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Activation

Running head: Pioglitazone and vocal fold fibrosis

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Abstract

Macrophages aid in wound healing by changing their phenotype and can be a key driver of fibrosis. However, the contribution of macrophage phenotype to fibrosis following vocal fold injury remains unclear. Peroxisome proliferator-activated receptor- γ (PPAR γ) is expressed mainly by macrophages during early wound healing and regulates the macrophage phenotype. This study aimed to evaluate the effects of pioglitazone, a PPARy agonist, on the macrophage phenotype and fibrosis following vocal fold injury in rats. Pioglitazone was injected into the rats' vocal folds on days 1, 3, 5, and 7 after injury, and the vocal fold lamina propria was evaluated on days 4 and 56 after injury. Moreover, THP-1-derived macrophages were treated with pioglitazone, and the expression of pro-inflammatory cytokines under lipopolysaccharide/interferon-y stimulation was analyzed. The results revealed that pioglitazone reduced the expression of Ccl2 both in vivo and in vitro. Furthermore, pioglitazone decreased the density of inducible nitric oxide synthase⁺ CD68⁺ macrophages and inhibited the expression of fibrosis-related factors on day 4 after injury. On day 56 after injury, pioglitazone inhibited fibrosis, tissue contracture, and hyaluronic acid loss in a PPARy-dependent manner. These results indicate that PPARy activation could inhibit accumulation of inflammatory macrophages and improve tissue repair. Considered together, these findings imply that inflammatory macrophages play a key role in vocal fold fibrosis.

Introduction

Voice disorders worsen the quality of life by impairing communication and social functioning, which negatively affect mental health^{1,2}. Vocal fold (VF) scarring, which occurs following inflammation, injury, and surgical resection, can cause irreversible voice impairment^{3,4}. This results from excessive extracellular matrix (ECM) collagen deposition in the lamina propria, causing scarring and abnormal VF vibration^{5,6}. Fibroblasts are the main producers of ECM⁷, and previous studies targeting fibrosis prevention have primarily focused on them^{8–11}. Treatments using growth factors and stem cells prevent fibrosis to some extent^{10,12–14}, however, they have not resulted in ideal tissue repair.

Fibrosis is an outcome of inflammatory and repair responses, and recent studies have demonstrated that macrophages are involved in both^{15–17}. They change from pro- to antiinflammatory phenotype during the wound healing process¹⁸, contributing to the development of pathological fibrosis in various organs and injury models¹⁶. Therefore, we hypothesized that fibrosis could be regulated by controlling the macrophage phenotype, which functions as the key driver. Macrophages also interact with VF fibroblasts¹⁹ and VF-derived mesenchymal stem cells^{20,21} *in vitro*.

Peroxisome proliferator-activated receptor- γ (PPAR γ) is a transcription factor that is prominently expressed by macrophages during early wound healing²², regulating the phenotype. PPAR γ activation suppresses the macrophage pro-inflammatory (M1-like) phenotype under Th1 and Th17 polarizing conditions and promotes the anti-

inflammatory/reparative (M2-like) phenotype under Th2²³.

Pioglitazone (PIO), a PPARγ agonist, is an antidiabetic drug in clinical use and has anti-inflammatory and anti-fibrotic effects in various disease models^{24,25}. However, no studies have focused on the macrophage phenotype that prevents VF scar formation *in vivo*. Therefore, this study aimed to evaluate the effect of PIO, a macrophage phenotype regulator, on fibrosis in a rat model of VF injury.

Materials and Methods

<u>Animals</u>

Male Sprague-Dawley rats (350–450 g) were purchased from Shimizu Laboratory Supplies Co., Ltd. (Kyoto, Japan). The 13-week-old rats underwent surgical and injection procedures, while naïve (uninjured and untreated) ones were age-matched in each experiment. The rats were housed in a specific pathogen-free environment under controlled temperature, humidity, and light, and were given standard rodent food and water *ad libitum*. No rats died during the observation period. All the protocols and procedures were compliant with the Guidelines for the Proper Conduct of Animal Experiments of the Science Council of Japan. Ethical approval was obtained from the Animal Research Committee of Kyoto University (approval number: MedKyo20137).

Surgical procedure

The rat VF injury model is often used as a preclinical model for VF fibrosis^{5,26}. Bilateral VF mucosal injuries were performed transorally, as previously described^{26,27}. The rats were anesthetized with an intraperitoneal injection of 2.5 mg/kg butorphanol (Meiji Seika Pharma Co., Ltd., Tokyo, Japan), 2.0 mg/kg midazolam (Astellas Pharma Inc., Tokyo, Japan), and 0.15 mg/kg medetomidine (Nippon Zenyaku Kogyo Co., Ltd., Fukushima, Japan). They were then placed in a near-vertical position on a custom-made operating table. The larynx was observed with a 1.9 mm diameter, 30-degree rigid endoscope (Hopkins Telescope 1232BA, Karl Storz, Tuttlingen, Germany). The bilateral VFs were stripped with a custom-made 0.2 mm diameter tip needle²⁸ until the VF thyroarytenoid muscle was exposed. Drug treatment

PIO (P1901, Tokyo Chemical Industry, Tokyo, Japan) and GW9662 (M6191; Sigma-Aldrich, St. Louis, MO, USA) were dissolved in saline containing 0.5% dimethyl sulfoxide. The rats were randomly assigned to treatment with 5 μ M PIO, 50 μ M PIO, 50 μ M PIO with GW9662 or vehicle (0.5% dimethyl sulfoxide). Bilateral VF injections were administered on days 1, 3, 5, and 7 post-injury (Figure 1), using 10 μ L microsyringes with a 33 G needle (Hamilton Company, Reno, NV). Animal positions and injection techniques were based on previous studies²⁹. VFs were observed using a rigid endoscope under general anesthesia, as described above.

Histology and indirect immunofluorescence

Larynges were harvested at days 4 and 56 post-injury, corresponding to the inflammatory and scar maturation phases of wound healing in the rat model, respectively^{26,27,30}. Rats were deeply anesthetized with pentobarbital and transcardially perfused with phosphate-buffered saline (PBS) followed by 4% paraformaldehyde. Larynges were dissected, placed in 4% paraformaldehyde overnight at 4 °C, and were then incubated in graded sucrose solutions overnight at 4 °C again. Tissues were snap-frozen in the optimum cutting temperature compound (Sakura Finetek, Torrance, CA). Serial 8 µm sections were cut in a cryostat and stored at -80 °C. Coronal sections containing laryngeal alar cartilage were used to assess the center of the injury, as previously described²⁷. Five rats each were assigned to all the experimental groups, and one sample per rat was employed for each analysis. One sample each from the PIO 5 µM and PIO 50 µM groups at 56 days after injury was excluded from the analysis owing to insufficient dehydration and fixation before sectioning.

Routine hematoxylin and eosin (H&E), Masson's trichrome, and Alcian blue (pH 2.5, with and without hyaluronidase digestion) histological staining were performed to evaluate morphology, collagen abundance, and hyaluronic acid (HA), respectively.

For indirect immunofluorescence, the sections were boiled in citrate buffer (0.01 M, pH 6.0) for antigen retrieval, permeabilized, and blocked with 5% normal donkey serum (NDS), 1% bovine serum albumin (BSA), and 0.2% Triton X-100 in PBS for 1 hour at room

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temperature. They were subsequently incubated with primary antibodies in PBS containing 2% NDS and 1% BSA overnight at 4 °C. Sections were washed three times, 5 minutes each, with PBS and incubated with secondary antibodies diluted in blocking buffer (2% NDS and 1% BSA) at 1:500 for 1 hour at room temperature. Nuclei were stained with 4',6-diamidino-2-phenylindole (DAPI). The slides were washed three times with PBS and mounted with Fluoromount-G (Thermo Fisher, Waltham, MA, USA).

Antibodies

Indirect immunofluorescence was performed using CD68 as a pan-macrophage marker, inducible nitric oxide synthase (iNOS) as pro-inflammatory (M1-like), and CD206 as anti-inflammatory (M2-like) macrophage markers, as previously described^{31,32}. The three primary antibodies (CD68, iNOS, and CD206) were applied simultaneously.

The primary antibodies used were anti-rat CD68 (mouse, 1:1000, MCA341GA, Serotec, Oxford, UK), anti-iNOS (rabbit, 1:50, PA1-036, Invitrogen, Carlsbad, CA, USA), anti-mouse MMR/CD206 (goat, 1:250; AF2535, R&D Systems, Minneapolis, MN, USA), anti-collagen I (rabbit, 1:100, ab34710, Abcam, Cambridge, UK), and anti-fibronectin (mouse, 1:100, ab6328, Abcam). The secondary antibodies used were donkey anti-mouse Alexa 488, donkey anti-rabbit Alexa 488, donkey anti-rabbit Alexa 568, donkey anti-mouse Alexa 568, and donkey anti-goat Alexa 647 (Invitrogen).

Microscopy and image analysis

Images were obtained at × 40 magnification using a BZ-X710 all-in-one microscope (Keyence Co., Osaka, Japan). The investigators were blinded to the experimental group during image analysis. The lamina propria was defined as the area between the basement membrane and the thyroarytenoid muscle in the coronal section of the VFs. BZ-X analyzer software (Keyence) was used to count the number of stained cells and measure the area with the same threshold determined for each staining. The threshold value for a positive result was determined based on the background signal intensity and the negative control for each staining.

To evaluate the polarization of M1 and M2, we counted the number of cells with specific marker positivity. M1-like macrophages were defined as iNOS⁺ CD68⁺ cells, whereas M2-like ones were defined as CD206⁺ CD68⁺cells. The number of macrophages, including M1-like and M2-like ones, were manually counted. H&E staining was used to measure the area of VF lamina propria. The positive area ratio of Masson trichrome staining, Alcian blue staining, and immunostaining for collagen I and fibronectin were analyzed. <u>RNA isolation and quantitative real-time polymerase chain reaction (RT-PCR) (*in vivo*)</u>

VF mucosa samples were collected as previously described³³. Total RNA was isolated using the RNeasy Micro kit (Qiagen, Valencia, CA, USA) with on-column DNase I (Qiagen) digestion according to the manufacturer's instructions. RNA concentration and purity were determined using a spectrophotometer (ND-1000; NanoDrop, Wilmington, DE,

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USA). cDNA was synthesized using a High-Capacity cDNA Reverse Transcription Kit (Applied Biosystems, Foster City, CA, USA) according to the manufacturer's instructions. Five rats each were assigned to all the experimental groups, and one sample per rat was used for each analysis. One sample each from the vehicle and PIO 50 μ M + GW9662 groups was excluded from the analysis because of suspected contamination during RNA isolation. Quantitative RT-PCR was performed on a StepOnePlus RT-PCR system (Applied Biosystems) using THUNDERBIRD Probe qPCR Mix (Toyobo, Osaka, Japan) and TaqMan gene expression assay (Thermo Fisher), according to the manufacturer's instructions. Each assay was performed in duplicate or triplicate. The expression levels of each gene were normalized to the levels of $Sdha^{34}$ and analyzed using the $\Delta\Delta$ Ct method³⁵. The following probes were used: Illb (R n00580432 m1), Tnf (Rn99999017 m1), Arg1 (Rn00691090 m1), Tgfb1 (Rn00572010 m1), Cd68 (Rn01495634 g1), Cd36 (Rn00580728 m1), Ccl2 (Rn00580555 m1), Collal (Rn01463848_m1), Fn1 (Rn00569575_m1), Acta2 (Rn01759928 g1), Il10 (Rn00563409 m1), Has1 (Rn00597231 m1), Has2 (Rn00565774 m1), and Sdha (Rn00590475 m1).

Cell culture

Human monocytic THP-1 (American Type Culture Collections, Rockville, MD, USA) cells were maintained in suspension culture in RPMI 1640 (30264-85, Nacalai Tesque, Kyoto, Japan) supplemented with 10% fetal bovine serum (CCP-FBS-BR-500, Cosmo Bio

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Co., Tokyo, Japan), 100 U/ml penicillin and 100 μ g/ml streptomycin solution (26253-84, Nacalai Tesque), and 0.05 mM 2-mercaptoethanol (21985-023, Thermo Fisher). THP-1 cells were differentiated into M0 macrophages by treatment with 100 ng/mL phorbol 12-myristate 13-acetate (PMA) (P1585; Sigma-Aldrich) for 96 h. These cells were divided into four groups: M0, M1, M1 + PIO, and M1 + PIO + GW9662. M0 macrophages were preincubated with GW9662 (10 μ M; M6191; Sigma-Aldrich) or vehicle for 1 hour, then with PIO (10 μ M; E6910; Sigma-Aldrich) or vehicle for another 2 hours. After 3 hours of preincubation, cells were co-stimulated with 1 μ g/ml lipopolysaccharide (LPS) (E. coli O111: B4 L2630, Sigma-Aldrich) plus 20 ng/ml interferon (IFN)- γ (AF-300-02, PeproTech, Cranbury, NJ, USA) for 6 hours to generate inflammatory macrophages³⁶. Cells were lysed with RNA lysis buffer to perform quantitative PCR analyses. All the experiments were performed as 5 independent experiments.

RNA isolation and quantitative PCR (in vitro)

Total RNA from the cells was isolated using the RNeasy mini kit (Qiagen) according to the manufacturer's instructions. The RNA concentration and purity were assessed by an ultraviolet spectrophotometer. Total RNA (1 µg) was reverse-transcribed using ReverTra Ace qPCR RT Master Mix with gDNA Remover (Toyobo). Quantitative PCR was performed on a QuantStudio 3 Real-Time PCR system (Applied Biosystems) using THUNDERBIRD Probe qPCR Mix (Toyobo) and TaqMan gene expression assay (Thermo Fisher), according to

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the manufacturer's instructions. Each assay was performed in triplicate. The expression levels of each gene were normalized to the levels of *Gapdh* and analyzed using the $\Delta\Delta$ Ct method. The following probes were used: *Il1b* (Hs01555410_m1), *Tnf* (Hs00174128_m1), *Ccl2* (Hs00234140_m1), and *Gapdh* (Hs99999905_m1).

Statistical Analysis

Statistical analyses and graphic representations were conducted using GraphPad Prism version 9.3.0 (GraphPad Software, San Diego, CA, USA). The mean values and standard error of the mean were calculated. Each data point derived from the quantitative PCR assays represents an average of 2 or 3 technical replicates, and data were averaged over independently replicated experiments (n = 4–5 independently collected samples) and analyzed by one-way analysis of variance followed by post-hoc Dunnett's test for comparison with the vehicle-treated control group. *P* values of **P* < 0.05, ***P* < 0.01, ****P* < 0.001, were selected to indicate statistical significance.

Results

PIO reduces injury-induced macrophage accumulation

In the rat VF injury model, macrophages increased in the lamina propria on days 1 – 5 after injury¹⁰. The effects of PIO on macrophage infiltration were observed at day 4 after injury in the VF lamina propria. Indirect immunofluorescence revealed that PIO reduced the

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injury-induced increase in density of CD68+ macrophages (Figure 2A, B). Quantitative RT-PCR demonstrated that PIO inhibited the injury-induced increase in *Cd68* and *Ccl2* gene expression. This effect of PIO was abolished by GW9662 (Figure 2C).

PIO inhibits injury-induced increase in M1-like macrophages

To assess the effects of PIO on macrophage phenotypes at day 4 after injury, proinflammatory M1-like and reparative M2-like macrophages were immunostained with iNOS and CD206, respectively (Figure 3A, D). PIO attenuated the injury-induced increase in density of iNOS⁺ macrophages (Figure 3B). Moreover, 5 μM PIO, but not 50 μM PIO, suppressed the injury-induced increase in density of CD206⁺ macrophages (Figure 3C). In the naïve VF, the majority of macrophages consisted of iNOS⁻ CD206⁺ macrophages (60%), and there were few iNOS⁺ CD206⁻ macrophages (4.1%). Following injury, the percentage of iNOS⁻ CD206⁺ macrophages markedly decreased (20.1%), and that of iNOS⁺ CD206⁻ macrophages (19.7%) and iNOS⁻ CD206⁻ macrophages (36.1%) increased. PIO treatment increased the percentage of iNOS⁻ CD206⁺ macrophages (PIO 50 μM, 33.7%) and decreased that of iNOS⁺ CD206⁻ macrophages (PIO 50 μM, 9.9%). iNOS⁺ CD206⁺ macrophages were present in all the experimental groups (Figure 3E).

Quantitative RT-PCR was performed to examine the expression of genes related to pro- and anti-inflammatory mediators. Gene expression of *Il1b*, *Tnf*, *Il10*, *Arg1*, and *Cd36* was not significantly different in the PIO (50 μ M) and GW9662+PIO (50 μ M) groups compared to the vehicle group (Figure 4A).

PIO treatment attenuates fibrosis

To evaluate the anti-fibrotic effect of PIO, the expression of fibrosis-related factors and ECM components in the VF lamina propria was evaluated. Quantitative RT-PCR revealed that PIO inhibited the injury-induced increase in expression of Tgfb1 and Acta2 and the effect of PIO was inhibited by GW9662. PIO inhibited the injury-induced increase in Fn1expression, however, the effect of PIO was not abrogated by GW9662 (Figure 4B). The effect of PIO on *Collal* expression was not significant.

Indirect immunofluorescence indicated that PIO inhibited the injury-induced increase in the percentage of collagen type 1 (Figure 5A) and fibronectin (Figure 5B)-positive areas, and the effect of PIO was reversed by GW9662. Masson's trichrome staining revealed that PIO inhibited the injury-induced increase in the percentage of collagen deposition, and the effect of PIO was abrogated by GW9662 (Figure 5C).

PIO prevents contracture and hyaluronic acid level reduction in the VF lamina propria

The degree of atrophy in the VF lamina propria was assessed by the cross-sectional area on day 56 after injury. H&E staining revealed that PIO inhibited the injury-induced decrease in cross-sectional area, and GW9662 reversed the effect of PIO (Figure 5D). Alcian blue staining revealed that 50 μ M PIO inhibited the injury-induced decrease in HA level, and the effect of PIO was reversed by GW9662 (Figure 5E). Moreover, 50 μ M PIO inhibited the

injury-induced increase in Has1 expression, and GW9662 reversed the effect of PIO (Figure 4B).

PIO prevents LPS/IFN-γ-induced upregulation of Ccl2 in THP-1-derived macrophages

To evaluate the effect of PIO on human macrophages, the expression of *Ccl2*, *Il1b*, and *Tnf* in THP-1-derived macrophages under LPS/IFN γ stimulation was examined by quantitative PCR. PIO inhibited the LPS/IFN- γ -induced increase in *Ccl2* expression in THP-1-derived macrophages, but the effect of PIO was not reversed by GW9662. PIO enhanced the LPS/IFN- γ -induced increase in *Il1b* expression in THP-1-derived macrophages, and GW9662 reversed the effect of PIO (Figure 6).

Discussion

This study evaluated the effects of PIO, a PPAR γ agonist, on macrophage infiltration and phenotype, and tissue repair following VF injury. Results showed that PIO suppressed inflammatory macrophage accumulation and ECM-related gene expression in the acute phase. Furthermore, PIO inhibits fibrosis, contractures, and HA loss in the VF lamina propria during the scar maturation phase via PPAR γ . These findings suggest that treatments targeting inflammatory macrophages or PPAR γ activation in macrophages may prevent VF fibrosis.

Macrophages contribute to both inflammation and repair in the wound healing process. Depletion experiments in various organs have been performed using knockout mice

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or liposomal clodronate, and revealed that macrophages are involved in fibrosis^{37–39}. Although drug-induced fibrosis has been ameliorated by macrophage withdrawal during the fibrotic phase, injury-induced fibrosis, which has been tested primarily in the skin, has been reduced within 7 days during the acute phase^{40,41}. Macrophage function in VFs has not been investigated in detail, however, a transient increase in iNOS-positive inflammatory macrophages has been reported from days 1 to 5 after injury⁴². Therefore, the therapeutic strategy in this study was focused on controlling inflammatory macrophages within 7 days of the acute phase. As expected, PIO treatment reduced the accumulation of macrophages, especially inflammatory macrophages, in the acute phase. However, unexpectedly, PIO treatment did not suppress the expression of *Il1b* and *Tnf*, which are upregulated in M1 macrophages⁴³.

PIO inhibited the injury-induced upregulation of *Ccl2* expression via PPARγ *in vivo* and LPS/IFNγ-induced upregulation of *Ccl2* expression in a PPARγ-independent manner *in vitro*. These results are consistent with previous findings that PIO reduces CCL2 production^{44,45}. *Ccl2* is transcriptionally regulated by NF κ B^{46,47}. PPARγ ligands repress the transcriptional activity of NF κ B and NF κ B target genes in macrophages in a PPARγ-dependent^{48,49} or PPARγ-independent manner^{50,51}, depending on their concentration and the stimulus applied to macrophages⁵². It is noteworthy that PIO repressed the expression of *Ccl2* in both *in vivo* experiments in rats and *in vitro* experiments in human macrophages, however,

noted that the signaling pathway of PIO could be different owing to the diverse conditions of the two experiments.

PIO inhibited the injury-induced upregulation of fibrosis-related genes and fibrosis, and Ccl2 may be associated with this effect of PIO. Ccl2 is reported to recruit and locally proliferate bone marrow-derived monocytes and macrophages via activation of C-C chemokine receptor type 2 (CCR2)^{53,54}. iNOS⁺ macrophages differentiate from bone marrowderived monocytes in a CCR2-dependent manner⁵⁵. These findings suggest that PIO reduces the accumulation of iNOS⁺ macrophages by inhibiting *Ccl2* expression. Furthermore, studies using lung and heart injury models have revealed that bone marrow-derived macrophages lack repair capacity, whereas tissue-resident macrophages exhibit excellent repair capacity^{56,57}. Based on these findings, PIO could exhibit improved repair capacity by preventing the accumulation of bone marrow-derived macrophages that lack repair capacity, resulting in reduced fibrosis and preserved HA. Further studies using genetic phylogenetic tracking techniques are needed. Ccl2 has also been reported to amplify the expression of *Tgfb1* in human monocytes through autocrine and paracrine effects with TGF- β 1⁵⁸. In the VFs, TGF-β1 differentiates fibroblasts into myofibroblasts⁵⁹, induce a contractile phenotype⁷, and upregulate the expression of Fn1, $Acta2^{60}$, and $Has1^{61}$. Thus, suppression of $TGF-\beta I$ expression via Ccl2 by PIO may have inhibited the initiation of the fibrotic cascade, resulting in reduced tissue contraction. It is important to note that VF atrophy can cause dysphonia,

based on the size of the glottal gap during phonation⁶².

This study had several limitations. PIO treatment may affect cells other than macrophages, as PPARγ agonists have been reported to act on adipocytes^{63,64}, fibroblasts^{65,66}, T cells⁶⁷, and vascular smooth muscle cells⁶⁸. However, PPARγ is mainly expressed by macrophages during the inflammatory phase of wound healing²². Furthermore, studies using macrophage-specific PPARγ knockout mice have revealed that PIO treatment exerts its antiinflammatory^{69–71} and anti-fibrotic^{71,72} effects by acting on PPARγ in macrophages. Therefore, the effect of PIO treatment in this study appears to be mainly due to its effects on the macrophages. Flow cytometry analysis of VF mucosal cells may be useful in assessing the changes in the functional phenotype of macrophages and other cell populations localized to the wound. Moreover, adoptive transfer of treated macrophages may be useful to evaluate the effects of PIO focusing only on macrophages.

In this study, sex differences were not examined. Previous studies have suggested that the pharmacokinetics of PIO may differ between males and females⁷³. Furthermore, PPAR γ expression is affected by sex hormones⁷⁴ and the sexual cycle⁷⁵. Therefore, only male rats were used in this study to eliminate the effects of sex differences and sexual cycle. It remains unclear whether the effect of PIO on the macrophages differs between males and females. Hence, further research is warranted.

In summary, results of this study suggest that PIO inhibits the accumulation of

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inflammatory macrophages in the early stage of wound healing and consequently prevents fibrosis during the scar maturation phase in a rat VF injury model. These effects may be due to the modulation of *Ccl2* expression in macrophages by PIO. Inflammatory macrophages and PPAR γ activation may be potential targets for the treatment of VF fibrosis.

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Figure legends

Figure 1. Experimental timeline

Schematic illustration of the injury, drug injection, and tissue collection schedule. Vehicle, pioglitazone 5 μ M, pioglitazone 50 μ M, or pioglitazone 50 μ M with GW9662 were injected into the rat vocal folds (VFs) on days 1, 3, 5, and 7 after injury. VFs were harvested on days 4 and 56 after injury for indirect immunofluorescence, histology, and quantitative real-time polymerase chain reaction. Uninjured and untreated VFs were used as naïve controls.

Figure 2. Pioglitazone decreases macrophage accumulation in the vocal fold (VF) lamina propria on day 4 after injury. (A) Coronal sections of VF lamina propria on day 4 after injury were stained for CD68 (green) by indirect immunofluorescence. Nuclei were counterstained with 4',6-diamidino-2-phenylindole (blue). Representative images of CD68⁺ macrophages are shown. The bottom panels show magnified images of the regions indicated by dashed boxes in the middle panels. (B) Quantification of the CD68⁺ cells. (C) Quantitative real-time polymerase chain reaction analysis of mRNA levels of *Cd68* and *Ccl2* on day 4 after injury. Scale bars: 100 µm (A, top and middle panels); 50 µm (A, bottom panels). All data are presented as means \pm standard error of the mean. n = 4–5 rats per group. **P* < 0.05, ***P* < 0.01, ****P* < 0.001 compared with the vehicle-treated group.

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Figure 3. Pioglitazone reduces the percentage of M1-like macrophages in the vocal fold (VF) lamina propria on day 4 after injury. Indirect immunofluorescence analysis of macrophage phenotypes in the coronal sections of the VF lamina propria on day 4 after injury. (A) Representative images of $CD68^+$ inducible nitric oxide synthase (iNOS)⁺ (M1like) macrophages. iNOS (red), CD68 (green), and 4',6-diamidino-2-phenylindole (DAPI) (blue). The bottom panels show magnified images of the regions indicated by the dashed boxes in the middle panels. (B) Quantitative analysis of the density of CD68⁺ iNOS⁺ (M1like) macrophages. (D) Representative images of CD68⁺ CD206⁺ (M2-like) macrophages. CD206 (red), CD68 (green), and DAPI (blue). The bottom panels show magnified images of the regions indicated by the dashed boxes in the middle panels. (C) Quantitative analysis of the density of CD68⁺ CD206⁺ (M2-like) macrophages. (E) Stacked bar graphs show the proportions of iNOS⁻ CD206⁻, iNOS⁻ CD206⁺, iNOS⁺ CD206⁺, and iNOS⁺ CD206⁻ macrophages. Scale bars: 100 µm (A and D, top and middle panels); 20 µm (A and D, **bottom panels**). All the data are presented as means \pm standard error of the mean. n = 5 rats per group. *P < 0.05, ***P < 0.001 compared with the vehicle-treated group.

Figure 4. Effect of pioglitazone on the expression of genes related to inflammation and wound healing on day 4 after injury. (A) Quantitative real-time polymerase chain reaction (RT-PCR) analysis of gene expression related to M1/M2 polarization. mRNA expression levels of IL-1 β , TNF, Arg1, IL-10, and CD36 were measured. (B) Quantitative RT-PCR analysis of gene expression related to wound healing. mRNA expression levels of TGF- β 1, Fn1, Acta2, Col-1a1, Has1, and Has2 were determined. All the data are presented as means ± standard error of the mean. n = 4–5 rats per group. *P < 0.05, **P < 0.01, ***P < 0.001compared with the vehicle-treated group.

Figure 5. Pioglitazone inhibits fibrosis, tissue contracture, and hyaluronic acid loss in vocal fold (VF) lamina propria on day 56 after injury. Coronal sections of the VF lamina propria on day 56 after injury were analyzed. (A) Indirect immunofluorescence of type I collagen (red). (B) Indirect immunofluorescence of fibronectin (red). (C) Masson's trichrome staining for collagen (blue). (D) Hematoxylin and eosin staining. (E) Alcian blue staining of hyaluronic acid (light blue). Representative images (left four panels) and quantification of the percentage of the positive area (right panel) are shown. Scale bars: 200 μ m. All the data are presented as means \pm standard error of the mean. n = 4–5 rats per group. **P* < 0.05, ***P* < 0.01, ****P* < 0.001 compared with the vehicle-treated group.

Figure 6. Effects of pioglitazone on lipopolysaccharide (LPS) + interferon (IFN)- γ induced expression of *Ccl2*, *Il1b*, and *Tnf* in THP-1-derived macrophages

THP-1-derived macrophages (M0) were pretreated with or without GW9662 (10 $\mu M)$ for 1

hour and then treated with pioglitazone (10 μ M) for another 2 hours, followed by LPS + IFN- γ treatment for 6 hours. Expression of Ccl2, Il1b, and Tnf was assessed by quantitative polymerase chain reaction. All data are presented as means \pm standard error of the mean. n = 5 per group. *P < 0.05, ***P < 0.001 compared with the vehicle-treated M1 macrophage group.

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