

Inducible Nitric Oxide Synthase (iNOS) Expression Is Increased in Lipopolysaccharide (LPS)-Stimulated Diabetic Rat Glomeruli: Effect of ACE Inhibitor and Angiotensin II Receptor Blocker

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Previously, we reported that high glucose enhanced cytokine-induced nitric oxide (NO) production by rat mesangial cells (MCs), and that the enhanced expression of the iNOS pathway may promote extracellular matrix accumulation by MCs. The present study was designed to examine whether the iNOS pathway is pathologically altered in experimental diabetic nephropathy, and whether therapy with angiotensin converting enzyme (ACE) inhibitor (imidapril: I) or angiotensin II type I receptor (AT1) blocker (L-158,809: L), ameliorates these changes. Male Sprague-Dawley rats were injected with diluent (control: C) or streptozotocin. At sacrifice after 4, 8 and 12 weeks, rats underwent either a 4 hour placebo or an intraperitoneal lipopolysaccharide (LPS, 2 mg/kg) challenge. Systolic blood pressure (SBP) and urinary protein excretion (UPE) increased significantly in diabetic (D) rats compared with C. The basal expression of glomerular iNOS mRNA was increased in D rats compared with that of C rats, by reverse-transcription (RT)-polymerase chain reaction (PCR), whereas there was no significant difference in the level of protein by Western blot analysis. Upon LPS stimulation, the iNOS mRNA and protein expression was significantly elevated in D rats. In D rats, this up-regulation, of LPS-stimulated iNOS expression, was equally ameliorated both by I and L in mRNA and protein levels. From immunohistochemistry (IHC), there was a nega-

tive staining for the iNOS within the glomeruli of five C rats without LPS treatment, but one of four rats, with LPS treatment, showed minimal iNOS staining in the glomeruli. In D rats, the glomerular mesangium and podocytes were positive for iNOS in each of three out of five rats with, and without, LPS treatment.

In conclusion, LPS-stimulated glomerular iNOS expression was enhanced in diabetic nephropathy, and the activation of angiotensin II may play a role in this enhancement.

Key Words: iNOS, diabetic nephropathy, angiotensin II

INTRODUCTION

Diabetic nephropathy is the leading cause of end-stage renal disease and one of the serious microvascular complications in diabetes mellitus (DM). Despite numerous studies, the pathophysiology of DM nephropathy is not completely understood. One of the mediators implicated in the development of DM nephropathy is the NO system.¹ NO acts as a vascular and neuronal messenger activating soluble guanylate cyclase, resulting in increased levels of cGMP. The family of NOS proteins are classified into; the constitutive type, neuronal NOS (nNOS), endothelial NOS (eNOS), and the inducible type. Within the kidney, high levels of nNOS are located in the macula densa. eNOS is found in glomerular afferent and efferent arterioles, with iNOS being localized in the cells of the mesangium.² Since there is enhanced production of a variety of

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cytokines including TNF- α and IFN- γ in diabetes³ and macrophages infiltrated into the glomeruli of rats in the earliest stages of diabetes,⁴ it can be postulated that the cytokine-induced iNOS pathway may play an important role in glomerular injury seen in diabetic kidneys. However, the exact role of iNOS in mediating DM nephropathy has not been fully elucidated, and the results of previous studies have been inconsistent. Some authors have reported an inhibition of induced NO production in cultured rat MCs by certain proinflammatory stimuli.⁵ Others were able to demonstrate an augmented LPS- or cytokine-induced NO production in murine MCs.⁶ In our own study, we reported that exposure of rat MCs to high glucose led to modest, but significant increases in iNOS mRNA, protein expression and NO production.⁷

The aim of this study was to assess whether the iNOS pathway is pathologically altered in experimental diabetic nephropathy, and if therapy with ACE inhibitor or AT1 blocker ameliorates these changes.

MATERIALS AND METHODS

Animals and experimental protocol

All animal experiments described in this article were conducted in accordance with the Yonsei University Medical College guide for the care and use of laboratory animals. Male Sprague-Dawley rats, weighing 250 to 300 g, were made diabetic using a single intraperitoneal injection of 65 mg/kg streptozotocin (Sigma Chemical Co., St. Louis, MO, USA). Twenty-four hours later, blood glucose levels were determined using tail blood samples. The rats were housed in a temperature-controlled room, were given free access to water and standard laboratory chow during the study period.

Every 4 weeks, body weight (BW), SBP, and UPE were measured. At week 4, 8 and 12, the rats received either a 4 hour placebo, or an intraperitoneal LPS (2 mg/kg, Sigma Chemical Co.) challenge, and were then sacrificed. Kidney weights were measured at the time of sacrifice. Coronal slices of kidney, for routine light micros-

copy, were processed in the standard fashion, and the kidney remnants were prepared for the isolation of glomeruli. Glomeruli were prepared using a sieving technique as previously described,⁸ and the samples stored at -70°C in preparation for the analytical procedures described later in this article. Thin kidney slices, from the rats sacrificed at 12 weeks, were processed for IHC studies.

The effect of ACE inhibitor, or AT1 blocker, on the expression of glomerular iNOS was also examined in LPS-stimulated D rats. Diabetes was induced by the same method as above, and all the rats received LPS stimulation at sacrifice. Four groups were studied for 12 weeks. The first (n=6) and second groups (n=7), were C and D rats without drug treatment respectively; the third group (n=7) were D rats treated with ACE inhibitor (I, 2 mg/kg/day, Donga Pharmaceutical); and the fourth group (n=7) were D rats treated with AT1 blocker (L, 1 mg/kg/day, Merck Sharp and Dohme). I or L was given by gavage. Every 4 weeks, BW, SBP, and UPE were measured. At week 12, the expression of the iNOS mRNA and protein was examined.

Analytical methods

Blood glucose was measured by a glucometer, and 24 hour UPE was determined by the Bradford method, using the Bio-Rad assay (Bio-Rad Laboratories, Inc., Hercules, CA, USA). SBP was measured by the tail-cuff method.

Total RNA extraction

Total RNA was extracted by the addition of 100 μ L of RNA STAT-60 reagent (Tel-Test, Inc., Friendswood, TX, USA) to the glomeruli, followed by glomerular lysis by freezing and thawing three times. Another 700 μ L of RNA STAT-60 reagent was added, and the mixture was vortexed, and stored for five minutes at room temperature. Four hundred microliters of chloroform were added, and the mixture shaken vigorously for 30 seconds. After three minutes, the mixture was centrifuged at 12,000 \times g for 15 minutes at 4°C. The upper aqueous phase, containing the extracted RNA, was transferred to a new tube. RNA was precipitated from the aqueous phase by 400 μ L isopro-

panol, and pelleted by centrifugation at $12,000 \times g$ for 30 minutes at 4°C . The RNA precipitate was washed with 70% ice-cold ethanol, dried using Speed Vac, and dissolved in DEPC-treated distilled water. The RNA yield and quality were assessed based on spectrophotometric measurements at 260 and 280 nm for OD.

RT-PCR

First-strand cDNA was made using a Boehringer Mannheim cDNA synthesis kit (Boehringer Mannheim GmbH, Mannheim, Germany). Total RNA from glomeruli was reverse transcribed using $10 \mu\text{mol/L}$ of random hexanucleotide primer, 1mmol/L dNTP, 8mmol/L MgCl_2 , 30mmol/L KCl, 50mmol/L Tris-HCl, pH 8.5, 0.2mmol/L dithiothreitol, 25 U RNase inhibitor, and 40 U AMV reverse transcriptase. The mixture was incubated at 30°C for 10 minutes and 42°C for one hour, followed by inactivation of the enzyme at 99°C for five minutes.

PCR amplification was performed at a final concentration of $1 \times$ PCR buffer (10mmol/L Tris-HCl, pH 8.3, 50mmol/L KCl, 1.5mmol/L MgCl_2), $20 \mu\text{mol/L}$ dNTP, sense and antisense primers (20pmol for GAPDH; 30pmol for iNOS), Taq polymerase (1U for GAPDH) or Taq-Gold polymerase (1.5U for iNOS) in a total volume of $50 \mu\text{L}$. Sense primer for iNOS was 5'-GGTGAAAGCGGTGTCTTTGC-3', and antisense primer was 5'-GGTGAGACAGTTTCTGGTCG-3', which yielded a 389 bp PCR product. Sense primer for GAPDH was 5'-GACAAGATGGTGAAGGTCGG-3', and antisense primer was 5'-CATGGACTGTGGTCATGAGC-3', which yielded a 538 bp PCR product. The PCR conditions were as follows: (1) GAPDH, 38 cycles, denaturing at 94°C for 30 seconds, annealing at 60°C for 30 seconds, and extension at 72°C for 1 minute; (2) iNOS, 43 cycles, denaturing at 94.5°C for 1 minute, annealing at 60°C for 1 minute, and extension at 72°C for 2 minutes. Initial heating at 95°C for 9 minutes, and final extension at 72°C for 7 minutes, were performed for all PCRs. Each cDNA was amplified with 30 to 50 cycles using specific primers, and the amplification cycles were selected during the exponential phase. The RT-PCR products were separated by electrophoresis, and the band densities

analyzed by laser densitometry (Helena Laboratories, Beaumont, TX, USA).

Western blot analysis

Isolated rat glomeruli were lysed in a sodium dodecyl sulfate (SDS) sample buffer [2% SDS, 10mmol/L Tris-HCl, pH 6.8, 10% (vol/vol) glycerol]. The lysate was centrifuged at $12,000 \times g$ for 15 minutes at 4°C , and the supernatant stored at -70°C until all the rats were sacrificed. Protein concentrations were determined with a Bio-Rad kit. Aliquots, $30 \mu\text{g}$ of protein, were mixed with sample buffer, containing SDS and β -mercaptoethanol, heated at 100°C for five minutes, and underwent electrophoresis in a 7.5% acrylamide denaturing SDS-polyacrylamide gel. Proteins were transferred to a Hybond-ECL membrane (Amersham Life Science, Inc., Arlington Heights, IL, USA) using a Hoeffer semidry blotting apparatus (Hoeffer instruments, San Francisco, CA, USA). The blots were then blocked, using 5% non-fat milk, and probed with iNOS-specific (mouse monoclonal anti-rat antibody; Transduction Laboratories, Lexington, KY) antibody for 1 hour at room temperature. The membrane was washed with PBS-Tween-20 for 1 hour, then probed with peroxidase-conjugated secondary antibody (horseradish peroxidase-linked anti-mouse IgG; Santa Cruz, CA, USA) for 1 hour at room temperature. After washing, immunoblots of the iNOS protein were visualized by enhanced chemiluminescence (Amersham Life Science, Little Chalfont, England, UK). Positive immunoreactive bands were quantified densitometrically.

Immunohistochemistry

Deparaffinized and rehydrated $4 \mu\text{m}$ thick sections were treated with 3% H_2O_2 , to block endogenous peroxidase, and incubated with normal blocking serum, for 30 minutes, to block non-specific staining. Sections were incubated for 1 hour at room temperature, and then overnight at 4°C in a humidity chamber with polyclonal rabbit anti-mouse iNOS antibody ($1:500$ dilution, Affinity Bioreagents, Golden, CO, USA). After washing in PBS (pH=7.5), the sections were sequentially incubated with biotinylated link antibody and

peroxidase-labeled streptavidine (Dako, Santa Barbara, CA, USA). The substrate reaction was carried out using 3-amino-9-ethylcarbazole. The sections were counterstained with hematoxylin and mounted in an aqueous mounting media. As a positive control, sections with pyelonephritis were used. For a negative control, PBS was replaced with primary antibody.

Statistical analysis

Results are presented as the mean \pm standard error (SE). The means between the two groups were compared using the Mann-Whitney U-test. Analysis of variance was used to assess the differences between multiple groups. A *p*-value < 0.05 was used as the criterion for a statistically significant difference.

RESULTS

Clinical parameters

Tables 1 and 2 show the time course data of clinical parameters. Compared with the C group, the mean blood glucose levels of the D rats were significantly higher throughout the 12-week period. The C rats gained weight over the 12-week experiment, however, the D rats failed to gain

weight. The ratio of kidney to body weight was significantly higher in the D rats. The treatment with I or L did not affect these parameters. SBP in the D rats rose significantly at week 4 compared with the C rats, and kept significantly higher levels throughout the observation period of 12 weeks. The treatment with I or L, at the dose administered in this study, resulted in no significant reduction in SBP compared with that in the untreated D rats.

Twenty four-hour UPE was significantly greater in the D rats over the 12-week experiment. The treatment with I or L partly reversed this enhancement of UPE in the D rats.

Expression of iNOS mRNA and protein

Without LPS stimulation, the iNOS mRNA and protein expression were not apparent at week 4 and 8, but were found at week 12. At 12 weeks, the basal expression of glomerular iNOS mRNA was increased in the D rats compared with that of the C rats by RT-PCR (Fig. 1), whereas the Western blot analyses showed no significant difference in the level of protein between the two groups (Fig. 2).

Upon LPS stimulation, significantly higher level of iNOS mRNA expression in the D group was noted from week 4 (1.29 \times), and was maintained throughout the observation period (2.70 \times at

Table 1. Blood Glucose, Body Weight, and Kidney Weight/Body Weight in 4 Groups

Parameter	Control	DM	DM+I	DM+L
Blood glucose (mg/dl)				
Week 0	93.4 \pm 1.3	480.3 \pm 6.1 ^a	487.3 \pm 7.8 ^a	477.0 \pm 6.0 ^a
Week 4	117.3 \pm 5.1	468.7 \pm 9.7 ^a	462.0 \pm 13.1 ^a	473.5 \pm 25.5 ^a
Week 8	138.7 \pm 6.5	465.0 \pm 11.2 ^a	463.5 \pm 18.3 ^a	458.0 \pm 16.0 ^a
Week 12	121.1 \pm 8.1	495.5 \pm 1.2 ^a	498.0 \pm 2.9 ^a	486.0 \pm 12.0 ^a
Body weight (g)				
Week 0	252.9 \pm 4.7	261.1 \pm 4.1	269.8 \pm 4.7	277.0 \pm 3.0
Week 4	397.7 \pm 5.5	250.8 \pm 8.0 ^a	285.8 \pm 17.1 ^a	297.03 \pm 3.0 ^a
Week 8	471.2 \pm 11.4	258.1 \pm 11.8 ^a	264.6 \pm 26.8 ^a	296.5 \pm 55.5 ^a
Week 12	510.3 \pm 24.2	266.3 \pm 18.5 ^a	258.8 \pm 34.0 ^a	284.5 \pm 91.5 ^a
Kidney weight / Body weight ($\times 1000$)				
Week 4	7.1 \pm 0.3	15.3 \pm 0.4 ^a	—	—
Week 8	6.2 \pm 0.2	15.2 \pm 0.8 ^a	—	—
Week 12	6.5 \pm 0.2	15.1 \pm 0.3 ^a	16.8 \pm 2.0 ^a	14.3 \pm 2.3 ^a

Data are expressed as mean \pm SEM, I: imidapril, L: L-158,809, ^a*p* < 0.01 vs. control.

Table 2. Systolic Blood Pressure and 24-Hour Urinary Protein Excretion in 4 Groups

Parameter	Control	DM	DM+I	DM+L
SBP (mmHg)				
Week 0	113.5 ± 3.0	115.5 ± 5.6	113.5 ± 5.7	108.6 ± 11.1
Week 4	122.7 ± 2.9	132.4 ± 3.1 ^a	130.1 ± 11.1	132.5 ± 23.8
Week 8	122.2 ± 1.7	136.9 ± 4.0 ^a	136.6 ± 3.4 ^a	137.2 ± 2.9
Week 12	131.5 ± 4.9	148.1 ± 5.4 ^a	137.3 ± 6.8	142.0 ± 1.7
Urinary protein excretion (mg/d)				
Week 4	23.0 ± 1.5	78.8 ± 7.2 ^a	45.4 ± 1.9 ^{a,b}	66.3 ± 6.0 ^a
Week 8	23.9 ± 1.2	73.8 ± 7.2 ^a	50.2 ± 3.7 ^{a,b}	62.5 ± 3.6 ^a
Week 12	19.8 ± 3.2	90.6 ± 11.3 ^a	59.0 ± 2.8 ^{a,b}	61.6 ± 6.6 ^{a,b}

Data are expressed as mean ± SEM.

I: imidapril, L: L-158,809, ^a*p* < 0.05 vs. control, ^b*p* < 0.05 vs. DM.

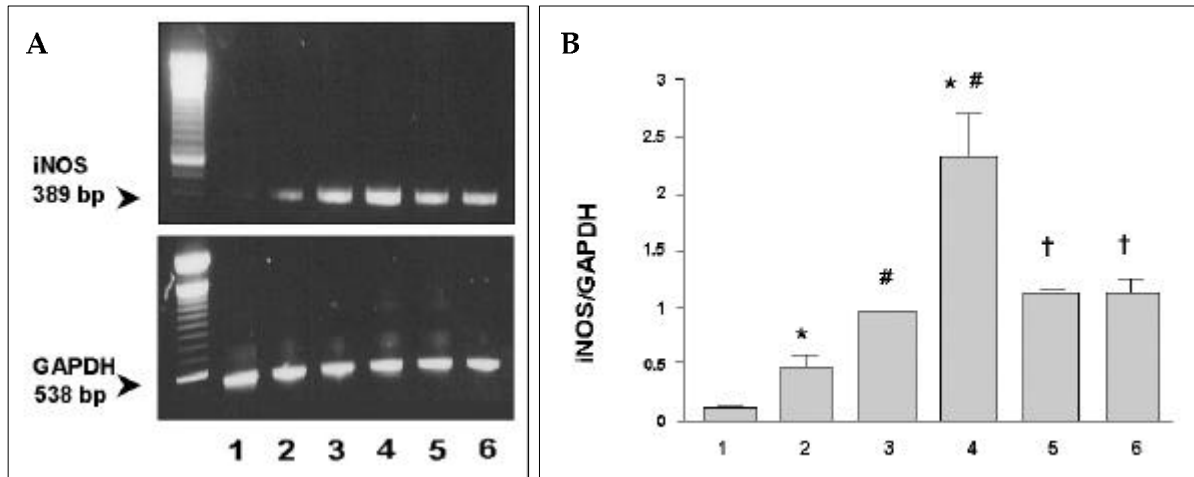


Fig. 1. (A) A representative RT-PCR at 12 weeks. (B) Values are the intensity of densitometric readings of iNOS mRNA corrected for GAPDH mRNA. Lane 1, control (C); lane 2, DM; lane 3, C + LPS; lane 4, DM + LPS; lane 5, DM + LPS + I; lane 6, DM + LPS + L. Values are expressed as a relative to that of C + LPS from four experiments. I: imidapril, L: L-158,809. **p* < 0.05 vs. paired C; #*p* < 0.05 vs. corresponding values without LPS; †*p* < 0.05 vs. DM+LPS.

week 8, 2.42 × at week 12) compared with the corresponding C group (Fig. 3). Administration of I or L in the D rats attenuated iNOS mRNA expression (1.12 ×, 1.12 × of the C group) at week 12 (Fig. 1). At the level of protein, D glomeruli expressed significantly higher levels of iNOS at week 8 (6.16 ×) and 12 (1.86 ×) compared with the corresponding C group (Fig. 3). Administration of I (0.77 × of the C group) or L (0.94 × of the C group) for 12 weeks, restored the enhanced expression of the iNOS protein observed in the D glomeruli compared to that seen in the glomeruli from the C rats (Fig. 2).

Histology and IHC staining for iNOS protein

Cytoplasmic clear cell change of the distal tubular epithelium was present in the D rats regardless of LPS treatment from week 4 onwards. The glomeruli of the D rats were hypertrophied compared with the C rats, but were normocellular. Mesangiolytic was present in five of the twenty LPS-treated D rats.

There was negative staining for iNOS within the glomeruli of five C rats without LPS treatment, but one of four rats with LPS treatment showed minimal iNOS staining in the glomeruli. In the D rats, the glomerular mesangium and

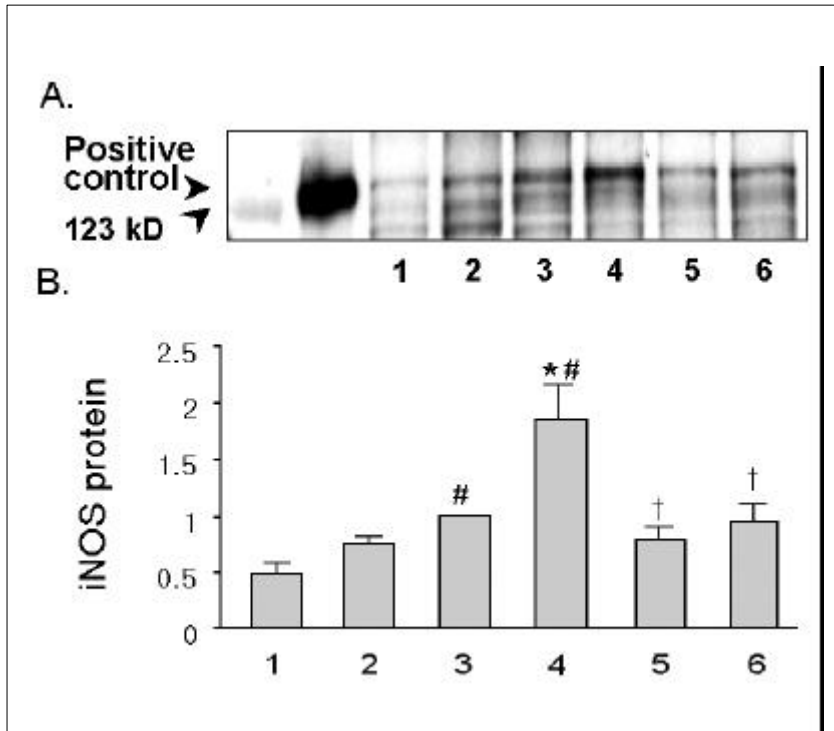


Fig. 2. (A) A representative Western blot at 12 weeks. (B) Values are the intensity of densitometric readings of iNOS protein. Lane 1, control(C); lane 2, DM; lane 3, C+LPS; lane 4, DM + LPS; lane 5, DM+LPS+I; lane 6, DM +LPS+L. Values are expressed as a relative to that of C+LPS from four experiments. I: imidapril, L: L-158,809. * $p < 0.05$ vs. C + LPS; # $p < 0.05$ vs. corresponding values without LPS; † $p < 0.05$ vs. DM + LPS.

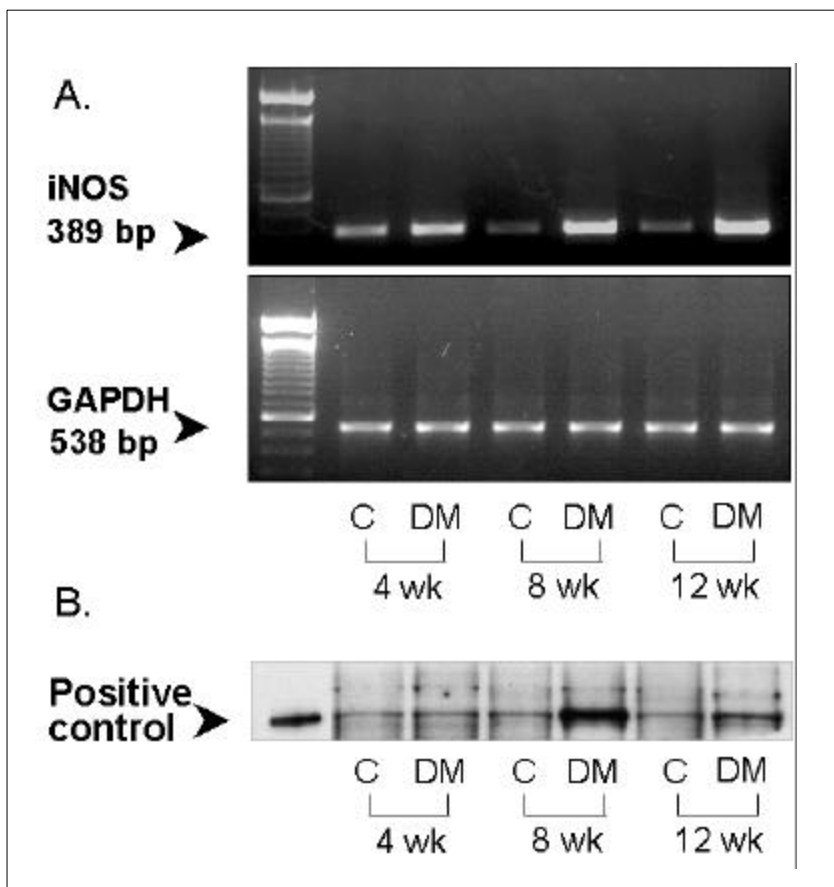


Fig. 3. (A) A representative RT-PCR of four separate experiments at indicated times. There was 1.29-fold, 2.70-fold and 2.42-fold increases in glomerular iNOS mRNA expression in DM rats compared with corresponding control rats at 4, 8 and 12 weeks. (B) A representative iNOS Western blot of four separate experiments at indicated times. Densitometric quantitation revealed that there was 6.16-fold and 1.86-fold increases in DM rats compared with corresponding control rats at 8 and 12 weeks. All the rats of these experiment were treated with LPS.

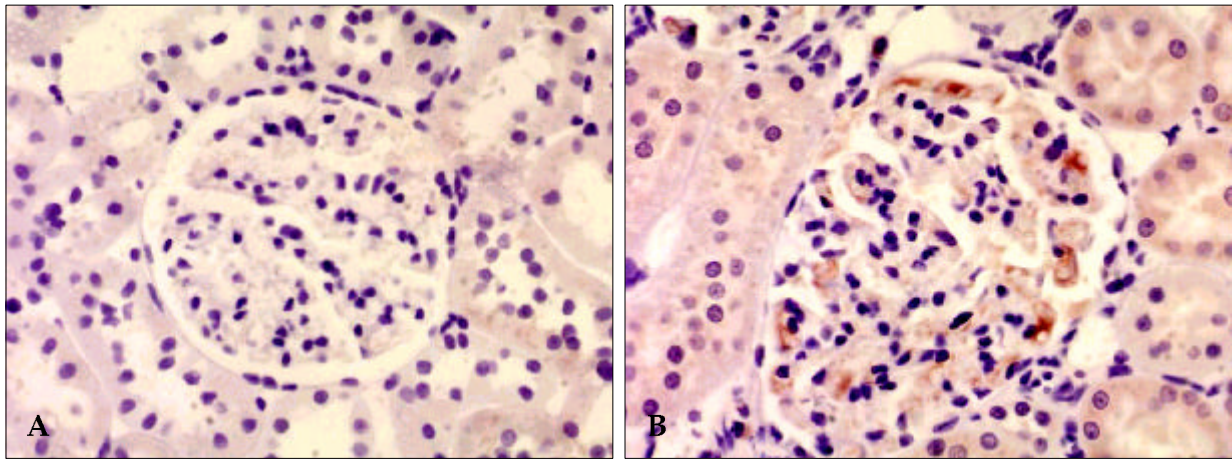


Fig. 4. Immunohistochemical staining for iNOS is negative in a glomerulus of control rats (A), but positive in the mesangium and capillary loop of diabetic rats (B).

podocytes were positive for iNOS in three of the five rats with, and without, LPS treatment (Fig. 4). No rats showed more than 10% positivity of the glomeruli.

DISCUSSION

Although several studies have suggested the alteration of the NO system could contribute to the pathogenesis of DM nephropathy,⁹⁻¹³ they primarily focused on the constitutive NOS (cNOS) pathway, with the role of the iNOS pathway remaining unclear. It was logical to consider the possibility that iNOS may be involved in the pathogenesis of DM nephropathy because iNOS is increased in the beta cells of the pancreas after streptozotocin treatment,¹⁴ and high glucose concentrations have been demonstrated to increase iNOS mRNA expression in macrophages and glomerular MCs.⁶⁻⁷

Our study shows that LPS-stimulated iNOS expression is increased in diabetic glomeruli at the levels of mRNA and protein. Since previous studies described the expression of iNOS was not detected in the basal condition,¹⁵ and also reported enhanced production of a variety of cytokines including TNF- α and IFN- γ in diabetes,³ the present study examined iNOS expression with, and without, LPS stimulation. The basal expression of iNOS mRNA and protein was not

apparent at 4 and 8 weeks but was found at week 12. Although the Western blot analyses have not confirmed the difference in basal expression of iNOS protein between C and D glomeruli, the results of the RT-PCR and IHC suggests a greater expression of iNOS in D glomeruli compared with that of the C. With LPS stimulation, the D rats had more iNOS mRNA and protein expression from 4 weeks onwards. Recently, Sugimoto et al. reported that the expression of iNOS significantly increased in the D rat glomeruli at 52 weeks after the induction of diabetes.¹⁶ They suggested the sequential pathway of advanced glycation end-products (AGE)-cytokine-iNOS in the development of diabetic nephropathy. Because we stimulated the rats with LPS in the early stages of diabetes, the role of AGE in the induction of iNOS was not evaluated in this study. The results of this study have confirmed our earlier *in vitro* study, in which exposure of rat MCs to high glucose enhanced cytokine-induced iNOS mRNA and protein with increased NO production.⁷

Angiotensin II leads to renal vasoconstriction, regulating glomerular perfusion, as well as filtration, and directly influences the mesangial constrictive response.¹⁷ Angiotensin II is a potent regulator of MC phenotypical changes, such as migration,¹⁸ proliferation,¹⁹ or protein kinase C expression.²⁰ This supports its role as a growth factor, profibrogenic and proinflammatory media-

tor,²¹ and regulator of additional proinflammatory mediators, such as: transforming growth factor- β (TGF- β)²² or monocyte-chemoattractant protein-1 (MCP-1).²³

Angiotensin II and NO have been postulated to interact closely. Conflicting data has been published regarding the effect of angiotensin II on iNOS expression in various experimental models. Some authors report a down-regulation, or inhibition, of the induced NO production in cultured rat astroglial cells by certain proinflammatory stimuli,²⁴ and similarly, in renal cells such; as mouse proximal tubular cells²⁵ and rat MCs.²⁶ Others were able to demonstrate an augmented LPS- or cytokine-induced NO production in rat cardiac myocytes.²⁷⁻²⁸ Mehta et al. reported that the manipulation of the renin-angiotensin system, with antisense-oligodeoxynucleotides directed at ACE mRNA, inhibits the expression of the iNOS in the myocardial ischemia model.²⁹

Recently, Schwobel et al. reported that angiotensin II receptor subtypes determine the influence of angiotensin II on the production of inducible NO in rat MCs.³⁰ They demonstrated that induced NO production is negatively controlled by the angiotensin II type 2 receptor (AT2), whereas AT1 stimulation enhanced NO synthesis in MCs. Our data shows that the up-regulation of LPS-stimulated iNOS expression was ameliorated by ACE inhibitor or AT1 blocker, at the levels of mRNA and protein, suggesting angiotensin II mediates the enhancement of iNOS expression via AT1 in diabetic glomerular injury, and are in agreement with the work of Schwobel.³⁰ To our knowledge, this is the first study providing evidence of the linkage between angiotensin II and glomerular iNOS pathway in DM nephropathy.

It is possible the balance of cNOS/iNOS is more important than the levels of cNOS, or iNOS, themselves, in the pathophysiological conditions. Recently, Kobayashi et al.³¹ and Bachetti et al.³² reported, by maintaining a correct ratio between eNOS and iNOS, ACE inhibitors are likely to protect the endothelium, and reduce the progression of atherosclerosis. Also, NO has been demonstrated to exert the capacity to autoinhibit NOS via specific binding sites on the enzyme.³³ Schwarts et al. have shown that local overproduc-

tion of NO, by iNOS, may down-regulate cNOS by autoinhibition.³⁴ From the results of our studies, taken together with previously published data showing that constitutive NO production is reduced in DM nephropathy,¹²⁻¹³ further studies are necessary to investigate the role of the cNOS/iNOS balance in the development of diabetic glomerular injury.

In summary, this study is the first report on the alteration of the iNOS pathway in diabetic glomerulopathy and the contribution of angiotensin II in the changes of the iNOS system. Based on the previous studies, showing the enhancement of the iNOS pathway may promote the accumulation of extracellular matrix in various experimental models,^{7,35,36} these results may implicate the iNOS pathway as a potential mediator of DM nephropathy.

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