

Cytokine-induced Expression of a Nitric Oxide Synthase in Rat Renal Tubule Cells

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Abstract

Nitric oxide (NO[•]) has been implicated in the regulation of renal vascular tone and tubular sodium transport. While the endothelial cell is a well known source of NO[•], recent studies suggest that tubular epithelial cells may constitutively generate NO[•]. An inducible isoform of nitric oxide synthase which produces far greater quantities of NO[•] exists in some cell types. We sought to determine whether kidney epithelial cells exposed to cytokines could express an inducible nitric oxide synthase. Primary cultures of rat proximal tubule and inner medullary collecting duct cells generated NO[•] on exposure to TNF- α and IFN- γ . NO[•] production by both cell types was inhibited by N^G-monomethyl-L-arginine; this inhibition was partially reversed by the addition of excess L-arginine. Stimulation of kidney epithelial cells with TNF- α and IFN- γ dramatically increased the level of inducible nitric oxide synthase mRNA. In summary, renal proximal tubule and inner medullary collecting duct cells can produce NO[•] via expression of an inducible isoform of nitric oxide synthase. (*J. Clin. Invest.* 1993. 91:2138–2143.) Key words: kidney • proximal tubule • collecting duct • interferon • tumor necrosis factor

Introduction

Nitric oxide (NO[•])¹ exerts a wide variety of effects on renal blood flow and function (for recent review, see [1]). NO[•] mediates, for example, bradykinin and acetylcholine-induced renal vasodilation (2, 3) and inhibits renin release (4). Furthermore, NO[•] decreases renal vascular responsiveness to vasoconstrictors (5, 6), while inhibitors of NO[•] synthesis reduce renal blood flow (7). In addition to regulating vascular tone, NO[•] affects tubule function. Endothelial cell-derived NO[•], for instance, inhibits sodium transport by cortical collecting tubule cells (8).

The source of NO[•] in the kidney has been assumed to be endothelial cells. Recent studies, however, indicate that tubule epithelial cells are capable of NO[•] synthesis (9, 10). These studies, however, have only identified NO[•] production by a constitutive form of nitric oxide synthase (NOS_c).

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1. Abbreviations used in this paper: IFN- γ , recombinant rat IFN- γ ; IMCD, inner medullary collecting duct; L-NMMA, N^G-monomethyl-L-arginine monoacetate; NO, nitric oxide; NOS_c, constitutive form of nitric oxide synthase; NOS_i, inducible nitric oxide synthase; PT, proximal tubule.

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Inducible forms of NOS have been found in macrophages, endothelial cells, hepatocytes, Kupffer cells, smooth muscle cells, fibroblasts, and tumor cells (11–21). In these cells, cytokines, immunomodulators, and lipopolysaccharide induce the expression of inducible nitric oxide synthase (NOS_i) (22). NOS_i produces NO[•] for prolonged periods and in concentrations that far exceed the amount produced by the constitutive enzyme (14). Consequently, we conducted the current study to determine if renal epithelial cells contain NOS_i. We report that both proximal tubule (PT) and inner medullary collecting duct (IMCD) cells are capable of producing NO[•] via induction of NOS_i.

Methods

Reagents

Recombinant human TNF- α was purchased from Genzyme Corp. (Cambridge, MA); recombinant rat interferon- γ (IFN- γ), Moloney leukemia virus reverse transcriptase, DMEM, Ham's F-12 media, RPMI 1640, FBS, and bovine calf serum from Gibco Laboratories (Grand Island, NY); penicillin-streptomycin solution from Irvine Scientific (Santa Ana, CA); N^G-monomethyl-L-arginine monoacetate (L-NMMA) from Chem-Biochem Research (Salt Lake City, UT); glacial acetic acid, formic acid, chloroform, and CaCl₂ from Mallinckrodt Specialty Chemicals (Paris, Kentucky); types II and IV collagenase from Worthington Diagnostic Systems (Freehold, NJ); insulin, transferrin, and selenium from Collaborative Research (Bedford, MA); RNAase A and Taq polymerase from Promega Corp. (Madison, WI); random hexamers from Boehringer Mannheim (Indianapolis, IN); dNTP from Perkin-Elmer Cetus (Norwalk, CT); guanidine isothiocyanate from Calbiochem (La Jolla, CA); acrylamide from Aldrich Chemical Co., (Milwaukee, WI); ammonium hydroxide reagent from Braun-Knecht-Heimann (Salt Lake City, UT); [³²P]dCTP from Amersham Corp. (Arlington Heights, IL); liquid scintillation cocktail from Beckman Instruments (Fullerton, CA); BCA protein assay reagent from Pierce Scientific (Rockford, IL); Primaria plates from Becton Dickinson & Co. (Lincoln Park, NJ); and 96-well assay plates from Costar Corp. (Cambridge, MA). Rabbit anti-human vWf antibody was obtained from Accurate Chemical & Scientific Co. (Westbury, NY). All other reagents, unless stated otherwise, were purchased from Sigma Chemical Co. (St. Louis, MO).

Cell isolation

Male Sprague-Dawley rats (Sasco, St. Louis, MO) weighing 150–250 g were used for all studies. The following tubule segments were studied:

(a) IMCD cells were obtained by a modification of the method of Grenier (23, 24), as previously described by this laboratory (25–27). Briefly, rat renal papillae were minced and incubated in Krebs solution (145 mM NaCl, 5 mM KCl, 1 mM Na₂HPO₄, 2.5 mM CaCl₂, 1.8 mM MgSO₄, 5 mM glucose, and 10 mM Hepes, pH 7.3) containing 0.1% type II collagenase and 0.01% type I deoxyribonuclease for 1 h at 37°C. Distilled water was then added to give an osmolality of 120 mOsm, a procedure shown to disrupt all cells except those of the collecting duct (23). The cells were washed in PBS containing 10% albumin, and suspended in modified medium K1 (see below).

(b) PT cells were isolated by a modification of the method of Vinay (28), as previously described by this laboratory (29, 30). Rat renal cor-

tex was minced and incubated in a balanced salt solution (118 mM NaCl, 4.7 mM KCl, 2.5 mM CaCl₂, 1.1 mM KH₂PO₄, 1.2 mM MgSO₄, 10 mM glucose, 20 mM Na cyclamate, 10 mM HEPES, pH 7.3) containing 1% type II collagenase, 0.01% type I deoxyribonuclease, and 0.1% soybean trypsin inhibitor for 45 min at 37°C. The suspension was strained over a 1,000- μ m pore mesh, and the tissue going through the mesh was washed three times in balanced salt solution and once in PBS with 5% BSA. The cells were centrifuged on a Percoll density gradient (50% final concentration of Percoll in balanced salt solution). The F4 fraction, corresponding to a density of 1.22, was collected, washed, and suspended in culture media (see below).

Cell culture

(a) *IMCD cells*. IMCD cells were suspended in modified media K1 (50:50 DMEM:Ham's F12 containing 5 μ g/ml insulin, 5 μ g/ml transferrin, 5 ng/ml selenium, 50 nM hydrocortisone, and 25 ng/ml PGE₁) and plated into 24-well Primaria plates. All studies were performed at confluence, ~ 4–6 d after incubation at 37°C in a 5% CO₂ environment. As expected, confluent monolayers of IMCD cells increased cAMP in response to 10 nM ADH and increased cGMP in response to 10 nM atrial natriuretic peptide.

(b) *PT cells*. PT cells were plated into 24-well Primaria plates in media consisting of 50:50 DMEM:Ham's F12 containing 5 μ g/ml insulin, 5 μ g/ml transferrin, 5 ng/ml selenium, 50 nM hydrocortisone, and 5% FBS. All studies were performed at confluence, ~ 3–5 d after incubation at 37°C in a 5% CO₂ environment. Greater than 95% of the cells stained positive for alkaline phosphatase, a marker for PT brush border. As anticipated, confluent monolayers of PT cells increased cAMP in response to 100 nM PTH, but not 1 μ M ADH.

(c) *Mesangial cells*. Mesangial cells were isolated from rats as previously described (26). Briefly, glomeruli were obtained using 60, 100, and 200 mesh sieves, incubated in 750 U/ml type IV collagenase and 0.01% type I deoxyribonuclease for 30 min at 37°C. Digested glomeruli were washed twice and cultured in RPMI 1640 containing 20% BCS and insulin (0.5 U/ml) at 37°C in a 5% CO₂ incubator. Cultures were used between passages 5 and 10.

(d) *Macrophages*. The peritoneum of Sprague-Dawley rats was lavaged with ice-cold saline. Cells were plated in RPMI 1640 + 10% FBS and vigorously washed after 1 h. Lipopolysaccharide (1 μ g/ml) was added for 12 h to half the wells. Cells were studied 48 h after plating.

Immunofluorescence

Confluent IMCD and PT cell cultures were examined for the presence of endothelial cells with anti-Factor VIII-related antigen (vWF) antibody as previously described (27). Briefly, monolayers grown on coverslips were fixed in ice-cold acetone and incubated with 20% goat serum for 10 min. The cells were then incubated with rabbit anti-human vWF antibody (1:20 dilution in 20% goat serum) for 30 min, washed in PBS, and incubated in FITC goat anti-rabbit IgG (1:10 dilution in 20% goat serum) for 30 min. After final washing in PBS for 15 min, the coverslips were mounted on glass slides and observed for fluorescence using a Zeiss fluorescent microscope. Endothelial cells scraped from the rat aorta were stained in an identical manner and served as positive controls. To exclude contamination with smooth muscle cells or mesangial cells, immunofluorescence for myosin was conducted in a similar manner as previously described (30).

Experimental protocol

Confluent cultures of PT and IMCD cells were incubated for up to 96 h at 37°C in a 5% CO₂ environment under the following conditions: (a) PT or IMCD media alone (5% FBS was added to the modified media K1 for IMCD cells); (b) 150 U/ml TNF- α + 500 U/ml IFN- γ in media (concentrations shown to induce NO[•] production in another cell line) (31); (c) 150 U/ml TNF- α + 500 U/ml IFN- γ + 0.5 mM L-NMMA; and (d) 150 U/ml TNF- α + 500 U/ml IFN- γ + 0.5 mM L-NMMA + 5 mM L-arginine. In another series of experiments, condition (b) above was repeated using 300 U/ml TNF- α .

Determination of nitrite levels. After incubation, aliquots of the supernatants from the above conditions were removed and immediately tested for nitrite (NO₂⁻) and nitrate (NO₃⁻) levels (the stable breakdown products of NO[•]) (32). 50 μ l of the conditioned medium from each well was aliquoted in duplicate into a 96-well plate. To this, 100 μ l of Greiss reagent (1% sulfanilamide in 30% acetic acid and 0.1% *N*-(1-naphthyl)ethylenediamine dihydrochloride in 60% acetic acid in a 1:1 mixture) was added at room temperature. The plates were shaken for 60 s at 80 rotations per min. The NO₂⁻ reacts with the Greiss reagent to form a chromophore (33, 34), and its absorbance at 570 nm was immediately measured in a microplate reader (MR 700; Dynatech Laboratories Inc., Alexandria, VA). The NO₂⁻ level was determined by comparing values obtained for samples with a standard curve established by adding known quantities of sodium nitrite to the media used for the experiment (31).

Determination of NO₃⁻ levels. 50 μ l of conditioned media from each well was aliquoted in duplicate into a 96-well plate. To this, 50 μ l of a mixture containing 1 M HEPES (pH 7.25), 2.4 M ammonium formate (pH 7.2), and an *Escherichia coli* homogenate with nitrate reductase activity (generous gift from Dr. John Hibbs, Jr., University of Utah) was added in a ratio of 1:1:0.05. The plate was incubated for 60 min at 37°C. The plate was then centrifuged for 10 min at 2,000 rpm at 4°C. 50 μ l of the supernatant was then aliquoted into a new 96-well plate and the NO₂⁻ level was measured as described above to determine the total NO₂⁻/NO₃⁻ level (33, 35). Standards comprised of sodium nitrite in media were put through the same procedure. The previously determined NO₂⁻ level was subtracted from the total amount of NO₂⁻/NO₃⁻ to obtain the NO₃⁻ level. After extraction in 1% lauryl sulfate, the protein level in each well was measured using the BCA protein assay reagent as previously described (36).

Isolation of RNA. RNA was obtained from IMCD and PT cells exposed for 96 h to media alone or media containing 150 U/ml TNF- α + 500 U/ml IFN- γ . Confluent monolayers were overlaid with 4 M guanidinium thiocyanate, 1% β -mercaptoethanol, and 1% sarcosyl (pH 7.0). The cells were homogenized and 1/10th volume 2 M Na acetate added. The RNA was extracted using acid phenol (pH 4.0) and chloroform, precipitated in isopropanol, and resuspended in water. Each sample was quantified spectrophotometrically.

Quantitation of RNA. 5 μ g of total RNA from each sample was reverse transcribed by incubating with 250 pmol random hexamers, 3 mM MgCl₂, 400 U murine Moloney leukemia virus reverse transcriptase, 500 μ M dNTP, 0.01 mM dithiothreitol, 75 mM KCl, and 50 mM Tris-Cl (final pH 8.3, final volume 50 μ l) for 1 h at 37°C. After incubation, 5 μ g RNAase A was added for 5 min at 37°C, followed by addition of 20 μ l 5 M NaCl and 200 μ l water. The cDNA was then extracted in phenol (pH 7.8) and chloroform, ethanol precipitated, and resuspended in water.

The cDNA was amplified by polymerase chain reaction. Each sample was measured for NOS_i and β -actin cDNA in separate tubes using specific primers. The upstream and downstream primers for NOS_i were 5'-GCATGGACCAGTATAA GGCAAGCA-3' and 5'-GCTTCTGGT-CGATGTTCATGAGCAA-3', respectively. These yielded a single band corresponding to a 222-bp fragment. The sequence for the 222-bp fragment was analyzed by Margaret Robinson in Dr. Ray White's laboratory at the University of Utah using cycle sequencing with fluorescein-labeled primer ends. Analysis of the sequence revealed that it was identical to position 1693–1915 in mouse macrophage NOS_i cDNA (37). These primers have < 40% homology with rat brain NOS_c cDNA (37, 38). Furthermore, these primers, if hybridized to rat brain NOS_c, would be predicted to yield a 348-bp product after PCR (38). PCR of rat genomic DNA with the primers for NOS_i yields a 690-bp product, indicating that this primer set spans an intron.

The upstream and downstream primers for β -actin were TGGA-GAAGAGCTATGAGCTGCCTG and GTGCCACCAGACAGCA-CTGTGTTG, respectively, which yielded a single band corresponding to a 201-bp cDNA fragment. PCR of rat genomic DNA with the β -actin primers yields a 289-bp product which is complementary to position 2499–2788 in the β -actin gene, confirming that this primer set spans an

intron. PCR was performed by incubating 5 μ l (~ 0.5 μ g) of sample cDNA with 50 mM KCl, 10 mM Tris-Cl, .01% gelatin, 1.5 mM MgCl₂, 2.5% formamide, 2 U Taq polymerase, 200 μ M dNTP, 100 pmol of β -actin or NOS_i primers, and 1 μ Ci [³²P]dCTP in 50 μ l final volume (final pH 8.3 at room temperature). PCR using β -actin primers was carried out for 20 cycles (15 s at 94°C, 15 s at 65°C, 30 s at 72°C) using a Perkin-Elmer Cetus 9600 Gene-Amp System. PCR using NOS_i primers was carried out for 30 cycles under identical conditions. NOS_i and β -actin primers were never combined in the same tube. 20 μ l of the final PCR reaction was electrophoresed using an 8% acrylamide gel in 1 \times TBE buffer. Gels were stained with 1 μ g/ml ethidium bromide, the bands corresponding to the cDNA product excised, mixed with scintillation cocktail, and counts per minute determined on a beta counter (Beckman Instruments).

NOS_i and β -actin cDNA obtained from PCR of reverse transcribed RNA were used to generate standard curves. The cDNA was amplified by PCR, the resultant amplified product divided into small fractions that were, in turn, reamplified. The purity of the final product was confirmed by electrophoresis. If a single band of the appropriate size was obtained, the final product was cleaned using Magic PCR Prep (Promega Corp.) to remove the primers. The cleaned product was again electrophoresed to confirm that it contained only the desired cDNA. If pure, the cDNA was quantitated spectrophotometrically. Standard curves for β -actin or NOS_i were made by simultaneously amplifying sample cDNA and, in separate tubes, standard cDNA (10⁻¹-10⁻⁸ ng/tube). Every PCR amplification included a standard curve. All PCR consisted of simultaneous amplification (in separate tubes) of cDNA for NOS_i and β -actin. All results are expressed as atograms NOS_i cDNA/pg β -actin cDNA to control for the amount of RNA initially reverse transcribed.

Statistical analysis. Unless otherwise stated, all data points represent the mean of values obtained from eight experiments. Data for NO₂ and NO₃ production, when involving more than two groups, were analyzed by analysis of variance. All other data, involving only two groups, were analyzed by the unpaired Student's *t* test. Statistical significance was taken as *P* < 0.05. Values are presented as mean \pm standard error of the mean.

Results

Cell culture. Confluent IMCD and PT monolayers exhibited dome formation characteristic of salt- and water-transporting epithelia. None of the cell cultures exhibited multivacuolated cells characteristic of interstitial cells. Only the typical cobblestone pattern was apparent on light microscopy; no fibroblasts were seen. No IMCD or PT cells phagocytosed serum-treated zymosan particles; in contrast, 96% of peritoneal macrophages, whether treated with lipopolysaccharide or not, took up three or more particles. Cultures were negative for myosin immunofluorescence; in contrast, cultured mesangial cells were 99% positive. Both cell types were completely negative for Factor VIII-related antigen immunofluorescence; 95% of endothelial cells were brightly positive. In summary, the IMCD and PT cells were highly purified and did not contain significant interstitial, fibroblast, macrophage, mesangial, or endothelial cell contamination.

NO^o production by inner medullary collecting duct cells. Unstimulated IMCD cells produced total NO₂/NO₃ at levels near the threshold for detection (7 \pm 3 nmol/mg protein). Addition of 150 U/ml TNF- α + 500 U/ml IFN- γ for 96 h markedly increased total NO₂/NO₃ accumulation in the supernatants (52 \pm 4 nmol/mg protein, *P* < 0.0001 vs. control) (Fig. 1). L-NMMA (0.5 mM) inhibited the stimulatory effect of TNF- α and IFN- γ (5 \pm 2 nmol/mg protein). Finally, the addition of 5 mM L-arginine partially reversed the inhibitory effect of L-

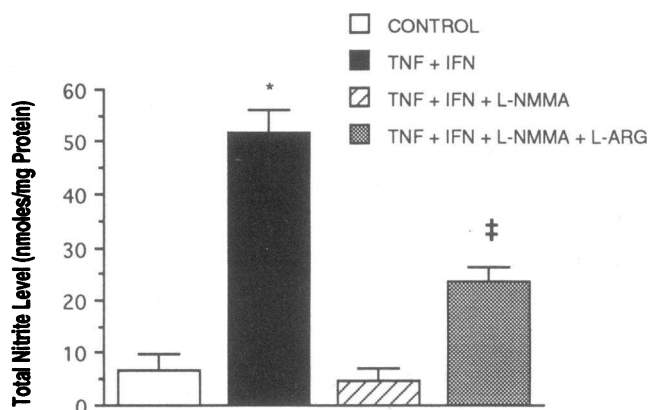


Figure 1. Total nitrite (NO₂/NO₃) production by IMCD cells. Total nitrite was measured in the supernatants of rat IMCD cells after 96 h of incubation with the reagents noted in the figure. TNF, 150 U/ml TNF- α ; IFN, 500 U/ml IFN- γ ; L-NMMA, 0.5 mM L-NMMA; L-ARG, 5 mM L-arginine (*n* = 8 in each group; **P* < 0.0001 vs control; [†]*P* < 0.001 vs control).

NMMA on TNF- α and IFN- γ -stimulated NO^o production (Fig. 1).

TNF- α augmented NO^o production in a dose-dependent manner. Raising the TNF- α dose from 150 U/ml to 300 U/ml (keeping the IFN- γ dose constant at 500 U/ml) increased NO₂/NO₃ concentration from 118 \pm 7 to 158 \pm 12 μ M/mg protein (*P* < 0.03) (Fig. 2 a). In addition, TNF- α and IFN- γ stimulated NO^o production by IMCD cells in a time-dependent manner. NO₂ was not detected in the supernatants after 48 h of exposure to these cytokines. 72 h of incubation with TNF- α and IFN- γ increased total NO₂/NO₃ to 23 \pm 5 nmol/mg protein; 96 h of exposure further augmented total NO₂/NO₃ accumulation (52 \pm 4 nmol/mg protein, *P* < 0.0006) (Fig. 2 b). At 96 h, 75% of total NO₂/NO₃ was NO₃ and 25% was NO₂.

NO^o production by proximal tubular cells. PT cells also synthesized NO^o in response to cytokine stimulation. There were no detectable breakdown products of NO^o in the supernatants of unstimulated cells, while cell exposed to 150 U/ml TNF- α and 500 U/ml IFN- γ had a total NO₂/NO₃ concentration of 164 \pm 10 nmol/mg protein (*P* < 0.0001 vs. control). Concurrent exposure to 0.5 mM L-NMMA inhibited TNF- α and IFN- γ -stimulated production of NO^o. The inhibitory effect of L-NMMA could be partially overcome by addition of 5 mM L-arginine (Fig. 3). In this cell population as well, NO^o was preferentially broken down to NO₃ (90%) rather than NO₂ (10%).

NOS_i mRNA. Traditional Northern analysis and RNAase protection assays were not sensitive enough to detect NOS_i mRNA under unstimulated conditions, hence a quantitative PCR method was used. The accuracy of the quantitative PCR used was determined as previously described by this laboratory (39). In the first control, the reproducibility of reverse transcription was examined. Three separate reverse transcriptions of the same RNA sample, followed by PCR of the resultant cDNA (using the same NOS_i and β -actin standard curves), yielded < 6% variability in measured NOS_i and β -actin cDNA; the ratio of NOS_i to β -actin cDNA varied by < 7%. In the second control, the reproducibility of quantitation of cDNA was evaluated. Three separate PCR amplifications, each using independently made standard curves, were performed on the same sample of cDNA. The calculated amount of NOS_i and

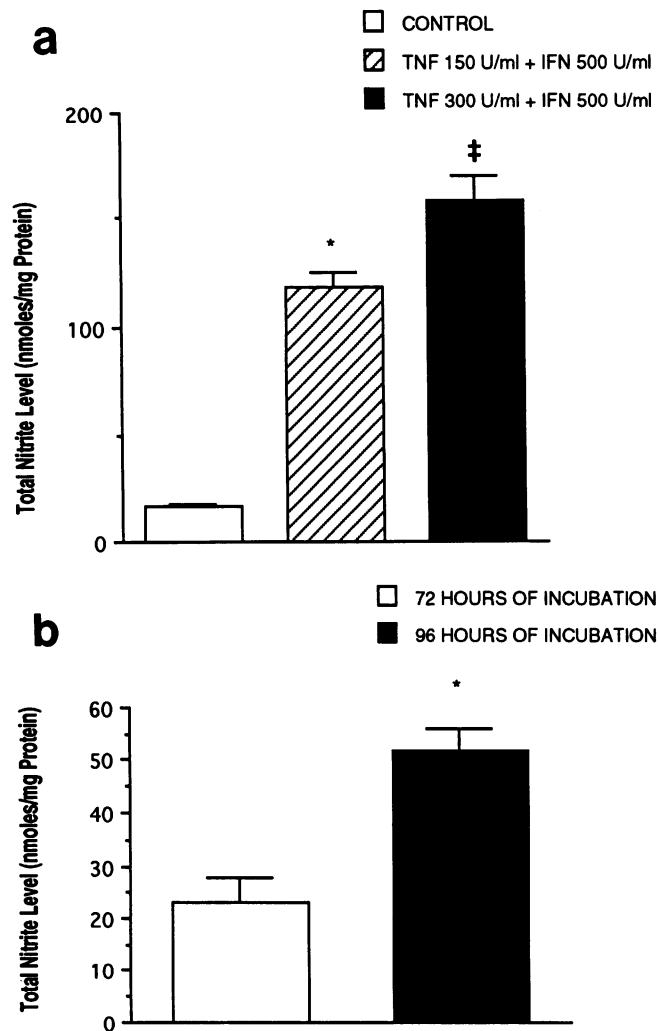


Figure 2. (a) Effect of TNF- α dose on total nitrite production ($\text{NO}_2^-/\text{NO}_3^-$) by IMCD cells. Total nitrite was measured in the supernatants of rat IMCD cells after 96 h of incubation with increasing doses of TNF- α ($n = 4$ in each group; * $P < 0.0001$ vs control; ‡ $P < 0.03$ vs 150 U/ml of TNF- α). (b) Effect of time on total nitrite production ($\text{NO}_2^-/\text{NO}_3^-$) by IMCD cells. Total nitrite was measured in the supernatants of rat IMCD cells after 72 and 96 h of incubation with 150 U/ml of TNF- α and 500 U/ml of IFN- γ ($n = 8$ in each group; * $P < 0.0006$ vs 72 h).

β -actin cDNA each varied by $< 16\%$; while the ratio of NOS_i to β -actin cDNA varied by $< 12\%$. Standard curves using 0.1–100 pg of β -actin cDNA (which encompasses all sample values) gave correlation coefficients invariably > 0.99 . Similarly, standard curves using 0.1–100 fg of NOS_i cDNA (which encompasses all sample values) gave correlation coefficients invariably > 0.97 . Finally, PCR of identically processed samples but without reverse transcriptase, or PCR in which water was substituted for cDNA, yielded no detectable product.

The results of reverse transcription and polymerase chain reaction of RNA from PT and IMCD cells are shown in Fig. 4. Unstimulated PT and IMCD cells contained extremely low levels of NOS_i mRNA (both cell types had a ratio of NOS_i/β -actin of 1:27,000). Addition of 150 U/ml TNF- α + 500 U/ml IFN- γ for 96 h resulted in a marked stimulation of PT and IMCD NOS_i mRNA (Fig. 4). The magnitude of cytokine-stimulated NOS_i mRNA appears to be greater in IMCD cells

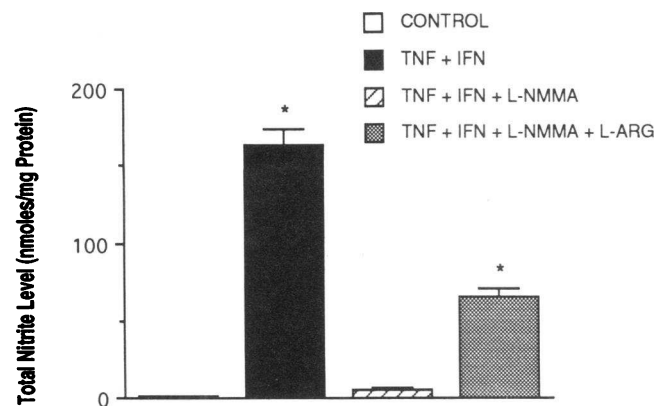


Figure 3. Total nitrite production ($\text{NO}_2^-/\text{NO}_3^-$) by PT cells. Total nitrite was measured in the supernatants of rat PT cells after 96 h of incubation with the reagents noted in the figure. TNF, 150 U/ml of TNF- α ; IFN, 500 U/ml of IFN- γ ; L-NMMA, 0.5 mM L-NMMA; L-ARG, 5 mM L-arginine ($n = 7$ or 8 for each group; * $P < 0.0001$ vs control).

than in PT cells. However, given that the levels of total $\text{NO}_2^-/\text{NO}_3^-$ appeared to be greater in the supernatants of PT than in IMCD cells, the significance of higher cytokine-induced NOS_i mRNA levels in IMCD cells is unclear. The increase in NOS_i/β -actin mRNA ratio after cytokine exposure was not due to a decrease in β -actin mRNA: PT β -actin mRNA was 38 ± 25 pg in unstimulated cells and 73 ± 34 pg in TNF- α and IFN- γ treated cells; while IMCD β -actin mRNA was $1.2 \text{ pg} \pm 0.2 \text{ pg}$ in unstimulated cells and $1.4 \pm 0.6 \text{ pg}$ in TNF- α and IFN- γ treated cells.

Discussion

This study demonstrates that PT and IMCD cells stimulated with TNF- α and IFN- γ produce large amounts of NO^* . L-NMMA, an inhibitor of both L-arginine transport and NOS activity (40), blocks NO^* production in cytokine-stimulated epithelial cells. Excess L-arginine partially reverses this inhibition. Finally, cytokine treatment of these cells markedly increases NOS_i mRNA levels, confirming that these cells can express NOS_i and may serve as a source for the increased NO^* generation caused by inflammatory mediators.

NO_2^- levels were not detectable in the supernatants of renal

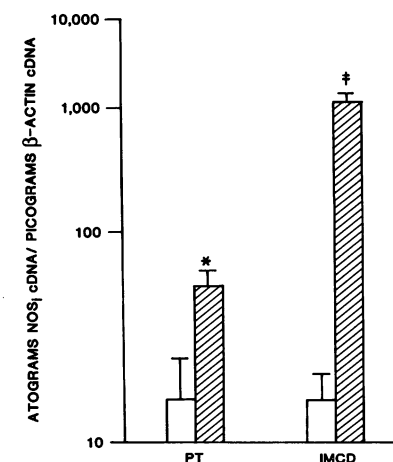


Figure 4. Inducible NOS_i mRNA levels in IMCD and PT cells. NOS_i mRNA was measured in PT and IMCD cells 96 h after incubation in media alone (open bar) and after stimulation with 150 U/ml of TNF- α and 500 U/ml of IFN- γ (cross-hatched bar). ($n = 3$ for each data point; * $P < 0.05$ vs control; ‡ $P < 0.005$ vs control).

epithelial cells exposed to TNF- α and IFN- γ for 48 h. In contrast, TNF- α induction of NOS activity in bovine endothelial cells occurs after 8 h (41). This previous study used a more sensitive assay of NO $^{\circ}$ detection; if such a method had been used in the current study, induction of renal epithelial cell NO $^{\circ}$ synthesis may have been observed at an earlier time. While delineation of the time of onset of cytokine-stimulated NO $^{\circ}$ production by renal tubule cells is of interest, the current study focused on the time required to produce micromolar amounts of NO $^{\circ}$ in the culture supernatant. This amount of NO $^{\circ}$ clearly separates constitutive from inducible production and underscores the potential physiologic differences between the two systems.

IMCD cells appeared to have a higher NOS $_i$ / β -actin mRNA ratio, however, PT cells appeared to produce more NO $^{\circ}$. Since, due to logistic considerations, IMCD and PT cells were not obtained from the same rats, direct comparison of NOS $_i$ activity between the two cell types was not possible. While precise determination of the relative activity of NOS $_i$ in nephron segments would be of interest, the present study was not designed to address this issue. Rather, two renal tubule cell types were chosen that had the greatest difference in baseline arginine synthesis: synthesis of arginine by the unstimulated rat nephron is almost entirely limited to the PT, while the IMCD produces virtually no detectable arginine (42). Despite differences in baseline arginine synthesis, PT and IMCD cells could both be induced to express NOS $_i$ and generate substantial amounts of NO $^{\circ}$. Hence, it appears that once IMCD cells are stimulated, ample arginine precursors are available. The source of this arginine has yet to be determined, but likely reflects an increase in arginine uptake and/or upregulation of arginine synthetic ability. It should be noted that the absolute level of NO $^{\circ}$ produced by a tubule cell may not solely determine the magnitude of the local response. The renal medulla, for example, has been shown to produce more cGMP in response to NO $^{\circ}$ than does the renal cortex (43). The amount of NO $^{\circ}$ produced by cytokine stimulated PT and IMCD cells is similar to that reported for IL-1-stimulated rat aortic smooth muscle cells (19), but less than can be produced by murine peritoneal macrophages which can generate $\sim 1.5 \mu\text{mol}/\text{mg}$ protein per 20 h (44).

NOS $_e$ has been demonstrated in LCC-PK $_1$, a porcine proximal tubule-like cell line (9, 10). The NO $^{\circ}$ produced by the constitutive enzyme is believed to function as an intra- and extracellular messenger by activating soluble guanylate cyclase (10). In the tubular epithelial cell, NO $^{\circ}$ has also been proposed to have a regulatory role in sodium absorption (10, 43). The inducible isoform of NOS, however, has not been previously shown to exist in kidney epithelial cells. The NOS $_i$ requires L-arginine or an L-arginine containing compound as a substrate and is dependent on NADPH, tetrahydrobiopterin, flavin adenine dinucleotide, and flavin mononucleotide; a strict requirement for calcium has not been demonstrated (11). The inducible L-arginine:NO $^{\circ}$ pathway appears to be involved in defense against intracellular microorganisms, activation of cytotoxic macrophages, and the hypotension induced by endotoxin (14, 40, 45). Whether inducible NO $^{\circ}$ production by renal tubular epithelial cells affects renal vascular tone, sodium absorption, or resistance to cancer cells or infection remains to be determined.

In addition to a possible physiologic role, inducible NO $^{\circ}$ production may play a role in renal pathophysiology. Cells stimulated to generate NO $^{\circ}$ may themselves be injured by this

reactive species because it can inhibit key iron containing enzymes involved in the citric acid cycle, mitochondrial respiration, and DNA synthesis (33). Experimental studies suggest that NO $^{\circ}$ may partly mediate immune complex induced tissue injury. Inhibiting NO $^{\circ}$ generation, for example, prevents the development in rats of immune complex alveolitis and dermal vasculitis (46). An NOS inhibitor also suppresses adjuvant induced arthritis in rats (47). Glomerular NO $^{\circ}$ production, presumably arising from macrophages, is enhanced in immune complex glomerulonephritis, raising the possibility of a role for NO $^{\circ}$ in producing or modulating the injury (48). Our results suggest that renal tubule epithelial cells may also be an important source of NO $^{\circ}$ in the setting of inflammatory renal diseases. In addition, renal tubule NO $^{\circ}$ generation may contribute to the pathophysiology of other renal diseases characterized by tubule damage, such as acute tubular necrosis.

Other epithelial cells may contain the inducible isoform of NOS. EMT-6 cells, a murine mammary adenocarcinoma, produce micromolar amounts of nitrite and nitrate after cytokine stimulation (21). In addition, cultured hepatocytes obtained from rats injected with *Corynebacterium parvum* generate $\sim 400 \text{ nmol}$ nitrate and nitrite/mg protein per 24 h. This nitrate and nitrite production were calcium independent, indicative of NOS $_i$ -mediated production of NO $^{\circ}$ (17). Finally, insulin-producing HIT cells release NO $^{\circ}$ and express NOS $_i$ mRNA after exposure to interleukin-1 (49).

In summary, we have demonstrated the presence of inducible nitric oxide synthase and inducible NO $^{\circ}$ production in proximal tubule and inner medullary collecting duct cells. To our knowledge, this is the first report of inducible NO $^{\circ}$ production in renal epithelial cells. The role of this pathway in the physiology and pathophysiology of kidney function remains to be determined.

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