

Relationship between peroxisome proliferator-activated receptor- γ and renal ischemia-reperfusion injury

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Abstract. The pathogenesis of ischemia-reperfusion (I/R) injury is known to involve cytokines and, in particular, surface adhesion molecules, the expression of which initiates inflammatory cell attachment. It has been suggested that peroxisome proliferator-activated receptor (PPAR)- γ is an important immunomodulatory factor as well as a regulator of fatty acid. In this study, we investigated the expression of PPAR- γ in a renal I/R injury rat model. The right kidney was harvested and the left renal artery and vein were clamped by means of a laparotomy. The kidney was reperfused following 90 min of ischemia. Rats were sacrificed at 0, 1.5, 3, 5, 12 and 24 h after reperfusion. PPAR- γ expression was analyzed by immunohistochemical staining using monoclonal antibody. PPAR- γ staining was weak in the endothelial cells, interstitial cells and collecting ducts in the normal kidney. From 1.5 to 5 h after reperfusion, PPAR- γ staining was strong. Twelve hours after reperfusion, necrosis had extended throughout the kidney, and nearly all the tubular epithelial cells were destroyed. However, 12 h after reperfusion, PPAR- γ staining was weak in the endothelial cells and its expression was moderate in the interstitial cells and collecting ducts. PPAR- γ was induced in the endothelial cells, including the mesangial cells, interstitial cells and collecting ducts in a rat model of renal I/R injury.

Introduction

Renal transplantation is an acceptable therapeutic approach for patients with end-stage renal disease. Renal ischemia-

reperfusion (I/R) injury is a clinically significant problem and an invariable consequence of renal transplantation that results from aortic cross-clamping and resuscitation after systemic hypotension. Previous studies have focused on the function of neutrophils, the action mechanisms of inflammatory cytokines, tissue factors, intercellular adhesion molecule-1, oxygen-free radicals, vascular plugging, edema and other complications (1).

Peroxisome proliferator-activated receptors (PPARs) are members of the nuclear hormone receptor superfamily of ligand-activated transcriptional factors, which includes receptors for steroids, thyroid hormones, vitamin D₃ and retinoic acid. PPAR binds to a peroxisome proliferator responsive element to form a heterodimer with the retinoic receptor in the regulation of PPAR target genes (2).

There are three known PPAR isoforms: PPAR- α , - β and - γ . PPAR- α primarily regulates the fatty acid metabolism and has an anti-inflammatory effect; its ligands are used in fibrate drugs. PPAR- β is ubiquitously expressed throughout the body, but its physiological functions are unknown. PPAR- γ has anti-inflammatory effects and controls the functioning of the immune system, having inhibitory action on the production of nitric oxide, cytokine and monocyte chemoattractant protein-1 and on B-cell growth, vascularization and cell growth, and differentiation-inducing action on dendritic cells and helper T-cells (3).

PPAR- γ has also been implicated in inflammation and tumorigenesis (4), and significant evidence from many experimental models suggests that it plays a role in carcinogenesis. We demonstrated that PPAR- γ is upregulated in malignant tissue, and that its ligands induce terminal differentiation in human urological cancers, inhibiting their growth (4-7). However, few reports have addressed the relationship between PPAR- γ and renal I/R injury. For this reason, we investigated their relationship in a rat model.

Materials and methods

Ischemia-reperfusion model. Male Lewis rats (180-230 g) were used. During a laparotomy using pentobarbital sodium

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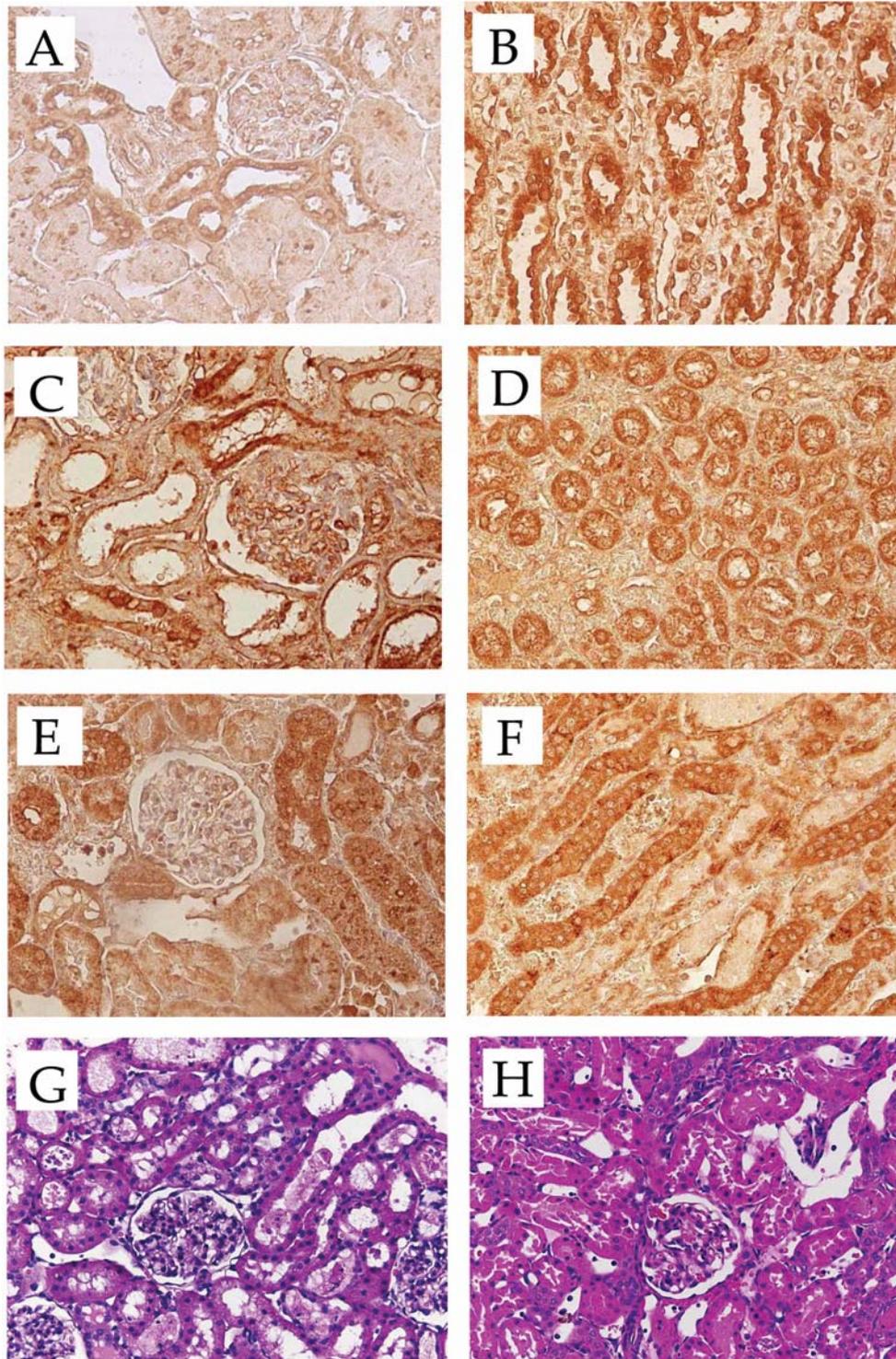


Figure 1. PPAR- γ immunohistochemistry and H&E staining. In the normal kidney, PPAR- γ staining was weak in the endothelial cells, including the mesangial cells (A), weak in the interstitial cells and strong in the collecting ducts of the medulla (B). From 1.5 to 5 h after reperfusion, PPAR- γ staining was strong in the endothelial cells (C), moderate in the interstitial cells and strong in the collecting ducts (D). It was, however, weak in the tubules (C). Five hours after reperfusion, the internal spaces of the tubular epithelial cells were expanded and slight destruction of the tubular epithelial cells was apparent (G). The necrotic area extended and encompassed nearly all of the ischemic kidney within 12 h after reperfusion (H). At 12 h after reperfusion, PPAR- γ staining was weak in the endothelial cells (E), moderate in the interstitial cells and strong in the collecting ducts (F).

anesthesia, the right kidney was harvested and the left renal artery and vein were clamped with a hemostasis clip for 90 min. The clip was subsequently removed to permit reperfusion, and the abdomen was closed during I/R. The rats were sacrificed at 0, 1.5, 3, 5, 12 and 24 h after reperfusion, and their kidneys harvested. Samples of ischemic and non-

ischemic kidney tissue were fixed in 10% buffered formalin for 24 h for immunohistochemistry and H&E staining.

Immunohistochemical staining. Immunohistochemical staining was performed with the Vectastain avidin-biotin peroxidase complex kit (Vector Laboratories, Burlingame, CA, USA).

Time after reperfusion injury	PPAR- γ of the renal cortex	PPAR- γ of the renal medulla	ATN
0 h	1.5 \pm 0.5	3.0 \pm 0.7	0.1 \pm 0.2
1.5 h	3.4 \pm 0.6 ^a	3.4 \pm 0.6	0.2 \pm 0.3
3.0 h	3.2 \pm 0.7 ^a	3.5 \pm 0.6	0.6 \pm 0.4
5.0 h	2.8 \pm 0.7 ^a	3.3 \pm 0.5	1.2 \pm 0.4
12.0 h	2.2 \pm 0.8	3.4 \pm 0.5	2.5 \pm 0.4
24.0 h	1.7 \pm 0.8	3.3 \pm 0.6	2.9 \pm 0.3
Before ischemia	1.2 \pm 0.7	2.9 \pm 0.6	N.P.

The acute tubular necrosis (ATN) score was graded as 0-3 on coded sections by two observers in a blinded manner. 0, no destruction; 3, maximum destruction. The PPAR- γ score was graded as 0-4 on coded sections by two observers in a blinded manner. 0, no staining; 4, maximum intensity. Statistical analysis was performed using the analysis of variance (ANOVA); * p <0.01. Values represent the mean \pm SD.

Kidney tissues sectioned onto microscope slides were deparaffinized. The slides were immersed for 45 min in 0.3% peroxide in methanol to deplete endogenous peroxidase activity. Non-specific binding sites were saturated with 0.2% bovine serum albumin and normal goat serum diluted to 1:66.7 in PBS for 20 min. Primary antibodies against goat PPAR- γ (Santa Cruz, CA, USA) (1:50 dilution in PBS), or control normal goat serum (Vector Laboratories) were used at a dilution of 1:50, applied to tissue sections and incubated in a humidified chamber at room temperature for 30 min. The sections were then washed with PBS for 10 min. Biotinylated rabbit anti-goat IgG (Vector Laboratories) was applied to the tissue sections, which were then incubated at room temperature for 30 min. After washing with PBS for 10 min, the slides were incubated with avidin DH-biotinylated peroxidase (Vector Laboratories) for 45 min. Finally, the color was developed by immersion of the sections in a peroxidase substrate solution including 0.02% peroxide, 3,3' diaminobenzidine tetrahydrochloride, 0.04% nickel chloride and 0.01% hydrogen peroxide in 0.05 M Tris-HCl, pH 7.2, for 2-7 min. The sections were counterstained with hematoxylin (Sigma, St. Louis, MO, USA).

Analysis of acute tubular necrosis and PPAR- γ expression score. To quantify the degree of acute tubular necrosis (ATN), a scale of 0-3 was assigned by two observers in a blinded manner on two separate occasions using coded slides. The average of the scores was evaluated, and the degree of ATN was defined as: no, mild, moderate or severe necrosis. Necrosis, capillary congestion, interstitial edema, cast, destruction, and flat and extended areas of tubular epithelial cells were also evaluated. Similarly, PPAR- γ expression was classified in the renal cortex and renal medulla. PPAR- γ staining was graded on a scale of 0-4 according to the intensity of staining, with a score of 4 considered to be maximum. The same two pathologists assessed ATN and PPAR- γ expression throughout the study.

Statistical analysis. All results are presented as the means \pm standard deviation. Analysis of the data was performed using the analysis of variance (ANOVA) (8).

Results

H&E and immunohistochemical staining of PPAR- γ . In the normal kidney, PPAR- γ staining was weak in the endothelial cells, including the mesangial cells (Fig. 1A). On the other hand, PPAR- γ staining was weak in the interstitial cells and strong in the collecting ducts of the medulla (Fig. 1B). From 1.5 to 5 h after reperfusion, PPAR- γ staining was strong in the endothelial cells including mesangial cells (Fig. 1C), moderate in the interstitial cells and strong in the collecting ducts (Fig. 1D). However, PPAR- γ staining was weak in the tubules (Fig. 1C). Five hours after reperfusion, the internal spaces of the tubular epithelial cells were expanded, and slight destruction of the tubular epithelial cells was apparent (Fig. 1G). The necrotic area extended and encompassed nearly all of the ischemic kidney within 12 h after reperfusion (Fig. 1H). Twelve hours after reperfusion, PPAR- γ staining was weak in the endothelial cells (Fig. 1E), moderate in the interstitial cells and strong in the collecting ducts (Fig. 1F). However, PPAR- γ staining was very weak in the proximal tubules, distal tubules and necrotic tubules at each time point after reperfusion.

Statistical analysis of PPAR- γ expression and acute tubular necrosis score. ATN scores (0 h, 0.1 \pm 0.2; 1.5 h, 0.2 \pm 0.3; 3 h, 0.6 \pm 0.4; 5 h, 1.2 \pm 0.4; 12 h, 2.5 \pm 0.4; 24 h, 2.9 \pm 0.3) were gradually higher at time flow after reperfusion. PPAR- γ scores of the renal cortex (0 h, 1.5 \pm 0.5; 1.5 h, 3.4 \pm 0.6; 3 h, 3.2 \pm 0.7; 5 h, 2.8 \pm 0.7; 12 h, 2.2 \pm 0.8; 24 h, 1.7 \pm 0.8; non-ischemia: 1.2 \pm 0.7) were significantly higher at 1.5, 3 and 5 h after reperfusion than at 0, 12 and 24 h after reperfusion. PPAR- γ scores in the renal medulla (0 h, 3.0 \pm 0.7; 1.5 h, 3.4 \pm 0.6; 3 h, 3.5 \pm 0.6; 5 h, 3.3 \pm 0.5; 12 h, 3.4 \pm 0.5; 24 h, 3.3 \pm 0.6, non-ischemia: 2.9 \pm 0.6) were higher than PPAR- γ scores in the renal cortex (Table I).

Discussion

Renal I/R injury is a clinically significant problem and an invariable consequence of renal transplantation involving the

onset of acute tubular necrosis. This occurs when transplantation is prolonged by a long ischemic interval due to the use of a kidney from a cardiac arrest donor. The longer the ischemic interval, the higher the incidence rate of ATN. The reduction of I/R injury (3) is thus of extreme clinical importance.

I/R injury is best alleviated by the blocking of macrophage-derived cytokines such as tumor necrosis factor- α and interleukin-1, which are abundant in I/R injury tissues. PPAR- γ ligands in part inhibit the expression of nitric oxide, cytokines such as tumor necrosis factor- α and interleukin-1, chemokines and adhesion molecules by antagonizing the activities of transcription factors such as activator protein-1 and nuclear factor- κ B. Among the inducible transcription factors involved in I/R injury, intercellular adhesion molecule-1 and nuclear factor- κ B play important roles (2).

Several reports have demonstrated the efficacy of PPAR- γ ligands in renal I/R injury. Naito *et al* demonstrated that the PPAR- γ ligand pioglitazone ameliorated reperfusion-induced intestinal injury in rats and inhibited the increase in neutrophil accumulation associated with tumor necrosis factor- α expression (9). Sivarajah *et al* demonstrated that the PPAR- γ ligands rosiglitazone and ciglitazone reduced the renal dysfunction and injury associated with I/R of the kidney (10). Yue *et al* demonstrated that rosiglitazone reduced myocardial infarction and improved contractile dysfunction caused by I/R injury (11). Besides I/R injury, Buckingham *et al* demonstrated that the PPAR- γ ligand rosiglitazone protected against nephropathy and pancreatic islet abnormalities in rats (12). However, there have been no reports on the correlation between the progression of time and PPAR- γ expression in I/R injury.

Generally, PPARs modulate the activities of different immune cell types, such as monocytes/macrophages, lymphocytes and endothelial cells. Braissant *et al* demonstrated that PPAR- γ is highly expressed in hepatocytes, cardiomyocytes, enterocytes and the proximal tubule cells of the kidney. PPAR- β is expressed ubiquitously and often at higher levels than PPAR- α or - γ . PPAR- γ is predominantly expressed in the adipose tissue and the immune system of rats (13). Kawahito *et al* found markedly enhanced expression of PPAR- γ in macrophages, as well as modestly enhanced expression in the synovial lining layer, fibroblasts and endothelial cells in a rat adjuvant-induced arthritis model, in which PPAR- γ was localized predominantly to the perinuclear region and the cytoplasm (2). Using RT-PCR, Yang *et al* demonstrated that PPAR- γ was abundant in the renal inner medulla and localized to the inner medullary collecting duct and renal medullary interstitial cells in normal kidney tissues (14). Iwashima *et al*, also using RT-PCR, demonstrated that PPAR- γ was expressed in cultured mesangial cells (15). Asano *et al* found that PPAR- γ was expressed in rat mesangial cells and that its differentiation was modulated by PPAR- γ ligands (16). Besides I/R injury, Paueksakon *et al* demonstrated that PPAR- γ expression was increased in areas of sclerosis in arteries and glomeruli, with expression of both in glomerular mesangial, parietal and visceral epithelial cells. Infiltrating macrophages in glomeruli were PPAR- γ negative, in contrast with their positivity in macrophages, in control cases of carotid artery plaque and in renal interstitial macrophages by immunostaining in human diabetic nephropathy (17).

In the present study, we found PPAR- γ to be strongly expressed in the collecting ducts, interstitial cells and endothelial cells including mesangial cells. The extent and intensity of PPAR- γ in these cells in the renal cortex were most intense 1.5-5 h after renal I/R injury. Twelve hours after renal I/R injury, we found that PPAR- γ was weakly expressed in these cells. In the tubules, PPAR- γ was weakly expressed at each time point after renal I/R injury. On the other hand, PPAR- γ expression in the renal medulla was strong at each time point after renal I/R injury and in the normal kidney. These results suggest that the relationship between PPAR- γ and renal I/R injury is stronger in the renal cortex than in the renal medulla.

It is possible, therefore, that PPAR- γ ligands control PPAR- γ expression in the collecting ducts, interstitial cells and endothelial cells including mesangial cells in renal I/R injury. In conclusion, PPAR- γ is induced in the endothelial cells including the mesangial cells, in the interstitial cells and in the collecting ducts in a rat model of renal I/R injury.

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