

Oleuropein-induced acceleration of CYP-catalyzed drug metabolism: central role for nuclear receptor PPAR α

Foteini Malliou^{1#}, Christina E. Andriopoulou^{1#}, Frank J. Gonzalez², Aristeidis Kofinas¹, Alexios-Leandros Skaltsounis³, Maria Konstandi^{1*}

¹University of Ioannina, School of Health Sciences, Faculty of Medicine, Department of Pharmacology, Ioannina GR-45110, Greece;; ²Laboratory of Metabolism, National Cancer Institute, National Institutes of Health, Bethesda 20892, Maryland, USA; ³Faculty of Pharmacy, School of Health Sciences, University of Athens, Greece

Corresponding author: Maria Konstandi, E-mail: mkonstan@uoi.gr; Tel: +3026510 07554, Fax: +3026510 07859.

#These authors have equally contributed

Keywords: CYP induction; CYP1A; CYP2C; CYP2D; CYP3A; peroxisome proliferator-activated receptors; drug metabolism

Running title: Oleuropein induces major CYPs via PPAR α activation

Number of text pages: 22

Number of tables: 1

Number of figures: 8

Number of references: 105

Number of words in abstract: 218

Number of words in introduction: 925

Number of words on discussion: 1824

Abbreviations: ACN, acetonitrile; CYP, cytochrome P450; C, control; CAR, constitutive androstane receptor; CREB, cAMP responsive element-binding protein; FOXO1, Forkhead box protein O1; HNF4 α , hepatocyte nuclear factor 4 α ; JNK, c-Jun N-terminal kinase; PRL, prolactin; PXR, pregnane X receptor; RXR, retinoic X receptor; rCYP, recombinant CYP; WORT, wortmannin. EIA, Elisa; OLE, oleuropein; FEN, fenofibrate; PKA, protein kinase A; PPAR α , peroxisome proliferator-activated receptor α ; PCR, polymerase chain reaction; RIA, radioimmunoassay.

Abstract

Oleuropein (OLE), the main constituent of *olea oleuropea*, displays pleiotropic beneficial effects in health and disease, which are mainly attributed to its antiinflammatory and cardioprotective properties. Several food supplements and herbal medicines contain OLE and are available without a prescription. This study investigated the effects of OLE on the main CYPs catalyzing the metabolism of many prescribed drugs. Emphasis was given on the role of peroxisome proliferator-activated receptor alpha (PPAR α), a nuclear transcription factor regulating numerous genes including *CYPs*. 129/Sv wild-type and *Ppara*-null mice were treated with OLE for six weeks. OLE induced *Cyp1a1*, *Cyp1a2*, *Cyp1b1*, *Cyp3a14*, *Cyp3a25*, *Cyp2c29*, *Cyp2c44*, *Cyp2d22* and *Cyp2e1* mRNAs in liver of wild-type mice, whereas no similar effects were observed in *Ppara*-null mice, indicating that the OLE-induced effect on these *CYPs* is mediated by PPAR α . Activation of the PI3k/AKT/FOXO1, JNK, AKT/p70 and ERK related pathways participates in CYP induction by OLE. These data indicate that consumption of herbal medicines and food supplements containing OLE could accelerate the metabolism of drug substrates of the above mentioned CYPs, thus reducing their efficacy and the outcome of pharmacotherapy. Therefore, OLE-induced activation of PPAR α could modify the effects of drugs due to their increased metabolism and clearance, which should be taken into account when consuming OLE-containing products with certain drugs, in particular those of narrow therapeutic window.

Significance statement

This study indicated that oleuropein, which belongs to the main constituents of the leaves and olive drupes of *Olea europaea*, induces the synthesis of the major CYPs metabolizing the majority of prescribed drugs via activation of PPAR α . This effect could modify the pharmacokinetic profile of co-administered drugs-substrates of the CYPs, thus altering their therapeutic efficacy and toxicity.

Introduction

Globally, there is an upsurge in the use of medicines coming from herbs as many people claim that they are non-toxic based on their natural origin and their use in popular medicine for centuries. Nonetheless, herbs contain many active substances that could induce adverse effects, toxicity or even cancer (Bensoussan et al., 2002; De Smet, 1995; Deng, 2002; Eisenberg et al., 1993; Ernst and Pittler, 2002a; Ernst and Pittler, 2002b; Greensfelder, 2000; Haller and Benowitz, 2000; Kennedy and Seely, 2010; Klepser and Klepser, 1999; Koh and Woo, 2000; Malliou et al., 2018; Mckenna et al., 2012; McRae et al., 2002; Stedman, 2002). When all foreign substances (xenobiotics) enter the body, they undergo biotransformation primarily in the liver, which accelerates their elimination. In particular, the metabolism of xenobiotics, such as drugs, pre-carcinogens and toxic agents during Phase I is mainly catalyzed by CYPs and can result either in activation of pro-drugs or in inactivation of pharmacologically active drugs, or in activation of pre-carcinogens to carcinogens (Gonzalez and Gelboin, 1994b; Ingelman-Sundberg, 2004a). It is estimated that CYPs belonging to the CYP3A, CYP2C, CYP2D and CYP1A subfamilies catalyze hepatic metabolism of over than 95% of the most widely prescribed drugs (Daskalopoulos et al., 2012a; Guengerich, 2003; Ingelman-Sundberg, 2004a; Konstandi, 2013). From a toxicological point of view, it is worth noting that CYP1A1/2 and CYP1B1 catalyze the bioactivation of the major groups of pre-carcinogens, the polycyclic aromatic hydrocarbons (PAHs), polycyclic arylamines and aflatoxin B1 to electrophilic DNA-binding derivatives (Cheng and Morgan, 2001; Flint et al., 2010; Harkitis et al., 2015; Kawajiri, 1999; Konstandi et al., 2006; Konstandi et al., 2005; Pasanen and Pelkonen, 1994). Interestingly, the

biotransformation of steroids, fatty acids and several other endogenous compounds is also catalyzed by CYPs (Guengerich, 2003; Spatzenegger and Jaeger, 1995).

It is well-documented that the traditional Mediterranean diet has various beneficial effects in health and longevity, and olive oil and olives are substantial ingredients of this diet (Impellizzeri et al., 2012). The main compounds found in the leaves and olive drupes of *Olea europaea* are oleuropein (OLE) and its hydrolysis product, hydroxytyrosol. There is accumulating evidence based on preclinical studies that OLE displays significant cardioprotective properties (Andreadou et al., 2006; Briante et al., 2001; Malliou et al., 2018; Oi-Kano et al., 2008; Tuck and Hayball, 2002; Visioli and Galli, 1994; Visioli et al., 2002a; Visioli et al., 2002b), which could be attributed to activation of peroxisome proliferator-activated receptor α (PPAR α) (Harkitis et al., 2015; Malliou et al., 2018), a ligand-activated nuclear receptor that controls lipid homeostasis (Fruchart and Duriez, 2006; Hansen and Connolly, 2008; Kuusisto et al., 2007; Robillard et al., 2005). The anti-inflammatory properties of OLE may also contribute to the drug's cardioprotective effects (Shimada et al., 1996; Spatzenegger and Jaeger, 1995). Preliminary data also indicated a pleiotropic effect of OLE on several vital functions of the body including stimulation of neural plasticity and protection against neurodegenerating disorders, among others, effects potentially mediated by activation of PPAR α (data not shown).

It is well defined that OLE activates PPAR α (Malliou et al., 2018; Spatzenegger and Jaeger, 1995), which controls several genes participating in the regulation of inflammatory responses, the metabolism of lipids and glucose, as well as the adipose differentiation and cancer, among others (Malliou et al., 2018; Yang et al., 2008; YUMUK, 2006). It is of interest that PPAR α apparently holds a substantial regulatory role for CYPs (Choi and Waxman, 1999; Tauber et al., 2020). In

particular, the hepatic sexual dimorphism of the CYP expression pattern is largely regulated by PPARs (Leuenberger et al., 2009).

Drug-drug interactions are of critical clinical significance, because they markedly determine the outcome of pharmacotherapy, the side effects of the drugs and pharmacotoxicity (Konstandi, 2013). They are usually dependent on the drugs' effect on CYPs acting either as inducers, inhibitors or substrates. On one hand, induction of the most important CYPs that catalyze the metabolism of the majority of prescribed drugs, may accelerate the biotransformation of their drug-substrates and in most cases, result in their reduced pharmacological efficacy (Daskalopoulos et al., 2012a; Daskalopoulos et al., 2012b; Ingelman-Sundberg et al., 2007; Konstandi et al., 2020; Konstandi et al., 2005; Zhou et al., 2004), whereas in some cases, in perturbation of several endogenous regulatory circuits, often associated with pathophysiological states (Choi and Waxman, 1999). On the other hand, inhibition of CYPs may lead to accumulation of their drug-substrates in the blood, followed by adverse side effects of varying severity that may reach the level of toxicity and this is of particular clinical interest when drugs of low therapeutic index are administered (Daskalopoulos et al., 2012b; Konstandi et al., 2020; Spatzenegger and Jaeger, 1995). Therefore, it should be noted that food supplements or herbal medicines containing pharmacologically active compounds, such as OLE, which act either as substrates of CYPs, or even as their inducers or inhibitors (Zhou et al., 2004), may modify the efficacy of co-administered drugs in multi-drug therapeutic schemes and potentially, induce pharmacotoxicity (Daskalopoulos et al., 2012a; Daskalopoulos et al., 2012b; Gonzalez and Gelboin, 1994a; Guengerich, 2003; Harkitis et al., 2015; Konstandi et al., 2014; Pelkonen et al., 2008).

In the light of the above considerations, current study investigated the potential regulatory role of OLE for the main CYPs that are involved in drug metabolism, emphasizing the role of PPAR α in this regulation. To approach this issue, wild-type (WT) and *Ppara*-null mice were treated with OLE and *Cyp* mRNA and protein expressions were analyzed. OLE markedly upregulated several genes encoding the most significant drug metabolizing CYP isozymes in the liver, a process profoundly mediated by PPAR α activation.

Materials and Methods

Animals and treatment: Adult male 129/Sv wild-type (WT) and *Ppara*-null mice that were used in this study were bred in the Animal House of the University of Ioannina (Ioannina, Greece) and were housed up to five mice per cage. All animals (5-6 per group of treatment) had access to a standard chow diet for rodents (1324 TPF, Altromin Spezialfutter GmbH & Co. KG) and water *ad libitum*. Throughout the experiments, all mice were housed in their cages under a standard 12-h light/12-h dark cycle (lights on at 6:00 AM). The experimental protocols employing procedures with animals received the approval of the Ethics Committee of the University of Ioannina- Faculty of Medicine. The procedure followed was in compliance with the European Commission ethical standards for the care and use of experimental animals (Directive 86/609-EEC). Both, WT and *Ppara*-null mice were treated with food pellets containing OLE (100 mg/kg) daily for 6 consecutive weeks (Andreadou et al., 2006; Andreadou et al., 2014; Impellizzeri et al., 2012). For the isolation of OLE the leaves of *Olea europaea* were used following a previously described method (Andreadou et al., 2014). The choice of the dose of OLE that was used in this study was based on information from the literature and our previous findings (Andreadou et

al., 2006; Andreadou et al., 2014; Impellizzeri et al., 2012). It corresponds approximately to the average consumption of olive oil and olive drups in the Mediterranean countries during a day (Abdel-aleem et al., 1997), and represents total polyphenol consumption from these olive products and is estimated to be approximately 100mg/day (Abdel-aleem et al., 1997; Del Boccio et al., 2003). Mice were treated with OLE in food pellets, because it is known that the drug is slightly absorbed in intestinal lumen even under normal iso-osmotic conditions. The absorption of OLE can be markedly improved by solvent flux through paracellular junctions, an effect that is facilitated by hypotonic luminal conditions (Edgecombe et al., 2000). A significant factor stimulating the water flux through the opening of paracellular junctions is the postprandial presence of glucose or amino acids in the intestinal lumen. This mechanism regulating the absorption of OLE in intestinal lumen appears to have similar effects with those of hypotonic solutions (Pappenheimer and Reiss, 1987). The pharmacokinetic profile of OLE in mice has not been determined, but only that in rats, when one dose of OLE (100 mg/kg per os) reached 200 ng/ml in tmax of 2 h (Boccio et al., 2003). Controls received regular rodent food for 6 weeks. Wild type and *Ppara*-null mice also recieved intraperitoneally either fenofibrate (FEN, 100mg/kg), a selective PPAR α agonist (Ghonem et al., 2015; Hu et al., 2019) or normal saline for 7 consecutive days. Upon completion of the experiments, all mice were euthanized using CO₂ asphyxiation, and parts from the liver were dissected for the extraction of total RNA, nuclear/cytosolic and microsomal proteins. All liver tissue samples were preserved at -80°C until assayed.

Isolation of microsomal proteins: Liver tissue was homogenized in homogenization buffer containing 0.15 M KCl, 10 mM K₂EDTA and 1 mM dithiothreitol (pH 7.4) at +4°C, for the isolation of microsomal fractions. The homogenates were then centrifuged for 20 min at 15,000 rpm (+4°C). The upper phase was transferred into clean vials followed by centrifugation at 27,500 rpm (+4°C) for 60 min. The formed microsomal pellet was resuspended by homogenization in the specific ice-cold homogenization buffer and centrifuged at 27,500 rpm for 45 min. Temperature was always kept at +4°C. Finally, the formed microsomal pellet was resuspended in the specific ice-cold storage buffer containing K₂HPO₄/KH₂PO₄ (pH 7.4), K₂EDTA (1 mM), dithiothreitol (0.1 mM) and 20% glycerol. Aliquots of microsomal suspensions were stored at -80°C until assayed (Lang et al., 1981).

CYP2D Activity, 1'-Bufuralol hydroxylation: 1'-Bufuralol hydroxylation is mainly catalysed by CYP2D isozymes (Matsunaga et al., 1990). Liver microsomal proteins (40 µg ~ 20µl of sample) were pre-incubated in a 180 µl reaction mixture containing potassium phosphate (0.1 M, pH 7.4) at 37°C for 5 min, in the presence of 50 µM bufuralol (substrate) and NADPH (0.5 mM, Sigma-Aldrich). The duration of the reaction was 7.5 min and was terminated using 20 µl of perchloric acid (60%). Following a ten min centrifugation at 14,075g, the supernatant containing the main metabolite of bufuralol (1'-hydroxy-bufuralol) was analysed using an HPLC method. The fluorescence detection was set at 252 nm (excitation wavelength) and 302 nm (emission wavelength) and a specific column (reverse-phase Luna C18, 5 µm, 150X 3 mm; Phenomenex, Torrance, CA) was used for this purpose. The mobile phase consisted of 30% acetonitrile/70% perchlorate buffer (20 mM, pH 2.5). The elution of each sample took place at a flow rate of 1 ml/min for 14 min. The recombinant rat

CYP2D1 and CYP2D2, enriched with CYP reductase BD Supersomes, was used as a positive control of bufuralol 1'-hydroxylation (BD Gentest, Woburn, MA). Bufuralol (50 μ M) was used as substrate, which was pre-incubated at 37°C for 2 min with approximately, 200 μ l of 0.1 M potassium phosphate buffer (pH 7.4). Recombinant CYPs (50 μ M) along with NADPH (1 mM) were added in the mixture, which was incubated at 37°C for 30 min. The termination of the reaction was achieved with 20 μ l acetonitrile. Then, all samples were placed on wet ice and left undisturbed for 10 min, before their injection into the HPLC for analysis.

Western Blot Analysis: Alterations in CYP apoprotein levels were assessed with immunoblot analysis using liver microsomes, while nuclear extracts were used for PPAR α and HNF4 α analysis. pJNK, pP38, pERK, pP70, PXR, CAR and pFOX01 were analyzed using total cellular extracts. For the preparation of nuclear and cytosolic extracts, the corresponding cytosolic and nuclear extraction kit of Thermo Fisher Scientific, Waltham, MA was used. The content of proteins in the sample was determined using the bovine serum albumin (BSA) assay (Thermo Fisher Scientific). All proteins run on SDS-polyacrylamide gel electrophoresis followed by immunoblotting and for this purpose the following antibodies were used: mouse/rat polyclonal CYP3A, CYP2C and CYP2D IgGs, mouse/rat monoclonal HNF4 α , CAR, PXR IgGs and mouse polyclonal PPAR α IgG (Santa Cruz Biotechnology). The rabbit phosphorylated polyclonal IgGs CREB-1 (Ser133), P38 (Thr180/ Tyr182) (Santa Cruz Biotechnology), MAPK (Erk1/2) (Thr202/Tyr204) (Cell Signaling Technology), p70S6K (Thr389) (Cell Signaling Technology), FOX01 (Ser256) (Santa Cruz Biotechnology), AKT (Ser473) (Santa Cruz Biotechnology) and the mouse phosphorylated monoclonal JNK (Thr183 and Tyr185) IgG (Santa Cruz

Biotechnology). The secondary antibodies used in this study were conjugated with horseradish peroxidase (Santa Cruz Biotechnology). For the detection of proteins in the blot the enhanced chemiluminescence detection kit (GE Healthcare, Chalfont St. Giles, Buckinghamshire, UK) was used. As a loading control, immunoblotting employing the specific antibodies, Histone H3 or actin (Santa Cruz Biotechnology), and the secondary antibody, anti-mouse horseradish peroxidase-conjugated IgG was used.

Quantitative Real-Time Polymerase Chain Reaction Assays: TRIzol reagent (Invitrogen) was used for the isolation of total RNA from liver tissue following the instructions in the manufacturer's protocol. Following a spectrophotometric method the concentration of total RNA in each sample was determined. Total RNA (1 μ g) and a SuperScript II reverse transcriptase kit (Invitrogen) were used to generate cDNA, which was used in quantitative and reverse transcription-polymerase chain reaction (PCR) assays. Table 1 shows all sequences of the forward and reverse gene-specific primers that were used in this study. For the real-time reactions, the SYBR Green PCR Master Mix (Applied Biosystems, Foster City, CA) was used, and these reactions were performed employing the C1000 Touch thermal cycler with a real-time detection system (Bio-Rad Laboratories, Hercules, CA). Relative mRNA expression was estimated using β -actin mRNA levels to normalize mRNA expression levels of each gene (QuantiTect primer assay; QIAGEN, Valencia, CA). The comparative threshold cycle method was used to quantify all values.

Statistical Analysis: For the analysis of data, the one-way analysis of variance followed by multiple comparisons employing Bonferroni's and Tukey's least honest significant difference methods was used. All values are presented as mean \pm S.E. and p values of <0.05 were considered significant.

Results

OLE-induced effect on CYP1s

OLE markedly increased hepatic *Cyp1a1* and *Cyp1a2* mRNA expression and CYP1A protein expression in WT mice (Fig. 1A and 1B, respectively); *Cyp1b1* mRNA levels were also increased by OLE (Fig. 1C).

Oleuropein-induced effect on CYP2 and CYP3s

Oleuropein markedly increased hepatic *Cyp2c29* and *Cyp2c44* mRNA and CYP2C protein expression in WT mice (Fig. 2A & 2B, respectively). Similarly, OLE increased the hepatic *Cyp2d22* mRNA, CYP2D protein and activity levels (Fig. 2C) and the hepatic *Cyp2e1* (Fig. 3A), *Cyp3a14* and *Cyp3a25* expression (Fig. 3B) at mRNA and protein levels.

Assessment of OLE effect on transcription factors and signal transduction pathways related to CYP regulation

The OLE-induced effect on aryl hydrocarbon receptor (AHR), aryl hydrocarbon receptor repressor (AhrR) and heat shock protein 90 (HSP90) (Fig. 4A, 4B and 4C,

respectively), critical transcription factors in *Cyp1a1*, *Cyp1a2* and *Cyp1b1* regulation (Nebert et al., 2013), is apparently followed by upregulation of these CYPs, an effect profoundly associated with the OLE-induced PPAR α activation (Malliou et al., 2018), because OLE did not increase *Ahr*, *AhrR* and *Hsp90* mRNAs in *Ppara*-null mice, but the drug rather slightly repressed the expression of these transcription factors (Fig. 4A-4C).

The OLE-induced effect on *Cyp2c29*, *Cyp2c44* and *Cyp3a25* mRNA expression in the liver of WT mice is apparently mediated by induction of *Pxr*, *Car*, *Rxra* and *Rxrb*, which encode critical transcription factors regulating the above CYP genes (Daskalopoulos et al., 2012a) (Fig. 5A-5D). The drug's upregulating effect on these transcription factors is probably mediated by PPAR α , because, OLE had a repressing effect on *Pxr*, *Car* and *Rxrb* in *Ppara*-null mice (Fig. 5A, 5B and 5D, respectively).

Long term treatment of WT mice with OLE induced hepatic *Ppara* mRNA expression by 4-5 fold, a stimulating effect that was also evident at protein level (Malliou et al., 2018), (Supplemental Fig. 1). Further analysis employing molecular docking experiments suggests that OLE is a PPAR α ligand and the luciferase reporter gene assay revealed significant activation of this nuclear receptor and transcription factor by OLE (Malliou et al., 2018), followed by upregulation of various PPAR α target genes including *Acox1*, *Acot1*, *Cyp4a10* and *Cyp4a14* (Malliou et al., 2018). *Hnf4a*, encoding a nuclear transcription factor, which holds a determinant role in the regulation of several CYP genes (Daskalopoulos et al., 2012a; Konstandi et al., 2020), was also upregulated by OLE in the liver of mice (Malliou et al., 2018), (Supplemental Fig. 1). It is assumed that the OLE upregulating effect on all Cyps analyzed in this study is profoundly mediated by PPAR α , because no similar effects

on these *Cyps* were observed in *Ppara*-null mice (Fig. 1A-1C, Fig. 2A-2C, Fig. 3A-3C). In particular, OLE either repressed *Cyp1a1*, *Cyp1a2*, *Cyp1b1* (Fig. 1A-1C) *Cyp2c29*, *Cyp2c44*, *Cyp2d22* (Fig. 2A-2C) and *Cyp3a25* (Fig. 3C) or had no effect on *Cyp2e1* (Fig. 3A) and *Cyp3a14* (Fig. 3B) in the liver of *Ppara* deficient mice. The determinant role of PPAR α in CYP induction was also confirmed in mice treated with the selective PPAR α agonist, FEN (Ghonem et al., 2015; Hu et al., 2019), which increased CYP1A1, CYP1A2, CYP1B1, CYP2C, CYP2D, CYP2E1 and CYP3A protein expression in the liver of WT mice, whereas FEN had no similar effects in the liver of PPAR α -deficient mice (Fig 6).

The role of nitric oxide synthases, endothelial NOS (NOS3) and inducible NOS (NOS2), in the regulation of hepatic *Cyp1a*, *Cyp1b*, *Cyp3a*, *Cyp2c*, *Cyp2d* and *Cyp2e1* by OLE in WT mice appears to be less significant, because although OLE induced *Nos3* and *Nos2* mRNA expression (Fig. 7A and 7B), this effect was not followed by inhibition of the afore-mentioned CYPs (Hara and Adachi, 2002). Instead, OLE induced the synthesis of these CYPs (Fig 1A-1C, Fig. 2A-2C and Fig. 3A-3C), indicating that the NOS inhibiting effect on CYPs was probably overlapped by the inducing effect of other regulatory factors.

At signal transduction level, OLE increased FOXO1 and JNK phosphorylation, along with that of p70 in the livers of WT mice (Fig. 8). These findings indicate that the OLE-induced activation of PI3k/AKT/FOXO1, JNK and AKT/p70 pathways (Fig. 8) is profoundly related to the induction of CYPs that belong to the CYP1A, CYP1B, CYP3A, CYP2C, CYP2D and CYP2E subfamilies. Activation of ERK could also participate in the regulation of CYP induction by OLE (Fig. 8).

Comparatively, OLE and FEN at the doses given, induced hepatic protein expression of CYP1A1, CYP2C and CYP3A to a similar extent, whereas the induction of CYP1A2, CYP1B1, CYP2D and CYP2E1 by FEN was markedly higher than that by OLE (Figs 1-3 and Fig. 6).

Discussion

There is an accumulating amount of evidence that supports the beneficial effects of OLE in preservation of good health and in the outcome of various disease states, including those related to the cardiovascular system (Ahamad et al., 2019; Andreadou et al., 2006; Araki et al., 2019; Lockyer et al., 2017; Vogel et al., 2014), which are mainly attributed to the drug's anti-inflammatory properties (Malliou et al., 2018), to its effects on lipid homeostasis (Andreadou et al., 2014; Araki et al., 2019; Lockyer et al., 2017; Malliou et al., 2018) and the protection of myocardium in conditions related to ischemia (Andreadou et al., 2014). Based on these OLE-related beneficial effects, the pharmaceutical industry produced several food supplements and herbal medicines containing the drug, which are available in the market without a prescription, a potentially high risk practice for health and disease.

It is well documented that cytochromes belonging to CYP families 1-3 are involved in the metabolism of a plethora of diverse endogenous and exogenous compounds, such as drugs, pre-carcinogens, carcinogens, environmental pollutants, food additives, prostaglandins, fatty acids, lipid hydroperoxides, steroid hormones, biogenic amines and numerous other xenobiotics (Cribb et al., 2005; Gonzalez, 2005; Gonzalez and Gelboin, 1994b; Gonzalez and Yu, 2006; Guengerich, 2003; Ingelman-Sundberg, 2004b; Xu et al., 2005), in order to increase their water solubility and

prepare them for the subsequent conjugation and elimination (Gonzalez, 2005). Their biotransformation takes place primarily in the liver and usually, results in inactive metabolites, or in some cases, in active molecules, a process that usually depends on the structure of the parent compound. These active metabolic products may induce several serious toxic manifestations, such as teratogenesis, carcinogenicity, cell death, and oxidative stress, among others (Cribb et al., 2005; Gonzalez, 2005; Gonzalez and Gelboin, 1994b; Gonzalez and Yu, 2006; Guengerich, 2003; Ingelman-Sundberg, 2004a; Konstandi, 2013; Xu et al., 2005). The findings of the current study clearly showed that OLE induced the expression of several CYPs in the liver of WT mice, including CYP1A1, CYP1A2, CYP1B1, CYP3A14, CYP3A25, CYP2C29, CYP2C44, CYP2D22 and CYP2E1 via activation of major nuclear transcription and other cellular factors, such as AHR, CAR, RXR and PXR (Dalton et al., 2000; Daskalopoulos et al., 2012a; Harkitis et al., 2015). Induction of CYPs is a part of the regulatory mechanisms aimed at maintaining homeostasis. It is important for the adaptation of the organism to a modified biological and chemical internal milieu and environment. Nonetheless, it is not always feasible to predict the extent of the effect of CYP induction on drug efficacy and toxicological risk assessment, determinant parameters in clinical drug therapy (Pelkonen et al., 2008). According to the regulatory standards of the Food and Drug Administration (FDA), a drug is considered as CYP inducer with clinical significance if the fold change of *CYP* mRNA expression relative to the vehicle control is ≥ 2 -fold and then, is efficient to accelerate the metabolism of CYP-substrates (Fahmi and Ripp, 2010); <https://www.fda.gov/drugs/drug-interactions-labeling/drug-development-and-drug-interactions-table-substrates-inhibitors-and-inducers>). It is considered that induction of CYP1A2, CYP3A14, CYP3A25, CYP2C29, CYP2C44, CYP2D22 and CYP2E1

may result in reduced concentrations of their drugs-substrates in blood, potentially below their therapeutic levels and therefore, in failure of pharmacotherapy (Dalton et al., 2000; Kawajiri and Fujii-Kuriyama, 2007; Konstandi, 2013; Okey, 1990; Zhang et al., 1999), whereas induction of CYP1A1, CYP1A2 and CYP1B1 is followed by bioactivation of the major groups of pre-carcinogens, the polycyclic aromatic hydrocarbons, polycyclic arylamines and aflatoxin B1, to DNA-binding metabolites (Cheng and Morgan, 2001; Kawajiri, 1999; Mulder et al., 2001; Pasanen and Pelkonen, 1994). In contrast to the present findings related to the induction of CYP3A14/25 by OLE, previous *in vitro* studies indicated that OLE and its metabolite hydroxytyrosol, inactivated androstenedione 6 β -hydroxylase (CYP3A4-dependent) activity in microsomal preparations of human liver (Stupans et al., 2001; Stupans et al., 2000; Zhou et al., 2007). This discrepancy could be explained primarily, on the basis of a species-specific regulation of the main CYPs (Konstandi et al., 2020; Visioli et al., 2002a) and the different experimental approaches followed in these studies. Current *in vivo* study evaluated the OLE effect on CYP3A14/25 in mouse liver microsomes, whereas the *in vitro* studies mentioned above used human liver microsomes (Stupans et al., 2001; Stupans et al., 2000).

In order to investigate the mechanisms underlying the OLE-induced upregulating effect on CYPs, the role of the nuclear transcription factor, PPAR α that regulates a variety of genes encoding CYPs was assessed (Tauber et al., 2020). For this purpose, WT and *Ppara*-null mice were employed, which received OLE in their diet, for six weeks. The findings indicated that, in contrast to what happened in WT mice, OLE did not induce the expression of CYP1A1, CYP1A2, CYP1B1, CYP3A14, CYP3A25, CYP2C29, CYP2C44, CYP2D22 and CYP2E1 in the liver of *Ppara*-null mice. It is apparent that the OLE-mediated induction of these CYPs in

the liver of WT mice is mediated by the drug's stimulating effect on PPAR α activation. In order to further evaluate the involvement of PPAR α in the regulation of CYPs and in particular, of CYP induction, WT mice were treated with the selective PPAR α agonist, fenofibrate (Ghonem et al., 2015; Hu et al., 2019), which induced CYP1A1, CYP1A2, CYP1B1, CYP2C, CYP2D, CYP2E1 and CYP3A protein expression in their livers. No similar effects were observed in the liver of *Ppara*-null mice following treatment with fenofibrate, and these findings confirm the role of PPAR α in CYP induction. This hypothesis is also supported by other studies reporting the up-regulating effects of PPARs on several CYPs, and in particular of PPAR α on CYPs displaying epoxigenase activities on polyunsaturated fatty acids, including isozymes that belong to CYP1A, CYP3A, CYP2C and CYP2E subfamilies (Tauber et al., 2020; Thomas et al., 2013). Interestingly though, findings from the current and previous studies (Shi et al., 2017) showed that basal CYP expression levels in the liver of *Ppara*-null mice are markedly higher than those in WT mice. This observation indicates a distinct role of PPAR α in the regulation of constitutive and inducible CYP expression. It is well established that CYP induction and constitutive expression are regulated by distinct mechanisms (Hahn et al., 2009; Nebert, 2000; Zanger and Schwab, 2013), including various transcription factors, co-activators, and co-repressors, with both positive and negative regulatory roles that cross-talk between several regulatory pathways (Pelkonen et al., 2008). In this regard, the diverse role of PPAR α in CYP regulation at basal and induced states may be attributed in the cross-talk between this transcription factor and AhR, CAR, PXR and/or other transcription factors with significant regulatory roles in CYP induction by xenobiotics (Honkakoski and Negishi, 2000; Pelkonen et al., 2008; Zanger and Schwab, 2013). This hypothesis is supported by the finding that several functional

PPAR α -binding regions within the CYP promoters were detected (Makia and Goldstein, 2016; Oshida et al., 2016a; Thomas et al., 2013; Yao et al., 2007) (Oshida et al., 2016b). It should be noted that the mechanism underlying the PPAR α -mediated CYP induction by OLE could also include modifications at PPAR α phosphorylation status. It is well established that ligand-induced PPAR α activation is mediated by increased expression of this nuclear receptor at transcriptional and protein level along with alterations at its phosphorylation state (Ning et al., 2016; Tamasi et al., 2008)((Barger et al., 2001; Passilly et al., 1999).

The nuclear transcription factor, HNF4 α , profoundly participates in the OLE-induced upregulation of the above mentioned CYPs, because OLE induced hepatic *Hnf4a* expression in WT mice. It is known that the transcription factors, HNF4 α , RXR and HNF1 α (Weltman et al., 1998; Wiwi and Waxman, 2004), along with FOXO1 and the nuclear receptors, CAR and PXR (Kodama et al., 2004), belong to a complex cross-talk mechanism displaying central regulatory roles in the expression of various CYPs.

In CYP regulation there is also an interplay between PPAR α and inducible nitric oxide synthase (iNOS or NOS2), efficient enough to modify *CYP* expression. In particular, it has been reported that PPAR α ligands reduced the LPS-induced iNOS expression and subsequent nitric oxide (NO) synthesis in macrophages by increasing the proteasome pathway mediated iNOS protein degradation (Paukkeri et al., 2007). Notably, NO displays down-regulating effects on several CYP genes, including CYP1A1/2, CYP2B1/2, CYP2D6, CYP2E1 and CYP3A4, but the underlying mechanism of this regulation remains blurred (Gergel et al., 1997; Hara and Adachi, 2002; Wink et al., 1993). Current findings indicated that the OLE-induced effect on NOS has a weak impact on CYPs compared to other regulatory factors. It appears

that the up-regulating effect of OLE on NOS, which should be followed by CYP downregulation (Gergel et al., 1997; Hara and Adachi, 2002; Wink et al., 1993), was overlapped by the upregulating effects of other transcription factors. Further analysis indicated that at signal transduction level, activation of the PI3k/AKT/FOXO1, JNK, AKT/p70S6K and ERK related signaling pathways profoundly participate in CYP induction by OLE (Kim and Novak, 2007). It is well documented that upon activation, AKT stimulates the phosphorylation of the nuclear transcription factor, forkhead box O1 (FOXO1), which in turn translocates into the cytoplasm. This process is usually followed by termination of FOXO1 transcriptional activity in CYP genes. But then, the OLE-mediated activation of c-Jun N-terminal kinases (JNK) promotes the nuclear localization of FOXO1, an effect that restores its transcriptional activity in CYP genes and counteracts the downregulating effect of AKT (Daskalopoulos et al., 2012a; Hay, 2011). Current findings indicated that OLE decreased JNK phosphorylation in the liver of *Ppara*-null mice, an effect that likely participates in the OLE-mediated downregulation of CYPs in these mice.

Taken together the above data clearly show the upregulating effect of OLE on various CYPs encoding the main drug-metabolizing enzymes of Phase I, an effect that is apparently mediated by PPAR α activation. Current and previous findings indicate that diverse and distinct mechanisms participate in the regulation of constitutive and inducible CYP expression by PPAR α agonists. Apparently, PPAR α displays a negative regulatory role in constitutive CYP expression and a positive role in CYP induction by OLE. Although, it is not always feasible to accurately predict the clinical impact of the OLE-induced effect on CYP3A4, CYP3A5, CYP2C9, CYP2C19, CYP2D6, CYP2E1, CYP1A1, CYP1A2 and CYP1B1, under certain conditions it could modify the pharmacokinetic profile of drug-substrates of these

CYPs and subsequently, affect the outcome of pharmacotherapy and the severity of drug-related adverse reactions. Although these data come from a preclinical study, and cannot be extrapolated directly to humans, they underscore the necessity of taking into consideration the consumption of herbal medicines and food supplements containing drugs, such as OLE that may affect the pharmacokinetic profiles of co-administered drugs. This parameter should be considered predominately, in multiple drug therapeutic schemes, in particular those of vital significance for the patient, or in drugs with narrow therapeutic windows or with severe side effects. This concern is of particular importance, because herbal medicines and food supplements are widely used and freely available in the market. All of these concerns indicate that further investigation, pharmacovigilance, better regulatory control and seek of advice from health professionals is essential to ensure the safety when using herbal medicines and food supplements.

Acknowledgments

This study has been funded by the European Union (European Regional Development Fund- ERDF) and Greek national funds coming from the Operational Program “THESSALY- MAINLAND GREECE AND EPIRUS-2007-2013” of the National Strategic Reference Framework (NSRF 2007-2013, Grant 346985/80753).

Author contributions

Participated in research design: Konstandi, Gonzalez

Conducted experiments: Malliou, Andriopoulou, Konstandi, Skaltsounis

Performed data analysis: Konstandi, Andriopoulou, Malliou

Wrote or contributed to the writing of the manuscript: Konstandi, Gonzalez

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Footnotes

Funding: This study was funded by the European Union (European Regional Development Fund- ERDF) and Greek national funds coming from the Operational Program “THESSALY- MAINLAND GREECE AND EPIRUS-2007-2013” of the National Strategic Reference Framework (NSRF 2007-2013, Grant 346985/80753). We certify that there is no involvement of the funding sources in the research conduct and/or preparation of this article, in study design, in the collection, analysis and interpretation of data, in the writing of the article, and the decision to submit the article for publication in DMD. The authors also state that there is no conflict of interest.

Figure legends

Figure 1. PPAR α -mediated regulation of hepatic CYP1A1, CYP1A2 and CYP1B1. Assessment of the effect of OLE, a PPAR α agonist, on (A) CYP1A1, (B) CYP1A2 and (C) CYP1B1 protein levels employing western blotting, and relative gene expression with quantitative PCR analysis. In wild type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein; Values are expressed as mean \pm SE, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. wild-type mice (n=11), *Ppara* null mice (n=10). Numbers below bands in western blot captures show the ratio of the density of the sample band to that of α -tubulin.

Figure 2. PPAR α -mediated regulation of hepatic CYP2C29/2C44 and CYP2D22. Assessment of the effect of OLE, a PPAR α agonist, on (A) CYP2C29, (B) CYP2C44 and (C) CYP2D22 protein levels employing western blotting, and on relative gene expression with quantitative PCR analysis. The OLE-mediated effect on 1'-bufuralol hydroxylation was performed using HPLC. In wild-type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein; Values are expressed as mean \pm SE, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. wild-type mice (n=11), *Ppara* null mice (n=10). Numbers below bands in western blot captures show the ratio of the density of the sample band to that of α -tubulin.

Figure 3. PPAR α -mediated regulation of hepatic CYP2E1 and CYP3A14/3A25.

Assessment of the effect of OLE, a PPAR α agonist, on (A) CYP2E1, (B) CYP3A14 and (C) CYP3A25 protein level employing western blotting, and on relative gene expression with quantitative PCR analysis. In wild-type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein; Values are expressed as mean \pm SE, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. wild-type mice (n=11), *Ppara*-null mice (n=10). Numbers below bands in western blot captures show the ratio of the density of the sample band to that of α -tubulin.

Figure 4. Assessment of the effect of OLE, a PPAR α agonist, on various factors involved in CYP1A and CYP1B regulation (Matsunaga et al., 1990). (A) aryl hydrocarbon receptor (Ahr), (B) aryl hydrocarbon receptor repressor (AhRR), and (C) heat shock protein 90 (HSP90) relative mRNA expression was estimated with quantitative PCR analysis. In wild-type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein; Values are expressed as mean \pm SE, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. wild-type mice (n=11), *Ppara*-null mice (n=10).

Figure 5. Assessment of the effect of OLE, a PPAR α agonist, on various factors involved in CYP3A and CYP2C regulation (Konstandi et al., 2006). (A) Pregnane X receptor (*Pxr*), (B) Constitutive androstane receptor (*Car*), (C) Retinoid X receptor α (*Rxra*) and (D) Retinoid X receptor β (*Rxrb*) relative mRNA expression with quantitative PCR analysis. In wild type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein; Values are expressed as mean \pm SE, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$. wild-type mice (n=11), *Ppara* null mice (n=10). Numbers below bands in western blot captures show the ratio of the density of the sample band to that of Histone H3.

Figure 6. *In vivo* assessment of the effects of FEN, a selective PPAR α agonist, on CYP protein expression by Western blotting using microsomal proteins extracted from liver samples of wild type and PPAR α -null mice. FEN-induced hepatic CYP1A1, CYP1A2, CYP1B1, CYP2C, CYP2D, CYP2E1 and CYP3A protein expression. In wild type and *Ppara*-null mice, comparisons were made between Controls and FEN-treated mice. C: Control, FEN: fenofibrate. The samples in the western blot captures are representative of other three that were analyzed in separate blots. Numbers below bands in western blot captures show the ratio of the density of the sample band to that of GAPDH.

Figure 7. Assessment of the effect of OLE, a PPAR α agonist, on (A) endothelial NOS (*Nos3*) and (B) inducible nitric oxide synthase (*Nos2*) relative mRNA expression with quantitative PCR analysis. In wild type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein; Values are expressed as mean \pm SE, ** $p < 0.01$, *** $p < 0.001$. wild-type mice (n=11), *Ppara* null mice (n=10).

Figure 8. *In vivo* assessment of the effects of OLE, a PPAR α agonist, on several signal transduction pathways involved in CYP regulation by Western blotting using total cellular proteins extracted from liver samples of mice. OLE-induced activation of FOXO1, JNK, p70 and ERK but not to p38-MAPK. In wild type and *Ppara*-null mice, comparisons were made between Controls and OLE-treated mice. C: Control, OLE: oleuropein. The samples in the western blot captures are representative of other three that were analyzed in separate blots. Numbers below bands in western blot captures show the ratio of the density of the sample band to that of α -tubulin.

Graphic abstract. Oleuropein (OLE) induced the expression of various important drug metabolizing CYP genes in the liver of mice, including *Cyp1a1*, *Cyp1a2*, *Cyp1b1*, *Cyp2c29*, *Cyp2c44*, *Cyp2d22*, *Cyp2e1*, *Cyp3a14* and *Cyp3a25*. OLE acts as a ligand of the nuclear transcription factor, PPAR α , and the complex binds to peroxisome proliferator activated receptor responsive element (PPRE) in the promoter of the CYP gene, thus stimulating its transcription. PXR: pregnane X receptor is a drug activated transcription factor, involved both in constitutive and/or inducible expression of CYP3A CYP2C and CYP2A cytochromes; RXR: retinoid X

receptor and CAR: constitutive androstane receptor, are two drug activated transcription factors involved in both constitutive and/or inducible *CYP3A* and *CYP2C* expression Ahr: Aryl Hydrocarbon Receptor and ARNT: Aryl Hydrocarbon Receptor Nuclear Translocator are two transcription factors regulating both, constitutive and inducible *CYP1A* and *CYP1B* expression (Daskalopoulos et al., 2012a; Daskalopoulos et al., 2012b; Harkitis et al., 2015; Nebert et al., 2013).

Table 1: Oligonucleotide sequences used as primers for quantitation of gene mRNA levels through quantitative PCR assays

Gene	Primer sequence
<i>Ppara</i>	F: CAGTGGGGAGAGAGGACAGA R: AGTTCGGAACAAGACGTTG
<i>Cyp2d22</i>	F: ACCGGTAAAGGTAGCTGGAGT R: CATAGGGCCTGGAGGGTAGT
<i>Hnf4a</i>	F: CGGAGCCCCTGCAAAGT R: ACTATCCAGTCTCACAGCCCATTG
<i>Cyp2e1</i>	F: TGGTCCTGCATGGCTACAAG R: CGGGCCTCATTACCCTGTTT
<i>Cyp2c29</i>	F: TGTTACAAACCCCCGTGACT R: GGATGTGGATAAAGACCTGAGAC
<i>Cyp2c44</i>	F: CCTAAAGGCTCTGGTGGAGC R: GAAACAAATGCCCCACGTGCT
<i>Cyp3a14</i>	F: GGCCCAGTGGGGATAATGAG R: GGTGCCTTATTGGGCAGAGT
<i>Cyp3a25</i>	F: TAGAAACCTGGGTGCTGCTG R: GGATGTGGATAAAGACCTGAGAC
<i>Pxr</i>	F: AAGAAGCAGACTCTGCCTTGGA R: GTGGTAGCCATTGGCCTTGT
<i>Car</i>	F: CCTCTTCTCCCCTGGTTTCTG R: TCATTGCCACTCCCAAGCTC
<i>Rxra</i>	F: CAGTACGCAAAGACCTGACCTACA R: GTTCCGCTGTCTCTTGTCGAT
<i>Rxrb</i>	F: AAGTGTCTGGAGCACCTGTTCTT R: CTCCATGAGGAAGGTGTCAATG
<i>Cyp1a1</i>	F: GAAGTGGAAGGGCATAGGCAG R: GGCCAAAGCATATGGCACAG
<i>Cyp1a2</i>	F: ACTTCGAACCAGTCAGCCAG R: GTGCTTGAACAGGGCACTTG
<i>Cyp1b1</i>	F: CCAGCTTTTTGCCTGTCACC R: TGCACTGATGAGCGAGGATG
<i>Ahr</i>	F: TTCAGAACTGACTCCACCGC R: CCGGGTGTGATATCGGGAAG
<i>Ahrr</i>	F: AGTGTACATACGCCGGTAGG R: CAAGACTGGTGCCACAATGC
<i>Hsp90</i>	F: CAGACCATGGTGAGCCCATT R: TCAACCACACCGCGGATAAA

<i>Nos3</i>	F: GCAGAAGAGTCCAGCGAACA R: GGCAGCCAAACACCAAAGTC
<i>Nos2</i>	F: GTGTTCCACCAGGAGATGTTG R: CTCCTGCCCACTGAGTTCGTC

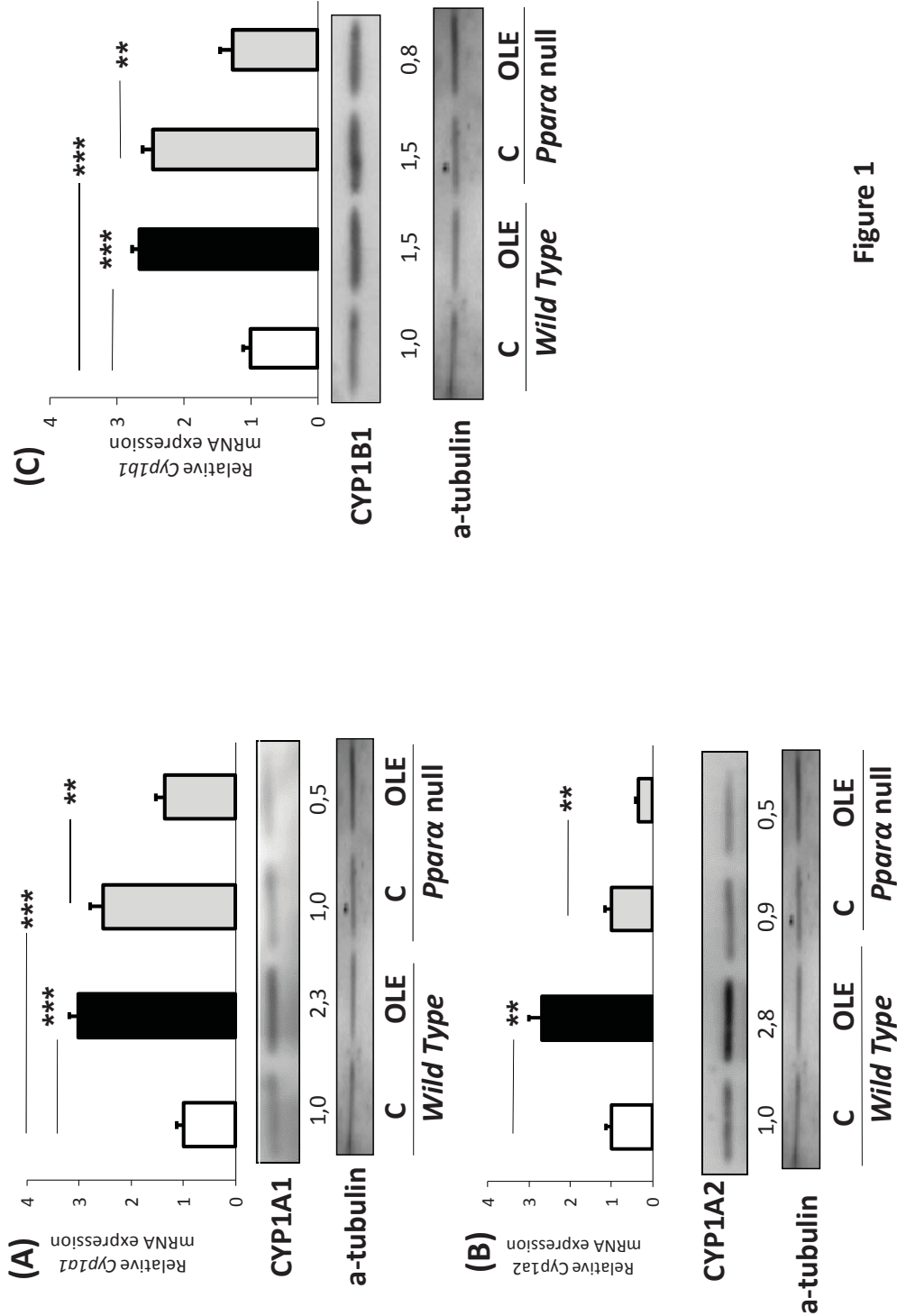


Figure 2

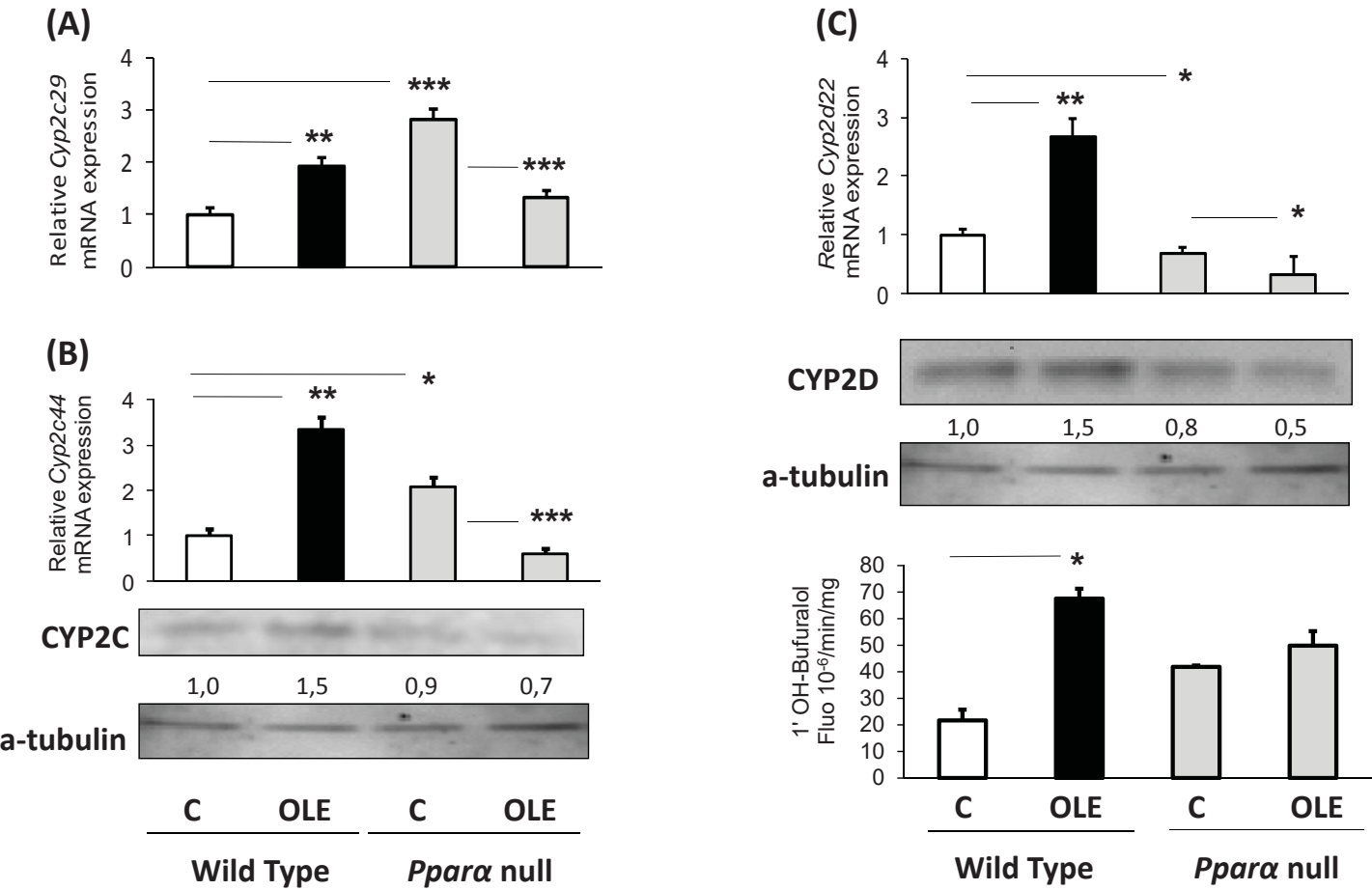
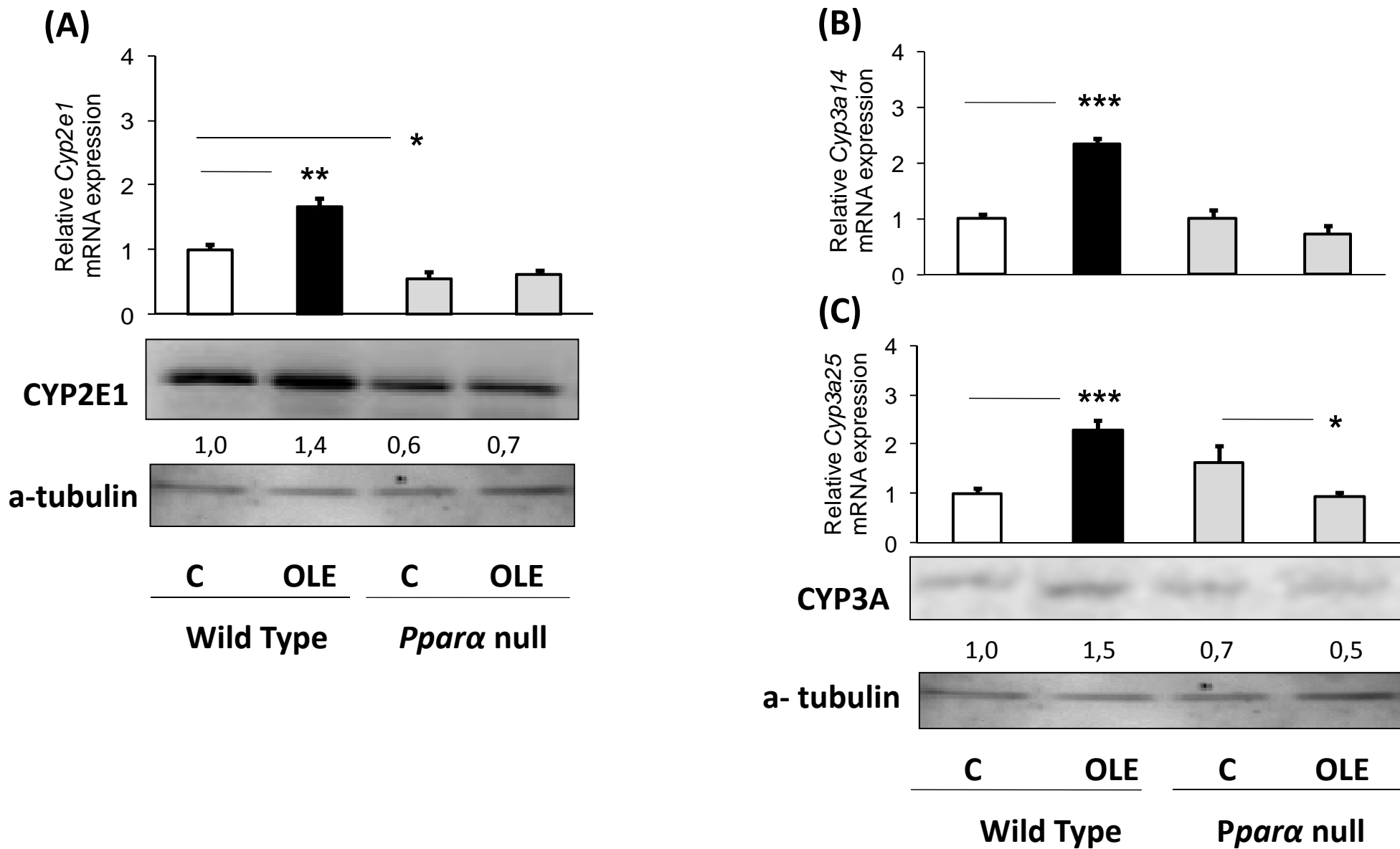


Figure 3



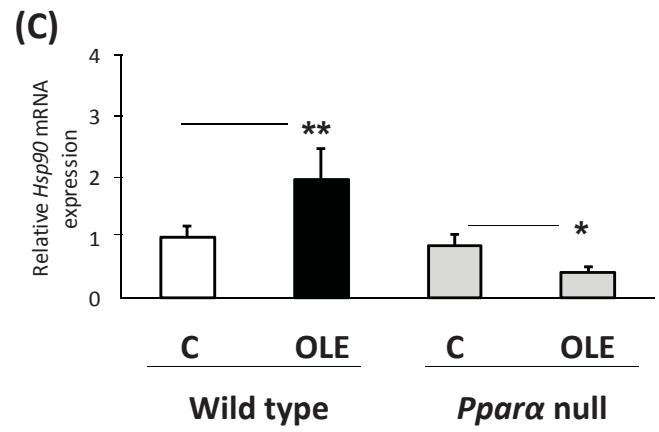
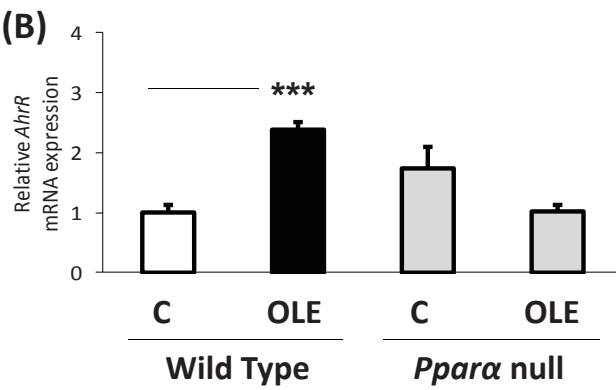
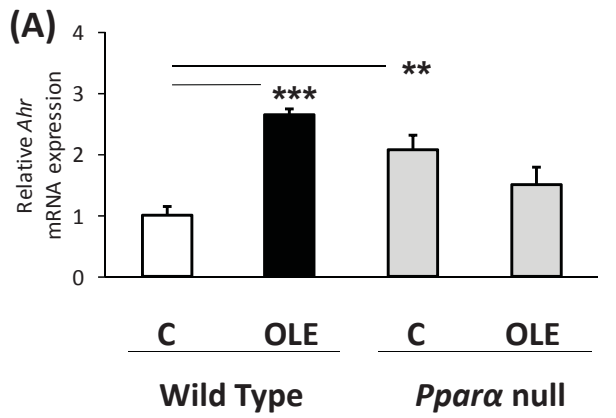


Figure 4

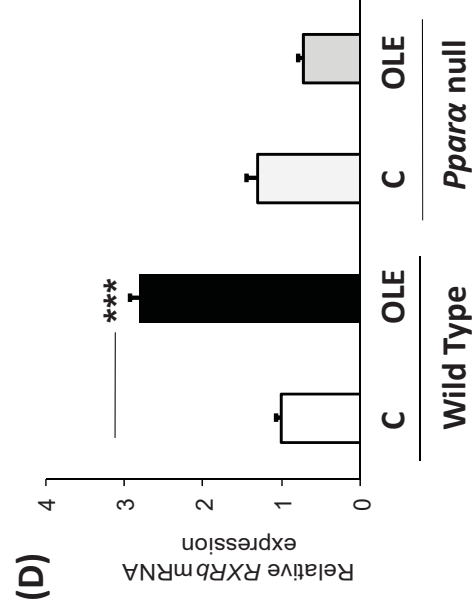
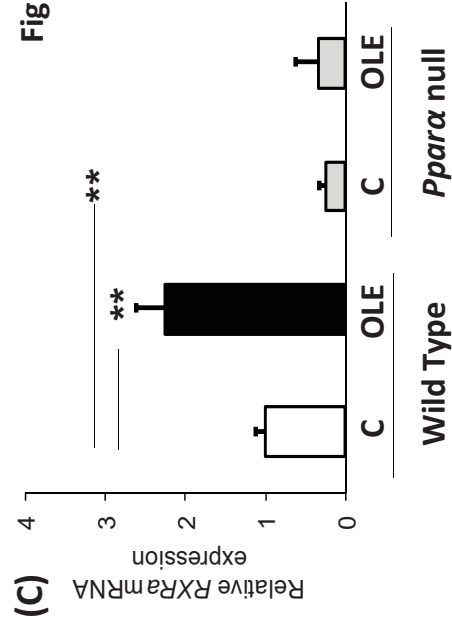
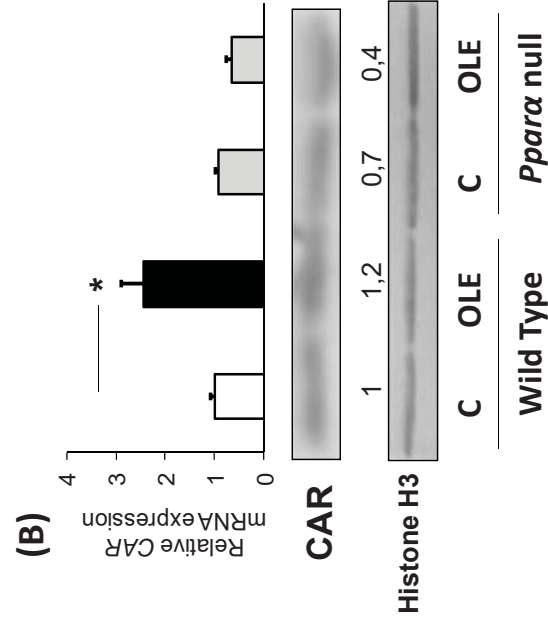
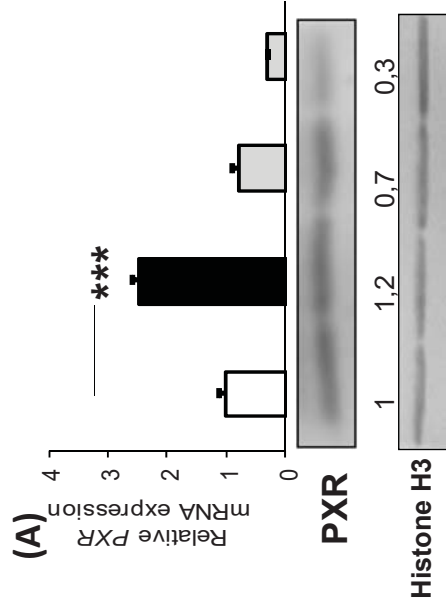


Figure 5

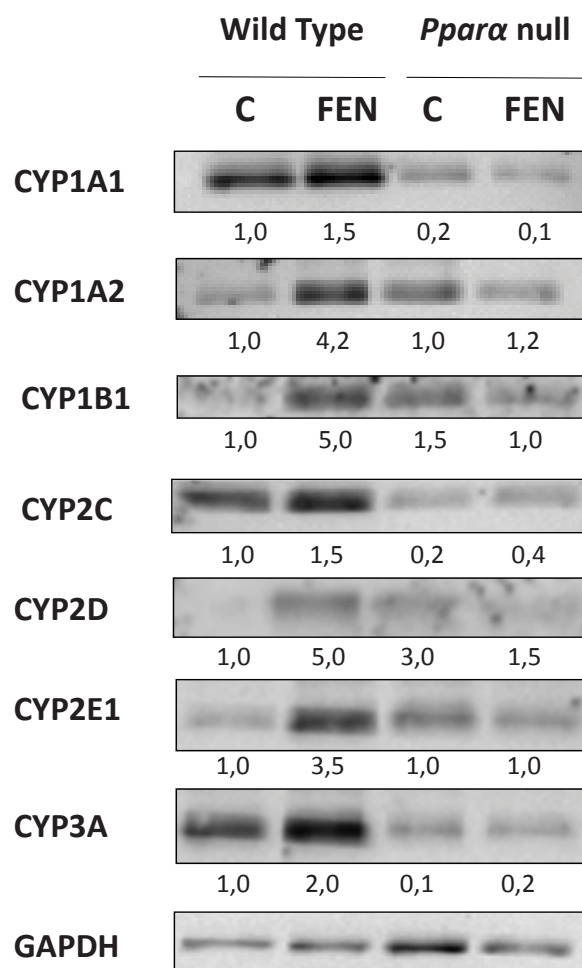


Figure 6

Figure 7

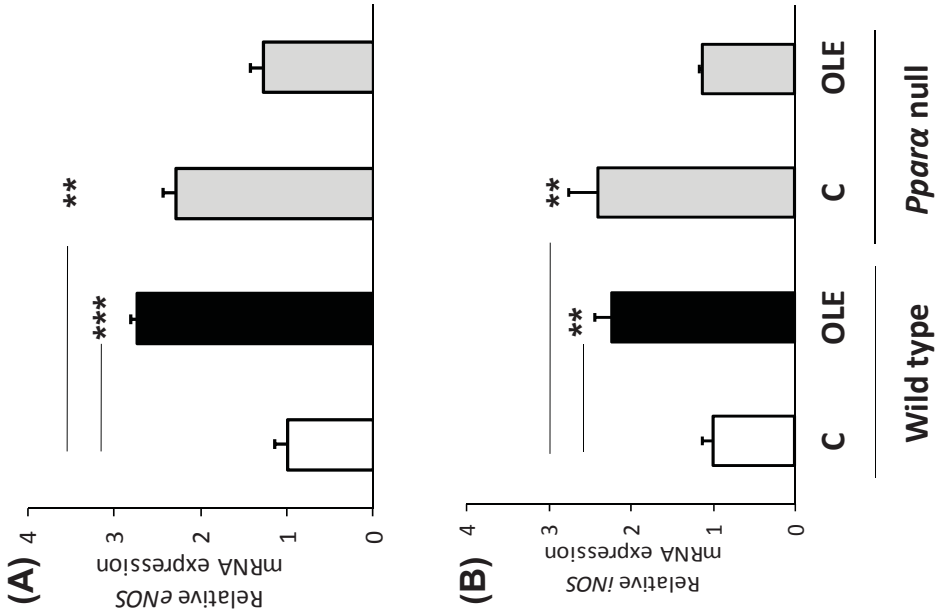
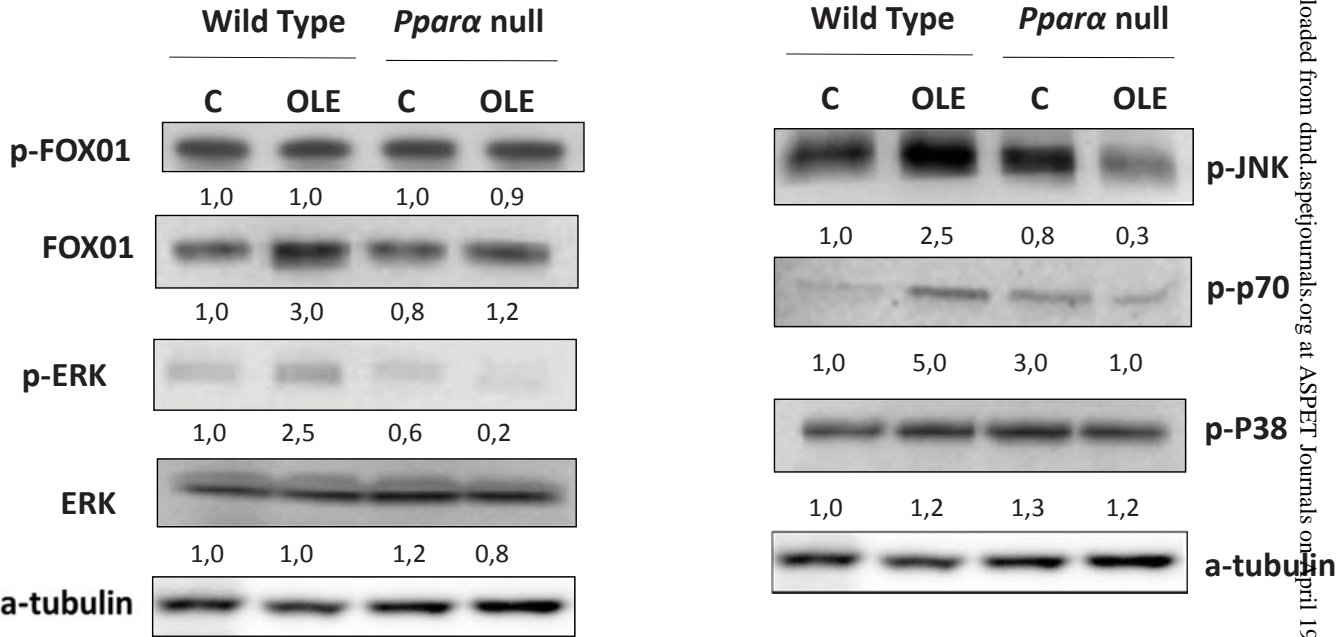
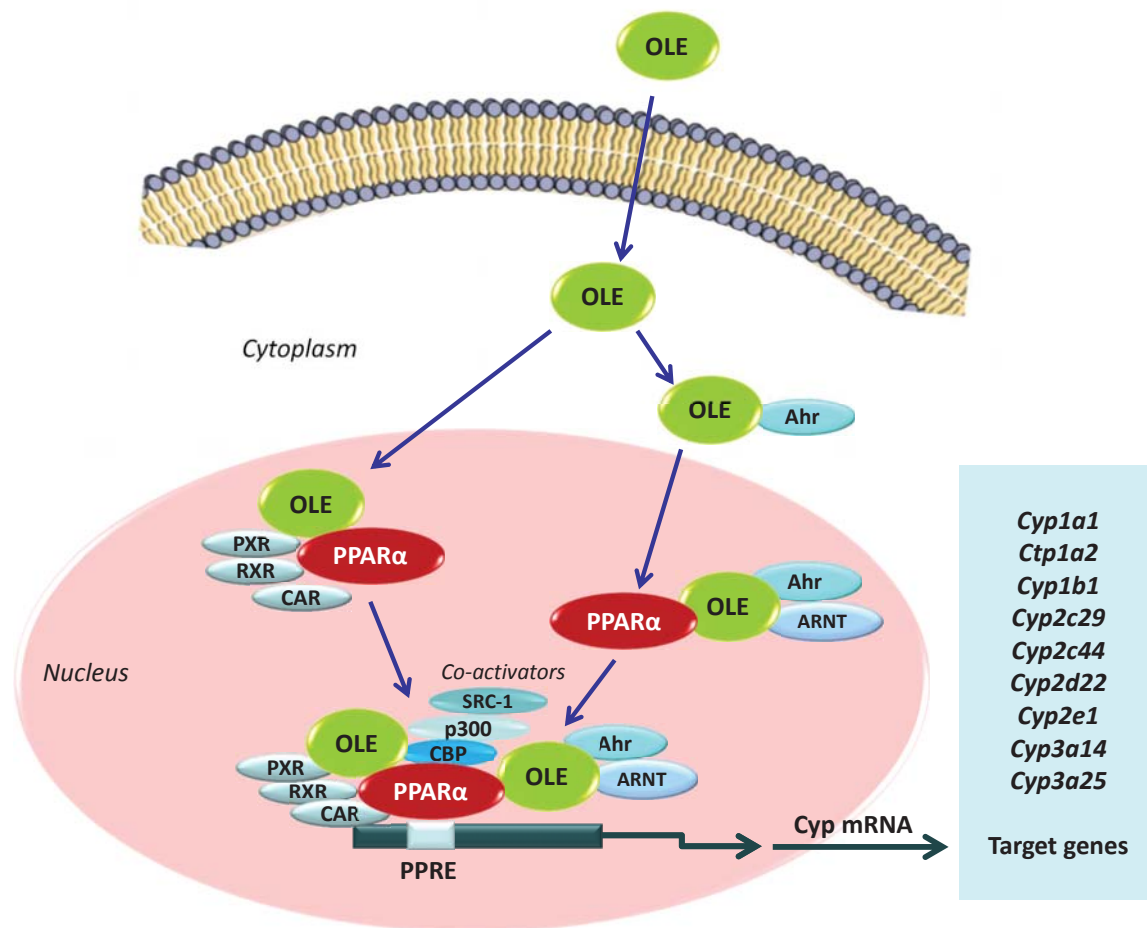


Figure 8



Total cellular proteins



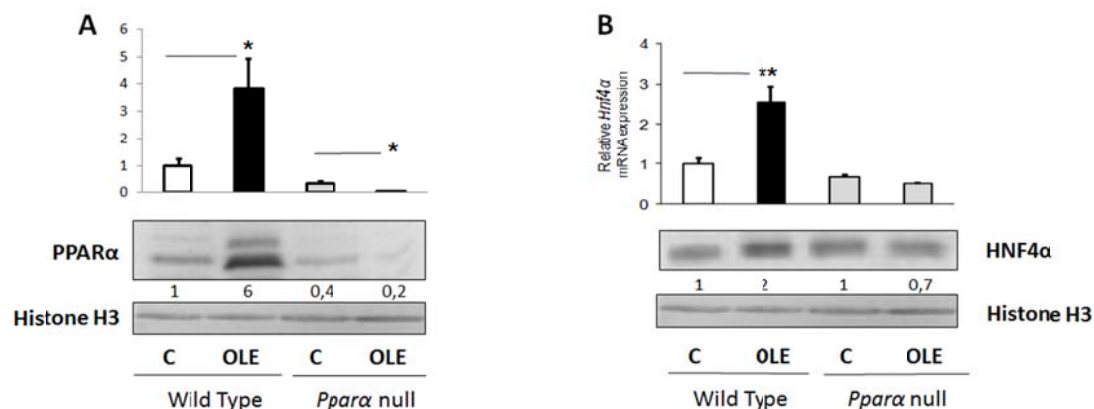
Oleuropein-induced acceleration of CYP-catalyzed drug metabolism: central role for nuclear receptor PPAR α

Foteini Malliou^{1#}, Christina E. Andriopoulou^{1#}, Frank J. Gonzalez², Aristeidis Kofinas¹, Alexios-Leandros Skaltsounis³, Maria Konstandi^{1*}

¹University of Ioannina, School of Health Sciences, Faculty of Medicine, Department of Pharmacology, Ioannina GR-45110, Greece;; ²Laboratory of Metabolism, National Cancer Institute, National Institutes of Health, Bethesda 20892, Maryland, USA; ³Faculty of Pharmacy, School of Health Sciences, University of Athens, Greece

Journal: Drug Metabolism & Disposition

MS #DMD-AR-2020-000302



Supplemental Figure 1. *In vivo* assessment of the effects of OLE, a PPAR α agonist, on *Ppara* and *Hnf4a* mRNA expression using quantitative PCR analysis. PPAR α and HNF4 α protein levels were analyzed by western blot. In wild type and *Ppara*-null mice, comparisons took place between Controls and OLE-treated mice. C: Control, OLE: oleuropein, WT: wild type; Values are expressed as mean \pm SE, ** $p < 0.01$, *** $p < 0.001$. wild type mice (n=11), *Ppara*-null mice (n=10).