Phosphorylation of signal transducer and activator of transcription 3 induced by hyperglycemia is different with that induced by lipopolysaccharide or erythropoietin via receptor-coupled signaling in cardiac cells

YU-HSIN CHIU¹, PO-MING KU², YUNG-ZE CHENG³,⁴, YINGXIAO LI⁴, JUEI-TANG CHENG⁴,⁵ and HO-SHAN NIU⁶

¹Division of Infectious Diseases; ²Cardiovascular Center, Department of Internal Medicine, Chi-Mei Medical Center-Liouying, Tainan 73601; Departments of ³Emergency Medicine and ⁴Medical Research, Chi-Mei Medical Center, Tainan 71003; ⁵Institute of Medical Sciences, Chang Jung Christian University, Tainan 71101; ⁶Department of Nursing, Tzu Chi University of Science and Technology, Hualien 97005, Taiwan, R.O.C.

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Abstract. The signal transducer and activator of transcription 3 (STAT3) is known to be involved in hypertrophy and fibrosis in cardiac dysfunction. The activation of STAT3 via the phosphorylation of STAT3 is required for the production of functional activity. It has been established that lipopolysaccharide (LPS)-induced phosphorylation of STAT3 in cardiomyocytes primarily occurs through a direct receptor-mediated action. This effect is demonstrated to be produced rapidly. STAT3 in cardiac fibrosis of diabetes is induced by high glucose through promotion of the STAT3-associated signaling pathway. However, the time schedule for STAT3 activation between LPS and high glucose appears to be different. Therefore, the difference in STAT3 activation between LPS and hyperglycemia in cardiomyocytes requires elucidation. The present study investigated the phosphorylation of STAT3 induced by LPS and hyperglycemia in the rat cardiac cell line H9c2. Additionally, phosphorylation of STAT3 induced by erythropoietin (EPO) via receptor activation was compared. Then, the downstream signals for fibrosis, including the connective tissue growth factor (CTGF) and matrix metalloproteinase (MMP)-9, were determined using western blotting, while the mRNA levels were quantified.

LPS induced a rapid elevation of STAT3 phosphorylation in H9c2 cells within 30 min, similar to that produced by EPO. However, LPS or EPO failed to modify the mRNA level of STAT3, and/or the downstream signals for fibrosis. High glucose increased STAT3 phosphorylation to be stable after a long period of incubation. Glucose incubation for 24 h may augment the STAT3 expression in a dose-dependent manner. Consequently, fibrosis-associated signals, including CTGF and MMP-9 protein, were raised in parallel. In the presence of tiron, an antioxidant, these changes by hyperglycemia were markedly reduced, demonstrating the mediation of oxidative stress. Therefore, LPS- or EPO-induced STAT3 phosphorylation is different compared with that caused by high glucose in H9c2 cells. Sustained activation of STAT3 by hyperglycemia may promote the expression of fibrosis-associated signals, including CTGF and MMP-9, in H9c2 cells. Therefore, regarding the cardiac dysfunctions associated with diabetes and/or hyperglycemia, the identification of nuclear STAT3 may be more reliable compared with the assay of phosphorylated STAT3 in cardiac cells.

Introduction

The signal transducer and activator of transcription 3 (STAT3) has been demonstrated to be one of the regulators in cardiac dysfunction (1). STAT3 possesses multiple functions, with its central role described as a transcription factor. Moreover, STAT3 has been demonstrated to function as a signaling molecule, as a factor involved in cellular respiration, and as a protein interacting with the mitochondrial pore (2-5). Therefore, in cardiomyocytes, STAT3 plays an important role in survival, growth, sarcomere architecture, energetics, and metabolism (6-8).

Hyperglycemia is important in the pathogenesis of diabetic disorders. Hyperglycemia was found to increase the STAT3 either through the gene expression or the phosphorylation (9). STAT3 is known as a cytoplasmic transcription factor that transmits extracellular signals to the nucleus (10). Activated
STAT3 in the nucleus binds to specific DNA promoter sequences to regulate the gene expression (11). Recent studies have indicated that hyperglycemia increases STAT3 activation, thereby contributing to the pathophysiology of tissue injury (12). STAT3 activation, increased phosphorylated STAT3 (p-STAT3) and p-STAT3 nuclear translocation, are reportedly some of the underlining mechanisms of STAT3 under high glucose condition. However, p-STAT3 was induced at Y705 and S727 in cells for STAT3 activation by high glucose levels (13). STAT3 has been demonstrated to shuttle between the cytoplasm and nucleus independently of tyrosine phosphorylation (14) while unphosphorylated STAT3 in nucleus also can drive gene expression (15).

Lipopolysaccharide (LPS) is mainly obtained from the outer membrane of gram-negative bacteria, and the inflammatory cytokines produced as a consequence of LPS exposure are implicated in cardiac dysfunction (16,17). The rapid activation of STAT3 by LPS through phosphorylation in cardiomyocytes has been identified (18), and it is suggested as a direct receptor-mediated activation (19). However, STAT3 activation by LPS in hepatocytes is slower than in cardiomyocytes (20). Toll-like receptor 4 (TLR4) is known as the binding site of LPS (21). Activation of TLR4 by LPS has also been indicated to induce an inflammatory response that decreases cardiomyocytes contractility (22). Moreover, the Janus-activated kinase 2 (JAK2) and the STAT3 pathway (JAK2/STAT3 pathway) is also coupled to the signaling of cytokine receptors including TLR4 (23). Otherwise, erythropoietin (EPO) is also produced effectiveness through activation of the specific cell-surface receptor, erythropoietin receptor (EPOR) (24). It has been established that JAK2/STAT3 signaling pathway is also coupled to EPOR (25). Interestingly, agent improves left ventricular performance via activation of JAK2/STAT3 pathway in rats (26). Therefore, we included the effects of EPO in this study, because EPO produced actions also through an activation of receptors, EPOR, which is similar to the action of LPS (27).

Additionally, STAT3 is introduced to involve in cardiac fibrosis of diabetes (28), while high glucose increased STAT3 activated by angiotensin II has been demonstrated to be produced mainly through a reactive oxygen species (ROS)-dependent mechanism (29). Recently, the ROS-activated STAT3 pathway has been characterized in early reperfusion of heart (30). High glucose is known as the main factor for inducing diabetes-associated cardiovascular dysfunctions. It seems that activation of STAT3 through phosphorylation by hyperglycemia differs from the promotion of STAT3 via receptor-coupled signaling in the regulation of cardiac function. However, variations in the phosphorylation of STAT3 between high glucose-induced change and promotion by receptor-coupled signaling remained unclear in cardiomyocytes.

In the present study, we focused on STAT3 phosphorylation that is important in regulation of cardiac function. Also, we are interested to know the difference whether STAT3 phosphorylation induced by receptor signaling is varied with that induced by pathologic disorders such as hyperglycemia. Therefore, we used the embryonic rat cardiomyoblast cell line H9c2 which offers the advantage of being an animal-free alternative (31). Moreover, H9c2 expressed TLR4 (32) and EPOR (33). Therefore, it is suitable to apply in the present study.

Additionally, we link it to signals-associated fibrosis, including connective tissue growth factor (CTGF) and metalloproteinase (MMP)-9, to determine its association in cardiac disorders.

Materials and methods

Cell cultures. It has been confirmed that H9c2 cells possess the advantage of being an animal-free alternative (31). The H9c2 cells (BCRC, no. 60096) were cultured according to a previous method (34). In brief, H9c2 cells were maintained in Dulbecco's modified Eagle's medium (pH 7.2) supplemented with 10% fetal bovine serum. The H9c2 cells were plated at a density of 6,000 cells/cm² and allowed to proliferate in growth medium. The medium was changed every 48 h.

Drug treatment. The cultured H9c2 cells were treated at indicated times with Salmonella typhosa LPS (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany), as described previously (18). The stock solution of EPO containing epoetin beta (Recormon, 5,000 IU/0.3 ml), purchased from Roche Diagnostics (Mannheim, Germany), was diluted in culture medium. A fresh solution diluted to the indicated dose was applied to treat the H9c2 cells. Incubation of hyperglycemia with H9c2 cells was also performed according to our previous report (35).

Western blot analysis. Protein was extracted and separated by SDS-PAGE, following our previous method (27). Proteins were detected using antibodies (1:1,000) against p-STAT3, STAT3, CTGF and MMP-9, while antibody against β-actin serving as the internal control. After comparing with the marker, the immunoblots of TLR4 (95 kDa), EPO (55 kDa), p-JAK2 (130 kDa), JAK2 (130 kDa), p-STAT3 (88 kDa), STAT3 (88 kDa), CTGF (38 kDa), MMP-9 (92 kDa) and β-actin (43 kDa) were then quantified.

Quantitative reverse transcription-polymerase chain reaction (qRT-PCR). Total RNA was extracted from cell lysates with TRIzol (Qiagen, Hilden, Germany). Two microgram of total RNA was used for the reverse transcription reaction, along with Superscriptase II (Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA), oligo-dT, and random primers. Web-based assay-design software from the Universal Probe Library Assay Design Center (http://www.roche-applied-science.com/sis/rtpcr/upladc.jsp) was utilized to design TaqMan primer pairs and to select appropriate hybridization probes (Table I). For quantification, real-time PCR analysis was performed using Light Cycler 480 SYBR-Green I Master on a Light Cycler 480 II (Roche Diagnostics). The relative fold changes were quantified using the comparative threshold cycle method, and β-actin was used as a control, according to our previous reports (36,37).

Statistical analysis. Data were indicated as the mean ± standard error of the mean (SEM) from the sample number (n) of each group. The differences between two groups were analyzed using a Student’s two-sided t-test. A value of P<0.05 was considered to indicate a statistically significant difference.
Results

Effects of LPS on the phosphorylation of STAT3 in H9c2 cells. Incubation of LPS dose-dependently induced a marked elevation of STAT3 phosphorylation within 30 min in H9c2 cells, as shown in Fig. 1A. The results were obtained from TLR4 through phosphorylated JAK2 (p-JAK2) to STAT3 phosphorylation (Fig. 1A and B). However, the expression of STAT3 was still not modified in terms of both protein and mRNA levels (Fig. 1C). Consequently, the downstream signals both CTGF and MMP-9 were also not activated by LPS in this condition (Fig. 1A and B).

Effects of EPO on the phosphorylation of STAT3 in H9c2 cells. The same incubation of EPO with H9c2 cells also produced a similar change in STAT3 phosphorylation, as shown in Fig. 2A, except that the EPOR was activated by EPO. Additionally, the expression of STAT3 was also not changed by EPO in terms of both protein and mRNA levels (Fig. 2B and C). Similarly, the consequent downstream signals, both CTGF and MMP-9, were also not activated by EPO in this condition.

Effect of hyperglycemia on the phosphorylation of STAT3 in H9c2 cells. Incubation of high glucose (30 mM) with H9c2 cells at the time same as LPS failed to induce changes in STAT3 phosphorylation. Therefore, we incubated H9c2 cells for a longer time with high glucose in the medium. As shown in Fig. 3A, changes in STAT3 phosphorylation were not stable except at 24 h post-incubation. Both protein and mRNA levels of STAT3 were also markedly elevated after 24-h incubation with high glucose (Fig. 3B).

Additionally, increase of STAT3 expression, both in terms of protein (Fig. 4A) and mRNA (Fig. 4B) levels, was produced in a dose-dependent manner by hyperglycemia after 24-h incubation. However, changes were not observed in H9c2 cells that received similar incubation with manitol (30 mM), which produced the same osmolarity as high glucose (30 mM), as described in our previous report (35). Thus, the possible influence of osmolarity in the changes of STAT3 expression can be excluded.

Moreover, the downstream signals for fibrosis, including CTGF and MMP-9, were also enhanced by hyperglycemia in the same dose-dependent fashion (Fig. 4A). This change was also not related to osmolarity as shown in manitol-treated cells. However, fibrosis-related signals were not modified in H9c2 cells treated with LPS or EPO at the effective dose in above.

Mediation of oxidative stress in hyperglycemia increased expressions of STAT3 in H9c2 cells. We applied the anti-oxidant, tiron, to examine the role of oxidative stress in the changes of STAT3 expression by hyperglycemia. As shown in Fig. 5A, tiron inhibits the elevation of STAT3 activation in a dose-related manner. Moreover, the promotion in mRNA level of STAT3 by hyperglycemia was also reduced by tiron in the same fashion (Fig. 5B). Consequently, fibrosis-associated signals, including CTGF and MMP-9, elevated by hyperglycemia were markedly reduced by tiron in the same manner (Fig. 4A). Mediation of oxidative stress can thus be confirmed.

Discussion

In the present study, we demonstrated that STAT3 phosphorylation occurred within 30 min after exposure to LPS in H9c2 cells. Additionally, the dose-dependent effect of LPS was produced through TLR4 to link the p-JAK2 to STAT3 phosphorylation; it is fully consistent with the previous reports (22,26). However, LPS did not influence the expressions of STAT3 and the downstream signals, including CTGF and MMP-9. Similar results were also obtained in EPO-treated H9c2 cells except that EPOR was involved in the STAT3 phosphorylation by EPO (24). Otherwise, high glucose increased STAT3 phosphorylation after a longer time, particularly 24 h post-incubation. Additionally, expression of STAT3 was also augmented by high glucose in a dose-dependent manner after 24-h incubation. In parallel, fibrosis-related signals, including CTGF and MMP-9, were both elevated. Therefore, STAT3 phosphorylation induced by LPS or EPO is quite different from that by hyperglycemia. The possible reason might be due to the treatment of H9c2 with LPS or EPO may stimulate cardiac mitochondrial function through a highly regulated, receptor-mediated, eNOS/Akt1 and JAK-STAT-dependent cascade that activates the transcriptional program of mitochondrial biogenesis in a short time (38-40). However, hyperglycemia-induced oxidative stress attenuated the mitochondrial function and affected the proliferation and survival of cells. Increased ROS in diabetes is implicated in the development of diabetic cardiomyopathy (41). To the best of our knowledge, the present study is the first to conduct this finding.

Inflammatory cytokines produced as a consequence of LPS exposure are implicated in myocardial dysfunction (17,42). Paradoxically, sub-lethal doses of LPS provided cardioprotective effects against ischemia-reperfusion injury (43). In addition, STAT3 is demonstrated to be a key modulator of an integrated signaling network in the heart (1). Increase of STAT3 phosphorylation by LPS in a CD14-independent manner has been indicated in cardiomyocytes (18). Moreover, EPO is known to protect heart from chemical damage (44) and to improve experimental heart failure (45). In the present study, we observed that STAT3 phosphorylation is raised by EPO in a way that is similar to LPS in H9c2 cells. Interestingly,

<table>
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<th>Target</th>
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<tr>
<td>STAT3</td>
<td>F</td>
<td>5’-GGCTTCAGCCCGAGAGC-3’</td>
</tr>
<tr>
<td></td>
<td>R</td>
<td>5’-CTCCAGTGGCGCTGC-3’</td>
</tr>
<tr>
<td>CTGF</td>
<td>F</td>
<td>5’-ATGCTTGGAGAAGGCTGTA-3’</td>
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<td></td>
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<td>5’-GGGCAATTGTGTCCG-3’</td>
</tr>
<tr>
<td>MMP-9</td>
<td>F</td>
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<td></td>
<td>R</td>
<td>5’-GAAGTCCGTGGGTC-3’</td>
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<tr>
<td></td>
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F, forward; R, reverse; STAT3, signal transducer and activator of transcription 3; CTGF, connective tissue growth factor; MMP-9, matrix metalloproteinase-9.

Table I. Primers used for targets amplification in this study.
both effects were induced by activation of each specific receptors, LPS via TLR4 (21) and EPO through specific receptor EPOR (25). Moreover, both effects on STAT3 phosphorylation were produced through p-JAK2, as described previously (23). Therefore, a rapid increase of p-STAT3 seems helpful in protection against cardiac damage. This view is consistent to previous reports that demonstrated the cardio-protective effects of propofol (46) and morphine (47) through an increase in STAT3 phosphorylation.

In contrast, as shown in Fig. 3, STAT3 phosphorylation was not immediately increased by high glucose, but rather changed in unstable way. After a longer time of incubation with
H9c2 cells, approximately 24 h later, STAT3 phosphorylation was stable and markedly increased in high glucose medium. Moreover, expression of STAT3 was also promoted after the same incubation. Basically, phosphorylation is known as the major way for the activation of STAT. Moreover, p-STAT3 has been shown to enter the nucleus easily (48). However, nuclear accumulation of STAT3 without phosphorylation has also been demonstrated (49), and the unphosphorylated STAT3 in nucleus may activate gene expression both in cancer and in responses to cytokines (15). Therefore, STAT3 is effective to promote transcription in cardiac cells that have been characterized in this study. We demonstrated that fibrosis-related
signals, including CTGF and MMP-9, were both enhanced in high glucose medium with the increased STAT3. This novel view is useful to explain the role of cardiac fibrosis in diabetes (50,51).

Cardiac fibrosis is a prominent component of diabetic cardiomyopathy (52,53). High glucose has been demonstrated to promote fibrosis in vitro, including changes in MMP activity (28). Hyperglycemia may sustain the progression of heart failure through excessive interstitial myocardial collagen accumulation, thus leading to impaired diastolic and systolic function (54). STAT3 is also shown to mediate the proliferation of cardiac fibroblasts and collagen synthesis induced by high glucose (55). Although STAT3 may participate in the transcription of target genes in ischemia/reperfusion injury (54) and pressure overload hypertrophy (55), the role of STAT3 in cardiac fibrosis induced by hyperglycemia is critical.

The pathogenesis of cardiovascular diseases almost invariably involves, the occurrence of oxidative stress (56), a major cause of progressive cellular sufferance and death. Moreover, diabetic cardiomyopathy is a condition in which oxidative stress seems to play a major pathogenic role (57). Therefore, we focused on the role of oxidative stress in the changes of STAT3 induced by hyperglycemia in H9c2 cells. In the presence of the antioxidant, tiron (58), increase of STAT3 by high glucose was markedly reduced in H9c2 cells. Additionally, the augmented fibrosis-related signals, including CTGF and MMP-9, were also attenuated in parallel. Different to the effects induced by stimulation of receptors, such as LPS and EPO, phosphorylation of STAT3 by hyperglycemia needs a longer time in incubation. It seems that enough oxidative stress induced by hyperglycemia needs a time to accumulate and STAT3 phosphorylation via oxidative stress is varied with that rapidly induced via the p-JAK2. Taken together, we found that expression of STAT3 increased by hyperglycemia is mainly through oxidative stress to promote the expressions of fibrosis-related signals, including CTGF and MMP-9, in H9c2 cells. This mechanism is quite different with that induced by receptor activation, both LPS and EPO.

In conclusion, we identified that STAT3 phosphorylation is rapidly raised by LPS or EPO via receptor-mediated signaling, but different from high glucose, in H9c2 cells. Additionally, STAT3 increased by hyperglycemia needs a longer time because it is mainly through an accumulation of

![Figure 3. Changes of p-STAT3 and STAT3 expressions in H9c2 cells after exposure to high glucose at indicated time points. (A) The representative changes in p-STAT3 and STAT3 expressions from H9c2 cells after incubation with high glucose (30 mM) at the indicated times (h=0-24). The protein levels, using p-STAT3 over STAT3 or STAT3 over β-actin, are indicated as mean ± SEM (n=4 per group) in each column. (B) Related mRNA expression as detected using RT-PCR and the quantified mRNA level of STAT3 is represented as mean ± SEM (n=6 per group). *P<0.05 compared to the control (at 0 h). p-, phosphorylated; STAT3, signal transducer and activator of transcription 3; SEM, standard error of the mean.]
oxidative stress which is effective to promote the transcription of downstream signals for fibrosis, including CTGF and MMP-9, in H9c2 cells. Therefore, we suggest that phosphorylation of STAT3 seems suitable for rapidly identification of

Figure 4. Effects of high glucose on changes in expressions of STAT3 and associated downstream signals in H9c2 cells. (A) The representative changes in p-STAT3, STAT3, CTGF and MMP-9 expressions after a 24-h incubation with high glucose at the indicated concentration are shown in the upper panel. Similar incubation with manitol (30 mM) was demonstrated to produce the same osmolarity as high glucose (30 mM). The protein levels, using p-STAT3 over STAT3 or each signal over β-actin, are indicated as mean ± SEM (n=4 per group) in each column. (B) Related mRNA expression as detected using RT-PCR and the quantified mRNA level of STAT3 is represented as mean ± SEM (n=6 per group). *P<0.05 compared to the control incubated in normal medium (at 0 concentration of glucose). STAT3, signal transducer and activator of transcription 3; p-, phosphorylated; CTGF, connective tissue growth factor; MMP-9, matrix metalloproteinase-9; SEM, standard error of the mean.
receptor-mediated signaling while the nuclear STAT3 is more reliable in cardiac cells receiving hyperglycemic stress. Furthermore, nuclear STAT3 may be a potential clinical indicator of cardiac fibrosis and heart dysfunction. The developments of new drugs that not only prevent myofibroblast formation but also alleviate the hyperglycemia-induced STAT3 phosphorylation may be useful to prevent cardiac dysfunction.

Figure 5. Effects of antioxidant tiron on the hyperglycemia-induced changes in expressions of STAT3 and associated downstream signals in H9c2 cells. (A) The representative changes in signal expressions modified by tiron at indicated two concentrations after a 24-h incubation with high glucose (30 mM) shown in the upper panel and the protein levels are indicated as mean ± SEM (n=4 per group) in each column. (B) Related expression of the quantified mRNA level for STAT3 by RT-PCR is represented as mean ± SEM (n=6 per group). *P<0.05 and **P<0.01 compared to the control that incubated in normal medium (first column), respectively. STAT3, signal transducer and activator of transcription 3; p-, phosphorylated; CTGF, connective tissue growth factor; MMP-9, matrix metalloproteinase-9; SEM, standard error of the mean.
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