Deletion of PPAR-γ in immune cells enhances susceptibility to antiglomerular basement membrane disease

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Abstract: Activation of the nuclear hormone receptor peroxisome proliferator-activated receptor gamma (PPAR-γ) has been shown to be immunoregulatory in autoimmune diseases by inhibiting production of a number of inflammatory mediators. We investigated whether PPAR-γ gene deletion in hematopoietic cells would alter disease pathogenesis in the antiglomerular basement membrane (anti-GBM) mouse model. PPAR-γ⁺/⁺ and PPAR-γ⁻/⁻ mice were immunized with rabbit antimouse GBM antibodies and lipopolysaccharide and evaluated for two weeks. Although both the PPAR-γ⁺/⁺ and PPAR-γ⁻/⁻ mice had IgG deposition in the glomerulus and showed proteinuria two weeks after injection, glomerular and tubulointerstitial disease in PPAR-γ⁻/⁻ mice were significantly more severe compared with the PPAR-γ⁺/⁺ animals. We observed that the PPAR-γ⁻/⁻ mice had decreased CD4⁺CD25⁺ regulatory T cells and an increased CD8⁺:CD4⁺ ratio as compared with the PPAR-γ⁺/⁺ mice, suggesting that PPAR-γ has a role in the regulation of T cells. Furthermore, plasma interleukin-6 levels were significantly increased in the PPAR-γ⁻/⁻ mice at two weeks as compared with the PPAR-γ⁺/⁺ animals. Taken together, these studies show that the lack of PPAR-γ expression enhances inflammatory renal disease in the anti-GBM antibody-induced glomerulonephritis mouse model and suggests targeting PPAR-γ may have therapeutic efficacy.

Keywords: PPAR-γ, antiglomerular basement membrane disease

Introduction

Antiglomerular basement membrane (anti-GBM) disease is an autoimmune disorder that affects the glomeruli of the kidneys.¹ In this disease, glomerular capillaries become targets of autoantibodies, or anti-GBM antibodies, which are directed against an antigen normally present in the GBM and alveolar basement membrane.² The GBM antigen that is responsible for this disease is a component of the alpha-3 chain of Type IV collagen. The resulting clinical syndrome encompasses a spectrum ranging from mild or no renal involvement to rapidly progressive glomerulonephritis. Most patients have both pulmonary and kidney involvement.³ Clinically, anti-GBM disease that induces both glomerulonephritis and pulmonary hemorrhage is termed Goodpasture’s syndrome.¹

Lupus nephritis is an immune-mediated disease in which T cells, B cells, and innate immune cells have been shown to have pathogenic roles. In mouse models of spontaneous lupus nephritis, disease takes 6–12 months to manifest, necessitating the development of models that will induce lupus-like disease over a quicker timeframe. One such
model to study spontaneous lupus nephritis is the experimental anti-GBM mouse.4,5 GBM glomerulonephritis, characterized by crescent formation and necrotizing inflammation of the glomerular capillary, is the most severe form of glomerulonephritis. When given anti-GBM antibodies and inflammatory stimulation, these animals develop glomerular basement nephritis that in many ways resembles lupus nephritis and Goodpasture’s disease. The studies of spontaneous lupus nephritis in mouse models and experimental anti-GBM disease have provided valuable insights into the underlying mechanisms of human lupus nephritis.4,6

Peroxisome proliferator-activated receptors (PPARs) belong to the 48-member superfamily of nuclear receptor proteins that function as transcription factors, regulating gene expression. PPARs are receptors for endogenous lipid molecules and are the molecular targets for drugs against Type 2 diabetes.7,8 They represent promising new targets for the treatment and prevention of inflammatory and autoimmune disorders, such as inflammatory bowel disease and systemic lupus erythematosus.9–11 Treatment with rosiglitazone, a pharmacologic agonist for PPAR-γ of the thiazolidinedione class of insulin-sensitizing drugs, has therapeutic efficacy: it reduces autoantibody production, atherosclerosis, and renal injury in lupus nephritis mice, showing reduced glomerular scarring and reduced inflammation in the renal cortex.12,13 There are three known PPARs (α, β, and γ) that each differ in their tissue distribution and functional activity.14 PPAR-γ is expressed in T and B cells, monocytes/macrophages, dendritic cells, and epithelial cells.15,16 There are three isoforms of PPAR-γ:17 the first and third are identical when fully translated and only differ in their splice variants, whereas the second differs from the other isoforms at the N-terminus.18 All three isoforms have been identified in adipocytes. Additionally, PPAR-γ1 is present in virtually all other tissues, including smooth muscle and splenic tissue. PPAR-γ3 is expressed in macrophages and in the colon.19

Whole body deletion of PPAR-γ causes embryonic lethality, but several conditional knockout mouse models have been developed using homozygous floxed PPAR-γ mice that result in epithelial and hematopoietic cells lacking PPAR-γ.20–23 PPAR-γ conditional knockout peritoneal macrophages have markedly reduced expression of ABCG1, reduced cholesterol efflux, and more atherosclerosis when crossed to a proatherogenic mouse model, such as a low-density lipoprotein receptor knockout.19 In our current studies, we sought to determine if the deletion of PPAR-γ from hematopoietic cells would result in more severe development of disease in the anti-GBM mouse model, given the anti-inflammatory and immunomodulatory effects of PPAR-γ expressed in immune cells.

Materials and methods

Mice

In this project we used PPAR-γ floxed mice expressing the Cre transgene ((PPAR-γ fl/fl; MMTV-Cre- or PPAR-γ+/−) and PPAR-γ fl/fl; MMTV-Cre− littermate mice (PPAR-γ+/+). These mice undergo premature termination of translation due to the enzymatic activity of recombinase in genomic DNA on hematopoietic and epithelial cells.20,24,25 We have previously used these mice to cause conditional deletion of PPAR-γ in mouse models of irritable bowel disease, obesity and diabetes, and inflammation-driven colorectal cancer.25–28 All experimental procedures were approved by the Institutional Animal Care and Use Committee of Virginia Polytechnic Institute and State University, and met or exceeded requirements of the Public Health Service/National Institutes of Health and the United Stated Animal Welfare Act as amended.

Anti-GBM sera

Anti-GBM serum was provided by John Zhang (Medical University of South Carolina, Charleston, SC). Essentially, glomeruli from C57BL/6 mice were isolated with a series of grading sieves (150, 106, and 63 μm mesh) and sonicated for seven minutes. Rabbits were next immunized with glomerular sonicates in complete Freund’s adjuvant, followed by two injections of antigen in incomplete Freund’s adjuvant (2 mg antigen per rabbit per injection), three weeks apart. Sera harvested from these rabbits seven weeks after the initial injection were tested by direct immunofluorescence to demonstrate strong glomerular binding.

Induction of lupus-like disease

Six- to eight-week-old mice were injected with a single dose of 100 μg lipopolysaccharide (Sigma-Aldrich, St Louis, MO) in phosphate-buffered saline intraperitoneally and administered 250 μg anti-GBM sera in phosphate-buffered saline intravenously on day 0 to induce disease. At day 14, the mice were euthanized and the tissues were collected and analyzed for pathology.5

Measurement of proteinuria

As a measure of renal function, urine was collected at days 0, 7, and 14 and tested for proteinuria by a standard semiquantitative test using Bayer Multistix dipsticks (Bayer, Fernwald, Germany). Results were graded according to the
manufacturer’s instructions as negative, trace (15–20 mg/dL of albumin), 1+ (30 mg/dL), 2+ (100 mg/dL), 3+ (300 mg/dL), or 4+ (>500 mg/dL).

Cytokine enzyme-linked immunosorbent assay
Interleukin-6 (IL-6) levels in the sera were quantified by an enzyme-linked immunosorbent assay according to the manufacturer’s instructions (R&D Systems Inc., Minneapolis, MN, USA).

Flow cytometric analysis
Flow cytometric analysis was performed using various combinations of antibodies, including fluorescein isothiocyanate (FITC)-conjugated CD44, R-phycoerythrin-conjugated CD4, FITC-conjugated CD8, and PerCP-CY5.5-conjugated CD25 rat antimouse monoclonal antibodies (BD Pharmingen, San Diego, CA). Splenic cells were isolated as previously described.29 Briefly, spleen lymphocytes from PP AR-γ−/− and PP AR-γ−/+ mice were aseptically dissociated, treated with Tris-ammonium chloride lysis buffer (pH 7.2) to remove erythrocytes, washed, and placed in complete medium consisting of 10% heat-inactivated fetal bovine serum, 200 mM L-glutamine, 5000 IU/mL penicillin, 5000 µg/mL streptomycin, and 100 × nonessential amino acids. Cell numbers were adjusted to 5 × 10⁶ cells/mL, stained with monoclonal antibodies or appropriate fluorochrome-tagged isotype antirat IgG₂a control antibodies, and analyzed on a Coulter Epics XL/MXL flow cytometer (Hialeah, FL).

Kidney pathology
As we previously described, renal pathology was assessed by a veterinary pathologist who was blinded to the treatment groups.30 At day 14, the mice were euthanized for pathologic evaluation. At the time of euthanasia, the mice were weighed and the kidneys were removed and divided into sections. One portion was placed in neutral buffered formalin for subsequent embedding in paraffin, sectioning, and hematoxylin and eosin and periodic acid-Schiff staining. Sections were assessed via light microscopy for glomerular proliferation, glomerular inflammation, glomerular size, number of nuclei per glomerulus, crescents, necrosis, and fibrosis. Each of these parameters was graded as 0 (normal), 1 (mild increase in mesangial matrix and cellularity), 2 (moderate increase in mesangial matrix and cellularity), 3 (focal endocapillary hypercellularity, obliterated capillary lumen, and marked thickening of glomerular basement membrane), or 4 (crecent formation, segmental necrosis, marked hypercellularity, and hyalinized glomeruli), and an overall glomerular score was derived.31 One portion of the kidney was frozen in optimal cutting temperature media and cut into 5 µm sections and stained with FITC-conjugated antibodies (goat antimouse IgG diluted 1:100, Pierce, Rockford, IL).

Statistical analysis
Statistical analysis was performed by Student’s t-test. A P value < 0.05 was considered statistically significant.

Results
Rabbit IgG deposition in kidneys after anti-GBM antibody injection
Before assessing the impact of PPAR-γ gene deletion on susceptibility to immune-mediated glomerular basement nephritis, we first sought to determine if the kidneys from the PP AR-γ++/− mice and PP AR-γ−/+ mice showed similar glomerular staining patterns to anti-GBM treatment. Fourteen days after the induction of disease, the anti-GBM antibodies were observed to bind exclusively to the glomerular capillary wall in a linear pattern (Figure 1). Additionally, the kidneys from both the PP AR-γ+++/− and PP AR-γ−/+ mice showed similar amounts of rabbit antimouse IgG antibodies in their glomeruli.

Determination of proteinuria
Prior to the induction of disease (day 0), both the PP AR-γ++/− and PP AR-γ−/+ mice showed minimal proteinuria (Figure 2). At day 14, both the PP AR-γ++/− and PP AR-γ−/+ animals had significantly greater amounts of proteinuria by dipstick analysis compared with the baseline. The differences were not statistically significant between the PP AR-γ++/− and PP AR-γ−/+ mice at day 14.

Effect of anti-GBM sera on splenic tissue
At day 14, the animals were sacrificed and the spleen weights were measured. The spleen weights and spleen to body weights were not statistically different between PP AR-γ++/− and PP AR-γ−/+ mice (0.056±0.14 g versus 0.054±0.14 g for spleen and 22.5±2.3 g versus 23.4±3.2 g body weight, respectively).

Figure 1 Representative picture of IgG staining in the glomerulus at D14. Renal deposition of anti-GBM antibodies between PP AR-γ++/− (right) and PP AR-γ−/+ mice (left). Abbreviations: GBM, glomerular basement membrane; PPAR-γ, peroxisome proliferator-activated receptor gamma.
To characterize the splenic phenotype, we isolated the dissociated splenocytes. There was no difference in total splenocyte numbers between the PPAR-\(\gamma^{+/+}\) and PPAR-\(\gamma^{-/-}\) mice (data not shown). Next, we assessed the T cell populations by flow cytometry. We examined the CD4\(^+\):CD8\(^+\) ratio, CD4\(^+\)CD25\(^+\) (T regulatory cells), and CD4\(^+\)CD44\(^+\) (activated memory T cells) due to prior reports showing PPAR-\(\gamma\) expression or activation modulates T cell profiles by altering the CD4\(^+\):CD8\(^+\) ratio and the T regulatory cell population.\(^32,33\) Our results showed that there was no difference in the CD4\(^+\)CD44\(^+\) expression profiles in PPAR-\(\gamma^{+/+}\) mice compared with PPAR-\(\gamma^{-/-}\) mice. Intriguingly, we found that the PPAR-\(\gamma^{-/-}\) mice exhibited a significant decrease in the CD4\(^+\):CD8\(^+\) T cell ratio, as well as a decrease in CD4\(^+\)CD25\(^+\) cells compared with the PPAR-\(\gamma^{+/+}\) mice (Figure 3 and Table 1).

**Kidney pathology**

Both PPAR-\(\gamma^{+/+}\) and PPAR-\(\gamma^{-/-}\) mice developed renal disease with anti-GBM antibody administration characterized by relatively mild glomerular inflammation. However, the PPAR-\(\gamma^{-/-}\) mice showed significantly more severe disease compared with the PPAR-\(\gamma^{+/+}\) mice (Figure 4). In addition to increased glomerular inflammation, the PPAR-\(\gamma^{-/-}\) mice had increased mesangial matrix that obliterated the glomerular architecture (as shown in the periodic acid-Schiff-stained sections of the kidney). Furthermore, several of the PPAR-\(\gamma^{-/-}\) mice showed severe interstitial abnormalities.

**Evaluation of cytokine levels**

We measured serum levels of the proinflammatory cytokine IL-6 as a measure of inflammation (Figure 5). Prior to induction of anti-GBM disease, IL-6 levels were not detectable. Two weeks after anti-GBM antibody administration, IL-6 levels in the PPAR-\(\gamma^{-/-}\) mice were significantly elevated compared with PPAR-\(\gamma^{+/+}\) mice.

**Discussion**

The activation of PPAR-\(\gamma\) was originally shown to block the proinflammatory effects of lipopolysaccharide and various cytokines in both monocytes and macrophages by antagonizing the activities of AP-1, STAT, and NF-xB.\(^34,35\) Specifically, activation of PPAR-\(\gamma\) decreased the production of nitric oxide, IL-6, and tumor necrosis factor-alpha. Since these initial observations, numerous studies have further characterized the expression, regulation, and mechanism of PPAR-\(\gamma\)'s anti-inflammatory activity.

In our current studies, we sought to determine the effects of PPAR-\(\gamma\) gene deletion in hematopoietic and epithelial cells using a mouse model of induced glomerulonephritis. The administration of anti-GBM antibodies and lipopolysaccharide stimulation serves as an acceptable model for murine glomerulonephritis with characteristics of Goodpasture’s syndrome as well as lupus nephritis.\(^4,36\) Our studies demonstrate that PPAR-\(\gamma\) has a role in the maintenance of regulatory T cell numbers. We found that PPAR-\(\gamma\) gene deletion resulted in a decreased CD4\(^+\)CD25\(^+\) T cell population, along with an increase in disease in mice challenged with lipopolysaccharide and anti-GBM antibodies. This coincides with previous reports demonstrating that the deletion of PPAR-\(\gamma\) in T regulatory cells abrogates their ability to prevent CD4\(^+\) T cell-induced colitis in adoptive transfer studies and ameliorate graft versus host disease.\(^26,33\) We have previously shown the importance of T regulatory cells in decreasing disease severity in lupus mice.\(^37\) Together, these studies indicate that a decrease in T regulatory cell numbers and/or function contributes to the development of autoimmune diseases.

Several lines of evidence suggest that PPAR-\(\gamma\) exerts anti-inflammatory effects by negatively regulating the expression of proinflammatory genes induced during macrophage differentiation/activation and the production of proinflammatory cytokines.\(^38\) We observed a significant increase in the production of Th1/Th17 cytokine IL-6 after PPAR-\(\gamma\) gene deletion. IL-6 is not only important for the differentiation of Th1/Th17 phenotypes, but is also critical for inhibiting the differentiation of T regulatory cells.\(^39\) This is consistent with our flow cytometry analyses showing a decreased T regulatory cell population of mice deficient in PPAR-\(\gamma\). The general skewing of immune cell phenotypes may play a significant role during the pathogenesis of renal nephritis and lupus in mice and humans. Moreover, the deletion of PPAR-\(\gamma\) in CD4\(^+\) T cells results in enhanced antigen-specific proliferation and overproduction of interferon-\(\gamma\) in response to IL-12, indicating that adequate expression of PPAR-\(\gamma\) in CD4\(^+\) T cells is required to downregulate excessive Th1 responses.\(^26\)
Perhaps the most striking difference we observed in the PPAR-γ−/− mice was the more severe development of glomerular and interstitial lesions compared with the PPAR-γ+/+ mice, demonstrating the role for PPAR-γ gene expression in modulating renal disease. Direct binding of antiglomerular antibodies to glomerular antigens plays a key role in disease pathogenesis in the GBM model. All specimens of the anti-GBM antibody-treated mice showed linear deposition of IgG along the GBM accompanied by complement C3 deposits. Our observations by light microscopy revealed a significant overall increase in glomerular pathology, including crescent formation in the PPAR-γ−/− mice, while the PPAR-γ+/+ mice showed less pathology. Interestingly, we found that proteinuria was similar in both the PPAR-γ−/− and PPAR-γ+/+ mice.

Table 1 Percentage of T cell markers in the spleens of mice with autoimmune anti-GBM glomerulonephritis (n = 10)

<table>
<thead>
<tr>
<th>Parameter</th>
<th>CD4+ T cells</th>
<th>CD8+ T cells</th>
<th>CD8+/CD4+ ratio</th>
<th>CD4+CD44+ cells</th>
<th>CD4+CD25+ cells</th>
</tr>
</thead>
<tbody>
<tr>
<td>PPAR-γ−/−</td>
<td>13.95 ± 0.4+</td>
<td>15.86 ± 2.6</td>
<td>1.16 ± 0.08</td>
<td>17.62 ± 2.8</td>
<td>2.84 ± 0.31</td>
</tr>
<tr>
<td>PPAR-γ+/+</td>
<td>10.98 ± 0.3</td>
<td>14.95 ± 2.4</td>
<td>1.36 ± 0.13*</td>
<td>17.17 ± 3.4</td>
<td>1.96 ± 0.28*</td>
</tr>
</tbody>
</table>

*P < 0.05.

Abbreviations: GBM, glomerular basement membrane; PPAR-γ, peroxisome proliferator-activated receptor gamma.
This finding could be due to the single time point at which we collected urine and our method of determination of proteinuria. A 24-hour urine collection may have shown a better representation of urine output with respect to proteinuria.

In lupus nephritis mouse models, PPAR-γ activation has shown therapeutic effects, leading to the reduction of disease. Recently Venegas-Pont et al reported that PPAR-γ agonists showed beneficial effects on renal function in the NZB/W lupus mouse. Others have suggested that PPAR-γ modulates renal disease in lupus nephritis through induction of adiponectin. These studies support our findings for a therapeutic benefit of PPAR-γ in autoimmune nephritis. We have previously demonstrated that mesangial cells produce endogenous PPAR-γ ligands. The results presented in this paper are in line with the effect of endogenous PPAR-γ ligands on PPAR-γ expressed in immune cells, because the

![Figure 4](image)

**Figure 4** Renal pathology of PPAR-γ+/- and PPAR-γ-/- two weeks after anti-GBM administration. A Representative photomicrograph of a kidney from a PPAR-γ+/- (upper left) and PPAR-γ-/- (upper right) with hematoxylin and eosin stain (insert 25 × magnification of glomeruli). Note the severe cellular infiltrate and crescent formation in the PPAR-γ-/- mouse kidney. Representative photomicrograph of a kidney from a PPAR-γ+/- (lower left) and PPAR-γ-/- (lower right) with periodic acid-Schiff stain. Note the increased deposition of periodic acid-Schiff-positive mesangial matrix in the PPAR-γ-/- mouse kidney. B Tabulation of renal tubular and C glomerular interstitium. **Abbreviations:** GBM, glomerular basement membrane; PPAR-γ, peroxisome proliferator-activated receptor gamma.
deletion of PPAR-γ resulted in increased kidney pathology in autoimmune mice. Additionally, endogenously generated lipid molecules can also activate PPAR-γ, suggesting pharmacologic or dietary intervention may not be required for PPAR-γ modulation of disease. Taken together, these observations support a critical role for PPAR-γ expression in the maintenance of kidney homeostasis and support the notion that PPAR-γ may be a therapeutic avenue to target for inhibition of autoimmune inflammatory kidney diseases.

Disclosure
The authors report no conflicts of interest in this work.

References


