Phenotypic Modulation of Mesenteric Vascular Smooth Muscle Cells from Type 2 Diabetic Rats is Associated with Decreased Caveolin-1 Expression

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Abstract
Aims: Diabetes-induced vascular complications are associated with vascular smooth muscle cell (VSMC) phenotypic modulation, switching from a contractile to a synthetic-proliferative phenotype. Loss of caveolin-1 is involved with proliferation of VSMCs. We tested the hypothesis that mesenteric VSMCs from type 2 diabetic Goto-Kakizaki (GK) rat undergo phenotypic modulation and it is linked to decreased caveolin-1 expression. Methods: VSMCs were isolated from mesenteric arteries from GK rats and age-matched control Wistar rats. Western blotting was used to determine expression of target proteins such as caveolin-1, calponin (marker of differentiation), and proliferating cell nuclear antigen (PCNA, marker of proliferation). In addition, we measured intracellular reactive oxygen species (ROS) production using H\textsubscript{2}DCF-DA and activation of extracellular signal-regulated kinase (ERK1/2) by western blotting in VSMCs from GK stimulated with lipopolysaccharide (LPS), an endotoxin upregulated in diabetes. Results: Mesenteric VSMCs from diabetic GK rats exhibited decreased caveolin-1 and calponin expression and increased PCNA expression compared to control. Increased levels of ROS and phospho-ERK1/2 expression were also found in GK VSMCs. LPS augmented ROS and phosphorylated ERK1/2 levels to a greater extent in GK VSMCs than in control. Likewise, high glucose decreased caveolin-1 and calponin expression, increased PCNA expression and augmented ROS production in control mesenteric VSMCs. Conclusion: These results suggest that mesenteric VSMCs from diabetic GK rats undergo phenotypic modulation and it is associated with decreased caveolin-1 expression. These alterations may be due to enhanced inflammatory stimuli and glucose levels present in diabetic milieu.
Introduction

Vascular smooth muscle cells (VSMCs) from adult blood vessels display a highly specialized and differentiated phenotype. Its central function is contraction and control of vessel diameter to regulate blood flow distribution and blood pressure [1, 2]. VSMCs in native contractile phenotype express markers of differentiation such as calponin, one of the essential proteins of the contractile apparatus in VSMCs [2, 3]. VSMCs exhibit remarkable plasticity switching from a contractile to a proliferative phenotype characterized by decreased expression of contractile proteins and increased proliferation [2, 4]. In the vascular environment, mechanical and chemical changes lead to a rapid and reversible phenotypic modulation of VSMCs in order to maintain vascular homeostasis. It is well-established that phenotypic modulation of VSMCs is involved in the pathogenesis of atherosclerosis, one of the complications found in type 2 diabetes mellitus (T2DM) [5].

Emerging findings have revealed that patients with T2DM exhibit elevated circulatory levels of LPS [6]. LPS is an endotoxin known to stimulate reactive oxygen species (ROS) production in vascular cells [7], which is involved in the phenotypic modulation of VSMCs [8]. T2DM often is associated with obesity and hypertension and exhibits low-grade inflammatory states [9]. These complications challenge the understanding of onset of vasculopathy in T2DM. Goto-Kakizaki (GK) rats are a suitable model of nonobese T2DM [10]. We and others investigators reported that GK rats display impaired relaxation and increased contraction in various vascular beds [11, 12]. Remodeling in resistance arteries and cerebral arteries was seen in GK rats [13-15]. Recently, Chettimada et al. [16] found that ROS lead to vascular wall remodeling and dysfunction by changing contractile protein expression in aortas from GK rats. To date, it is unknown whether or not VSMCs from mesenteric resistance arteries of GK rats display phenotypic modulation.

Caveolin, a 21-24 kDa integral membrane protein, consists of a family of at least three different isoforms: caveolin-1, -2, and -3, which function in the organization of caveolae and in signal-transduction [17, 18]. Caveolin-1 has been implicated in multiple functions such as cell-cycle arrest as well as senescence, prevention of cell transformation, and anti-proliferative effects [17-19]. Changes of caveolin-1 expression are associated with pathophysiological conditions [20, 21]. For example, decreased caveolin-1 expression was correlated with increased cell cycle entry in VSMCs from wild type mice and over-expression of caveolin-1 inhibited proliferation of VSMCs [22]. Therefore, caveolin-1 may play a role in the development of VSMCs phenotypic changes. Thus far, association between caveolin-1 and phenotypic changes in VSMCs in diabetes has not been reported.

In the present study, we hypothesized that mesenteric arterial VSMCs isolated from GK rats undergo phenotypic modulation and it is associated with changes in caveolin-1 expression and ROS levels.

Materials and Methods

Animals and Experimental Procedures

Twenty weeks-old male Wistar and GK rats were maintained on a 12-hour light and dark cycle with access to standard rodent chow and water ad libitum. The presence of diabetes was confirmed by measuring blood glucose levels using a One Touch Ultra Glucometer (Monitoring System Abbott). All experimental protocols were approved by the Institutional Animal Care and Use Committee of Georgia Regents University and conducted in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals. The mean plasma glucose levels in control and GK rats were 94.1±11 and 151.5±7 mg/dl, respectively. We previously reported mean arterial blood pressure to be comparable in control (103±5 mmHg) and in GK (103±4 mmHg) rats as measured by telemetry at this age [23].

Cell Cultures

VSMCs were isolated from rat mesenteric arteries by the explant method, as previously described [24]. Briefly, second and third order mesenteric arteries were cleaned of adipose and connective tissues.
Branches of mesenteric arteries were placed in a culture dish and were maintained in Dulbecco Modified Eagle's Medium (DMEM) containing 10% fetal bovine serum (FBS) and antibiotics in a humidified incubator at 37°C, 5% CO₂, and atmospheric O₂. After 48 hours, mesenteric arteries were removed from the culture dish and cells attached to plate were maintained in cultures. VSMCs exhibited the typical “hill and valley” growth morphology and were confirmed positive (>95%) for smooth muscle α-actin. Cells at early passage (Passage 3) were used in all experiments. After 80% of confluence, VSMCs were serum starved for 24 hours in order to reach a quiescent state, followed by treatments with high glucose (HG, 25 mM) for 24, 48, and 72 hours or with normal glucose (NG, 5 mM) for 24 hours. To investigate the effect of HG on ROS generation, quiescent mesenteric VSMCs were treated with HG or NG for 12 hours. In another set of experiments, to measure ROS generation and ERK1/2 activation, quiescent cells were stimulated with LPS (100 ng/mL) for 15 minutes [25, 26].

Western blotting

Western blotting was performed to determine caveolin-1, proliferating cell number antigen (PCNA), calponin, phosho-extracellular signal-regulated kinase (ERK1/2), total ERK1/2 and β-actin protein expression. Briefly, 10 µg of cellular protein was resolved by 10% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), transferred to polyvinylidene difluoride (PVDF) membranes (Thermo Scientific, Rockford, IL, USA) as previously described [24] and probed with specific antibodies. Primary antibodies were follows: caveolin-1 (1:1,000) (BD Biosciences, San Jose, CA, USA); PCNA (1:1,000) (Abcam, Cambridge, UK); calponin (1:2,000) (Santa Cruz Biotechnology, Santa Cruz, CA, USA); phospho-ERK1/2 (1:1,000) (Cell Signaling Technology, Danvers, MA); total ERK1/2 (1:1,000) (Cell Signaling Technology, Danvers, MA). The stripped membranes were later probed for β-actin (1:20,000) (Sigma Aldrich, St. Louis, MO, USA) antibody as a loading control. Data are presented as fold-induction normalized to β-actin.

Measurement of reactive oxygen species (ROS)

ROS production was measured in VSMCs by 2,7-dichlorofluorescein diacetate (H2DCF-DA), as previously described [27]. Quiescent mesenteric VSMCs from GK and Wistar rats were incubated in the dark with H2DCF-DA (10 µM) for 30 minutes, followed by stimulation with LPS (100 ng/mL) for 15 min. Additionally, quiescent mesenteric VSMCs from control rats were treated with HG or NG for 12 hours then intracellular ROS were measured. A set of ten images per each field were acquired at 20x magnification using a fluorescence microscopy (Axiovert 200) fitted with a camera. Then, semi-quantitative analyses were performed to detect changes in H2DCF-DA fluorescence in living VSMCs using Image Pro Plus software.

Statistical Analysis

Data are expressed as mean ± SD. Data were analyzed with Student’s t test or 1-way analysis of variance (ANOVA) followed by using Tukey post hoc test. p<0.05 was considered statistically significant.

Results

Morphology of mesenteric VSMCs from GK and Wistar rats

When isolated in cultures, mesenteric VSMCs from control Wistar rats exhibited spindle shape (Fig. 1A) and organized arrangement at confluence (Fig.1B). Conversely, mesenteric VSMCs from GK rats lost the typical spindle shape and at confluence and displayed disarrangement (Fig. 1C and D, respectively).

Expression of caveolin-1, PCNA, and calponin in mesenteric VSMCs from GK and Wistar rats

As shown in Figs. 2A & 2B, decreased protein expression of caveolin-1 was found in mesenteric VSMCs from the GK group compared to the Wistar group. To evaluate whether mesenteric VSMCs undergo phenotypic changes, we measured protein expression of PCNA, a marker of cell proliferation [28-30] (Figs. 2C & 2D) and calponin (Figs. 2E & 2F). The expression of PCNA was increased whereas the expression of calponin was decreased in GK group (vs. Wistar).
Since phenotypic modulation of VSMC is associated with ROS production [1] and ERK1/2 activation [3], we next examined whether LPS, which is found in elevated levels in diabetic subjects, causes alterations in ROS production and ERK1/2 activation in mesenteric VSMCs from control Wistar and diabetic GK rats.
VSMCs as shown in Figs. 3A and 3B. At baseline, mesenteric VSMCs from GK exhibit higher levels of ROS than mesenteric VSMCs from control rats. LPS treatment increased ROS fluorescence detection greater in mesenteric VSMCs from GK rats than in those from Wistar rats. Additionally, elevated levels of phosphorylated ERK were found in mesenteric VSMCs from GK rats and it was exacerbated after LPS treatment (Figs. 3C and D).

**Effect of high glucose on caveolin-1, differentiation and proliferation in mesenteric VSMCs**

High blood glucose levels are the hallmark of diabetes. In order to determine whether HG has direct effect on VSMCs phenotype and caveolin-1 expression, we evaluated caveolin-1 and phenotype-related proteins (PCNA and calponin) in mesenteric VSMCs isolated from control Wistar rats treated with HG (25 mM) for 24 (caveolin-1) and 24 to 72 (PCNA and calponin) hours. VSMCs treated with HG for 24 h significantly reduced caveolin-1 expression compared to the control (treated with 5 mM glucose) (Figs. 4A & B). Furthermore, HG significantly increased PCNA (Figs. 4C & D) and decreased calponin (Figs. 4C & E) expression in mesenteric VSMCs in a time-dependent manner.
To determine whether increased ROS levels found in mesenteric VSMCs from GK may arise from hyperglycemia, we measured ROS in mesenteric VSMCs isolated from Wistar rats. Quiescent mesenteric VSMCs were treated with high glucose (HG, 25 mM) for 24 hours (A, B) or for 24, 48, 72 hours (C, D, E) and normal glucose (NG) (control, 5 mM) for 24 hours and protein expressions were measured by western blotting. (A, C) Representative immunoblots showing caveolin-1 (cav-1) (A), PCNA (C), calponin (C) and β-actin. (B, D, E) Bar graph showing relative specific bands densities of proteins to β-actin. Values (mean ± SD, n=4-5) are expressed as fold of control NG group. *P<0.05 vs. NG (control).

**Fig. 5.** Effect of high glucose on ROS generation in mesenteric VSMCs. Quiescent mesenteric VSMCs from control rats were treated with high glucose (HG, 25 mM) or normal glucose (NG) (control, 5 mM) for 12 hours and intracellular ROS was measured by H2DCF-DA. (A) Photomicrographs showing changes in fluorescence (white) indicative of intracellular ROS production and relative changes in fluorescence intensity were quantified and normalized by number of cells (DAPI stain nucleus). (B) Data are shown as mean ± SD, n=3. *P<0.05 vs. NG.

**ROS production in response to high glucose in mesenteric VSMCs**

To determine whether increased ROS levels found in mesenteric VSMCs from GK may arise from hyperglycemia, we measured ROS in mesenteric VSMCs isolated from Wistar rats.
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Discussion

Phenotypic modulation of VSMCs plays a crucial role in vascular dysfunction in T2DM; however, the mechanisms involved have not been well explored.

Although it has been difficult to mimic symptoms of diabetes, significant advances in treatment for diabetes may come from studies of suitable animal models [32, 33]. A number of T2DM animal models exhibit metabolic syndrome other than DM itself including hypertension, hyperlipidemia, or obesity [32, 33]. Thus, it may be difficult to clarify the relative role of each of these confounding factors in the development of DM-associated vascular dysfunction in these animal models. GK rats are a relatively unique strain as a T2DM model which does not develop obesity [10]. Studies have demonstrated that vascular dysfunction in GK rats comprise large and small arteries [11, 12] and remodeling process [13-15]. However, the impact of diabetes on caveolin-1, phenotype interactions and changes in the signaling pathways in resistance arterial VSMCs in this model remain unknown.

The main finding of this study is that mesenteric VSMCs from rats with T2DM undergo phenotypic modulation and it is associated with decreased levels of caveolin-1 in parallel with enhanced levels of ROS (Fig. 6). This was supported by our results showing that GK VSMCs exhibited increased proliferation, reduced contractile protein content and augmented ROS generation.

Caveolin-1 plays pivotal roles in physiological and pathophysiological conditions [17-22]. It has been demonstrated that deletion of caveolin-1 promotes VSMCs proliferation increasing intimal hyperplasia upon carotid injury [34, 35]. Additionally, reduced caveolin-1 levels have been reported for plaques from hypercholesterolemic rabbits and humans [22, 36]. Lower caveolin-1 levels were also associated with features of plaque instability [37]. Therefore, caveolin-1 exerts an anti-atherosclerotic effect and may control function in VSMCs in disease conditions. In the present study, caveolin-1 and calponin were down-regulated while PCNA was up-regulated in VSMCs from diabetic resistance arteries. These results suggest that the reduction of caveolin-1 expression might be closely associated with phenotypic change in GK mesenteric VSMCs.

High levels of glucose are the hallmark of diabetes. Exposure of VSMCs with HG results in increased ROS production and inflammatory agents [28, 31, 38]. In the present study, HG-treated VSMCs caused early-elevated ROS production and down-regulation of caveolin-1, which preceded alterations of phenotype-related proteins (viz. PCNA up-regulation and calponin down-regulation). Therefore, we demonstrated that HG directly leads to modulation of caveolin-1 and phenotype-related protein expressions in mesenteric VSMCs. This conclusion is supported by the evidence showing that HG leads to a change in expression of caveolin-1 (down-regulation) in lens epithelial cells [21] and PCNA (up-regulation) in rat aortic

stimulated with HG for 12 hours. As shown in Fig. 5A & 5B, production of ROS was increased by HG in mesenteric VSMCs, in accordance with previous findings in the literature [28, 31].
VSMCs [27]. Moreover, several studies have reported that ROS and ERK1/2 are key players in the phenotypic changes observed in VSMCs [3, 5, 28, 31, 38]. ERK1/2 activation has been shown to be downstream of ROS release and previous studies demonstrated that phenotypic modulation of VSMCs is associated with ERK1/2 activation [3, 30]. Indeed, in our present study, baseline ROS production and ERK activation were increased in mesenteric VSMCs isolated from GK rats. LPS led to further ROS generation and ERK activation, which was more pronounced in GK cells. Taking our present data and other relevant evidence together, HG exposure and/or chronic inflammatory stimuli accompanied with hyperglycemia might be causal factors of caveolin-1 down-regulation and phenotypic modulation in VSMCs and these alterations were associated with ROS generation and ERK1/2 activation (Fig. 6). Although the precise mechanisms underlying the alterations of phenotype-related proteins in GK rat mesenteric VSMCs are not certain, our data suggests that these alterations might be, at least in part, due to down-regulation of caveolin-1. Future experiments will be needed to address how down-regulation of caveolin-1 play role in increased ROS levels and ERK1/2 activation.

These findings bring a new perspective to diabetic research which has been investigating alternative therapeutic targets to treat diabetic vascular complications since the actual anti-diabetic drugs do not prevent the development of vascular disorders in diabetic patients.

In summary, we found that the expression of caveolin-1 was decreased in the GK mesenteric VSMCs which might be associated with phenotypic change. These alterations of GK mesenteric VSMCs are possibly attributable to increased susceptibility to inflammatory stimuli and chronic exposure of HG levels which is characteristic of diabetes. Clarifying the signal-transduction and control of caveolin-1 expression is needed to understand the pathophysiology of diabetic vascular complications, as well as advance the treatment of T2DM-associated vascular complications.

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Disclosure Statement

The authors have no conflict of interest.

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