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## Renal (pro)renin receptor upregulation in diabetic rats through enhanced angiotensin AT<sub>1</sub> receptor and NADPH oxidase activity

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### Abstract

Recent studies have demonstrated the presence of the (pro)renin receptor (PRR) in the glomerular mesangium and the subendothelial layer of the renal arteries. We hypothesized that diabetes upregulates PRR expression through enhanced angiotensin subtype 1 (AT<sub>1</sub>) receptor–NADPH oxidase cascade activity. Using real-time polymerase chain reaction, Western blot analysis and immunostaining, we studied renal localization of the PRR in the streptozotocin-induced diabetic rat model and in response to 1 week of treatment with the AT<sub>1</sub> receptor blocker valsartan (10 mg kg<sup>-1</sup> day<sup>-1</sup>), the angiotensin AT<sub>2</sub> receptor blocker PD123319 (0.5 mg kg<sup>-1</sup> day<sup>-1</sup>) or the NADPH oxidase inhibitor diphenylene iodonium (DPI; 0.5 mg kg<sup>-1</sup> day<sup>-1</sup>) 6 weeks post-induction of diabetes. Both PRR mRNA and protein were expressed constitutively in the kidneys of normal rat renal cortex and medulla, mainly in glomerular mesangium, proximal, distal and collecting tubules. Compared with normal rats (100%), diabetic rats demonstrated an increase in renal PRR mRNA (184%), protein (228%) and immunostaining. Valsartan and DPI prevented the increase in the PRR mRNA (106 and 126%, respectively), protein (97 and 140%, respectively) and immunostaining that was seen in the kidneys of diabetic rats. The AT<sub>2</sub> blocker PD123319 did not have significant effects on PRR mRNA (157%) or protein expression (200%) in the kidneys of diabetic rats. These results demonstrate that the PRR is constitutively expressed in renal glomeruli and tubules. Expression of the PRR is upregulated in diabetes via enhancement of AT<sub>1</sub> receptor–NADPH oxidase activity.

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All components of the renin–angiotensin system (RAS) are present within the kidney (Siragy, 2006). Previous studies have demonstrated the contribution of this system to the development of kidney disease in diabetes (Andersen *et al.* 2000; Chan *et al.* 2000; Brenner *et al.* 2001; Lewis *et al.*, 2001; Carey & Siragy, 2003). However, the mechanisms involved in the pathophysiology of this clinically relevant problem are still not well elucidated.

Recently, a (pro)renin receptor (PRR) was discovered as a new component of the RAS (Nguyen *et al.* 2002). This receptor has been associated with prorenin/renin uptake, non-proteolytic activation of prorenin and local production of angiotensin I (Nguyen *et al.* 2002; Nabi *et al.* 2006; Nguyen, 2007). The PRR was reported to be expressed in the glomerular mesangial cells and subendothelial space of the renal vasculature (Nguyen *et al.* 2002) but not in renal tubules.

Angiotensinogen and angiotensin-converting enzyme are present in the renal tubular cells (Tang *et al.* 1995; Imig *et al.* 1999; Ingelfinger *et al.* 1999; Harrison-Bernard *et al.* 2002). The presence of PRR in renal tubules may contribute to tubular generation of angiotensin II through enhanced conversion of locally produced angiotensinogen to angiotensin I (Kobori *et al.* 2007). Angiotensin subtype 1 receptors (AT<sub>1</sub> receptors) are also present in renal glomeruli and tubules (Siragy, 2006) and in close proximity to sites of angiotensin II formation. Excess of

AT<sub>1</sub> receptors and oxidative stress activities contribute to development of renal and cardiac diseases (Onozato *et al.* 2002; Privratsky *et al.* 2003).

A recent study demonstrated that PRR blockade improves kidney disease in diabetes (Ichihara *et al.* 2006). However, the interaction between diabetes and renal expression of the PRR is not known. Similarly, it is not known whether oxidative stress associated with diabetes influences PRR expression. It is plausible that both AT<sub>1</sub> receptors and oxidative stress lead to increased expression of the PRR in different renal regions, such as glomeruli and tubules, a process that could further enhance the development of renal disease in diabetes. In the present study, we hypothesized that in addition to glomerular mesangium and renal vasculature, PRR is expressed in the renal tubules and is upregulated in diabetes via enhanced angiotensin AT<sub>1</sub> receptor and oxidative stress activities. Our results confirm that PRR is expressed not only in glomerular mesangium but also in renal tubules. The PRR expression is upregulated in diabetes, a process that can be suppressed by angiotensin AT<sub>1</sub> receptor blockade and inhibition of NADPH oxidase activity.

## Methods

### Animal preparation and treatment

The experiments were approved by the University of Virginia Animal Research Committee and conducted in accordance with institutional guidelines. Sprague–Dawley rats were divided randomly into control and diabetic groups. Diabetes was induced by streptozotocin (STZ; 65 mg/kg i.p.). Studies were conducted for 6 weeks. At the end of the fifth week, an osmotic minipump filled with vehicle (0.9% NaCl; control group,  $n = 8$ ), valsartan (10 mg kg<sup>-1</sup> day<sup>-1</sup>; diabetes + valsartan group,  $n = 8$ ), PD123319 (0.5 mg kg<sup>-1</sup> day<sup>-1</sup>; diabetes + PD123319 group,  $n = 8$ ) or diphenylene iodonium (DPI; 0.5 mg kg<sup>-1</sup> day<sup>-1</sup>; diabetes + DPI group,  $n = 8$ ) was implanted into the peritoneal cavities of the studied rats. At the end of the sixth week of the study, rats' body weight was recorded and blood samples were collected. The kidneys were removed under anesthesia (Ketamine 60 mg kg<sup>-1</sup>, I.P.) and stored in -80°C for quantitative real-time reverse transcriptase-polymerase chain reaction (RT-PCR), Western blot analysis and immunostaining.

### Quantitative real-time RT-PCR

After removal of the kidneys, the renal tissue was weighed promptly and homogenized on ice. The total renal RNA was extracted using RNeasy Kit (Qiagen, Valencia, CA, USA). The quality of RNA was confirmed by ethidium bromide staining in 1% formaldehyde agarose gel. Single-stranded cDNA was synthesized using iScript cDNA Synthesis Kit (Bio-Rad, Hercules, CA, USA). Gene-specific primers were as follows: for (pro)renin receptor, forward sequence 5'-GAGGCAGTGACCCTCAACAT-3' and reverse sequence 5'-CCCTCCTCACACAACAAGGT-3'; and for  $\beta$ -actin, forward sequence 5'-AGCCATGTACGTAGCCATCC-3' and reverse sequence 5'-ACCCTCATAGATGGGCACAG-3'. The specificities of the primers were verified by melting curves (iCycler; Bio-Rad) and amplified product size using agarose gel electrophoresis. Quantitative real-time RT-PCR was performed using iCycler (Bio-Rad), and threshold cycle number was determined using iCycler software version 3.0 (Bio-Rad). Reactions were performed in triplicate, and threshold cycle numbers were averaged. Non-template control was used as negative control. The mRNA results were normalized to  $\beta$ -actin mRNA. Fold-downexpression or -upexpression was calculated according to the formula  $2^{(Rt - Et) / 2^{(Rn - En)}}$ , where  $Rt$  is the threshold cycle number for the reference gene observed in the test sample,  $Et$  is the threshold cycle number for the experimental gene observed in the test sample,  $Rn$  is the threshold cycle number for the reference gene observed in the control sample and  $En$  is the threshold cycle number for the experimental gene observed in the test sample.

## Western blot analysis

Kidney tissue ( $n = 8$ ) was homogenized in lysis buffer [50 mM Tris-HCl (pH 8.0), 150 mM NaCl, 2 mM EDTA, 0.1% sodium dodecyl sulphate, 1% IGEPAL CA-630, 0.5% deoxycholate sodium, 20  $\mu$ M MG132 (CalBiochem, La Jolla, CA, USA), 50 mM sodium fluoride (NaF), 2 mM sodium orthovanadate, 1 mM phenylmethylsulphonic fluoride (PMSF) and 1  $\times$  dilution protease inhibitor cocktail (Roche Applied Science, Indianapolis, IN, USA)] to form a crude homogenate. Crude lysates were cleared by centrifugation at 22 000 g, 4°C for 20 min. Protein quantification of whole cell lysate was performed using the BCA Protein Assay Kit (Pierce Biotechnology, Rockford, IL, USA). A total of 50  $\mu$ g of cell lysate was loaded for each sample and separated on a 4–20% Tris-HCl Criterion precast gel (Bio-Rad, Hercules, CA, USA), followed by blotting of the proteins on polyvinylidene fluoride membrane (Bio-Rad). Antibody to PRR (Abcam, Cambridge, MA, USA) was used in the Western blot (Kaneshiro, 2007). Signal detection was carried out by using Super Signal West Pico Chemiluminescent Substrate (Pierce Biotechnology, Rockford, IL, USA). The blot was treated with Restore Western Blot Stripping Buffer (Pierce Biotechnology) according to the manufacturer's recommendation, followed by reprobing with a monoclonal antibody against  $\beta$ -actin (Sigma, St Louis, MO, USA). Densitometry of the bands was done using Image Master™ TotalLab version 2.0 (Amersham, Piscataway, NJ, USA). The band density of PRR was normalized to the corresponding density of  $\beta$ -actin. The arbitrary unit of band densities was represented as the expression level of PRR.

## Immunostaining for (pro)renin receptor

Immunostaining was performed for PRR localization in the kidney. Briefly, the frozen kidneys were taken out from  $-80^{\circ}\text{C}$  storage and placed into a  $-18^{\circ}\text{C}$  cryostat chamber for 1 h. Each tissue sample was mounted with OCT (an embedding medium) on a microtome block with designed cutting orientation. The block was held firmly in the cryostat microtome and 2–4- $\mu\text{m}$ -thick sections were cut. Sections were then picked up on histological slides and immediately dipped into Streck Tissue Fixative (NE68128; Streck Laboratories, La Vista, NE, USA) at  $-4^{\circ}\text{C}$  for 15 min, transferred into phosphate-buffered saline at  $-4^{\circ}\text{C}$  for 30 min, and preabsorbed in 5% goat serum for 20 min before the immunostaining process. The immunostaining was performed by incubating with polyclonal rabbit anti-(pro)renin receptor antibody at  $4^{\circ}\text{C}$  overnight, followed by 1 h of incubation with secondary antibody conjugated with biotin at room temperature (Sigma). Immunoreactive signal was detected with an avidin–biotin immunoperoxidase reaction (Sigma) and visualized by exposure to diaminobenzidine (Sigma). The non-specific binding was controlled by using the mouse immunoglobulin G isotype (Sigma) as a primary antibody for negative control.

## Statistical analysis

Comparisons among different treatment groups were examined by ANOVA, including a repeated-measures term, using the general linear models procedure of the Statistical Analysis System (SAS Software, SAS Institute, Cary, NC, USA). Data are expressed as means  $\pm$  S.E.M.  $P < 0.05$  was considered statistically significant.

## Results

### Body weight and blood glucose levels

After 6 weeks of STZ-induced diabetes, body weights of rats in the diabetes groups ( $227.4 \pm 21.37$  g) were significantly less ( $P < 0.01$ ) compared with normal control rats ( $392.3 \pm 16.31$  g). There were no significant differences in body weight among different treatments of the diabetes groups: the AT<sub>1</sub> receptor blocker valsartan ( $230.9 \pm 17.28$  g); the AT<sub>2</sub> receptor blocker PD123319 ( $219.8 \pm 23.78$  g); and the NADPH oxidase inhibitor DPI ( $231.6 \pm 17.34$  g).

The blood glucose levels were  $69.4 \pm 19.3$  mg dl<sup>-1</sup> in normal rats,  $323.3 \pm 134.3$  mg dl<sup>-1</sup> in diabetic rats ( $P < 0.01$ ),  $312.2 \pm 109.8$  mg dl<sup>-1</sup> in diabetic rats treated with valsartan ( $P < 0.01$ ),  $331.9 \pm 124.2$  mg dl<sup>-1</sup> in diabetic rats treated with PD123319 ( $P < 0.01$ ) and  $309.1 \pm 98.39$  mg dl<sup>-1</sup> in diabetic rats treated with DPI. There were no significant differences in blood glucose levels among diabetic rats treated with valsartan, PD123319 or DPI.

### Expression of PRR in the kidney of diabetic rats

The expression of PRR mRNA (Fig. 1A) and protein (Fig. 1B) was significantly higher in diabetic compared with normal rats. Valsartan and DPI significantly decreased PRR mRNA and protein expressions. Treatment with PD123319 had no significant effect on PRR mRNA or protein expressions in diabetic rats.

### Effect of valsartan, PD123319 and DPI on glomerular and tubular expression of PRR

Figure 2 shows a representative immunostaining for PRR in renal cortex and medulla. In normal animals, PRR immunostaining showed a distribution pattern in glomeruli and renal tubules in cortex and medulla. Compared with normal rats, immunostaining for PRR increased significantly in glomeruli and tubules of the diabetic animals. The PRR immunostaining was significantly decreased in the renal cortex and medulla of diabetic rats treated with valsartan or DPI compared with untreated diabetic animals. Treatment of diabetic rats with PD123319 did not influence PRR immunostaining and was not different from PRR immunostaining in untreated diabetic rats.

## Discussion

Our studies demonstrated that in basal conditions, PRR is expressed in renal glomeruli and tubules. This finding was confirmed by RT-PCR, Western blot analysis and immunostaining. The localization of these receptors to specific kidney sites may suggest their involvement in physiological regulation of renal haemodynamic and excretory functions and possible contribution to the development of renal diseases. At the present time, since there are no available PRR knockout animal models, the exact functions of these receptors remain largely unknown. Inhibitors PRR, although not commercially available, can be constructed (Ichihara *et al.* 2006). Recent studies reported the contribution of the PRR to the conversion of angiotensinogen to angiotensin I at the cell surfaces which express these receptors (Nguyen *et al.* 2002, 2007; Nabi *et al.* 2006). This solid-state formation of angiotensin I seems to be more efficient than the process of its formation in the circulation (Danser & Deinum, 2005).

Our data also show an upregulation of PRR in the STZ-induced diabetic rat model. Expression of PRR mRNA and protein was significantly increased 6 weeks after development of diabetes. The renal upregulation of PRR in diabetes suggests that this receptor may play a role in the development of renal complications of this disease. Using a decoy peptide to block the PRR in a diabetic mouse model led to significant reduction in the development of glomerulosclerosis (Ichihara *et al.* 2006). In the present study, we could not conclude whether the increased blood glucose or the decrease in insulin availability was responsible for the increased expression of the PRR in our STZ-induced diabetic rat model.

Our study demonstrated that AT<sub>1</sub> receptor blockade with valsartan significantly inhibits the renal expressions of PRR mRNA and protein in the diabetic rats. The exact mechanisms responsible for the reduction of renal PRR expression by this treatment are unknown at the present time. The AT<sub>1</sub> receptor regulates renin production through what is known as the short-loop negative feedback mechanism, with decreased renin production during stimulation of this receptor (Kurtz & Wagner, 1999). Thus, it is possible that the increased production of prorenin/renin during AT<sub>1</sub> receptor blockade contributes to the downregulation of PRR expression.

Another possibility that could contribute to upregulation of the PRR is related to increased angiotensin II production and expression of AT<sub>1</sub> receptors in diabetes (Siragy *et al.* 2003; Sodhi *et al.* 2003), leading to enhanced oxidative stress by increasing NADPH oxidase activity (Privratsky *et al.* 2003; Wilcox, 2003). Blockade of AT<sub>1</sub> receptors reduces oxidative stress and superoxide production in the kidney of STZ-induced diabetic rats (Onozato *et al.* 2002; Onozato & Toji, 2005). Our study clearly demonstrated that blockade of NADPH oxidase activity led to downregulation of PRR expression in the kidney. In contrast to the AT<sub>1</sub> receptor, AT<sub>2</sub> receptor blockade does not seem to influence PRR expression in our diabetic rat model and could be related to a reduction in expression of this receptor in diabetes (Wehbi *et al.* 2001).

In conclusion, this study demonstrated that PRR is constitutively expressed in renal glomeruli and tubules. In diabetes, PRR expression is upregulated through enhancement of AT<sub>1</sub> receptor and NADPH oxidase activities.

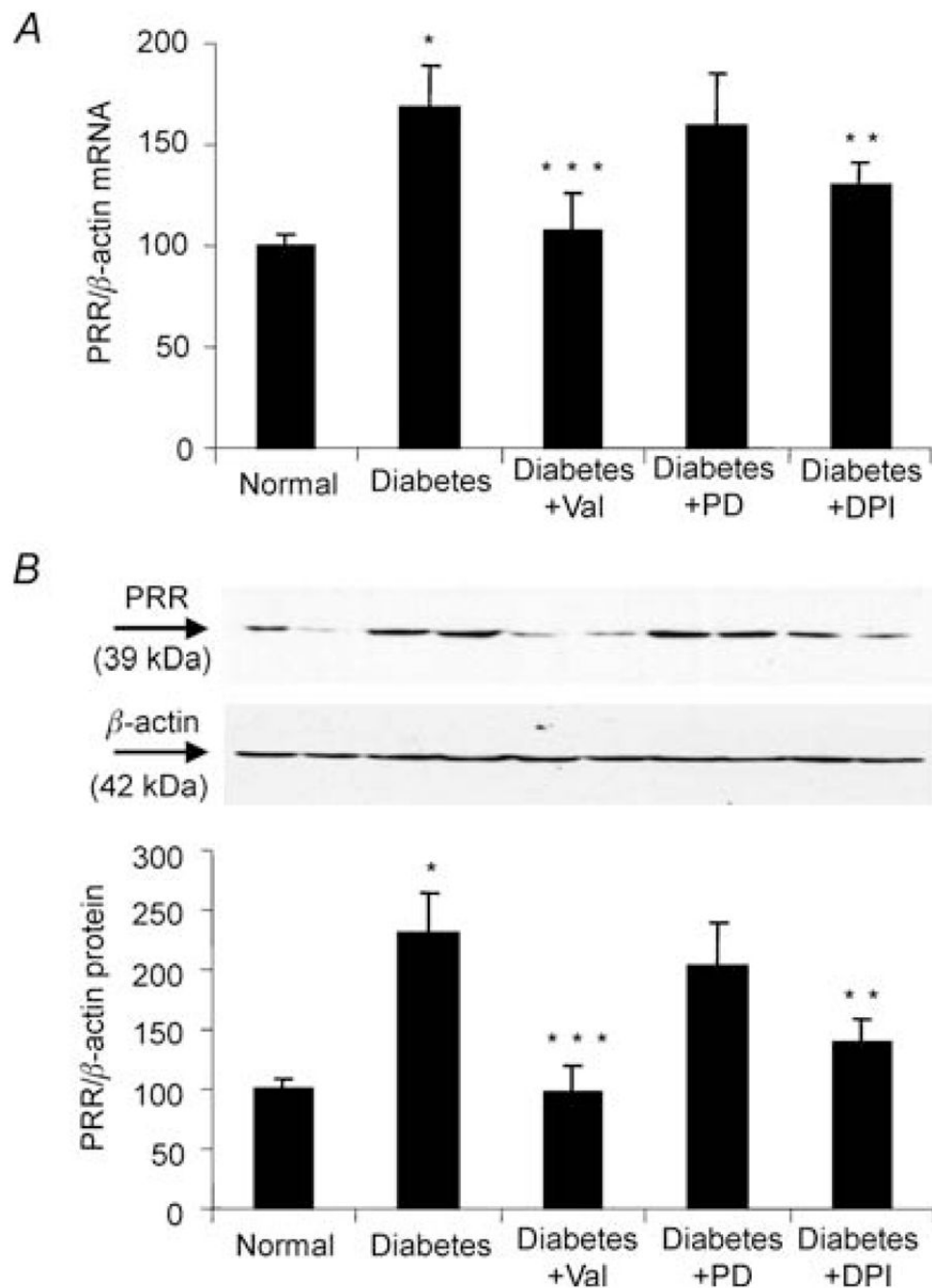
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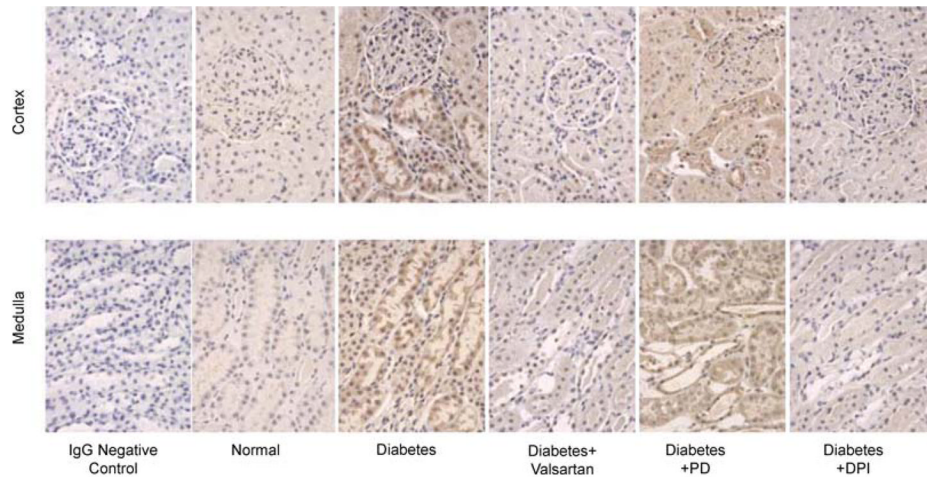
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**Figure 1. Expressions of renal PRR mRNA (A) and protein (B) in normal and diabetic rats and in response to AT<sub>1</sub> receptor blockade with valsartan (Val), AT<sub>2</sub> receptor blockade with PD123319 (PD) or NADPH oxidase inhibition with DPI**

Data represent two rats from each group, while the total number of animals studied is 8 per group. Lower panel represents densitometric analysis of the PRR Western blots ( $n = 8$  each group). \* $P < 0.05$  versus normal rats; \*\* $P < 0.05$  and \*\*\* $P < 0.01$  versus untreated diabetic rats.



**Figure 2. Representative PRR immunostaining**

Upper panel represents rat renal cortex and lower panel represents renal medulla. Brown color indicates PRR immunostaining in glomerular mesangium and tubules. Diabetes increased PRR immunostaining in the renal cortex and medulla, while valsartan and DPI treatments were associated with reduced immunostaining of this receptor in diabetic rats. There were no significant changes in PRR immunostaining in response to PD123319 (PD) treatment.