i>mPPARα</i>	21.0 ± 0.9 (8)	20.3 ± 2.2 (8)	20.7 ± 1.6 (9)
<i>Ppara</i> -null	21.1 ± 2.5 (9)	21.1 ± 2.8 (9)	21.8 ± 1.5 (9)
<i>hPPARα</i>	20.3 ± 1.9 (9)	19.4 ± 1.0 (9)	20.3 ± 2.5 (10)
Body weight (g) after APFO treatments			
<i>mPPARα</i>	24.8 ± 1.0 (8)	24.7 ± 2.2 (8)	23.5 ± 1.7 (9)
<i>Ppara</i> -null	24.2 ± 2.4 (9)	23.6 ± 2.8 (9)	25.6 ± 2.5 (9)
<i>hPPARα</i>	23.5 ± 1.3 (9)	22.3 ± 1.7 (9)	23.5 ± 2.1 (10)
Liver weight (g)			
<i>mPPARα</i>	1.05 ± 0.06 (8)	1.98 ± 0.22 (8) [*]	2.72 ± 0.29 (9) ^{**#}
<i>Ppara</i> -null	1.10 ± 0.10 (9)	1.57 ± 0.29 (9) [*]	3.35 ± 0.48 (9) ^{**#}
<i>hPPARα</i>	1.03 ± 0.08 (9)	1.48 ± 0.14 (9) [*]	2.49 ± 0.35 (10) ^{**#}
Liver/body ratio (%)			
<i>mPPARα</i>	4.23 ± 0.23 (8)	8.02 ± 0.62 (8) [*]	11.54 ± 0.71 (9) ^{**#}
<i>Ppara</i> -null	4.56 ± 0.38 (9)	6.68 ± 1.27 (9) [*]	13.06 ± 1.16 (9) ^{**#}
<i>hPPARα</i>	4.37 ± 0.30 (9)	6.67 ± 0.42 (9) [*]	10.56 ± 0.91 (10) ^{**#}
Plasma ALT (IU/L)			
<i>mPPARα</i>	6.7 ± 0.9 (8)	7.6 ± 2.6 (8)	15.6 ± 8.6 (9) ^{**#}
<i>Ppara</i> -null	7.2 ± 1.6 (9)	8.6 ± 2.5 (9)	9.3 ± 1.8 (9) [*]
<i>hPPARα</i>	6.5 ± 1.3 (9)	8.1 ± 2.7 (9)	8.9 ± 1.5 (10) [*]
Plasma TG (mg/dl)			
<i>mPPARα</i>	156.4 ± 43.6 (8)	107.3 ± 27.4 (8) [*]	77.6 ± 19.0 (9) [*]
<i>Ppara</i> -null	168.7 ± 80.6 (9)	183.2 ± 67.2 (9)	164.8 ± 57.7 (9)
<i>hPPARα</i>	195.9 ± 69.2 (9)	209.5 ± 58.7 (9)	142.6 ± 36.9 (9) [#]

		APFO	
		Control	5.0 mg/kg
Hepatic TG (mg/g)			
mPPAR α	13.3 \pm 1.5 (8)	23.4 \pm 5.5 (8)*	11.8 \pm 3.8 (9) [#]
Ppara-null	19.6 \pm 11.0 (9)	58.7 \pm 38.6 (9)*	106.3 \pm 34.9 (9) ^{##}
hPPAR α	17.4 \pm 4.4 (9)	34.4 \pm 15.7 (9)*	51.7 \pm 11.9 (10) ^{##}
Plasma TC (mg/dl)			
mPPAR α	66.9 \pm 22.3 (8)	63.8 \pm 17.6 (8)	39.4 \pm 11.9 (9) ^{##}
Ppara-null	109.5 \pm 27.7 (9) [†]	84.5 \pm 16.3 (9)	57.8 \pm 21.5 (9) ^{##}
hPPAR α	106.6 \pm 13.3 (9) [†]	99.2 \pm 27.2 (9)	83.5 \pm 25.4 (10)
Hepatic TC (mg/g)			
mPPAR α	3.7 \pm 0.8 (8)	3.6 \pm 0.8 (9)	5.3 \pm 1.3 (9)
Ppara-null	4.5 \pm 0.6 (9)	6.8 \pm 2.5 (9)	10.4 \pm 4.4 (9)*
hPPAR α	5.4 \pm 0.8 (9)	6.7 \pm 1.5 (9)	6.1 \pm 1.0 (10)

Data represent mean \pm SD. Data in parentheses are the numbers of mice

* Significantly different from respective control (0 mg/kg) group ($P < 0.05$)

[#] Significantly different from respective low-dose (1.0 mg/kg) group ($P < 0.05$)

[†] Significantly different from control of mPPAR α mice ($P < 0.05$)

Table 2

Changes in livers after APFO treatment

Genotype	mPPAR α		Ppara-null		hPPAR α	
	Control	1.0 mg/kg APFO	5.0 mg/kg APFO	Control	1.0 mg/kg APFO	5.0 mg/kg APFO
Macrovesicular steatosis	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0	1.2 \pm 0.8*	2.0 \pm 0.7*
Microvesicular steatosis	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0	0.6 \pm 0.5	0.6 \pm 0.5
Lobular inflammatory cells	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0	2.0 \pm 0*,#
Single-cell necrosis	0 \pm 0	0 \pm 0	0.8 \pm 0.4	0 \pm 0	0 \pm 0	0.2 \pm 0.4
Hydropic degeneration	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0	2.0 \pm 0*,#

Steel–Dwass method was conducted to compare the effects of APFO treatment among each treated group of each genotyped mouse

Data represent mean \pm SD for 5 mice per group* Significantly different from respective control (0 mg/kg) group ($P < 0.05$)# Significantly different from respective low-dose (1.0 mg/kg) group ($P < 0.05$)