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RESEARCH ARTICLE

The anti-angiogenic herbal extracts Ob-X from *Morus alba*, *Melissa officinalis*, and *Artemisia capillaris* suppresses adipogenesis in 3T3-L1 adipocytes

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Abstract

Context: Growing adipose tissue is thought to require adipogenesis, angiogenesis, and extracellular matrix (ECM) remodeling. Close examination of developing adipose tissue microvasculature reveals that angiogenesis often precedes adipogenesis. Since our previous study demonstrated that Ob-X, the anti-angiogenic herbal composition composed of *Melissa officinalis* L. (Labiatae), *Morus alba* L. (Moraceae), and *Artemisia capillaris* Thunb. (Compositae), reduced adipose tissue mass in obese mice, we hypothesized that adipogenesis can be inhibited by Ob-X.

Objective: To investigate the effects of the anti-angiogenic herbal extracts Ob-X on adipogenesis in 3T3-L1 adipocytes.

Materials and methods: After differentiated 3T3-L1 adipocytes were treated with Ob-X, we studied the effects of Ob-X on triglyceride accumulation and expression of genes involved in adipogenesis, angiogenesis, and ECM remodeling.

Results: Treatment of cells with Ob-X inhibited lipid accumulation and adipocyte-specific gene expression caused by troglitazone or monocyte differentiation-inducing (MDI) mix. Ob-X reduced mRNA levels of angiogenic factors (vascular endothelial growth factor-A, -B, -C, -D, and fibroblast growth factor-2) and matrix metalloproteinases (MMPs; MMP-2 and MMP-9), whereas it increased mRNA levels of angiogenic inhibitors [(thrombospondin-1, tissue inhibitor of metalloproteinase-1 (TIMP-1), and TIMP-2)] in differentiated cells. MMP-2 and MMP-9 activities were also decreased in Ob-X-treated cells.

Discussion and conclusion: These results suggest that the anti-angiogenic herbal composition Ob-X inhibits differentiation of preadipocytes into adipocytes. These events may be mediated by changes in the expression of genes involved in lipogenesis, angiogenesis, and the MMP system. Thus, by reducing adipogenesis, anti-angiogenic Ob-X provides a possible therapeutic approach for the prevention and treatment of human obesity and its related disorders.

Keywords: Angiogenesis, MMP, 3T3-L1, anti-angiogenic plants

Introduction

Obesity and the related metabolic disorders, such as dyslipidemia, atherosclerosis, and type 2 diabetes, have become global health problems. Obesity is characterized by increased adipose tissue mass that results from both increased fat cell number (hyperplasia) and increased fat cell size (hypertrophy) (Couillard et al., 2000). Fat mass can be regulated by various factors including adipogenesis, angiogenesis, and remodeling of the extracellular matrix (ECM) (Crandall et al., 1997).

During adulthood, most tissues normally do not grow and the supporting vasculature is quiescent. Exceptionally, adipose tissue can grow throughout the life span and is highly vascularized. In growing adipose tissue, each adipocyte is nourished by an extensive capillary network (Silverman et al., 1988; Crandall et al., 1997; Bouloumié et al., 2002; Cao, 2010). Thus, it can be suggested that the growth and differentiation of adipocytes are angiogenesis-dependent.

Extensive changes in ECM remodeling have also been shown to occur during adipose tissue growth. The

matrix metalloproteinases (MMP) play important roles in the development of adipose tissue and microvessel maturation by modulating the ECM (Galaray et al., 1994; Bouloumié et al., 2001; Lijnen et al., 2002). In most cases, MMPs are expressed at very low levels, but expression is rapidly induced at times of active tissue remodeling associated with adipogenesis. Several lines of evidence suggest that endogenous and exogenous MMPs regulate adipogenesis (Kawaguchi et al., 1998; Bouloumié et al., 2001; Christiaens & Lijnen, 2006). During obesity, MMP expression is modulated in adipose tissue, and MMPs (e.g., MMP-2 and MMP-9) potentially affect adipocyte differentiation (Bouloumié et al., 2001; Maquoi et al., 2002; Chavey et al., 2003).

Adipocytes produce angiogenic factors, such as vascular endothelial growth factors (VEGFs) and fibroblast growth factor (FGF)-2, contributing to the formation of new blood vessels inside the fat pad (Claffey et al., 1992; Voros et al., 2005; Cao, 2007). VEGFs and FGF-2 stimulate proliferation and migration of endothelial cells (ECs) and enhance adipocyte differentiation (Carmeliet et al., 1996; Bikfalvi et al., 1997; Kawaguchi et al., 1998; Lijnen, 2008). Adipocytes also secrete several MMPs, including MMP-2 and MMP-9 (Bouloumié et al., 2001). Indeed, it is well established that degradation of the ECM represents the first step in the angiogenic process, and MMP-2 and MMP-9 have been shown to be necessary for this event (Sang, 1998).

We screened anti-angiogenic activity of water extracts from medicinal herbs and plants which have been used for a long period of time, and selected three herbal extracts *Melissa officinalis* L. (Labiatae), lemon balm, *Morus alba* L. (Moraceae) white mulberry, and *Artemisia capillaris* Thunb. (Compositae) injin, to make Ob-X (Kim et al., 2006). Moreover, Ob-X reduces adipose tissue mass in both genetically and nutritionally obese mice (Kim et al., 2010; Yoon and Kim, accepted). Accordingly, we hypothesized that adipocyte growth and differentiation can also be effectively regulated by Ob-X. Treatment of 3T3-L1 adipocytes with Ob-X reduced lipid accumulation and adipocyte-specific gene expression. The expression of angiogenic factors, MMPs, and their inhibitors in adipose tissue was markedly modulated by Ob-X. These studies suggest that Ob-X can inhibit adipogenesis by targeting adipocytes.

Materials and methods

Preparation of Ob-X

Ob-X was prepared from food grade aqueous extracts of the three herbs *M. officinalis* (Frutarom, Wädenswil, Switzerland), *M. alba* (Segae FL, Buyeo, Korea), and *A. capillaris* (Segae, FL) as previously described (Lee et al., 2008). The quality of each herbal extract in Ob-X was controlled by standardization with reference compounds by high-pressure liquid chromatography. The corresponding reference compounds were rosmarinic

acid (*M. officinalis*), 1-deoxynojirimycin (*M. alba*), and 6,7-dimethylesculetin (*A. capillaris*).

Induction of preadipocyte differentiation

Mouse 3T3-L1 cells (ATCC) were grown proliferated in Dulbecco's modified Eagle's medium (DMEM) containing 10% bovine calf serum (Invitrogen, Carlsbad, CA). After cells were kept confluent for 2 days, they were incubated in monocyte differentiation-inducing (MDI) medium (day 0) containing 0.5 mM 1-methyl-3-isobutyl-xanthin, 1 μ M dexamethasone, and 1 μ g/mL insulin in DMEM with 10% fetal bovine serum (FBS; Invitrogen). The cultures were continued for 2 more days to induce adipocyte differentiation. Thereafter, cells were cultured in DMEM with 10% FBS for the rest of the differentiation process. All other treatments were administered on day 0 to day 2 only, and the medium was changed every other day. Cells were stained at day 6 with Oil-red O and photographed.

Reverse transcription-polymerase chain reaction

Total cellular RNA from 3T3-L1 cells was prepared using the Trizol reagent (Gibco-Brl, Grand Island, NY). After 2 μ g of total RNA was reverse-transcribed using Moloney murine leukemia virus reverse transcriptase (MuLV-RT) and an antisense primer, cDNA was generated. The RNA was denatured for 5 min at 72°C and then immediately placed on ice for 5 min. Denatured RNA was mixed with MuLV-RT, MuLV-RT buffer, and a dNTP mixture, and incubated for 1 h at 42°C. Synthesized cDNA fragments were amplified by polymerase chain reaction (PCR) in an MJ Research Thermocycler (Waltham, MA). The PCR primers used for gene expression analysis are shown in Table 1. The cDNA was mixed with PCR primers, *Taq* DNA polymerase (Solgent, Daejeon, Korea), and a dNTP mixture. The reaction consisted of 30 cycles of denaturation for 1 min at 94°C, annealing for 1 min at 58°C, and elongation for 1 min at 72°C. The PCR products were analyzed by electrophoresis on a 1% agarose gel. Relative expression levels were presented as a ratio of target gene cDNA versus β -actin cDNA. PCR products were quantified from agarose gels using the GeneGenius (Syngene, Cambridge, UK).

Zymography

MMP activity was determined by gelatin zymography. 3T3-L1 Cells were extracted with 10 mM sodium phosphate buffer (pH 7.2) containing 150 mM NaCl, 1% Triton X-100, 0.1% SDS, 0.5% sodium deoxycholate, and 0.2% NaN_3 at 4°C. Protein concentrations were quantified using Bicinchonnic acid protein assay reagent (Pierce, Rockford, IL). Cell extracts were mixed with zymography sample buffer (63 mM Tris HCl, 10% glycerol, 2% SDS, 0.0025% bromophenol blue, pH 6.8) without heat denaturation. Culture medium of HT1080 cells was used for the molecular weight markers for MMP. Electrophoresis was performed on 10% SDS polyacrylamide gels containing 0.1% gelatin at

Table 1. PCR primers used for cDNA synthesis by RT-PCR.

Gene	Gene bank	Primer sequence	cDNA size (bp)
MMP-2	M84324	Forward: 5'-agatcttcttctcaagaccgggtt-3' Reverse: 5'-ggctggtcagtggttgggga-3'	224
MMP-9	NM_013599	Forward: 5'-tgcgaccacatcgaacttcg-3' Reverse: 5'-gagaagaagaaaacctcttgg-3'	683
TIMP-1	NM_001044384	Forward: 5'-ggcatcctctgtgtgtatcactg-3' Reverse: 5'-gtcatctgtatctcatacgtgg-3'	170
TIMP-2	NM_021989	Forward: 5'-gagatcaagcagataaagatg-3' Reverse: 5'-gaccagctccatccagaggc-3'	320
TSP-1	M62470	Forward: 5'-cctcattgtgtgtgactgagtaa-3' Reverse: 5'-ttgtgttctgtacataagaaac-3'	556
FGF-2	NM174056	Forward: 5'-aactacaactcaagcagaagagaga-3' Reverse: 5'-ttaagatcagctcttagcagacat-3'	293
VEGF-A	NM009505	Forward: 5'-gctctcttggtgactgga-3' Reverse: 5'-caccgcttggctgtcaca-3'	561
VEGF-B	NM_011697	Forward: 5'-gtcaacaactagtgtccag-3' Reverse: 5'-tgtctgggtgagcttaag-3'	407
VEGF-C	NM_053653	Forward: 5'-ccaaccagtaacaatcag-3' Reverse: 5'-attcacaggcacatttc-3'	461
VEGF-D	D89628	Forward: 5'-acctctacatctccaac-3' Reverse: 5'-tccagacttcttgcac-3'	385
PPAR γ	NM_013124	Forward: 5'-attctggccaccaacttcgg-3' Reverse: 5'-tggaagcctgatgctttatccca-3'	338
aP2	K02109	Forward: 5'-caaaatgtgtgatccttgg-3' Reverse: 5'-ctctccttggctcatgcc-3'	415
β -Actin	NM_00793	Forward: 5'-tggaatcctgtgcatccatgaac-3' Reverse: 5'-taaaacgcagctcagtaacagtcg-3'	348

aP2, adipocyte fatty acid binding protein; FGF, fibroblast growth factor; MMP, matrix metalloproteinases; PCR, polymerase chain reaction; PPAR γ , peroxisome proliferator-activated receptor γ ; RT, reverse transcription; TIMP, tissue inhibitor of metalloproteinase; TSP, thrombospondin; VEGF, vascular endothelial growth factor.

125 V. After electrophoresis, the gels were incubated in a renaturing buffer of 0.25% Triton X-100 for 30 min at room temperature, and equilibrated in developing buffer (50 mM Tris base, 40 mM HCl, 200 mM NaCl, 5 mM CaCl₂, 0.2% Brij-35) for 30 min at room temperature. The gels were then incubated in developing buffer overnight at 37°C. The gels were stained with 0.1% Coomassie Blue R-250 and destained with 10% acetic acid in 40% methanol.

Statistics

Unless otherwise indicated, all values are expressed as mean \pm SD. All data were analyzed by the unpaired Student's *t*-test for statistically significant differences between groups.

Results

Effects of Ob-X on 3T3-L1 differentiation

Accumulation of triglyceride droplets in 3T3-L1 cells was evident on the sixth day following 2 days of treatment with MDI differentiation mix (MDI) (Figure 1B) or the peroxisome proliferator-activated receptor γ (PPAR γ) agonist troglitazone (Figure 1D) compared with non-differentiated control (Figure 1A), as shown by positive staining with Oil-red O. Treatment of cells with Ob-X,

however, inhibited triglyceride accumulation; the percentage of differentiated cells was ~63% in MDI plus Ob-X-treated cells (Figure 1C) and 69% in troglitazone plus Ob-X-treated cells (Figure 1E) compared with cells treated with MDI or troglitazone alone, respectively. These results suggest that Ob-X effectively inhibits adipogenesis in 3T3-L1 cells.

Effects of Ob-X on mRNA expression of genes involved in lipogenic pathways

To quantify the changes in differentiation degree by Ob-X, we analyzed mRNA expression of genes involved in adipogenic and lipogenic pathways. PPAR γ and adipocyte fatty acid binding protein (aP2) mRNA levels were substantially upregulated by 61 and 62% at day 6 following MDI treatment compared with non-differentiated controls. Importantly, co-administration of Ob-X significantly downregulated MDI-induced PPAR γ and aP2 mRNA levels by 34 and 37%, respectively (Figure 2A). Similarly, troglitazone treatment increased PPAR γ and aP2 mRNA levels by 112 and 56% compared with non-differentiated controls, whereas Ob-X significantly decreased troglitazone-induced PPAR γ and aP2 mRNA levels by 30 and 16%, respectively (Figure 2B). Thus, Ob-X inhibited MDI- or troglitazone-induced expression of genes involved in adipocyte differentiation.

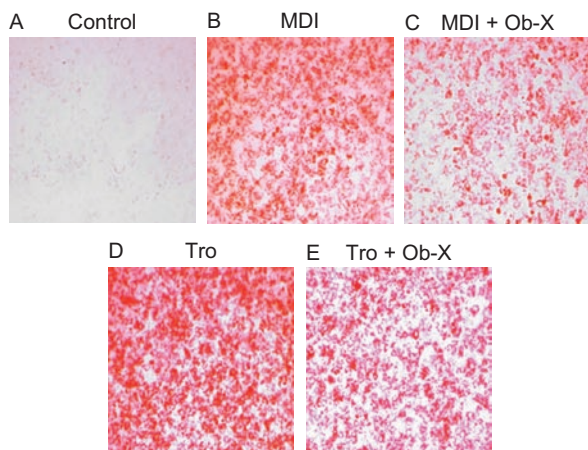


Figure 1. Effects of Ob-X on triglyceride accumulation in 3T3-L1 cells. 3T3-L1 preadipocytes were differentiated into mature adipocytes as described in "Materials and Methods". 3T3-L1 cells were treated with monocyte differentiation-inducing (MDI) differentiation mix (MDI), MDI plus 10 μ g/mL Ob-X, 10 μ M troglitazone (Tro), or 10 μ M Tro plus 10 μ g/mL Ob-X. At day 6 post-induction, cells were fixed and stained for neutral lipids with Oil-red O.

Effects of Ob-X on mRNA expression of angiogenic factors

The effects of Ob-X on the expression patterns of genes involved in angiogenesis were investigated in 3T3-L1 cells. The mRNA expression of angiogenic factors was downregulated, and anti-angiogenic factors were upregulated, in Ob-X-treated cells compared with non-differentiated control cells. After MDI treatment, the mRNA levels of angiogenic factors VEGF-A, -B, -C, -D, and FGF-2 were elevated by 516, 275, 65, 42, and 72%, respectively (Figure 3A). However, Ob-X treatment reduced the mRNA levels of VEGF-A, -B, -C, -D, and FGF-2 by 86, 73, 36, 20, and 47%, respectively, in MDI-treated cells. Ob-X also decreased these levels by 29, 43, 47, 29, and 50%, respectively, in troglitazone-treated cells (Figure 3B). In contrast, the mRNA level of the anti-angiogenic molecule thrombospondin-1 (TSP-1) was elevated by 86% in MDI-treated cells, and by 25% in troglitazone-treated cells.

Effects of Ob-X on mRNA expression of MMPs

During differentiation of 3T3-L1 cells, MMP-2 and MMP-9 mRNA expression was stimulated by 42 and 10%, respectively, following MDI treatment (Figure 4A), and 125 and 62%, respectively, following troglitazone treatment (Figure 4B). In contrast, Ob-X increased tissue inhibitor of metalloproteinase-1 (TIMP-1) and TIMP-2 mRNA levels by 28 and 65%, respectively, in MDI-treated cells and by 136 and 29%, respectively, in troglitazone-treated cells.

Effects of Ob-X on MMP activity

MMP activity was examined using zymography on gelatin-containing gels. Gelatin zymography revealed that the activity of proMMP-2 and proMMP-9 was significantly reduced in Ob-X-treated cells. Ob-X treatment reduced

the proMMP-2 and proMMP-9 activities in MDI-treated cells by 26 and 61%, respectively (Figure 5A). Ob-X also decreased the proMMP-2 and proMMP-9 activities in troglitazone-treated cells by 8 and 50%, respectively (Figure 5B).

Discussion

The development of fat cells from preadipocytes, or adipogenesis, includes morphological changes, expression of many lipogenic enzymes, and extensive lipid accumulation (Rosen and Spiegelman, 2000), contributing to the growth and expansion of adipose tissue. Because our previous study demonstrated that Ob-X, a mixture of three herbs with anti-angiogenic and MMP-inhibiting properties, reduced adipose tissue mass (Kim et al., 2010; Yoon and Kim, accepted), this study was undertaken to investigate whether Ob-X inhibits adipogenesis and to determine the expression patterns of genes involved in lipogenesis, angiogenesis, and MMPs using an *in vitro* cell culture system.

We clearly demonstrate that Ob-X is capable of suppressing adipogenesis and adipocyte-specific gene expression. In early stage development of adipose tissue, angiogenesis is tightly associated with angiogenesis, and EC proliferation occurs in expanding adult adipose tissue (Crandall et al., 1997; Lijnen, 2008). MMP inhibitors are also known to block adipocyte conversion only when present in the cell culture medium during the initial stages of the differentiation program (Croissandeau et al., 2002), suggesting that this is when angiogenesis and MMP activities play a critical role. Based on these previous results, we treated cells with Ob-X for the first 2 of 6 days. As expected, MDI or troglitazone increased accumulation of triglyceride droplets in 3T3-L1 cells compared with undifferentiated cells. However, Ob-X treatment prevented this MDI- or troglitazone-induced lipid accumulation, indicating that Ob-X has an inhibitory effect on troglitazone- or MDI-induced adipogenesis.

Adipogenesis is initiated by the production of the key transcription factor PPAR γ , which is responsible for inducing the expression of adipocyte-specific genes (Rosen et al., 1999). Consistent with the effects of Ob-X on lipid accumulation, Ob-X decreased the expression of PPAR γ and the PPAR γ target gene aP2, which are directly implicated in lipogenic pathways in 3T3-L1 adipocytes. Our results are supported by the finding that blocking differentiation in 3T3-L1 cells is associated with diminished PPAR γ expression (Croissandeau et al., 2002). MMP inhibitors are not thought to affect the mitotic clonal expansion, but inhibit adipocyte differentiation through diminished expression of PPAR γ (Croissandeau et al., 2002). PPAR γ expression is controlled by members of the CCAAT/enhancer-binding protein (C/EBP) (Wu et al., 1996) and sterol regulatory element binding protein (Fajas et al., 1999) families. In this process, MMP inhibitors decrease the DNA binding capacity of C/EBP β to its recognition element in the PPAR γ gene promoter,

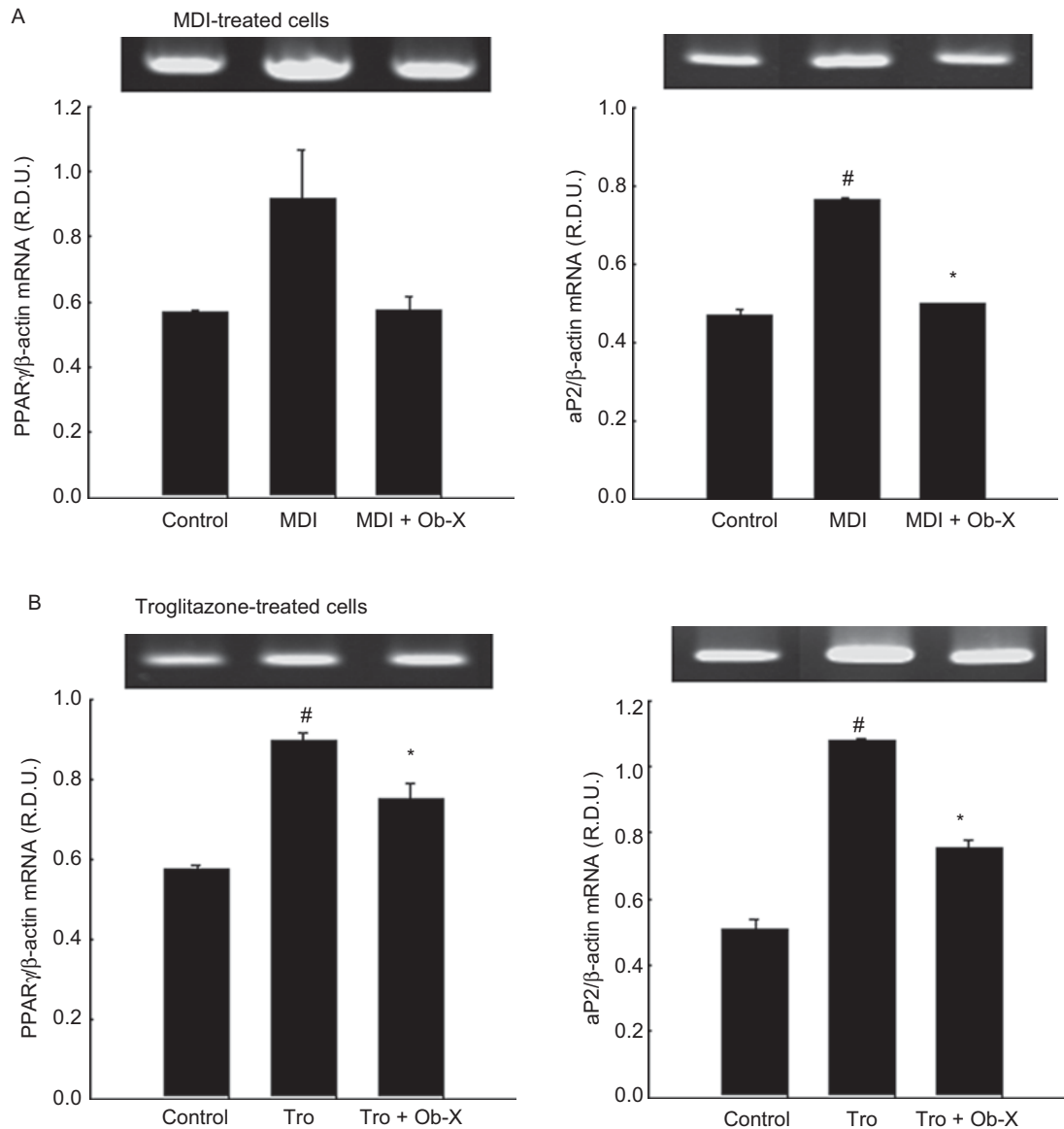


Figure 2. Effects of Ob-X on mRNA expression of adipose-specific genes in 3T3-L1 cells. 3T3-L1 preadipocytes were differentiated into mature adipocytes as described in "Materials and Methods". 3T3-L1 cells were treated with monocyte differentiation-inducing (MDI) differentiation mix (MDI), MDI plus 10 μ g/mL Ob-X, 10 μ M troglitazone (Tro), or 10 μ M Tro plus 10 μ g/mL Ob-X, and the effects of Ob-X on (A) MDI- or (B) troglitazone-induced expression of adipocyte-specific genes were investigated. Total cellular RNA was extracted from differentiated cells on day 6, and mRNA levels of peroxisome proliferator-activated receptor γ , aP2, and β -actin were measured using reverse transcription-polymerase chain reaction (RT-PCR). All values are expressed as the mean \pm SD. Insets show representative RT-PCR bands used for quantitation. *Significantly different versus MDI or Tro, respectively, $P < 0.05$; #significantly different versus control, $P < 0.05$.

resulting in the impairment of PPAR γ transcription (Croissandeau et al., 2002). Collectively, angiogenesis and MMP inhibitors are capable of blocking adipose conversion at the early stages of differentiation when C/EBP β induction is maximal. Thus, Ob-X seems to be an effective angiogenesis and MMP inhibitor to inhibit adipocyte differentiation. Interestingly, inhibition of adipocyte differentiation by overexpression of a dominant-negative PPAR γ construct leads to impaired development of both adipose tissue and angiogenesis (Fukumura et al., 2003).

Growing adipocytes produce multiple angiogenic factors and their inhibitors that regulate adipose angiogenesis. Angiogenic factors such as VEGF-A, -B, -C, and -D, as well as FGF-2, promote the proliferation, differentiation,

and migration of ECs within fat and enhance adipocyte differentiation *in vivo* (Carmeliet et al., 1996; Bikfalvi et al., 1997; Kawaguchi et al., 1998) whereas TSP-1 inhibits angiogenesis *in vivo* and impairs migration and proliferation of cultured microvascular ECs (Armstrong and Bornstein, 2003). Blockage of the VEGF receptor 2 (VEGFR2) signaling system by a neutralizing antibody inhibits both angiogenesis and preadipocyte differentiation, suggesting that VEGFs act on ECs to regulate preadipocyte differentiation (Fukumura et al., 2003). In addition, angiogenesis inhibitors, such as TNP-470 and VEGFR2-specific inhibitors, have been shown to prevent the development of obesity in genetic mouse models and studies based on high-fat diets (Rupnick et al., 2002;

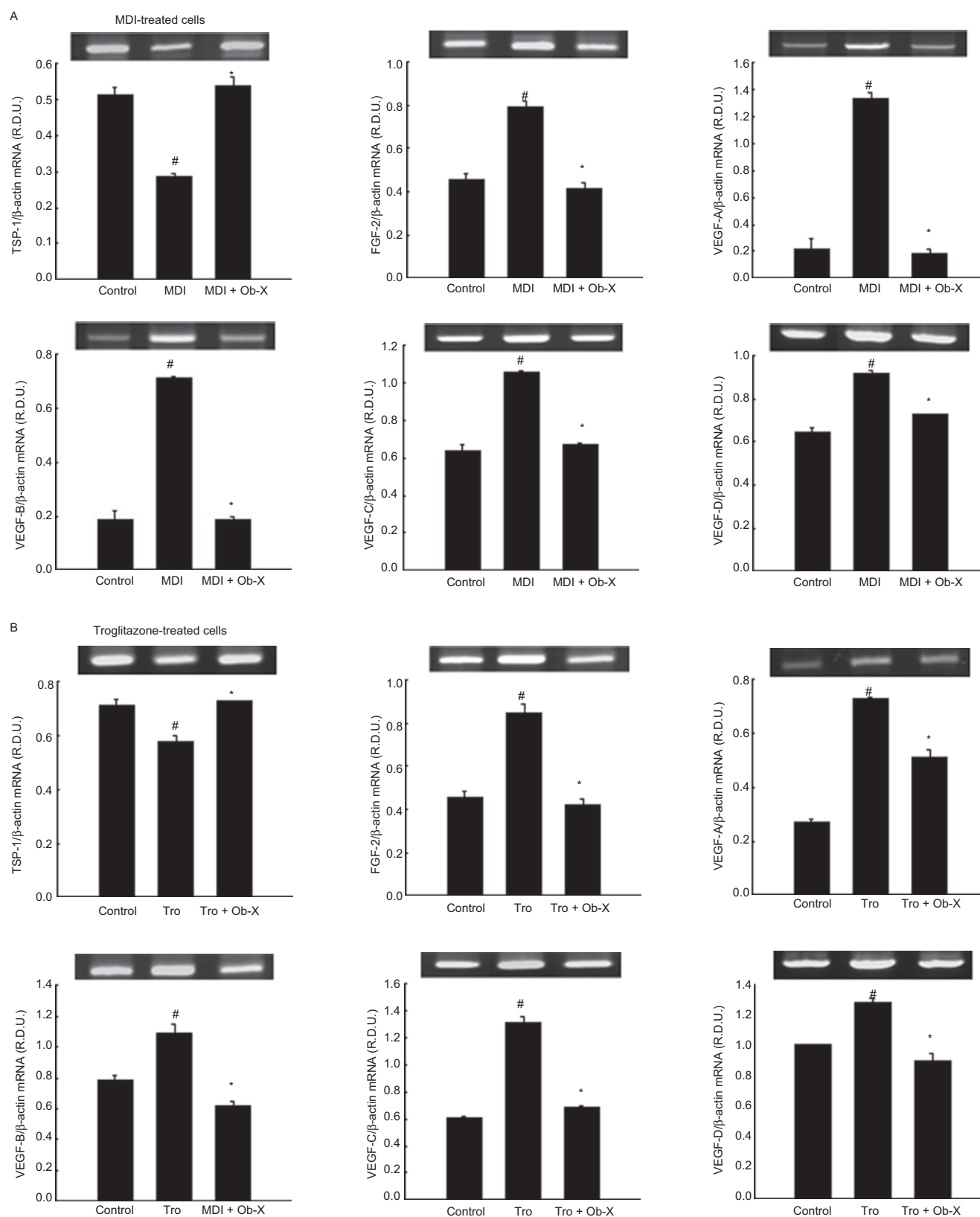


Figure 3. Effects of Ob-X on mRNA expression of genes involved in angiogenesis in 3T3-L1 cells. 3T3-L1 preadipocytes were differentiated into mature adipocytes as described in "Materials and Methods". 3T3-L1 cells were treated with monocyte differentiation-inducing (MDI) differentiation mix (MDI), MDI plus 10 μ g/mL Ob-X, 10 μ M troglitazone (Tro), or 10 μ M Tro plus 10 μ g/mL Ob-X, and the effects of Ob-X on (A) MDI- or (B) troglitazone-induced expression of genes involved in angiogenesis were investigated. Total cellular RNA was extracted from differentiated cells on day 6, and mRNA levels of VEGF-A, -B, -C, and -D, thrombospondin-1 (TSP-1), and β -actin were measured using reverse transcription-polymerase chain reaction (RT-PCR). All values are expressed as the mean \pm SD. Insets show representative RT-PCR bands used for quantitation. *Significantly different versus MDI or Tro, respectively, $P < 0.05$; #significantly different versus control, $P < 0.05$.

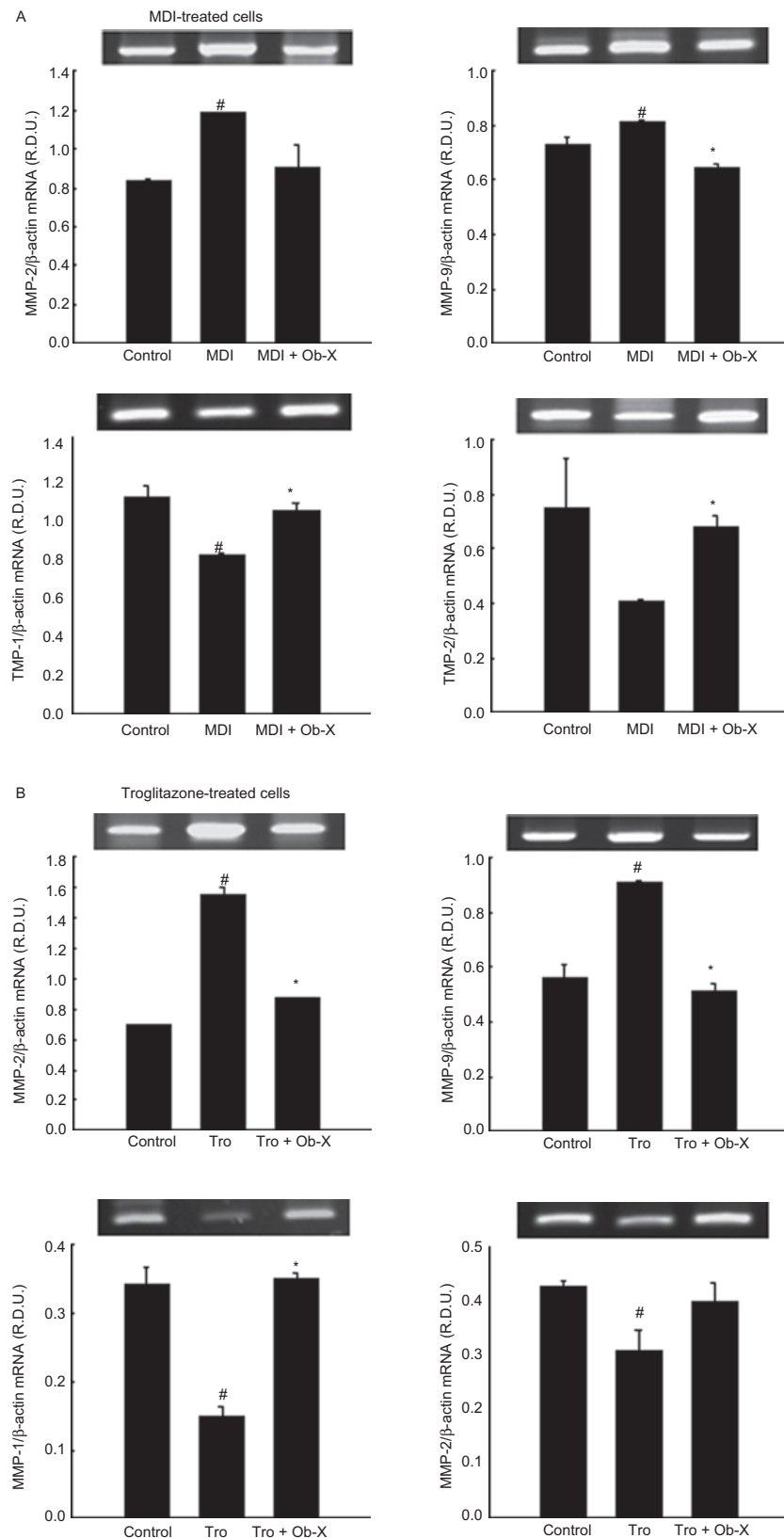


Figure 4. Effects of Ob-X on mRNA expression of matrix metalloproteinases (MMPs) and their inhibitors in 3T3-L1 cells. 3T3-L1 preadipocytes were differentiated into mature adipocytes as described in "Materials and Methods". 3T3-L1 cells were treated with monocyte differentiation-inducing (MDI) differentiation mix (MDI), MDI plus 10 μ g/mL Ob-X, 10 μ M troglitazone (Tro), or 10 μ M Tro plus 10 μ g/mL Ob-X, and the effects of Ob-X on (A) MDI- or (B) troglitazone-induced expression of MMPs and their inhibitors were investigated. Total cellular RNA was extracted from differentiated cells on day 6, and mRNA levels of MMP-2, MMP-9, tissue inhibitor of metalloproteinase-1 (TIMP-1), TIMP-2, and β -actin were measured using reverse transcription-polymerase chain reaction (RT-PCR). All values are expressed as the mean \pm SD. Insets show representative RT-PCR bands used for quantitation. *Significantly different versus MDI or Tro, respectively, $P < 0.05$; #significantly different versus control, $P < 0.05$.

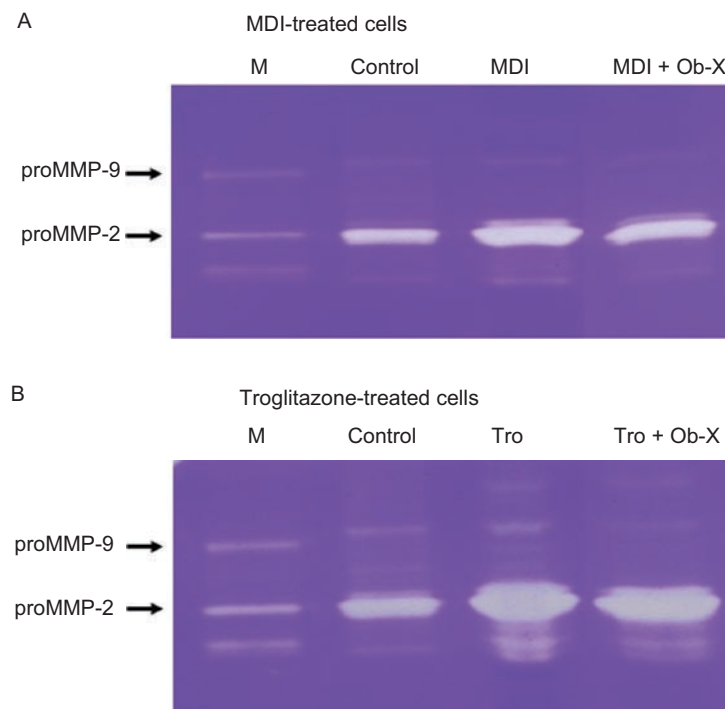


Figure 5. Zymographic analysis of 3T3-L1 cells. (A) Extracts from 3T3-L1 cells treated with monocyte differentiation-inducing (MDI) differentiation mix (MDI) or MDI plus 10 μ g/mL Ob-X were applied to a gelatin-containing gel. (B) Extracts from 3T3-L1 cells treated with 10 μ M troglitazone (Tro) or 10 μ M Tro plus 10 μ g/mL Ob-X were applied to a gelatin-containing gel. Gelatinolytic activity was measured by zymography. M is the molecular weight marker for matrix metalloproteinases.

Fukumura et al., 2003; Bråkenhielm et al., 2004; Tam et al., 2009). In our present study, Ob-X decreased the mRNA expression of four kinds of VEGFs and FGF-2 that were increased in differentiated 3T3-L1 cells compared with non-differentiated cells. Moreover, Ob-X treatment the increased mRNA level of the anti-angiogenic TSP-1 in 3T3-L1 cells, suggesting that the anti-angiogenic agent Ob-X reduces adipogenesis and can be used for treatment of obesity.

Adipocytes also produce MMPs and MMP inhibitors that are differentially expressed in adipose tissue during obesity in murine obesity models (Maquoi et al., 2002; Chavey et al., 2003; Voros et al., 2005). Furthermore, the secretion of MMP-2 and MMP-9 increases during adipocyte differentiation in both human adipocytes and mouse preadipocyte cell lines (Bouloumié et al., 2001; Maquoi et al., 2002; Chavey et al., 2003), suggesting that MMP-2 and MMP-9 are important for adipocyte conversion. Our reverse transcription-PCR analysis showed that Ob-X decreased MMP-2 and MMP-9 mRNA levels, but increased TIMP-1 and TIMP-2, indicating that Ob-X exerts a specific regulatory effect on genes involved in angiogenesis and the MMP system in 3T3-L1 cells. Several studies demonstrated that MMPs have a novel function in adipogenesis, modulating adipocyte differentiation independent of angiogenesis and therefore, MMP inhibitors can block the adipocyte differentiation process (Bouloumié et al., 2001; Croissandeau et al., 2002; Maquoi et al., 2002; Chavey et al., 2003). Treatment with MMP inhibitors impairs adipose tissue development in mice

fed a high-fat diet (Lijnen et al., 2002). Consistent with the inhibitory effects of Ob-X on the mRNA expression of MMP-2 and MMP-9, zymographic analysis revealed that Ob-X suppressed MMP-2 and MMP-9 gelatinolytic activities, since proMMP-2 and proMMP-9 activities were markedly reduced in Ob-X-treated 3T3-L1 cells. It has also been reported that *in situ* zymography with gelatin-containing gels on cryosections of adipose tissue confirmed lower MMP activity in tissues of the MMP inhibitor galardin-treated animals. Our present results indicate that the reduction of adipogenesis by Ob-X may be due to its anti-angiogenic and MMP-inhibiting actions.

These studies demonstrate that Ob-X, which inhibits angiogenesis and MMP activity, suppresses adipogenesis in 3T3-L1 adipocytes. These events may be mediated by changes in the expression of genes involved in lipogenesis, angiogenesis, and the MMP system. Thus, by reducing adipogenesis, anti-angiogenic Ob-X provides a possible therapeutic approach for the prevention and treatment of human obesity and its related disorders.

Declaration of interest

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