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Molecular Mechanisms of the Antiproliferative Effect of Beraprost, a Prostacyclin Agonist, in Murine Vascular Smooth Muscle Cells

HENG LIN,¹ JA-LING LEE,² HSIN-HAN HOU,³ CHIH-PENG CHUNG,⁴ SUNG-PO HSU,⁵ AND SHU-HUI JUAN^{2,6,7,8}*

¹ Institute of Pharmacology and Toxicology, Tzu Chi University, Hualien, Taiwan

 2 Graduate Institute of Medical Sciences, Taipei Medical University, Taipei, Taiwan

³ Institute of Biomedical Sciences, Academia Sinica, Taipei, Taiwan

⁴ Graduate Institute of Cell and Molecular Biology, Taipei Medical University, Taipei, Taiwan

⁵ Institute of Physiology, College of Medicine, National Taiwan University, Taipei, Taiwan

6 Department of Physiology, Taipei Medical University, Taipei, Taiwan

 7 Graduate Institute of Neuroscience, Taipei Medical University, Taipei, Taiwan

⁸Topnotch Stroke Research Center, Taipei Medical University, Taipei, Taiwan

Prostacyclin (PGI₂) has been shown to inhibit proliferation in vascular smooth muscle cells. To clarify the underlying molecular mechanism, we investigated the vasoprotection of beraprost (a PGI₂ agonist) both in vivo and in vitro. Beraprost eliminated increases in proliferation of rat aortic smooth muscle cells (RASMCs) by 12-O-tetradecanoylphorbol 13-acetate, and enhanced the peroxisome proliferator-activated receptor-delta (PPARd) and inducible nitric oxide synthetase (iNOS) expressions, which were associated with the antiproliferative action of beraprost according to inhibition experiments by [³H]thymidine incorporation. Additionally, elimination of iNOS activity by PPAR₀ antagonists suggested that iNOS is the downstream target of PPAR₀. Furthermore, beraprost increased both consensus PPAR₀-responsive element (PPRE)-driven luciferase activity and the binding activity of the PPAR₀ to the putative PPRE in the iNOS promoter; nevertheless, it was abolished by PPAR δ antagonists. Deletion of PPRE (-1,349/-1,330) in the iNOS promoter region (1,359/þ2) strongly reduced promoter-driven activity, representing a novel mechanism of iNOS induction by beraprost. Consistent with this, PPAR₀ and the concomitant iNOS induction by beraprost were also evident in vivo. Beraprost-mediated protection in a murine model of balloon angioplasty was significantly attenuated by 13S-HODE, a PPAR₀ antagonist. Taken together, the results suggest that the causal relationship between PPAR₀ and iNOS contributes to the vasoprotective action of beraprost in RASMCs.

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Prostacyclin (PGI₂), produced by vascular endothelium and smooth muscles, has therapeutic effects on the relaxation of smooth muscle and reduction of vascular resistance by direct vasodilation (Hiatt, 2001). Administration of $PGI₂$ or its stable analogue, iloprost, limits coronary artery disease and infarct size in various animal models (Reynolds et al., 1967; Chiariello et al., 1988). $PGI₂$ is synthesized by a series of enzymes: cytosolic phospholipase A2 cleaves arachidonic acid (AA) from the sn-2 position of phospholipids, cyclooxygenase (COX) converts AA to $PGH₂$, and $PGI₂$ synthetase (PGIS) converts PGH₂ to PGI₂ (Matijevic-Aleksic et al., 1995). Beraprost sodium (BPS), a stable $PGI₂$ analogue, has been reported to have therapeutic effects on the treatment of primary pulmonary hypertension and obstructive peripheral arterial disease (Lievre et al., 2000; Galie et al., 2002). Similar to the biological properties of PGI₂, BPS can activate adenylate cyclase and increase intracellular cAMP levels through activation of the PGI₂ receptor. On account of its chemical characteristics, BPS is more stable and has higher affinity to the PGI₂ receptor than does natural PGI₂ (Kainoh et al., 1991).

PGI₂ functions through cell surface G protein-coupled receptors linked to different cytoplasmic signaling pathways Abbreviations: $PGI₂$, prostacyclin; BPS, beraprost; PPAR δ , peroxisome proliferator-activated receptor δ ; PPRE, PPAR δ responsive elements; RASMCs, rat aortic smooth muscle cells; TPA, 12-O-tetradecanoylphorbol 13-acetate.

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*Correspondence to: Shu-Hui Juan, Graduate Institute of Medical Sciences and Department of Physiology, Taipei Medical University, 250 Wu-Hsing Street, Taipei 110, Taiwan. E-mail: juansh@tmu.edu.tw

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which exert their effects by interacting with a nuclear hormone receptor, peroxisome proliferator-activated receptor (PPAR) (Vane and Botting, 1995; Prescott and White, 1996). PPARs are members of the nuclear hormone receptor superfamily and are ligand-activated transcription factors. PPARs modulate target gene expression in response to ligand activation after heterodimerization with the retinoid X receptor and binding to peroxisome proliferator-responsive elements (PPREs) of target genes. There are three different PPAR isoforms, designated PPAR α , PPAR β (also referred to as PPAR δ), and PPAR γ , each with distinct physiological functions (Evans et al., 2004; Kim et al., 2005). PPAR₀ has been shown to be involved in the regulation of lipid metabolism, and potential effects of PPAR δ activation are involved in the development of atherosclerotic lesions (Kersten et al., 1999). Moreover, Graham et al. (2005) showed that the potent and selective PPAR₀ agonist, GW0742X, reduces atherosclerosis in $LDLR-/-$ mice. These observations support an atheroprotective effect of PPAR δ agonists in vivo. By contrast, Dressel et al. (2003) showed that PPAR₀ plays a contrasting role in cardiovascular diseases associated with rat aortic smooth muscle cell (RASMC) proliferation such as atherosclerosis and restenosis. Although there is increased interest in characterizing the roles of PPAR δ in numerous disease models, the function of PPAR δ remains an enigma.

Numerous studies have focused on the effects of PPAR δ on cell viability. For instance, activation of endogenous PPAR δ by intracellular $PGI₂$ produced by expressing $PGIS$ in the human embryonic kidney epithelial 293 cell line promotes apoptosis (Hatae et al., 2001). By contrast, neutralization of PPAR δ by sulindac, a nonsteroidal anti-inflammatory drug, suppressed colorectal tumorigenesis in a nude mouse model (He et al., 1999). Therefore, the above-described discrepancies might be due to the effects of PPAR_o being tissue and species dependent.

Increasing evidence has shown a close link between PGI₂ and NO. For example, both PGI₂ and NO possess cardioprotective effects in animal models of myocardial infarction. Administration of $PGI₂$ or its stable analogue, iloprost, or an infusion of authentic NO or NO-donating drugs appears to reduce infarct size in most species studied (Sturzebecher et al., 1989). $PGI₂$ and NO can interact synergistically and influence each other's synthesis and release. The synergistic antiplatelet activity of $PGI₂$ and NO in rabbit and human platelets has been demonstrated by a number of workers (Radomski et al., 1987; Macdonald et al., 1988). Haider et al. (2003) demonstrated that COX-2 regulates tumor necrosis factor-mediated G1 shortening, but on the other hand, $PGI₂$ derived from COX-2 causes nitric oxide-mediated inhibition of vascular smooth muscle cell proliferation. Interestingly, this synergy was not seen in vascular smooth muscles (Lidbury et al., 1989).

In a murine model of carotid artery balloon injury, the activated vascular endothelium produces cytokines and growth factors, which promote the growth and migration of RASMCs, key events in the formation of atherosclerotic lesions in humans. Thus, in addition to the primary culture system, we also examined the role of PPAR δ in the proliferation of smooth muscle cells following surgery. Given that $PGI₂$ and NO influence each other's synthesis and $PGI₂$ is a ligand of PPAR δ , we attempted to delineate the relationship of PPAR δ and inducible nitric oxide synthetase (iNOS) induced by $PGI₂$ and their involvements in the vasoprotective effects of $PGI₂$ in the vasculature.

Materials and Methods

Isolation and primary culture of RASMCs

RASMCs were isolated from the thoracic aorta of male Sprague–Dawley rats (275–325 g) using the explant technique (Fisher-Dzoga et al., 1973). Briefly, after removal of the

endothelium and adventitia, the aortic explants were cultured in DMEM and supplemented with 10% fetal bovine serum, penicillin (100 U/ml), streptomycin (100 μ g/ml), and 25 mM HEPES (pH 7.4). After 2 weeks, cells that had migrated out of the explants were removed by trypsinization and successively subcultured. The purity and identity of cells were examined by immunostaining using an antibody specific against smooth muscle cell α -actin. Cells from passages 5 to 12 were used for the experiments.

[³H] Thymidine incorporation

As previously described (Jain et al., 1996), cells at a density of 1×10^4 cells/cm³ were applied to 24-well plates in DMEM plus 10% fetal bovine serum from Invitrogen (Carlsbad, CA). After the cells had grown to 70–80% confluence, they were rendered quiescent by incubation for 24 h in DMEM containing 2% fetal bovine serum. Chemicals including beraprost (a $PGI₂$ agonist), 13Shydroxyoctadecadienoic acid (13S-HODE) (a PPAR_o antagonist), and N^G -nitro-L-arginine methyl ester HCL (L-NAME) and NG-nitro-L-arginine (L-NNA) (NO inhibitors) were purchased from Cayman (Ann Arbor, MI), and 12-O-tetradecanoylphorbol 13-acetate (TPA) (a mitogen), sulindac (a PPAR_o antagonist), and sodium nitroprusside dehydrate (SNP) (a NO donor) were purchased form Sigma Chemical (St. Louis, MO). Chemicals as indicated or dimethyl sulfoxide (DMSO) in 10% fetal bovine serum were added to the cells, and the mixture was allowed to incubate for 24 h. During the last 4 h of the incubation with or without beraprost, $[^{3}H]$ thymidine was added at 1 μ Ci ml $^{-1}$ (1 μ Ci = 37 kBq). The incorporated [³H]thymidine was extracted in 0.2 N NaOH and measured in a liquid scintillation counter.

MTT assay

The viability of cells was determined based on the activity of mitochondrial dehydrogenase to reduce 3-(4,5-dimethylthiazol-2 yl)-2,5-diphenyl tetrazolium bromide (MTT) (Sigma) to formazan. Briefly, cells were plated in 24-well plates at a density of 1 \times 10⁴ cells/well and treated with DMSO or varying concentrations of beraprost for 24 or 48 h. MTT was then added to each well, and incubation continued for 4 h in culture. The formazan formed was dissolved by overnight incubation with 10 mM HCl containing 10% SDS, and then measured at a wavelength of 570 nm.

Electrophoretic mobility shift assay

The electrophoretic mobility shift assay (EMSA) was performed as previously described (Juan et al., 2005). To prepare nuclear protein extracts, cells in 10 -cm² dishes after treatment with indicated chemicals for 4 h were washed twice with ice-cold PBS and scraped off into 1 ml of PBS. After centrifugation of the cell suspension at 500g for 3 min, the supernatant was removed, and cell pellets were subjected to NE-PERTM nuclear extraction reagents (Pierce, Rockford, IL) with the addition of 0.5 mg/ml benzamidine, $2 \mu g/ml$ aprotinin, $2 \mu g/ml$ leupeptin, and $2 \mu M$ PMSF. The subsequence procedures of nuclear protein extraction followed the manufacturer's instructions. The fraction containing the nuclear protein was used for the assay or was stored at -80° C until use. The sequence of the oligonucleotide used was GTGGAGGTCAGGGGACAATT for PPRE binding. The oligonucleotide was end-labeled with $[\gamma^{32}P]$. Extracted nuclear proteins (10 μ g) were incubated with 0.1 ng of 32 P-labeled DNA for 15 min at room temperature in 25 μ l of binding buffer containing $\perp \mu$ g of poly(deoxyinosine-deoxycytidine) (dl-dC). For competition with unlabeled oligonucleotides, a 100-fold molar excess of unlabeled oligonucleotide relative to the ³²P-labeled probe was added to the binding assay. Mixtures were electrophoresed on 5% non-denaturing polyacrylamide gels. Gels were dried and imaged by means of autoradiography.

Transient transfection experiment and luciferase activity assay

Two DNA fragments with or without PPRE of rat iNOS gene promoter sequences from $-1,359$ to $+2$ and $-1,329$ to $+2$ bp, respectively, were obtained by a polymerase chain reaction (PCR) using rat genomic DNA as the template. The sense and antisense primers were as follows: sense (with PPRE) 5'-TGGTACACATGTGGAGGTCAGGGGACAATT-3', sense (without PPRE) 5'-TATGGGAGTTTGTTTTCTCCTTCCA-CCGTG-3', and antisense 5'-CAACTCCCTGTAAAGCTGT-GGCCCTGACAG-3'. The PCR product was flanked with dA and subcloned into the yTA clone vector, which was digested with Kpn I and Sal I restriction enzymes and subcloned into the KpnI-XhoI site of the PGL3-basic vector. The identities of the sequences were confirmed using an ABI PRISM 377 DNA Analysis System (Perkin-Elmer, North Point, Hong Kong).

For the reporter activity assay, cells were seeded in six-well plates at a density of 1×10^5 cells/well. In brief, RASMCs were transiently transfected with $1.05 \mu g$ of plasmid DNA containing 0.05 μ g of the Renilla luciferase construct, pRL-TK (Promega, Madison, WI), to control transfection efficiency and μ μ g of the appropriate iNOS promoter firefly luciferase (FL) construct. The next day, cells were transfected with pGL3/iNOS promoter variants $(-1,359/+2$ or $-1,329/+2)$ and pRL-TK (the internal control plasmid) using LipofectAMINE 2000TM (Invitrogen). After transfection for 4 h, the medium was replaced with complete medium, and incubation continued for another 20 h. Transfected cells were then treated with drugs for 12 h, and cell lysates were collected. Luciferase activities were recorded in a TD-20/20 luminometer (Turner Designs, Sunnyvale, CA) using the dual luciferase assay kit (Promega) according to the manufacturer's instructions. Luciferase activities of the reported plasmids were normalized to luciferase activities of the internal control plasmid.

RNA Isolation and analysis of gene expression by real-time-PCR

Total RNA was prepared from cultures by directly lysing cells in Trizol buffer (Life Technologies, Gaithersburg, MD), and mRNAs were reversed-transcribed into cDNA using oligo-dT and reverse transcriptase (Invitrogen). After first-strand cDNA synthesis, it was used as a template and amplified by pairs of primers derived from PPAR_o and iNOS genes for RT-PCR and quantitative real-time-polymerase chain reaction (RT-PCR) analysis. Sequences of the primer pairs for amplification of each gene were 5'-AACATCCCCAACTTCAGCAG-3' and 5'-TACTGCGCAAGAACTCATGG-3' (for the PPAR_o gene); 5'-ACCTGAAAGAGGAAAAGGAC-3' and 5-CTCTGGTC-AAACTCTTGGAG-3' (for the iNOS gene); and 5'-CCACCATGGAGAAGGCTGGGGCTCA-3' and 5'-ATCACGCCACAGTTTCCCGGAGGGG-3' (for the GAPDHgene). PCR amplifications were conducted using QPCRM aster Mixture and SYBER Green-based detection T^M systems (ABI; Applied Biosystems, Foster City, CA) according to the manufacturer's instructions, with 100 nM primers and 100 ng of $cDNA$ template in a 20 - μ l reaction volume. Thermocycling was initiated by 2 min of decontamination at 50° C and 10 min of hot start at 95 $^{\circ}$ C, followed by 40 cycles of 95 $^{\circ}$ C for 15 sec, 55 $^{\circ}$ C for 30 sec, and 60° C for 60 sec, with a single fluorescence reading taken at the end of each cycle. Each run was completed with a melting curve analysis to confirm the specificity of amplification and lack of primer dimers. Ct values were determined by the ABI System Software using a fluorescence threshold manually set to 0.0160. Individual gene expression Ct values at various time points were normalized by subtracting the respective Ct value of a housekeeping gene, GAPDH, to obtain a Δ Ct calibrated value. Experimental samples are presented as multiples of induction with respect to each control group at various time points.

Western blots

Western blotting was carried out with a rabbit polyclonal anti-PPAR_o antibody, rabbit polyclonal anti-iNOS antibody (Santa Cruz; Santa Cruz, CA), and mouse polyclonal anti-GAPDH antibody (Biogenesis; Kingston, NH). Cell lysate (50 μ g) was electrophoresed on an 8% SDS–polyacrylamide gel and then transblotted onto a Hybond-P membrane (GE Healthcare, Mongkok, Hong Kong). The blot was blocked in PBS containing 0.1% Tween-20 and 5% skim milk at room temperature for 1 h, followed by incubation with the first antibody (diluted 1: 1,000) for another hour in PBS containing 0.1% Tween-20 and 5% skim milk. After 2 washes, the blot was incubated with peroxidase-conjugated goat anti-rabbit immunoglobulin G (IgG)/anti-mouse IgG (1: 2,000) (Sigma) (diluted 1: 2,000) for an additional 1 h. Antigen was detected with an enhanced chemiluminescence system (GE Healthcare).

Nitrite assay

RASMCs were plated at a density of 1×10^5 cells/ml in 24-well plates for 16 h, followed by treatment with beraprost (1 μ M) or the addition of various concentrations of PPAR_o antagonists for a further 24 h. The amount of NO production in the medium was detected with the Griess reaction. One hundred microliters of each supernatant was mixed with the same volume of Griess reagent (1% sulfanilamide in 5% phosphoric acid and 0.1% naphthylethylenediamine dihydrochloride in water). The absorbance of the mixture at 530 nm was determined with an enzyme-linked immunosorbent assay (ELISA) plate reader (Dynatech MR-7000; Dynatech Laboratories).

Carotid injury

Animal care and treatment were conducted in conformity with the protocols of the Animal Center, Taipei Medical University. Male C57B6 mice (6–10 weeks old, 25.8 ± 1.8 g) were used in this study. Animals were anesthetized intramuscularly with a combination of ketamine (8 mg/100 g body weight), xylazine (2 mg/100 g), and atropine (0.16 mg/100 g). The right carotid artery was exposed, and endothelial denudation of the common carotid was induced with a 29-gauge needle (0.35 mm in diameter), the tip of which was roughened and coated with an α -cyanoacrylate bead (0.5 mm in diameter). This modified needle was then introduced through the internal carotid artery and moved into the common carotid artery. The common carotid artery was abraded four times. The arteriotomy site was ligated, and the skin was closed using sutures. The external carotid was then tied off, and blood flow was restored through the common carotid artery. The mice recovered from anesthesia and were allowed free access to food and water. Nalbuphine (1–2 mg/kg intramuscularly) was given as the postoperative analgesia. After carotid balloon injury, mice were divided into four groups: saline as the control, beraprost alone, PPAR δ antagonist (13S-HODE), and beraprost combined with 13S-HODE (with six mice in each group). Balloon-injured mice were subcutaneously given beraprost (1 mg/kg) on day 2 or 13S-HODE (100 μ g/kg) on day 3 following surgery, and separate chemicals were given every other day for 2 weeks. Furthermore, mice without balloon injury were given saline, beraprost, or 13S-HODE as the controls.

Histological analysis and quantification of atherosclerotic lesions

Animals were killed 2 weeks after vascular injury. The operated carotid arteries were excised, fixed in 4% paraformaldehyde, embedded in paraffin, and serially sectioned at $7 \mu m$ for histological staining or other experiments. Totally, 100 sections from each injured artery were collected. In total, 11 sections sampled from every 10 consecutive sections were stained with Verhoeff van Gieson elastic stain and used for lesion estimation (Lee et al., 1999).

Immunohistochemistry

Immunostaining was carried out using the same antibodies as in the Western blot analysis. Tissue sections were pretreated with 3% H₂O₂ for 15 min at room temperature. After incubation in PBS containing 2% bovine serum albumin (BSA) at 37° C for 30 min, sections were incubated with the primary antibody at 37° C for 30 min, followed by three washes in PBS. Sections were then incubated with a horseradish peroxidase-conjugated secondary antibody for another 30 min. After three washes, color was developed with 0.1% 3,3'-diaminobenzidine/0.01% H_2O_2 . Pictures were taken at a 200 \times magnification using a CCD camera (DP70, Olympus, Melville, NY) attached to a microscope system (BX51, Olympus).

Statistical analysis

Values are expressed as the mean \pm SD The significance of the difference from the control groups was analyzed by Student's t-test or one-way analysis of variance (ANOVA) and Bonferroni's method as a post hoc test. A value of $P < 0.05$ was considered statistically significant.

Results

Effects of beraprost on cell proliferation and cell viability

To examine the specificity of a direct effect of PGI_2 on cellular proliferation, we performed [3H]thymidine incorporation in RASMCs in the presence of various concentrations of beraprost, a PGI₂ agonist. Beraprost not only concentration-dependently inhibited [³H]thymidine incorporation into RASMCs by about 36%, but also reversed TPA-induced DNA synthesis by about 50%, as shown in Figure 1A. In addition, to examine the effect of cell

Fig. 1. Effects of beraprost (BPS) on DNA synthesis and viability of rat aortic smooth muscle cells (RASMCs). A: BPS concentrationdependently inhibited DNA synthesis and reversed the 12-Otetradecanoylphorbol 13-acetate (TPA)-induced cell proliferation by [3 H]thymidine incorporation in RASMCs. Cells were pretreated with TPA for 1 h prior to additional BPS treatment. Data are shown as the mean \pm SD of five independent experiments. P < 0.05 and $*P$ < 0.01 versus the control group; # P < 0.01 versus TPA treatment alone. B: BPS decreased cell viability at 2 days after treatment. The cell viability of cells in 24-well plates treated with BPS was determined by the MTT assay at the indicated time points. Data shown are the $mean \pm SD of five independent experiments. Significantly different$ $(P < 0.05$ and $* P < 0.01$ versus the DMSO-treated group at day 2).

viability by beraprost, an MTT assay was carried out. Beraprost concentration-dependently inhibited cell viability on day 2 by approximately 30%, but not on day 1, as illustrated in Figure 1B.

Time-dependent inductions of $PPAR\delta$ and iNOS, and an increase in NO production by beraprost

To unravel the close linkages among $PGI₂$, NO, and $PPAR₀$, we analyzed the time-dependent inductions of PPAR₀ and iNOS by beraprost in RASMCs at both the mRNA and protein levels. RNA and total cell lysate were harvested from RASMCs treated with $1 \mu M$ beraprost for 2, 4, 6, 12, or 24 h, as analyzed by real-time-PCR, RT-PCR, and Western blotting. As illustrated in Figure 2A,B, beraprost significantly increased induction of PPAR δ at as early as 2 h, whereas induction of iNOS was observed at 4 h after treatment. Time-course inductions of PPAR δ and iNOS by BPS from 0 to 12 h increased by approximately 1.5-fold to 3.6-fold and 1.2-fold to 2.9-fold, respectively. Additionally, the nitrite assay in Figure 2C demonstrated that beraprost significantly stimulated NO production no earlier than 4 h after treatment; this time point is consistent with the findings in Figure 2A,B, regarding iNOS induction by beraprost at both the RNA and protein levels. NO production was increased approximately twofold at 24 h after 1μ M BPS treatment in RASMCs.

Decrease in iNOS expression and NO production by PPAR_o antagonists

Results in Figure 2 show that the time frame of PPAR_o induction was approximately 2 h earlier than that of iNOS by bepraprost. In order to examine the causal relationship between $PPAR\delta$ and iNOS, we used the PPAR_o antagonists, 13S-HODE and sulindac, to investigate their effects on iNOS expression. A nitrite assay analysis (Fig. 3A) demonstrated that PPAR δ antagonists statistically inhibited beraprost-induced NO production in an amount similar to the control group after 24 h of treatment, suggesting that iNOS is the downstream target of PPAR_o. Moreover, cells pretreated with PPAR_o antagonists (sulindac or 13S-HODE) for 1 h prior to the addition of beraprost, showed decreased protein levels of iNOS as a result of a decrease in PPAR_o protein expression, as shown in Figure 3B by Western blotting.

Transcriptional regulation of iNOS by PPAR₀ in RASMCs treated with beraprost

As it is known that PGI_2 exerts its function through PPAR δ and has a close link with nitric-oxide synthetase (NOS) (Sturzebecher et al., 1989; Vane and Botting, 1995; Narumiya et al., 1999), we also demonstrated in Figure 2 that beraprost induced both PPAR δ and iNOS expression; the time frame of PPAR δ 's action was earlier than that of iNOS. Therefore, we constructed two repeats of the consensus PPAR δ -responsive element (PPRE) in the pGL2-promoter vector. Beraprost concentration-dependently enhanced the luciferase activity by approximately 1.6-fold to 5.6-fold in cells transfected with the PPRE construct as shown in Figure 4A. The putative PPRE in the iNOS promoter region as analyzed by the MOTIF Search software was synthesized and used as a probe for an EMSA. The results presented in Figure 4B show that beraprost increased the DNA binding activity of PPAR δ to the putative PPRE, whereas the activity was alleviated by $PPAR\delta$ antagonists.

Fig. 3. Effects of the PPAR δ antagonists, sulindac (10 and 100 μ M) and I3S-HODE (0.1 and 1 µM), on NO production in RASMC exposed
to beraprost (A) and protein levels of PPAR δ and iNOS by Western blots (B). Six samples were analyzed in each group, and values are presented as the mean ± SD. Significantly different (*P < 0.05 vs. the
DMSO-treated group, *P < 0.05 vs. BPS alone). GAPDH was used as an internal control, and results shown were reproduced in three separate experiments.

Fig. 4. Binding activity of PPAR δ to the PPRE in relation to beraprost or its antagonists. A: A concentration-dependent increase in consensus PPRE-driven luciferase activity by beraprost. Luciferase activities of the reported plasmid were normalized to those of the internal control and are presented as the mean \pm SD. Significantly different (*P<0.05 vs. the DMSO-treated group, #P<0.05 and
##P<0.01 vs. the piNOS-1,329 group). B: A decrease in the binding
##P<0.01 vs. the piNOS-1,329 group). B: A decrease in the binding
activity of PPAR δ to the PP cultured and pretreated with PPAR₀ antagonists for 1 h prior to the addition of $I \mu M$ beraprost. The putative PPRE binding activity derived from the iNOS promoter region of nuclear proteins was assayed by EMSA. 100× cold denotes a 100-fold molar excess of
unlabeled oligonucleotides relative to the ³²P-labeled probe; this was added to the binding assay for competition with the unlabeled oligonucleotide. The mobility of specific PPRE complexes is indicated. Representative results of three separate experiments are shown. C: Importance of the putative PPRE_o in iNOS promoter-driven luciferase activity by beraprost. Luciferase constructs of iNOS promoter variants were obtained as described in Section ''Materials and Methods.'' The experimental reporter luciferase activity was calculated by subtracting the intrinsic activity as measured by samples corresponding to the pGL3-basic vector and then normalizing them to the transfection efficiencies as measured by the activity derived from pRL-TK. Data are presented as the mean \pm SD. Significantly different ($\frac{N}{P}$ < 0.05 vs. the DMSO-treated group or $\frac{N}{P}$ \mathbb{P} <0.05 and \mathbb{P} <0.01 vs. the piNOS-1,359 group treated with the indicated concentrations of beraprost).

Another approach, consisting of luciferase constructs of the rat iNOS promoter regions $(-1,359/+2)$ with or without the putative PPRE $(-1,349/-1,330)$ by deletion mutation, was used to confirm that the binding of PPAR_o to the PPRE is associated with transactivation of iNOS by beraprost. The wild-type construct was transiently transfected into RASMCs for 24 h followed by the addition of the indicated concentrations of beraprost. After incubation for another 12 h, cells were

harvested and analyzed for luciferase activity. As shown in Figure 4C, luciferase activity increased by about twofold to 2.6-fold in response to increased concentrations of beraprost of from 0.5 to 5 μ M. In contrast, the concentration-dependent increase in promoter activity by beraprost was abolished when the putative PPRE was removed, as shown in Figure 4C.

Reversal of beraprost-induced inhibition of RASMCs DNA synthesis by PPAR δ and NO inhibitors, and a mimicking of the effect of beraprost by sodium nitroprusside dehydrate (SNP), a NO donor

To examine whether PPAR_o and iNOS are responsible for the antiproliferative effects of beraprost in RASMCs, their individual antagonists or inhibitors were utilized to investigate this postulation. We show in Figure 5A, B, that the PPAR δ antagonist and NO inhibitors significantly reversed the decrease in DNA synthesis by beraprost in the [3H]thymidine incorporation assay, suggesting that beraprost's actions are exerted through PPAR δ and NO. Conversely, SNP, a NO donor, was used to mimic this effect of beraprost in RASMCs. As shown in Figure 5C, SNP at concentrations of $0.1-10 \mu M$, similar to those of beraprost, caused a statistically significant decrease in DNA synthesis in the [³H]thymidine incorporation experiment.

Fig. 5. Abolishment of beraprost-induced decrease in cell proliferation by the PPAR₀ antagonists, sulindac and 13S-HODE (A), and by the NO inhibitors, L-NNA and L-NAME (B), and sodium nitroprusside dehydrate (SNP), a NO donor, mimicking the effect of
beraprost in RASMCs (C). The effects of PPAR δ antagonists or iNOS inhibitors on berapost-induced DNA synthesis inhibition, or SNP mimicking the effects of BPS were assessed by [3H]thymidine incorporation, as described in Section ''Materials and Methods.'' RASMCs were pretreated with PGI2 antagonists or NO inhibitors for I h prior to the addition of $I \mu M$ beraprost, or cells were treated with increasing concentrations of SNP and then incubation continued for another 24 h. Six samples were analyzed in each group, and values
represent the mean ± SD. Significantly different (*P< 0.05 and
**P< 0.01 vs. the DMSO-treated group, *P< 0.05 vs. BPS alone). $*$ P<0.01 vs. the DMSO-treated group, $*$ P<0.05 vs. BPS alone).

Therapeutic effects and mechanisms of beraprost in mice subjected to carotid balloon injury

An animal model of carotid balloon injury was established to examine the antiproliferative effects of beraprost in vivo. Carotid balloon injury is the commonest procedure to simulate the pathology of atherosclerosis; the key feature is smooth muscle cell overproliferation. We demonstrate in Figure 6A,B that balloon-injured mice given beraprost as therapeutic treatment showed decreased progression of restenosis by about 70% compared to injured mice alone. Additionally, 13S-HODE, a PPAR δ antagonist, was used to examine whether PPAR δ is the key player exerting the antiproliferation effects of beraprost in RASMCs. Indeed, the addition of the PPAR δ antagonist blocked the antiproliferative effect of beraprost, resulting in increased progression of restenosis compared with

Fig. 6. Effect of beraprost (BPS) or in combination with 13S-HODE on neointimal formation. The carotid arteries of mice were subjected to angioplasty and subsequent treatment with saline, BPS alone, or BPS in combination with 13S-HODE. At 2 weeks after injury, neointimal formation was examined. A: Verhoeff van Gieson staining of injured arterial segments. B: Intima/media (I/M) ratios of injured arteries in various treated groups. The thicknesses of the intima and media of injured arteries were quantified as described in Section ''Materials and Methods.'' The number of animals in each group The saling was six. ^{We} p < 0.01 versus the saline group and ##P < 0.01 versus the BPS-treated group. C: The extents of BPS-induced PPAR δ and iNOS expressions were examined by immunostaining with the indicated antibodies. [Color figure can be viewed in the online issue, which is available at www.interscience.wiley.com.]

the experimental group treated with beraprost alone, which suggests the important role of PPAR_ô in the antiproliferative action of beraprost in balloon injury. Furthermore, histoimmunochemical staining showed that beraprost treatment caused the induction of PPAR δ and iNOS (Fig. 6C) which agrees with the in vitro findings.

Discussion

In this study, we provide both in vitro and in vivo data to support the roles of PPAR δ and iNOS in the antiproliferative action of beraprost, a PGI₂ agonist, in RASMCs. Antagonists or inhibitors of PPAR δ and iNOS were utilized to reverse a decrease in DNA synthesis of RASMCs by beraprost. We also clearly delineate the causal relationship between PPAR₀ and iNOS, as PPAR₀ antagonists decreased the protein levels of iNOS and NO production. Furthermore, a deletion mutation identified a functional PPRE located in the promoter region of the murine iNOS gene by the promoter luciferase assay, suggesting that $PPAR\delta$ is an upstream effecter of iNOS induction by beraprost.

In the present study, we examined the potential effects of PPAR_b induced by beraprost on the inhibition of vascular smooth muscle proliferation and its implications in the prevention of restenosis after carotid balloon injury in vivo. Apparently, the present finding is contradictory to a recent report showing that $PPAR\delta$ induced by PDGF or by carotid balloon injury promoted cell proliferation in RASMCs (Zhang et al., 2002). The discrepancy of PPAR δ on RASMC proliferation might be attributed to whether it is properly ligand-bonded (i.e., beraprost, a PGI₂ agonist, a natural ligand of $PPAR\delta$) or what its inducers are, although this speculation remains to be further investigated. Nevertheless, a very recent study by Graham et al. (2005) with outcomes similar to our findings demonstrated the atheroprotective effect of the PPAR δ agonist, GW0742, in LDLR^{-/-} mice by showing a decrease in the chemotactic and proinflammatory molecules (Graham et al., 2005).

It has been documented that an intracellular or extracellular source of PGI₂ produces different cell fates. For instance, Hatae et al. (2001) demonstrated that intracellular $PGI₂$ formed by expressing PGIS in human embryonic kidney 293 cells promotes apoptosis by activating PPAR₀. Conversely, iloprost (an extracellular PGI_2) or dibutyryl cAMP (a cAMP agonist) reduced apoptosis. Furthermore, Ii et al. (2001) demonstrated that beraprost, an extracellular PGI₂ source, reduces RASMCs proliferation through cAMP signaling by preventing downregulation of p27. In our study, we provide another line of evidence showing that even though beraprost is an extracellular $PGI₂$ source, it can activate PPAR δ with a concomitant induction of iNOS to exert the antiproliferative action of $PGI₂$ in RASMCs. Likewise, the antiproliferative effect of beraprost in RASMCs was limited to cell cycle arrest, as no apoptotic cell death was observed (data not shown) in our study, similar to the findings of li et al. (2001).

Additionally, a deletion mutation of the putative PPRE revealed that a functional PPRE in the promoter region of iNOS is associated with the induction of iNOS by the luciferase promoter assay (Fig. 4C). This causal relationship between PPAR and NO is in accord with a recent study by Crosby et al. (2005) showing that a novel PPAR γ -responsive element in the promoter region of iNOS has been identified in rat mensangial cells, which contributes positive basal expression and negative expression of iNOS in response to inflammatory stimuli. Likewise, Niwano et al. (2003) demonstrated that beraprost transcriptionally stimulates eNOS expression in vascular endothelial cells, although it is transcriptionally activated by a c AMP-responsive element. The synergistic activity of $PGI₂$ and NO has been manifested in various tissues (Radomski et al.,

1987; Macdonald et al., 1988), but not in vascular smooth muscle cells (Lidbury et al., 1989); nevertheless, we demonstrate that their causal relationship is involved in the antiproliferation of RASMCs. In view of the still not clearly determined role of PPAR_o in the physical and pathological remodeling of the vasculature, we demonstrate a correlation of PPAR δ with iNOS induction by PGI₂ to exert its antiproliferative effect in RASMCs both in vivo and in vitro.

In summary, the data presented herein provide evidence to support the essential role of PPAR δ in the antiproliferative function of beraprost, although the cellular signal pathway by which beraprost induces $PPAR\delta$ expression remains to be determined. PPAR_o appears to transcriptionally mediate the upregulation of iNOS in RASMCs. Identifying the proteins $including PPAR_o and iNOS responsible for the antiproliferation$ of RASMCs by beraprost will provide an important molecular basis for the design of new therapeutic strategies to treat atherosclerosis, restenosis or stroke.

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