

Up-regulation of TGF- β via the activation of extracellular signal-regulated kinase 1 and 2 induced by prorenin in human renal mesangial cells

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Abstract. Prorenin is thought to be an inactive precursor of renin. This study investigated whether human prorenin was capable of activating the (pro)renin receptors [(P)RRs], leading to the phosphorylation of extracellular signal-regulated kinase 1 and 2 (ERK1/2) in cultured human renal mesangial cells (HRMCs). HRMCs cultured *in vitro* were pretreated with an AT₁ and AT₂ blocker prior to stimulation by prorenin, PD98059 (an inhibitor of ERK1/2) and handle-region peptide (HRP). Phosphorylated ERK1/2 was evaluated using Western blot analysis, and the concentration of TGF- β was measured by ELISA. The mRNA of TGF- β was evaluated by RT-PCR. It was found that prorenin activated the (P)RR in cultured HRMCs, which in turn increased p-ERK1/2. Prorenin induced rapid phosphorylation of ERK1/2 and increased p-ERK1/2 in a time- and dose-dependent manner. The protein levels of TGF- β increased significantly with the stimulation of prorenin. PD98059 significantly decreased p-ERK1/2 and then downregulated TGF- β . HRP did not inhibit either ERK1/2 phosphorylation or the increase in TGF- β . ERK1/2 phosphorylation induced by prorenin led to a marked increase in TGF- β . The regulation of TGF- β was highly dependent on ERK1/2. Thus, ERK1/2 may play a key role in the development of kidney disease. HRP neither affects the ERK1/2 signaling nor the level of TGF- β in HRMCs.

Introduction

It is traditionally thought that the biologically active peptide angiotensin II plays a key role in the development of diseases

associated with renin-angiotensin system (RAS) activation. This hypothesis was supported by Nguyen *et al* (1), who first reported (pro)renin receptors [(P)RRs] in 2002. Two (P)RRs have been identified thus far. The first, the mannose-6-phosphate (M6P) receptor, is a clearance receptor. Studies showed that it binds and internalizes renin and prorenin (2). Such binding, however, does not result in intracellular angiotensin generation, and it has been concluded that the M6P/IGF2R is a clearance receptor for prorenin (3). The specific (P)RR is a 350-amino acid protein with a single transmembrane domain that binds renin and prorenin equally well (4,5). It is highly conserved across species and has no homology with other known proteins (6). Nguyen *et al* first detected the (P)RR in renal mesangial cells and in vascular smooth muscle cells of the coronary and renal arteries using an immunofluorescence method (1). The (P)RR is highly expressed in cardiovascular, kidney, brain and placenta tissue (7). (P)RR activation activates certain signal transduction pathways and upregulates the expression of certain genes as well as enhancing the enzymatic activity of prorenin (8). (P)RR has become the focus of much research since it was recognized as being associated with diseases involving RAS activation.

Suzuki *et al* developed a decoy peptide according to the epitope of the prorenin prosegment, which was termed the handle-region peptide (HRP) (9). The HRP ostensibly blocks the binding of prorenin to the (P)RR, but there is no adequate evidence to support the hypothesis that the HRP is the blocker of the (P)RR. In this study, we aimed to demonstrate whether the HRP inhibits (P)RR in HRMCs.

The (P)RR is highly conserved across species (10). Several studies carried out on genetic model animals have shown that the (P)RR plays a pivotal role in the processes of myocardial fibrosis and glomerular sclerosis (11-14). However, no evidence of the (P)RR pathological mechanism in humans exists. Proliferation of human renal mesangial cells HRMCs is a common indicator of hypertension nephropathy and diabetic nephropathy. The proliferation of HRMCs and the cytokines produced by them play key roles in hypertensive nephropathy and diabetic nephropathy. The purpose of this study was to investigate the effect of the (P)RR on the activation of ERK1/2 and the fibrotic factor TGF- β on HRMCs. Furthermore, the study aimed to evaluate whether the putative (P)RR blocker HRP inhibits this pathway in HRMCs.

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Materials and methods

Cell culture. The HRMCs and modified Eagle's medium (MEM) were purchased from ScienCell Research Laboratories (San Diego, CA, USA). The cells were incubated in MEM supplemented with 10% bovine calf serum (BCS) in a humidified atmosphere of 95% air plus 5% CO₂. When the cells reached approximately 80% confluence, they were digested by 0.025% trypsinase at 37°C for 2 min. Medium that contained 10% fetal bovine serum was used to terminate the reaction. The third to fifth generations were used to carry out the study, and the cells were cultured in serum-deprived medium for 24 h prior to the experiment.

Reverse transcription-polymerase chain reaction (RT-PCR) to detect the TGF- β mRNA. Total RNA was extracted from HRMCs using TRIzol reagent (Invitrogen, Carlsbad, CA, USA). The total RNA was solubilized in RNase-free water and quantified by measuring the optical density (OD) at 260 nm. The RNA purity was determined by obtaining a 260/280 nm OD ratio of >1.70. A sample of total RNA (2 μ g) was reverse transcribed into cDNA in a 20 μ l reaction mixture. Primers for TGF- β (forward, 5'-GCATGGAGTCTGTGGCAT-3'; reverse, 5'-CTAGAAGCATTTGCGGTGG-3') and β -actin (forward, 5'-CCGCAAGGACCTCGGCTGGAA-3'; reverse, 5'-GATCATGTTGGACAGCGCTC-3') were designed based on previously reported cDNA sequences. A primer designer (Primer Premier 5.0 software) was used to design primers for TGF- β . The primers were synthesized by Takara, Japan. The 25 μ l PCR solution contained 1.5 mM MgCl₂, 0.2 mM deoxynucleotide triphosphates and 1 unit of Taq DNA polymerase (Gibco-BLR, Gaithersburg, MD, USA). The PCR program was as follows: 94°C for 2 min for the initial step; 94°C for 45 sec, 58°C for 45 sec and 72°C for 1 min for 20 cycles; and a final extension at 72°C for 10 min. Following amplification, the reaction products were electrophoresed on 1% agarose TAE gels and then EB-stained. TGF- β mRNA was normalized with the housekeeping gene β -actin mRNA and was analyzed using the UVP Bioimaging System (LabWorks™, Version 4.6, USA).

Western blot analysis of ERK1/2. Western blot analyses were performed as previously reported. Briefly, the cultured HRMCs were lysed in lysis buffer containing Tris (50 mM), NaCl (100 mM), NAF (50 mM), EDTA (1 mM), 0.1% sodium dodecyl sulfate (SDS), 0.5% deoxycholic acid sodium salt, 1% Triton X-100 and protease inhibitor (1 tab/9 ml buffer). Following centrifugation at 15,000 rpm for 15 min at 4°C, the supernatant was collected and subjected to SDS polyacrylamide gel electrophoresis. The proteins were transferred to polyvinylidene difluoride membranes. After blocking the blots for 3 h at room temperature with TBS-T containing 5% bovine serum albumin and 0.5% Tween-20, they were incubated for 24 h with rabbit monoclonal anti-phosphorylated ERK antibody (1:200 dilutions, Santa Cruz Biotechnology, Japan). Immunoreactivity was detected by horseradish-peroxidase-conjugated goat anti-rabbit secondary antibody. Protein bands were detected with an ECL detection kit (Pierce, Rockford, IL, USA) and recorded on X-ray film. The band densities were quantified

by a LabWorks system. The change in ERK1/2 phosphorylation was reflected by the relative abundance of p-ERK1/2 to that of ERK1/2.

Statistical analysis. Standard statistical methods from the SPSS Statistical Analysis System V-16.0 (SPSS, Chicago, IL, USA) were used. Statistical comparisons were made by using a two-way ANOVA test. Values of $P < 0.05$ were considered to indicate statistical significance. The data are reported as the means \pm SEM. Duplicate wells were analyzed for each experiment, and each experiment was performed independently at least three times.

Results

The impact of various concentrations of prorenin on phosphorylated ERK1/2. To investigate the impact of various concentrations of prorenin respectively on the activation of ERK1/2, the cells were treated with 5×10^{-9} , 5×10^{-10} , 5×10^{-11} , 5×10^{-12} or 5×10^{-13} M prorenin. It was found that phosphorylation of ERK1/2 is concentration-dependent. p-ERK1/2 increased at 5×10^{-10} M and markedly increased at 5×10^{-9} M. Thus, 5×10^{-9} M prorenin was used in the subsequent experiments (Fig. 1).

Effect of prorenin on ERK1/2 activation induced by (P)RR at various time points. Cells were harvested at 0, 2.5, 5, 10, 20, 35 or 60 min following the administration of prorenin at 5×10^{-9} M. p-ERK was measured using the Western blot method. No difference was found between 0 and 2.5 min, and although p-ERK increased 5 min after the addition of prorenin, there was no disparity after 20 min of stimulation. Therefore, human recombinant prorenin induced long-lasting phosphorylation of ERK1/2 by activating (P)RR in cultured HRMCs (Fig. 2).

Effects of HRP and PD98059 on ERK1/2 activation induced by human recombinant prorenin in HRMCs. To evaluate whether HRP and prorenin induced the phosphorylation of ERK1/2 in cultured HRMCs, angiotensin II type 1 and 2 receptors were blocked using olmesartan and PD123319, respectively, at 5×10^{-5} M. The same concentration of prorenin was used as mentioned previously. PD98059 (an inhibitor of ERK1/2) was used to inhibit the phosphorylation of ERK1/2, and 10 μ M HRP was added. The cells were harvested following a 35 min incubation. The protein levels of p-ERK increased, in addition to the activation of (P)RR induced by prorenin. PD98059 inhibited the increase of p-ERK1/2, but the HRP did not have the same effect (Fig. 3).

Effects of the HRP and PD98059 on TGF- β secretion induced by human recombinant prorenin in HRMCs. To demonstrate the role of prorenin and HRP on the levels of TGF- β , the protein concentration of TGF- β in the medium of cultured cells was measured using ELISA following stimulation with these factors. The bar chart shows that TGF- β markedly increased compared with the control when treated with prorenin. PD98059 was not markedly different from the control. The HRP again failed to reduce the protein levels of TGF- β induced by the (P)RR (Fig. 4).

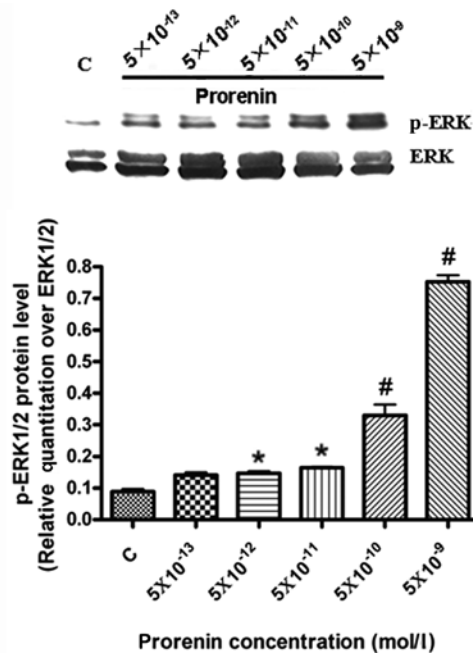


Figure 1. The activation of p-ERK1/2 following stimulation by prorenin at various concentrations. The bar chart shows the relative density of phosphorylated p-ERK1/2 to ERK1/2. * $P < 0.05$ vs. control, # $P < 0.01$ vs. control (C, control; n=3).

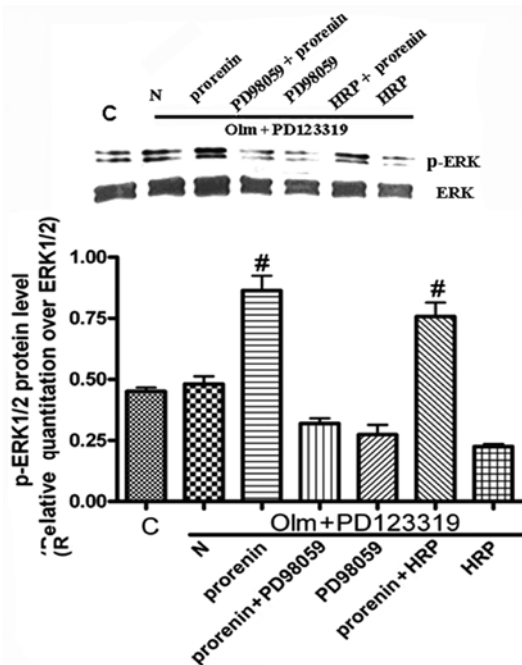


Figure 3. The effect of prorenin, HRP and the inhibitor of ERK1/2 on p-ERK1/2 in HRMCs. HRMCs were pretreated with the AT1 blocker olmesartan (olm) and the AT2 blocker PD123319 at 5×10^{-5} mol/l and then co-incubated with other reagents. A Western blot analysis was performed. The graph shows the relative levels of p-ERK1/2 compared with ERK1/2 in each group (C, control; N, no additive; n=2; # $P < 0.01$).

Effects of the HRP and PD98059 on TGF- β mRNA induced by human recombinant prorenin in HRMCs. To study whether prorenin and the HRP affect the TGF- β mRNA levels, the mRNA of TGF- β in the harvested cultured cells

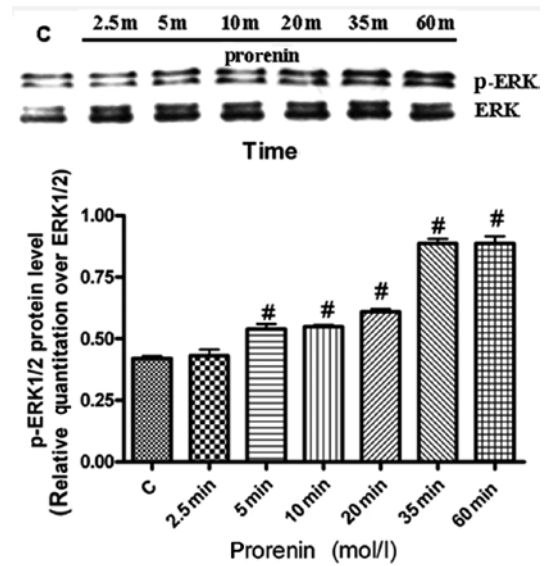


Figure 2. The time course of p-ERK1/2 following the addition of human prorenin at 5×10^{-9} M to HRMCs. Western blot analyses were carried out. The bar chart of the band shows the density of phosphorylated p-ERK1/2 to ERK1/2 bands. # $P < 0.01$ vs. control (m, min; C, control; n=3).

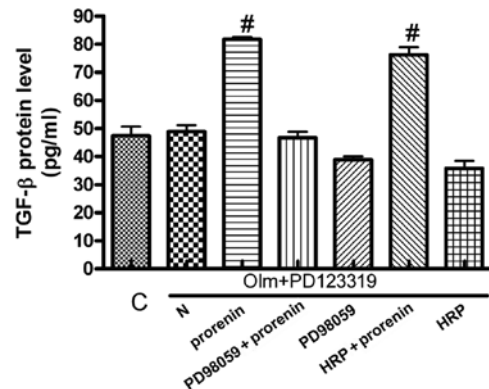


Figure 4. The effects of prorenin, HRP and inhibitor of ERK1/2 on TGF- β production in HRMCs. HRMCs were pretreated as mentioned previously. The graph shows the relative levels of TGF- β evaluated by ELISA in each group. The value is expressed relative to no additive (N) and control (C). # $P < 0.01$ compared with the control (n=4).

was measured with the RT-PCR method following stimulation with prorenin and HRP. We found that the TGF- β mRNA levels markedly increased when compared with the control. PD98059, but not HRP, inhibited the TGF- β mRNA expression induced by the (P)RR. The protein level of TGF- β was consistent with its mRNA level. This result indicates that the regulation of TGF- β is highly dependent on ERK1/2 signaling transduction (Fig. 5).

Discussion

Target-organ protection from hypertension is a controversial issue in clinical research. Studies have shown that the early stage of hypertensive nephropathy includes glomerular hypertrophy and cell proliferation; however, most of the proliferative cells are renal mesangial cells. HRMCs cultured

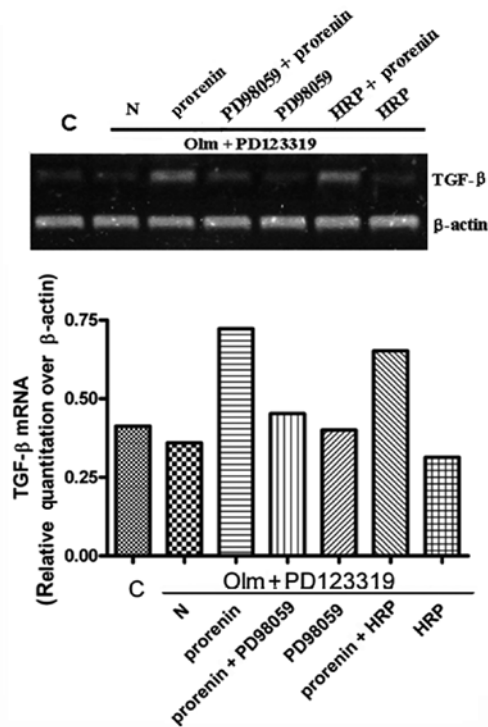


Figure 5. The effects of prorenin, HRP and the inhibitor of ERK1/2 on TGF- β mRNA expression in HRMCs. HRMCs were pretreated. RT-PCR was performed. The graph shows the relative levels of TGF- β mRNA compared with β -actin in each group ($n=1$).

in vitro were treated with various concentrations of human recombinant prorenin. The results demonstrated that prorenin rapidly induced ERK1/2 phosphorylation of MAPKs in a dose-dependent manner. p-ERK1/2 was measured at various time points after the administration of prorenin; the p-ERK increased 5 min following the addition of prorenin. No disparity occurred after 20 min. A previous study reported that phosphorylation of ERK1/2 may be involved in the development of nephropathy (15). Therefore, the study shows that the mechanism through which the (P)RR induced the rapid phosphorylation of ERK1/2 is involved in the development of nephropathy.

There are two pathways for the phosphorylation of ERK1/2 induced by prorenin. First, it is well known that prorenin-renin conversion occurs in the kidney (16). Prorenin, following its local conversion to renin, contributes to angiotensin (Ang) generation. ERK1/2 is activated when angiotensin II binds to its type 1 receptor. Second, ERK1/2 activation by (P)RR is independent of angiotensin II generation (8,17,18). To evaluate the function of (P)RR, angiotensin II type 1 and 2 receptors were blocked with olmesartan and PD123319. The cells were then incubated with prorenin. The results demonstrated that p-ERK1/2 markedly increased from the stimulation of prorenin and decreased sharply when co-incubated with the ERK1/2 inhibitor PD98059. Since AT₁ and AT₂ were blocked, it appears that the phosphorylation of ERK1/2 was due to (P)RR activation induced by prorenin.

The TGF- β level was observed and it was found that prorenin increased TGF- β mRNA levels and the secretion of TGF- β in HRMCs. PD98059, which is an inhibitor of ERK1/2, reduced the phosphorylation of ERK1/2 and the level

of TGF- β . RT-PCR and ELISA were used to test the mRNA and protein level of TGF- β , respectively; both revealed the same results. It was concluded that the regulation of TGF- β is highly dependent on ERK1/2 signaling transduction. TGF- β is the key mediator of renal tissue fibrosis (15). It enhances extracellular matrix synthesis and extracellular disposition, promoting mesangial cell proliferation (19). These processes play a significant role in diabetic nephropathy and kidney complications in hypertension. These results suggest a novel mechanism for the development of renal fibrosis in the presence of a high prorenin concentration, including in diabetic nephropathy. The MAPK cascade plays a pivotal role in the induction of cell differentiation, proliferation and cell stress by G protein-coupled receptors in response to extracellular stimulation (20). Once activated, ERK1/2 translocates to the nucleus, where it is thought to regulate the expression of transcription factors and thereby regulate cell differentiation and proliferation (21). Inhibiting ERK1/2 phosphorylation may shed light on therapies for renal disease.

In the present study, it was found that prorenin activated (P)RR, which led to the activation of ERK1/2 and increased TGF- β in cultured HRMCs. According to a previous study, (P)RR is involved in cardiac fibrosis (22), nephrosclerosis (14) and microvascular complications (23). As a result, it was concluded that the activation of (P)RR induced by prorenin in HRMCs may be involved in the development of human renal disease. Blockers of the RAS are widely used for the treatment of cardiovascular and renal diseases. It is generally assumed that the beneficial effects of these drugs are due, at least in part, to the blockade of the generation or action of angiotensin II at the tissue site (24). This effect of the (P)RR is prevented by the angiotensin-converting enzyme (ACE) inhibitors, angiotensin II type 1 receptor blockers and aliskiren, which is a direct renin inhibitor that was introduced in the clinical arena in 2007 (25). However, the high plasma and local tissue renin and prorenin levels induced by therapeutic AT₁ receptor blockers and aliskiren could conceivably induce (P)RR signaling independent of Ang II and may contribute to vascular sclerosis and renal fibrosis, thereby limiting the effectiveness of this therapy.

Since we aimed to examine a new method of blocking the effect of (P)RR, HRP was used to block the (P)RR. It was found that the HRP inhibited neither the phosphorylation of ERK1/2 nor the increase in TGF- β induced by prorenin. The effect of the putative blocker HRP is controversial. Ichihara *et al* revealed that HRP normalized renal angiotensin I and angiotensin II levels. Proteinuria was also normalized, and the glomerulosclerosis was completely inhibited in double transgenic model animals. HRP effectively reversed myocardial fibrosis and reduced the collagenase type I and III levels to almost normal values (12-14,26). However, HRP failed to block p-ERK1/2 in monocytes and vascular smooth muscle cells and did not affect prorenin binding to its receptor or Ang II generation (27-29). HRP did not affect blood pressure, cardiac hypertrophy and renal damage in two-kidney, one-clip Goldblatt hypertensive rats (30). However, no solid proof is currently available that clearly establishes HRP as a (P)RR blocker. The evidence shows that HRP failed to inhibit ERK1/2 phosphorylation and TGF- β expression in HRMCs. Additional evidence is required before the role of HRP is established.

Experimental evidence regarding the role of prorenin and (P)RR in renal damage suggests that blocking (P)RR is a new therapeutic target for tissue protection. As an inhibitor of ERK1/2, PD98059 raises the possibility that blocking signal transduction may lead to increased tissue protection. However, whether the benefits outweigh the drawbacks have yet to be fully determined as it is unknown whether there is cross-talk between MAPK signaling pathways.

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