Pioglitazone inhibits the expression of matrix metalloproteinase-9, a protein involved in diabetes-associated wound healing

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Abstract. Matrix metalloproteinase-9 (MMP-9) is a protein involved in diabetes-associated wound healing. The present study aimed to determine whether pioglitazone, an agonist of peroxisome proliferator-activated receptor- γ (PPAR- γ), inhibits the expression of MMP-9. HaCaT cells at a density of 6x10⁵ cells/well were seeded into 6-well plates in medium and were cultured for 24 h. The cells were then treated with bovine serum albumin (BSA) only or advanced glycation end-product (AGE)-BSA (50, 100, 200, 300 or 400 µg/ml), with or without pioglitazone (0.5 or 1 μ M). The effects of AGE-BSA on cell viability were determined by the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay. The levels of MMP-9 secreted into the medium were detected by an enzyme-linked immunosorbent assay. The mRNA and protein levels were analyzed by quantitative polymerase chain reaction (qPCR) and western blot analysis, respectively. AGEs are able to increase the level of MMP-9 mRNA in HaCaT cells and the levels of MMP-9 protein secreted into the medium. Pioglitazone (0.5 or $1 \,\mu\text{M}$) significantly inhibited the levels of MMP-9 in the treated HaCaT cells. Pioglitazone (0.5 or $1 \mu M$) also suppressed the levels of MMP-9 in the cell culture medium. Pioglitazone at concentrations of 0.5 and 1 μ M significantly suppressed the levels of MMP-9 mRNA to 20 or 8%, respectively. These results suggest that pioglitazone is able to effectively suppress the expression of MMP-9 via a transcriptional mechanism.

Introduction

Impaired wound healing is frequently reported in patients with diabetes. The increase in matrix metalloproteinase-9 (MMP-9) expression is associated with diabetes-associated

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wound healing (1). Previous experimental results indicate that MMP-9 expression is elevated during wounding in diabetic rats (2). Furthermore, exogenous MMP-9 expression is able to exacerbate chronic wounding (3). Since increased levels of MMP-9 are a factor contributing to poor wound healing in diabetic foot ulcers (4,5), reagents that decrease MMP-9 expression levels may be a useful method to cure the impaired wound healing in diabetic patients.

Pioglitazone is an agonist of the peroxisome proliferator-activated receptor-γ (PPAR-γ) that is a ligand-dependent transcription factor (6). The activation of PPAR-γ protects pancreatic β-cells from cytotoxicity by preventing nuclear factor (NF)- κ B activation (7-10). Pioglitazone is an antidiabetic agent, which improves insulin production in patients with diabetes. Pioglitazone also increases insulin sensitivity, thus it elevates glucose uptake and inhibits hepatic glucose output (11). It has been reported that pioglitazone is also able to reduce oxidative stress (12-16).

Advanced glycation end-products (AGEs) are a group of heterogeneous compounds that are derived from the non-enzymatic reaction of reducing sugars with proteins, lipids or nucleic acids (17). AGEs contribute to the development of various vascular diabetic complications through increasing the production of reactive oxygen species (ROS), the formation of cross-links between molecules in the basement membrane and the extracelluar matrix, and by affecting various cellular signaling pathways through the receptor for AGEs (RAGE) (18). The binding of AGEs to RAGE triggers oxidative stress and activates the transcription factor NF- κ B, thus, promoting the expression of pro-inflammatory mediators and local cellular responses (19,20).

In the present study, human keratinocytes were treated with pioglitazone in the presence of AGEs. It was demonstrated that pioglitazone decreases the expression of MMP-9 induced by the treatment of AGEs. The results suggest that pioglitazone may have therapeutic effects on impaired wound healing associated with diabetes via a mechanism of inhibiting the expression level of MMP-9.

Materials and methods

Cells and reagents. Human HaCaT keratinocytes, which were provided by Xiangya Hospital (Changsha, China), were cultured in Dulbecco's modified Eagle's medium (DMEM)

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supplemented with 10% fetal bovine serum (Gibco-BRL, Carlsbad, CA, USA), penicillin (100 U/ml) and streptomycin (100 μ g/ml). For the serum-starving experiments, cells at 80% confluence were cultured by overnight incubation in serum-free DMEM containing 0.5 mg/ml bovine serum albumin (BSA; Calbiochem, La Jolla, CA, USA). Pioglitazone and AGE-BSA were purchased from Sigma (St. Louis, MO, USA). Antibodies against MMP-9 and β -actin were purchased from Santa Cruz Biotechnology, Inc. (Santa Cruz, CA, USA).

Cell treatments and the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. HaCaT cells at a density of 6×10^5 cells/well were seeded into 6-well plates in medium and were cultured for 24 h. The cells were then treated with BSA only or AGE-BSA (50, 100, 200, 300 and 400 μ g/ml), with or without pioglitazone (0.5 μ M). At the end of each experiment, cells were incubated with 0.5 mg/ml MTT at 37°C for 4 h. The MTT kit was purchased from Invitrogen Life Technologies (Carlsbad, CA, USA). The supernatants were discarded and 50 μ l dimethylsulfoxide was added into each well. The 96-well plates (Asahi Glass Corp, Tokyo, Japan). were agitated for 10 min. The growth status and morphological changes of the cells were detected under an inverted microscope (Olympus, Tokyo, Japan). The absorbance was determined at 540 nm using a Synergy HT microplate reader (Molecular Devices, Sunnyvale, CA, USA). The viability of treated cells was expressed relative to the control cells treated with BSA (relative viability).

Quantitative polymerase chain reaction (qPCR). Total RNA was harvested from cells using the RNeasy mini kit (Qiagen, Valencia, CA, USA) according to the manufacturer's instructions. RNA (1 μ l) was reverse transcribed into cDNA using random primers with a Reverse Transcription II system purchased from Promega Corporation (Madison, WI, USA) according to the manufacturer's instructions. qPCR was conducted using an ABI Prism Sequence Detection system (Applied Biosystems, Foster City, CA, USA). An assay reagent containing premixed primers and a VIC-labeled probe (Applied Biosystems; cat. no. 4310884E) was used to quantify the expression of endogenous GAPDH mRNA. The amplification of the MMP-9 cDNA and the endogenous GAPDH cDNA was determined with FAM and VIC fluorescent intensities, respectively. The relative quantity of MMP-9 transcripts was normalized to the quantity of GAPDH mRNA at the same conditions. The primers used were as follows: Forward: 5'-GCACGACGTCTTCCAGTACC-3' and reverse: 5'-CAGGATGTCATAGGTCACGTAGC-3' for MMP-9. The experiments were repeated independently at least three times.

Immunoblotting assays. Total proteins were harvested from supernatants or from cells, separated on 10% SDS-PAGE gels and then subjected to immunoblot analysis. The primary antibodies against MMP-9 and β -actin were purchased from Santa Cruz Biotechnology, Inc. (anti-MMP-9; cat. no. sc-21733; 1:200; anti- β -actin; cat. no. sc-130301; 1:10,000). The secondary antibodies used in the present study were goat anti-mouse IgG conjugated to horseradish peroxidase antibodies (cat. no. sc-2005; 1:10,000; Santa Cruz Biotechnology, Inc.). Bound antibodies were detected using an enhanced chemiluminescence system (Pierce Biotechnology, Inc., Rockford, IL, USA). The experiments were repeated independently at least three times. Image quantifications were performed using ImageQuant software (GE Healthcare Life Sciences, Piscataway, NJ, USA).

Enzyme-linked immunosorbent assay (ELISA). The MMP-9 concentrations in the medium of the HaCaT keratinocytes treated with BSA, AGE-BSA or in the presence or absence of pioglitazone (0.5 or 1 μ M) were determined using commercially available ELISA kits (MMP-9 ELISA kit; Raybiotech, Norcross, GA, USA). The experiment was repeated independently at least six times. The values are expressed as the mean \pm standard deviation (SD).

Statistical analysis. The experimental data are expressed as the mean \pm SD. Statistical software (SPSS 10.0; SPSS, Inc., Chicago, IL, USA) was used for independent sample t-tests. P<0.05 was considered to indicate a statistically significant difference.

Results

AGEs increase the expression of MMP-9 transcripts in HaCaT cells. AGEs contribute to the development of various vascular diabetic complications and high MMP-9 expression exacerbates chronic wounds. To determine whether AGE-BSA affects the expression of MMP-9 in HaCaT cells, the HaCat cells were treated with BSA only or AGE-BSA (50, 100, 200, 300 and 400 μ g/ml) for 24 h. As shown in Fig. 1A, AGE-BSA treatment was not observed to significantly affect the cell viability, indicating that such dosages of AGE-BSA do not result in non-specific cytotoxicity. However, the qPCR results (Fig. 1B) indicated that AGE-BSA at concentrations of 300 or 400 μ g/ml markedly increased the transcript levels of MMP-9 in the cells. These results suggest that AGEs are able to increase the level of MMP-9 mRNA in HaCaT cells.

AGEs increase the levels of MMP-9 protein secreted into the medium. To determine whether AGEs increase the levels of MMP-9 protein secreted into the medium, HaCaT cells were seeded into 6-well plates at a density of 6x10⁵ cells/well, serum-starved overnight and then cultured for 24 h with various concentrations (50, 100, 200, 300 and 400 μ g/ml) of AGE-BSA or unmodified BSA. The medium was collected 24 h after treatment with AGE-BSA or unmodified BSA. The levels of MMP-9 in the culture were detected by ELISA. As shown in Fig. 2A, the ELISA results suggested that AGE-BSA increased the levels of MMP-9 in the medium. The levels of MMP-9 secreted into the medium were upregulated by up to four-fold upon treatment with AGE-BSA at a concentration of 400 μ g/ml. The western blot analysis results (Fig. 2B) also indicated that AGE-BSA (300 or 400 μ g/ml) significantly increased the levels of the secreted MMP-9 proteins. These results suggest that AGEs are able to increase the levels of MMP-9 protein secreted into the medium.

Pioglitazone is able to reduce the high levels of MMP-9 protein induced by AGE. In order to determine whether



Figure 1. AGE-BSA induces the expression of MMP-9 in HaCaT cells. HaCaT cells at a density of $6x10^5$ cells/well were seeded into 6-well plates, serum-starved overnight and then cultured for 24 h with various concentrations (50, 100, 200, 300 and 400 μ g/ml) of AGE-BSA or unmodified BSA. (A) Cell viability was measured using the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide assay following the treatment. (B) Total RNA was harvested from HaCaT cells treated with AGE-BSA or unmodified BSA only. Quantitative polymerase chain reaction was performed to analyze the MMP-9 mRNA levels in the cells treated with various concentrations (50, 100, 200, 300 and 400 μ g/ml) of AGE-BSA orouge to analyze the MMP-9 mRNA levels in the cells treated with various concentrations (50, 100, 200, 300 and 400 μ g/ml) of AGE-BSA compared with the BSA only control. The levels (mean value) of MMP-9 transcripts in cells were calculated. Error bars show the mean ± standard deviation (P<0.05). The experiments were repeated at least three times. *P<0.05 vs. the corresponding control. MMP-9, matrix metalloproteinase-9; AGE, advanced glycation end-products; BSA, bovine serum albumin.



Figure 2. Levels of MMP-9 protein in the medium as determined by ELISA and western blot analysis. HaCaT cells at a density of $6x10^5$ cells/well were seeded into 6-well plates, serum-starved overnight and then cultured for 24 h with various concentrations (50, 100, 200, 300 and 400 µg/ml) of AGE-BSA or unmodified BSA. (A) Medium was collected 24 h post-treatment with AGE-BSA or unmodified BSA. The levels of MMP-9 in the culture were detected by ELISA. The data (mean ± standard deviation) are from six independent experiments. (B) Medium was collected and concentrated. The total proteins were isolated from the medium and subjected to western blot analysis. Primary antibodies against MMP-9 and β -actin were purchased from Santa Cruz Biotechnology, Inc, Santa Cruz, CA, USA). The secondary antibodies used in the present study were goat anti-mouse IgG conjugated to horseradish peroxidase. Bound antibodies were detected using an enhanced chemiluminescence system (Pierce Biotechnology, Inc.). The experiments were repeated independently at least three times. Image quantifications were performed using ImageQuant software. MMP-9, matrix metalloproteinase-9; ELISA, enzyme-linked immunosorbent assay; AGEs, advanced glycation end-products; BSA, bovine serum albumin.

pioglitazone is able to decrease MMP-9 expression, HaCaT cells at a density of $6x10^5$ cells/well were serum-starved overnight and then cultured for 24 h with 300 μ g/ml AGE-BSA in the absence or presence of pioglitazone (0.5 or 1 μ M). The total proteins were harvested from the cells and then subjected to western blot analysis. As shown in Fig. 3A, pioglitazone (0.5 μ M) significantly inhibited the MMP-9 level. A higher concentration of pioglitazone (1 μ M) resulted in a greater inhibitory effect on the MMP-9 level. The levels of MMP-9 in the medium were also measured by ELISA. As shown in Fig. 3B, pioglitazone (0.5 or 1 μ M) significantly suppressed the levels of MMP-9 in the medium. These results suggest that pioglitazone significantly inhibits the expression of MMP-9.

Pioglitazone reduces the protein expression of MMP-9 induced by AGEs. To further investigate the mechanisms underlying the inhibitory effect of pioglitazone on the increased MMP-9 expression induced by AGEs, HaCaT cells at a density of $6x10^5$ cells/well were serum-starved overnight and then cultured for 24 h with 300 µg/ml AGE-BSA in the absence or presence of pioglitazone (0.5 or 1 µM). Total RNA was harvested from HaCaT cells and qPCR was performed to analyze the mRNA levels of MMP-9. As shown in Fig. 4, the qPCR results indicated that pioglitazone at concentrations of 0.5 and 1 µM significantly suppressed the levels of MMP-9 mRNA to 20 and 8%, respectively. These results suggest that pioglitazone suppresses the expression of MMP-9 via a transcriptional mechanism.



Figure 3. Effects of pioglitazone on the high level of MMP-9 protein induced by AGE. HaCaT cells at a density of $6x10^5$ cells/well were serum-starved overnight and then cultured for 24 h with 300 µg/ml AGE-BSA in the absence or presence of pioglitazone (0.5 or 1 µM). (A) Total proteins were harvested from the cells and then subjected to western blot analysis. (B) The levels of MMP-9 in the medium were also measured by enzyme-linked immunosorbent assay. The experiments were repeated at least six times. *P<0.05, compared with the cells treated with AGE-BSA. MMP-9, matrix metalloproteinase-9; AGEs, advanced glycation end-products; BSA, bovine serum albumin.



Figure 4. qPCR detection of MMP-9 mRNA. HaCaT cells at a density of $6x10^5$ cells/well were serum-starved overnight and then cultured for 24 h with 300 µg/ml AGE-BSA in the absence or presence of pioglitazone (0.5 or 1 µM). Total RNAs were harvested from HaCaT cells. qPCR was performed to analyze the MMP-9 mRNA levels in the cells. The levels (mean value) of MMP-9 transcripts in the cells were calculated. Error bars show the mean ± standard deviation (P<0.05). The experiments were repeated at least three times. *P<0.05, when compared with the cells treated with AGE-BSA. qPCR, quantitative polymerase chain reaction; MMP-9, matrix metalloproteinase-9; AGEs, advanced glycation end-products; BSA, bovine serum albumin.

Discussion

As a PPAR agonist, pioglitazone has demonstrated promise as a therapeutic agent due to its ability to increase the functional recovery of wounds and decrease lesion sizes following injury (21). The therapeutic effects of pioglitazone are considered to be a result of the regulation of multiple pathways (11-16). Since the increased expression of MMP-9 contributes to poor wound healing in diabetic foot ulcers (4,5), the present study examined whether pioglitazone acts via a mechanism associated with the regulation of the expression of MMP-9 in human keratinocytes treated with AGEs. The results revealed that pioglitazone at concentrations of 0.5 or 1 μ M suppressed the levels of MMP-9 mRNA to 20 or 8%, respectively. Since the increased expression of MMP-9 contributes to poor wound healing (4,5), the results of the present study suggest that pioglitazone may have therapeutic effects on impaired wound healing associated with diabetes via a mechanism of inhibiting MMP-9 expression. This finding provides novel evidence for the application of pioglitazone in the field of wound healing.

It is reported that pioglitazone may inhibit the TGF- β -induced myofibroblast differentiation (22). Pioglitazone also attenuates TGF- β -induced type I collagen and fibronectin mRNA and protein production, which are involved in burn wound healing (22,23). It was revealed that PPAR- γ -dependent and PPAR- γ -independent mechanisms were involved in the action of pioglitazone (22-24), although PPAR- γ agonists do not prevent the activation of quiescent hepatic stellate cells *in vitro*, nor hepatic fibrogenesis in mice (24).

In addition to the diabetes-associated wound healing, pioglitazone may be useful for other types of wound healing, including gastric ulcer healing. The involvement of PPAR- γ in inflammatory responses during pioglitazone-mediated gastric ulcer healing has been reported (25). In the present study, the findings suggest that use of pioglitazone has potential in the wound healing therapy and it may be a promising approach upon further study. The results provide novel evidence for understanding the molecular mechanisms underlying the action of pioglitazone.

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