

## **Short Communication**

# IL-1beta sensitizes rat intervertebral disc cells to Fas ligand mediated apoptosis in vitro

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# **Key words**

intervertebral disc cell; apoptosis; Fas ligand; flow cytometry; interleukin-1beta

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#### **Abstract**

Aim: To determine the apoptotic effect of recombinant rat Fas Ligand on rat intervertebral disc cells pre-treated with IL-1beta in vitro, and the expression of Fas in cultured rat intervertebral disc cells. Methods: Cells were isolated from the inner annulus fibrosus and transition zones of lumbar discs from Sprague-Dawley rats. The cells were grown in monolayer and divided in 5 treatment groups. IL-1beta (10 ng/mL), FasL (5, 20 ng/mL) with/without IL-1beta (10 ng/mL) pre-treatment was respectively added in Dulbecco's modified Eagle's medium and Ham's F-12 medium with 1% fetal bovine serum. After 32 h, the cells were stained with annexin V-FITC and propidium iodide to evaluate apoptosis using flow cytometry and to analysis transcription of Fas using RT-PCR. Results: Compared with control group, FasL (20 ng/mL), IL-1β (10 ng/mL)+FasL (5 ng/mL), and IL-1β (10 ng/mL) +FasL (20 ng/mL) induced significant apoptosis of the disc cells (P<0.01). Apoptosis was also induced by FasL 5 ng/mL (P<0.05); whereas, apoptosis was not induced by IL-1 $\beta$  (10 ng/mL) (P>0.05). IL-1 $\beta$  (10 ng/mL) enhanced the apoptosisinducing effects of FasL (5 ng/mL) and FasL (20 ng/mL) in disc cells. Fas gene transcription in all groups and Fas expression in the 5 treatment groups were approximately 1.2–2.1-fold greater than control group (respectively, P<0.05). Additionally, Fas expression in FasL with IL-1β pre-treatment groups were significantly up-regulated than in FasL groups (P<0.01). Conclusion: The results of this study showed disc cells pre-treated with IL-1beta increased apoptotic rate in response to FasL in vitro and provided insights to understand Fas/FasL systemmediated apoptosis in disc cells which would be enhanced due to inflammation factor in degenerative disc.

# Introduction

Low back pain is a leading cause of morbidity. It is estimated that about 70% of the population will experience low back pain during their lives<sup>[1]</sup>. In recent years, intervertebral disc (IVD) disorders and age-related degeneration have been significant contributors to low back pain and spine-related disability. There is great interest in understanding the complex pathogenesis of IVD diseases and recent reports of the existence of apoptotic cells in IVD have provided a new insight into the pathophysiology of IVD degeneration<sup>[2–8]</sup>.

Fas/FasL system-mediated apoptosis is thought to play an important role in the loss of disc cells that leads to diminished generation, organization, and repair of the extracellular matrix in the herniated lumbar disc tissues. The disc cells, after herniation, undergo apoptosis via autocrine or paracrine FasL mechanisms by the disc cells themselves<sup>[2,5-8]</sup>. Fas (CD95) and Fas ligand (FasL) belong to the TNF family. The binding of FasL with Fas triggers the formation of the death-inducing signaling complex by recruiting an adaptor molecule Fasassociated death domain (FADD) to the cytoplasmic tail of Fas (C-terminal region). The subsequent autocatalytic acti-

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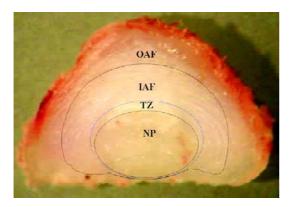
vation of a downstream cascade of caspases leads to the cleavage of specific substrates and thus, the activation of the apoptotic executioners<sup>[9–12]</sup>.

Herniated IVD tissue has been shown to produce proinflammatory cytokines, including matrix metalloproteinases (MMPs), interleukin (IL)-1 $\beta$ , interleukin-1 $\alpha$ , interleukin-6, tumor necrosis factor-alpha (TNF- $\alpha$ ), nitric oxide(NO), and prostaglandin E2 (PGE-2)<sup>[13-18]</sup>.

We proposed that normal disc cells can upregulate apoptosis in an inflammation microenvironment attributed to the production of a large number of inflammatory cytokines in the disc degeneration process. Therefore, the goal of the current study was to investigate the apoptotic effect and the Fas gene expression on cultured rat IVD cells which were stimulated by Fas ligand and IL-1 $\beta$ .

#### Materials and methods

Primary disc cell isolation All cell culture supplies were purchased from Gibco BRL (Gaithersburg, MD, USA) unless otherwise noted. Lumbar IVD (L3 to L6) from Sprague-Dawley rats (aged 3 months, male, 455±29 g in weight) were harvested immediately in a sterile environment after they were killed. The nucleus pulposus (NP) was removed and the inner annulus, including the transition zone (TZ), was separated through an operating microscope. The determination between the outer and inner annulus was based on the amount of hydrated ground substance between the lamellae. The outer annulus is a dense, fibrous tissue, with little space between the oriented lamellar layers. The inner annulus, including the TZ, contains more ground substance which causes the lamellae to distend and become less distinct and organized. For most discs, the outer one-third of the annulus is designated as the outer annulus, and the inner



**Figure 1.** Cross-section of a Sprague–Dawley rat lumbar IVD. The boundary of the outer annulus fibrosus (OAF), IAF, TZ, and NP are shown.

two-thirds are designated as the inner annulus (Figure 1).

The inner annulus fibrosus (IAF), including the TZ between the annulus and the nucleus, was dissected and placed in a humidified incubator with 5% CO<sub>2</sub> at 37 °C in Dulbecco's modified Eagle's medium and Ham's F-12 medium (DMEM/ F-12) with 10% fetal bovine serum (FBS) (Hyclone, Utah, USA), 100 U/mL penicillin, and 100 µg/mL streptomycin for 8 h. To isolate the cells, the disc tissues in the DMEM/ F-12 medium were digested with 0.25% trypsin\_(including 0.02% EDTA) for 40 min followed by another treatment with 0.1% collagenase for 8 h. After enzyme digestion, the suspension was filtered through a 70 µm mesh. The filtered cells were then washed with the DMEM/F-12 medium and a primary cell culture was started. About  $1.5 \times 10^6$ cells were extracted from each rat lumbar disc. The rat disc cells from the IAF including the TZ, were used for this study.

Cell culture in selected concentrations of recombinant rat IL-1β or the recombinant rat Fas ligand. When the primary cell culture became confluent, the cells were trypsinized and subcultured into 6-well plates at 3×10<sup>5</sup> cells/ well. The cells were cultured in DMEM/F-12 medium with 10% FBS, 100 U/mL penicillin, and 100 µg/mL streptomycin in a humidified atmosphere with 5% CO<sub>2</sub> at 37 °C. When the confluence in each well was over 80%, the medium was replaced with medium containing 1% FBS without penicillin and streptomycin. There were 5 treatment groups. In 3 treatment groups, the disc cells were cultured in the medium (1% FBS) for 8 h. The medium was refreshed and 10 ng/mL IL-1 $\beta$ (Cytolab/Peprotech Asia, Rehovot, Israel), 5 ng/mL FasL (R&D Systems, Minneapolis, MN, USA), and 20 ng/mL FasL was respectively added to the medium (1% FBS); the cells were cultured for up to another 24 h. In the other 2 treatment groups (10 ng/mL IL-1β+5 ng/mL FasL, and 10 ng/mL IL- $1\beta+20$  ng/mL FasL), the cells were pretreated with 10 ng/mL IL-1 $\beta$  for 8 h in the medium (1% FBS). The medium was then refreshed and FasL (5, and 20 ng/mL) was respectively added to the medium (1%FBS); the cells cultured for up to another 24 h. Cultures without addition of IL-1β or FasL acted as the controls. After 32 h, the cells were double stained with Annexin V-fluorescein isothiocyanate (Annexin V-FITC) and propidium iodide (PI) (Bender MedSystems, Vienna, Austria), and the cells were harvested for RNA extraction.

**Flow cytometry**(**FCM**) Apoptosis was determined by staining cells with both Annexin V–FITC and PI, according to the manufacturer's instructions. Annexin V–FITC is used to quantitatively determine the percentage of cells undergoing apoptosis. It relies on the property of cells to lose membrane asymmetry in the early phase of apoptosis. In apoptotic

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cells, the membrane phospholipid phosphatidylserine is translocated from the inner leaflet of the plasma membrane to the outer leaflet, thereby exposing phosphatidylserine to the external environment. Cells that were positively stained with Annexin V–FITC and negatively stained for PI were considered apoptosis. Cells that were positively stained for both Annexin V–FITC and PI were considered necrosis [19,20]. To quantitate apoptosis, the cells were washed with cold phosphate-buffered saline solution and then resuspended in binding buffer (10 mmol/L HEPES (N-2-hydroxyethylpiperazine-N,-2-ethanesulphonic acid)/NaOH [pH 7.4], 140 mmol/L NaCl, and 2.5 mmol/L CaCl<sub>2</sub>). The cells were stained with 5  $\mu$ L Annexin V–FITC and 10  $\mu$ L PI and then analyzed with EpicsAltra (Beckman Coulter, CA, USA) FCM.

RT-PCR analysis of Fas gene transcription Total RNA was isolated from the disc cells with TRIzol reagent (Invitrogen, Carlsbad, CA, USA), according to the manufacture's directions. Single-strand cDNA templates were prepared from 2 µg total RNA using oligo (dT)<sub>18</sub> and RevertAid M-MuLV reverse transcriptase (Fermentas, Vilnius, Lithuania). Specific cDNA were then amplified by PCR using the following primers (Sangon, Shanghai, China): Fas sense primer: 5'-GCATCTTTGAGGGTTTGGA-3', antisense primer: 5'-CATTTGGTGTTGCTGGTTC-3' and GAPDH sense primer: 5'-ACCACAGTCCATGCCATCAC-3', antisense primer: 5'-TCCACCACCCTGTTGCTGTA-3'. PCR amplification (PCR System 2700, PE, CA, USA) from cDNA was performed in a final volume of 50 µL containing 15 mmol/L MgCl<sub>2</sub>, 1.25 U Takara Taq, and 0.3 μmol/L specific primers (TaKaRa, Dalian, China). The cycling conditions were: denaturation at 94 °C for 30 s, annealing (Fas 50 °C, GAPDH 58°C) for 30 s, and elongation at 72 °C for 60 s. The optimum cycle number was 30 cycles for Fas and 25 cycles for GAPDH. All PCR products were determined by 2% agarose gel electrophoresis by ethidium bromide staining and visualized by UV transillumination. Gel images were analyzed by densitometry using Scion Image (Scion Corp, Frederick, MD, USA). Fas gene expression data are presented as normalized to GAPDH expression.

**Statistical analysis** All experiments were performed at least 3 times to ensure consistency. SPSS 11.0 software (Chicago, IL, USA) was used for the statistics. Data were compared using unpaired 2-tailed Student's *t*-test analysis, with a *P*-value of 0.05 or less considered significant.

## Results

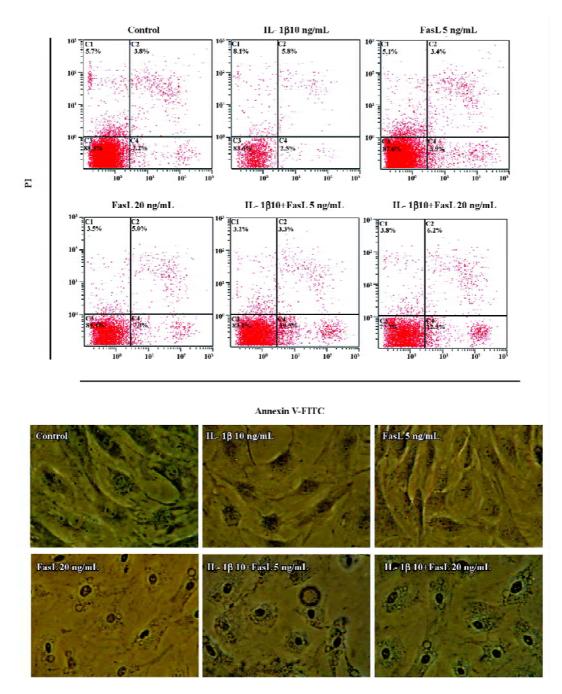
**Establishment of cultures in monolayer** The primary cells from the IAF, including the TZ of rat lumbar IVD, became confluent after 9 d in the monolayer. Then the primary cells were trypsinized and subcultured into 6-well plates with  $3\times10^5$  cells/well. The first passage cells displayed a uniform, rounded, chondrocyte-like morphology and achieved 80% confluence after 7 d.

**Evaluation of apoptosis** When treated and cultured for 32 h, the apoptosis of the 6 groups was determined by double staining with Annexin V–FITC and PI. The apoptotic ratio of rat IAF and TZ cells were calculated as a percentage of apoptotic cells/total cells (Table 1). Compared with the control group, FasL (20 ng/mL), IL-1 $\beta$  (10 ng/mL)+FasL (5 ng/mL), and IL-1 $\beta$  (10 ng/mL)+FasL (20 ng/mL) induced significant apoptosis of the disc cells (P<0.01). Apoptosis was also induced by FasL 5 ng/mL (P<0.05); whereas, apoptosis was not induced by IL-1 $\beta$  (10 ng/mL) (P>0.05). IL-1 $\beta$  (10 ng/mL) enhanced the apoptosis-inducing effects of FasL (5 ng/mL) and FasL (20 ng/mL) in disc cells (Figure 2, P<0.01).

Transcription of Fas When treated and cultured for 32 h, RNA was extracted from the monolayer-cultured rat disc cells. RT-PCR was used for determining the transcription of the Fas gene. The Fas gene was transcripted in the 5 treatment and control groups (Figure 3). And the transcription levels of the Fas gene in the 5 treatment groups were approximately 1.2–2.1-fold greater than the control group (respectively, P<0.05). Additionally, group of IL-1 $\beta$  (10 ng/mL)+FasL (5 ng/mL) compared with group of FasL (5 ng/mL) and group of IL-1 $\beta$  (10 ng/mL)+FasL (20 ng/mL) compared with group of FasL (20 ng/mL) significantly upregulated transcription of Fas (respectively, P<0.01).

Table 1. Effect of IL-1 $\beta$  and FasL on disc cells. Percentage of apoptotic cells was expressed as mean±SD. <sup>1</sup>Groups with IL-1 $\beta$  (10 ng/mL) pretreatment.

Group	Control	IL-1β (ng/mL)	FasL (ng/mL)		FasL <sup>1</sup> (ng/mL)	
		10	5	20	5	20
Apoptosis (%)	2.1±0.2	2.6±0.6	3.6±0.6	7.0±0.9	10.7±1.2	12.3±1.3



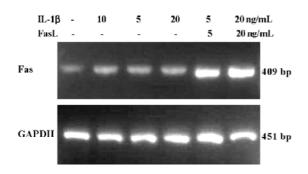
**Figure 2.** Evaluation of apoptosis. After 32 h, cell death was assayed with FCM after double staining with Annexin V–FITC and PI. Lower right quadrant was defined apoptosis which stained positive for Annexin V–FITC and negative for PI. (×300).

# **Discussion**

Fas is a membrane-bound receptor that is activated by the binding of FasL and results in programmed cell death/apoptosis. Park  $et\ al^{[6]}$  reported the effect of Fas on disc cells in human herniated disc tissues and found that the percentage of Fas-positive cells correlated significantly with

patients' age, but not with the degree of disc degeneration on magnetic resonance imaging. Wang  $et\ al^{[8]}$  also found that post-operative samples had an increased number of Faspositive cells in rat cervical degenerative disc models. Anderson  $et\ al^{[21,22]}$  detected a high expression of Fas following both annular laceration in a rabbit model and fibronectin fragment coculturing with rabbit IVD cells  $in\ vitro$ . Fas is

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**Figure 3.** RT-PCR analysis of Fas transcription. After 32 h, mRNA samples were extracted from control group and 5 treatment groups stimulated respectively with IL-1 $\beta$  (10 ng/mL) and FasL (5 and 20 ng/mL) with/without IL-1 $\beta$  (10 ng/mL) pre-treatment. The Fas gene was transcripted in the 5 treatment and control groups. GAPDH served as a control gene.

widely expressed in numerous different cell types throughout the body, whereas Fas ligand expression appears to be more restricted. The expression of Fas ligand in disc cells could be detected in developing embryos<sup>[23]</sup>, degenerative discs<sup>[7]</sup>, normal discs<sup>[24]</sup>, and scoliotic discs<sup>[25]</sup>. Intervertebral discs with their extensive extracellular matrix are largely avascular tissues and display anatomically isolation from the hosts' immune system. Many studies have demonstrated that Fas ligand should play a key role in the potential molecular mechanism to maintain immune privilege of the disc<sup>[7,24]</sup>. However, in degenerative discs<sup>[7]</sup> and scoliotic discs<sup>[25]</sup>, Fas ligand had a close relationship with the apoptosis of disc cells.

The inflammatory cytokine IL-1 plays an important role in disc degeneration. IL-1 has been shown to increase the synthesis of matrix-degrading enzymes (MMP-2, MMP-3, MMP-13, and ADAMTS-4 (A Disintegrin-like and metalloprotease with thrombospondin motifs 4) and to decrease the synthesis of proteoglycan, collagen I and collagen II, and to induce the expression of IL-6, cyclooxy-genase-2, stromolysin-1, and PGE2<sup>[26–30]</sup>. IL-1β can also induce the production of endogenous IL-1β by disc cells in vitro. There is evidence to support that the positive feedback loop of IL-1 $\beta$ exists in degenerative disc cells which upregulate the production of mediators and thus can cause the cessation of symptoms in intervertebral disc herniation<sup>[30]</sup>. Our study demonstrated that FCM found no significant apoptosis after the disc cells were treated with IL-1 $\beta$  (10 ng/mL) for 24 h, however, the apoptotic rate could not deny the changes which occurred to the disc cells. The effect of IL-1β on disc degeneration has been unknown until now.

This study is the first to document normal disc cells *in vitro* response to FasL with/without IL-1 $\beta$  pre-treatment. Disc

cells with IL-1 $\beta$  pre-treatment have a significant apoptotic rate compared with control and disc cells without IL-1 $\beta$  pre-treatment. This implies that the sensitivity of intervertebral discs to FasL increased after IL-1 $\beta$  treatment, which led to a high apoptotic rate at a low level of FasL in normal disc cells.

The present study, using RT-PCR, demonstrates that the transcription of Fas in rat lumbar disc cells increased significantly in the 5 treatment groups. It is important to note that we detected the transcription of Fas on control cells using RT-PCR. Park  $et\ al^{[6]}$  and Wang  $et\ al^{[8]}$  showed a similar result by means of immunohistochemistry, but Anderson  $et\ al^{[22]}$  reported no apparent transcription of Fas on control discs based on the RT-PCR result.

There are 3 limits of the current study. One is that rat disc cells only from the inner annulus fibrosus and TZ were used, because rat NP primary cells could not proliferate and disappeared after 3 weeks. The second limit was that our study stimulated disc cells with IL-1 $\beta$  only for 24 h. It is necessary to prolong the observation time to study whether IL-1 $\beta$  can significantly induce apoptosis of disc cells. The third limit was that we cultured the disc cells in the monolayer, which can not completely represent the cells *in vivo*, so it is necessary to culture disc cells in a 3-D culture system.

In conclusion, the results of this study showed that the apoptotic rate of disc cells pretreated with IL-1 $\beta$  increased in response to FasL *in vitro* and provided insights into understanding the Fas/FasL system-mediated apoptosis in disc cells which would be enhanced due to the inflammation factor in degenerative discs.

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