Effect of High Glucose on Basal Intracellular Calcium Regulation in Rat Mesangial Cell

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Abstract

Background: A number of cellular mechanisms are critically dependent on intracellular Ca2+ homeostasis. A sustained increase in the intracellular Ca2+ concentration ([Ca2+]i) is capable of activating a number of potentially harmful processes including phenotype change to secretory type, dysregulated cell proliferation, and cell injury and death. Mesangial cells (MCs) play an important role in the pathophysiology of diabetic nephropathy.

Methods: We evaluated the effect of high glucose on basal [Ca2+]i in the unstimulated state and identified its contributing pathways. MCs were isolated and cultured from Sprague-Dawley rats. [Ca2+]i, was measured by fluoroimetric technique with fura-2AM. Results: In a dose-dependent manner, superfusion of MCs with Tyrode’s solution containing high glucose (30 and 50 mM) induced a delayed spontaneous increase in [Ca2+]i, which was not found in those with normal (5.5 mM) glucose or mannitol. The high glucose-induced increase in [Ca2+]i occurred through transmembrane influx of extracellular Ca2+ and was blocked by SKF96365, an inhibitor of store-operated Ca2+ influx. Na+-Ca2+ exchanger (NCX) activity, a major channel regulating basal [Ca2+]i, and the clearing ability of intracellular Ca2+ were depressed after MCs were cultured in high-glucose medium. Western blot analysis revealed the decreased expression of a 70-kD NCX protein in MCs cultured in high-glucose medium.

Conclusions: A high-glucose concentration induced a spontaneous increase in basal [Ca2+]i of MCs without stimulation. There was a decrease in the activity of NCX in the high-glucose condition, which seems to occur at the level of protein expression. The present results provide a novel insight into the mechanisms of diabetic nephropathy in that intracellular Ca2+ homeostasis is an important secondary messenger and a mediator in hormonal signaling.
Cytosolic calcium (Ca\(^{2+}\)) activity, or intracellular Ca\(^{2+}\) concentration ([Ca\(^{2+}\)]), is an important biological signal in the control of protein secretion, contraction, development, and apoptosis in a wide variety of cells. Many cellular functions critically depend on [Ca\(^{2+}\)], [4–6] and a sustained elevation in [Ca\(^{2+}\)], is capable of activating a number of potentially harmful cellular processes including phenotypic change, dysregulated cell proliferation, cell injury, apoptosis, and death [6–8].

Several studies have been done on Ca\(^{2+}\) regulation in MCs. Most of these studies focused on the [Ca\(^{2+}\)] responses of MCs to vasoactive hormones, such as angiotensin II, arginine vasopressin, thromboxane analogues, which are related to the contractile properties of MCs. Loss of the contractile responsiveness to vasoactive hormones in MCs is related to hyperfiltration and microalbuminuria in the early course of diabetic nephropathy [9, 10]. Recent investigations revealed that the alteration in vasoconstrictor-induced Ca\(^{2+}\) signaling by high glucose contributes to this phenomenon [11–13].

However, these findings are related solely to the early part of the pathophysiological process in diabetic nephropathy. As the disease progresses, a variety of cellular dysfunctions in the MCs incorporate into the whole complex feature of diabetic nephropathy, such as the phenotypic change to the secretory and proinflammatory cells, an abnormal cell cycle, or cell injury and death. Since the change in [Ca\(^{2+}\)], is comprehensively related to these pathophysiological processes [6], we have focused on the basal intracellular Ca\(^{2+}\) homeostasis of the MCs in a condition without any stimulation by vasoactive hormones, which has rarely been investigated in MCs. The aims of the present study are to evaluate the effect of high glucose on basal [Ca\(^{2+}\)], of MCs, and to identify the contributing Ca\(^{2+}\) pathways, and to focus on Na\(^{+}\)-Ca\(^{2+}\) exchanger (NCX) function.

**Materials and Methods**

**Isolation and Cell Culture of Rat Mesangial Cells**

The kidneys of Sprague-Dawley rats (80–100 g body weight) were removed and homogenized under sterile conditions. The glomeruli were isolated from the homogenate by sequential sieving and collected on a 75-µm sieve. After incubation with 0.25% trypsin and 0.02% EDTA in PBS (pH 7.4) for 30 min, the glomeruli were plated in tissue culture dishes containing Dulbecco’s modified Eagle’s medium/F-12 nutrient mixture (DMEM/F-12) supplemented with 5.5 mM glucose, 20% heat-inactivated fetal bovine serum, 100 U/ml penicillin, and 100 µg/ml streptomycin. To characterize cell outgrowths, immunohistochemical staining was performed for vimentin, desmin, and cytokeratin. To evaluate the basal [Ca\(^{2+}\)], change and transmembrane Ca\(^{2+}\) flux, superfusion with various experimental solutions was performed on the cells at subconfluence or at confluence with 0.05% trypsin/0.02% EDTA. Cells between 5 and 10 passages were used. To obtain quiescent cells, MCs were maintained in DMEM/F-12 media supplemented with 0.5% fetal bovine serum for 24 h. To evaluate the change in function and protein expression of NCX, subconfluent quiescent MCs were further incubated in DMEM/F-12 media containing 5.5 or 30 mM glucose or 30 mM mannitol for 3 days.

**Experimental Solutions**

The changes in basal [Ca\(^{2+}\)], and transmembrane Ca\(^{2+}\) flux were examined by superfusion of the solutions containing variable glucose concentrations. Tyrode’s solution was used as basal solution. The composition of Tyrode’s solution in the present study was as follows: 5 mM HEPES, 140 mM NaCl, 4 mM KCl, 2 mM CaCl\(_2\), 1 mM MgCl\(_2\), 1 mM NaH\(_2\)PO\(_4\), 5 mM HEPES, and 5.5 mM glucose (pH 7.4). To evaluate the effect of the glucose concentration, the glucose concentration was increased to 30 and 55 mM. Mannitol, replacing glucose at equivalent osmolarities, was used as an osmotic control. These solutions, containing 5.5, 30, or 55 mM glucose or 30 or 55 mM mannitol, were superfused at a flow rate of 2 ml/min. The measurement of [Ca\(^{2+}\)] during superfusion is described below.

In order to investigate the source of Ca\(^{2+}\), we superfused 30 and 55 mM glucose Tyrode’s solution with the removal of Na\(^{+}\). This nominally 0 Ca\(^{2+}\) solution was prepared by chelating Ca\(^{2+}\) with 1 mM EGTA from Tyrode’s solution. Next, we investigated the role of store-operated Ca\(^{2+}\) influx (SOCCI) using its inhibitor, SKF96365 (Tocris, Bristol, UK). In the preliminary experiment, the inhibitory effect was maximal at a dose of 50 µM in our cells (refer to the result in fig. 3B, SKF96365 at this dose was added to 30 and 55 mM glucose Tyrode’s solution containing 2 mM Ca\(^{2+}\) and superfused.

NCX function and expression were evaluated using MCs previously cultured in 5.5 or 30 mM glucose or 30 mM mannitol for 3 days. NCX activity was observed by activating it in the reverse mode by acute removal of extracellular Na\(^{+}\). Na\(^{+}\) was removed from Tyrode’s solution with isotonic replacement of N-methyl-D-glucamine. This so-called Na\(^{+}\)-free solution was superfused and the intracellular Ca\(^{2+}\) spike representing NCX activity was measured.

We evaluated Ca\(^{2+}\) clearance via NCX. We first elevated intracellular Ca\(^{2+}\) by activating NCX in the forward mode with Na\(^{+}\)-free solution. When MCs achieved a sufficient level of [Ca\(^{2+}\)], Ca\(^{2+}\) channels other than NCX were then blocked by adding 5 mM caffeine, 1 µM thapsigargin (Sigma Chemical Co., St Louis, Mo., USA), and 250 µM La\(^{3+}\). Under this condition, Na\(^{+}\) was acutely increased to 140 mM in the superfusion solution to elicit NCX in forward mode causing Ca\(^{2+}\) removal from the intracellular space.

**Fig. 1.** Experiments show a typical response of mesangial cells to angiotensin II (AT-II) in 5.5 mM glucose (A). Superfusion with 30 and 50 mM glucose increases [Ca\(^{2+}\)], spontaneously (C, D). The increases occurred in a dose-dependent manner to the glucose concentration. [Ca\(^{2+}\)], is not changed by superfusion with 5.5 mM glucose (B) and 30 and 50 mM mannitol (E, F). Each fluorescence tracing is a representative of: 6 experiments (A), 21 experiments (B), 18 experiments (C), 16 experiments (D), 20 experiments (E), and 19 experiments (F).
Basal [Ca^{2+}]i Regulation in Rat Mesangial Cell

A. 1 µM AT-II

B. 5.5 mM glucose

C. 30 mM glucose

D. 55 mM glucose

E. 30 mM mannitol

F. 55 mM mannitol
Fig. 2. The increases in \([\text{Ca}^{2+}]_i\) are significant at 1,500 s in MCs superfused with 50 mM glucose. At 3,500 s, \([\text{Ca}^{2+}]_i\) in MCs superfused with 30 mM glucose also has a significant increase. \([\text{Ca}^{2+}]_i\) is higher in MCs superfused with 55 mM glucose as compared to those with 30 mM glucose. * \(p < 0.05\) to 5.5, 30 mM glucose and 30, 55 mM mannitol. † \(p < 0.05\) to 55 mM glucose at 500 and 1,500 s. + \(p < 0.05\) to 5.5, 50 mM glucose and 30, 55 mM mannitol. Each column is representative of: 21 experiments with 5.5 mM glucose; 18 experiments with 30 mM glucose; 16 experiments with 55 mM glucose; 20 experiments with 30 mM mannitol, and 19 experiments with 55 mM mannitol.

**Intracellular Calcium Measurement**

MCs were washed with PBS and incubated in 2 ml of buffer (0.05% trypsin and 0.02% EDTA). They were then resuspended in Tyrode’s solution containing 5 mM glucose and loaded with 3 \(\mu\)M fura-2 AM (Molecular Probe, Eugene, Oreg., USA), a \(\text{Ca}^{2+}\)-sensitive intracellular probe, for 30 min at 37 °C. After loading, the MCs were washed with Tyrode’s solution, spun down for 5 min at 580 g. They were resuspended in Tyrode’s solution and transferred to a recording chamber on an epifluorescence-inverted microscope (Nikon Diaphot 300, Tokyo, Japan). During the superfusion of experimental solutions as described above, fluorescence was measured using a cooled CCD camera (Photometrics PXL37, Tucson, Ariz., USA) and cellular \(\text{Ca}^{2+}\) imaging was processed using the Axon Imaging Workbench v.2.2 (Axon Instrument, Foster City, Calif., USA). The \([\text{Ca}^{2+}]_i\) is presented as \(R_{340/380}\), a ratio of fluorescence intensities excited by alternating illumination of 340- and 380-nm wavelength beams with emission collected at 500 nm.

**Western Blot Analysis**

After being cultured in media containing 5.5 or 30 mM glucose or 30 mM mannitol for 3 days, MCs were homogenized in 4 ml of a buffer solution containing 50 mM Tris/HCl (pH 7.5), 2 mM EDTA, 10 mM EGTA, 5 mM dithiothreitol, 250 mM sucrose and 1% Triton X-100 and the protease inhibitors (0.5 \(\mu\)g/ml leupeptin, 5 \(\mu\)g/ml peptatin, 1 mM PMSF, 1 mM benzamidine, 2 mM iodoacetamide, and 1 \(\mu\)g/ml aprotinin). The homogenate was then centrifuged at 10,000 g for 15 min, and the resulting supernatant was centrifuged at 100,000 g for 90 min. The crude microsomal pellet was resuspended in the same buffer as described above, frozen in liquid nitrogen, and stored at −70°C. Protein concentrations were determined by the Lowry method. Protein samples were separated by SDS/PAGE (8–16% gradient gel) and probed with rabbit anti-mouse sodium-calci
cion exchange monoclonal antibody (R3F1; Swant, Bellinzona, Switzerland). Bands were stained with R3F1 and developed by enhanced chemiluminescence (Amersham Pharmacia Biotech AB, Uppsala, Sweden) using horseradish peroxidase-conjugated goat anti-rabbit IgG and enhanced chemiluminescence. Each lane was loaded with 30 \(\mu\)g of total protein.

**Statistical Analysis**

Each fluorescence tracing is representative of the indicated number of experiments conducted on a cell from separated cell cultures. \(R_{\text{max}}\), the slope of \(\text{Ca}^{2+}\) clearance (\(R_{\text{max}}/\text{min}\)) and NCX protein expression (arbitrary units) are reported as the mean ± SE for the indicated number of experiments. Statistical differences among the data were determined with Student’s t test and considered significant at \(p < 0.05\).

**Results**

**Characteristics of Cells**

The cell studies showed the typical characteristics of the MCs: positive immunohistochemical staining for vimentin and desmin and negative staining for cytokeratin. They showed the typical characteristics of the \([\text{Ca}^{2+}]_i\) of MCs in response to angiotensin II, which was identical to the previous studies (fig. 1A).

**Change in Basal \([\text{Ca}^{2+}]_i\), and Role of Transmembrane Influx and SOCI**

Superfusion of the MCs with Tyrode’s solution containing 30 or 55 mM glucose increased \([\text{Ca}^{2+}]_i\), spontaneously. The increase was not found with a normal glucose concentration (5.5 mM) or mannitol at equivalent osmolarity (30 and 55 mM; fig. 1B–F). The increases started at about 3,000–3,500 s after superfusion with 30 mM glucose and at 1,000–1,500 s with 55 mM glucose.
[Ca\textsuperscript{2+}], increased significantly in the MCs superfused with 55 mM glucose at 1,500 s. At 3,500 s, the [Ca\textsuperscript{2+}], of the MCs superfused with 30 mM glucose increased significantly (fig. 2). The [Ca\textsuperscript{2+}], at 3,500 s was higher with 55 mM glucose (2.32 ± 0.46) than with 30 mM glucose (1.87 ± 0.51; p < 0.05).

The removal of extracellular Ca\textsuperscript{2+} by chelation with 0.1 mM EGTA completely prevented the increase in [Ca\textsuperscript{2+}], both with 30 and 55 mM glucose (fig. 3A). This suggests that the high-glucose-induced increase in [Ca\textsuperscript{2+}], originates from the transmembrane influx but not from intracellular Ca\textsuperscript{2+} stores. The addition of SKF96365 to the normal Ca\textsuperscript{2+} solution also completely blocked the high-glucose-induced increase in [Ca\textsuperscript{2+}],. The result suggests that transmembrane influx occurs through SOCI (fig. 3C).
Fig. 4. Ca²⁺ spike representing Na⁺-Ca²⁺ exchanger activity is normally elicited by acute removal of extracellular Na⁺ in MCs cultured in 5.5 mM glucose medium (A) and 30 mM mannitol medium (C). The elicitation of Na⁺-Ca²⁺ exchanger activity did not occur in MCs cultured in 30 mM glucose medium (B). Peak [Ca²⁺] (Rmax) is significantly lower in MCs cultured in 30 mM glucose than in 5.5 mM glucose and 30 mM mannitol (D). Each fluorescence tracing and column is representative of: 32 experiments with 5.5 mM glucose; 37 experiments with 30 mM glucose, and 24 experiments with 30 mM mannitol.

The Activity, Ca²⁺ Clearance Ability, and Expression of Na⁺-Ca²⁺ Exchanger

When NCX was elicited in the reverse mode using Na⁺-free Tyrode’s solution, the Ca²⁺ spike that represents NCX activity was markedly depressed in MCs cultured in 30 mM glucose medium. This was not found in MCs cultured in 5.5 mM glucose medium (fig. 4). Peak [Ca²⁺], following NCX activation (Rmax) was 1.37 ± 0.06 in MCs cultured in 30 mM glucose, which was significantly lower than 2.61 ± 0.23 and 2.53 ± 0.20 in MCs cultured in 5.5 mM glucose and 30 mM mannitol respectively (p < 0.05; fig. 4D).

As described in Methods, we elevated intracellular Ca²⁺ using Na⁺-free solution to evaluate intracellular Ca²⁺
clearance. In this step, a sufficient intracellular Ca\(^{2+}\) level could be achieved in 20 of 21 (95.2%), 19 of 32 (59.4%), and 21 of 24 (87.5%) MCs cultured in 5.5 or 30 mM glucose, and 30 mM mannitol medium respectively. Ca\(^{2+}\) clearance was evaluated in these cells. As a result, Ca\(^{2+}\) clearance was decreased in MCs cultured in 30 mM glucose as compared to those cultured in 5.5 mM glucose or 30 mM mannitol (fig. 5). The slope of Ca\(^{2+}\) clearance (R/min) was –0.027 ± 0.004 in MCs cultured in 30 mM glucose, which was significantly lower than –0.076 ± 0.009 and –0.061 ± 0.006 in MCs cultured in 5.5 mM glucose and 30 mM mannitol, respectively (p < 0.05; fig. 5 D). In Western blot analysis, the protein expression of 70 kD NCX protein was decreased in MCs cultured

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Fig. 5. Ca\(^{2+}\) clearance via Na\(^{+}\)-Ca\(^{2+}\) exchanger is normal in MCs cultured in 5.5 mM glucose (A) and 30 mM mannitol medium (C), but decreased in MCs cultured in 30 mM glucose medium (B). The slope of Ca\(^{2+}\) clearance (R/min) is significantly lower in MCs cultured in 30 mM glucose than in 5.5 mM glucose and 30 mM mannitol (D). Each fluorescence tracing and column is a representative of: 20 experiments with 5.5 mM glucose; 19 experiments with 30 mM glucose, and 21 experiments with 30 mM mannitol.
Fig. 6. MCs cultured in high glucose show decreased expression of 70 kDa Na\(^+\)-Ca\(^{2+}\) exchanger (A). Densitometric analysis revealed that the expression in MCs cultured in 30 mM glucose is significantly depressed as compared to MCs in 5.5 mM glucose and 30 mM mannitol medium (B). Each fluorescence tracing and column is a representative of: 7 experiments with 5.5 mM glucose; 8 experiments with 30 mM glucose, and 7 experiments with 30 mM mannitol.

in 30 mM glucose medium. Densitometric analysis revealed that this was significant when compared to those cultured in 5.5 mM glucose and 30 mM mannitol medium (fig. 6).

**Discussion**

Altered regulation of Ca\(^{2+}\) may play a key role in multiple cellular dysfunctions. A sustained tonic upregulation of [Ca\(^{2+}\)], is capable of activating a number of potentially harmful processes. Some of these occur through the activation of hydrolytic enzymes such as phospholipase A\(_2\), Ca\(^{2+}\)-activated proteases, and Ca\(^{2+}\)-activated endonucleases, whereas others occur through the modulation of signaling pathways including ligand-activated kinases as well as a number of intracellular messengers [6–8]. To maintain intracellular Ca\(^{2+}\) homeostasis, a complex regulatory mechanism is required that integrates the release from internal stores, transmembrane influx from and efflux to the extracellular space, and sequestration into internal stores [14–18]. There are three kinds of Ca\(^{2+}\)-regulating pathways clearly identified in rat MCs: inositol trisphosphate (InsP\(_3\))-mediated release from intracellular stores; SOCI, and NCX. The voltage-operated L-type calcium channel, another important pathway, has rarely been investigated in these cells. One study reported that this channel is also present in MCs and, if this is the case, its function is expected to be related to agonist-induced contraction as in other cells [19].

Ca\(^{2+}\) release from intracellular stores requires stimulation of vasoconstrictive hormones, angiotensin II, arginine vasopressin or thromboxane mimetics which act as stimulators. Binding of these molecules to phospholipase C-coupling receptor induces the release of equimolar water-soluble InsP\(_3\) and diacylglycerol from membrane phosphoinositides. Consequently, InsP\(_3\) mobilizes Ca\(^{2+}\) from the intracellular stores, resulting in an explosive elevation in [Ca\(^{2+}\)], and subsequent influx of Ca\(^{2+}\) through SOCI. SOCI, also called ‘capacitative’ or Ca\(^{2+}\)-release-activated Ca\(^{2+}\) channel influx, sustains Ca\(^{2+}\) influx following InsP\(_3\)-mediated Ca\(^{2+}\) release [14–18, 20]. Recent investigations revealed that the [Ca\(^{2+}\)], response of MCs to vasoactive hormones is blunted in high-glucose conditions including the function of SOCI [10–13]. These changes contribute to the loss of contractility of MCs in response to vasoactive hormones and are relevant to hyperfiltration and microalbuminuria in early diabetic nephropathy.

As diabetic nephropathy progresses, the pathophysiological features other than contractile abnormality appear, such as phenotypic changes, deranged proliferative rate, apoptosis and death. These Ca\(^{2+}\)-dependent cellular dysfunctions are related to a sustained ‘tonic’ elevation, rather than a responsive increase in stimulators of [Ca\(^{2+}\)]. The adverse role of elevated basal [Ca\(^{2+}\)], have already been reported in many other cells in various pathophysiological conditions [21–25]. Two recent studies report that intracellular Ca\(^{2+}\) signals are related to the regulation of proliferation in MCs [26, 27]. Therefore, we focused on the effect of high glucose on basal [Ca\(^{2+}\)], and its regulation in MCs that are not stimulated.

The present study shows that high glucose can alter the basal intracellular Ca\(^{2+}\) pool. The feature of [Ca\(^{2+}\)], increase was rather gradual and sustained, which is quite different from the curve by stimulation. It does not match the [Ca\(^{2+}\)], curve by InsP\(_3\)-mediated release from intracellular stores.


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The increased Ca\textsuperscript{2+} seems to come entirely from transmembrane influx from the extracellular space in that 0 Ca\textsuperscript{2+} perfusion results in complete disappearance of a high-glucose-induced increase in [Ca\textsuperscript{2+}]. To identify the Ca\textsuperscript{2+} channel responsible for the high-glucose-induced transmembrane influx, we blocked SOCl with SKF96365 since it is the only kind of transmembrane Ca\textsuperscript{2+} channel confirmed in MCs\textsuperscript{[10, 28]}. As a result, the high-glucose-induced increase in [Ca\textsuperscript{2+}], was blocked by SKF96365, suggesting that the entry of Ca\textsuperscript{2+} occurs mainly through SOCl.

Two important points from our findings are that InsP\textsubscript{3}-mediated release from intracellular stores does not evolve and that SOCl is a main pathway in the process of the mediated release from intracellular stores. Primarily, SOCl is activated following the prior agonist stimulation through the release of Ca\textsuperscript{2+} from intracellular stores or a putative ‘influx factor’ coupled to the depletion in Ca\textsuperscript{2+} stores\textsuperscript{[13]}. In the stimulated MC models, the response of SOCl to the stimulators was decreased in the high-glucose condition\textsuperscript{[10–13]}. Another aspect is whether the activity of SOCl by itself is changed and, if so, in which direction will it be. The change in the resting activity of SOCl in the high-glucose condition is still unknown. Our results also cannot provide further information since the Ca\textsuperscript{2+} influx via SOCl occurred spontaneously and was beyond the normal regulating mechanism of SOCl in the present study. A simple malfunction of the channel might permit Ca\textsuperscript{2+} flux regardless of its activity. Therefore, the investigations on this issue will provide further interesting information on the mechanism of Ca\textsuperscript{2+} homeostasis in diabetic nephropathy.

In the next step, we focused on the NCX function. Intracellular Ca\textsuperscript{2+} homeostasis is not only maintained in resting conditions\textsuperscript{[29, 30]}, but also in stimulated conditions\textsuperscript{[31, 32]}. Normally it transports Na\textsuperscript{+} into and Ca\textsuperscript{2+} out of the cells, the so-called forward-mode activity, but it can also be activated by the reverse mode. To observe the presence and activity of NCX, the reverse mode of NCX activity is commonly elicited by the acute removal of extracellular Na\textsuperscript{+}. We evaluated the activity of NCX in the reverse mode using this popular experimental method. In our results, the activity of NCX was depressed in MCs cultured in the high-glucose condition. Peak [Ca\textsuperscript{2+}], following NCX stimulation was significantly decreased by high glucose as compared to normal glucose or mannitol. Therefore, we evaluated the actual Ca\textsuperscript{2+} clearance via NCX using a further complex method activating NCX in the forward mode. First, we increased intracellular Ca\textsuperscript{2+} by activating the reverse mode of NCX, as described above. Since NCX activity is depressed in MCs cultured in high glucose, achieving a sufficient intracellular Ca\textsuperscript{2+} was not easy in these cells. The addition of thapsigargin might be helpful in achieving sufficient intracellular Ca\textsuperscript{2+} since it causes a transient sustained [Ca\textsuperscript{2+}], elevation by depleting Ca\textsuperscript{2+} from the intracellular stores. In the condition of increased intracellular Ca\textsuperscript{2+}, extracellular Na\textsuperscript{+} was acutely restored to the physiological concentration to activate NCX in the forward mode. To exclude the possible sources of Ca\textsuperscript{2+} removal rather than NCX, La\textsuperscript{3+}, a PMCA inhibitor, and thapsigargin and caffeine, a blocker of reuptake to intracellular stores, were added during this procedure. As a result, intracellular Ca\textsuperscript{2+} was rapidly decreased in MCs cultured in normal glucose or mannitol. In contrast, the decrease in Ca\textsuperscript{2+} was blunted in MCs cultured in high glucose, showing a significantly decreased slope.

The alterations in NCX activity have frequently been reported in various pathological conditions. For example, it relates to contractile dysfunction, arrhythmogenesis and remodeling of cardiac muscles\textsuperscript{[33–35]}, to abnormal Ca\textsuperscript{2+} regulation in pancreatic cells\textsuperscript{[36, 37]}, and to the abnormal development of brain cells\textsuperscript{[38, 39]}. In cultured human and rat MCs, the expression of NCX was shown to be related to the rate of growth and life span\textsuperscript{[40–42]}. One of the important implications of the present study is that NCX function of MCs can also be altered in diabetic conditions. The change in NCX function may relate to a number of pathophysiological changes that depend on the intracellular Ca\textsuperscript{2+} signal. In Western blot analysis, the depression in NCX by high glucose was revealed to occur at the level of protein expression. The investigation on the pathophysiological role of altered NCX in MCs will be an interesting subject in the field of diabetic nephropathy. Further investigations are also required to define the mechanism by which high glucose decreases protein expression of NCX.

In conclusions, we found that high glucose by itself increases basal [Ca\textsuperscript{2+}], in unstimulated MCs. [Ca\textsuperscript{2+}], was increased in a dose-dependent manner with glucose concentrations. The entry of extracellular Ca\textsuperscript{2+}, perhaps mainly through SOCl, was the source of Ca\textsuperscript{2+} during the high-glucose-induced increase in [Ca\textsuperscript{2+}]. The activity of NCX and its Ca\textsuperscript{2+} clearance ability decreased in MCs cultured in the high-glucose condition. The inhibited function of NCX occurred at the level of protein expression.
The present study provides novel insight into the pathophysiological mechanisms of diabetic nephropathy since many cellular functions depend critically on intracellular Ca\(^{2+}\) homeostasis.

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