

The Effect of Interferon- γ and Tumor Necrosis Factor- α on COL1A2 Promoter Activity Elevated by Interleukin-4 in Human Skin Fibroblasts Cultures

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Background: IL-4 is known as a potent stimulating factor and, TNF- α and IFN- γ suppress collagen synthesis of dermal fibroblasts. However, relatively little is known about interaction of these cytokines.

Objective : The purpose of this study was to evaluate the interactional effect of TNF- α and IFN- γ for the type I collagen gene expression and promoter activation produced by IL-4.

Methods : Using dermal fibroblasts from the normal skin cultured with three cytokines, IL-4, TNF- α and IFN- γ , we examined Northern blot analysis with each cDNA and assayed CAT activity with human pro α 2(I)collagen(COL1A2)/CAT reporter gene chimeric construct.

Results : Compared to the control, treating with IL-4 resulted in prominent elevation of both type I procollagen mRNA levels and pro α 2(I) collagen promoter activity. IL-4 with TNF- α suppressed the IL-4 induced elevations, whereas IL-4 with IFN- γ did not reveal the obvious suppression of the elevations produced by IL-4. Fibroblasts treated with IL-4 together with IFN- γ and TNF- α completely abolished the type I procollagen gene expression and the activation of COL1A2 promoter gene elicited by IL-4.

Conclusion : IL-4 induced enhancement of type I collagen gene expression was synergistically suppressed by TNF- α and IFN- γ on transcriptional level. (Ann Dermatol 9:(2) 108~115, 1997).

Key Words : IL-4, TNF- α , IFN- γ , Type I collagen, Promoter activity

Fibroblasts are responsible for the production and maintenance of the connective tissue matrix of all organs including the dermis of the skin. Cultured fibroblasts derived from the early phases of traumatic wound tissue or from sites of pathological fibrosis, display activated phenotypes characterized by increased production of the connective tissue matrix

components: collagen and fibronectin and retain their activated phenotypes for many generations *in vitro*¹⁻³. Following tissue injury, some cytokines make the fibroblasts undergo a metabolic activation and exhibit increased growth and synthetic rates, or inhibit fibroblast growth and collagen production required to effect fibrotic repair of injured tissue.

A number of cytokines have been shown to modulate the growth and migration of connective tissue derived cells and the synthesis of extracellular matrix components, such as collagens⁴. Interleukin-4(IL-4) is a potent stimulating factor of collagen synthesis by fibroblasts especially in the initiation of the fibrotic process⁵. In contrast, tumor necrosis factor- α (TNF- α) and interferon- γ (IFN- γ) suppress collagen synthesis⁶⁻⁸. Short-term

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exposure of cultured sclerodermal fibroblasts to IFNs results in suppression of their abnormally elevated collagen production and in induction of a reduced/normalized-collagen-producing phenotype which is retained in the absence of IFN's⁷⁻⁹. When interferons bind to specific cell surface receptors, there are changes in steady-state levels of multiple mRNAs¹⁰. The role of TNF- α , IFN- γ and IL-4 in the pathogenesis of the fibrotic condition and cultured fibroblast has been elucidated¹¹⁻¹², but relatively little is known about modulation of the collagen gene expression and interaction of these cytokines. A synergistic action of IFN- α with TNF- γ has been shown for tumoricidal¹³ and antiviral activity¹⁴. Several studies have provided evidence that activation of type I collagen gene expression by fibroblasts plays a crucial role in the development of dermal fibrosis in diseases, such as scleroderma, eosinophilic fasciitis, and keloids^{1,15-16}.

To investigate the mechanism by which cytokines affect remodeling processes, the potential interaction of IL-4 with IFN- γ and/or TNF- α on modulating type I collagen gene expression was studied using Northern blot analysis, and type I collagen promoter gene activation was studied also by transfection experiments and chloramphenicol acetyltransferase (CAT) assay in cultured human skin fibroblasts. Human pro α 2(I)collagen/CAT reporter gene chimeric construct used in this study can mimic physiological expression of the gene when transiently expressed in human fibroblastic cells¹⁷.

MATERIALS AND METHODS

Fibroblasts culture and treatment of cytokines

For the collection of fibroblasts, skin specimens were taken from the clinically normal skin of identical anatomical sites on three normal subjects of unrelated cosmetic surgery. Normal human skin fibroblasts were cultured to confluency and maintained in Dulbecco's modified Eagle's medium (DMEM, Gibco BRL Co., Grand Island, NY, USA) with 10% fetal bovine serum (FBS), streptomycin (100 μ g/ml) and amphotericin B (1 μ g/ml) in the humid atmosphere of 37°C and 5% CO₂, and subcultured after trypsinization. Cultured fibroblasts were utilized in passages 3-6. The trypan blue test was used for the checking of cell viability. Confluent fibroblasts in 100 mm diameter

petri-dishes were treated with cytokines for 48 hr in the absence of FBS. The following cytokines and concentrations were used for this study: 10 ng/ml of IFN- γ (Genzyme Co., Cambridge, MA, USA), 1 μ g/ml of TNF- α (Sigma Co., St. Louis, MO, USA), 5 ng/ml of IL-4 (Genzyme Co., Cambridge, MA, USA). Most experiments were performed in triplicate and repeated twice.

cDNA probe preparation

The following human-sequence-specific cDNAs were used for Northern hybridizations: for type I collagen, a 1.8-kb pro α 1(I) cDNA; for fibronectin, a 1.3-kb cDNA; for glyceraldehyde-3-phosphate dehydrogenase (GAPDH), 1.2-kb cDNA. The cDNAs were labelled with [³²P]-dCTP (NEG 036H, New England Nuclear, Boston, MA, USA) by nick translation²² to a specific activity of approximately 1×10^8 counts/min/ μ g.

RNA isolation and Northern blot hybridization

Total RNA was isolated from cultured normal human skin fibroblasts by the methods of Chomczynski and Sacchi¹⁹. The RNA was subjected to Northern blot hybridization analysis as described. Briefly, total RNA was fractionated by 1% agarose gel electrophoresis (50 volt, 5 hr) after denaturing the samples with formaldehyde and formamide²⁰. RNA transcripts obtained were transferred to the nitrocellulose filter (Trans-Blot, BioRad, Hercules, CA, USA) in 20 x SSC at 4°C for 12 hr. After drying the filter at room temperature, RNA was immobilized by heating at 78°C for 90 min under vacuum. The filter was prehybridized for 12-18 hours at 42°C with prehybridization mixture (50% formamide, 0.1% sodium dodecylsulfate (SDS), 3x SSC, 1x Denhart's solution, 50 μ g/ml salmon sperm DNA (ss-DNA)) and hybridized with [³²P]-labelled cDNAs by nick translation at 42°C for 24 to 36 hr. After hybridization, washing and autoradiography were performed.

Transfection experiments and CAT assay

The construct containing the 3.5-kb COL1A2 promoter fused to the CAT gene (pMS-3.5/CAT) was derived from a 3.5-kb EcoRI/SphI genomic subclone that spans from position -3,500 to +58 of the COL1A2 promoter²¹. Cells were trypsinized for 24 hr prior to transfection and seeded at a density

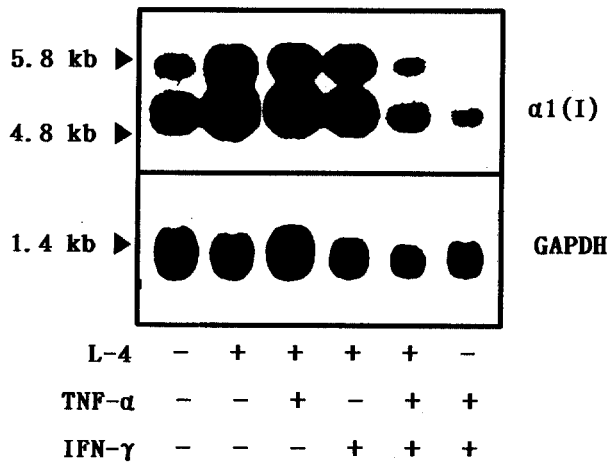


Fig. 1. Northern blot analysis of type I collagen mRNA transcripts from human skin fibroblasts cultures incubated for 48 hr in medium supplemented with IL-4(5 ng/ml), TNF- (10 ng/ml) and/or IFN- γ (1 μ g/ml). Abbreviations: IL-4, interleukin-4; TNF- α , tumor necrosis factor- α ; IFN- γ , interferon- γ ; α 1(I), pro α 1(I) collagen; GAPDH, glyceraldehyde-3-phosphate dehydrogenase.

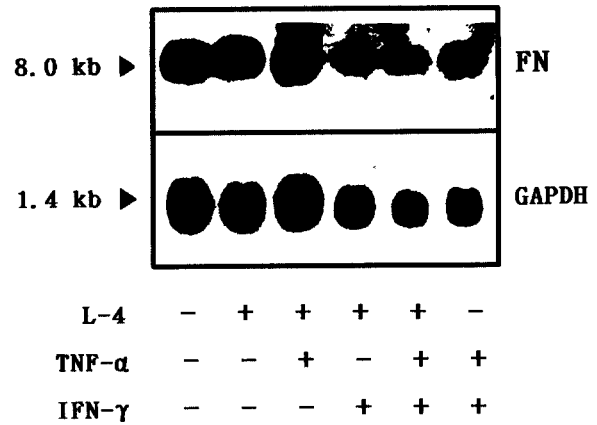


Fig. 2. Northern blot analysis of fibronectin(FN) mRNA transcripts from human skin fibroblasts cultures incubated for 48 hr in medium supplemented with IL-4(5 ng/ml), TNF- (10 ng/ml) and/or IFN- γ (1 μ g/ml). Abbreviations were described in fig. 1.

Table 1. Type I collagen and fibronectin(FN) mRNA abundance in human skin fibroblasts treated with IL-4 alone or in combination with TNF- α and/or IFN- γ

	Type I collagen/GAPDH	FN / GAPDH
Control	24.0 \pm 1.8 (1.00)	0.92 \pm 0.11 (1.00)
IL-4	67.8 \pm 2.1 (2.81)	0.97 \pm 0.09 (1.05)
IL-4 + TNF - α	35.8 \pm 3.9 (1.48)	0.71 \pm 0.11 (0.77)
IL-4 + IFN - γ	54.5 \pm 4.8 (2.26)	0.86 \pm 0.10 (0.94)
IL-4 + TNF - α + IFN- γ	20.3 \pm 1.9 (0.85)	0.75 \pm 0.15 (0.82)
IFN- γ + TNF - α	17.5 \pm 2.2 (0.73)	0.68 \pm 0.13 (0.74)

The values are mean SD from six separate experiments. Abbreviations were described in fig. 1. Values in the parentheses indicate fold induction.

of $1-3 \times 10^5$ cells/60 mm dish for the calcium chloride technique. The next day, cells were transfected with 20 μ g of reporter and 3.5 μ g of β -galactosidase DNA by the calcium-chloride / DNA co-precipitation method, followed by a 1 min of glycerol(15%) shock²². After 48 hr incubation with or without 10 ng/ml of IFN- γ , 1 μ g/ml of TNF- α and 5 ng/ml of IL-4, the cells were harvested and lysed by three cycles of freeze-thawing in 100 μ l of 0.25 M Tris-HCl, pH 7.8. The protein concentration of each extract was used for determination of CAT activity using acetyl CoA and [¹⁴C] chloramphenicol as substrate²³. The acetylated and non acetylated forms of radioactive chloramphenicol were separated by thin layer chromatography(TLC) and vi-

sualized by autoradiography. The enzyme activity was quantified by cutting out pieces of TLC plates containing different forms of [¹⁴C] chloramphenicol converted to its acetylated forms. Each piece of data was corrected by β -galactosidase activity. The values were expressed as the percentage of the [¹⁴C] chloramphenicol converted to its acetylated forms per 10 μ g of cell extraction protein.

RESULTS

Northern blot analysis of pro α 1(I) collagen and fibronectin mRNA

For Northern blot analysis of cultured normal skin fibroblasts with or without cytokines, ³²P la-

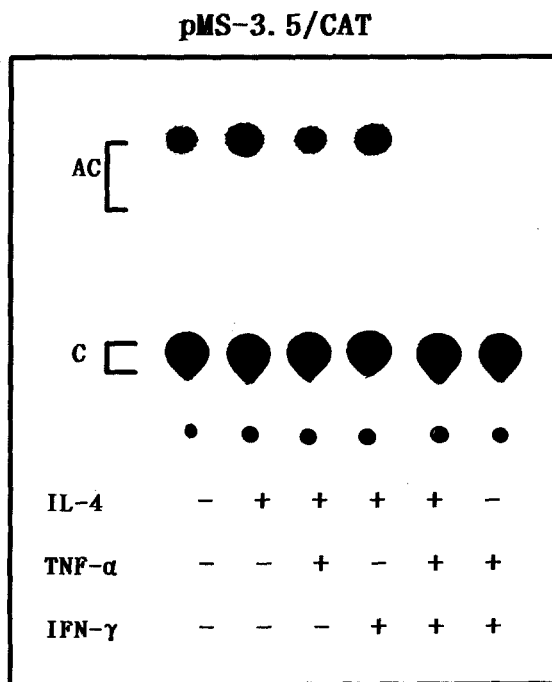


Fig. 3. Assay of COL1A2 promoter(pMS-3.5/CAT) activity in human skin fibroblasts cultured with IL-4 (5 ng/ml), TNF- α (10 ng/ml), and/or IFN- γ (1 μ g/ml). The figure shows CAT assay depicting the separation of acetylated (AC) and unacetylated (C) forms of [14 C]chloramphenicol by thin layer chromatography. Abbreviations were described in fig. 1.

beled Pro α 1(I) collagen, fibronectin, and GAPDH cDNA probes were hybridized with each mRNAs specifically. Pro α 1(I) collagen revealed two mRNA transcripts, the sizes of which were 5.8-kb and 4.8-kb(Fig.1). Fibronectin and GAPDH revealed one transcript each, and their sizes were 8.0-kb and 1.4-kb, respectively(Fig.1 & 2). There was no change of their sizes, which meant no alteration of

quality. For the quantitative assay of pro α 1(I) collagen mRNA steady-state level, a densitometer was used.

Effects of IL-4 on the steady-state accumulation of type I collagen and fibronectin mRNA

To examine the effect of IL-4 on the endogenous expression of type I collagen and fibronectin gene, the quantities of pro α 1(I) collagen and fibronectin mRNA were evaluated. Incubation of cultured human skin fibroblasts with IL-4(5 ng/ml) alone for a period of 48 hr resulted in a marked increase in the steady-state levels of type I collagen mRNA, which was the highest increase, 2.8-fold higher than the untreated control cells. This increase was selective in that the steady-state levels of fibronectin mRNA for GAPDH were not altered by IL-4(Table 1).

Effects of TNF- α and IFN- γ on type I collagen and fibronectin mRNA

Incubation with IL-4 together with TNF- α showed 1.5-fold increased pro α 1(I) collagen mRNA levels compared to the control, which showed suppressed levels by 47% compared to incubation with IL-4 alone. Incubation with IL-4 together with IFN- γ showed 2.3-fold increased pro α 1(I) collagen mRNA levels compared to the control, which showed suppressed levels by 19.6% compared to incubation with IL-4 alone. Incubation with IFN- γ together with TNF- α showed a suppressed mRNA production level, which was 73% of the control level. Co-treatment with all these three cytokines which completely abolished the expression of pro α 1(I) collagen gene increased by IL-4, revealed marked suppression to the level of 85% of

Table 2. Effects of TNF- α (10 ng/ml) and/or IFN- γ (1 μ g/ml) in combination with IL-4(5 ng/ml) on COL1A2 promoter activity in human skin fibroblasts

	CAT activity	Fold induction
Control	2.1 \pm 0.3	1.00
IL-4	5.2 \pm 0.6	2.48
IL-4 + TNF - α	2.8 \pm 0.2	1.33
IL-4 + IFN - γ	4.8 \pm 0.5	2.28
IL-4 + TNF - α + IFN- γ	0.8 \pm 0.3	0.38
IFN- γ + TNF - α	0.7 \pm 0.4	0.33

The values are mean SD from six separate experiments. Abbreviations were described in fig. 1. Values in the parentheses indicate fold induction.

the untreated control culture. Those increased by IL-4 and other decreases in fibronectin mRNA were not as sensitive as in type I collagen (Table 1).

Evidence for transcriptional regulation of the type I collagen gene at the promoter level

Fibroblasts were transfected with construct and cytokines were added to the culture media for 48 hr after glycerol shock. As the result of CAT assay, the percentage of acetylation were 2.1% in the untreated control, 5.2% in 5 ng/ml of IL-4, 2.8% in IL-4 with 10 ng/ml of TNF- α , 4.8% in IL-4 with 1 μ g/ml of IFN- γ , 0.8% in IL-4 with other two cytokines, and 0.7% in IFN- γ with TNF- α (Fig. 3). Compared to the control group, each promoter activity was increased by 2.5-fold, 1.3-fold, 2.3-fold, 0.4-fold and 0.3-fold in fibroblasts treated with IL-4 alone, IL-4 with TNF- α , IL-4 with IFN- γ , IL-4 with other two cytokines, and IFN- γ with TNF- α , respectively (Table 2).

DISCUSSION

Regulation of collagen production is an important step in biological situation such as tissue development, homeostasis, and wound healing. Collagen biosynthesis and collagen accumulation appears to result from increased production of collagen on a per-cell basis, and not from decreased turnover of the deposited collagen¹⁻².

The activation of fibroblast collagen production has been found to be associated with increased steady-state cellular levels of pro α 1(I) and pro α 1(III) collagen mRNA, suggesting activation of type I and type III procollagen gene expressions at the pretranslational or transcriptional level²⁴⁻²⁶. But the precise mechanisms leading to activation of collagen gene expressions are unknown so far. The role of cytokines from tissue-infiltrated inflammatory cells in the pathophysiology of fibrosis has been suggested²⁷. Mononuclear cells, predominantly lymphocytes and macrophages, secrete soluble factors capable of stimulating collagen synthesis by fibroblasts, suggesting that cytokines may stimulate collagen gene expression of fibroblasts^{25,28}.

A number of recently identified cytokines have shown significant influences on cell proliferation and collagen synthesis. The effect of IL-4 on fibroblast functions may be a "fibrogenic cytokine" that could be important in promoting biogenesis

of extracellular matrix proteins in normal wound healing and in pathological fibrosis in which mast cells and T lymphocytes play a central role²⁹. Elevation of steady-state levels of type I procollagen and fibronectin mRNAs in IL-4 treated cells suggests that this cytokine regulates extracellular matrix biogenesis by pretranslational mechanisms. IL-4 induced dose-dependent increase of collagen production, but non-collagen protein synthesis was not significantly altered, and a concomitant increased steady-state level of pro α 1(I) collagen mRNAs was observed, showing that IL-4 acts at a pre-translational level^{5,30}. Increased mRNAs may correspond either to an increased gene transcription or to an increased stability of the mRNAs. Incubation of cells transfected with pro α 2(I) collagen/CAT reporter gene chimeric construct with IL-4 resulted in marked up-regulation of the promoter activity in this study and other experiment¹⁵.

The interactions of IL-4 with fibroblasts could be modulated by other cytokines produced by inflammatory cells, for instance IFN- γ or TNF- α , which inhibits collagen synthesis⁸. In this study, incubation of human skin fibroblast with IL-4 (5 ng/ml) alone produced highest promoter activity, 2.5-fold higher increase than untreated control cells. Compared to the control, the steady-state levels of pro α 1(I) collagen mRNA showed a marked increase by 2.8 fold induction, but GAPDH were not affected. These results suggest that the expression of type I collagen gene can be up-regulated by IL-4 at the transcriptional level. Other two lymphokine factors, TNF- α and IFN- γ , have been shown to suppress collagen gene expression in cell cultures⁶. The decreased steady state levels of procollagen α (I) mRNA coupled with the decreased procollagen α 1(I) gene transcriptional activity indicates that TNF- α inhibits collagen synthesis at the transcriptional or pretranslational level^{6,31}.

According to Kähäri *et al.*¹⁵, the exposure of human skin fibroblasts transfected with pMS-3.5/CAT to human recombinant TNF- α resulted in inhibition of the pro α 2(I) collagen promoter activity, while TNF- α did not have a significant effect on the expression of the cells transfected with pSV2CAT. This observation suggested that TNF- α exerts its inhibitory effects on type I collagen gene expression at the transcriptional level. Inhibition of collagen production caused by TNF was

dose and time-dependent⁶. In this study, TNF- α had a more potent suppressive effect on the gene expression and CAT activity for the collagen gene than IFN- γ . On the effect of TNF- α on gene expression of fibronectin, normal dermal fibroblasts demonstrated decreased levels of fibronectin mRNA production. The effects of IFN- γ on human skin fibroblasts has demonstrated the decreased mRNA levels in type I collagen and fibronectin, and reduction of collagen accumulation in experimental fibrosis³². It is interesting how these cytokines affect the promoter activity and whether these cytokines modulate the *cis*-acting DNA sequences controlling the transcription of the type I collagen genes or not.

In this study, the modulation of type I procollagen promoter activity was examined by a variety of cytokines, including IL-4, TNF- α , and IFN- γ . The approach was made possible by previous studies which established that the human pro α 2(I) collagen/CAT reporter gene chimeric construct is able to mimic physiological expression of the gene when transiently expressed in human fibroblastic cells¹⁷. This construct contains -3.5 kb of 5' flanking DNA from human pro α 2(I) collagen gene, cloned in front of the structural gene for CAT. Incubation of cells transfected with this chimeric construct with IL-4 resulted in marked up-regulation of the promoter activity in cultured normal human skin fibroblasts, which was approximately 2.5-fold increase compared to the control. The increase or decrease noted in the promoter activity was in parallel with pro α 1(I) collagen mRNA levels according to combination of cytokines. Therefore, the increase noted in the type I procollagen mRNA steady-state levels appeared to reflect primarily activation of transcription for the gene³³.

Both TNF- α and IFN- γ showed counteraction to the IL-4-induced enhancement of type I procollagen gene expression. However, the mechanism of inhibition is known to be different. According to the observation of Kähäri et al¹⁵, TNF- α appeared to affect type I procollagen gene expression on the transcriptional level by suppressing the pro α 2(I)collagen promoter activity, while IFN- γ appeared to exert its effect primarily through destabilization of the mRNA and had only a minimal effect at the transcriptional level. This study showed that IL-4 with IFN- γ did not apparently decrease the IL-4 induced CAT activity, suggesting that IFN- γ

has the minimal effect on the transcriptional level. Treating with IL-4 alone showed the highest promoter activity, but the remarkable decreased activity, 38% of the control, was revealed when TNF- α and IFN- γ were added to IL-4. The activation of pro α 2(I) promoter gene produced by IL-4 was completely abolished by treating with IL-4 together with IFN- γ and TNF- α . The synergistic suppressive effect of TNF- α and IFN- γ on pro α 2(I)collagen promoter activity can be explained by the previous report which IFN- γ could increase the number of TNF- α receptors without causing a significant change in the apparent binding affinity in cultured cells, thus making these cells more susceptible to TNF- α modulation³⁴. IFN- γ leads to direct stimulation of TNF-receptor synthesis, possibly due to increased transcription of its mRNA. But the mRNA for collagenase was increased by TNF- α and no synergism with IFN- γ was obvious³⁵.

To understand how collagen transcription is selectively controlled under the physiological conditions, a number of studies have recently focused on the identification and characterization of *cis*-acting DNA regulatory elements in mammalian genes. Collagen expression is known to be regulated by unique arrangement of complex networks of interacting sequences which reside both downstream and upstream of the transcription initiation sites³⁶. The synergistic inhibition of collagen gene expression induced by TNF- α in combination with IFN- γ is already known in other systems^{13,14}. The effects of TNF- α and IFN- γ are also synergistic and potentiated the specific reduction of collagen synthesis. A better understanding of synergistic effects of cytokines might not only shed light on the pathogenetic event in fibrosis or wound healing defects, but also open new concepts for therapeutic approaches. This result suggests that combination of cytokines would provide a potential approach for a more efficient pharmacological suppression or enhancement of collagen formation than afforded by either one of the peptide factors alone.

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