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Effects of PGC-1 α on TNF- α -Induced MCP-1 and VCAM-1 Expression and NF- κ B Activation in Human Aortic Smooth Muscle and Endothelial Cells

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ABSTRACT

Increased oxidative stress in vascular cells is implicated in the pathogenesis of atherosclerosis. Reactive oxygen species (ROS) induce vascular inflammation via the proinflammatory cytokine/NF- κ B pathway. Several lines of evidence suggest that peroxisome proliferator-activated receptor- γ coactivator 1- α (PGC-1 α) is an important regulator of intracellular ROS levels. However, no studies have examined the effects of PGC-1 α on this process. We investigated the effects of PGC-1 α on inflammatory molecule expression and activity of the redox-sensitive transcription factor, NF- κ B, in vascular cells. PGC-1 α expressed in human aortic smooth (HASMCs) and endothelial cells (HAECs) is upregulated by AMP-activated protein kinase activators, including metformin, rosiglitazone and α -lipoic acid. Tumor necrosis factor- α (TNF- α), a major proinflammatory factor in the development of vascular inflammation, stimulates intracellular ROS production through an increase in both mitochondrial ROS and NAD(P)H oxidase activity. Adenovirus-mediated overexpression of the PGC-1 α gene in HASMCs and HAECs leads to a significant reduction in intracellular and mitochondrial ROS production as well as NAD(P)H oxidase activity. Consequently, NF- κ B activity and MCP-1 and VCAM-1 induced by TNF- α are suppressed. Our data support the possibility that agents stimulating PGC-1 α expression in the vasculature aid in preventing the development of atherosclerosis. *Antioxid. Redox Signal.* 9, 301–307.

INTRODUCTION

ATHEROSCLEROSIS is a chronic inflammatory disease that involves the interplay of various soluble mediators, monocytes, endothelial cells, and vascular smooth muscle cells (9, 24). The initial step in atherosclerosis is the recruitment of mononuclear cells into subendothelial space (24). One of the underlying mechanisms of leukocyte recruitment is a series of complex interactions between circu-

lating leukocytes and vascular cells. Stimulation of vascular cells by cytokines initiates a signaling cascade leading to NF- κ B-dependent expression of genes encoding chemokine and adhesion molecules, including vascular cell adhesion molecule-1 (VCAM-1) and monocyte chemoattractant protein-1 (MCP-1) (2, 4). These changes accelerate vascular endothelial cell damage and vascular smooth muscle cell proliferation and migration. Thus, vascular expression of MCP-1 and VCAM-1 induced by the proinflammatory cy-

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tokine/NF- κ B pathway is a key mechanism in the development of atherosclerosis.

It is well established that increased oxidative stress in vascular cells plays an important role in the pathogenesis of atherosclerosis (6, 8). Reactive oxygen species (ROS) induce a local inflammatory response through release of various cytokines, including tumor necrosis factor (TNF)- α , from monocytes (12). Generation of intracellular ROS in vascular cells by TNF- α initiates a signaling cascade leading to NF- κ B-dependent gene expression, and induces a local inflammatory response through production of various chemokines and adhesion molecules (3, 14). Although the relative contribution of the individual ROS generating systems in the vasculature is currently ambiguous, both cell membrane NAD(P)H oxidase and the mitochondrial electron-transport chain play significant roles in ROS generation by TNF- α (10, 13, 14, 21). Previous studies demonstrate that adenovirus-mediated overexpression of uncoupling protein-2 (UCP-2) in human aortic smooth muscle cells (HASMCs) decreases ROS generation by inhibiting both pathways (22) and suppresses NF- κ B activation by fatty acids in human aortic endothelial cells (HAECs) (19). Moreover, peroxisome proliferator-activated receptor- γ coactivator 1- α (PGC-1 α) induces UCP-2 expression in vascular cells (26). Collectively, these results support the hypothesis that PGC-1 α decreases TNF- α -induced NF- κ B activation and MCP-1 and VCAM-1 expression in vascular cells by inhibiting intracellular ROS generation.

PGC-1 α is a transcriptional coactivator, identified as an upstream regulator of mitochondrial number and function (26). Several reports show that PGC-1 α is involved in the control of ROS production by mitochondria (16, 25, 26). Recently, a direct role for PGC-1 α in vascular endothelial cells was proposed from the finding that its expression is associated with an increase in mitochondrial antioxidant defense (16, 26).

Vallea *et al.* (26) showed that overexpression of PGC-1 α in vascular endothelial cells trigger an increase in mitochondrial antioxidative enzymes and suppress ROS production. Moreover, Kukidome and colleagues (16) suggested that active AMP-activated protein kinase (AMPK) reduces ROS generation via PGC-1 α expression under hyperglycemic conditions.

In this study, we investigated the effects of PGC-1 α on TNF- α -stimulated MCP-1 and VCAM-1 expression, as well as NF- κ B activity in the vascular cell lines, HASMC and HAEC.

MATERIALS AND METHODS

Materials

Recombinant human TNF- α was purchased from R&D systems (Minneapolis, MN). Radiochemicals ([α -32P]dCTP, [γ -32P]ATP) were from Amersham Biosciences (Little Chalfont, U.K.). Metformin was acquired from Sigma. Rosiglitazone and α -lipoic acid were provided by GlaxoSmithKline (U.K.) and Viartis GmbH & Co. KG (Frankfurt, Germany), respectively.

Cell culture

HASMCs were isolated from the thoracic aorta of kidney transplantation donors by using the explant method, as described previously (1). Tissue collection was approved by the Ethics Committee of the Institution. Cells were cultured in DMEM (Gibco BRL, MD) containing 20% fetal bovine serum (FBS) (Hyclone, Logan, UT). In each preparation, HASMC's purity was determined by positive staining with smooth muscle-specific α -actin monoclonal antibodies (Santa Cruz). All cells were used within passages 5 and 6. At 90% confluence, cells in 100-mm dishes were serum-starved for 24 h with DMEM containing 0.5% serum and 5.5 mM D-glucose. HAECs were obtained from BioWhittaker Inc. (Walkersville, MD) and maintained in endothelial basal medium (EBM; BioWhittaker) supplemented with various growth factors and 2% FBS. Cells were passaged more than 3 times before use in experiments, and subsequently processed for nuclear protein or RNA extraction, as described later.

Preparation of recombinant adenovirus

Full-length human PGC-1 α cDNA was inserted into the *HindIII/BamHI* site of the pAd-YC2 shuttle vector (7). Shuttle vectors containing human PGC-1 α and the rescue vector, pJM17, were co-transfected into human embryonic kidney 293 (HEK-293) cells, which were cultured on 24-well plates the day before transfection. After 12–15 days, recombinants were identified by polymerase chain reaction (PCR) (7), after which they were amplified in HEK-293 cells, and purified and isolated by using CsCl (Sigma). Preparations were collected and desalted, and titers were determined by measuring plaque counts. Control adenovirus devoid of PGC-1 α (Ad-Null) was additionally prepared and identified by using this protocol.

Northern blot analysis

Total RNA was isolated from cells by using TRI reagent (Sigma), according to the manufacturer's instructions. Aliquots (20 μ g) of total RNA from each sample were loaded onto 1.0% formaldehyde-agarose gels. After electrophoresis, RNA was transferred to Hybond-N⁺ nylon membrane (Amersham). After transfer, membranes were cross-linked by using UV crosslinker 1800 (Stratagene, La Jolla, CA), and hybridized to random-primed ³²P-labeled probes at 65°C overnight. The membrane was washed with wash solution I (2 \times SSC, 0.05 % SDS solution) and wash solution II (0.1 \times SSC, 0.1 % SDS solution), air-dried, and exposed to autoradiography film.

Measurement of intracellular H₂O₂ production

Cells were seeded onto a six-well plate. At 90% confluence, cells were incubated in DMEM medium containing 1% FBS for 24 h. Cells treated with 100 MOI (multiplicity of infection) of adenovirus containing the PGC-1 α gene (Ad-PGC-1 α) or Ad-Null were cultured for 24 h. After exposure to TNF- α for 30 min, 10 μ M 2',7'-dichlorofluorescein diacetate (DCF-DA; Sigma), a H₂O₂-sensitive fluorescent

probe, was added, and cells were cultured for 30 min. H₂O₂ production was quantified by using an AxioCam MRc5 Carl Zeiss fluorescence microscope (Thornwood, NY) at an excitation wavelength of 488 nm and emission wavelength of 515 nm.

Measurement of mitochondrial ROS

To evaluate the direct production of mitochondrial ROS in HASMCs and HAECs, specific staining with the reduced MitoTracker Red probe (CM-H2XROS; Molecular probe) was performed. In brief, cells were cultured at 37°C for 24 h under each condition (with 1% FBS), and loaded with 0.2 μ M CM-H2XROS at 37°C for 30 min. A fluorescence microscope (AxioCam MRc5; Carl Zeiss) equipped for equifluorescent illumination was used.

Measurement of NAD(P)H oxidase activity

To measure NAD(P)H oxidase activity, cells were washed twice with PBS, lysed with lysis buffer (50 mM Tris-HCl, 150 mM NaCl, 1 mM EDTA, 1% Triton X-100, 10% glycerol, 1 mM PMSF, 1 mg/ml aprotinin, 1 mg/ml leupeptin), and incubated for 1 h on ice. The lysate was centrifuged at 12,000 g for 20 min, and the supernatant obtained. Protein content was determined by using the Bradford method (Bio-Rad, Hercules, CA). NAD(P)H oxidase activity was measured by using lucigenin chemiluminescence (17).

Electrophoretic mobility shift assay

Nuclear extracts were prepared from cells and incubated (6 μ g) with ~60,000 cpm of the ³²P-labeled NF- κ B-binding site oligomer, 5'-AGTTGAGGGGACTTTCCAGGC-3' (Santa Cruz), for 20 min at RT. In brief, DNA probes, such as those for NF- κ B, were labeled by using [γ -³²P]ATP and T4 polynucleotide kinase. After end-labeling, ³²P-labeled NF- κ B was purified with a NAP-5 column (Amersham, Little Chalfont, U.K.). Protein-DNA-binding reactions were performed at room temperature for 20 min in a total volume of 30 μ l. Reaction mixtures contained 6 μ g nuclear extract, 100 μ g/ml poly dI:dC, 10 mM Tris/HCl (pH 7.5), 50 mM NaCl, 0.5 mM EDTA, 0.5 mM DTT, 1 mM MgCl₂, 4% glycerol, and 60,000 cpm ³²P-labeled primer DNA. After incubation, samples were loaded onto 4% native polyacrylamide gels in 0.5 M Tris-borate-EDTA buffer and run at 150 V for 2 h. Gels were dried and visualized by autoradiography.

Real-time quantitative RT-PCR

After incubation for 24 h in each condition, total cellular RNA was isolated from cells by using TRIzol reagent, according to the manufacturer's protocol (Life Technologies, Gaithersburg, MD). The LightCycler System (Roche Molecular Biochemicals, Indianapolis, IN) was used to quantify PGC-1 α transcripts. PCR reactions were performed by using SYBR Green I master mix, and specific primers for human PGC-1 α (5'-TCAGTCCTCACTGGTGGACA-3' and 5'-TGCTTCGTCAAAAACA G-3') and human β -actin (5'-CAC-CCACACTGTGCCCATCTACGA-3' and 5'-CAGCGGAACCGCTCA TTGCCAATGG-3'). To assess the specificity of the

amplified products, a melting curve analysis was performed after the last cycle.

Statistical analysis

Results were expressed as mean values \pm SEM. Variance analysis with a subsequent Duncan's test was used to determine significant differences in multiple comparisons. A *p* value of <0.05 was considered statistically significant. All experiments were performed at least three times.

RESULTS

Adenovirus-mediated PGC-1 α gene overexpression inhibits TNF- α -induced MCP-1 and VCAM-1 expression in vascular cells

In view of the lack of reports on PGC-1 α expression in HASMCs. We initially examined whether PGC-1 α is basally expressed in this cell line. RT-PCR results disclose that primary cultured HASMCs express PGC-1 α mRNA (data not shown). To determine whether PGC-1 α affects MCP-1 and VCAM-1 expression in vascular cells, we generated an adenoviral vector containing human PGC-1 α cDNA (Ad-PGC-1 α). HASMCs and HAECs were infected with Ad-PGC-1 α at concentrations of 10, 50, and 100 MOI. Notably, Ad-PGC-1 α suppressed TNF- α -induced MCP-1 and VCAM-1 mRNA expression in HASMCs and HAECs in a dose-dependent manner, whereas Ad-Null had no effect (Fig. 1).

Ad-PGC-1 α inhibits TNF- α -induced intracellular ROS production

We determined the effects of PGC-1 α on TNF- α -induced ROS production in HASMCs and HAECs. TNF- α (5 ng/ml) stimulated the production of H₂O₂ in HASMCs and HAECs, compared with that observed under basal conditions. Infection with Ad-PGC-1 α (100 MOI) led to complete inhibition of TNF- α -induced H₂O₂ production to the basal level. In contrast, Ad-Null did not influence the H₂O₂ level (Fig. 2).

Ad-PGC-1 α inhibits TNF- α -stimulated mitochondrial ROS production and NAD(P)H oxidase activity

To elucidate the mechanism by which PGC-1 α inhibits TNF- α -induced ROS production, we examined the effects of Ad-PGC-1 α on mitochondrial ROS and NADPH oxidase activity, two major sources of TNF- α -induced intracellular ROS. As shown in Fig. 3, MitoTracker Red fluorescence in HASMCs and HAECs significantly increased with 5 ng/ml TNF- α , compared with the control. TNF- α -induced fluorescence was completely suppressed in the presence of Ad-PGC-1 α , indicating abolishment of mitochondrial ROS production. In addition, infection of HASMCs and HAECs with Ad-PGC-1 α partially but significantly decreased TNF- α -induced NAD(P)H oxidase activity. However, infection with Ad-Null had no effects on this parameter (Fig. 4).

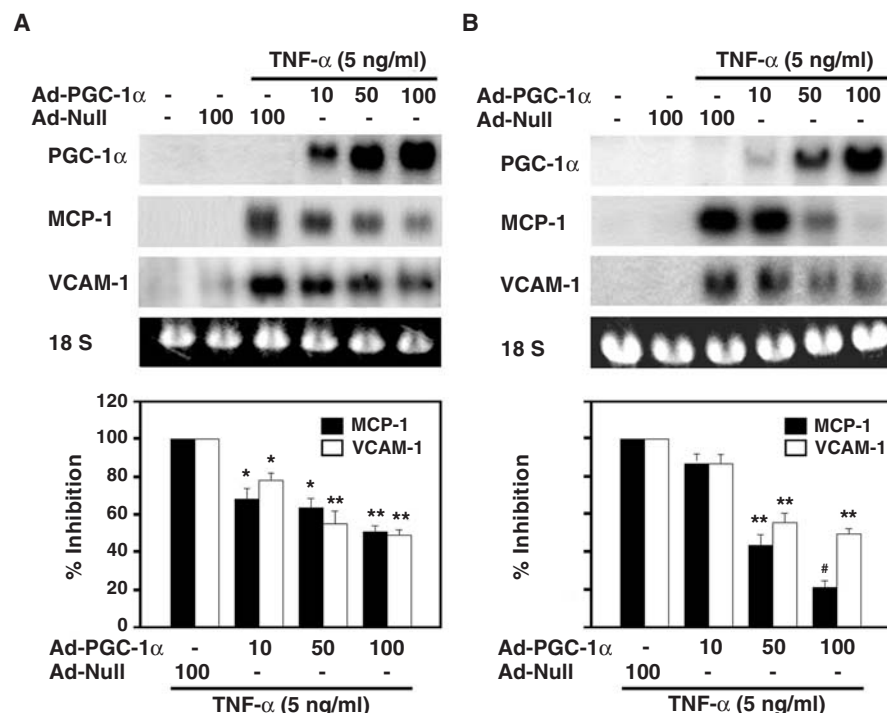


FIG. 1. Effects of Ad-PGC-1 α on MCP-1 and VCAM-1 mRNA expression. HASMCs (A) and HAECs (B) were infected with indicated doses (MOI) of Ad-PGC-1 α or Ad-Null for 2 h and treated with TNF- α (5 ng/ml) for 24 h. Total RNA (10 μ g) was hybridized with MCP-1, VCAM-1, and PGC-1 α probes for Northern blot analysis. The bars represent fold increase, compared with TNF- α -treated control. Data are expressed as mean \pm SEM of three separate measurements (lower panel). Statistical significance was determined as * p < 0.05, ** p < 0.01, and # p < 0.001, compared with TNF- α and Ad-Null.

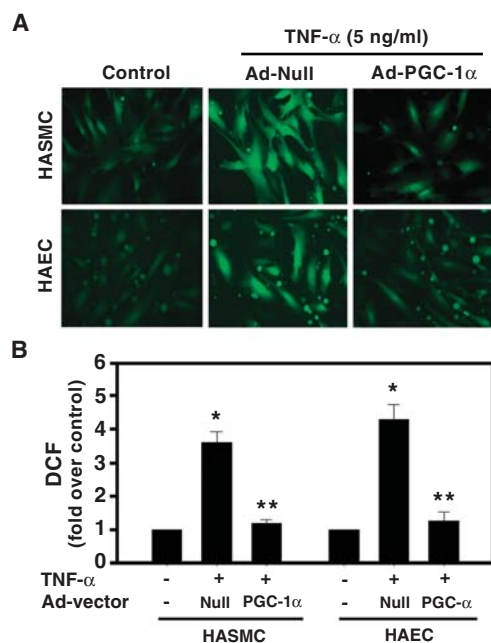


FIG. 2. Effects of Ad-PGC-1 α on H₂O₂ production in response to TNF- α . (A) HASMCs and HAECs were infected with 100 MOI of Ad-PGC-1 α or Ad-Null. Cells were further treated with TNF- α (5 ng/ml) for 30 min and processed for fluorescence microscopy by using the oxidant-sensitive probe 2',7'-dichlorofluorescein diacetate. Fully confluent fields of HASMCs and HAECs were randomly selected, and fluorescence was quantified with the NIH Image program at a magnification of $\times 100$. (B) Data are expressed as mean \pm SEM of three separate measurements. Statistical significance was determined as * p < 0.001, compared with basal conditions, and ** p < 0.001, compared with TNF- α and Ad-Null.

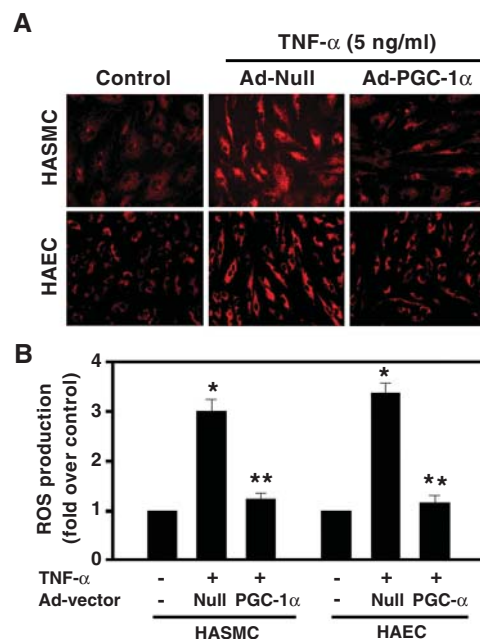


FIG. 3. Effects of Ad-PGC-1 α on mitochondrial ROS production. (A) HASMCs and HAECs were infected with 100 MOI of Ad-PGC-1 α or Ad-Null for 2 h and treated with TNF- α (5 ng/ml) for 24 h. Mitochondrial ROS production was detected by using MitoTracker Red CM-H2XRos. Fully confluent fields of HASMCs or HAECs were randomly selected, and fluorescence was quantified by using the NIH Image program. The magnification was $\times 100$. (B) Data are expressed as mean \pm SEM of three separate measurements. Statistical significance was determined as * p < 0.001, compared with basal, and ** p < 0.001, compared with TNF- α and Ad-Null.

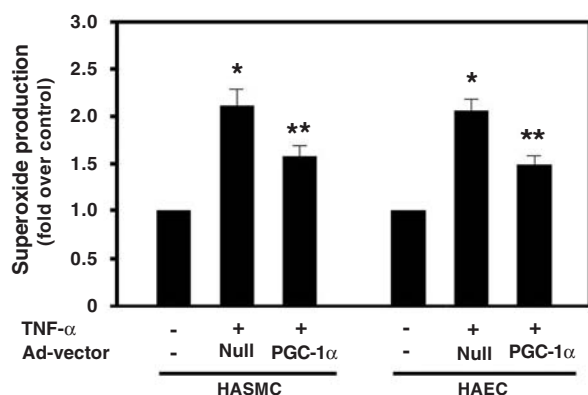


FIG. 4. Effects of Ad-PGC-1 α on NAD(P)H oxidase activity. (A) HASMCs and HAECs were infected with 100 MOI of Ad-PGC-1 α or Ad-Null for 2 h and treated with TNF- α (5 ng/ml) for 24 h. NAD(P)H oxidase activity was determined, as described in Materials and Methods. (B) Data are expressed as mean \pm SEM of three separate measurements. Statistical significance was determined as * p < 0.01, compared with basal conditions, and ** p < 0.05, compared with TNF- α and Ad-Null.

Ad-PGC-1 α inhibits TNF- α -stimulated NF- κ B activation

Because NF- κ B activation by intracellular oxidative stress is critical for the expression of MCP-1 and VCAM-1 in vascular cells, we examined whether Ad-PGC-1 α inhibits TNF- α -stimulated NF- κ B activity. Notably, TNF- α (5 ng/ml) markedly increased NF- κ B-DNA binding activity in HASMCs and HAECs. This binding activity was attenuated

by Ad-PGC-1 α in a dose-dependent manner, whereas Ad-Null had no effect (Fig. 5).

AMPK activators increase PGC-1 α mRNA expression in HASMCs

Metformin, rosiglitazone, and α -lipoic acid activate AMPK (11, 15, 28). Furthermore, recent studies disclose that activation of AMPK leads to enhanced PGC-1 α expression in vascular endothelial cells (16). Previously, we reported that α -lipoic acid stimulates AMPK activity in HAECs (20). The data collectively suggest that metformin, rosiglitazone, and α -lipoic acid increase PGC-1 α expression in vascular endothelial cells. Real-time RT-PCR findings from the present study indicate that metformin, rosiglitazone, and α -lipoic acid significantly increase PGC-1 α mRNA expression in HASMCs in a dose-dependent manner (Fig. 6). Additionally, we observed that metformin, rosiglitazone, and α -lipoic acid increased PGC-1 α expression in HASMCs after exposure to TNF- α (data not shown).

DISCUSSION

We demonstrated that PGC-1 α expressed in HASMCs is upregulated by AMPK activator including metformin, rosiglitazone, and α -lipoic acid. Adenoviral transfer of the PGC-1 α gene to HASMCs significantly suppressed intracellular ROS production, NF- κ B activation, and expression of MCP-1 and VCAM-1 induced by TNF- α . Similar data were obtained with HAECs.

While findings on ROS production in response to TNF- α are inconsistent in general, recent studies show that TNF- α

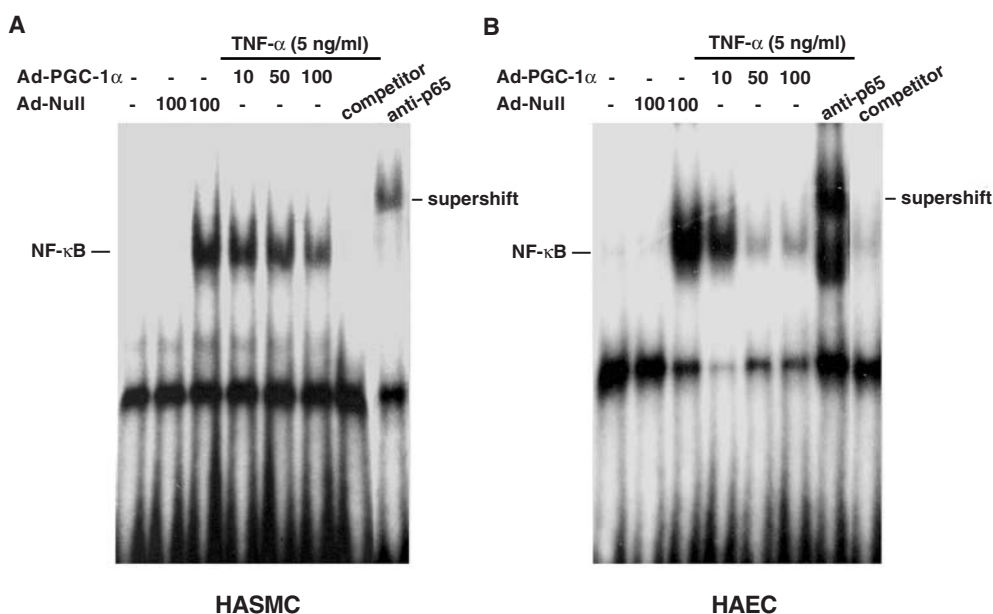


FIG. 5. Effects of Ad-PGC-1 α on NF- κ B binding activity. HASMCs (A) and HAECs (B) were infected with indicated doses (MOI) of Ad-PGC-1 α or Ad-Null for 2 h and treated with TNF- α (5 ng/ml) for 24 h. Nuclear extracts were prepared, as described in Materials and Methods, and protein (6 μ g) was incubated with radiolabeled NF- κ B probe.

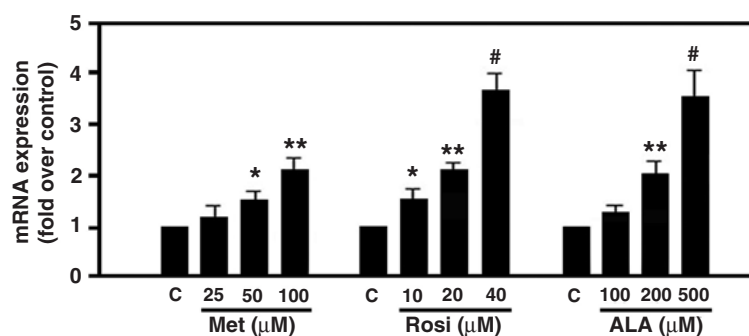


FIG. 6. Effects of metformin, rosiglitazone, and α -lipoic acid on PGC-1 α mRNA expression in HASMCs. HASMCs were treated with the indicated doses of metformin, rosiglitazone, or α -lipoic acid for 24 h. PGC-1 α mRNA expression was quantified by using real-time RT-PCR ($n = 5$). C, control; Met, metformin; rosi, rosiglitazone; ALA, α -lipoic acid. Data are expressed as mean \pm SEM of three separate measurements. Statistical significance was determined as * $p < 0.05$, ** $p < 0.01$, and # $p < 0.001$, compared with control (basal expres-

stimulates ROS generation from mitochondria (10, 13, 14) and plasma membrane NAD(P)H oxidase (21). In accordance with previous data, we showed that TNF- α induces ROS from both mitochondria and NAD(P)H oxidase. PGC-1 α in vascular endothelial cells increases mitochondrial antioxidant gene expression and reduces ROS generation (26). As expected, overexpression of PGC-1 α was associated with inhibition of TNF- α -induced mitochondrial ROS. Moreover, PGC-1 α partially but significantly decreased TNF- α -increased NAD(P)H oxidase activity. The mechanism by which PGC-1 α decreases NAD(P)H oxidase activity in HASMCs and HAECs remains to be established. However, our current findings are consistent with previous data showing that overexpression of UCP-2 in HASMCs decreases NADPH oxidase activity (22). Additionally, a recent publication shows that PGC-1 α in vascular cells increases UCP-2 levels (26). Therefore, the effect of PGC-1 α on NAD(P)H oxidase activity may be partly explained by the increase in UCP-2 expression in the presence of PGC-1 α .

NF- κ B, a major transcription factor in the development of atherosclerosis, is activated by intracellular ROS induced by various stimuli. NF- κ B activity in vascular cells is associated with the activation of genes responsible for increased transcription of adhesion molecules, cytokines, and chemokines (3, 4, 23, 25). Previously, we demonstrated that inhibition of intracellular ROS production by UCP-2 in HAECs blocked lysophosphatidylcholine and linoleic acid-induced NF- κ B activation (19). Because UCP-2 is a target gene of PGC-1 α in vascular cells (26), these data raise the possibility that PGC-1 α inhibits NF- κ B activation, which is stimulated by ROS. To date, no reports on the relation between PGC-1 α and NF- κ B activation and chemokine expression induced by TNF- α have been documented. Here, we show that PGC-1 α expression in vascular cells leads to the inhibition of TNF- α -induced NF- κ B activation and MCP-1 and VCAM-1 expression. These findings collectively suggest that upregulation of PGC-1 α prevents TNF- α -induced vascular inflammation by reducing intracellular oxidative stress and NF- κ B activation.

Several lines of evidence disclose that metformin, rosiglitazone, and α -lipoic acid inhibit vascular inflammatory molecule expression, including MCP-1 and VCAM-1 (5, 18, 27). In addition, metformin, rosiglitazone, and α -lipoic acid activate AMPK (11, 15, 28). Recently we demonstrated that α -lipoic acid activates AMPK in HAECs and prevents endothelial dysfunction (20). More recently, Kukidome *et al.* (16) demonstrated that metformin increases PGC-1 α gene expres-

sion and normalizes hyperglycemia-induced mitochondrial ROS production via AMPK activation. The present study showed that agents that activate AMPK, metformin, rosiglitazone, and α -lipoic acid additionally enhance PGC-1 α expression in vascular smooth muscle cells. Although possibly another pathway exists by which metformin, rosiglitazone, or α -lipoic acid inhibits TNF- α -induced activation of NF- κ B and expression of MCP-1 and VCAM-1 (18), the present data support one possible mechanism by which these AMPK activators prevent vascular inflammation.

In summary, we show that adenovirus-mediated transfer of PGC-1 α inhibits TNF- α -induced NF- κ B activation as well as MCP-1 and VCAM-1 expression in both HASMCs and HAECs. Based on the results, we propose that agents stimulating PGC-1 α expression in the vasculature aid in preventing the development of atherosclerosis.

ACKNOWLEDGMENTS

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ABBREVIATIONS

AMPK, AMP-activated protein kinase; HAECs, human aortic endothelial cells; HASMCs, human aortic smooth muscle cells; MCP-1, monocyte chemoattractant protein-1; NF- κ B, nuclear factor- κ B; ROS, reactive oxygen species; PGC-1 α , peroxisome proliferator activated receptor γ coactivator-1 α ; TNF- α , tumor necrosis factor- α ; VCAM-1, vascular cell adhesion molecule-1; UCP-2, uncoupling protein-2.

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