

Original Article

Biophysically stressed vascular smooth muscle cells express MCP-1 via a PDGFR- β -HMGB1 signaling pathway

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ABSTRACT Vascular smooth muscle cells (VSMCs) under biophysical stress play an active role in the progression of vascular inflammation, but the precise mechanisms are unclear. This study examined the cellular expression of monocyte chemoattractant protein 1 (MCP-1) and its related mechanisms using cultured rat aortic VSMCs stimulated with mechanical stretch (MS, equibiaxial cyclic stretch, 60 cycles/min). When the cells were stimulated with 10% MS, MCP-1 expression was markedly increased compared to those in the cells stimulated with low MS intensity (3% or 5%). An enzyme-linked immunosorbent assay revealed an increase in HMGB1 released into culture media from the cells stimulated with 10% MS compared to those stimulated with 3% MS. A pretreatment with glycyrrhizin, a HMGB1 inhibitor, resulted in the marked attenuation of MCP-1 expression in the cells stimulated with 10% MS, suggesting a key role of HMGB1 on MCP-1 expression. Western blot analysis revealed higher PDGFR- α and PDGFR- β expression in the cells stimulated with 10% MS than 3% MS-stimulated cells. In the cells deficient of PDGFR- β using siRNA, but not PDGFR- α , HMGB1 released into culture media was significantly attenuated in the 10% MS-stimulated cells. Similarly, MCP-1 expression induced in 10% MS-stimulated cells was also attenuated in cells deficient of PDGFR- β . Overall, the PDGFR- β signaling plays a pivotal role in the increased expression of MCP-1 in VSMCs stressed with 10% MS. Therefore, targeting PDGFR- β signaling in VSMCs might be a promising therapeutic strategy for vascular complications in the vasculatures under excessive biophysical stress.

INTRODUCTION

The vessel wall is continuously exposed to biomechanical stressors that elicit functional and adaptative responses. Among multiple cells in the blood vessel walls, vascular smooth muscle cells (VSMCs) are critical cells involved in vascular remodeling in the injured vasculatures [1]. VSMCs show proliferative and migratory activities in response to physiological stress or pathological insults such as inflammation and hypertension [2,3]. When blood pressure increases, the physical force continually exposes the blood vessel walls to mechanical stretch (MS) [4].

In the process of vascular remodeling, VSMCs in the media move into the intima and then proliferate in response to various mediators secreted by inflammatory cells [5,6]. In atherosclerotic plaques, high levels of high mobility group box 1 (HMGB1) are detected, which are suggested to be a major mediator involved in vascular inflammation [7,8]. The contractile phenotype of VSMCs in the normal vasculatures switches to the synthetic phenotype in vascular inflammation [9,10], leading VSMCs to migrate from the media to the intima, ultimately causing intimal hyperplasia and vascular restenosis [11]. Although HMGB1 was suggested to be a major contributor to the development of proliferative vascu-



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lar diseases through the phenotypic conversion of VSMCs [12,13], the precise role and molecular action mechanisms of HMGB1 are still unclear.

Inflammatory cytokines, such as monocyte chemoattractant protein 1 (MCP-1), play a vital role in the infiltration of monocytes into the vessel wall in the injured vasculatures. MCP-1 recruits various inflammatory cells including monocytes to the inflammation sites, indicating its importance in mediating neointima formation and vascular inflammation [14]. Although MCP-1 has been indicated as a major molecule in the initiation and progression of the vascular inflammatory process, the major cellular source and molecular mechanisms responsible for MCP-1 expression in VSMCs are unclear.

VSMCs under biophysical stress play an active role in vascular inflammation and remodeling, but the precise mechanisms are unclear. Therefore, this study investigated MCP-1 expression and its related mechanisms using VSMCs stressed with high intensity (10%) MS (equibiaxial cyclic stretch, 60 cycles/min). In addition, to investigate the individual importance of PDGFR isoforms on vascular remodeling in the vasculatures under excessive biophysical stress, this study identified the links between MCP-1 expression and PDGFR signaling in VSMCs stimulated with 10% MS.

METHODS

Ethics statements and animals

All animal procedures were complied with the Guide for the Care and Use of Laboratory Animals published by the United States National Institutes of Health (NIH Publication No.85-23, 2011 revision). The Pusan National University Institutional Animal Care and Use Committee approved all experimental protocols (PNU-2021-2965). Sprague-Dawley (SD) rats were purchased from Charles River Breeding Laboratories.

Materials

MCP-1 (PA5-34505) antibodies were purchased from Invitrogen. Glycyrrhizin (3-O-(2-O- β -D-Glucopyranuronosyl- α -D-glucopyranuronosyl)-18 β -glycyrrhetic acid) was obtained from Sigma-Aldrich Inc. β -Actin (sc-47778) antibody was acquired from Santa Cruz Biotechnology Inc. The PDGFR- α (3164S) and PDGFR- β (3169S) antibodies were supplied by Cell Signaling Technology. The secondary antibody used horseradish peroxidase (HRP)-conjugated IgG antibody obtained from Santa Cruz Biotechnology Inc.

Cell culture

Male SD rats (seven weeks old) were euthanized by CO₂ inhalation, and then dissected to isolate the thoracic aorta. The resected

aortas were cut and transplanted into a cell culture dish containing Dulbecco's Modified Eagle's Medium (DMEM, Gibco BRL) supplemented with 10% fetal bovine serum (FBS, Gibco BRL). The cells were then maintained at 37°C in DMEM with 10% FBS and an antibiotic-antimycotic solution (Gibco BRL).

Mechanical stretch

The VSMCs cultured from the rat thoracic aorta were seeded onto flexible-bottomed 6-well BioFlex culture plates (BF-3001C; Dunn Labortechnik) and incubated in CO₂ incubator at 37°C, 95% humidity and 5% CO₂ for 24 h. A Flexcell Tension Plus FX-4000T system (Flexcell International Corp.) was used to exert physiological equibiaxial cyclic stretch (1 Hz, 0%–10% strain, 60 cycles/min) for 0 to 12 h.

Western blot analysis

Ice-cold lysis buffer (Thermo Fisher Scientific) was used to collect and prepare the VSMC lysates. The same amounts of protein were separated on 10%–12% polyacrylamide gels under reducing conditions, and then transferred to nitrocellulose membranes (Amersham-Pharmacia Biotech). The membranes were blocked with 5% skim milk in Tris-buffered saline with Tween-20 (TBST) for 2 h at room temperature. The membranes were then incubated overnight with the primary antibodies in 5% skim milk at 4°C. TBST was used to wash the membranes, and then incubated with HRP-conjugated secondary antibody for 2 h at room temperature. The blots were developed using the enhanced chemiluminescence Western blotting detection reagents (Amersham-Pharmacia Biotech), and images of the blots were captured using image capturing software (Amersham, Imager 680 version. 2.0.). The signal bands were quantified using the UNSCAN-IT CEL 7.1 program. The membranes were reblotted with anti- β -actin antibody as an internal control.

Enzyme-linked immunosorbent assay

Rat aortic VSMCs seeded onto flexible-bottomed 6-well BioFlex culture plates were stimulated by MS, and the HMGB1 released into culture media was measured using a rat HMGB-1 ELISA kit (Elabscience Biotechnology Inc) according to the manufacturer's instructions.

Preparation and transfection of small interfering RNA (siRNA)

PDGFR- α and PDGFR- β siRNA oligonucleotides were synthesized at Bioneer to prepare cells deficient of PDGFR isoforms. The siRNA negative control duplex was used as a control. To transfect all the siRNA molecules, lipofectamine 2000 (Invitrogen) was used. Briefly, VSMCs were seeded on 6-well plates and grown

for 24 h before siRNA transfection. After replacing the culture medium with Optimized-Minimal Essential Medium (Opti-MEM, Gibco BRL), the cells were transfected with siRNAs for PDGFR- α , PDGFR- β and negative control using Lipofectamine 2000 and then incubated for 6 h at 37°C. The Opti-MEM was then replaced with DMEM, and the cells were incubated at 37°C for 24 h.

Statistical analysis

All data are expressed as the means \pm SEMs. One-way analysis of variance followed by Dunnett multiple comparison test or student's t-test was used to determine the significant differences. Statistical significance was accepted for p-values less than 0.05.

RESULTS

Increased MCP-1 expression in VSMCs stimulated with MS

The effects of MS on MCP-1 expression in VSMCs were identified by stimulating cultured rat aortic VSMCs seeded onto BioFlex 6-well culture plates with 0, 3, 5 and 10% MS for 12 h. As shown in Fig. 1A, MCP-1 expression in cells stimulated with 10% MS was markedly increased compared to those in the cells stimulated with 3% or 5% MS. The MCP-1 expression in 10% MS-stimulated cells was increased at 6 h of stimulation, and increased further up to 12 h of MS stimulation (Fig. 1B).

Increased HMGB1 release in VSMCs stimulated with MS

The effects of MS on HMGB1 secretion in VSMCs were examined by stimulating cells seeded onto BioFlex 6-well culture plates with 3 or 10% MS for 0–12 h. The levels of HMGB1 released into culture media of 10% MS-stimulated cells were significantly higher than those in 3% MS-exposed cells. In the cells stimulated with 10% MS, the secretion of HMGB1 was markedly increased at 6 to 12 h, but not in the cells stimulated with 3% MS (Fig. 2A). Since 10% strain showed the most substantial increase in HMGB1 secretion in this study, this level of strain was used for the subsequent experiments.

HMGB1 mediates MCP-1 expression in VSMCs stimulated with MS

The role of HMGB1 secreted into culture media on MCP-1 expression in MS-stimulated VSMCs was examined by pretreating the cells with 1, 3 and 10 nM glycyrrhizin, a HMGB1 inhibitor. The MCP-1 expression was determined in the cells stimulated with 10% MS for 12 h. In cells stimulated with 10% MS, MCP-1 expression was increased markedly compared to that in the control cells. The increased MCP-1 expression in VSMCs stimulated with 10% MS was attenuated by pretreatment with glycyrrhizin in a dose-dependent manner (Fig. 2B).

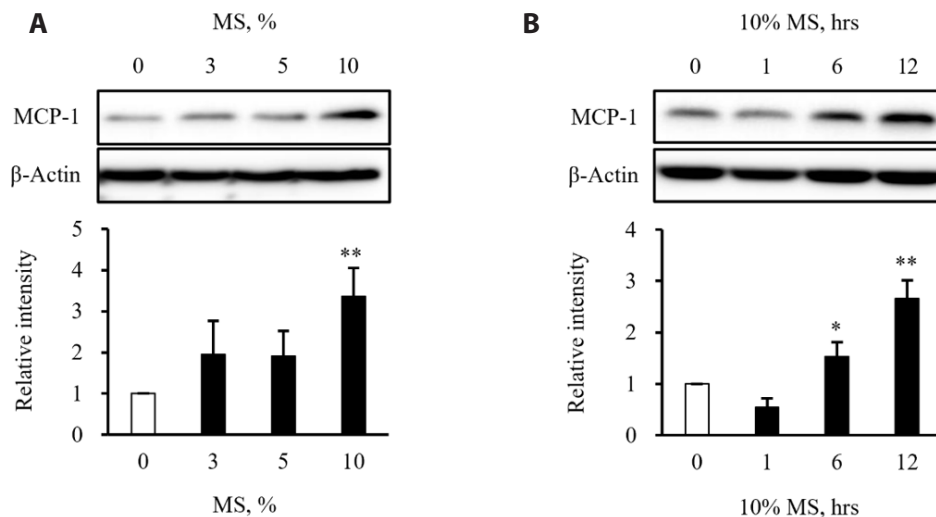


Fig. 1. Force- and time-dependent effects of mechanical stretch (MS) on monocyte chemoattractant protein 1 (MCP-1) expression in vascular smooth muscle cells (VSMCs). (A) Rat aortic VSMCs were stressed with MS (0, 3, 5 and 10% strain, 60 cycles/min) using Flexcell Tension Plus FX-4000T system for 12 h. MCP-1 expression in VSMCs was measured by Western blot, and the blots are representative of 5 independent experiments. β -Actin was used as an internal control. Quantitative results were expressed as the mean \pm SEM of 5 independent experiments. ** $p < 0.01$ vs. control. (B) The cultured VSMCs were stressed with 10% MS for 12 h, and MCP-1 expression in VSMCs was measured by Western blotting. β -Actin was used as an internal control. The blots are representative of 4 independent experiments. Quantitative results were expressed as the means \pm SEM of 4 independent experiments. * $p < 0.05$; ** $p < 0.01$ vs. control.

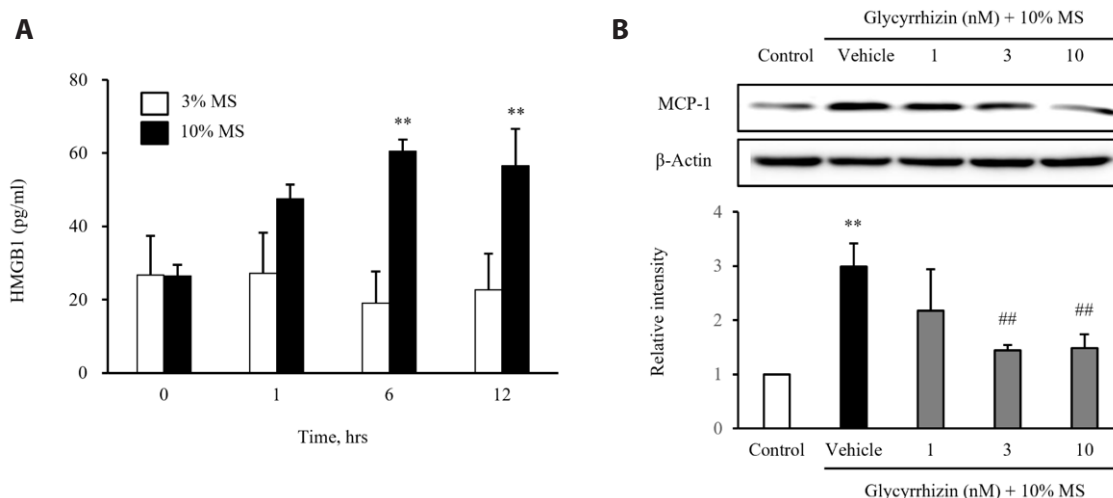


Fig. 2. High mobility group box 1 (HMGB1) release and its role on monocyte chemoattractant protein 1 (MCP-1) expression in vascular smooth muscle cells (VSMCs) stimulated with mechanical stretch (MS). (A) Force- and time-dependent effects of MS on HMGB1 release in VSMCs. Rat VSMCs were stimulated with 3% and 10% MS for 0–12 h, and then HMGB1 released into cell culture media was measured by ELISA. The concentration of HMGB1 released into culture media was quantified and expressed as the means \pm SEM of 4 independent experiments $**p < 0.01$ vs. corresponding value in the 3% MS group. (B) Role of HMGB1 on MCP-1 expression in VSMCs stimulated with MS. Rat VSMCs were pretreated with an inhibitor for HMGB1 (glycyrrhizin, 0–10 nM) for 24 h, and the cells were then stimulated with 10% MS for 12 h. The expression of the MCP-1 protein was determined by Western blot. β -Actin was used as an internal control. The blots are representative of 4 independent experiments. Relative intensities to β -actin were quantified and expressed as the means \pm SEM of 4 independent experiments. $**p < 0.01$ vs. control. $##p < 0.01$ vs. vehicle.

Individual role of PDGFR isoforms on MCP-1 expression in VSMCs stimulated with MS

The involvement of PDGFR signaling on MS-induced MCP-1 expression in VSMCs was examined by measuring the effects of MS on the expression of PDGFR isoforms. In this study, VSMCs were stimulated with 3% and 10% MS for 6 h, and then the expression of PDGFR- α and PDGFR- β was determined using Western blot. As shown in Fig. 3, PDGFR- α and PDGFR- β expression in 10% MS-stimulated cells were increased significantly compared to those in the cells stimulated with 0% or 3% MS.

To investigate the individual role of PDGFR isoforms on HMGB1 release in VSMCs stimulated with 10% MS, VSMCs deficient of PDGFR- α or PDGFR- β were produced by transfection of their specific siRNA, and then stimulated with 10% MS for 6 h. As shown in Fig. 4, the expression of PDGFR- α and PDGFR- β was markedly attenuated in cells transfected with their specific siRNA (200 nM) compared to cells transfected with the negative control. In the cells deficient of PDGFR- β , the increased release of HMGB1 in cells stimulated with 10% MS was markedly attenuated, but not in cells deficient of PDGFR- α . Similarly, MCP-1 expression in 10% MS-stimulated VSMCs was markedly attenuated in the cells deficient of PDGFR- β (Fig. 5).

DISCUSSION

The results of this study shows that MCP-1 expression and HMGB1 release were significantly higher in VSMCs exposed to

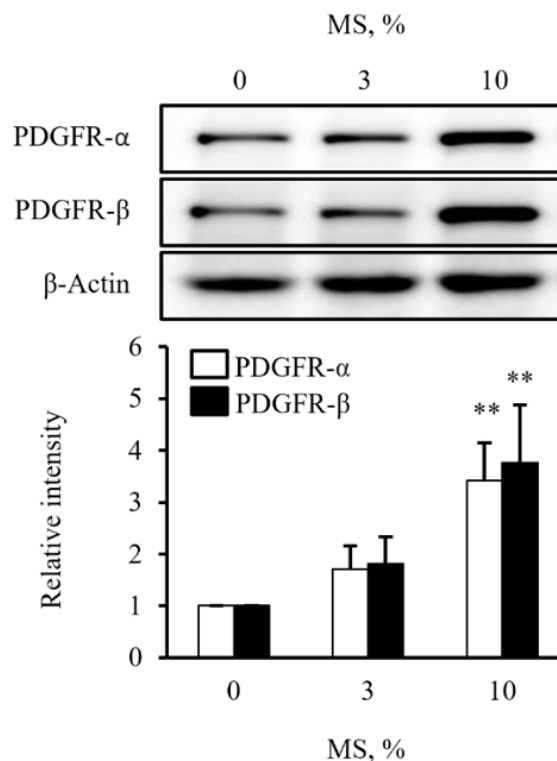


Fig. 3. Effects of mechanical stretch (MS) on PDGFR expression in vascular smooth muscle cells (VSMCs). The cultured rat VSMCs were stimulated with MS (3 and 10%) for 6 h, and then the expression levels of PDGFR- α and PDGFR- β were determined by Western blot. β -Actin was used as an internal control. The blots are representative of 5 independent experiments. Relative intensities to β -actin were quantified and expressed as the means \pm SEM of 5 independent experiments. $**p < 0.01$ vs. corresponding value in control.

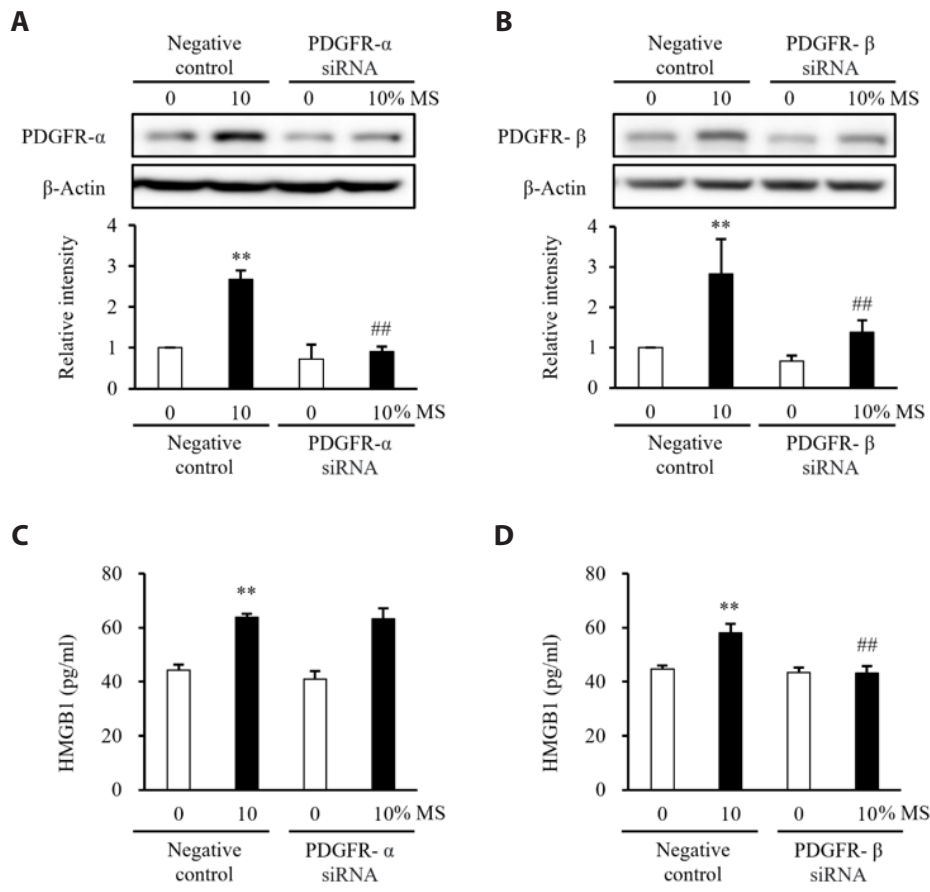


Fig. 4. Individual roles of the PDGFR isoforms on high mobility group box 1 (HMGB1) release in mechanical stretch (MS)-stimulated vascular smooth muscle cells (VSMCs). The cultured VSMCs were transfected with siRNAs (200 nM) for PDGFR- α (A) or PDGFR- β (B) for 24 h, and then stimulated with 10% MS for 6 h. The expression levels of PDGFR- α and PDGFR- β isoforms were determined by Western blotting using β -actin as an internal control. Relative intensities were quantified and expressed as the means \pm SEM of 4 independent experiments. ** $p < 0.01$ vs. control. ## $p < 0.01$ vs. corresponding value in negative control. (C, D) VSMCs transfected with PDGFR- α or PDGFR- β siRNA (200 nM) for 24 h were stimulated with 10% MS for 6 h. The HMGB1 released in the culture media were quantified by ELISA. The concentration of HMGB1 were quantified and expressed as the means \pm SEM of 4 independent experiments. ** $p < 0.01$ vs. control. ## $p < 0.01$ vs. corresponding value in negative control.

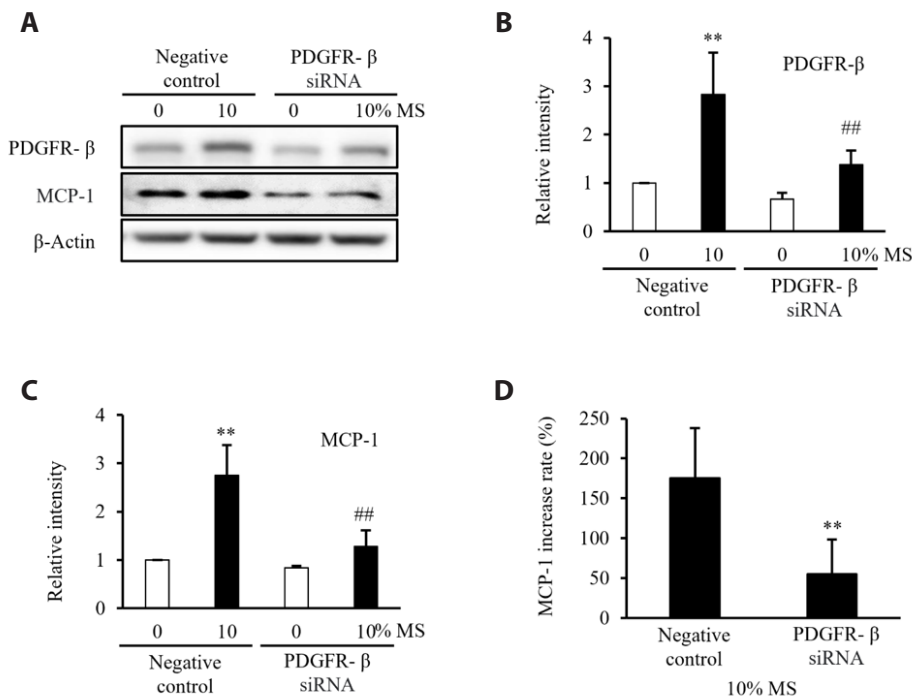


Fig. 5. Role of PDGFR- β signaling on monocyte chemoattractant protein 1 (MCP-1) expression in vascular smooth muscle cells (VSMCs) stressed with mechanical stretch (MS). The cultured VSMCs were transfected with PDGFR- β siRNA (200 nM) for 24 h, and then stimulated by 10% MS for 6 h. The expression levels of PDGFR- β and MCP-1 were determined by Western blotting using β -actin as an internal control. The blots are representative of 4 independent experiments (A). Relative intensities of PDGFR- β (B) and MCP-1 (C) to β -actin were quantified and expressed as the means \pm SEM of 4 independent experiments. ** $p < 0.01$ vs. control. ## $p < 0.01$ vs. corresponding value in negative control. (D) MCP-1 expression in 10% MS-stimulated cells was compared with that in control cells, and the relative increase rate was quantified. Data were expressed as the means \pm SEM of 4 independent experiments. ** $p < 0.01$ vs. negative control.

10% MS than to low-intensity MS (3% and 5%). These increases in MCP-1 expression and HMGB1 secretion in 10% MS-stimulated cells were significantly attenuated in the cells transfected with PDGFR- β siRNA, but not with PDGFR- α siRNA. In addition, the increased MCP-1 expression in 10% MS-stressed VSMCs was attenuated in the cells treated with glycyrrhizin, a HMGB1 inhibitor, suggesting the crucial role of PDGFR- β signaling on MCP-1 expression in VSMCs undergoing excessive biophysical stress via the increased secretion of HMGB1.

An increase in VSMC proliferation is an essential event inducing various vascular diseases including atherosclerosis and restenosis after angioplasty [15]. In response to various mediators secreted by inflammatory cells, VSMCs in the media move into the intima and then proliferate within the vascular lesions [5,16]. Therefore, vascular inflammation was suggested as a key event involved in the progression of vascular remodeling. In the injured vasculatures, the increased migration and infiltration of inflammatory cells into the damaged tissues were followed by an increased production of HMGB1, a major inflammatory mediator mediating various cardiovascular diseases [8,17,18].

To investigate cellular release of HMGB1 in VSMCs stressed with excessive biophysical stress, VSMCs cultured from rat thoracic aorta were seeded onto flexible-bottomed 6-well plates and then stimulated with equibiaxial cyclic stretch (0%–10% strain, 60 cycles/min). Considering the previous reports that 10% intensity of MS mimics hypertension [19,20] and the assumption that they serve as an *in vitro* model similar to hypertension [21], we stimulated VSMCs with 10% MS in this study. In an ELISA analysis, the concentration of HMGB1 released into culture media was increased markedly in VSMCs stimulated with 10% MS, but not in the cells stimulated with a low intensity (3% and 5%) of MS. Although the precise links between HMGB1 and vascular inflammation in the injured vasculatures have not been fully clarified in VSMCs, the results of this study suggests a pivotal role of HMGB1 in the progress of vascular complications associated with hypertension.

Among the numerous injury-induced mediators involved in the progression of cardiovascular diseases, HMGB1 is considered as a pivotal player [8,18,22]. A high level of HMGB1 was observed in atherosclerotic plaque, and was suggested to be involved in vascular remodeling by promoting inflammatory processes [8,23]. In line with the previous results indicating a pivotal role of HMGB1 in developing neointimal lesions following vascular injury [24,25], the cellular migration was increased markedly in VSMCs treated with HMGB1 in our previous study [26]. The results of the present study also shows that the levels of HMGB1 released into culture media were higher in VSMCs exposed to 10% MS than to low-intensity MS (3% and 5%). Based on these results, it was suggested that HMGB1 released in the injured vasculatures might be a key player in the development and progression of vascular remodeling.

In the process of vascular inflammation, MCP-1/CCR2 sig-

nalizing plays a major role in recruiting monocytes to the sites of tissue injury [27,28]. MCP-1 was reported to be one of the most important chemoattractants for monocytes, and enhances VSMC migration and proliferation in the process of vascular wall remodeling [29]. Increased MCP-1 expression has been detected in atherosclerotic lesions but not in normal arteries [30], suggesting a major role in vascular inflammation. Although MCP-1 is considered as an important factor that induces the inflammatory process in the injured vasculatures [31], the cellular source in the injured vasculatures is poorly understood.

The results of this study shows that MCP-1 expression was significantly higher in VSMCs stimulated with 10% MS than in cells stimulated with 3% and 5% MS. In addition, HMGB1 release into culture media were higher in VSMCs exposed to 10% MS than to low levels of MS, suggesting links between MCP-1 expression and HMGB1 release in VSMCs stressed with biophysical stretch. Therefore, this study measured MCP-1 expression in cells stimulated with 10% MS in the presence of HMGB1 inhibitor. The results clearly show that the increased expression of MCP-1 in VSMCs stimulated with 10% MS was attenuated in the cells pretreated with glycyrrhizin, an inhibitor for HMGB1, suggesting a pivotal involvement of HMGB1 in MCP-1 expression in VSMCs exposed to excessive biophysical stress.

PDGF synthesized in various cell types, including VSMCs, induces the proliferation and migration of VSMCs [32,33] and is suggested to be a major mediator in the progression of atherosclerosis and hypertension [4,21,34]. Among various growth factor receptors, PDGFR was suggested to be an essential signaling mediating vascular remodeling in hypertensive patients [35]. In line with our previous study showing that PDGFR- β signaling plays a vital role in cell migration in mechanically stressed VSMCs [36], the present study revealed the increased expression of PDGFR- α and PDGFR- β in VSMCs stressed with 10% MS compared to those in cells stressed with 3% MS, suggesting a potential role in vascular inflammation.

The relationship between PDGFR isoforms and MCP-1 expression in VSMCs exposed to biomechanical stress was examined by determining MCP-1 expression in the cells deficient of PDGFR isoforms. In the present study, the increased MCP-1 expression in 10% MS-stimulated cells was attenuated in cells transfected with PDGFR- β siRNA, but not with PDGFR- α siRNA, suggesting the critical involvement of PDGFR- β in MS-induced MCP-1 expression in VSMCs. In addition, the increased HMGB1 release in 10% MS-stressed VSMCs was attenuated in PDGFR- β -deficient cells, suggesting a key role of PDGFR- β signaling on MCP-1 expression in VSMCs undergoing excessive biophysical stress via an increased release of HMGB1.

Overall, the PDGFR- β signaling plays a crucial role in increasing the expression of MCP-1 in VSMCs stressed with 10% MS via the increased release of HMGB1. Thus, targeting the PDGFR- β -HMGB1-MCP-1 signaling axis in VSMCs undergoing excessive biophysical stress might be a promising therapeutic strategy for

vascular remodeling in the injured vasculatures under hypertensive conditions.

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CONFLICTS OF INTEREST

The authors declare no conflicts of interest.

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