



Published in final edited form as:

J Immunol. 2012 December 1; 189(11): 5411–5420. doi:10.4049/jimmunol.1201487.

Endothelial Cell Peroxisome Proliferator-activated Receptor γ Reduces Endotoxemic Pulmonary Inflammation and Injury

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Abstract

Bacterial endotoxin (LPS)-mediated sepsis involves severe, dysregulated inflammation that injures the lungs and other organs, often fatally. Vascular endothelial cells are both key mediators and targets of LPS-induced inflammatory responses. The nuclear hormone receptor peroxisome proliferator-activated receptor γ (PPAR γ) exerts anti-inflammatory actions in various cells, but it is unknown whether it modulates inflammation through actions within endothelial cells. To determine whether PPAR γ acts within endothelial cells to diminish endotoxemic lung inflammation and injury, we measured inflammatory responses and mediators in mice with endothelial-targeted deletion of PPAR γ . Endothelial cell PPAR γ (ePPAR γ) knockout exacerbated LPS-induced pulmonary inflammation and injury as shown by several measures, including infiltration of inflammatory cells, edema, and production of reactive oxygen species and pro-inflammatory cytokines, along with upregulation of the LPS receptor TLR4 in lung tissue and increased activation of its downstream signaling pathways. In isolated LPS-stimulated endothelial cells *in vitro*, absence of PPAR γ enhanced the production of numerous inflammatory markers. We hypothesized that the observed *in vivo* activity of the ligand-activated ePPAR γ may arise in part from nitrated fatty acids (NFAs), a novel class of endogenous PPAR γ ligands. Supporting this idea, we found that treating isolated endothelial cells with physiologically relevant concentrations of the endogenous NFA 10-nitro-oleate reduced LPS-induced expression of a wide range of inflammatory markers in the presence of PPAR γ , but not in its absence, and also inhibited neutrophil mobility in a PPAR γ -dependent manner. Our results demonstrate a key protective role of ePPAR γ against endotoxemic injury, and a potential ePPAR γ -mediated anti-inflammatory role for NFAs.

Keywords

LPS; Nrf2; PPAR; ALI; nitrated fatty acid

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¹INTRODUCTION

Sepsis is a major cause of morbidity and mortality, frequently involving acute lung injury, in hospitalized patients (1, 2). It is often caused by endotoxins, the lipopolysaccharide (LPS) cell wall components of Gram-negative bacteria. LPS acts via Toll-like receptor-4 (TLR4) (3) to trigger an innate immune response that is usually protective, but that during endotoxemia can become hyperactivated and dysregulated, thus causing excessive inflammation and widespread organ injury. TLR4 stimulation is a multistep process, activating a cascade of downstream pathways that, via the transcription factors NF- κ B and AP-1, upregulate expression of pro-inflammatory cytokines, chemokines, and adhesion molecules. Endogenous anti-inflammatory signaling pathways serve to protect the host by restraining such inflammatory responses. Sepsis represents a failure of these counter-regulatory mechanisms, which normally keep inflammation in check and allow it to perform its protective functions without excessive tissue injury. Understanding these endogenous anti-inflammatory mechanisms is key to developing new tools to control pathogenic inflammation and sepsis.

Cells of the vascular endothelium are among the first to encounter circulating bacteria and their products including LPS, and they play key roles as both initiators and targets of the innate immune response. Activation of TLR4 in endothelial cells promotes increased production of neutrophil-attracting chemokines, along with integrins and adhesion molecules that promote neutrophil adhesion to the endothelium and migration into surrounding tissue, where they promote further inflammation. These actions contribute to the clinical features of severe endotoxemic sepsis, including increased vascular permeability that causes tissue edema, thereby restricting pulmonary gas exchange, and hypotensive shock (4).

The present study seeks to determine whether the ligand-activated nuclear hormone receptor peroxisome proliferator-activated receptor γ (PPAR γ), which has anti-inflammatory activity (5), acts within pulmonary vascular endothelial cells to protect against endotoxemic lung damage. PPAR γ promotes transcription of genes for anti-inflammatory factors, apparently including the antioxidant nuclear factor (erythroid-derived 2)-like 2 (Nrf2) (6), and also inhibits activity of NF- κ B, AP-1, and other pro-inflammatory transcription factors, in part by competing for essential coactivators (7). PPAR γ is expressed in endothelial cells (8), and PPAR γ agonists exert beneficial effects in animal models of inflammatory diseases (9–14) and sepsis (15), but it is unknown whether PPAR γ acts within endothelial cells to protect against endotoxemia and sepsis. Also, the identities of physiologically relevant endogenous PPAR γ ligands are unresolved. Here we used a transgenic mouse model of targeted endothelial cell PPAR γ deficiency (16) to test the idea that endothelial PPAR γ protects against endotoxemic pulmonary inflammation and lung damage. We also discovered that the nitrated fatty acid (NFA) 10-nitro-oleic acid (OA-NO₂), a candidate physiological PPAR γ

¹Abbreviations used in this paper: BAL, bronchoalveolar lavage; EBD, Evans Blue Dye; eNOS, endothelial nitric oxide synthase; H₂O₂, hydrogen peroxide; iNOS, inducible nitric oxide synthase; IRAK-4, interleukin-1 receptor-associated kinase 4; MDA, malonyldialdehyde; MLEC, mouse lung endothelial cells; NFA, nitrated fatty acid; Nrf2, nuclear factor (erythroid-derived 2)-like 2; PBST, PBS + Tween-20; OA-NO₂, 10-nitro-oleic acid; PPAR- γ , peroxisome proliferator-activated receptor γ ; ROS, reactive oxygen species.

agonist, exerts a broad range of anti-inflammatory actions within pulmonary endothelial cells that are mediated via PPAR γ .

MATERIALS AND METHODS

Animals

Wild-type (ePPAR $\gamma^{+/+}$) and isogenic homozygous endothelial cell PPAR γ knockout (ePPAR $\gamma^{-/-}$) mice on a C57BL/6 genetic background were expanded from breeding pairs. Mice were housed in microisolator cages under specific pathogen-free conditions and fed autoclaved food. Male mice aged 6–8 weeks (20–25 g) were used. All studies were performed according to protocols reviewed and approved by the Atlanta Veterans Affairs Medical Center Institutional Animal Care and Use Committee.

Cells

Mouse lung endothelial cells (MLEC) were isolated as described previously (17). Endothelial identity was confirmed by cobblestone morphology, immunofluorescence staining, and positive Western blotting for PECAM-1, eNOS, vascular endothelial cadherin (BD Pharmingen, San Diego, CA) and negative blots for α -smooth muscle actin (BD Pharmingen). Cells were cultured and grown to confluent monolayers in Vasculife Basal Medium (Lifeline Cell Technology, Frederick, MD) supplemented with 2% FBS, 10 mM L-glutamine, 0.2% EnGS, 5 ng/mL rhEGF, 1 μ g/mL hydrocortisone hemisuccinate, 0.75 units/mL heparin sulfate (Lifeline Cell Technology), 10,000 units/mL penicillin, and 10,000 μ g/mL streptomycin (HyClone, Logan, UT), at 37°C in a humidified atmosphere of 5% CO₂ – 95% air in tissue culture flasks, plates, or dishes coated with 2% gelatin. All cells were used at passages 3–5.

LPS Administration

Mice were injected i.p. with 10 mg/kg of LPS prepared from *Escherichia coli* O111:B6 (Sigma-Aldrich, St. Louis, MO). After a further 12 h the mice were euthanized, blood was obtained by cardiac puncture and plasma prepared, and lung and bronchoalveolar lavage (BAL) fluid samples were collected.

Lung Wet:Dry Weight Ratio

As an index of lung edema, the proportion of extravascular lung water was calculated. The lower lobe of the right lung was ligated, excised and the wet weight was recorded. The lung was then placed in an incubator at 60°C for 24 h, the dry weight recorded, and the wet:dry ratio calculated.

BAL Fluid Collection and Cell Count

Following removal of the lower right lung lobe, BAL fluid was collected by flushing 3 \times 1 ml of PBS containing 0.1 mM EDTA into the lung via a tracheal cannula. The pooled BAL fluid was centrifuged at 500 \times g at 4°C for 5 min. Pelleted cells were then resuspended in 1 ml of PBS. Total cell number was counted by hemocytometer and a differential cell count was performed by cytosin staining with Diff-Quik stain (Siemens, Newark, DE).

BAL Fluid Protein

Increase in BAL fluid protein concentration was taken as a measure of increased permeability of alveolar-capillary barriers. Total protein concentration in the supernatant following BAL fluid centrifugation was determined using the BCA Protein Assay kit (Pierce, Rockford, IL).

Assessment of Capillary Leakage

To assess lung capillary permeability, Evans Blue dye (EBD; 50 mg/kg; Sigma-Aldrich) dissolved in 200 μ l PBS was injected into the tail veins of mice following LPS injection. After 30 min, the animals were euthanized and the lungs perfused with 5 ml of PBS, and the lungs then excised *en bloc* and snap-frozen in liquid nitrogen. The frozen lungs were then homogenized in 2 ml PBS and the homogenates diluted with 2 vol of formamide, then incubated at 60°C for 18 h followed by centrifugation at 5,000 \times g for 30 min. Supernatants were collected and absorbance measured at 620 and 740 nm. The EBD concentration was determined from standard absorbance curves evaluated in parallel. Correction for contaminating heme pigments was calculated by the formula $E_{620}(\text{EBD}) = E_{620} - (1.426 \times E_{740} + 0.030)$. The EBD concentration was expressed as μ g/g lung tissue.

Measurement of Myeloperoxidase Activity

Myeloperoxidase (MPO) activity was determined in BAL fluid and tissue as indices of neutrophil infiltration. Frozen lungs were thawed, weighed, homogenized, and sonicated on ice in radioimmunoprecipitation assay buffer. After centrifugation at 10,000 \times g at 4°C for 20 min, supernatants were collected. MPO activity in supernatants of tissue and BAL fluid were measured by a fluorometric assay (700160; Cayman Chemical, Ann Arbor, MI) and expressed as nmol/min-ml.

Measurement of Oxidant Stress

Hydrogen peroxide (H₂O₂) production in lung tissue was determined using the Amplex Red Hydrogen Peroxide Assay kit (Molecular Probes, Eugene, OR) according to the manufacturer's directions. The concentrations of nitrate and malondialdehyde (MDA) in lung homogenates were measured using colorimetric assay kits (Cayman Chemical).

Measurement of Lung and BAL Fluid Cytokine Levels

Lung and BAL fluid levels of TNF- α , IL-6, KC, and MIP-2 were measured using ELISA kits (R&D Systems, Minneapolis, MN) according to the manufacturer's instructions.

Transcription Factor DNA-binding Activity Assay

Nuclear proteins were extracted using a nuclear extraction kit (Active Motif, Carlsbad, CA) and their concentration was determined using the BCA Protein Assay kit (Pierce). Nuclear extracts were used to quantify DNA-binding activity of PPAR γ , Nrf2, and the p65 subunit of NF- κ B using ELISA-based TransAM kits (40096, 40696, and 50296; Active Motif) according to the manufacturer's instructions.

Western Blotting

Total protein extracts were prepared by lysing cells and homogenizing lung tissue in 500 μ l of ice-cold radioimmunoprecipitation assay buffer supplemented with Halt protease inhibitor cocktail (Pierce). Extracts were incubated for 20 min at 4°C followed by centrifugation at 14,000 \times g, for 15 min. The supernatants were collected and stored at -80°C. Samples were mixed with sample buffer, separated on a 10% SDS-PAGE gel and electroblotted onto a polyvinylidene difluoride membrane (Bio-Rad Laboratories, Hercules, CA). The membrane was blocked with blocking solution (3% BSA in TBST) for 1 h at room temperature. Blots were then incubated overnight at 4°C with primary antibody against the target protein (1:1000). Antibodies against NF κ B-p65, PPAR γ , Nrf2, lamin B1, IRAK-4, TRAF6, and β -actin were from Santa Cruz Biotechnology (Santa Cruz, CA). Antibodies against TLR4, MyD88, p-ERK, p-Akt, p-IKK α / β , p-JNK and p-p38 MAPK were from Cell Signaling Technology (Beverly, MA). The membrane was then washed in TBST and incubated with secondary antibodies consisting of donkey anti-mouse IR 680 (red; LI-COR, Lincoln, NE) and goat anti-rabbit IR 780 (green; LI-COR), diluted 1:5000, for 1 h at room temperature. The infrared signal was detected using an Odyssey Infrared Imager (LI-COR).

Lung Histopathology

The lungs were inflated and fixed with 10% neutral formalin overnight at room temperature. Lung tissue was dehydrated with increasing ethanol concentrations and then embedded in paraffin. Five-micrometer-thick paraffin sections were stained with H&E.

Immunofluorescence Staining and Confocal Imaging

MLEC were cultured on 2% gelatin-coated glass-bottom dishes (MatTek, Ashland, MA) in VasculLife Basal Medium (Lifeline Cell Technology) without growth factors and treated with 100 ng/ml LPS for 6 h. Cells were then washed twice with PBS and subsequently fixed in 10% neutral buffered formalin for 15 min at 37°C. Following fixation, cells were permeabilized with Target Retrieval Solution (Dako, Carpinteria, CA) for 10 min at 95°C, allowed to cool to room temperature, and blocked at 37°C for 1 h with 1% BSA in PBS containing 0.05% Tween-20 (PBST). After being washed, cells were incubated at 37°C for 1 h with primary antibodies to ICAM-1 or PECAM-1 (BD Pharmingen), diluted 1:50 in PBST-1% BSA. After washing with PBST, cells were incubated at 37°C for 1 h with FITC-conjugated goat anti-rat secondary antibody (Jackson ImmunoResearch, West Grove, PA) diluted 1:50 in PBST-1% BSA and then washed three times with PBST. Cover slips were retrieved and mounted on glass slides with Vectashield mounting medium with DAPI (Vector Laboratories, Burlingame, CA). The slides were viewed by an Olympus Fluoview FV1000 confocal microscope (Olympus, Center Valley, PA) using a 60 \times fluorescence lens along with Fluoview confocal software (FV10-ASW v1.7, Olympus).

Determination of Cellular ROS

MLEC were cultured and treated with LPS as described under *Immunofluorescence Staining*. Production of intracellular ROS in live cells was determined using the Amplitude Intracellular Fluorimetric Hydrogen Peroxide Assay kit (AAT Bioquest, Sunnyvale, CA). Cover slips were retrieved and mounted on glass slides with Vectashield mounting medium

(Vector Laboratories, Burlingame, CA). The slides were viewed by an Olympus Fluoview FV1000 confocal microscope (Olympus) using a 60 × fluorescence lens along with Fluoview confocal software (FV10-ASW v1.7, Olympus).

***In vitro* Neutrophil Transmigration Assay**

MLEC were seeded at a density of 5×10^5 cells in the upper chamber of each Transwell (6.5 mm diameter, 3 μm pore size; Costar, Corning, NY) and the cells were grown to form a confluent monolayer. In some experiments cells were treated with 100 nM OA-NO₂ or DMSO along with 100 ng/ml LPS or with PBS and incubated for 6 h. Neutrophils isolated from BAL fluid of LPS-stimulated ePPAR $\gamma^{+/+}$ mice were added to the top chamber at 4×10^5 cells/well and, where indicated, chemoattractant was added to the lower chamber. The neutrophils were incubated at 37°C for 1.5 h, after which neutrophils that had transmigrated into the lower chamber were collected, counted, and transmigration expressed as % of neutrophils added.

RNA Isolation and Quantitative Real-Time RT-PCR

MLEC were cultured and treated with LPS as described under *Immunofluorescence Staining*, but were pre-treated with 100 nM OA-NO₂ or DMSO for 1 h prior to LPS exposure. After 6 h, cells were lysed and RNA was isolated using RNeasy Mini kit (Qiagen, Valencia, CA), and cDNA generated from 100 ng of total RNA using MultiScribe reverse transcriptase (Applied Biosystems, Foster City, CA) employing random and oligo-dT primers. Real-time quantitative PCR was performed using 100 ng cDNA with 2X SYBR Green Master mix (Applied Biosystems) and specific primers for the genes of interest (Supplemental Table 1). These experiments were performed on an AB 7500 fast thermal cycler using a three-step protocol employing the melting curve method. The average of each gene cycle threshold (C_t) was determined for each experiment. Relative cDNA levels (2^{-C_t}) for the genes of interest were determined by using the comparative C_t method, which generates the C_t as the difference between the gene of interest and the housekeeping genes GAPDH and 9s rRNA for each sample. Each averaged experimental gene expression sample was compared to the averaged control sample, which was set to 1.

Statistical Analysis

Data are presented as mean ± SD. Differences between groups were analyzed using ANOVA, followed by a Bonferroni multiple comparison test using GraphPad Prism 5.03 (GraphPad Software, La Jolla, CA). $P < 0.05$ was considered significant.

RESULTS

Endotoxemia-induced Capillary Permeability Increases and Lung Injury Is More Severe in Endothelial PPAR γ KO Mice

To test our hypothesis that endothelial cell PPAR γ modulates endotoxemia-induced lung injury, we compared lung edema and inflammation in endothelial PPAR γ knockout (ePPAR $\gamma^{-/-}$) and wild-type (ePPAR $\gamma^{+/+}$) control mice 12 h after treatment with LPS (10 mg/kg, i.p.). ePPAR γ deficiency markedly increased all measures of lung injury-associated pulmonary edema and capillary leakage. LPS-induced increases in protein content of BAL

fluid (Fig. 1A), an indicator of elevated capillary permeability, and lung wet:dry tissue weight ratio (Fig. 1B) were nearly doubled in ePPAR γ ^{-/-} vs. ePPAR γ ^{+/+} mice, as was extravasation of Evans Blue dye 30 min after i.v. injection (Fig. 1C). Both wild-type (Fig. 1D, left) and ePPAR γ KO mice (Fig. 1D, right) treated with LPS exhibited leukocyte infiltration, alveolar distortion and edema on histological examination, but these were more severe in ePPAR γ KO animals.

ePPAR γ Deletion Elevates Production of ROS and Pro-inflammatory Cytokines

To assess the influence of ePPAR γ on endotoxemia-associated mediators of pulmonary inflammation *in vivo*, we compared pulmonary LPS-induced elevations of relevant ROS and pro-inflammatory cytokines in ePPAR γ ^{-/-} and ePPAR γ ^{+/+} mice. ROS production was assessed in lung tissue and BAL fluid by measuring 3 different indices: H₂O₂ production (Fig. 2A, D); concentrations of nitrate, which is an end product of the key vasodilator NO generated by a different pathway than H₂O₂ (Fig. 2B, E); and MDA, a lipid oxidation product that provides an index of overall oxidant stress (MDA:protein ratio; Fig. 2C, F). LPS treatment elevated the levels of all these markers in both mouse strains, but the LPS-induced increases seen were significantly greater in ePPAR γ KO mice for all ROS-related analytes tested. We also measured LPS-induced elevations in expression within lung tissue and BAL fluid of the pro-inflammatory cytokines IL-6 (Fig. 2G, J) and TNF- α (Fig. 2H, K), for which endothelial cells are important sources, as well as the chemokine KC (Fig. 2I, L). Again, LPS treatment elicited increased cytokine/chemokine levels in both mouse strains, but the increases seen in all markers were significantly greater in ePPAR γ KO than in wild-type mice. These results indicate that the greater severity of inflammation seen in ePPAR γ KO mice is associated with elevations in a range of ROS and key inflammatory cytokines.

Endotoxemia-induced Neutrophil Infiltration into Alveolar Space Is Increased in ePPAR γ KO Mice

Endotoxemia-induced inflammation is associated with infiltration of neutrophils into target tissues including the lung. We found that multiple measures of LPS-induced neutrophil infiltration were markedly increased in ePPAR γ KO vs. in wild-type mice, including: total numbers of cells in alveolar space, assessed by measuring those recovered in BAL fluid (Fig. 3A); numbers of neutrophils identified by differential staining (Fig. 3B); and MPO levels in BAL fluid (Fig. 3C) and whole lung tissue (Fig. 3D). Microscopic examination of BAL fluid with differential staining similarly showed that ePPAR γ deficiency led to greater LPS-induced increases in both total cell numbers and proportion of neutrophils (Fig. 3E).

Genetic Deletion of ePPAR γ Upregulates LPS Signaling

To determine the mechanisms underlying the exaggerated inflammatory responses seen in ePPAR γ KO mice, we measured the levels of LPS-associated intermediate signaling proteins following induction of endotoxemia, using Western blots of lung tissue extracts. After LPS treatment, ePPAR γ ^{-/-} mice exhibited greater expression of TLR4 and the TLR signaling proteins MyD88, IRAK-4, and TRAF6 than did ePPAR γ ^{+/+} mice (Fig. 4A). ePPAR γ KO mice similarly exhibited greater levels of the phosphorylated (activated) forms

of downstream signaling mediators which serve to activate the transcription factors AP-1 and NF- κ B, than did ePPAR $\gamma^{+/+}$ mice.

Absence of ePPAR γ Exaggerates LPS-induced Changes in Activity of NF- κ B and Nrf2

Endotoxemia-induced inflammation involves upregulation of pro-inflammatory transcription factors including NF- κ B, which is transrepressed by PPAR γ , as well as downregulation of the antioxidant transcription factor Nrf2, which evidence suggests is upregulated by PPAR γ (6). To test the influence of PPAR γ on LPS-stimulated expression and activity of these signaling proteins *in vivo*, we compared their levels in ePPAR γ KO and wild-type control mice. Following LPS treatment, ePPAR γ KO mice exhibited elevated nuclear p65 and decreased nuclear Nrf2 protein levels vs. those in wild-type controls (Fig. 4C), paralleled by an LPS-induced upregulation of NF- κ B activity (Fig. 4F) and a decrease in Nrf2 activity (Fig. 4E). Knockout of PPAR γ specifically in endothelial cells did not significantly decrease baseline whole-lung PPAR γ activity (Fig. 4D), but it abolished the LPS-induced increase.

PPAR γ Deletion Exaggerates Inflammatory Responses of Endothelial Cells *in Vitro*

To ascertain whether ePPAR γ acts within lung endothelial cells to modulate inflammatory responses, we compared expression of inflammatory markers induced by LPS *in vitro*, in MLEC isolated from ePPAR $\gamma^{-/-}$ and ePPAR $\gamma^{+/+}$ mice. Endothelial identity of the isolated cells was confirmed microscopically (Fig. 5A) and by Western blotting (Fig. 5B). Endothelial cells exhibited the expected abundant expression of the endothelial isoform of nitric oxide synthase and the adhesion molecules vascular endothelial cadherin, the related PECAM-1, and near-absence of α -smooth muscle actin, while isolated fibroblasts exhibited an opposite expression profile. We confirmed the absence of PPAR γ mRNA and of PPAR γ protein in endothelial cells from ePPAR γ KO mice by real-time RT-PCR and Western blotting respectively (Fig. 5C, E). After stimulation of MLEC from wild-type and ePPAR γ KO mice *in vitro* with LPS (100 ng/ml) for 6 h, PPAR γ -deficient MLEC exhibited increased expression of the inflammation-associated adhesion molecules ICAM-1 (Fig. 5F) and PECAM-1 (Fig. 5G), together with increased ROS production (Fig. 5H) vs. wild type MLEC. These results indicate that PPAR γ acts within MLEC to reduce LPS-induced inflammatory markers.

To determine the effects of endotoxemia on PPAR γ , endothelial cells were isolated from ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice 12 h after stimulation *in vivo* with PBS or LPS. While there was a significant reduction in PPAR γ protein expression (Fig. 5E), a significant increase in PPAR γ activity (Fig. 5E) was seen in endothelial cells from LPS-stimulated ePPAR $\gamma^{+/+}$ mice compared to PBS-stimulated ePPAR $\gamma^{+/+}$ mice. These data support the concept that LPS induces a PPAR γ response via increased production of activating endogenous ligands.

ePPAR γ Deletion Abolishes Anti-inflammatory Effect of a Nitrated Fatty Acid

PPAR γ is ligand-activated, but it is unknown which of several classes of candidate endogenous ligands may be physiologically relevant. Nitrated fatty acids, including NO-derived positional isomers of nitro-oleic acid and nitrolinoleic acid, comprise one such group of candidate ligands (18, 19). Their concentrations in plasma are similar to their

potencies as PPAR γ activators, suggesting a physiological role as PPAR γ agonists (18). As agonists of the anti-inflammatory PPAR γ , NFAs would be expected to exert anti-inflammatory effects, but some findings suggest that the anti-inflammatory effects of NFAs may not be mediated exclusively by PPAR γ activation. We tested these ideas by assessing the effects of OA-NO₂ on LPS-induced inflammatory responses within MLEC *in vitro* and the extent of these effects' dependence on PPAR γ .

We first tested the ability of OA-NO₂ to inhibit LPS-induced cytokine production and expression of pro-inflammatory mediators and adhesion molecules in MLEC. MLEC cultured on 2% gelatin-coated culture dishes without growth factors were pre-treated with OA-NO₂ at a concentration similar to that found normally in blood (100 nM) or with DMSO vehicle for 1 h, then treated with 100 ng/ml LPS and incubated for another 6 h. OA-NO₂ markedly inhibited the robust LPS-induced transcription of all the following inflammation-associated genes in cells from wild-type mice (Fig. 6A–L): the adhesion-related molecules E-selectin, ICAM-1, PECAM-1, and VCAM-1; the enzymes NOS3, NOS2, NADPH oxidase 4, and cyclooxygenase-2; the chemokine MCP-1; the cytokines IL-6 and TNF- α ; and the p65 subunit of NF- κ B. The ability of OA-NO₂ to reduce LPS-induced expression of each of these genes was completely abolished in MLEC from PPAR γ KO mice, indicating strong dependence on the presence of ePPAR γ . This was likewise true for expression of the adhesion molecule VE-cadherin (Supplemental Fig. 1A) and the two isoforms of IL-12 (Supplemental Fig. 1B, C). We also found that in wild-type cells OA-NO₂ upregulated expression of PPAR γ , of the scavenger receptor CD36, which is involved in resolution of inflammation, of mitochondrial uncoupling protein 2, and of the antioxidant enzyme heme oxygenase-1 (Fig. 6M–P). These upregulating actions of OA-NO₂ were likewise abolished by PPAR γ KO. These findings indicate that OA-NO₂ at a physiologically relevant concentration exerts a range of anti-inflammatory effects directly on endothelial cells via a PPAR γ -dependent mechanism.

To assess the influence of NFAs on a major cellular route of lung inflammation, we also tested the ability of OA-NO₂ to inhibit neutrophil transmigration across MLEC monolayers *in vitro*. Endothelial cells were isolated from wild-type and PPAR γ KO mice and cultured in monolayers in the upper chambers of transwells. Transmigration was assessed after pretreating the MLEC monolayers with LPS along with OA-NO₂ or vehicle (Fig. 7A). Wild-type neutrophils were then placed in the upper chamber, the chemoattractant stimulus (LPS + TNF- α) added to the lower chamber, and the number of neutrophils entering each lower chamber in 1.5 h were determined. In wild-type cells, OA-NO₂ significantly inhibited neutrophil transmigration (by 45–65%), irrespective of whether LPS was present in the upper chamber (Fig. 7B, C). The extent of OA-NO₂-induced inhibition of neutrophil transmigration was substantially reduced (to 9–13%) in ePPAR γ ^{-/-} MLEC, although it did remain statistically significant. The results suggest that the anti-inflammatory effects of OA-NO₂ include inhibition of neutrophil transmigration across the endothelium, and are predominantly PPAR γ -dependent.

DISCUSSION

Our findings indicate that ePPAR γ plays an important protective role in pulmonary endothelial cells during endotoxemia, substantially mitigating lung injury and inflammation. Deficiency of ePPAR γ exacerbated LPS-induced pulmonary damage (edema, capillary leakage) and inflammation *in vivo*, and intensified the activation by LPS of numerous pro-inflammatory genes, signaling proteins, cytokines and other mediators both *in vivo* and *in vitro*. In isolated MLEC, robust LPS-induced increases seen in adhesion proteins and pro-inflammatory cytokines, chemokines and ROS-generating enzymes were similarly enhanced by ePPAR γ deficiency. This indicates that ePPAR γ modulates LPS responses by acting within endothelial cells in a manner that is not dependent on participation of other cell types. We found that ePPAR γ suppressed LPS-induced expression of endothelial pro-inflammatory mediators, including cytokines and adhesion molecules that facilitate migration of neutrophils into alveolar spaces, as well as LPS-induced pathophysiological deficits in endothelial cell function that lead to increased capillary permeability and lung edema. Together, these results clearly underscore the roles of endothelial cells as robust mediators and pathogenic targets of LPS during endotoxemia, and point to ePPAR γ acting within endothelial cells as a strong suppressor of endothelial LPS-induced cellular responses.

The exaggerated LPS effects we saw in ePPAR γ -deficient mice encompassed upregulated expression of LPS receptors along with increased activation of their downstream LPS signaling pathways. The first step in LPS activation of TLR4 is formation of a ternary complex with LPS binding protein and CD14, which then interacts with TLR4 and the adaptor protein MD2, leading to activation of MyD88, which activates IRAK-4 and TRAF6 (20). We found that ePPAR γ deficiency increased the robust LPS-induced upregulation of all these proteins in lungs of mice and that increased expression of additional signaling intermediates with ePPAR γ deficiency was associated with increased activation of NF- κ B. The enhanced activity of the latter pro-inflammatory transcription factors seen in ePPAR γ -deficient mice probably accounts for the enhanced expression we saw in a variety of pro-inflammatory cytokines, chemokines, and adhesion molecules. In addition to these observed effects on TLR signaling pathways and intermediates, PPAR γ also inhibits activity of NF- κ B, AP-1, and similar pro-inflammatory transcription factors directly, by competing for essential coactivators (7) and by other mechanisms. Although not measured directly in our present study, these known actions likely also contributed to the anti-inflammatory effects we observed.

We found that ePPAR γ represents a highly specific, LPS-inducible compartment of total PPAR γ activity in lungs, because ePPAR γ knockout had no effect on total baseline tissue PPAR γ activity levels in lung but abolished the LPS-induced increase in whole-lung PPAR γ activity seen in wild-type mice. This implies ePPAR γ activity is highly recruited in endothelial cells during sepsis and probably accounts for the entire LPS-induced elevation of whole-lung PPAR γ activity. Alternatively, it is conceivable that LPS-activated ePPAR γ somehow recruits expression of PPAR γ in other cell types, contributing part of the total response seen in PPAR γ activity. Considering the broad range and strength of the downregulating actions of ePPAR γ upon LPS responses that we observed, our results support the idea that LPS-induced upregulation of ePPAR γ activity constitutes an important

endogenous pathway for adaptive restraint of inflammatory responses, particularly under the relatively severe conditions (LPS dose) tested here.

ROS contribute both to the adaptive (antibacterial) and maladaptive (tissue-damaging) effects of inflammation. The transcription factor Nrf2 acts to reduce ROS-induced tissue damage by promoting expression of proteins that have antioxidant effects. Additionally, Nrf2 activates the PPAR γ transcriptional promoter when transfected into airway epithelial cells, while global deficiency of Nrf2 blocked the ability of PPAR γ ligands to suppress hyperoxia-induced lung inflammation and injury (6). Presently, we found that LPS reduced Nrf2 activity, and that ePPAR γ deficiency exaggerated the LPS-induced fall in Nrf2 levels, which would thus tend to reduce endogenous protection against deleterious effects of the high ROS levels occurring during sepsis. These findings indicate that the promotion of Nrf2 expression by PPAR γ is not limited to hyperoxia but also occurs in the high-ROS context of inflammation and sepsis. Also, these effects of PPAR γ are seen specifically in endothelial cells. Thus, the more pronounced LPS-induced decline in lung tissue Nrf2 levels caused by ePPAR γ -deficiency probably contributed to the elevated LPS-induced ROS production seen in isolated PPAR γ -deficient MLEC.

We found that ePPAR γ deficiency in endothelial cells increased their LPS-induced expression of adhesion proteins including ICAM-1, VCAM-1 and E-selectin, indicating that ePPAR γ suppresses induction of adhesion proteins. This action may profoundly influence pulmonary inflammation, because production of ICAM-1 and VCAM-1 is required for transmigration of neutrophils into the lungs as well as into other organs. Similarly, Wang and colleagues found that constitutively active PPAR γ suppressed TNF- α - and PMA-induced VCAM-1, ICAM-1 and E-selectin expression and neutrophil adhesion to HUVECs (21). Also, PPAR γ agonists (22–25) and compounds that upregulate PPAR γ expression (26, 27) have generally been found to inhibit neutrophil adhesion and VCAM-1 expression elicited by pro-inflammatory stimuli. Not all studies have concurred about whether ICAM-1 is downregulated by PPAR γ agonists, and the reported effects of individual agonists on adhesion protein expression have been variable (23, 24, 26). The reasons for such apparent discrepancies among reported actions of PPAR γ agonists are unknown, but are not thought to be attributable to differences in cell origin or in pro-inflammatory stimulus, although they may reflect differential dosage effects. In any case, the observed results of altered PPAR γ expression have been highly consistent, supporting the idea that PPAR γ activation serves to inhibit expression of inflammation-promoting adhesion proteins.

Our findings also point to a role of NFAs as potential endogenous agonists that might contribute to the observed actions of the agonist-driven transcriptional activator PPAR γ . A variety of endogenous compounds can bind to and activate PPAR γ and have thus been proposed as endogenous agonists, but the physiological relevance of most such agents is doubtful due to insufficient potencies in relation to *in vivo* concentration ranges. The most plausible identified candidates to date are NFAs, especially positional isomers of nitro-oleic and nitrolinoleic acids (18, 19), which are produced by nonenzymatic reactions of NO and its inorganic reaction products with endogenous unsaturated fatty acids (28). A role of endogenous NFAs as anti-inflammatory mediators has been thought plausible because they are present in body fluids (29), are active at physiological concentrations, and the production

of NFA-generating NO is increased during inflammation. We found that exposing MLEC to the NFA OA-NO₂ partially or completely blocked LPS-induced expression of adhesion molecules, cytokines, chemokines, and inflammation-associated enzymes in MLEC, in a manner that was completely dependent upon ePPAR γ expression. All these wide-ranging anti-inflammatory effects were elicited by a concentration of NFA OA-NO₂ that is well within its endogenous concentration range (30). Also, OA-NO₂ upregulated the expression by MLEC of several anti-inflammatory molecules, including PPAR γ , indicating that the observed inhibitory effects on the expression of pro-inflammatory LPS-induced proteins and mediators are not attributable to any nonspecific toxicity. In addition, we discovered that OA-NO₂ inhibited the transmigration of neutrophils across endothelial cell monolayers. Neutrophil transmigration is a function that is fundamental to inflammation, and we have previously found that it is inhibited by PPAR γ activation (31). We now show that NFA-mediated inhibition of neutrophil transmigration occurred under both baseline and LPS-stimulated conditions, and is partially dependent on expression of PPAR γ by the MLEC. These findings are consistent with, but qualitatively expand, the reported anti-inflammatory actions of NFAs, including inhibition of TNF- α -induced macrophage VCAM-1 and monocyte rolling and adhesion (30); inhibition of superoxide generation, calcium influx, elastase release, and CD11b expression by human neutrophils (32); and upregulation of the anti-inflammatory enzyme heme oxygenase-1 in human aortic endothelial cells (33). Our observations thus support a role for NFAs as physiologically relevant endogenous PPAR γ agonists. Further research is required to confirm this possibility *in vivo*.

Inflammatory responses to pathogens can be destructive and often lethal to the host if unchecked. Endogenous systems operating at many levels have evolved to restrain such damage, while permitting effective antimicrobial defense. Our findings indicate that PPAR γ acts within pulmonary endothelial cells during endotoxemia to exert such a key mitigating effect, inhibiting the many pro-inflammatory responses elicited by LPS in endothelial cells, thereby restraining inflammation and reducing pulmonary injury. We also addressed the unresolved question of which candidate molecule(s) serve as endogenous PPAR γ agonists by finding unequivocal and novel evidence that the endogenous NFA OA-NO₂ completely blocks a wide range of LPS-induced pro-inflammatory responses in MLEC by via a mechanism specifically dependent on PPAR γ . As our studies were confined to pulmonary endothelial cells and lung tissue, further research will be needed to determine whether they are also broadly applicable to inflammatory pathogenesis involving endothelial cell populations in other organs. These results may help point the way to new pharmacological interventions for sepsis and lung injury.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgment

This work was supported by NIH Grant HL093196 (R.C.R.).

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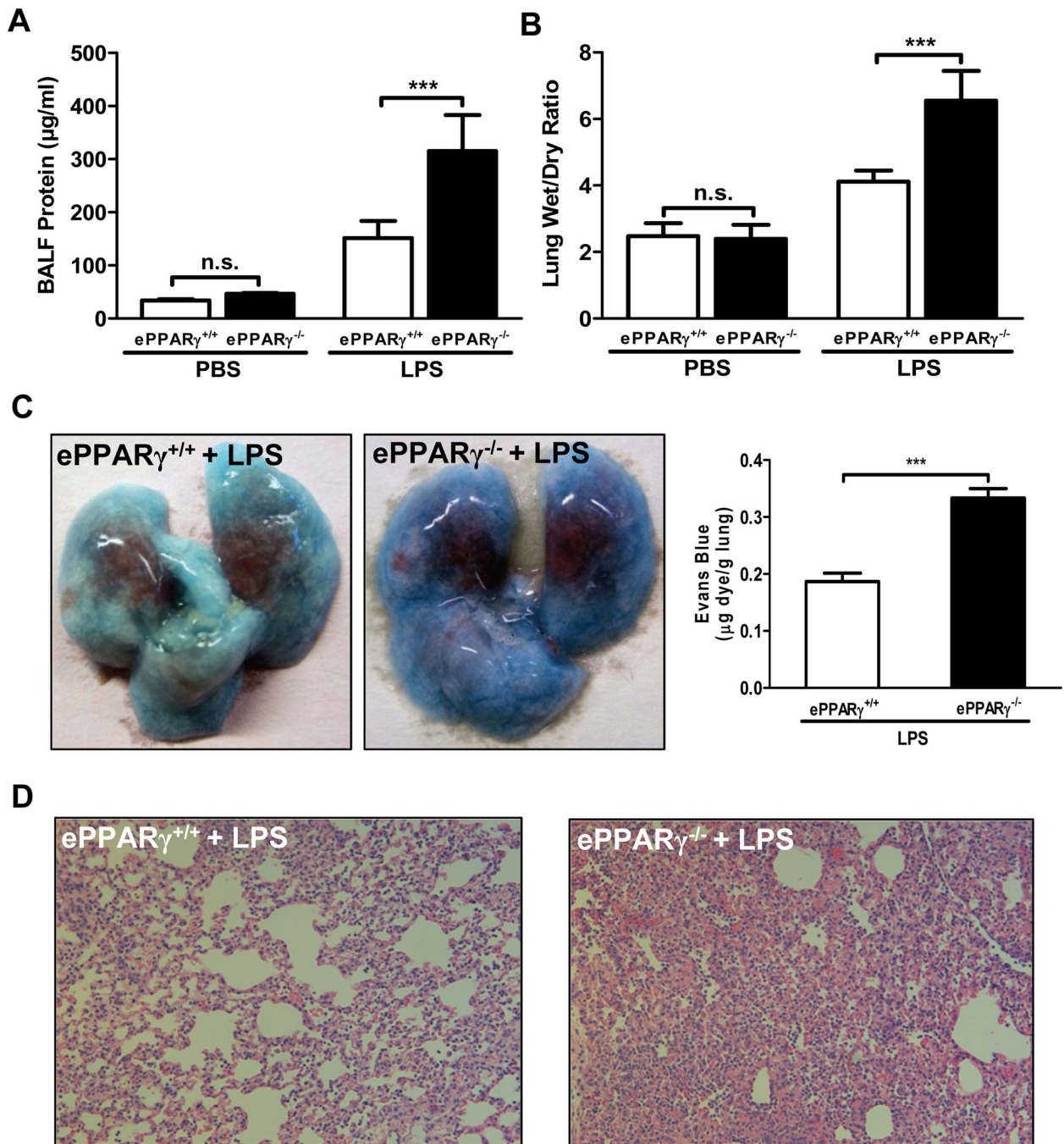


Figure 1. ePPAR γ Deficiency Exacerbates LPS-induced Lung Injury

All measures of LPS (10 mg/kg, i.p.)-induced (A, C) capillary leakage, (B) edema, and (D) inflammation were markedly elevated in ePPAR $\gamma^{-/-}$ mice vs. ePPAR $\gamma^{+/+}$ controls. Mice were euthanized, BAL fluid collected, and lungs excised 12 h after injection. (A) Protein concentration in BAL fluid. (B) Ratio of lung tissue wet:dry weight. (C) Extravasation of Evans Blue dye into the lung after i.v. injection, photographed (*left panel*; note lighter color) and quantitated by spectrophotometry (*right panel*). (D) Histological appearance with H&E staining, showing intensified infiltration by inflammatory cells in ePPAR $\gamma^{-/-}$ mice (*right*

panel). Each data set is representative of 2–3 independent experiments with $n = 6–8$ mice per group; *** $P < 0.001$.

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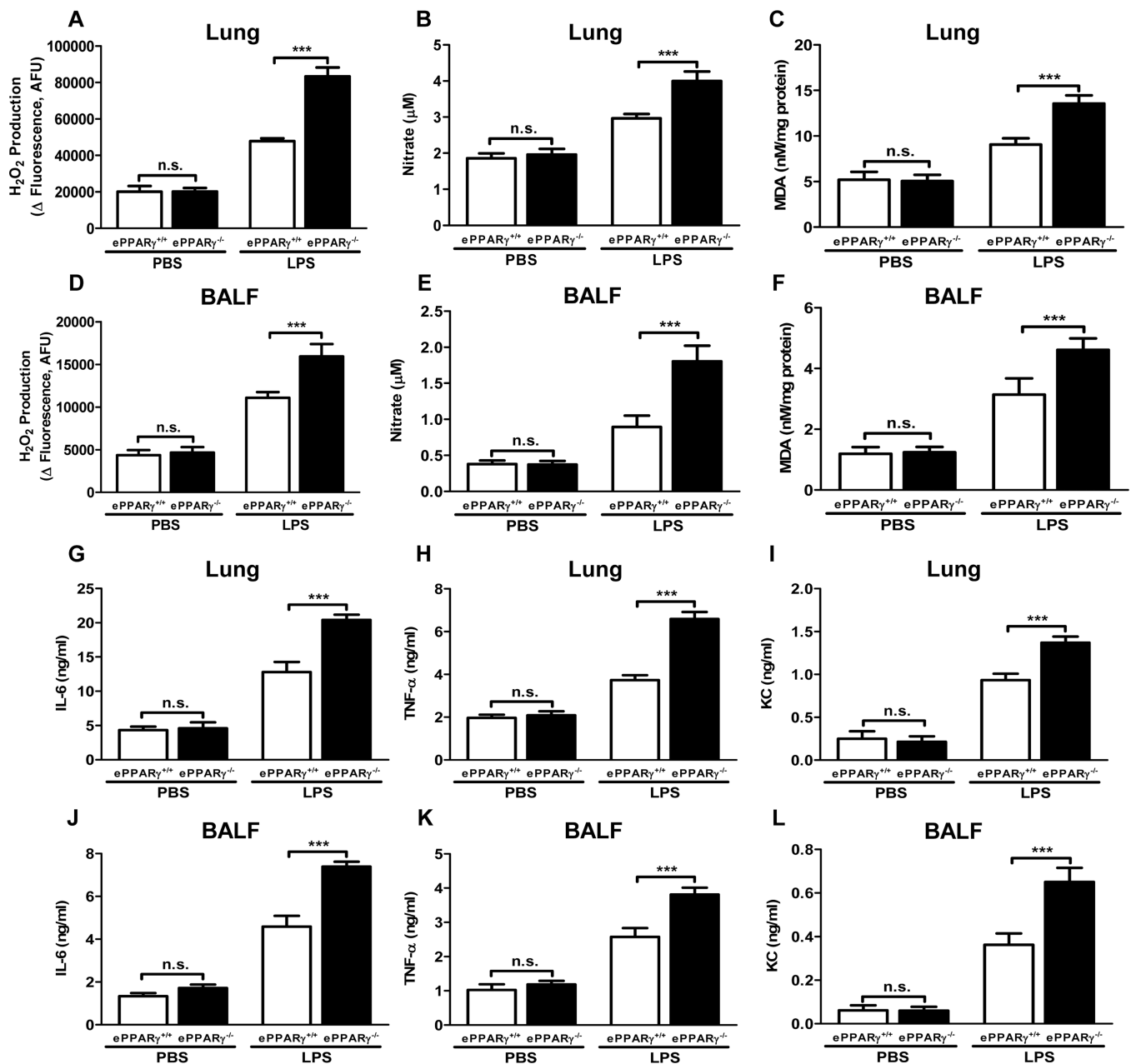


Figure 2. Production of ROS and Pro-inflammatory Cytokines Is Elevated Following ePPAR γ Deletion

ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice were treated with LPS (10 mg/kg, i.p.), and after 12 h ROS and cytokine levels were measured in BAL fluid and lung tissue extracts. LPS-induced elevations in ROS and cytokine levels were increased by ePPAR γ KO in lung and BAL fluid (BALF) for all analytes: (A, D) H₂O₂; (B, E) total nitrate concentration; (C, F) malonaldehyde (MDA)/protein ratio; (G, J) IL-6, (H, K) TNF- α ; and (I, L) KC. Data are representative of 3 independent experiments with $n = 6-8$ mice per group; *** $P < 0.001$.

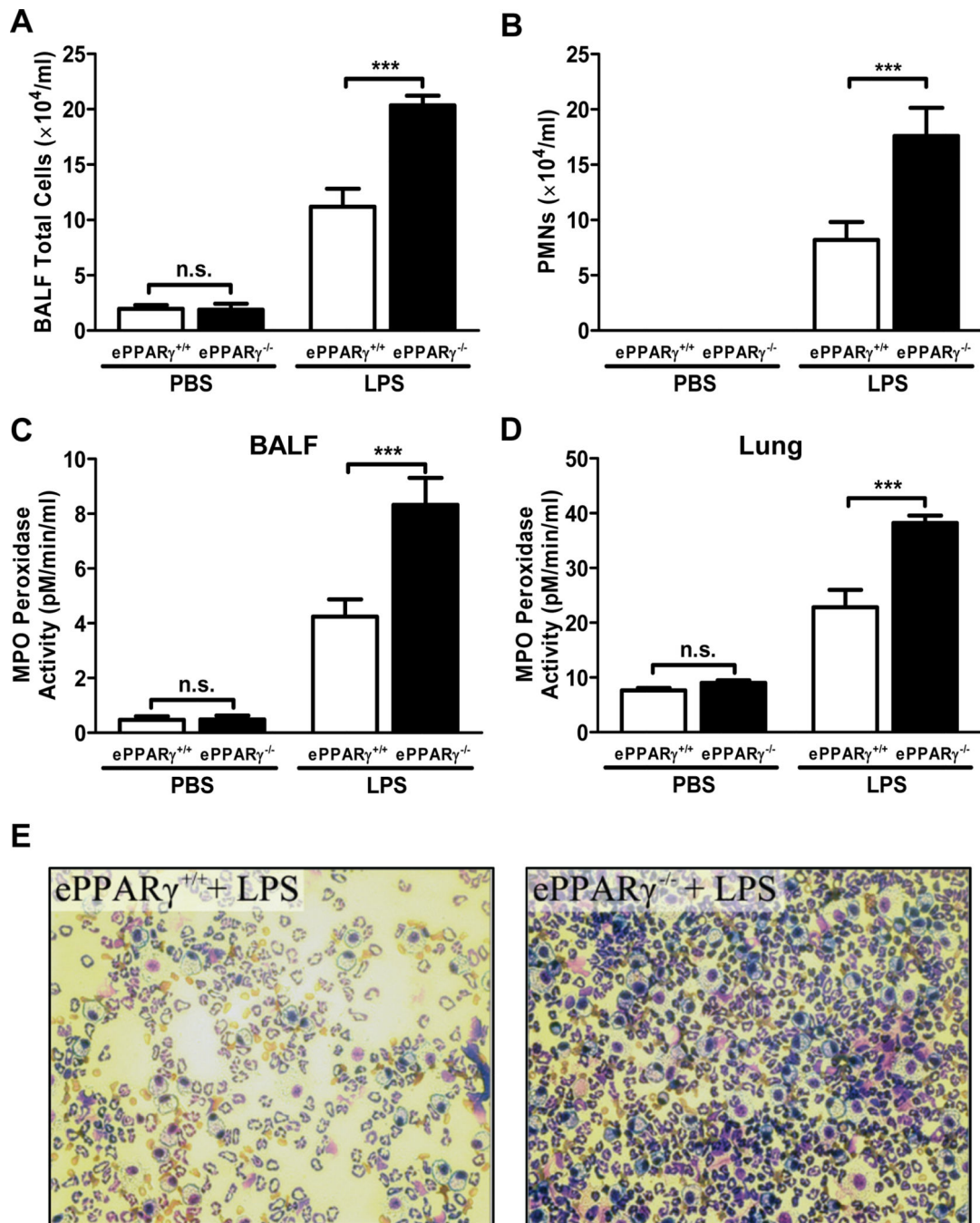


Figure 3. Migration of Neutrophils into the Lung Is Increased in ePPAR γ KO Mice
 ePPAR $\gamma^{-/-}$ mice exhibited greater elevations of total numbers of (A) cells, (B) neutrophils (PMNs), and (C) myeloperoxidase activity (MPO) in BAL fluid (BALF), as well as (D) MPO activity in lung tissue 12 h after injection of LPS (10 mg/kg, i.p.). (E) Differential staining of cells in BAL fluid revealed increased numbers of PMNs in ePPAR $\gamma^{-/-}$ (*right panel*) vs. that seen in ePPAR $\gamma^{+/+}$ mice (*left panel*). Data are representative of 2 independent experiments with $n = 6-8$ mice per group; *** $P < 0.001$.

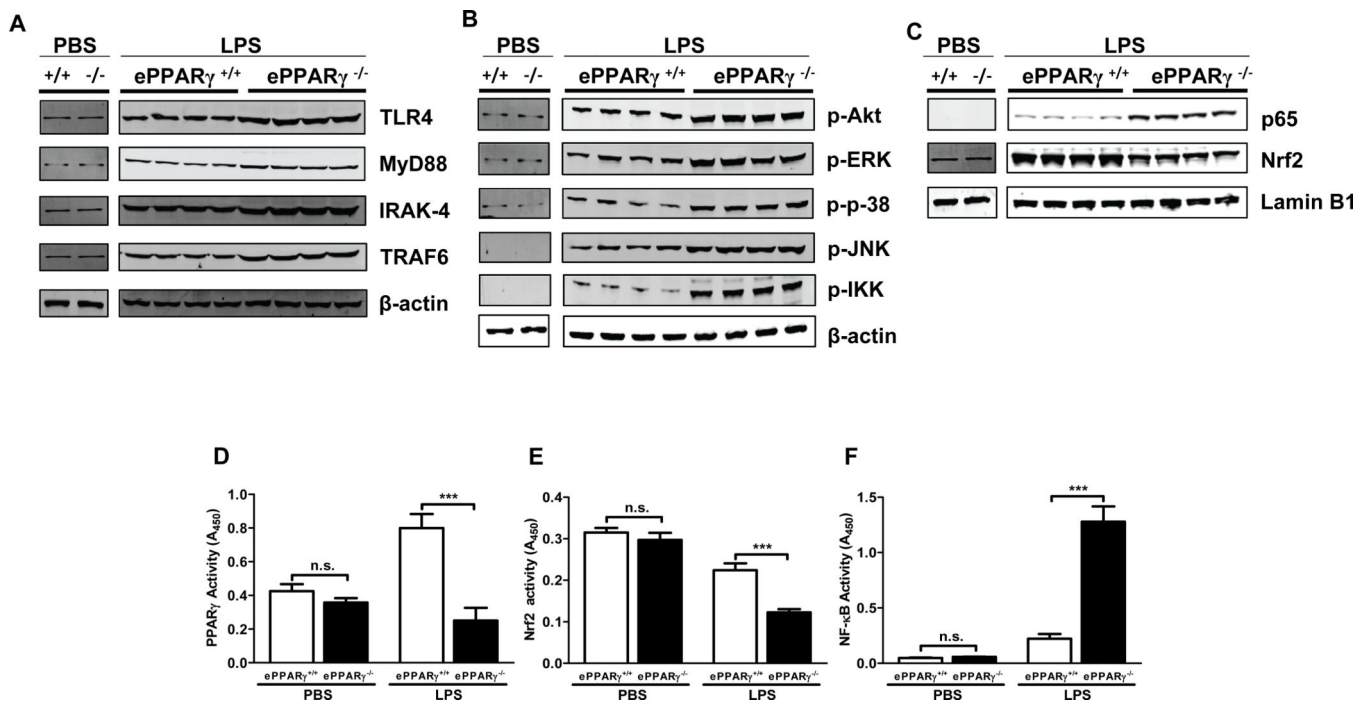


Figure 4. Genetic Deletion of ePPAR γ Upregulates *in vivo* LPS Signaling
 ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice were treated with LPS (10 mg/kg, i.p.). Twelve h later lung tissue protein extracts were prepared and the levels of LPS signaling proteins analyzed. Lungs of ePPAR $\gamma^{-/-}$ mice exhibited higher levels of (A) the LPS receptor TLR4 and its downstream signaling molecules MyD88, IRAK-4, and TRAF6; B) the phosphorylated (activated) forms of intermediate signaling molecules further downstream (Akt, ERK, p38, JNK, IKK α/β); and (C) the transcription factors NF- κ B (p65) and Nrf2. Blots were simultaneously incubated with lamin B1 or β -actin antibody where indicated. Smaller blot represents PBS treated ePPAR $\gamma^{+/+}$ mice (left), ePPAR $\gamma^{-/-}$ mice (right). In larger blot left four lanes represent LPS treated ePPAR $\gamma^{+/+}$ mice, right four lanes represent LPS treated ePPAR $\gamma^{-/-}$ mice. In separate experiments we also determined DNA-binding activities of transcription factors in lung extracts. ePPAR γ KO abolished the LPS-induced component of (D) PPAR γ activity, while (E) enhancing the LPS-induced decrease in Nrf2 activity and (F) markedly elevating LPS-induced NF- κ B activity. Blots are representative of 3 independent experiments with each lane representing a single mouse lung, and activity data of 2 independent experiments ($n = 6-8$ mice per group). *** $P < 0.001$.

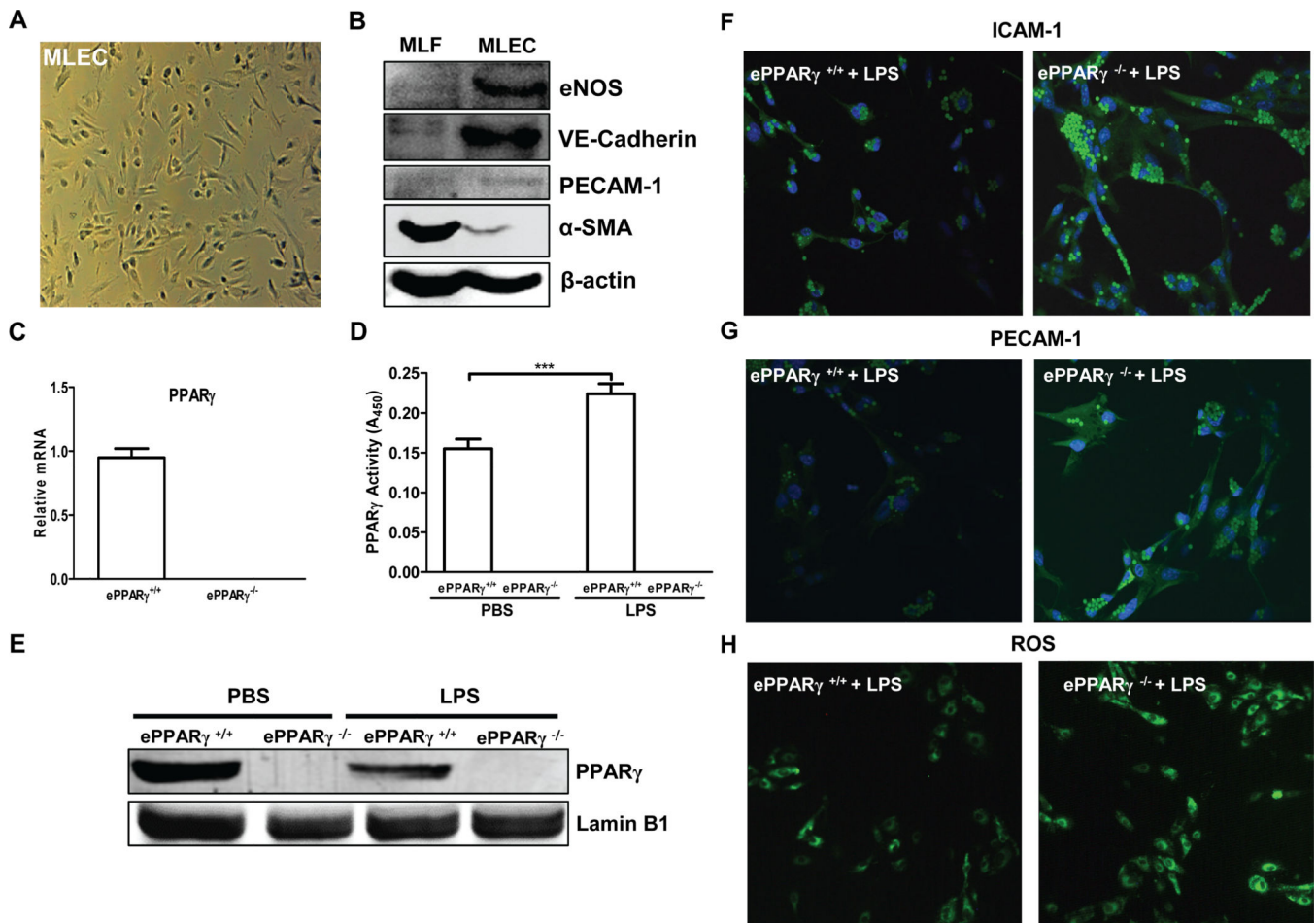


Figure 5. Absence of ePPAR γ Exaggerates Inflammatory Responses of MLEC *in Vitro*
 MLEC were isolated from ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice ($n = 7-8$ mice per group) and plated onto 2% gelatin-coated dishes. (A) MLEC exhibit expected endothelial cell morphology. (B) Western blotting showed the expected expression profiles in MLEC and similarly isolated fibroblasts (MLF), respectively, of the endothelium-specific proteins eNOS, vascular endothelial (VE) cadherin, and PECAM-1 and the fibroblast-specific protein α -smooth muscle actin (α -SMA). (C) Relative PPAR γ mRNA levels were determined by real-time RT-PCR. In some experiments MLEC were isolated from ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice 12 h after treatment with LPS (10 mg/kg, i.p.), and nuclear protein extracts prepared. (D) PPAR γ DNA-binding activity and (E) protein expression by Western blotting were determined. In other experiments after 2 h serum deprivation, monolayer cultures were treated with LPS (100 ng/ml) for 6 h. Following LPS treatment, immunofluorescence microscopy revealed increased ICAM-1 and PECAM-1 in MLEC isolated from (F,G; right panels) ePPAR $\gamma^{-/-}$ vs. those from (F,G; left panels) ePPAR $\gamma^{+/+}$ mice, and also elevated (H; right panel vs. left panel) intracellular ROS levels. Blots and images are representative of 3 independent experiments.

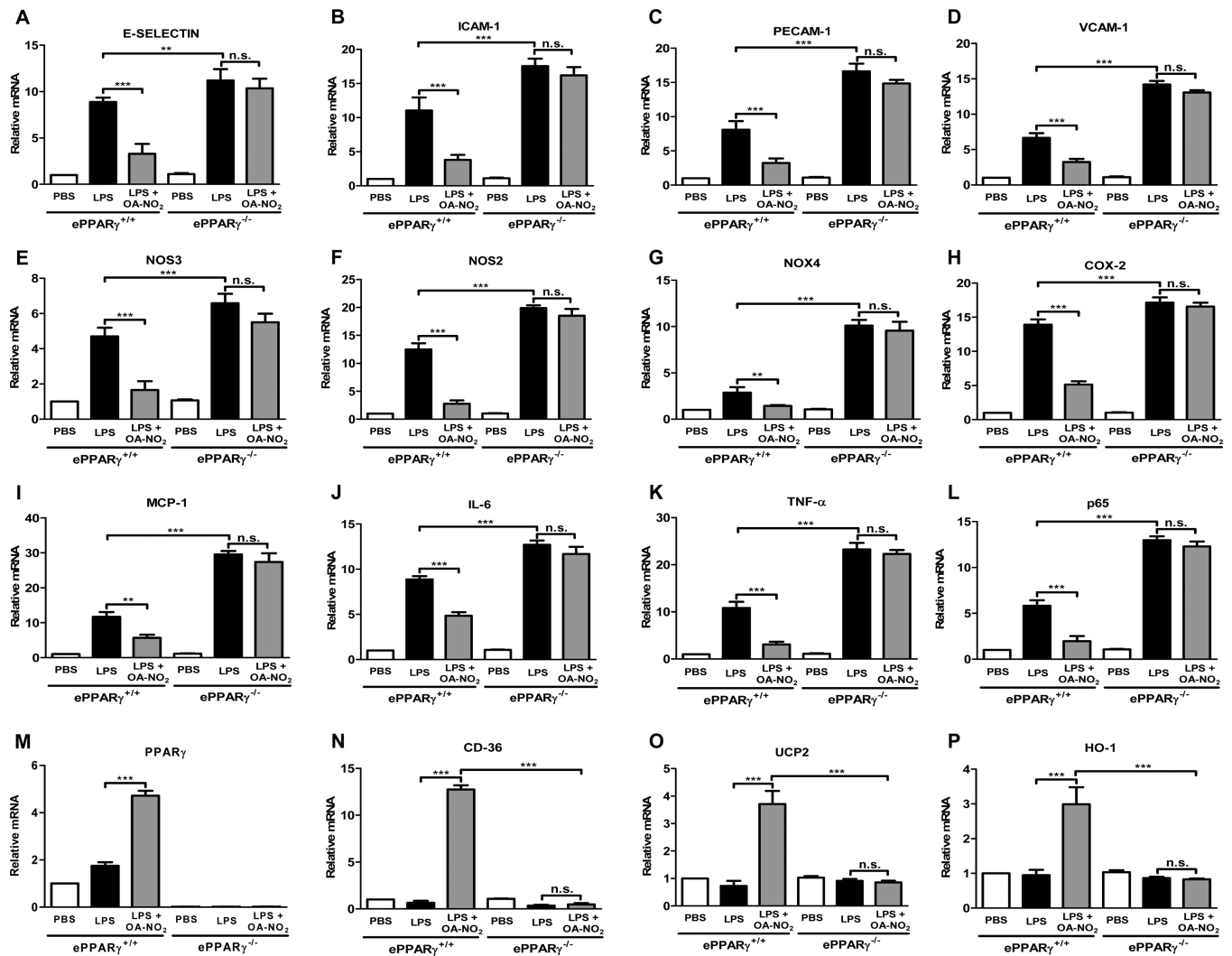


Figure 6. PPAR γ Deletion Eliminates OA-NO₂-Mediated Suppression of LPS-induced Expression of Inflammatory Markers in MLEC *in Vitro*
 MLEC were isolated from ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice ($n = 7-8$ mice per group) and plated onto 2% gelatin-coated dishes. Monolayer cultures were deprived of serum for 2 h, pre-treated with 100 nM OA-NO₂ or DMSO for 1 h, then exposed to 100 ng/ml LPS for 6 h. Cells were then harvested, RNA isolated, and expression of the indicated genes was determined using real-time PCR. Results are normalized to values for the housekeeping genes GAPDH and 9s rRNA. Data are representative of 3 independent experiments; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$.

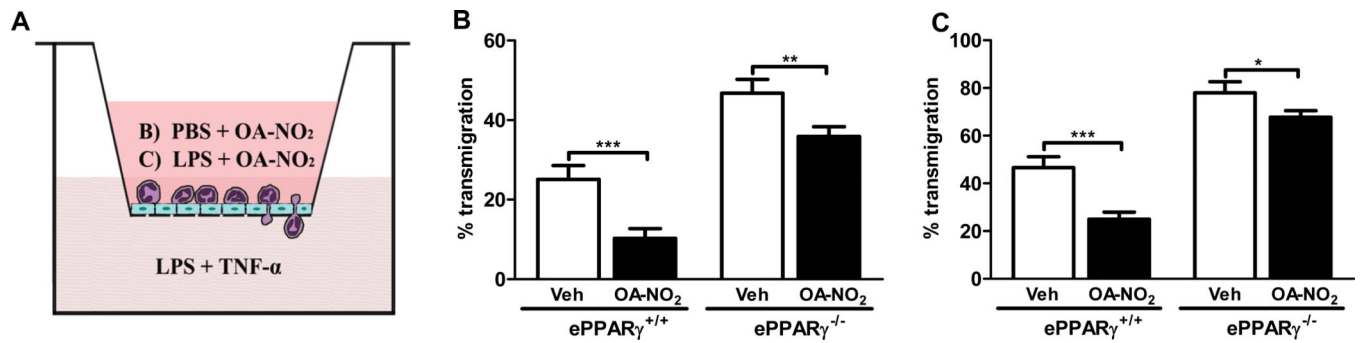


Figure 7. OA-NO₂ Inhibits LPS-induced Neutrophil Transmigration *in Vitro* via an ePPAR γ -dependent Mechanism

MLEC were isolated from ePPAR $\gamma^{+/+}$ and ePPAR $\gamma^{-/-}$ mice ($n = 7-8$ mice per group) and then grown to confluent monolayers in Transwell chambers. As shown schematically in (A), monolayers were treated with 100 nM OA-NO₂ or DMSO (Veh) along with (B) PBS or (C) LPS (100 ng/ml) and incubated for 6 h. After 6 h neutrophils (4×10^5) were added to each upper chamber, and a chemoattractant mixture of LPS (100 ng/ml) plus TNF- α (25 ng/ml) in RPMI 1640 was added to each lower chamber. The chambers were incubated at 37°C for 1.5 h, and neutrophils that had transmigrated into the lower chambers were collected and counted. Data are representative of 3 independent experiments; * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$.