Experimental research on the effect of microRNA-21 inhibitor on a rat model of intervertebral disc degeneration

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Abstract. Intervertebral disc degeneration is associated with angiogenesis and is the primary cause of disc-associated disease. Several studies have indicated the importance of microRNA (miR)-21 in angiogenesis. Thus, the present study aimed to validate the role and underlying mechanisms of miR-21 in a rat model of intervertebral disc degeneration. A total of 60 specific-pathogen-free Sprague-Dawley rats were used for in vivo experiments. A rat model of intervertebral disc degeneration was established and miR-21 inhibitor (antagomiR-21) was administered. The vertebral pulp and annulus fibrosus were isolated for immunohistochemical analysis of hypoxia inducible factor (HIF)-1α and vascular endothelial growth factor (VEGF) expression. Lumbar spine proteoglycan content was detected with the phloroglucinol method. Disc cell apoptosis was detected with terminal deoxynucleotidyl-transferase-mediated dUTP nick end labeling staining. It was revealed that antagomiR-21 treatment decreased the expression of HIF-1α and VEGF in the vertebral pulp and annulus fibrosus. Furthermore, antagomiR-21 treatment increased proteoglycan content and inhibited cell apoptosis in lumbar spines from model rats with intervertebral disc degeneration. In conclusion, antagomiR-21 treatment exerted a protective role in a rat model of intervertebral disc degeneration, which may provide the basis for a potential therapeutic approach in the treatment of disc-associated diseases.

Introduction

Intervertebral disc degeneration is a major pathological process that occurs in the lower back and is the main cause of disc-associated diseases, including disc herniation and spinal stenosis (1-4). Previous studies have indicated that the underlying cause of disc degeneration is tissue weakening, which occurs primarily due to genetic inheritance, aging, inadequate nutritional status and loading history (5). Several other factors may also influence the aging and degeneration of discs, including metabolite transport impairment, cell senescence and death, genetic inheritance, changes in matrix macromolecules and water content, alterations in enzyme activity, structural failure and neurovascular ingrowth (5).

Several studies have reported the important role of angiogenesis in degeneration of the intervertebral disc (6,7). Degenerative intervertebral disc disorders are thought to be characterized by angiogenesis and the increased expression of vascular endothelial growth factor (VEGF), an angiogenic factor (8). Wang et al (9) also concluded that degeneration of the intervertebral disc was accompanied by angiogenesis. David et al (10) demonstrated that angiogenesis influences the pain intensity of intervertebral disc hernias and negatively impacts postoperative pain improvement, mobility and overall quality of life.

MicroRNAs (miRNAs/miRs) are small non-coding RNAs composed of 20-22 nucleotides, which inhibit protein expression by binding to the 3′-untranslated region of target mRNAs, leading to transcriptional repression or degradation of the mRNA (11). MiR-21 is an oncogenic miRNA that is overexpressed in several human tumors and has the ability to modulate cancer-associated target gene expression (12). Notably, it has been reported that miR-21 overexpression impairs angiogenesis in normal epithelial cells (13). Zhao et al (12) demonstrated that arsenite-induced carcinogenesis involves angiogenesis mediated by miR-21. Liu et al (14) revealed that miR-21 overexpression induced tumor angiogenesis and increased the expression of hypoxia inducible factor (HIF). HIF-1α has an important role in angiogenesis and can stimulate the expression of VEGF (15); HIF-1α has also been reported to play an important role in the development of degenerative processes in the intervertebral discs of mice (16).

The aforementioned studies indicate the importance of miR-21 in angiogenesis. The present study aimed to investigate if miR-21 had a critical role in the progression of intervertebral disc degeneration through HIF-1α and VEGF expression regulation. A rat model of intervertebral disc degeneration was established and the model rats were administered miR-21 inhibitor (antagomiR-21). The vertebral pulp and annulus fibrosus were isolated for immunohistochemical analysis to
examine the effects of miR-21 on HIF-1α and VEGF expression. The proteoglycan content in the lumbar spines and disc cell apoptosis was also detected.

Materials and methods

Animals. A total of 60 1-year-old specific pathogen free female Sprague-Dawley rats (150-200 g) were used in strict accordance with the guidelines for the Care and Use of Laboratory Animals (17). The present study was approved by the Animal Ethics Association of The Affiliated Second Hospital of Soochow University (Suzhou, China). The rats were purchased from the Animal Laboratory of the Academy of Medical Sciences (Beijing, China). They were kept in separate cages with free access to food and water, and a 12/12 h light/dark cycle (temperature, 25±1˚C; humidity, 50%).

Experimental groups. Rats were randomly divided into four groups of 15. The normal control group received a skin incision, which were subsequently sutured. The rats in the three other groups underwent a previously described surgical procedure to induce the development of lumbar intervertebral disc degeneration (18). Briefly, the rats were anesthetized by an intraperitoneal injection of 350 mg/kg of 6.5% chloral hydrate. The sacrospinal muscles, spinous processes, supraspinous ligaments, interspinous ligaments and posterolateral halves of the bilateral zygapophysial joints of the lumbar spine were removed.

Subsequently, rats in the model (control) group received a tail vein injection of normal saline; rats in the scramble group received a tail vein injection of 80 mg/kg/day of control oligonucleotides (Guangzhou RiboBio Co., Ltd., Guangzhou, China); rats in the antagomiR-21 group received a tail vein injection of 80 mg/kg/day of antagomiR-21 (Guangzhou RiboBio Co., Ltd.). Injections were administered for 8 weeks. Following this, rats were euthanized by an intraperitoneal overdose of pentobarbital sodium. Lumbar spines, including the L4 to L6 discs, were removed en bloc; the paravertebral muscles and the posterior columns were fully removed. The vertebral pulp and annulus fibrosus were isolated for immunohistochemical analysis of HIF-1α and VEGF expression.

Immunohistochemical analysis. The metaphysis of the vertebral pulp and annulus fibrosus specimens were fixed in a 4% paraformaldehyde solution at room temperature for 30 min following three washes with PBS. Next, tissues were dehydrated with a graded series of ethanol, infiltrated with xylene, and then embedded in paraffin before being cut into 6-µm-thick sections. The slides were then deparaffinization and rehydration with a graded ethanol series. Following this, the sections were depleted of endogenous peroxidase activity through the addition of methanolic H₂O₂ for 15 min and blocked with 10% normal goat serum (Abcam, Cambridge, MA, USA) at 37°C for 30 min. The samples were then incubated at 37°C with anti-HIF-1α (cat. no. ab113642; 1:200; Abcam) and anti-VEGFA (cat. no. ab46154; 1:100; Abcam). VEGFA is a common variant of VEGF, generally referred to as VEGF. The samples were subsequently incubated with a biotinylated rabbit secondary antibody (cat. no. BA1100; 1:375; Vector Laboratories, Inc., Burlingame, CA, USA) at 37°C for 1 h. The bound secondary antibody was amplified using the Elite ABC kit (Vector Laboratories, Inc., Burlingame, CA, USA) at 37°C for 30 min. The slides were washed three times with PBS and incubated with diaminobenzidine (DAB) for 15 min at RT. Subsequently, the sections were counterstained with hematoxylin for 1 min at RT and then washed with distilled water. Subsequently, the slides were dehydrated with a graded series of ethanol and cleared with xylene before mounting with a cover slip with permount for light microscopy, allowing for the analysis of the immunohistochemical expression of HIF-1α and VEGFA.
USA). The antibody–biotin–avidin–peroxidase complex was visualized using 0.02% 3,3’-diaminobenzidine. The sections were mounted onto gelatin-coated slides, air-dried overnight at room temperature and the coverslips were mounted using Permount medium (Thermo Fisher Scientific, Inc., Waltham, MA, USA). The slides were viewed using a light microscope (Olympus BH-2; Olympus Corporation, Tokyo, Japan; magnification, x400) and the optical density (OD) was analyzed by Image Pro Plus Version 6.0 image analyzing system (Media Cybernetics, Inc., Rockville, MD, USA).

Detection of proteoglycan content. The proteoglycan content in the lumbar spines was detected using the phloroglucinol method, as previously described (19). Briefly, 0.6 g tissue was ground, 5 ml 3% NaOH was added and the solution was placed in a thermostatic oscillator at 40˚C for 3 h. Trypsin (5 ml) was subsequently added for 2 h at 37˚C. The saccharide standard concentration was formulated according to a Phloroglucinol solution (Shanghai Macklin Biochemical Co., Ltd., Shanghai, China) following the manufacturer’s protocol. Phloroglucinol solution (5 ml) was added and the sample was placed in a water bath at 100˚C for 8 min. The absorbance of the solution was determined at 554 nm with a UV-visible spectrophotometer (WFZ-UV 2800H; Unico, Shanghai, China). Distilled water was used as a blank sample; the standard tube A value was determined and a standard curve was drawn. The proteoglycan content (mg/g) in the lumbar intervertebral disc tissue was calculated relative to the value of A.

Apoptosis detection by terminal deoxynucleotidyl-transferase-mediated dUTP nick end labeling (TUNEL) staining. Lumbar spines were fixed in 4% paraformaldehyde at 4°C for 48 h, decalcified at 4°C in 20% EDTA for 5-7 weeks, embedded in paraffin and cut into 4-µm thick sections along the midsagittal plane. An in situ TUNEL reaction was performed on two serial sections using the MK1020 apoptosis detection kit (Wuhan Boster Biological Technology, Ltd., Wuhan, China), according to the manufacturer’s protocol. Apoptotic cells were imaged under a light microscope (magnification, x400). A total of 10 random fields were selected and the number of TUNEL-positive disc cells were compared with the total number of disc cells and expressed as a percentage.

Statistical analysis. Statistical analysis was performed with one-way analysis of variance followed by Tukey’s test, using
SPSS 11.5 software (SPSS, Inc., Chicago, IL, USA). The data are presented as the mean ± standard deviation from three independent experiments. Experiments were performed in triplicate. P<0.05 was considered to indicate a statistically significant difference.

Results

AntagomiR-21 treatment decreased the expression of HIF-1α in the vertebral pulp and annulus fibrosus. The vertebral pulp and annulus fibrosus from rats in each group were isolated for immunohistochemical analysis of HIF-1α expression. In the model and scramble groups, positive staining for HIF-1α (brown and yellow) was observed in the nucleus and cytoplasm (Fig. 1). The model and scramble groups exhibited significantly increased expression of HIF-1α in the vertebral pulp and annulus fibrosus compared with the control group. Compared with the scramble group, antagomiR-21 treatment significantly decreased HIF-1α expression in the vertebral pulp and annulus fibrosus.

AntagomiR-21 treatment decreased VEGF expression in the vertebral pulp and annulus fibrosus. The vertebral pulp and annulus fibrosus were also isolated for the immunohistochemical analysis of VEGF expression. Positive staining for the expression of VEGF (brown and yellow) was predominantly observed in the cytoplasm. Similar to the expression pattern of HIF-1α, VEGF expression in the model and scramble groups was significantly increased compared with the control group and VEGF expression was significantly decreased in the antagomiR-21 treatment group compared with the scramble group (Fig. 2).

AntagomiR-21 treatment increased the lumbar spine proteoglycan content. The phloroglucinol method was used to evaluate the effect of antagomiR-21 on the lumbar spine proteoglycan content. The results revealed that the proteoglycan content of the lumbar spine was significantly decreased in the model group compared with the control group. AntagomiR-21 treatment significantly increased the proteoglycan content in lumbar spines compared with the scramble group (Fig. 3).

Discussion

In the present study, antagomiR-21 treatment was demonstrated to decrease the expression of HIF-1α and VEGF in the vertebral pulp and annulus fibrosus of a rat model of intervertebral disc degeneration. Nucleus pulposus cell death mediated through apoptosis is involved in extracellular matrix degradation, which is a deleterious consequence of intervertebral disc degeneration (20). Liu et al (21) reported that miR-21 is upregulated in degenerative human nucleus pulposus tissues compared with normal tissues. Furthermore, Liu et al (21) demonstrated that miR-21 administration promotes nucleus pulposus cell proliferation.

To the best of our knowledge, the present study provided novel evidence to demonstrate the role of miR-21 in the regulation of angiogenesis, as evidenced by the decreased expression of HIF-1α and VEGF following antagomiR-21 treatment in a rat model of intervertebral disc degeneration. Consistent with these findings, Liu et al (14) reported that miR-21 overexpression increases HIF-1α and VEGF expression and induces tumor angiogenesis. Zhao et al (22) suggested that inhibition of angiogenesis with antagomiR-21 occurs through the HIF-1α/VEGF/VEGF receptor 2 signaling pathway.
HIF-1 is a key transcription factor expressed in response to hypoxic stress and is closely associated with angiogenesis. HIF-1 is a heterodimeric transcription factor containing HIF-1α and HIF-1β subunits (23). Under hypoxic conditions, HIF-1α accumulates in the cytoplasm and subsequently translocates into the nucleus. HIF-1α and HIF-1β can then dimerize and bind to hypoxia response elements to stimulate the transcription of a large number of genes, including prostaglandin synthase, angiopoietin, protein tyrosine phosphatase, erythropoietin and VEGF (24,25). Zhu et al (26) suggested that mutual promotion of HIF-1α expression occurs during the process of lumbar intervertebral disc degeneration and that the expression of HIF-1α is significantly associated with microvessel density, which provides evidence for the association between HIF-1α expression and angiogenesis in lumbar intervertebral disc degeneration of rats.

The present study also revealed that antagomiR-21 treatment increased proteoglycan content and inhibited cell apoptosis in lumbar spines. A decrease in proteoglycan content is consistently detected with degeneration, particularly in the center of the disc (27). The induction of disc cell apoptosis is closely associated with intervertebral disc degeneration (28). Thus, it was concluded that antagomiR-21 treatment exerted a protective role in the rat model of intervertebral disc degeneration. However, the present study only discussed the effect of antagomiR-21 treatment on expression of HIF-1α and VEGF in the vertebral pulp, annulus fibrosus, lumbar spine proteoglycan content, and lumbar spine cell apoptosis of a rat model of intervertebral disc degeneration. The association of an antagomiR-21-mediated decrease in the expression of HIF-1α and VEGF with proteoglycan content and/or cell apoptosis in lumbar spines was not fully elucidated, which was the limitation of the present study. In addition, the underlying mechanism by which antagomiR-21 treatment decreased expression of HIF-1α and VEGF also requires further investigation.

In conclusion, the present study demonstrated that antagomiR-21 treatment exerted a protective role in the rat model of intervertebral disc degeneration by increasing the proteoglycan content and inhibiting cell apoptosis, at least in part through HIF-1α and VEGF expression regulation. The findings of the current study demonstrate that antagomiR-21 may be a novel approach for the treatment of intervertebral disc degeneration.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors’ contributions
XS and YX conceived and designed the study. XS, QG, and JY performed the experiments. XS and QG wrote the paper. YX reviewed and edited the manuscript. All authors read and approved the manuscript.

Ethics approval and consent to participate

The present study was approved by the Animal Ethics Association of The Affiliated Second Hospital of Soochow University (Suzhou, China).

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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