

## Effect of high glucose concentrations on prostacyclin-stimulating factor mRNA expression in cultured aortic smooth muscle cells

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**Summary** Prostacyclin (PGI<sub>2</sub>) is a potent vasoactive prostanoid regulating vascular tone. We recently purified and cloned a PGI<sub>2</sub>-stimulating factor (PSF), which stimulates PGI<sub>2</sub> production by vascular endothelial cells (ECs). Previous study demonstrated that PSF is predominantly located in vascular smooth muscle cells (SMCs) and present in serum. PSF may act on vascular ECs to regulate PGI<sub>2</sub> synthesis for maintaining vessel wall homeostasis. Decreased PSF production in the vessel wall may result in an imbalance of prostanoid synthesis, leading to the development of vascular lesions such as diabetic angiopathy. In the present study, to investigate the regulatory mechanisms of PSF gene expression, we examined the effect of high glucose concentrations on PSF mRNA expression in cultured bovine aortic SMCs. Expression of PSF mRNA was significantly decreased to 66 ± 6 % of control value ( $p < 0.01$ ), when the glucose level was raised from 5.5 to 27.8 mmol/l. We also examined the effect of osmolarity on PSF

mRNA expression by addition of an appropriate dose of mannitol to the culture medium. We confirmed that high glucose concentration itself reduced the expression of PSF mRNA and glucose had much more effect than the osmolarity control. The expression of PSF mRNA was significantly decreased to 72 ± 5 % of control value ( $p < 0.05$ ) by a protein kinase C (PKC) activator, phorbol-12-myristate-13-acetate (PMA). The decreased expression of PSF mRNA in the presence of high glucose or PMA was restored by co-incubation with a PKC-specific inhibitor (GF109203X). These results suggest that PSF gene expression in vascular SMCs may be decreased via a specific effect of high glucose concentrations. High glucose-induced activation of PKC is suggested to participate partly in the regulation of PSF gene expression. [Diabetologia (1998) 41: 134–140]

**Keywords** Prostacyclin-stimulating factor, aortic smooth muscle cells, high glucose, protein kinase C.

Diabetic angiopathy is clinically important because it influences the prognosis and quality of life in patients with diabetes mellitus. The precise pathogenesis of diabetic angiopathy is still controversial, but hyper-

glycaemia is generally accepted as a significant factor in the development of both diabetic microangiopathy and macroangiopathy [1, 2]. The Diabetes Control and Complications Trial (DCCT) [3] recently demonstrated the clinical significance of hyperglycaemia in the development of diabetic angiopathy. Elevated glucose levels may cause a variety of metabolic abnormalities in the vascular cells and tissues of diabetic patients, including activation of the polyol pathway [4], myoinositol depletion [4], reduced Na<sup>+</sup>-K<sup>+</sup>-ATPase activity [4], formation of advanced glycation end-products [5], generation of superoxide and lipid peroxides [6, 7], alteration of the cellular redox balance [6], increased eicosanoid production [8], and activation of protein kinase C (PKC) [9–16].

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*Abbreviations:* SMC, Smooth muscle cell; EC, endothelial cell; PDS, plasma-derived serum; PGI<sub>2</sub>, prostacyclin; PSA, PGI<sub>2</sub>-stimulating activity; PSF, PGI<sub>2</sub>-stimulating factor; PG, prostaglandin; PKC, protein kinase C; PMA, phorbol-2 myristate-13-acetate; GFX, GF109203X.

Abnormalities of vasoactive substances have also been implicated in the pathogenesis of diabetic angiopathy [17, 18]. Prostacyclin (PGI<sub>2</sub>) is mainly produced by vascular endothelial cells (ECs) and is a potent vasoactive prostanoid which acts on vascular smooth muscle cells (SMCs) and regulates vascular tone. PGI<sub>2</sub> has a vasodilatory effect and also inhibits platelet aggregation, contributing to the maintenance of vessel wall homeostasis [19–23]. PGI<sub>2</sub> production by vascular ECs has been reported to be decreased in sera from diabetic animals [24]. Our previous studies [24, 25] demonstrated that plasma-derived serum (PDS) contains a PGI<sub>2</sub>-stimulating activity (PSA) and that this activity is significantly decreased in the PDS from experimental diabetic animals and patients with non-insulin-dependent diabetes mellitus. Thus, decreased PGI<sub>2</sub> production in diabetic patients may be due partly to a decreased level of PSA.

Recently, we purified and cloned a PGI<sub>2</sub>-stimulating factor (PSF) corresponding to PSA from the conditioned medium of cultured human fibroblasts [26, 27]. PSF is a novel peptide with a molecular weight of about 31 000 daltons, which stimulates PGI<sub>2</sub> production by both cultured bovine and human vascular ECs [26–28]. PSF mRNA is extensively expressed in tissues with abundant vascularity such as lung and kidney, which are known to be sites of high prostaglandin (PG) synthesis [29]. Our previous immunohistochemical study also demonstrated the dominant staining for PSF in arterial SMCs [30]. Furthermore, Northern blot analysis demonstrated that PSF mRNA expression was observed dominantly in cultured bovine aortic SMCs [29]. Therefore, it appears that PSF produced mainly by vascular SMCs may act on vascular ECs to regulate PGI<sub>2</sub> synthesis. Decreased PSF production may result in an imbalance of prostanoid synthesis in the vessel wall leading to the development of vascular lesions such as diabetic angiopathy.

The present study was conducted to elucidate the effect of a high glucose concentration on PSF mRNA expression in cultured bovine aortic SMCs. Furthermore, the regulatory mechanisms of PSF mRNA expression by glucose concentration was evaluated in relation to the activation of the diacylglycerol-PKC pathway by hyperglycaemia.

## Materials and methods

*Culture of smooth muscle cells.* Bovine aortic SMCs were obtained and subcultured according to the method described previously [31, 32]. In brief, the intima-media segments of bovine thoracic aorta were cut into small explants (1 mm<sup>2</sup>) under sterile conditions. Approximately 20 explants were placed in a culture dish 100 mm in diameter (Falcon 3003; Becton Dickinson, Lincoln Park, N.J., USA) and a cover glass was placed on top. The cells attaching to the dish were cultured in growth medium [Dulbecco's modified Eagle's medium (DMEM; Gibco Labo-

ratories, Grand Island, N. Y., USA) containing 10 % heat-inactivated fetal calf serum (FCS; Gibco), 100 µU/ml penicillin, and 100 µg/ml streptomycin (Gibco)] in an atmosphere of 95 % air and 5 % CO<sub>2</sub> at 37 °C. The initial outgrowth of SMCs was detected by phase-contrast microscopy after 1–2 weeks of culture. The medium was replaced with fresh medium twice a week. When the cells became confluent, the explants were carefully removed from the dishes with sterilized tweezers. The cells were then routinely subcultured by trypsinization with 0.02 % (w/v) EDTA (Katayama Chemical, Osaka, Japan) containing 0.05 % (w/v) trypsin (Katayama) until sufficient cells for the experiments had been produced. The cultured SMCs showed features that were consistent with the morphological characteristics described by Ross [31]. SMCs from the 10th subculture were used in the present study.

When the cells became almost (80–90 %) confluent in the culture dishes, the medium was exchanged for FCS-free DMEM. After serum-free culture for 24 h, the cells were exposed to a high glucose concentration (FCS-free DMEM with 11.1 or 27.8 mmol/l glucose) or a normal glucose concentration (FCS-free DMEM with 5.5 mmol/l glucose) for 1–4 days with daily medium changes. To evaluate the osmolar effect, the cells were also exposed to iso-osmolarity to 27.8 mmol/l glucose (FCS-free DMEM with 5.5 mmol/l glucose plus 22.3 mmol/l mannitol).

To evaluate the relationship between PKC activation and high glucose-induced alteration of PSF mRNA expression, further studies were conducted as follows. After culture with FCS-free DMEM for 24 h, almost confluent SMCs were exposed to either normal glucose (5.5 mmol/l) or high glucose (27.8 mmol/l) for 3 days. The final 8 h of 3 days incubation was done with or without a PKC activator, phorbol-12-myristate-13-acetate (PMA) (Sigma Chemical Co. St. Louis, Mo., USA) (100 nmol/l), or a specific PKC inhibitor, GF109203X (GFX) (Calbiochem-Novabiochem Corp., La Jolla, Calif., USA) (5 µmol/l). Before adopting 8 h as an incubation time in the present experiment, we performed preliminary studies. We confirmed that the time-dependent effect of PMA and GFX on PSF mRNA expression in the cells was observed and cell damage was not found in the present condition.

We also confirmed no significant increase of lactate dehydrogenase activity in each culture medium and more than 95 % of SMCs showing trypan blue exclusion in each culture dish. Therefore, there was no apparent cell damage in any experimental conditions of the present study.

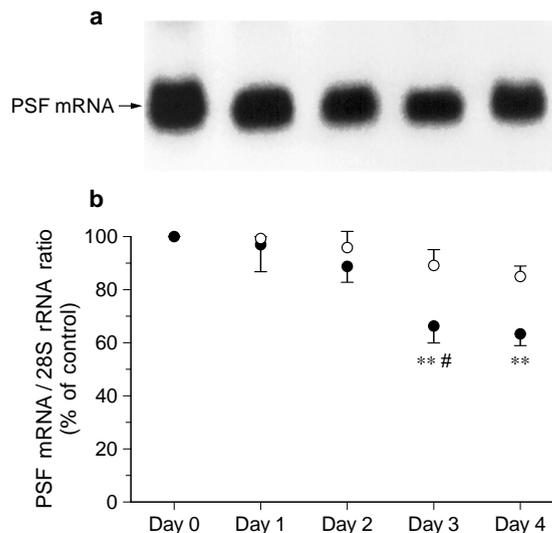
*Northern blot analysis.* To prepare RNA, the SMCs were lysed in 1 ml of ISOGEN solution (Wako Chemical, Tokyo, Japan) after washing with phosphate-buffered saline (pH 7.4). Total RNA was extracted according to the method described by Chomczynski and Sacchi [33] and according to the manufacturer's instructions for ISOGEN (Wako). In brief, after extraction with chloroform (Katayama Chemical, Osaka, Japan), precipitation in isopropanol (Katayama), and successive washes with 75 % ethanol (Katayama), the pellet was dissolved in diethylpyrocyanate (Sigma)-treated water. After quantification by measurement of the absorbance at 260 nm, 20 µg of total RNA was applied to each lane and electrophoresed on 1 % agarose gel (Sigma) containing 2.2 mol/l formaldehyde (Katayama) [34]. The RNA was then transferred to a Hybond N filter (Amersham, Buckinghamshire, UK) and fixed by baking at 80 °C for 2 h. PSF cDNA was obtained according to the method described previously by Yamauchi et al. [27]. The filter was hybridized with α-<sup>32</sup>P (deoxycytidine 5'-[α-<sup>32</sup>P] triphosphate; Amersham)-labelled probe (265-bp pair PvuII / SmaI-digested PSF cDNA fragment) using a multiprimer DNA-labelling system (Amersham) at 42 °C for 16 h in a solution

prepared with hybridization buffer tablets (Amersham) containing 50% formamide (Katayama). Filters were washed with  $2 \times$  SSPE ( $1 \times$  SSPE = 150 mmol/l NaCl, 10 mmol/l  $\text{NaH}_2\text{PO}_4$ , and 1 mmol/l EDTA, pH 7.4) plus 0.1% SDS at room temperature (three times for 10 min each) and then washed with  $1 \times$  SSPE plus 0.1% SDS at  $42^\circ\text{C}$  for 15 min and exposed to a Fuji imaging plate (Fuji Photo Film Co., Ltd., Tokyo, Japan) at room temperature for 30 min. The radioactivity of PSF mRNA was calculated by a Bio Imaging Analyzer (Fuji Photo Film). The same filter was stripped as described previously [35], and then rehybridized with  $\alpha$ - $^{32}\text{P}$ -dCTP labelled 28S ribosomal RNA probe (Oncogene, Cambridge, M.A., USA) as the RNA loading control [36]. The radioactivity ratio (PSF mRNA/28S rRNA) was used for comparison among each set of culture conditions.

**Statistical analysis.** All data are expressed as means  $\pm$  SEM. Comparisons between two groups were done by the Student's unpaired *t*-test. ANOVA and the Newman-Keuls test were used to assess the statistical significance of multiple comparisons. A *p* value less than 0.05 was taken to indicate a significant difference.

## Results

The effect of high glucose on PSF mRNA expression was evaluated by determining the radioactivity ratio of PSF mRNA to 28S rRNA in bovine aortic SMCs cultured with FCS-free DMEM containing 27.8 mmol/l glucose or 5.5 mmol/l glucose. The PSF mRNA / 28S rRNA ratio was decreased in a time-dependent fashion when vascular SMCs were cultured with 27.8 mmol/l glucose. After 3 days of culture, the ratio was significantly decreased to  $66 \pm 6\%$  of control value ( $p < 0.01$ ,  $n = 4$ ) in the vascular SMCs cultured with 27.8 mmol/l glucose as compared to the value on 0 day of culture. However, there was no significant change in the ratio when vascular SMCs were cultured with 5.5 mmol/l glucose (Fig. 1). After 3 days of culture, the PSF mRNA/28S rRNA ratio of vascular SMCs cultured with a high glucose level (27.8 mmol/l) was significantly lower than that of vascular SMCs with a normal glucose level (5.5 mmol/l) ( $66 \pm 6$  vs  $89 \pm 6\%$  of control value,  $n = 4$ ,  $p < 0.01$ , Fig. 1). After 4 days of culture, the ratio was also significantly lower in vascular SMCs cultured with high glucose, although the decrease was not significant as compared to that after 3 days of culture (data not shown). Therefore, the PSF mRNA/28S rRNA ratio was determined using vascular SMCs cultured for 3 days in subsequent experiments. As shown in Figure 2a, the PSF mRNA/28S rRNA ratio in vascular SMCs was decreased by elevated glucose levels in a dose-dependent fashion. When glucose level in the culture medium was elevated to 11.1 mmol/l, the ratio was decreased to  $88 \pm 6\%$  of control value. Whereas, there was no significant change in the ratio when the cells were cultured with 5.5 mmol/l glucose. When glucose level was elevated to 27.8 mmol/l, the ratio was significantly decreased to  $69 \pm 3\%$  of con-

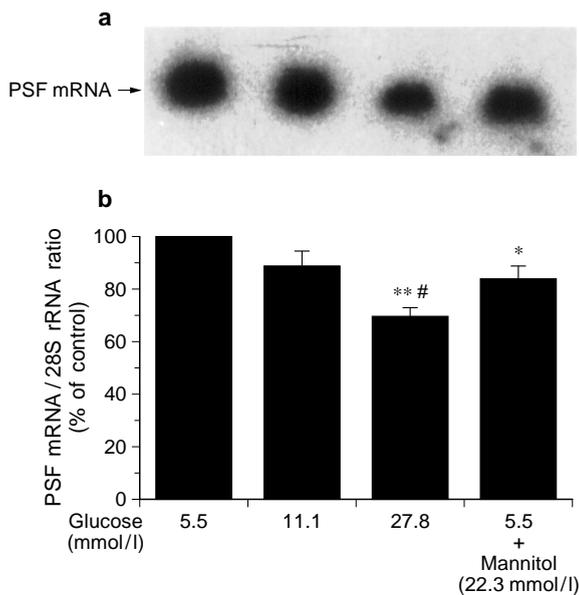


**Fig. 1a, b.** Effect of a high glucose concentration on PSF mRNA expression in cultured vascular SMCs. The cells were exposed to FCS-free DMEM containing 5.5 mmol/l ( $\circ$ ) or 27.8 mmol/l ( $\bullet$ ) glucose for 1 to 4 days. **a)** Northern blot analysis was performed using a PSF cDNA probe and the PSF mRNA/28S rRNA ratio was calculated as described in Materials and methods. **b)** Radioactivity ratio was found to decrease significantly in time-dependent fashion. Each experiment was performed in triplicate using 3 culture dishes ( $n = 4$ ). Results are expressed as the mean percentage of the control  $\pm$  SEM of the radioactivity ratio. \*\*  $p < 0.01$  vs 5.5 mmol/l glucose, #  $p < 0.01$  vs day 0

rol value ( $n = 4$ ,  $p < 0.01$ , Fig. 2b). Our preliminary study indicated that the decreased ratio was restored to the former level after 3 days of post-incubation with 5.5 mmol/l glucose. Therefore, the effect of glucose on PSF mRNA expression was thought to be reproducible.

In addition, to elucidate the effect of high osmolarity on PSF mRNA expression, the cells were cultured with iso-osmolar milieu to 27.8 mmol/l glucose (5.5 mmol/l glucose plus 22.3 mmol/l mannitol). The ratio was significantly decreased to  $83 \pm 5\%$  of control value ( $n = 4$ ,  $p < 0.05$ , Fig. 2b), when the cells were cultured with iso-osmolar milieu to 27.8 mmol/l glucose. However, there was still a significant difference between the ratios cultured with 27.8 mmol/l glucose and cultured with the iso-osmolar milieu ( $n = 4$ ,  $p < 0.05$ , Fig. 2b).

The mechanism of the high glucose-induced decrease of PSF mRNA expression in vascular SMCs was examined focusing on glucose-induced PKC activation in the diacylglycerol-PKC pathway. The effect of a specific PKC inhibitor, GFX, on the high glucose-induced decrease of PSF mRNA expression in vascular SMCs was examined (Fig. 3a). The decreased PSF mRNA/28S rRNA ratio of vascular SMCs cultured with 27.8 mmol/l glucose was increased to  $97 \pm 5\%$  of control value ( $n = 4$ ) by addition of GFX to the culture medium (Fig. 3b). In addi-

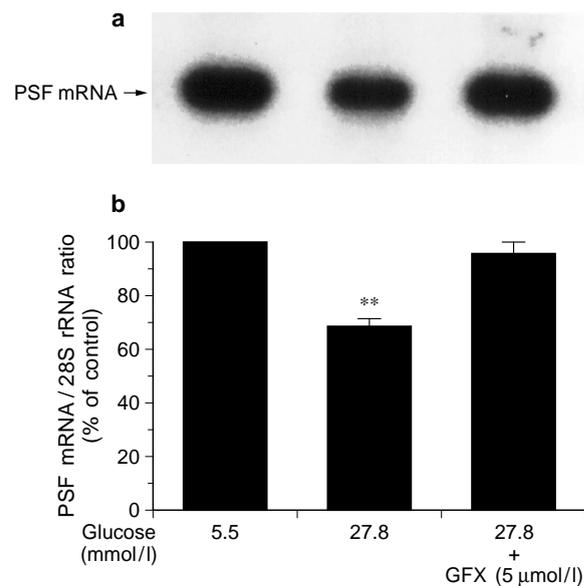


**Fig. 2a, b.** Dose-dependent effect of a glucose concentration and osmolar effect on PSF mRNA expression in cultured vascular SMCs. The cells were exposed to FCS-free DMEM containing 5.5 mmol/l glucose, 11.1 mmol/l glucose, 27.8 mmol/l glucose, and 5.5 mmol/l glucose plus 22.3 mmol/l mannitol for 3 days. Northern blot analysis was performed and the PSF mRNA/28S rRNA ratio was calculated as described in Materials and methods. **a)** Representative PSF mRNA message of cultured SMCs in each glucose concentration and glucose plus mannitol medium. **b)** Radioactivity ratios of each culture conditions are shown. Each experiment was performed in triplicate using 3 culture dishes ( $n = 4$ ). Results are expressed as the mean percentage of the control  $\pm$  SEM of the radioactivity ratio. \*  $p < 0.05$  vs 5.5 mmol/l glucose, \*\*  $p < 0.01$  vs 5.5 mmol/l glucose, #  $p < 0.05$  vs 5.5 mmol/l glucose plus 22.3 mmol/l mannitol

tion, the PKC activator, PMA, was found to mimic the effect of high glucose on PSF mRNA expression in vascular SMCs (Fig. 4a and b). The PSF mRNA/28S rRNA ratio was significantly decreased to  $72 \pm 5\%$  of control value ( $n = 4$ ,  $p < 0.05$ ) by addition of PMA to the normal glucose medium (Fig. 4b). The decreased radioactivity ratio of vascular SMCs in the presence of PMA was restored to  $96 \pm 7\%$  of control value ( $n = 4$ ) by addition of GFX (Fig. 4b).

## Discussion

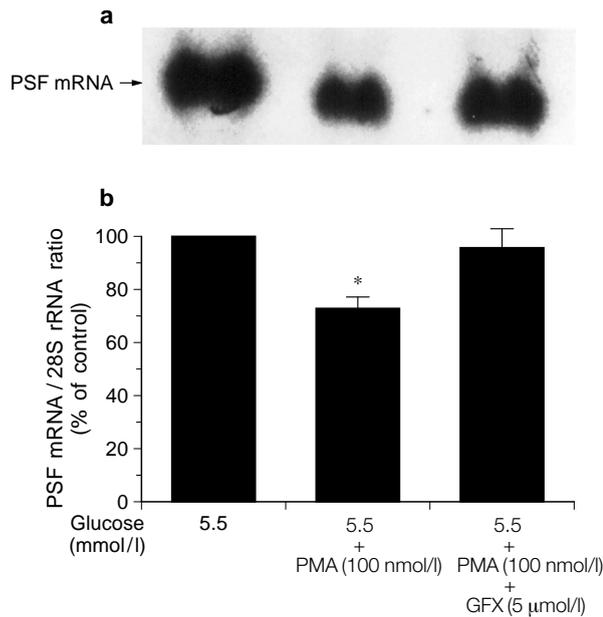
There have been several hypotheses proposed to explain the effect of hyperglycaemia on the development of diabetic vascular complications [1–12]. We have focused on the abnormalities of PG metabolism in diabetic patients and animals [37–43]. In particular, decreased PGI<sub>2</sub> synthesis commonly found in patients and experimental animals with diabetes might influence the progression of vascular damage such as diabetic microangiopathy and macroangiopathy.



**Fig. 3a, b.** Recovery of the high glucose-induced reduction of PSF mRNA expression in cultured vascular SMCs by a specific PKC inhibitor, GFX. The cells were exposed to FCS-free DMEM containing 27.8 mmol/l glucose for 3 days. The final 8 h incubation was performed with or without GFX. Northern blot analysis was performed and the PSF mRNA/28S rRNA ratio was calculated as described in Materials and methods. **a)** Representative PSF mRNA message in each culture condition. **b)** Decreased radioactivity ratio of cultured SMCs with high glucose was increased by the addition of GFX. Results are expressed as the mean percentage of the control  $\pm$  SEM of the PSF mRNA/28S rRNA ratio. \*\*  $p < 0.01$  vs 5.5 mmol/l glucose

The in vivo effect of hyperglycaemia on PG metabolism is complex, because diabetic patients show various kinds of abnormalities connected with circulating metabolic factors. Similarly, the in vitro effects of a high glucose concentration on PGI<sub>2</sub> production by cultured vascular ECs have also been controversial. Aanderud et al. [40] reported that high glucose had no effect on PGI<sub>2</sub> production by cultured human ECs. On the other hand, our previous studies have demonstrated that long-term culture of bovine aortic ECs in high glucose concentration reduces PGI<sub>2</sub> production, suggesting that the high glucose concentration is one of the important factors in the development of vascular dysfunction for PGI<sub>2</sub> synthesis [44].

Recently, we succeeded in cloning the cDNA of a novel bioactive peptide, PGI<sub>2</sub>-stimulating factor (PSF) [20]. The major biological in vivo action of PSF, which is mainly produced by vascular SMCs, is thought to stimulate the production of PGI<sub>2</sub> by vascular ECs for the maintenance of vascular homeostasis [28–30]. A decreased PSF expression has been found in the kidneys of diabetic rats and patients with diabetes and atherosclerosis [45, 46]. Therefore, the decreased PSF synthesis in the vessel wall might be a possible pathogenesis of an imbalance of prostanoid



**Fig. 4a, b.** Similar effect of a PKC activator, PMA, to high glucose on PSF mRNA expression in cultured vascular SMCs. The cells were exposed to 5.5 mmol/l glucose for 3 days. The final 8 h incubation was performed by addition of PMA, without PMA, with PMA plus GFX. Northern blot analysis was performed and the PSF mRNA/28S rRNA ratio was calculated as described in Materials and methods. **a)** Representative PSF mRNA message in each culture condition. **b)** Decreased radioactivity ratio by addition of PMA was increased by a co-incubation of GFX. Each experiment was performed in triplicate with 3 culture dishes ( $n = 4$ ). Results are expressed as the mean percentages of the control  $\pm$  SEM of the radioactivity ratio. \*  $p < 0.05$  vs 5.5 mmol/l glucose

synthesis involving the development of vascular damage such as diabetic angiopathy.

In the present study, we demonstrated that a high glucose concentration decreased the expression of PSF mRNA in cultured bovine aortic SMCs in a time- and dose-dependent fashion. A high osmolarity adjusted by the addition of mannitol also decreased the expression of PSF mRNA as compared with the normal glucose concentration (5.5 mmol/l). However, PSF mRNA level was still significantly lower ( $n = 4$ ,  $p < 0.05$ ) in the cells cultured in high glucose (27.8 mmol/l) than those with iso-osmolarity (5.5 mmol/l glucose plus 22.3 mmol/l mannitol) to high glucose. Although the precise mechanisms of osmolar effect by mannitol on PSF mRNA expression are unknown, it seems that the decreased expression of PSF mRNA in the cells cultured in high glucose may be due partly to an increased osmolarity. The effect of osmolarity was also confirmed using Western blotting and immunohistochemical study under similar experimental conditions [46]. Cell damage due to a high glucose concentration or mannitol was not observed in the present experimental conditions. Therefore, these findings suggest that a high glucose con-

centration itself reduced the expression of PSF mRNA and glucose had much more effect than the osmolarity control by addition of mannitol. Additionally, our previous Western blot analysis demonstrated that the intensity of specific band corresponding to PSF in cell lysate from high glucose culture was decreased compared to that of normal glucose [46]. Thus, high glucose-induced decrease of PSF mRNA expression might lead to the decreased production of PSF protein in vascular SMCs. We suggest that decreased PSF synthesis by vascular resident cells resulting from a high glucose concentration might be one reason for the decreased PGI<sub>2</sub> synthesis observed in vessels of diabetic patients.

Furthermore, to clarify the regulatory mechanisms of PSF gene expression by a glucose concentration, we focused on the diacylglycerol-PKC pathway. As regards the adverse effect of hyperglycaemia on the vasculature, there were some reports demonstrating a high glucose-induced PKC activation through increased de novo synthesis of diacylglycerol in vascular cells [9–16]. Since PKC modulates vascular permeability, contractility, and cell proliferation, abnormal activation of PKC in the vascular tissues of diabetic patients seems to be one possible cause of the development of diabetic angiopathy. We have previously confirmed the activated PKC levels in vascular SMCs cultured under high glucose conditions or in PMA added culture medium as well as the present experimental condition [11]. We also confirmed the viability of the cells as described in Materials and methods. In the present study, the PKC activator PMA mimicked the effect of high glucose levels on PSF mRNA expression in vascular SMCs. Furthermore, the effects of both high glucose levels and PMA on PSF mRNA expression in vascular SMCs were almost abolished by co-incubation with a PKC-specific inhibitor, GFX. These findings suggest that high glucose-induced PKC activation acts as a possible regulator for PSF gene expression in vascular SMCs. Further studies using other activators and inhibitors in correlation to the diacylglycerol-PKC pathway are needed, because there may be many factors which are altered by a high glucose concentration in this pathway.

Including the osmolar effect, there may be many possible candidates to regulate PSF gene expression. Therefore, to clarify the precise mechanisms of high glucose-induced regulation for PSF gene expression, we should study the relationship between the high glucose level-induced intracellular alterations such as transcription factor levels and abnormalities of PSF expression.

In conclusion, a high glucose concentration itself reduced the PSF gene expression in vascular SMCs and glucose had much more effect than appropriate osmolarity control. Decreased expression of PSF mRNA by high glucose level is suggested to be due

partly to the high glucose-induced PKC activation. The diacylglycerol-PKC pathway was suggested to participate in the regulation of PSF gene expression in part. We hypothesize that decreased expression of PSF mRNA, leading to the reduction of PSF synthesis by vascular resident cells, results in the decreased PGI<sub>2</sub> synthesis in vessel wall. These changes might be an important candidate leading to the development of diabetic angiopathy.

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