Effects of Smad4 on the expression of caspase-3 and Bcl-2 in human gingival fibroblasts cultured on 3D PLGA scaffolds induced by compressive force

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Received July 6, 2020; Accepted November 30, 2020

DOI: 10.3892/ijmm.2021.4858

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Key words: human gingival fibroblasts, poly(lactide-co-glycolide), Smad4, caspase-3, Bcl-2, compression pressure

Abstract. Human gingival fibroblasts (HGFs) are the main cells that comprise gingival tissue, where they transfer mechanical signals under physiological and pathological conditions. The exact mechanism underlying gingival tissue reconstruction under compressive forces remains unclear. The present study aimed to explore the effects of Smad4, caspase-3 and Bcl-2 on the proliferation of HGFs induced by compressive force. HGFs were cultured on poly(lactide-co-glycolide) (PLGA) scaffolds under an optimal compressive force of 25 g/cm². Cell viability was determined via Cell Counting Kit-8 assays at 0, 12, 24, 48 and 72 h. The expression levels of Smad4, caspase-3 and Bcl-2 were measured via reverse transcription-quantitative PCR and western blotting. The application of compressive force on HGFs for 24 h resulted in a significant increase in cell proliferation and Bcl-2 expression, but a significant decrease in the expression of Smad4 and caspase-3; however, inverse trends were observed by 72 h. Subsequently, a lentivirus was used to overexpress Smad4 in HGFs, which attenuated the effects of compressive force on HGF proliferation and Bcl-2 expression, but enhanced caspase-3 expression, suggesting that Smad4 may regulate compressive force-induced apoptosis in HGFs. In conclusion, these findings increased understanding regarding the mechanisms of compressive force-induced HGF proliferation and apoptosis, which may provide further insight for improving the efficacy and stability of orthodontic treatment.

Introduction

In orthodontic treatment, force is applied to the teeth using appliances and transmitted to the periodontal tissues, which eventually leads to tissue reconstruction of the gingiva, periodontal ligaments and alveolar bone, thereby moving the misaligned teeth to their normal positions (1). The adaptive reconstruction of periodontal tissue is the biological basis of tooth movement (2). As an important part of the periodontal tissue, gingiva is often involved in the process of orthodontic treatment, such as gingiva proliferation, recession and retraction. Such malformations persist for a number of years after treatment (3). Gingival reconstruction not only affects the rate of tooth movement and the stability after correction (4,5), but also leads to recurrence after orthodontic treatment (6).

Orthodontic pressure is converted into biological signals, which are transmitted from the extracellular to the intracellular environment, inducing a series of associated signal transduction reactions that affect the process of gingival tissue remodeling (7). Human gingival fibroblasts (HGFs) are the main cells in the gingival tissue, where they receive and transmit signals (8). In previous decades, although a number of groups have widely studied the reconstruction of alveolar bone and periodontal membrane under compressive force (9), mechanisms underlying gingival reconstruction remain largely unexplored. Our previous study showed that compressive force could promote the expression of transforming growth factor (TGF)-β1 and the proliferation of HGFs (10). TGF-β is an important growth factor that regulates cell proliferation and apoptosis (11,12); however, the factors and signaling pathways involving TGF-β that regulate HGF proliferation and apoptosis remain unknown.

Smad4, a key downstream factor of TGF-β signaling pathway, plays a crucial regulatory role (13). In the classical TGF-β pathway, activated Smad2/3 and Smad4 form the SMAD complex, which translocates to the nucleus (14). Smad4 is considered to be a tumor suppressor gene that is closely related to cell apoptosis (15-17). Imbalance of Smad4 expression in the TGF-β signaling pathway may lead to various diseases, including cancers, fibrous lesions, metabolic diseases and cardiovascular diseases (18-21). It has been found that Smad4 affects apoptosis by regulating apoptosis-related proteins; for example, the low expression of Smad4 in colon cancer can inhibit cell apoptosis by promoting Bcl-2 and Bcl-w expression, and decreasing caspase-3 and caspase-9 expression (22). Therefore, it is hypothesized that Smad4 may play a pivotal role in the proliferation and apoptosis of HGFs under compressive force.

Biodegradable poly(lactide-co-glycolide) (PLGA) has become one of the most widely used biological scaffold
materials (23-26). Compared with other scaffold materials, PLGA possesses good biological properties (27-29). Compared with a two-dimensional culture, a three-dimensional (3D) culture can more effectively mimic the natural environment within the human body, and improve the accuracy and reliability of experimental results. Therefore, a 3D HGF-PLGA culture model was established to explore the roles of Smad4, caspase-3 and Bcl-2 in regulating proliferation and apoptosis pathways in HGFs. The present study aimed to further understand the molecular mechanisms underlying orthodontic gingival reconstruction.

Materials and methods

3D culture of HGFs. HGFs were cultured in 3D PLGA as previously reported (10). Normal gingival tissues were collected from healthy teeth that received orthodontic extraction or gingival resection. Patients enrolled in the study had no systemic diseases, and were aged 10-14 years, with 17 males and 19 females. Normal gingival tissues were cut into small pieces (1x3 mm) and spread flat in culture flasks. Dulbecco's modified Eagle's medium (HyClone; Cytiva) with 20% fetal bovine serum (FBS; Gibco; Thermo Fisher Scientific, Inc.) and 100 U/ml antibiotics (Gibco; Thermo Fisher Scientific, Inc.) were put into the flasks for primary culture. After the primary cells had proliferated, they were digested with 0.25% trypsin (Sigma-Aldrich; Merck KGaA) containing 100 ng/ml DNase (Roche Diagnostics) for 1 min. The digestion was stopped by adding Dulbecco's modified Eagle's medium with 10% FBS and 100 U/ml antibiotics. Cells were cultured at 37°C with 5% CO₂. PLGA scaffolds (molar ratio, 75:25; pore diameter, 100-200 µm; porosity, 85%; size, 2x2 cm x 300 µm; Shandong Medical Instrument Research Institute) were cut into 1x1x0.2 cm sizes. All donors and their guardians provided written informed consent, and samples were collected between March 2018 and March 2019 at the Department of Oral and Maxillofacial Surgery of the Stomatological Hospital affiliated to Guangxi Medical University (Nanning, China).

Application of compressive force. After 4-6 generations, HGFs were digested, and then the cell count was adjusted to 1x10⁵/ml. HGF suspension (0.5-1 ml) was dripped on each piece of PLGA scaffold to enable more cells to attach to the scaffold. The scaffold was stored in an incubator for 2 h at 37°C. After confirming that the cells were stably attached to the PLGA, the PLGA were cultured in 6-well plates with 1 ml containing 10% fetal bovine serum medium for a further 24 h in an incubator with a constant temperature of 37°C. HGFs were digested, and the cell count was adjusted. HGFs were seeded in 96-well plates with 5x10³ cells/well and cultured for 12 h. Then, lentivirus (Lv-control or Lv-Smad4; plasmid backbone, GV492; Shanghai GeneChem Co., Ltd.) and transfection enhancement solution HiTrans A (Shanghai GeneChem Co., Ltd.) was added (MOI=50). After 24 h, aspirate the supernatant in the culture plate and replace with a new serum medium. Additionally, a blank control group constituting normal HGFs was included in the study. The infection efficiency was observed via green fluorescence using a fluorescence microscope (Olympus Corporation) 72 h after infection. Reverse transcription-quantitative (RT-q)PCR and western blotting (WB) were performed to analyze Smad4 expression at the mRNA and protein levels.

RT-qPCR. Total RNA from HGFs was isolated using TRIzol® reagent (Invitrogen; Thermo Fisher Scientific, Inc.). Then, RNA quality was tested using an RNA 6000 Nano kit (Agilent Technologies, Inc.). cDNA was synthesized using an ExScript RT reagent kit (Takara Bio, Inc.), according to the manufacturer's protocol (37°C for 15 min and 85°C for 5 sec, followed by storage at 4°C until further use. The RT-qPCR was performed on an ABI 7300 Real-Time PCR system (Applied Biosystems; Thermo Fisher Scientific, Inc.) with SYBR Premix Ex Taq (Takara Biotechnology Co., Ltd.) at 95°C for 30 sec, followed by 40 cycles at 95°C for 5 sec and 60°C for 31 sec, after which a melt curve analysis was performed at 95°C for 15 sec, 60°C for 1 min and 95°C for 15 sec. The primer sequences were: Smad4 forward, 5'-TGG TAT GAT TAT GAGATTACAG-3' and reverse, 5'-ACCACTCTAACAACACTAATTAGTC-3'; caspase-3 forward, 5'-CCTAAGCGGAGTGCTGTA-3' and reverse, 5'-CATGGGTTTGTCAGCTCACG-3'; Bcl-2 forward, 5'-AATATCCATCCCTGCTGCTA-3' and reverse, 5'-GGTTCCAGGTTTCTCT-3'; and GAPDH forward, 5'-TGTTGCTGCTGCACTTC-3' and reverse, 5'-TTGCTGTGAGTGCAGGAG-3'. All reactions were performed in triplicate and GAPDH was used as an internal control. Each experiment was repeated at least three times. The relative expression level was calculated using the 2-ΔΔCt method (30).

WB. Total protein was extracted from cells using RIPA lysis buffer with PMSF (1:100; Pierce; Thermo Fisher Scientific, Inc.). The protein concentration was detected using a BCA assay. Proteins (30 µg/sample) and marker (5 µl) were loaded onto 10% SDS-PAGE gels. Electrophoresis was conducted under a constant voltage of 200 V for 45 min. Proteins were transferred to PVDF membranes at 200 mA for 1.5 h. TBS-0.3% Tween-20 was used to wash the PVDF membranes 3 times (5 min/wash). Membranes were then blocked in 5% skimmed milk for 1 h at room temperature before being incubated with monoclonal antibodies against human Smad4 (1:1,000; cat. no. ab40759; Abcam), Bcl-2 (1:1,000; cat. no. 4223S; Cell Signaling Technology, Inc.), caspase-3 (1:5,000; cat. no. ab32351; Abcam) and GAPDH (1:5,000; cat. no. SAB4300645; Sigma-Aldrich; Merck KGaA) for overnight incubation at 4°C, followed by incubation for 1 h at room temperature with secondary anti-mouse/rabbit horseradish peroxidase-conjugated antibodies (1:5,000;
After several washes, protein bands were visualized using an Alpha Chemical Luminescent Gel Imaging System (Fluor Chem HD2; ProteinSimple) with ECL luminescence reagent (cat. no. abs920; Absin Bioscience, Inc.). The relative density of protein expression was analyzed quantitatively with Image Lab v3.0 software (Bio-Rad Laboratories, Inc.).

Statistical analysis. SPSS 21 (IBM Corp.) was used for statistical analysis of experimental data. Data are presented as the mean ± SD of at least three independent experiments. Data were analyzed using one-way or two-way ANOVA with post hoc tests (Dunnett’s or Tukey) for multiple comparisons. P<0.05 and α=0.05 were considered to indicate a statistically significant difference.

Results

Compressive force induces the proliferation of HGFs. A 3D PLGA-HGFs culture model was previously established by our group as showed in Fig. 1 (10). Compared with the control group (0 h), the proliferation ability of HGFs significantly increased under compressive force for 12, 24 and 48 h, peaking at 24 h. Subsequently, the proliferation of HGFs was inhibited and cell viability decreased by 72 h (P<0.05; Fig. 2).

Compressive force decreases the expression of Smad4 in HGFs. The gene and protein expression of Smad4 decreased significantly following exposure to compressive force, reaching its lowest level at 24 h. Subsequently, the expression levels increased at 48 and 72 h (P<0.05; Fig. 3).

Expression of caspase-3 and Bcl-2 changed in response to compressive force in HGFs. The gene and protein expression of caspase-3 decreased significantly, reaching its lowest level at 24 h. Subsequently, its expression was increased at 48 and 72 h. However, the gene and protein expression of Bcl-2 increased significantly compared with the control group, peaking at 24 h. Subsequently, the expression levels gradually decreased (P<0.05; Fig. 4).

Assessment of Smad4 mRNA and protein levels. At 72 h after transfection of HGFs with Lv-Smad4, the expression efficiency of enhanced green fluorescent protein was ~90% (Fig. 5). Furthermore, gene and protein expression analysis revealed that the levels of Smad4 (Fig. 6) were significantly upregulated in the Lv-Smad4 group compared with the Lv-control and blank control groups (P<0.01). The expression levels in the Lv-control group significantly decreased compared with the control cells (P<0.05), which indicated that the lentiviral system effectively delivered the exogenous Smad4 gene.

Smad4 overexpression inhibits the effects of compressive force on HGF proliferation. The CCK-8 assay results showed that the absorbance values of HGFs increased with the increased culture time (P<0.05; Fig. 7A). In the first two days, the difference in optical density (OD) values between the Lv-Smad4, Lv-control and control groups were not statistically significant. However, the OD value of the Lv-Smad4 group decreased compared with the Lv-control and control groups from day 3 onwards (P<0.05). Additionally, all three groups were significantly different to each other from day 5 onwards, with the highest OD values in the control group and the lowest in the Lv-Smad4 group (P<0.05). Furthermore, the CCK-8 results revealed the participation of Smad4 in HGF proliferation induced by compressive force, as shown in Fig. 7B. There was no significant increase in HGF proliferation in the Lv-Smad4 group after 24 h of compressive force compared with before force was applied (P>0.05; Fig. 7B). Conversely, in the Lv-control and Lv-Smad4 groups, stimulation with continuous compressive force for 24 h significantly increased HGF proliferation (P<0.05; Fig. 7B).

Smad4 overexpression reverses the expression of caspase-3 and Bcl-2 in HGFs. To determine whether Smad4 mediates caspase-3 and Bcl-2 expression under compressive force, Smad4 was overexpressed in HGFs. Although continuous compressive force stimulation for 24 h significantly decreased the expression of Smad4 in the Lv-Smad4 group, Smad4 expression was upregulated compared with in the Lv-control and control groups. Furthermore, it was demonstrated that with or without compressive force stimulation, the expression levels of Bcl-2 and caspase-3 were downregulated and upregulated, respectively, in the Lv-Smad4 group compared with the control group (Fig. 8). These findings indicated that Smad4 is
an important apoptosis-associated factor in HGFs; compressive force inhibited HGF apoptosis and promoted proliferation by downregulating the expression of Smad4.

**Discussion**

Clinically, gingival hyperplasia and accumulation of the stressed side are commonly found during orthodontic treatment (1-4). Exploring this phenomenon, Schwarz (31) reported that when the orthodontic force did not exceed the end capillary pressure of the periodontal ligament on the root surface of the tooth, it was most conducive to the movement of teeth, and exceeding this threshold results in the necrosis of periodontal tissue. In the human body and most mammals, this pressure is ~20-26 g/cm². Therefore, it has been proposed that under reasonable pressure, weight loading models can more effectively simulate changes in the periodontal tissue on the compressed side of the body (32,33). For example, Kook et al (34) and Li et al (35) used force-loading models to demonstrate that the proliferative ability of periodontal ligament fibroblasts is enhanced under suitable compressive pressure stimulation. In addition, Baker et al (36) reported that the elastic modulus of porous PLGA scaffolds is ~4 MPa, which is close to the elastic modulus of human gingival tissue (37). Therefore, the present study was conducted based on these findings and early experimental results of our research group, such as a PLGA-HGFs 3D culture model, an optimal value of 25 g/cm² and high expression of TGF-β1 (10).

Studies have found that Smad4 is closely related to cell proliferation and apoptosis (15-17). It has been found in tumor research that Smad4 is a tumor suppressor gene, and its decreased or absent expression in a variety of tumors has
Figure 5. Transfection efficiency of Lv‑Smad4 and Lv‑control. (A) EGFP fluorescence and (B) appearance under ordinary optical microscope of HGFs infected with Lv‑Smad4. (C) EGFP fluorescence and (D) appearance under ordinary optical microscope of HGFs infected with Lv‑control. Magnification, x40. EGFP, enhanced green fluorescent protein; HGF, human gingival fibroblast; Lv‑, lentivirus.

Figure 6. Smad4 gene and protein expression in human gingival fibroblasts following transfection with Lv‑Smad4. Relative expression of Smad4 (A) mRNA and (B) protein following transfection with Lv‑Smad4, Lv‑control or blank control treatment. One‑way ANOVA followed by Dunnett’s post hoc test was used to analyze data. *P<0.05 vs. Control. Lv‑, lentivirus.

Figure 7. Smad4 overexpression inhibits the effects of compressive force on the proliferation of HGFs. (A) Proliferation of HGFs following lentivirus transfection. Two‑way ANOVA with Tukey’s post hoc test was used to analyze data. *P<0.05 vs. day 1; †P<0.05 vs. Lv‑Smad4; ‡P<0.05 vs. Lv‑control. (B) Proliferation of HGFs under compressive force. Two‑way ANOVA with Tukey’s post hoc test was used to analyze data. *P<0.05 vs. Control; †P<0.05 vs. 0 h. HGF, human gingival fibroblast; Lv‑, lentivirus; OD, optical density.
been found to lead to the abnormal proliferation of tissues and cells (19,20); however, it has been found in some studies of cardiovascular diseases that TGF-β1/Smad4 can inhibit cell apoptosis. For example, Yin et al (38) silenced Smad4 in human aortic vascular smooth muscle cells (HASMCs) and found that low expression of Smad4 decreased the anti-angiotensin stimulation ability of HASMCs, leading to HASMC migration and apoptosis. Therefore, studies have suggested that the effect of TGF-β1/Smad4 signaling on cell apoptosis is dependent on the cell type and environment.

In the present study, HGF proliferation and expression of Smad4 were time-dependent. The proliferation ability of HGF enhanced and peaked after 24 h of compressive force, then began to decrease with longer exposure. Conversely, the expression of Smad4 decreased and reached a nadir at 24 h, then began to increase. These results suggested that HGFs underwent a series of proliferative and apoptotic processes under compressive force that involved Smad4.

To further explore the relationship between Smad4 and cell proliferation, Smad4 was overexpressed in HGFs. The results of CCK-8 assays revealed that proliferation in the Lv-Smad4 group was significantly decreased compared with in the Lv-control and the control groups from day 3, indicating that Smad4 inhibited the proliferation of HGFs compared with normal cells, which may be due to increased apoptosis. Additionally, the proliferation of HGFs under compressive force was significantly inhibited in the Lv-Smad4 group. These results suggested that Smad4 is associated with the proliferation and apoptosis of HGFs under compressive force, thus highlighting Smad4 as a pro-apoptotic factor in HGFs.

Apoptosis is a gene-regulated programmed cell death process; it is a physiological response of cells to adapt to the environment and maintain the stability of the internal environment of the body (39). There are two main pathways of apoptosis: The mitochondrial pathway and the death receptor pathway. Caspase and Bcl-2 family proteins are involved in both apoptosis pathways (40,41). Caspase-3 plays a key role in promoting apoptosis and is considered the executor of apoptosis (42). The Bcl-2 protein family is a large complex family, mostly located in the mitochondria, endoplasmic reticulum and continuous perinuclear membrane, that mediates the mitochondrial apoptosis pathway (43). Bcl-2 can inhibit apoptosis and is one of the most important oncogenes in apoptotic research (44). It has been reported that down-regulated expression of Smad4 in breast cancer can inhibit cell apoptosis by decreasing caspase-3 and caspase-9, and promoting Bcl-2 expression (45). Overexpression of Smad4 in human vascular endothelial cells has shown that Smad4 can inhibit endothelial cell proliferation by promoting caspase-3 activation and blocking the cell cycle (46). However, the role of compressive force and Smad4 in the apoptosis of HGFs, and the relationship between Smad4, caspase-3 and Bcl-2, remain undetermined.

In the present study, the expression of Bcl-2 and caspase-3 were also highly time-dependent. The expression of Bcl-2 was upregulated under compressive force and peaked at 24 h, at which point it began to decrease. Conversely, the expression of caspase-3 decreased, reaching its lowest point after 24 h of compressive force stimulation, at which point it began to increase. These results suggested that Bcl-2 and caspase-3 were
in involved in apoptotic processes in HGFs under compressive force.

It was further shown that, compared with normal cells, caspase-3 expression was increased in the Lv-Smad4 group, while Bcl-2 expression was downregulated. This suggested that Smad4 regulated the expression of caspase-3 and Bcl-2 in HGFs without compressive force. Furthermore, it was found that Lv-Smad4 cells exhibited downregulated expression of Bcl-2 and upregulated expression of caspase-3 compared with Lv-control and control group cells under compressive force, further suggesting that compressive force regulated the expression of caspase-3 and Bcl-2 via Smad4, thereby regulating HGF proliferation and apoptosis.

In conclusion, the present findings indicated that Smad4 promoted HGF proliferation by regulating caspase-3 and Bcl-2 under compressive force. Therefore, Smad4 may be an important target for preventing gingival hyperplasia after orthodontic treatment. However, there were certain limitations to the present study. For example, proliferation and apoptosis were only analyzed using CCK-8 assays; additional methods should be used, such as flow cytometry. In addition, the discussion on Smad4 was limited to enhancing the expression of Smad4; subsequent studies should further validate these findings by silencing Smad4.

Acknowledgements
Not applicable.

Funding
The present study was supported by grants from the National Natural Science Foundation of China (grant no. 81070860) and the Natural Science Foundation of Guangxi Province (grant no. 2013GXNSFDA019183).

Availability of data and materials
All data generated or analyzed during this study are included in this published article.

Authors' contributions
SZ conceived and coordinated the study, designed, performed and analyzed the experiments, and wrote the paper. LN, YW, LW and SM collected and analyzed data, and revised the manuscript. All authors approved the final version of the manuscript, LN and SM confirm the authenticity of all the raw data.

Ethics approval and consent to participate
The present study was approved by the Hospital Institutional Review Board (approval no. 20150304-22) of Guangxi Medical University. All donors and their guardians signed an informed consent form.

Patient consent for publication
Not applicable.

Competing interests
The authors declare that they have no competing interests.

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