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**Integrin-FAK 신호전달체계를 조절하여
폐 섬유화를 감소시키는 IL-32 γ 의
기전 연구**

IL-32 γ Attenuates Airway Fibrosis by Modulating
the Integrin-FAK Signaling Pathway in Fibroblasts

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의학과

박소영

Integrin-FAK 신호전달체계를 조절하여
폐 섬유화를 감소시키는 IL-32 γ 의
기전 연구

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이 논문을 의학박사 학위 논문으로 제출함

2018년 12월

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ABSTRACT

Background

Fibrosis in severe asthma often leads to irreversible organ dysfunction. However, the mechanism that regulates fibrosis remains poorly understood. Interleukin (IL)-32 plays a role in several chronic inflammatory diseases, including severe asthma. This study investigated whether IL-32 is involved in fibrosis progression in the lungs.

Methods

Murine models of chronic airway inflammation induced by ovalbumin and *Aspergillus melleus* protease and bleomycin-induced pulmonary fibrosis were employed. The degree of tissue fibrosis after treatment with recombinant IL-32 γ (rIL-32 γ) was evaluated. Expression of fibronectin and α -smooth muscle actin (α -SMA) was examined and the transforming growth factor (TGF)- β -related signaling pathways was evaluated in activated human lung fibroblasts (MRC-5 cells) treated with rIL-32 γ .

Results

rIL-32 γ significantly attenuated collagen deposition and α -SMA production in both mouse models. rIL-32 γ inhibited the production of fibronectin and α -SMA in MRC-5 cells stimulated with TGF- β . Additionally, rIL-32 γ suppressed activation of the integrin-FAK-paxillin signaling axis but had no effect on the Smad and non-Smad signaling pathways. rIL-32 γ localized outside of MRC-5 cells and inhibited the interaction between integrins and the extracellular matrix without directly binding to intracellular FAK and paxillin.

Conclusion

These results demonstrate that IL-32 γ has anti-fibrotic effects and is a novel target for preventing fibrosis.

Key words: Airway Inflammation; Asthma; Interleukin-32 γ ; Pulmonary Fibrosis; Subepithelial Fibrosis.

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INTRODUCTION

Fibrosis, characterized by the accumulation of fibroblasts and excess extracellular matrix, is a common feature of various pathological states in many organs, resulting in dysfunction. Interstitial lung diseases and chronic inflammatory airway diseases of the lungs, such as severe asthma and chronic obstructive pulmonary disease (COPD), lead to sub-bronchial fibrosis and pulmonary fibrosis, both of which result in irreversible structural changes that affect patient survival ¹⁻³). Because lung fibrosis is mainly a consequence of chronic inflammation, therapeutic strategies have focused on preventing inflammation by administering immunosuppressive agents or anti-inflammatory drugs, including corticosteroids ^{4,5}). However, recent studies have suggested that inflammation alone is not sufficient for inducing fibrosis development. Many studies showed that immunosuppressive therapies do not prevent lung fibrosis ⁶). To date, targeting fibrosis itself has been unsuccessful.

Interleukin (IL)-32, initially described as NK4 generated by activated T cells or NK cells ⁷), is produced by various cells, including epithelial cells, endothelial cells, and macrophages. IL-32 induces the production of several pro-inflammatory mediators, such as tumor necrosis factor (TNF)- α , IL-1 β , and IL-6, by activating the nuclear factor- κ B and p38 mitogen-activated protein kinase signaling pathways ^{8,9}). IL-32 is also involved in several chronic inflammatory diseases, such as rheumatoid arthritis and COPD ¹⁰⁻¹²). In addition to its role in inflammation, recent studies suggested that IL-32 is involved in liver fibrosis in patients with chronic hepatitis by affecting cytokine induction ¹³). Although the precise effects of IL-32 on tissue fibrosis are largely unknown, IL-32 contains an RGD motif, which is known to bind several integrins ¹⁴). Moreover, a 3-dimensional reconstruction model of IL-32 revealed that its structure was highly similar to that of the focal adhesion targeting (FAT) region of focal adhesion kinase (FAK). FAK-related non-kinase, a peptide with a structure similar to the FAT

region, inhibits FAK signal transduction ¹⁵). It is known that integrin-FAK signaling axis is critical for the development of tissue fibrosis ^{16,17}). Therefore, it was predicted that IL-32 interrupts the signaling pathway by binding to these molecules, thereby inhibiting FAK activation and alleviating fibrosis.

This study hypothesized that IL-32 γ modulates fibrosis in chronic airway and lung diseases by disrupting the integrin-FAK signaling pathway. Here, murine models were used for chronic airway inflammation and bleomycin-induced pulmonary fibrosis to examine the role of IL-32 γ in fibrosis of the airways and lungs, respectively. In addition, the role of IL-32 γ in mechanisms underlying fibroblast function was evaluated.

MATERIALS AND METHODS

1. Generation of murine models of airway inflammation and pulmonary fibrosis

To generate the bleomycin-induced pulmonary fibrosis model, mice were administered intratracheal injection of bleomycin (1 U/kg body weight) on day 2. To evaluate the effect of IL-32 γ treatment, mice were administered 500 ng of human recombinant IL-32 γ (rIL-32 γ) via intranasal injection on days 1, 2, 14, and 28. In this model, rIL-32 γ was injected intranasally; 1 h later, bleomycin was injected intratracheally. Mice were sacrificed at 30 days. To generate the chronic asthma model, wild-type (WT) mice were sensitized by intranasal administration of 22 μ g of ovalbumin (OVA) and 8 μ g of protease (*Aspergillus melleus* protease; Sigma, St. Louis, MO, USA) twice per week for 8 weeks, as previously described [23]. Mice were sacrificed at 58 days. To evaluate the effect of IL-32 γ treatment, mice were treated with 500 ng human recombinant IL-32 γ (rIL-32 γ) 2 h before each immunization. Bronchoalveolar lavage fluid (BALF) and lung tissues were obtained from mice 24 h after the last immunization for further analysis. IL-32 γ transgenic (TG) mice on a C57BL/6 background were generated as previously described ¹⁸). In brief, the ORF of IL-32 γ cDNA was transferred into pCAGGS. The entire sequence was linearized with *Sall* and microinjected into mouse zygotes. Transgenic mice showed no physical abnormalities and were screened by RT-PCR. Wild-type (WT) C57BL/6 and Balb/c mice were purchased from OrientBio (Gapyong, Gyeonggi-do, Korea) and were used as a control. All mice were bred and maintained in a specific pathogen-free animal facility. The Institutional Animal Care and Use Committee approved all experimental procedures (Animal Utilization Protocol 2014-14-013).

2. Histopathologic examination and quantification of tissue fibrosis

Lungs collected from mice (N=5) in each treatment group were perfused with 5 mL of PBS through the right ventricle. The lung was inflated by intratracheal infusion of 0.3% low-melting agar at 25 cm H₂O. The inflated lung was fixed with 10% neutral buffered formalin. Fixed lungs were embedded in paraffin and sectioned at 4 μm. Sections were deparaffinized, rehydrated, subjected to antigen retrieval by boiling in 10 mM sodium citrate buffer, pH 6.0, for 15 min, and blocked in 3% H₂O₂ (DaKo Peroxidase Blocking Solution) for 10 min at room temperature in a humidity chamber. To examine collagen deposition, lung sections were stained with Masson's trichrome staining. The slides were washed 3 times with PBS, blocked with 0.25% casein in PBS (DaKo Protein Block Serum-Free) for 15 min at room temperature in a humid chamber, and then incubated with antibodies against α-SMA (Cell Signaling Inc., Danvers, MA) overnight in the dark at 4 °C. After washing 3 times with PBS, the slides were incubated with FITC-conjugated goat anti-rabbit IgG, and cy3-donkeyAnti-goat IgG for 1 h at room temperature in the dark, and detected using a DaKo EnVision HRP/DAB system (Dako, Carpinteria, CA, USA). The lung sections used for staining were mounted with mounting media fortified with DAPI (Vector Laboratories Inc, Burlingame, CA). To quantify tissue fibrosis, hydroxyproline levels in the tissue were measured. To measure hydroxyproline in tissue, a hydroxyproline colorimetric assay kit (BioVision) per the manufacturer's protocol was used. Additionally, quantification graphs were drawn from intensity measurement data using the Image J program (NIH, Bethesda, MD, USA).

3. Cell culture and study design

The human lung fibroblast cell line MRC-5 was purchased from the American Type Culture Collection (Manassas, VA, USA). Mouse embryonic fibroblasts (MEFs) obtained from IL-32γ TG mice were also used. MRC-5 cells were seeded at 2×10^5 cells/well and

stimulated with recombinant proteins. These cells were cultured in MEM (Welgene, Seoul, Korea) with 10% FBS (Gibco, Carlsbad, CA) and 1% penicillin-streptomycin (Welgene). Cells were seeded at a density of 2×10^5 cells/well in 60-mm culture dishes and were stimulated with 5 ng/mL of TGF- β (R&D Systems, Minneapolis, MN), 10 ng/mL of TNF- α (R&D Systems), 1 μ g/mL of LPS (Sigma), 10 μ g/mL of Poly I: C (San Diego, CA), 10 ng/mL of IL-1 β (R&D Systems), 50–200 μ g RGD peptide (tripeptide Arg-Gly-Asp, R&D Systems), and 150 ng/mL of rIL-32 (YBDY, Seoul, Korea). To silence intracellular IL-32 γ expression, small interfering siRNA (siRNA) to IL-32 γ (antisense sequence: 5'-UCAUCAGAGAGGA CCUUCGUU-3') was used. Expression of various cellular molecules was measured by Western blotting, reverse transcription-PCR, and semi-quantitative PCR. All in vitro experiments were conducted at least 3 times.

4. Cell adhesion assay

For crystal violet staining, 96-well culture dishes were coated with collagen (Advanced BioMatrix, Inc., San Diego, CA, USA) and seeded with MRC-5 cells. Plates were incubated for 30, 60, or 180 min. Cells were washed with PBS to remove non-adherent cells, and adhered cells were stained with crystal violet. Crystal violet-stained cells were dissolved using 33% acetic acid and OD values were measured at a wavelength of 550 nm. For measuring spindle-shaped cells, cells were observed and counted at 30 min after staining under the microscope (magnification: 100 \times).

5. Western blotting

Cells were lysed on ice in lysis buffer (Cell signaling Inc, Danvers, MA) containing protease inhibitors for 30 min, and then centrifuged at 18,000 \times g for 20 min at 4 $^{\circ}$ C. Proteins

were detected with antibodies against fibronectin, smad3, p-smad3, TGF- β receptor1, JNK, p-JNK, p38, p-p38, Erk, p-Erk, paxillin, p-paxillin (all from Cell Signaling Inc.), smurf2 and FAK (Santa Cruz Biotechnology, Dallas, TX), α -SMA (Abcam, Cambridge, UK), β -actin (Bioworld, St. Louis Park, MN), IL-32 (Ybdy, Seoul, Korea), and p-FAK (Invitrogen, Carlsbad, CA). Anti-rabbit, -goat, and -mouse secondary antibodies were purchased from Bethyl Laboratories (Montgomery, AL). Protein bands were detected with ECL solution (GenedePot, Barker, TX). The quantification graphs are drawn through intensity measurements using the Image J program.

6. RT-PCR and semi-quantitative PCR

Total RNA was extracted from cells using Trizol reagent (Invitrogen, Carlsbad, CA) and purified using RNeasy Mini Kits (Qiagen, Hilden, Germany) according to the manufacturers' instructions. Purified RNA (1 μ g) was reverse-transcribed to cDNA using oligo(dT) primers and reverse transcriptase (Roche Applied Science, Mannheim, Germany). Target amplification was performed with following primers: IL-32 sense, GACAGTGGCGGCTTATTATGAG; IL-32 antisense, CCTCGGCACCGTAATCCA T; TNF- α sense, CGCTCTTCTGCCTGCTGCACTT; TNF- α antisense, AGGCTTGTCACCTCGGGGTTCGA; GAPDH sense, TGCACCACCAACTGCTTA; GAPDH antisense, GGCATGGACTGTGGTCAT; integrin α 2sense, GGAACGGGACTTTTCGCAT; integrin α 2 antisense, GGTACTTCGGCTTTCTCATCA; integrin α v sense, AATCTTCCAATTGAGGATATCAC; integrin α v antisense, AAAACAGCCAGTAGCAACAAT; integrin β 1sense, CGATGCCATCATGCAAGT; integrin β 1 antisense, ACACCAGCAGCCGTGTAAC; integrin β 3 sense, CCGTGACGAGATTGAGTCA; integrin β 3 antisense, AGGATGGACTTTCCACTAGAA; integrin β 5 sense, GGAGCCAGAGTGTGGAAACA; integrin β 5 antisense, GAAACTTTGCAAACCTCCCTC;

integrin β 8 sense, AATTTGGTAGTGGAAGCCTATC; integrin β 8 antisense, GTCACGTTTCTGCATCCTTC.

7. His pull-down assay and immunoprecipitation

His-tagged IL-32 γ (100 μ g) was incubated with Ni-NTA agarose (Qiagen, Valencia, CA) at 4 °C for 2 h. The beads were then washed 3 times with buffer containing imidazole (Sigma). TGF- β -stimulated MRC-5 cells were lysed and centrifuged; The supernatant of TGF- β -stimulated MRC-5 lysate was incubated with IL-32 pre-bound Ni-NTA Agarose at 4 °C for 2 h. Then, proteins associated with the beads were subjected to immunoblot analysis using anti-integrin β 3, anti-paxillin, or anti-His. For immunoprecipitation, flag-tagged IL-32 γ -overexpressing MRC-5 cells were lysed on ice in lysis buffer (20 mM Tris, pH 7.5, 150 mM NaCl, 0.25% NP-40, 1.5 mM MgCl₂, 1 mM phenylmethylsulfonyl fluoride). Anti-flag bound to protein G sepharose (GE Healthcare) was incubated with lysates at 4 for 3 h. Beads were washed 3 times with lysis buffer. Proteins were electrotransferred to PVDF membranes (GE Healthcare) and subjected to immunoblotting.

8. Live cell imaging of IL-32 γ

MRC-5 cells were cultured in a μ -Dish 35 mm, High, IbiTreat (Ibidi GmbH, Martinsried, Germany) and treated with Flamma496-labeled IL-32 γ . After 10 min, the cells were washed with medium. Fluorescence images were obtained under a Nikon Ti-E inverted I wamicroscope (Tokyo, Japan) equipped with PFS, iXon Ultra 897 EMCCD camera (Andor Technology, Belfast, UK), and excitation and emission filter wheels.

9. Statistical analysis

All data are reported as the mean \pm standard error of mean. Means were compared using the Mann–Whitney test in GraphPad Prism software (version 4.0; GraphPad, Inc., La Jolla, CA, USA). A value of $P < 0.05$ was considered statistically significant.

RESULTS

1. IL-32 γ modulates fibrosis in mouse models of airway inflammation and pulmonary fibrosis

First, histopathological analysis of bleomycin-induced lung fibrosis was conducted to determine the effect of IL-32 γ on pulmonary fibrosis. Treatment with rIL-32 γ significantly reduced collagen deposition and α -smooth muscle actin (SMA) expression (Fig. 1A and 1B). Hydroxyproline levels showed a tendency to be lower in the bleomycin-induced fibrosis group treated with rIL-32 γ than in the group without rIL-32 γ treatment (34.01 ± 7.24 vs. 25.52 ± 3.66 , $P = 0.048$; Fig. 1C). Next, to determine the effect of IL-32 γ on airway remodeling in chronic asthma, a murine model of chronic airway inflammation with subepithelial fibrosis was treated with rIL-32 γ . Treatment with rIL-32 γ reduced peribronchial collagen deposition (Fig. 1D). This was accompanied by reduced expression of α -SMA, a marker of activated fibroblasts, around the bronchi of treated mice (Fig. 1E). Figure 1F is a graph showing quantification of hydroxyproline. Hydroxyproline levels were significantly lower in the chronic asthma model treated with rIL-32 γ (32.35 ± 1.752 vs. 24.20 ± 1.344 , $P = 0.010$).

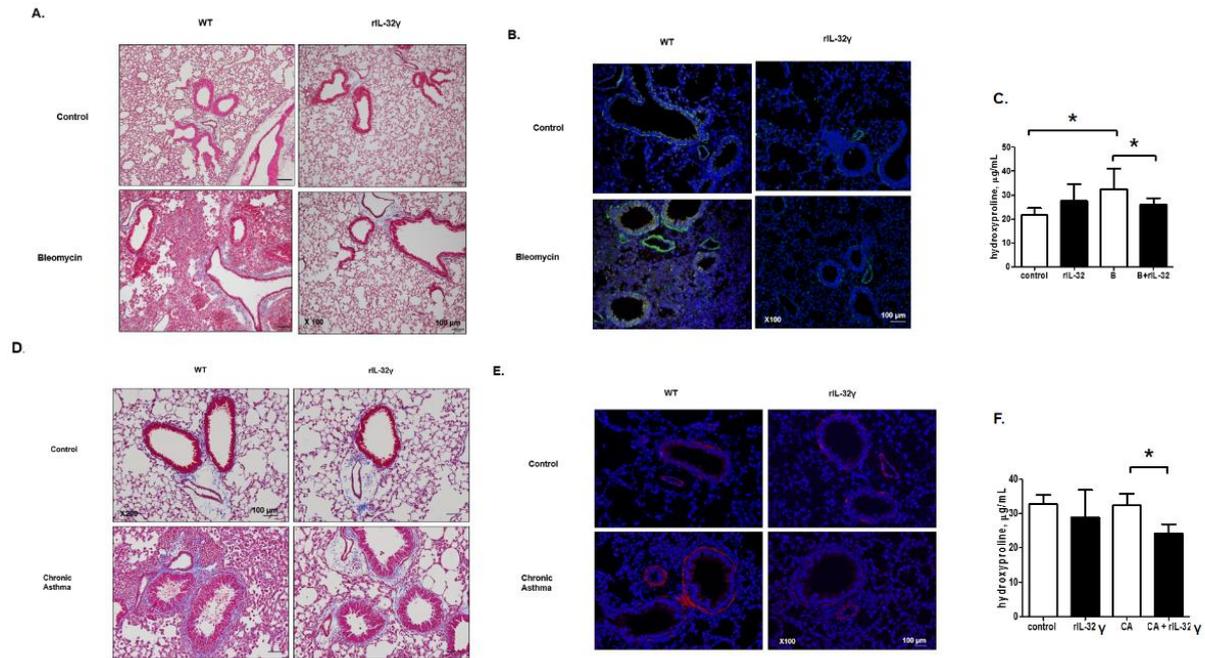


Fig. 1. Human IL-32 γ prevents fibrosis in chronic asthma and bleomycin-induced pulmonary fibrosis models. (A) Evaluation of collagen deposits in the lungs of bleomycin-induced mice using Masson's trichrome stain (original magnification: 100 \times). The quantification graphs of histological analysis in bleomycin-induced fibrosis groups. (B) Immunofluorescence analysis of α -SMA (green) expression in the lungs of bleomycin-induced mice. DAPI staining is blue (original magnification: 100 \times). (C) Hydroxyproline quantification. In the group with bleomycin-induced fibrosis treated with rIL-32 γ (N = 5, B + rIL-32 γ), hydroxyproline levels tended to decrease compared to in the non-rIL-32 γ -treated bleomycin-induced fibrosis model (N = 6, B) (32.40 ± 3.885 vs. 26.70 ± 1.287 , $P = 0.166$). (D) Evaluation of collagen deposition in the lungs of chronic asthmatic mice using Masson's trichrome stain (original magnification: $\times 200$). (E) Immunofluorescence analysis of α -SMA (red) expression in the lungs of mice with chronic asthma. DAPI staining is blue (original magnification: $\times 200$). (F) Hydroxyproline quantification graph. Similar results were obtained in each independent experiment, each using five mice per group (32.35 ± 1.752 vs. 24.20 ± 1.344 , $P = 0.010$). * $P < 0.05$.

2. rIL-32 γ attenuates fibroblast activation

Next, to determine whether IL-32 γ affects fibrosis by regulating fibroblast activation, expression of fibronectin and α -SMA was measured in the human fibroblast cell line MRC-5 after treatment with TGF- β in the presence or absence of rIL-32 γ . Fibronectin expression in rIL-32 γ -treated cells was significantly lower than that in untreated cells, whereas α -SMA expression was slightly lower at early time points (Fig. 2A). However, overexpression of endogenous intracellular IL-32 γ did not noticeably affect the production of fibronectin and α -SMA by MEFs from WT or IL-32 γ TG mice (Fig. 2B). Endogenous IL-32 expression is shown in Fig. 2C.

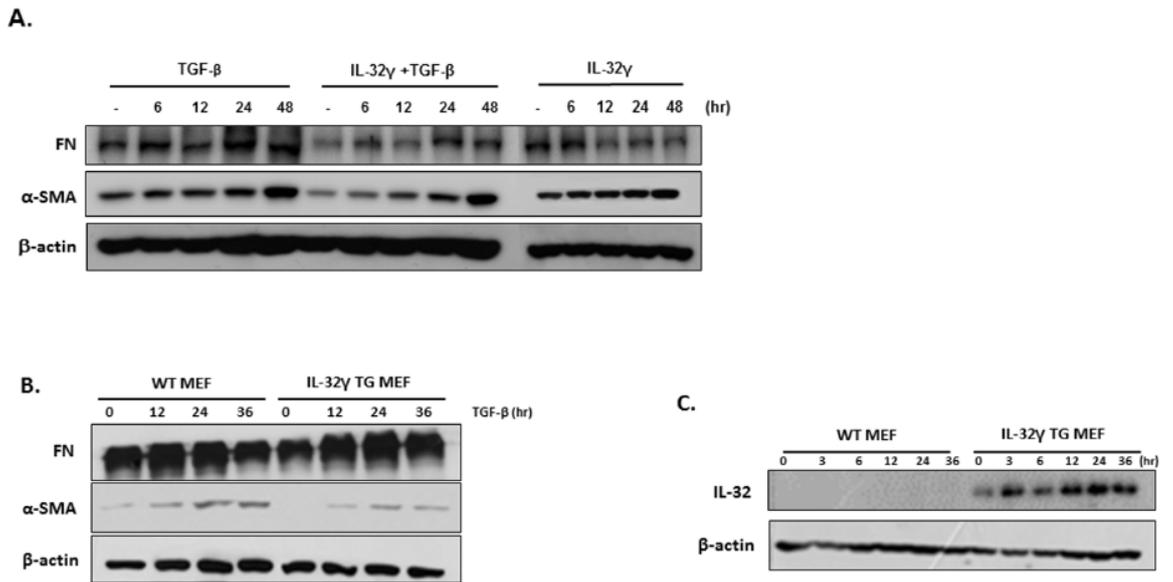


Fig. 2. Exogenous, but not endogenous, IL-32 γ attenuates fibroblast activation. (A) Fibronectin and α -SMA expression was detected in rIL-32 γ (150 ng/mL)-pretreated MRC-5 cells after TGF- β (5 ng/mL) stimulation. (B) Fibronectin and α -SMA expression are shown in IL-32 γ -expressing MEFs after TGF- β (5 ng/mL) stimulation. (C) Endogenous IL-32 expression. Data are representative of three independent experiments.

3. Anti-fibrotic effect of rIL-32 γ occurs independently of TNF- α

Because IL-32 induces the production of TNF- α and vice versa, this study examined whether IL-32 γ exerts anti-fibrotic effects by inducing TNF- α expression. First, it was found that significant expression of IL-32 γ mRNA was induced by TNF- α , although no significant change in TNF- α mRNA expression was observed (Fig. 3A and 3B). Similar to IL-32 γ , treatment with rTNF- α inhibited the expression of fibronectin and α -SMA in TGF- β -stimulated MRC-5 cells (Fig. 3C). However, IL-32 was not expressed by rTNF- α under IL-32 γ -knockdown conditions (Fig. 3D) and an anti-fibrotic effect of TNF- α was not observed in IL-32 γ -knockdown MRC-5 cells (Fig. 3E). Additionally, rIL-32 γ inhibited fibronectin and α -SMA expression after TNF- α inhibitor treatment (Fig. 3F).

Fig 3.

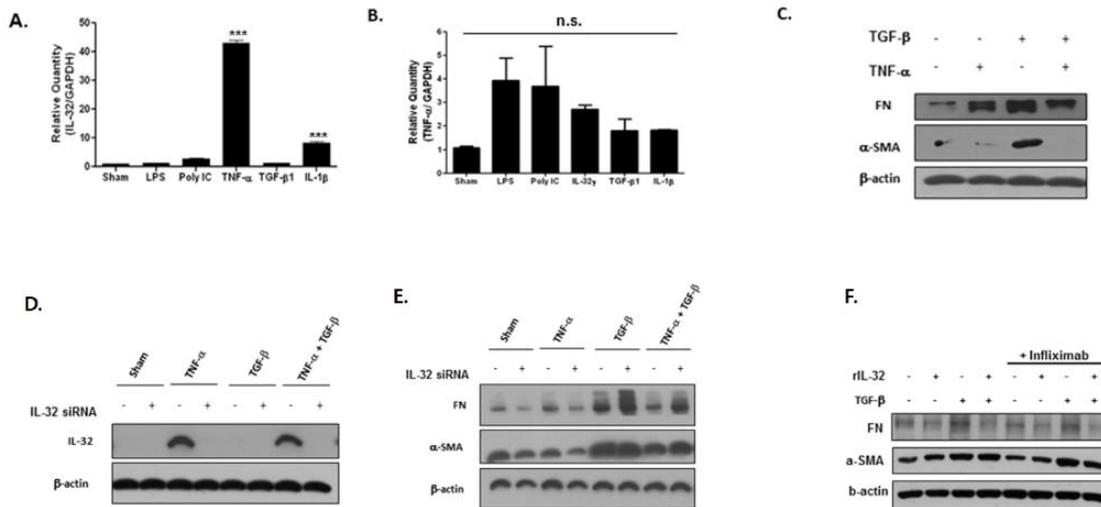


Fig. 3. Anti-fibrotic effects of rIL-32 γ are independent of TNF- α . (A) MRC-5 cells were stimulated with each cytokine including LPS (1 μ g/mL), Poly I: C (10 μ g/mL), TNF- α (10 ng/mL), IL-32 γ (150 ng/mL), TGF- β (5 ng/mL), and IL-1 β (10 ng/mL). After 24-h stimulation, mRNA level of IL-32 γ and TNF- α (B) were measured by quantitative PCR. (C) Fibronectin and α -SMA expression in MRC-5 cells after 24 h of stimulation with TNF- α (10 ng/mL) and TGF- β (5 ng/mL). (D) MRC-5 cells were transfected with IL-32 siRNA and then stimulated with TNF- α (10 ng/mL) or TGF- β (5 ng/mL). (E) Fibronectin and α -SMA expression in cell lysates was detected. (F) Infliximab-pretreated MRC-5 cells were stimulated with IL-32 γ (150 ng/mL) and TGF- β (5 ng/mL), and fibronectin and α -SMA expression in the cell lysate was detected. Results are representative of two independent experiments, each showing similar results.

4. rIL-32 γ does not appear to be involved in TGF- β -mediated Smad or non-Smad signaling

Next, the effect of IL-32 γ was examined on activation of the Smad pathway, a well-known TGF- β -mediated signaling pathway. There were no significant differences in the expression of Smad signaling molecules (p-Smad 3, smuf2, and TGF- β receptor 1), regardless of rIL-32 γ treatment (Fig. 4A). It was also examined whether the non-Smad pathway plays a role in the anti-fibrotic effects of rIL-32 γ . The result showed no significant differences in JNK, Erk, and p38 activation between MRC cells treated with rIL-32 γ and untreated cells (Fig. 4B).

Fig4.

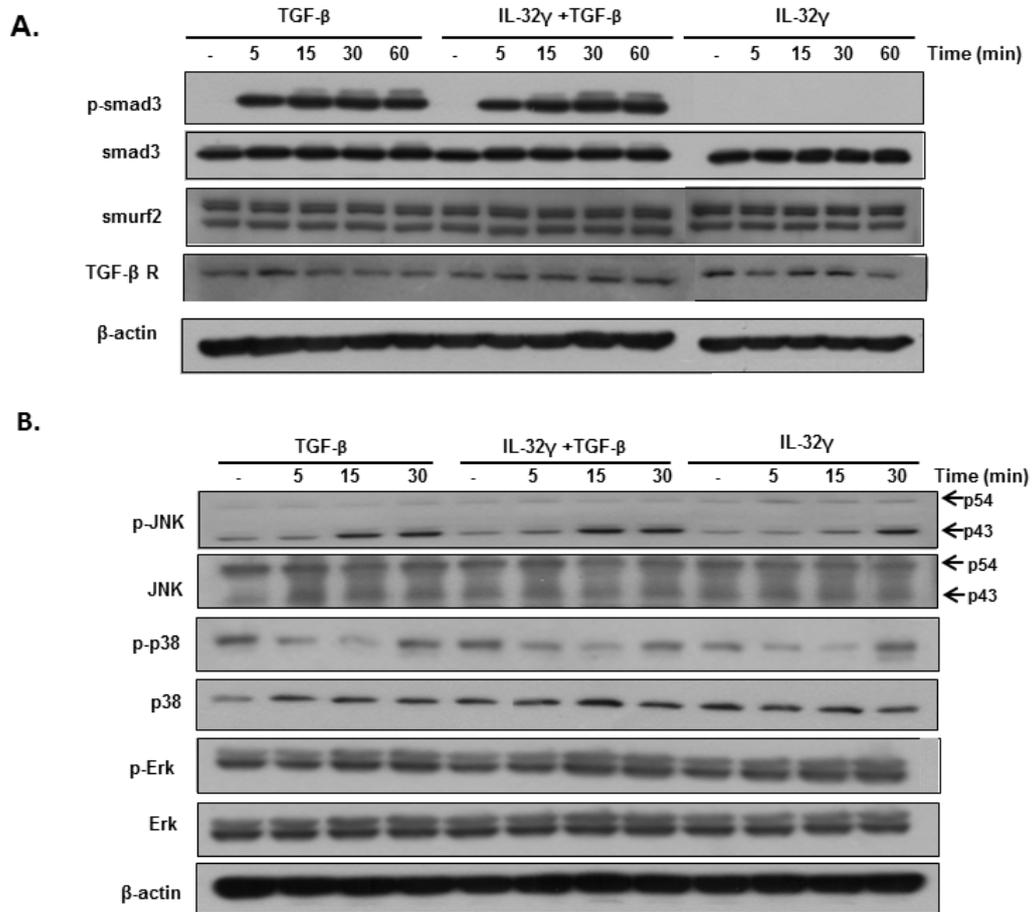


Fig. 4. rIL-32 γ has no effect on TGF- β -mediated Smad or non-Smad signaling pathways. MRC-5 wells were stimulated with TGF- β (5 ng/mL) in the presence or absence of rIL-32 γ and then harvested at the indicated times. Western blot analysis was performed to examine the expression of proteins in the Smad signaling (A) and non-Smad signaling pathways (B). Results are representative of three independent experiments.

5. rIL-32 γ inhibits integrin-mediated FAK/paxillin activation

Next, integrin-dependent activation of FAK and paxillin was evaluated, a critical pathway in fibroblast activation, after treatment with the RGD tripeptide and integrin blocker. RGD peptide inhibited signaling by both FAK and paxillin in MRC-5 cells stimulated with TGF- β (Fig. 5A). Interestingly, rIL-32 γ inhibited FAK and paxillin signaling in a manner similar to that of RGD peptide (Fig. 5B).

To investigate how IL-32 γ regulates the integrin-FAK-paxillin signaling pathway, it was performed a protein-protein binding assay to determine whether IL-32 directly binds to integrin β 3, paxillin, or FAK. Both integrin β 3 and paxillin were detected in the total cell lysate and flow-through lanes, but no bands were detected in the wash and elution fractions (Fig. 5C). This suggests that these proteins do not directly bind to IL-32 γ . Additionally, an anti-flag-IL-32 γ antibody did not immunoprecipitate with FAK (Fig. 5D).

Fig.5

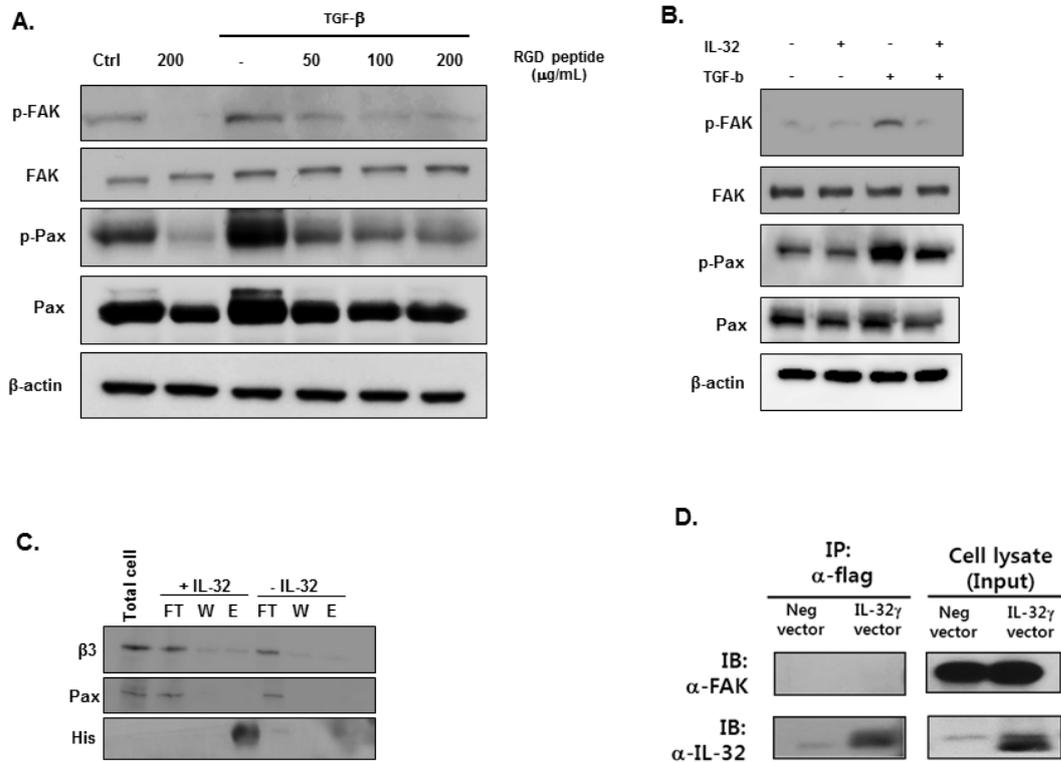


Fig. 5. rIL-32 γ inhibits integrin-mediated activation of FAK/paxillin. Phosphorylation of FAK and paxillin was detected in TGF- β (5 ng/mL)-stimulated MRC-5 cells pretreated with an RGD peptide (A) or rIL-32 γ (B). Activated FAK and paxillin were detected after 24 h. Results are representative of three independent experiments. (C) MRC-5 cells were stimulated with TGF- β for 24 h, and His-tagged rIL-32 γ was precipitated from cell lysates using Ni-NTA beads. Bound proteins were analyzed by Western blotting with antibodies specific for integrin β 3, paxillin, and the His-tag. (D) Flag-tagged IL-32 γ -overexpressing MRC-5 cells were stimulated with TGF- β and harvested at 24 h. Flag-tagged IL-32 γ was then immunoprecipitated from cell lysates using an anti-flag antibody followed by immunoblotting with an anti-FAK antibody. Similar results were obtained from two independent experiments. FT, flow-through; W, wash; E, elution.

6. rIL-32 γ is localized on the cell surface

To determine the mechanism by which rIL-32 γ inhibits activation of the FAK/paxillin pathway, the location of rIL-32 γ was examined by live cell imaging for 60 min. rIL-32 γ was located outside of MRC-5 cells after 60 min, suggesting that it does not enter cells by endocytosis and is not degraded; therefore, IL-32 γ acts extracellularly, at least during the period examined (Fig. 6).

Fig.6

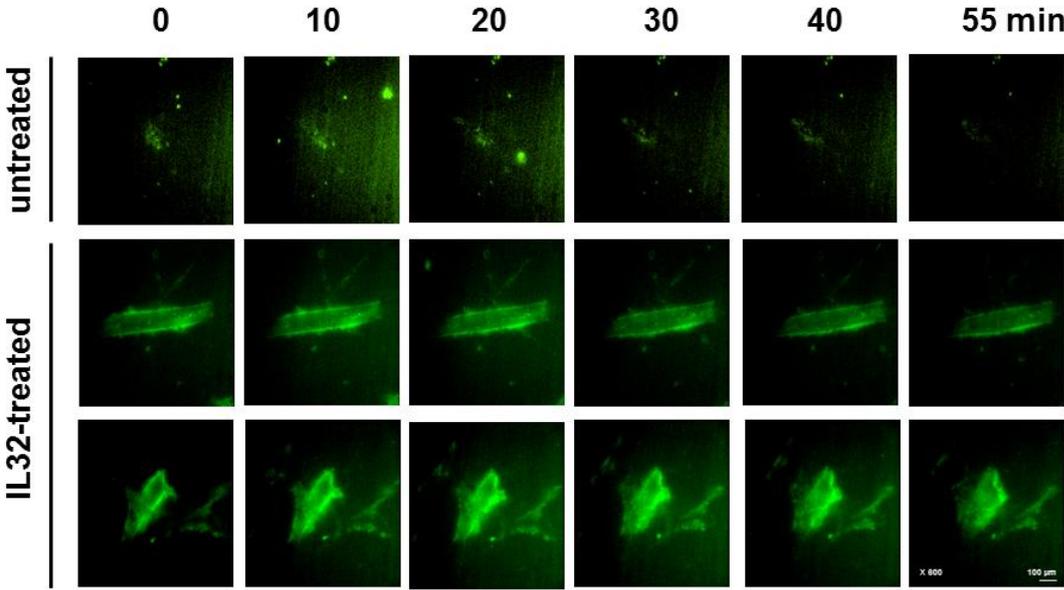


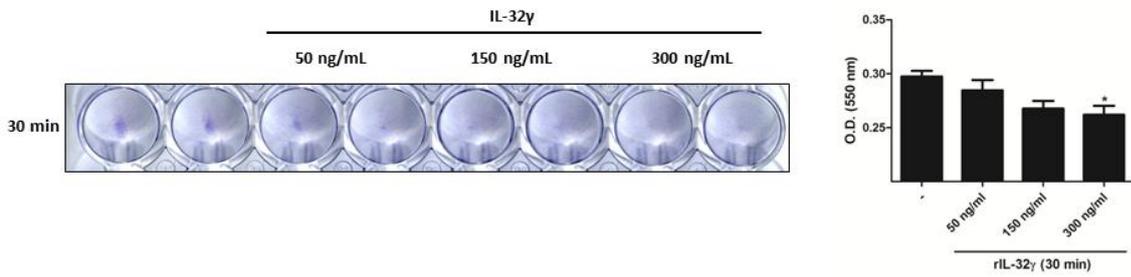
Fig. 6. rIL-32γ localizes extracellularly. Live cell imaging of MRC-5 cells at 10–60 min post-incubation with Flamma496-labeled IL-32γ (magnification, 600×; green color).

7. rIL-32 γ modulates the interaction between integrins and the extracellular matrix

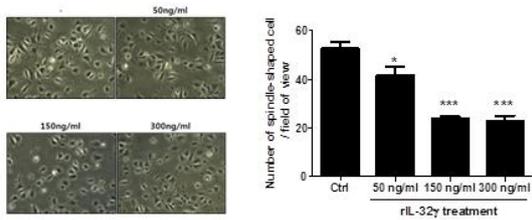
To examine the effect of IL-32 γ on integrin signaling, the adhesion of MRC-5 cells to collagen-coated plates was examined in the presence/absence of rIL-32 γ . MRC-5 cells adhered to collagen within 30 min in the absence of rIL-32 γ ; however, the process was impeded in the presence of rIL-32 γ (Fig. 7A). Moreover, the number of spindle-shaped MRC-5 cells was much lower in the presence of rIL-32 γ , even after 30 min (Fig. 7B). Interestingly, rIL-32 γ suppressed integrin/collagen-mediated activation of FAK and paxillin, which is typically induced by cell adhesion to collagen-coated plates in the absence of any other stimulation (Fig. 7C). Finally, the effect of IL-32 γ on integrin expression in MRC-5 cells was evaluated according to TGF- β upregulates integrin expression. Semi-quantitative PCR revealed increased expression of integrin β 3 and reduced expression of integrin β 8 following TGF- β stimulation. This pattern was not altered by IL-32 γ treatment (Fig. 7D).

Fig 7.

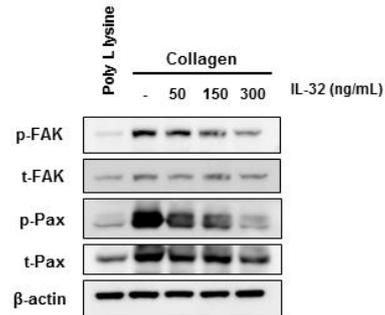
A.



B.



C.



D.

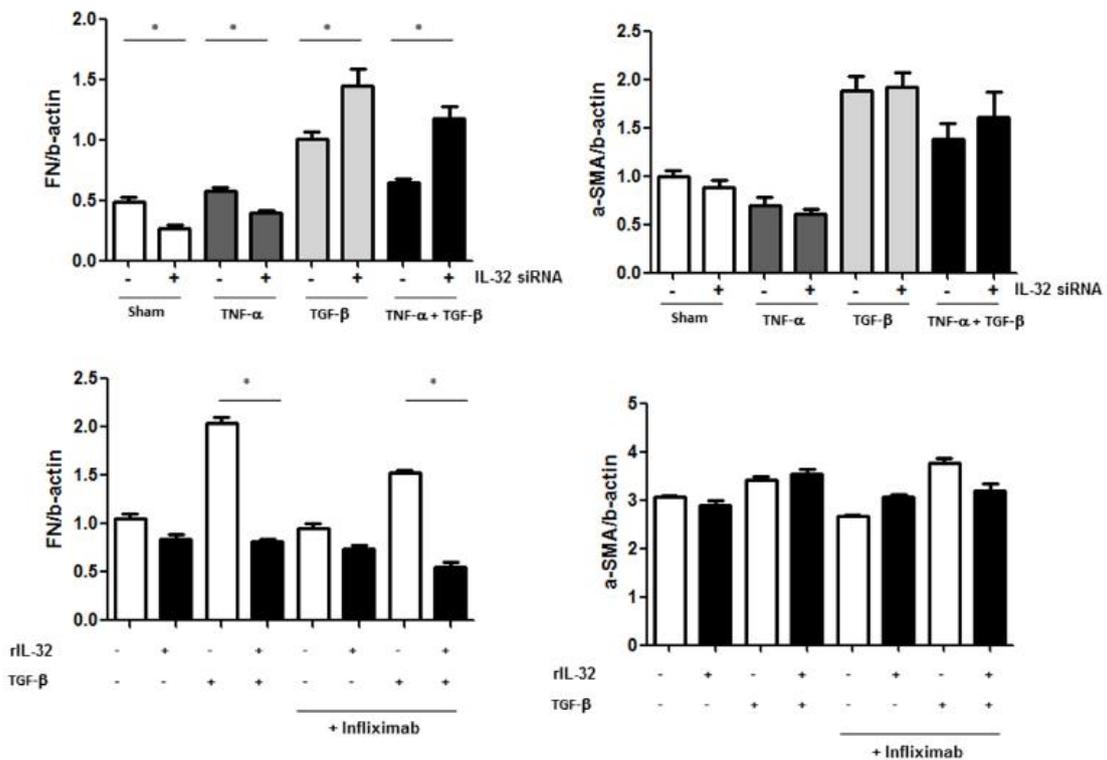


Fig. 7. rIL-32 γ modulates the interaction between integrins and the extracellular matrix. MRC-5 cells were plated on collagen-coated plates in the presence/absence of rIL-32 γ . (A) Adherent MRC-5 cells were stained with crystal violet immediately after the adhesion assay (left) and optical density values from the dissolved crystals are shown (right). Similar results were obtained from three independent experiments. *P < 0.05 (B) Adherent cells were observed at 30 min under a microscope (original magnification: 100 \times). Similar results were obtained from two independent experiments. *P < 0.05, ***P < 0.0001 (C) Phosphorylation of FAK and paxillin was detected after MRC-5 cells attached to collagen-coated plates for 24 h in the presence/absence of rIL-32 γ . Similar results were obtained from two independent experiments. (D) Anti-fibrotic effect of rIL-32 γ is independent of TNF- α . Anti-fibrotic effect of TNF- α was not observed in IL-32 γ -knockdown MRC-5 cells. rIL-32 γ suppressed the expression of fibronectin and α -SMA after TNF- α inhibitor treatment.

DISCUSSION

This study demonstrated the anti-fibrotic effect of IL-32 γ both in vitro and in vivo. It showed that rIL-32 γ regulates fibroblast activation by modulating the integrin-FAK signaling pathway. Thus, rIL-32 γ may be useful for inhibiting tissue fibrosis in the clinical setting.

The mechanism of tissue fibrosis is closely related to that of wound repair, which is a normal healing process in injured tissues. However, dysregulated fibrosis can lead to severe organ dysfunction, which is typically irreversible and has a fatal outcome in many disease states. In the lungs, for example, progressive parenchymal fibrosis is a consequence of serious pulmonary fibrotic diseases such as idiopathic pulmonary fibrosis, leading to high mortality. Additionally, bronchial subepithelial fibrosis can cause irreversible fixed airway obstruction, as observed in chronic inflammatory airway diseases such as chronic severe asthma and COPD, which can become critical if untreated.

Although lung fibrogenesis is thought to result from chronic inflammation, numerous studies have suggested that fibrosis is not completely dependent on inflammatory processes and that anti-inflammatory therapeutic strategies are not always effective. Thus, therapeutic trials have shifted their focus from anti-inflammatory targets to anti-fibrotic targets, as many studies demonstrated that such mechanisms underlie the development of fibrosis¹⁹⁻²²). However, therapeutic agents that effectively control fibrosis are lacking; therefore, there is an urgent need to identify novel molecules with potent anti-fibrotic activities.

IL-32, previously considered a pro-inflammatory cytokine, is a multifunctional protein with a potential role in lung diseases^{12,23-25}). The previous study showed that IL-32 γ modulates immune responses by recruiting IL-10-producing monocytic cells in a chronic asthma model [24]. Here, it observed that IL-32 γ also exhibits a strong anti-fibrotic effect in a model of sub-bronchial fibrosis. Because chronic inflammation is a major factor driving the progression of fibrosis, its apparent suppressive effect on airway fibrosis may be completely

dependent on the anti-inflammatory effects of IL-32 γ . Thus, the modulatory effects of IL-32 γ examined in a bleomycin-induced lung injury model, which is considered a prototype of tissue fibrosis but displays a lower accumulation of immune cells in the lungs. This is of interest because IL-32 γ is a putative immunomodulatory cytokine. The results of the current study suggest that IL-32 γ has a novel function in lung fibrosis, as well as anti-inflammatory effects on chronic airway inflammation.

The human fibroblasts used to further investigate the mechanism underlying the anti-fibrotic effect of IL-32 γ , as excessive accumulation of extracellular matrix produced by activated fibroblasts is a major pathological feature in tissue fibrosis, and any possible effects of inflammation in an animal can be excluded. MRC-5 cells were stimulated with TGF- β , which induces fibroblasts to differentiate into fibronectin- and α -SMA-expressing myofibroblasts. It found that IL-32 γ effectively inhibited expression of these activation markers upon TGF- β stimulation. Previous studies showed that TNF- α and IL-32 γ induce one another. Additionally, TNF- α inhibits the TGF- β -induced Smad signaling pathway²⁶⁻²⁹). Thus, the cells were used in which IL-32 γ had been silenced and a TNF- α -blocking agent to determine the exact mechanism underlying the suppressive effect of IL-32 γ on fibroblast activation. Furthermore, the intracellular pathways linked to the Smad and non-Smad signaling pathways were assessed. The mechanism shows underlying the role of IL-32 γ in fibrogenesis was not dependent on TNF- α expression, nor was it associated with activation of TGF- β downstream of the Smad or non-Smad signaling pathways.

Previous studies indicated that TGF- β -induced fibroblast activation depends on the integrin signaling pathway through FAK/paxillin activation^{16,30-32}). Protein structure modeling suggested that IL-32 γ is involved in integrin activation and downstream signaling pathways^{14,33}). In fact, IL-32 γ contains an RGD motif that binds to integrins; indeed, several isoforms of IL-32 bind to integrin α V β 3. In addition, IL-32 has a structure resembling the FAT region of

FAK (similar to an FAK-inhibitory peptide). However, these studies examined only IL-32 α and β , although IL-32 γ is considered the most active form ³⁴⁾.

rIL-32 γ inhibited the phosphorylation of FAK and paxillin in TGF- β -stimulated fibroblasts without directly binding to these molecules. Based on these results, extracellular rIL-32 γ regulates TGF- β -mediated fibroblast activation without entering the cell. Indeed, rIL-32 γ treatment inhibited integrin-mediated cell adhesion, although rIL-32 γ remained outside the cell. These results strongly suggest that IL-32 γ is involved in the development of tissue fibrosis, likely by disrupting the binding between integrins expressed in the cellular membrane and the extracellular matrix.

No study has fully identified an IL-32-associated pathway in the context of fibrosis, raising the question of whether IL-32 is released by dead cells or via a specific secretory pathway. Notably, in the early phase of several diseases, IL-32 is produced by activated T cells, monocytes, and NK cells and acts as a pro-inflammatory cytokine that stimulates TNF- α , IL-6, and IL-8 production ^{8,12,35,36)}. Because recent studies showed that IL-32 is not secreted ^{24,37)}, IL-32 γ released from injured epithelial cells in patients with chronic inflammatory diseases, including those with mycobacterium avium complex pulmonary disease and idiopathic inflammatory bowel disease ²²⁾, may play a regulatory role in inflammation or tissue remodeling. For instance, our previous study showed that rIL-32 γ suppresses chronic airway inflammation, which is closely associated with airway remodeling ²⁴⁾.

There were some limitations to the current study. First, to obtain more convincing and direct evidence to evaluate our hypothesis, mutations or deletions of the RGD motif of IL-32 γ should be used. Second, our results do not clearly define the precise function of intracellular and extracellular IL-32 γ . Further studies are necessary to resolve these questions.

In summary, IL-32 γ has anti-fibrotic effects likely by blocking the integrin-FAK-paxillin

pathway. Therefore, administration of rIL-32 γ may play a pivotal role in modulating both inflammation and fibrosis in patients in which inflammation-related fibrosis pathways are activated.

CONCLUSION

The present study suggested that IL-32 γ prevents tissue damage by regulating fibroblast activation in the chronic stage. The mechanism underlying this modulatory effect may involve disruption of integrin/FAK signaling cascades, without the need for IL-32 γ to directly bind molecules involved in these cascades. Thus, IL-32 γ is a new candidate for the treatment of lung fibrosis.

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국문초록

연구배경

중증 천식에서 기도의 섬유화는 비가역적으로 현재까지 적절한 치료가 없는 것이 사실이다. 그 이유는 이러한 기도 및 폐의 섬유화의 기전이 명확하게 밝혀지지 않았기 때문이다. IL-32는 중증 천식을 비롯한 다양한 만성 염증성 질환에서 주요한 역할을 하는 것으로 알려져 있다. 이 중 IL-32 γ 는 구조적으로 FAK (focal adhesion kinase)의 FAT 부분과 매우 유사하고, IL-32 γ 는 integrin과 결합하는 것으로 잘 알려져 있는 RGD motif를 가지고 있다. 그러므로 이 연구에서 IL-32 γ 가 기도 및 폐의 섬유화에 어떠한 기전으로 작용하는지 연구하였다.

연구방법

bleomycin 유발 폐 섬유화 모델과, ovalbumin 및 Aspergillus melleus protease 유발 만성 천식 모델을 구축하여 연구를 진행하였고, 재조합 rIL-32 γ 를 처리하여 섬유화 진행 정도를 비교하였다. rIL-32 γ 를 처리한 활성화 된 섬유아세포 (MRC-5)에서 fibronectin과 α -smooth muscle actin (α -SMA) 발현을 분석하고 TGF- β 신호전달 체계를 통한 폐 섬유화 정도를 분석하였다.

연구결과

rIL-32 γ 는 폐 섬유화 및 만성 천식 동물모델에서 조직학적으로 collagen 침착 및 α -SMA 발현을 유의하게 억제하였다. rIL-32 γ 를 처리한 경우 TGF- β 로 자극한 MRC-5세포에서도 fibronectin과 α -SMA 발현이 억제되었다. rIL-32 γ 는 integrin과 FAK-paxillin 신호 전달 축의 활성화를 억제하였지만, Smad 및 non-Smad 신호전달 경로에는 영향을 미치지 않았다. rIL-32 γ 는 MRC-5 세포의 바깥에서 작용하였으며, 세포 내부의 FAK와 paxillin에 직접적으로 결합하지 않고 외부에 국한되어 integrin과 ECM 간의 상호작용을 억제하는 것으로 확인되었다.

결론

이러한 결과는 IL-32 γ 가 세포 밖에서 작용하여 integrin과 ECM 간의 상호작용을 억제함으로써 섬유화 진행을 방해하는 것을 증명하며, 섬유화를 예방할 수 있는 새로운 치료 후보물질로의 개발 가능성을 증명하는 결과이다.

중심단어: 중증 천식, 기도 염증, 폐 섬유화, 인터류킨 32-감마