Angiotensin II inhibits chemokine CCL5 expression in vascular smooth muscle cells from spontaneously hypertensive rats

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Angiotensin II (Ang II) exerts some of its effects on the vasculature by stimulating chemokines and 12-lipoxygenase (12-LO). In addition, a high expression of chemokines by Ang II has been observed in vascular smooth muscle cells (VSMCs) in spontaneously hypertensive rats (SHR). In this study, the action mechanism of Ang II on CCL5 expression in SHR VSMCs was examined. Expression of CCL5 in SHR thoracic aorta tissues and VSMCs was lower than that in normotensive Wistar-Kyoto rats (WKY) thoracic aorta tissues and VSMCs. Moreover, Ang II inhibited CCL5 expression in SHR VSMCs, but not in WKY VSMCs. Inhibition of CCL5 by Ang II was mediated by both Ang II subtype 1 receptor (AT₁R) and subtype 2 receptor (AT₂R) activation in SHR VSMCs. However, Ang II did not inhibit CCL5 expression in SHR VSMCs that were transfected with 12-LO small interfering RNA. In addition, 12-LO metabolite, 12(S)-hydroxyeicosatetraenoic acid (HETE) inhibited CCL5 mRNA expression in SHR VSMCs. The expression of Ang II-induced 12-LO was also blocked by both AT₁R and AT₂R inhibitors. Mitogen-activated protein (MAP) kinase, extracellular signal-regulated kinase (ERK)1/2, p38 and Jun N-terminal kinase pathways all mediated the inhibitory action of Ang II on CCL5 expression in SHR VSMCs. Taken together, the inhibitory action of Ang II on CCL5 expression was shown to be mediated by the 12-LO pathway through the activation of both of AT₁R and AT₂R and this process was associated with MAP kinase pathways in SHR VSMCs. This result suggests that upregulation of 12-LO by Ang II leads to the downregulation of CCL5 expression in SHR VSMCs.

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INTRODUCTION

Angiotensin II (Ang II) is a potent vasoconstrictor and blood pressure regulator. In addition, Ang II has been shown to function as a potential mediator of inflammation and was reported to exert some of its effects on the vasculature by stimulating chemokines.^{1–7}

Chemokines have an important role in Ang II-induced vascular hypertension,^{8–11} and controlling chemokine production is important for regulating inflammatory reactions in hypertensive vascular walls. In fact, the suppression of chemokine-induced inflammatory cell infiltration has been shown to ameliorate hypertension in experimental animal models.^{12–15} Thus, up- and downregulation of chemokines has been considered a therapeutic strategy for blocking disease development or progress, including hypertension.¹⁶ However, the molecular and cellular mechanisms of most chemokines are not yet clearly understood.

The proinflammatory chemokine CCL5 (regulated upon activation, normally T-cell expressed and presumably secreted; RANTES) is a potent chemoattractant for monocytes/macrophages and memory T lymphocytes, and its production has been described in various cell

types, including human aortic vascular smooth muscle cells (VSMCs).^{17,18} It has a functional role in acute and chronic inflammatory responses in atherosclerosis, renal disease progression and vascular wall remodeling in pulmonary arterial hypertension.^{2,3,19,20} However, CCL5 downregulates LPS-induced cytokines expression in human peripheral blood monocytes,²¹ and has a possible neuroprotective role in the brains of patients with Alzheimer's disease.²²

Some chemokines upregulate Ang II-induced hypertension and a high expression of chemokines has been detected in VSMCs isolated from spontaneously hypertensive rats (SHR).^{8–11} The increase of chemokine CCL2 or CXCL8 expression by Ang II has been shown in SHR VSMCs and rat glomerular endothelial cells.^{6,8,9,23,24} The increase of CCL5 expression by Ang II has been also demonstrated in rat glomerular endothelial cells and the renal cortex.^{2,3} However, CCL5 has been shown to inhibit the expression of Ang II-induced 12-lipoxygenase (LO), a hypertensive modulator in SHR VSMCs.²⁵

Although the relationships between Ang II and some chemokines or chemokine receptors, including CCL2 and CXCL8, in renal disease and vascular hypertension have been studied,^{4,8,9,23,24} the relationship

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between Ang II and CCL5 has not yet been characterized and little is known in SHR VSMCs. Therefore, as a first step to understand the relationship between Ang II and CCL5 in Ang II-induced vascular hypertension, the effect of Ang II on CCL5 expression was evaluated in SHR VSMCs.

METHODS

Reagent

Trizol reagent for total RNA isolation was purchased from Invitrogen (Carlsbad, CA, USA). Dulbecco's phosphate-buffered saline, Dulbecco's modified Eagle's medium, penicillin-streptomycin and fetal bovine serum were purchased from Gibco/BRL (Life Technologies, Gaithersburg, MD, USA). Ang II was obtained from Calbiochem (San Diego, CA, USA). Losartan was obtained from MSD (Delaware, MD, USA). PD123,319 was obtained from Sigma Chemical (St Louis, MO, USA). LightCycler FastStart DNA SYBR Green I Mix was obtained from Roche (Mannheim, Germany). Bay11-7082 was obtained from Merck (Merck KGaA, Darmstadt, Germany). Baicalein was obtained from Sigma-Aldrich. 12(S)-HETE was purchased from Cayman Chemical (Ann Arbor, MI, USA). PD98059, PD169316 and SP600125 were purchased from Merck. The primer sequences for CCL5, 12-LO, AT1R, AT2R and β-actin were purchased from Bioneer technology (Daejeon, South Korea). The rat AT1R small interfering RNA (siRNA), AT2R siRNA, 12-LO siRNA, ERK1 siRNA, ERK2 siRNA and Jun N-terminal kinase (JNK) siRNA sequences were purchased from Bioneer technology. Rat p38 siRNA sequence was purchased from Invitrogen. All other reagents were pure-grade commercial preparations.

Experimental animals

Specific pathogen-free male inbred SHR or normotensive Wistar-Kyoto rats (WKY), 12–16 weeks of age, were purchased from Japan SLC (Shizuka, Japan). All experimental animals received autoclaved food and bedding to minimize exposure to viral and/or microbial pathogens. Rats were cared for in accordance with the Guide for the Care and Use of Experimental Animals of Yeungnam Medical Center.

Immunohistochemistry

Aortic cross-sections were immunohistochemically stained for CCL5. To assess CCL5 protein expression, slides fixed with aortic cross-sections were treated with pepsin for antigen retrieval, and endogenous peroxidase was blocked with 3% H₂O₂. After blocking with normal blocking serum for 30 min at room temperature (RT), sections were incubated with the anti-CCL5 antibody (1:50) for 1 h at RT and HRP-conjugated rabbit anti-mouse IgG antibody for 30 min at RT. The sections were then developed with the 3,3'-diaminobenzidine substrate-chromogen system for 5 min at RT.

VSMCs preparation

VSMCs were obtained from the thoracic aortas of 12–16 week-old-male SHR and WKY using the explant method as described by Kim *et al.*⁸ VSMCs were cultured in Dulbecco's modified Eagle's medium, which was supplemented with 10% fetal bovine serum and 1% penicillin–streptomycin. Cells were detached with 0.25% trypsin/EDTA and seeded onto 75-cm² tissue culture flasks at a density of 10⁵ cells ml⁻¹. All experiments were conducted between cell passages 3–7. Before stimulation, 95%-confluent VSMCs were serumstarved overnight by incubating in Dulbecco's modified Eagle's medium with 0.1% fetal bovine serum. Cell cultures were incubated in a humidified incubator at 37 °C and 5% CO₂ in the presence or absence of stimuli for the indicated times.

Preparation of total RNA, reverse transcriptase-PCR and real-time PCR

Total RNA was extracted using the Trizol reagent according to the manufacturer's instructions. The quantity of total RNA obtained was determined by measuring the optical density at 260 and 280 nm.

A total of $1\,\mu g$ RNA per sample was reverse transcribed using Moloney murine leukemia virus reverse transcriptase (Perkin Elmer, Norwalk, CT, USA)

and oligo dT priming at 42 °C for 15 min according to the manufacturer's instruction. Amplification with specific primers was performed in a Gene Amp PCR system 9600 (Perkin Elmer) for 35 cycles with a 30 s/94 °C denaturation, 30 s/62 °C annealing, 1 min/72 °C extension profile in the case of CCL5; for 35 cycles with a 20 s/94 °C denaturation, 10 s/55 °C annealing, 40 s/72 °C extension profile in the case of AT1R; for 35 cycles with a 20 s/94 °C denaturation, 10 s/ 60 °C annealing, 40 s/72 °C extension profile in the case of AT2R; for 30 cycles with a 20 s/95 °C denaturation, 30 s/60 °C annealing, 30 s/72 °C extension profile in the case of β-actin. The mRNA of the housekeeping gene β-actin was amplified and used as an internal quality standard. Amplified products were electrophoresed on 1.5–2% agarose gel and stained with $0.5\,\mu g\,ml^{-1}$ ethidium bromide. The primer sequences were as follows: B-actin (101 bp) sense, 5'-TACTGCCCTGGCTCCTAGCA-3', antisense, 5'-TGGACAGTGAG GCCAGGATAG-3'; CCL5 (110 bp) sense, 5'-CGTGAAGGAGTATTTTTACACC AGC-3', antisense, 5'-CTTGAACCCACTTCTTCTCTGGG-3'; AT₁R (445 bp) sense, 5'-CACCTATGTAAGATCGCTTC-3', antisense, 5'-GCACAATCGCCAT AATTATCC-3'; AT2R (65 bp) sense, 5'-CCGTGACCAAGTCTTGAAGATG-3', antisense, 5'-AGGGAAGCCAGCAAATGATG-3'.

12-LO, CCL5 was amplified by real-time PCR using the LightCycler (Roche). RNA was reverse transcribed to cDNA from 1 µg of total RNA and then subjected to real-time PCR. PCR was performed in triplicate. The total PCR volume was 20 µl and the PCR consisted of LightCycler FastStart DNA SYBR Green I mix (Roche), primer and 2 µl of cDNA. Before PCR amplification, the mixture was incubated at 95 °C for 10 min, and the amplification step consisted of 45 cycles of denaturation (10 s at 95 °C), annealing (5 s at the primerappropriate temperature) and extension (10s at 72 °C) with fluorescence detection at 72 °C after each cycle. After the final cycle, melting point analyses of all samples were performed over a temperature range of 65-95 °C with continuous fluorescence detection. β-actin expression levels were used for sample normalization. Results for each gene were expressed as the relative expression level compared with β-actin. The primers used for PCR were as follows: 12-LO (312 bp) sense, 5'-TGGGGGCAACTGGAAGG-3', antisense, 5'-AGAGCGCTTCAGCACCAT-3'; CCL5 (110 bp) sense, 5'-CGTGAAGGAGT ATTTTTACACCAGC-3', antisense, 5'-CTTGAACCCACTTCTTCTCTGGG-3'; β-actin (101 bp) sense, 5'-TACTGCCCTGGCTCCTAGCA-3', antisense, 5'-TGGACAGTGAGGCCAGGATAG-3'. The levels of 12-LO and CCL5 mRNA were determined by comparing experimental levels to standard curves and were expressed as relative fold expressions.

Electrophoretic mobility shift assay

Cells were washed three times with cold phosphate-buffered saline, then scraped and harvested by centrifugation. Cell pellets were resuspended and incubated on ice for 15 min in 400 µl of hypotonic buffer A (10 mmoll⁻¹ HEPES, 10 mmol l⁻¹ KCl, 1.5 mmol l⁻¹ MgCl₂, 0.5 mmol l⁻¹ Dithiothreitol 0.1 mmoll^{-1} phenylmethylsulfonyl fluoride, $10 \,\mu\text{g ml}^{-1}$ pepstatin, $10 \,\mu\text{g ml}^{-1}$ leupeptin, 10 µg ml⁻¹ autipain and 10 µg ml⁻¹ aprotinin). Nonidet P-40 was then added to a final concentration of 2.5%, and the cells were vortexed for 10 s. Nuclei were separated from the cytosol by centrifugation at 12000 g for 15 s. Pellets were resuspended in 40 µl of hypotonic buffer C (20 mmoll-1 HEPES, 25% glycerol, 0.4 mol l⁻¹ NaCl, 1 mmol l⁻¹ EDTA, 1 mmol l⁻¹ ethylene glycol tetraacetic acid, 0.5 mmol l⁻¹ Dithiothreitol, 0.1 mmol l⁻¹ phenylmethylsulfonyl fluoride, 10 µg ml⁻¹ pepstatin, 10 µg ml⁻¹ leupeptin, 10 µg ml⁻¹ autipain and $10 \,\mu g \,m l^{-1}$ aprotinin). Samples were sonicated for 2-3 s, then centrifuged for 10 min at 4 °C. The nuclear protein concentration was measured using the Bradford assay (Bio-Rad, Richmond, CA, USA). The consensus sequence of the NF-κβ DNA binding site (5'-AGTTGAGGGGACTTTAGGC-3') (sc-2505; Santa Cruz Biotechnology, Santa Cruz, CA, USA) was labeled with (\alpha-32P)dCTP using a random-primed DNA labeling kit (Roche). The mutant NF- $\kappa\beta$ binding sequence was identical to sc-2505 except for a 'G' \rightarrow 'C' substitution in the NF-κβ DNA binding motif (sc-2511; Santa Cruz Biotechnology). The labeled DNA was purified over a S-200 HR column (Pharmacia, Piscataway, NJ, USA) to remove unbound nucleotides. Nuclear protein extracts were incubated at RT for 20 min with \sim 50 000 c.p.m. of labeled oligonucleotides that were suspended in a binding buffer (200 mmoll⁻¹ HEPES, 500 mmol l⁻¹ KCl, 10 mmol l⁻¹ EDTA, 50% glycerol, 10 mmol l⁻¹ Dithiothreitol, 1 mg ml⁻¹ BSA and 1 μ g μ l⁻¹ poly (dI-dC)). Following this incubation step,

the samples were resolved on 4% polyacrylamide gels at $140\,\mathrm{V}$ and exposed to film.

Small interfering RNA

VSMCs were plated on 24-well plates and grown to 90% confluence. VSMCs were then transfected with AT2R siRNA oligomers (50 nmol l-1) using lipofectamine 2000 in accordance with the manufacturer's instructions. After 24 h of incubation, VSMCs were placed in growth medium for 24 h before the experiments. Cells were then cultured in the presence or absence of stimuli for 4 h. The sense and antisense oligonucleotides used in these experiments were as follows: AT1R siRNA sense, 5'-GUCACUGUUACUACACCUA-3', antisense, 5'-UAGGUGUAGUAACAGUGAC-3'; AT2R siRNA sense, 5'-GAGUGUUGAUAG GUACCAA-3', antisense, 5'-UUGGUACCUAUCAACACUC-3'; 12-LO siRNA sense, 5'-GUGUGUGACUAUGUUCCAA-3', antisense, 5'-UUGGAACAUAG UCACACAC-3'; ERK1 siRNA sense, 5'-CUAUGAUCCGACAGAUGAA-3', antisense, 5'-UUCAUCUGUCGGAUCAUAG-3'; ERK2 siRNA sense, 5'-CAGG AAAGCAUUACCUUGA-3', antisense, 5'-UCAAGGUAAUGCUUUCCUG-3'; JNK siRNA sense, 5'-GAGCAAAAGCAAGGUAGAU-3', antisense, 5'-AUCUA CCUUGCUUUUGCUC-3'; p38 siRNA sense, 5'-UACAUUUGCGAAGUUC AUCUUCGGC-3', antisense, 5'-GCCGAAGAUGAACUUCGCAAAUGUA-3'.

Statistical analysis

Results were expressed as means \pm s.e.m. from at least three or four independent experiments. For comparisons between multiple groups, statistical significance was determined by the Mann–Whitney test using SPSS version 17.0.

RESULTS

Ang II suppresses chemokine CCL5 expression in SHR VSMCs

First, CCL5 expression in SHR and WKY thoracic aorta tissues and VSMCs was examined. Expression of CCL5 in SHR thoracic aorta tissues and VSMCs was lower than that in WKY thoracic aorta tissues and VSMCs (Figure 1a). The expression of CCL5 mRNA by Ang II was then examined in SHR and WKY VSMCs. Ang II inhibited CCL5 mRNA expression in SHR VSMCs, but did not inhibit CCL5 expression in WKY VSMCs (Figure 1b). A decreased CCL5 level was also detected in SHR VSMCs that had been treated with Ang II (100 nM)

after a reaction time of 1 and 4 h (Figure 1c). The time course of Ang II-induced CCL5 inhibition was then determined in SHR VSMCs over a 0–16-h time period. In this experiment, inhibition of CCL5 mRNA induced by Ang II was detected 1 h after treatment and was sustained for up to 8 h after treatment and this effect disappeared at 16 h. The inhibition levels of CCL5 by Ang II were almost same from 1 h after treatment to 8 h after treatment (Figure 2a). The dose response of Ang II-induced CCL5 inhibition was also observed. Inhibition of CCL5 expression was initially detected from at a Ang II concentration of 10 nM Ang II. Although the inhibition by Ang II was shown in all three concentrations of 10, 100 and 1000 nM, there were no statistical differences among these concentrations (Figure 2b).

To understand the nature of the inhibitory effect of Ang II on CCL5 expression in SHR VSMCs, the role of NF- $\kappa\beta$ activation was examined. Bay11-7082 is known to selectively block the phosphorylation of IkB α ; thus, preventing activation and nuclear translocation of NF- κ B. Real-time PCR, (enzyme-linked immunosorbent assay) and electrophoretic mobility shift assay were performed on VSMCs after they had been treated or not treated with Ang II (100 nM) in the absence or presence of Bay11-7082 (10 μ M) for 1 h. Bay11-7082 prevented Ang II from inhibiting CCL5 mRNA expression, and the protein levels of CCL5 correlated to the mRNA levels (Figure 2c, upper). Increased NF- $\kappa\beta$ activity in SHR VSMCs treated with Ang II alone disappeared in cells treated with Ang II and Bay11-7082 (Figure 2c, lower).

To examine whether inhibition of CCL5 expression by Ang II was mediated by the Ang II subtype 1 receptor (AT_1R) or subtype 2 receptor (AT_2R) , SHR VSMCs were treated or not treated with Ang II (100 nM) in the presence or absence of an antagonist of the AT_1R , losartan (10 μ M) or an antagonist of the AT_2R , PD123,319 (10 μ M) for 1 or 4 h, and the total RNAs were analyzed by reverse transcriptase-PCR. Protein levels of CCL5 in the cell supernatants were evaluated by (enzyme-linked immunosorbent assay). The inhibitory effect of Ang II on CCL5 mRNA expression was not observed in the presence of both



Figure 1 Expression of CCL5 in SHR and WKY thoracic aorta tissues and VSMCs, and the inhibitory effect of Ang II on CCL5 expression in SHR VSMCs. (a) Aortic cross-sections were immunohistochemically stained for CCL5. Original magnification was $\times 200$ (Zeiss microscope, Oberkochen, Germany). After total RNAs were isolated from SHR or WKY VSMCs, real-time PCR was performed. Bars represent means ± s.e.m. from three independent experiments. *P < 0.05 vs. WKY VSMCs. (b) SHR and WKY VSMCs were treated or not treated (NT) with Ang II (0.1 µmol I⁻¹) for 1 h. After total RNAs were isolated, real-time PCR was performed. *P < 0.05 vs. untreated SHR VSMCs. (c) SHR VSMCs. (c) SHR VSMCs were treated or NT with Ang II (0.1 µmol I⁻¹) for 1 or 4 h. After cell supernatants were isolated, (enzyme-linked immunosorbent assay) was performed. Bars represent means ± s.e.m. from three independent experiments. *P < 0.05 vs. untreated SHR VSMCs. A full color version of this figure is available at the *Hypertension Research* journal online.

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Figure 2 Time course and dose response of CCL5 mRNA inhibition by Ang II in SHR VSMCs, and inhibitory action of Ang II on CCL5 expression is dependent on NF- $\kappa\beta$ activation in SHR VSMCs. (a) SHR VSMCs were treated with Ang II (0.1 µmol I⁻¹) for the indicated times and the total RNA was analyzed by real-time PCR. Symbols represent means ± s.e.m. from three independent experiments. (b) SHR VSMCs were treated or not treated (NT) with 10, 100 or 1000 nmol I⁻¹ of Ang II for 1 h. After total RNAs were isolated, real-time PCR was performed. Bars represent means ± s.e.m. from three independent experiments. **P*<0.05 *vs.* untreated SHR VSMCs. (c) SHR VSMCs were treated or NT with Ang II (0.1 µmol I⁻¹) in the absence or presence of Bay11-7082 (10 µm, a selective inhibitor of I κ Ba) for 1 h. After total RNAs and cell supernatants were isolated, real-time PCR and (enzyme-linked immunosorbent assay) were performed. Bars represent means ± s.e.m. from three independent experiments. ^a*P*<0.05 *vs.* untreated SHR VSMCs. ^b*P*<0.05 *vs.* SHR VSMCs treated with Ang II. Specific binding activity of NF- $\kappa\beta$ from nuclear extracts was assessed by electrophoretic mobility shift assay (EMSA). Aliquots of the nuclear extract were incubated with a 100-fold excess of the mutant probe (M) or with 2 µg of the anti-NF- $\kappa\beta$ antibody (Ab) before EMSA. Bay:Bay11-7082, Data shown are representative of three independent experiments.



Figure 3 Inhibitory effect of Ang II on CCL5 mRNA expression is related to the both of AT_1R and AT_2R activation in SHR VSMCs. (**a**, **b**) SHR VSMCs were treated or not treated (NT) with Ang II (0.1 μ mol I⁻¹) in the presence or absence of losartan (AT_1R antagonist, 10 μ mol I⁻¹) or PD123319 (AT_2R antagonist, 10 μ mol I⁻¹) for 1 or 4 h. After total RNAs and cell supernatants were isolated, reverse transcriptase-PCR and (enzyme-linked immunosorbent assay) were performed. Data shown are representative of three independent experiments. Bars represent means ± s.e.m. from three independent experiments. ^a*P*<0.05 *vs.* untreated SHR VSMCs. ^b*P*<0.05 *vs.* SHR VSMCs treated with Ang II. (**c**) SHR VSMCs were plated on 24-well plates, grown to 90% confluence and then transfected with AT_1R or AT_2R siRNA oligomers (50 nmol I⁻¹). Transfected or non-transfected SHR VSMCs were then treated or NT with Ang II (0.1 μ mol I⁻¹) for 1 h. Total RNAs were analyzed by reverse transcriptase-PCR to confirm successful transfection, Real-time PCR was then performed. Bars represent means ± s.e.m. from three independent experiments. The data are representative of three independent experiments.

losartan and PD123319 (Figure 3a) and the CCL5 levels increased to the CCL5 levels observed in the supernatant of untreated cells (Figure 3b). To further confirm these results, real-time PCR was performed in samples treated with AT_1R or AT_2R -directed siRNA.

Successful transfection of AT_1R or AT_2R siRNA into the SHR VSMCs was observed by reverse transcriptase-PCR and the inhibitory action of Ang II on CCL5 mRNA expression was not detected in SHR VSMCs transfected with AT_1R or AT_2R siRNA (Figure 3c).

The inhibitory effect of Ang II on CCL5 expression is mediated by 12-LO

Ang II is a potent positive regulator of 12-LO expression and activation in porcine and human VSMCs.^{26,27} In a previous study, CCL5 was shown to inhibit Ang II-induced 12-LO expression in SHR VSMCs.²⁵ Thus, to examine whether the inhibitory action of Ang II on CCL5 expression was related to 12-LO activation, SHR VSMCs were treated or not treated with Ang II (100 nM) and/or baicalein (a 12-LO inhibitor, 10 μ M) for 1 or 4 h. The inhibitory effect of Ang II on CCL5 mRNA expression was reduced in the presence of baicalein and the protein level of CCL5 increased significantly in the supernatant of cells treated with Ang II and baicalein together (Figure 4a). To further confirm these results, real-time PCR was performed on SHR VSMCs treated with 12-LO-directed siRNA. Successful transfection of 12-LO siRNA into SHR VSMCs was observed by reverse transcriptase-PCR and the inhibitory effect of Ang II on CCL5 mRNA expression was not observed in SHR VSMCs that had been treated with 12-LO-directed siRNA (Figure 4b).

The activities of 12-LO and the 12-LO metabolite, 12(S)-hydroxyeicosatetraenoic acid (12(S)-HETE) were shown to be higher in SHR,^{10,28} thus, to further understand the involvement of 12-LO in the inhibition of CCL5 expression in SHR VSMCs by Ang II, the direct effect of 12(S)-HETE on CCL5 expression was examined in SHR VSMCs. 12(S)-HETE inhibited CCL5 mRNA expression in SHR VSMCs (Figure 4c). In addition, the expression of Ang II-induced 12-LO was decreased by both AT_1R and AT_2R inhibitors (Figure 4d).

Inhibitory action of Ang II on CCL5 expression is mediated through MAPK signaling pathways in SHR VSMCs

The role of MAPK signaling pathways in the inhibitory effect of Ang II on CCL5 mRNA expression in SHR VSMCs was also examined. After SHR VSMCs were pretreated with the ERK1/2 inhibitor PD98059 (10 µM), the p38 mitogen-activated protein (MAP) kinase inhibitor PD169316 (10 µm) or the JNK inhibitor SP600125 (25 µm) for 0.5 h, the cells were treated or not treated with Ang II (100 nm) for 1 h. Realtime PCR and (enzyme-linked immunosorbent assay) were then performed. PD98059, PD169316 and SP600125 blocked the inhibitory action of Ang II on CCL5 mRNA expression in SHR VSMCs (Figure 5a). These (enzyme-linked immunosorbent assay) results showed the same patterns that were observed for gene expression (Figure 5b). In addition, to further confirm these results, real-time PCR was performed on SHR VSMCs treated with ERK1/2, p38 or JNK-directed siRNA. Inhibitory effect of Ang II on CCL5 mRNA expression was not observed in SHR VSMCs transfected with ERK1/2, p38 or JNK siRNA (Figure 5c).



Figure 4 12-LO is involved in the inhibitory action of Ang II on CCL5 expression in SHR VSMCs. (a) SHR VSMCs were treated or not treated (NT) with Ang II (0.1μ mol I⁻¹) in the absence or presence of baicalein (12-LO inhibitor, 10μ mol I⁻¹) for 1 h (for real-time PCR) or 4 h. After total RNAs and cell supernatants were isolated, real-time PCR and (enzyme-linked immunosorbent assay) were performed. Bars represent means ± s.e.m. from three independent experiments. ^a*P*<0.05 *vs.* untreated SHR VSMCs. ^b*P*<0.05 *vs.* SHR VSMCs treated with Ang II. (b) SHR VSMCs were plated on 24-well plates, grown to 90% confluence and then transfected with 12-LO siRNA oligomers (50 nmol I⁻¹). Transfected or non-transfected SHR VSMCs were then treated or NT with Ang II (0.1μ mol I⁻¹) for 1 h. Total RNAs were analyzed by reverse transcriptase-PCR to confirm successful transfection, real-time PCR was then performed. Bars represent means ± s.e.m. from three independent experiments. The data are representative of three independent experiments. (c) SHR VSMCs were treated or NT with Ang II (0.1μ mol I⁻¹) or 12(S)-HETE (500 nM) for 1 h. After total RNAs were isolated, real-time PCR was performed. Bars represent means ± s.e.m. from three independent experiments. **P*<0.05 *vs.* untreated SHR VSMCs. (d) SHR VSMCs were treated or NT with Ang II (0.1μ mol I⁻¹) or 12(S)-HETE (500 nM) for 1 h. After total RNAs were isolated, real-time PCR was performed. Bars represent means ± s.e.m. from three independent experiments. **P*<0.05 *vs.* untreated SHR VSMCs. (d) SHR VSMCs were treated or NT with Ang II (0.1μ mol I⁻¹) in the presence or basence of losartan (AT₁R antagonist, 10μ mol I⁻¹) or PD123,319 (AT₂R antagonist, 10μ mol I⁻¹) for 1 h. After total RNAs were isolated, real-time PCR was performed. Bars represent means ± s.e.m. from three independent experiments. **P*<0.05 *vs.* Untreated SHR VSMCs. (d) SHR VSMCs were treated or NT with Ang II (0.1μ mol I⁻¹) in the presence or a sesting experiment. *

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Figure 5 MAP kinases are involved in the inhibitory action of Ang II on CCL5 expression. (**a**, **b**) SHR VSMCs were untreated (NT) or pretreated with PD98059 (ERK inhibitor, 10μ M), PD169316 (p38 inhibitor, 10μ M) or SP600125 (JNK inhibitor, 25μ M) for 30 min, and then treated or not treated (NT) with Ang II (0.1μ mol I⁻¹) for 1 h. After total RNAs and cell supernatants were isolated, real-time PCR and (enzyme-linked immunosorbent assay) were performed. Bars represent means ± s.e.m. from three independent experiments. ^a*P*<0.05 *vs.* untreated SHR VSMCs. ^b*P*<0.05 *vs.* SHR VSMCs treated with Ang II. (**c**) SHR VSMCs were plated on 24-well plates, grown to 90% confluence and then transfected with ERK1, ERK2, ERK1/2, p38 or JNK siRNA oligomers (50 nmol I⁻¹). Transfected or non-transfected SHR VSMCs were then treated or NT with Ang II (0.1 µmol I⁻¹) for 1 h. Total RNAs were isolated and analyzed by real-time PCR. Bars represent means ± s.e.m. from three independent experiments.

DISCUSSION

In naive SHR thoracic aorta tissues, the expression of CCL5 was lower than that in naive WKY thoracic aorta tissues. Although some chemokines are constitutively transcribed and translated, constitutive production of most chemokines is usually low or absent. CCL5 transcription has been known to occur relatively late after the activation of naive cells and is coincident with the upregulation of cytolytic granules in cytotoxic T-cells.²⁹ However, CCL5 mRNA is constitutively transcribed and translated in peripheral blood natural killer cells³⁰ and the expression of CCL5 mRNA has been observed in naive WKY VSMCs.²⁵ Thus, constitutive expression of a chemokine may be dependent on cell type.

In this study, the low expression of CCL5 was demonstrated in SHR thoracic aortic tissues and VSMCs compared with WKY thoracic aortic tissues and VSMCs, and Ang II suppressed CCL5 expression in SHR VSMCs, but not in WKY VSMCs. However, high expression of CCL5 was demonstrated in pulmonary arterial hypertension and the endothelial cells within the pulmonary arterial wall were reported to be a major source of CCL5 production.¹⁹ Dorfmüller *et al.*¹⁹ reported that CCL5 may have a key role in arterial inflammatory processes, such as glomerulonephritis, Kawasaki disease and Takayasu's arteritis, including pulmonary arterial hypertension. Furthermore, Ang II was shown to increase CCL5 expression in rat glomerular endothelial cells from Sprague-Dawley rats and the renal cortex of Wistar rats.^{2,3} Thus, they suggested that Ang II-mediated induction of CCL5 may have an

important role in glomerular infiltration of monocytes/macrophages. Wolf *et al.*³ demonstrated that although Ang II increased CCL5 expression in rat glomerular endothelial cells, it did not induce CCL5 mRNA in mesangial cells. These discrepancies could be due to the different cell types and/or different experimental animals or the pleiotropic action of Ang II as a vasoactive cytokine.⁷

Ang II has two subtype receptors, the AT_1R and the AT_2R , and the density of the AT₂R is lower than that of the AT₁R in VSMCs.³¹ Ang II has been reported to increase AT₁R expression in SHR VSMCs, but Ang II was shown to only slightly affect AT₂R expression in SHR VSMCs.⁸ AT₁R mediates the major stimulatory actions of Ang II, including vasoconstriction, cell proliferation, aldosterone secretion and sodium retention.³² In contrast, AT₂R has been reported to antagonize the vascular actions of AT1R. However, several studies have reported growth and proinflammatory actions of AT₂R in VSMCs.^{31,33,34} Moreover, a positive role of the AT₂R in Ang IIinduced CCL5 expression has been demonstrated in rat glomerular endothelial cells and the rat renal cortex.^{2,3} The inhibitory action of CCL5 on Ang II-induced 12-LO expression is also mediated through AT₂R in SHR VSMCs.²⁵ In this study, the inhibition of CCL5 expression by Ang II in SHR VSMCs was found to be mediated through both AT₁R and AT₂R activation.

Ang II is a potent positive regulator of 12-LO expression and proliferation in porcine and human VSMCs.^{26,27} The activities of 12-LO and the 12-LO metabolite 12(S)-HETE are increased in



Figure 6 Flow diagram of the action of Ang II on CCL5 inhibition in SHR VSMCs.

SHR,^{10,28} and many studies have demonstrated that 12-LO and 12(S)-HETE have an important role in the pathogenesis of hypertension.^{10,28,35,36} This study demonstrated that the inhibitory action of Ang II on CCL5 expression in SHR VSMCs was mediated through 12-LO activation. Ang II did not inhibit CCL5 expression in 12-LO siRNA-transfected SHR VSMCs. In addition, the 12-LO metabolite, 12(S)-HETE directly inhibited CCL5 expression in SHR VSMCs. Moreover, the expression of Ang II-induced 12-LO mRNA was decreased by both AT₁R and AT₂R inhibitors. Taken together, these results suggest that activation of Ang II receptors by Ang II leads to the expression of 12-LO, which then acts to inhibit CCL5 in SHR VSMCs. Namely, the lower expression of CCL5 in SHR VSMCs compared with WKY VSMCs is likely because of the activity of 12-LO in SHR VSMCs.

Vascular structural changes in Ang II-induced hypertension are associated with cell proliferation and increased deposition of inflammatory cells and extracellular matrix components. Mechanisms underlying these processes are not still clear; however, the role of MAP kinases may be important. Ang II stimulates the activation of MAP kinases, including ERK1/2, p38 and JNK in VSMCs.31,37,38 Ang IIinduced ERK1/2 and p38 activation is increased in SHR vasculature³⁹ and ERK1/2 and p38 activation have important roles in Ang IIinduced VSMCs proliferation.^{37,40,41} In these previous studies, the downregulatory effect of CCL5 on Ang II-induced SHR VSMCs proliferation was mediated by inactivation of p38,²⁵ and the expression of CXCL8 by Ang II was mediated by ERK1/2 MAP kinase activation.8 A potent JNK-activating effect of Ang II was demonstrated in rat aortic cells including VSMCs.⁴⁰ Glomerular JNK activity was shown to be increased in Ang II-induced hypertension⁴² and cardiac JNK was found to be activated more sensitively than ERK1/2 in Ang II-induced hypertension.⁴³ However, Touyz et al.³⁹ reported that Ang II has no effect on JNK phosphorylation in SHR or WKY VSMCs. This discrepancy may be due to the relative distribution of Ang II receptors, where the density of the AT₂R is lower than that of the AT₁R in VSMCs,³¹ namely, Ang II-induced JNK activation occurs through AT₂R. Another reason for this discrepancy may be the different experimental reaction times; namely, JNK activation by Ang II is slower than ERK1/2 and p38 activation.44,45 In this study, all three MAPK signaling pathways were shown to be involved in the inhibitory action of Ang II on CCL5 mRNA expression in SHR VSMCs. The inhibitory effect of Ang II on CCL5 mRNA expression was blocked in SHR VSMCs transfected with ERK1/2, p38 or JNK siRNA.

These combined results indicate that the inhibitory action of Ang II on CCL5 expression was mediated by the 12-LO pathway through the activation of both AT₁R and AT₂R and was associated with MAP kinases, ERK1/2, p38 and JNK, pathways in SHR VSMCs (Figure 6). In SHR VSMCs, CCL5 downregulates Ang II-induced 12-LO expression and VSMCs proliferation²⁵ and has an upregulatory effect on DDAH-1 expression, which is a regulator of nitric oxide activity (data not shown). On the other hand, Ang II suppresses CCL5 production through 12-LO activity in SHR VSMCs. Taken together, CCL5 is more likely to have a negative role at some stages, but not a positive role as an inflammatory chemokine, in Ang II-induced vascular hypertension.

CONFLICT OF INTEREST

The authors declare no conflict of interest.

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