

The Compliance of Collagen Gels Regulates Transforming Growth Factor- β Induction of α -Smooth Muscle Actin in Fibroblasts

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Wound contraction is mediated by myofibroblasts, specialized fibroblasts that appear in large numbers as the wound matures and when resistance to contractile forces increases. We considered that the regulation of myofibroblast differentiation by wound-healing cytokines may be dependent on the resistance of the connective tissue matrix to deformation. We examined transforming growth factor- β_1 (TGF- β_1) induction of the putative fibroblast contractile marker, α -smooth muscle actin (α -SMA), and the regulation of this process by the compliance of collagen substrates. Cells were cultured in three different types of collagen gels with wide variations of mechanical compliance as assessed by deformation testing. The resistance to collagen gel deformation determined the levels of intracellular tension as shown by staining for actin stress fibers. For cells plated on thin films of collagen-coated plastic (ie, minimal compliance and maximal intracellular tension), TGF- β_1 (10 ng/ml; 6 days) increased α -SMA protein content by ninefold as detected by Western blots but did not affect β -actin content. Western blots of cells in anchored collagen gels (moderate compliance and tension) also showed a TGF- β_1 -induced increase of α -SMA content, but the effect was greatly reduced compared with collagen-coated plastic (<3-fold increase). In floating collagen gels (high compliance and low tension), there were only minimal differences of α -SMA protein. Northern analyses for α -SMA and β -actin indicated that TGF- β_1 selectively increased mRNA for α -SMA similar to the reported protein levels. In pulse-chase experiments, [35 S]methionine-labeled intracellular α -SMA decayed most rapidly in floating gels, less rapidly in anchored gels, and not at all in collagen plates after TGF- β_1 treatment. TGF- β_1 increased α_2 and β_1 integrin content by 50% in cells on collagen plates, but the increase was less marked on anchored gels and was undetectable in floating gels. When intracellular tension on collagen substrates was reduced by preincubating cells with blocking antibodies to the α_2 and β_1 integrin

subunits, TGF- β_1 failed to increase α -SMA protein content in all three types of collagen matrices. These data indicate that TGF- β_1 -induced increases of α -SMA content are dependent on the resistance of the substrate to deformation and that the generation of intracellular tension is a central determinant of contractile cytoskeletal gene expression. (*Am J Pathol* 1999, 154:871–882)

Wound contraction and remodeling of granulation tissue involve the differentiation of fibroblasts into myofibroblasts, cells that typically express high levels of α -smooth muscle actin (α -SMA^{1,2}). The myofibroblasts form tight adhesions to the substrate,³ and it appears that their differentiation is temporally associated with the resistance of the wound to contractile forces.² Myofibroblasts also exhibit the formation and arrangement of stress fibers along the long axis of the cells,^{4,5} which in turn facilitates tissue contraction.

To model wound contraction, hydrated collagen lattices have been used as floating or anchored matrices^{6–8} that resemble early and later stages of wound remodeling, respectively. Notably, the elevated breaking strength of mature skin wounds is due to increased reorganization and cross-linking of collagen,⁹ which can be modeled by varying the biophysical properties of collagen gels. Thus, contraction of floating collagen matrices provides a model for a mechanically relaxed tissue with low tensile strength comparable to resting dermis⁸ or to very early stages of wound healing,⁹ whereas anchored matrices develop into a more stressed tissue that resembles granulation tissue. Notably, in two variations of the *in vitro* collagen matrix reorganization model, the morphology and the behavior of fibroblasts in the gels reflect intracellular tension levels. In floating collagen matrices, fibroblasts develop a stellate morphology with long processes and a well developed subcortical actin meshwork.⁷ In marked contrast, cells in anchored matrices become bi-

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polar, orient along lines of tension, develop prominent stress fibers and fibronexus junctions, and resemble myofibroblasts.¹⁰

The molecular mechanisms of collagen contraction are largely unknown. Previous studies have suggested an essential role for cell surface collagen receptors.^{11–13} Indeed, a direct role for the collagen-binding integrin $\alpha_2\beta_1$ in fibroblast-mediated collagen gel contraction has been demonstrated.^{12,13} Members of the β_1 integrin family are known to mediate fibroblast interactions with collagen fibers,^{14,15} and wound-healing cytokines such as transforming growth factor (TGF)- β may enhance collagen gel contraction by increasing the expression of the $\alpha_2\beta_1$ integrin collagen receptor.¹⁶ This integrin-dependent force generation appears to require the cytoplasmic domain of the $\alpha_2\beta_1$ integrin as the extracellular domain is insufficient to mediate contraction.¹² Several growth factors (eg, TGF- β , platelet-derived growth factor (PDGF), and fibroblast growth factor) can modulate contraction of collagen matrices. TGF- β_1 is an extensively studied cytokine on the basis of its importance in wound healing^{17–19} and matrix formation^{19,20} and possibly in regulating α -SMA expression.²¹ It can stimulate contraction of both floating and anchored collagen matrices *in vitro*,²² but this mechanism is poorly understood.

We have assessed previously cell-mediated remodeling of floating collagen matrices and showed that α -SMA is functionally important for collagen gel contraction.²³ Although all fibroblasts probably exhibit some degree of contractile ability, fibroblast populations that express α -SMA are evidently more contractile. Furthermore, studies on the effects of TGF- β on fibroblasts from wounds at different stages of repair and normal skin show that cells from older granulation tissue contract collagen matrices faster than cells from younger granulation tissue or normal skin.²⁴ The source of this variability of contractility between fibroblast populations from different stages of wound development is not understood. In view of the increased mechanical strength and reduced compliance of tissues at later stages of wound healing,^{9,25} and because of the importance of integrins in gel contraction,^{12,13} we considered that fibroblast responsiveness to TGF- β is regulated by the compliance of the supporting matrix and modulation through integrins. Currently, it is unknown whether the compliance of the matrix affects TGF- β -induced up-regulation of α -SMA and how tension-dependent cell responses to TGF- β_1 are regulated. Based on *in vivo* results showing that the timing of the switch from the fibroblastic to the myofibroblastic phenotype² is temporally associated with the resistance of the wound to contraction,⁹ we studied the effect of TGF- β_1 on α -SMA expression by human fibroblasts cultured in collagen matrices with three different levels of resistance to deformation. Culture of cells in these three types of collagen substrates promoted the development of different levels of intracellular tension and allowed us to test the hypothesis that intracellular tension modulates the TGF- β_1 -induced expression of cytoskeletal genes associated with wound contraction.

Materials and Methods

Cell Culture and Collagen Substrates

Primary cultures of human gingival fibroblasts were obtained from biopsies of normal gingiva in patients aged 10 to 16 years as described.²⁶ Cells at passages 3 to 12 were used for all experiments. For all three types of collagen matrices (see below), cells were initially plated as monolayer cultures in α -minimal essential medium (α -MEM) plus 15% v/v fetal bovine serum on plastic tissue culture dishes. The medium was removed and cells were incubated with α -MEM and TGF- β_1 (R&D Systems, Minneapolis, MN) or vehicle (PBS) for 3 days. Cells were incubated with TGF- β_1 at concentrations of 1, 5, 10, and 20 ng/ml. Immunoblotting for α -SMA (see below) showed a dose-dependent response to TGF- β_1 , and the optimal dosage was 10 ng/ml TGF- β_1 . Subsequently, all experiments were performed at this dose. Notably, others^{17,27} have used similar doses for studies of TGF- β -induced actin gene expression that are equivalent to levels of TGF- β found in wound fluid.²⁸

TGF- β treatments consisted of two protocols. In the first protocol, cells on tissue culture plastic were treated with PBS vehicle (controls) or with TGF- β_1 (experimentals) at 10 ng/ml for 3 days in α -MEM without serum. Cells were then trypsinized and replated on either tissue culture plastic coated with collagen or in anchored or floating gels (see below) and incubated with either vehicle or TGF- β_1 for 3 to 4 days. In the second protocol, both controls and experimentals were treated with TGF- β_1 for 3 days on tissue culture plastic and then were trypsinized, inoculated onto collagen-coated plates, anchored collagen gels, or floating gels and incubated with either vehicle (controls) or TGF- β_1 (experimentals) for 3 to 4 days.

Collagen solutions were prepared as described.⁷ We used type 1 collagen as the substrate for all types of gels to validate comparisons based solely on the mechanical properties of the gels. Briefly, under sterile conditions, a collagen solution was prepared from 0.3 ml of 10X concentrated α -MEM, 0.3 ml of 0.26 mol/L NaHCO₃ buffer, 0.3 ml of fetal bovine serum (serum), 0.12 ml of 0.1 mol/L NaOH, and 1.5 ml of Vitrogen 100 (Collagen Corp., Santa Clara, CA). For relatively low-compliance (ie, rigid) collagen substrates, films of collagen (~10 μ m thick) were prepared on tissue culture plastic and polymerized by neutralizing the collagen films to pH 7.4. To block non-specific binding, plates were incubated with 1% (w/v) bovine serum albumin (BSA) for 4 hours at 4°C. Before plating the cells, dishes were rinsed with PBS to remove excess BSA. Cells were plated subsequently at 7.5×10^4 cells/cm² and incubated in α -MEM without serum in the presence or absence of TGF- β_1 (see below). For anchored or floating gels, a cell suspension of 4×10^5 cells/ml in α -MEM without serum was added to the solution. Gel solutions were pipetted into tissue culture or non-tissue-culture dishes to obtain anchored or floating gels, respectively, as described previously.²⁹ Collagen gels were incubated at 37°C in 95% air and 5% CO₂ until polymerization was completed. The gels were covered

with α -MEM and incubated with TGF- β_1 or vehicle for 3 to 10 days.

Physical assessments of gel compliance were performed with a mechanical deformation tester (Dynatek Dalta, Galena, MO) in which a pair of 5-mm-diameter disks was used to obtain measurements of gel compression in a 15-second deformation period. The tester was operated in the stroke mode, which in turn controlled the position of the actuator. For all gel types, the actuator function was a linear compression stroke of 1-mm displacement over a 15-second duration. The displacement of the compression disk was measured directly by a linearly variable differential transducer, and loads were measured by a load cell (50 g). The precision of the displacement transducer was $\sim 1 \mu\text{m}$. Measurements were conducted at 30°C in α -MEM. All samples were representative 5-mm-diameter circles cut out of the different types of gels that were prepared. Displacement and load were plotted separately over each 15-second cycle for each sample ($n = 5$ separate samples for each gel type), and a best-fit curve over the duration of the compression stroke was obtained by linear regression analysis to estimate displacement *versus* time (mm/second) and load *versus* time (g/second) functions. Gel compliance was expressed in terms of the slope of the individual displacement *versus* time and load *versus* time functions and expressed as the average load *versus* gel compression over the entire 15-second sampling period (g/mm). A full 1-mm compression stroke was achieved for the anchored and floating gels, but because of the thin gels on the collagen-coated plates ($\sim 10 \mu\text{m}$), only minimal displacements were recorded.

Gel Contraction Assay

Cells on tissue culture plastic were serum starved for 48 hours, pretreated with TGF- β_1 or vehicle for 3 days and were then incubated in the three different types of collagen matrices described above. We determined the effect of TGF- β_1 on remodeling of anchored and floating gels using contraction assays conducted on triplicate cultures. $^3\text{H}_2\text{O}$ was added to the culture media, and the radioactivity of the gels at certain time points was measured.³⁰ Samples of cell/collagen solutions (400 μl , 4×10^5 cells/ml) were pipetted into 35-mm tissue and non-tissue-culture dishes. After polymerization, gels were covered with α -MEM without serum containing 1 μCi of $^3\text{H}_2\text{O}$ (1 mCi/g; Dupont, Boston, MA) with or without TGF- β_1 . Equilibration of radioactivity in the gels required 30 minutes. At the times indicated, the medium was removed, and the gels were rinsed quickly and dissolved in 0.5 ml of 1 mol/L NaOH. The samples were neutralized with HCl, mixed with 2 ml of scintillation solution, and counted in a Beckman scintillation spectrometer. The initial gel volume was determined in gels without fibroblasts.

Fluorescence Microscopy

The effect of TGF- β_1 on development and reorganization of stress fibers was investigated using affinity labeling of

filamentous actin. Cells were plated on collagen-coated coverslips in 24-well multichamber slides (7.5×10^4 cells/cm²) and treated with TGF- β_1 as described above. Cells were washed with PBS and fixed with 1.5% paraformaldehyde, and F-actin was stained for 15 minutes with 5×10^{-6} mol/L tetramethylrhodamine isothiocyanate (TRITC)-phalloidin (Sigma Chemical Co., St. Louis, MO) in PBS containing 0.01% Nonidet. Finally, the coverslips were washed, air dried, and mounted with an anti-fade reagent. The total cell fluorescence due to (TRITC)-phalloidin was measured in a standardized area of cytoplasm (100 μm^2) using a Leitz MVP-SP spectrofluorimeter (Wetzlar, Germany) and a 25 \times Plan Apo objective with excitation at 530/30 nm and emission monochromators set at 600/3 nm. The photomultiplier tube voltage was set to 599 V and the amplifier gain to 4 \times . The fluorescence of standardized unstained cell areas was subtracted from each sample measurement to correct for background and dark current. To stain for α -SMA, cells were incubated with a mouse anti-human α -SMA monoclonal antibody (1:50 dilution, clone 1A4,³¹ Sigma) for 1 hour at 37°C followed by a fluorescein isothiocyanate (FITC)-conjugated goat anti-mouse antibody (1:100; Sigma) for 1 hour at 37°C. Preimmune mouse and goat sera were used to block nonspecific staining before antibody incubation. The cells were examined in a Leitz Orthoplan microscope equipped with specific filters for rhodamine and fluorescein. Cells were imaged on a confocal microscope (Leica CLSM, Heidelberg, Germany) with a 40 \times oil immersion lens (numerical aperture = 1.2). Optical sections were obtained at a nominal thickness of 1 μm . In some samples, computer-generated images were combined to reconstruct a composite image. For staining cells in collagen gels, fluorescence staining for α -SMA and F-actin was conducted as described for monolayer cultures.

Western Blotting

We quantified TGF- β_1 -induced modulation of α -SMA content by immunoblot analysis. For harvesting fibroblasts from the gels, each gel was rinsed thoroughly with Mg^{2+} , Ca^{2+} -free PBS and incubated for 10 minutes at 37°C with 0.3 ml of 0.05% trypsin/0.53 mmol/L EDTA solution, followed by a 20- to 30-minute incubation with 0.35 ml of collagenase (5 mg/ml). After cells were dispersed completely, enzymatic activity was blocked by the addition of 0.05 ml of serum. The dispersed cells were counted with a hemocytometer before collection by centrifugation at 14,000 RPM for 10 minutes at 22°C and 0.05 ml of extraction buffer (2% Triton X-100, 160 mmol/L KCl, 40 mmol/L Tris/HCl, 20 mmol/L EGTA, 10 mmol/L phenylmethylsulfonyl fluoride, 1 mmol/L leupeptin, 1 mmol/L benzamide; Sigma). Protein content was assessed by the BioRad assay. Samples were boiled for 3 minutes at 95°C, and equal amounts of protein were electrophoresed on a 10% SDS gel. Separated proteins were transferred to nitrocellulose filters and probed with the mouse monoclonal antibody for α -SMA followed by a horseradish-peroxidase-conjugated second antibody and devel-

oped with ECL reagents (Amersham, Arlington Heights, IL). Blots were stripped and probed with a monoclonal antibody for β -actin (clone AC-15, Sigma) for comparison. Subsequently, X-OMAT Kodak films were exposed to the blots, and the density of the bands was estimated by Scan Analysis (Biosoft, Cambridge, UK). The ratio of the density of α -SMA to β -actin was calculated for each sample.

Integrins

We studied the effect of TGF- β_1 on the expression of α_2 and β_1 integrin subunits. Immunoblotting for α_2 and β_1 integrins was performed with mouse monoclonal antibodies (clone P1E6 from Calbiochem, Temecula, CA, and 4B4-RD1 from Coulter, Burlington, Ontario, Canada). The importance of the $\alpha_2\beta_1$ integrin on TGF- β_1 -induced expression of α -SMA was examined using mouse monoclonal antibodies to block β_1 and/or α_2 integrin subunits. Fibroblasts (7.5×10^4 cells/cm²) were preincubated with mouse monoclonal antibodies for β_1 integrin (clone 4B4-RD1) and/or α_2 integrin (clone P1E6) at room temperature, and the cells were allowed to adhere to collagen-coated plastic petri dishes. The next day media were removed, cultures were covered with α -MEM containing 10 ng/ml TGF- β_1 and supplemented with β_1 integrin (1:30) and/or α_2 (1:50) integrin antibodies, and incubated for 3 days. The antibodies were replenished once during the 3-day incubation with the same concentrations used for preincubation. Each experimental condition was prepared in triplicate. To analyze the antibody-induced changes of cell morphology, cells were observed with a phase contrast microscope (Nikon, Mississauga, Ontario, Canada), and photographs were taken on the third day of incubation. At the end of treatment, immunoblotting for α -SMA was conducted as described before, and blot density was normalized to β -actin.

Flow Cytometry

Cell suspensions were prepared (0.01% trypsin), fixed with 3.7% formaldehyde, permeabilized in 0.02% Triton with PBS, and stained for α -SMA as described.²³ First (anti- α -SMA) and second antibody (FITC-conjugated goat anti-mouse) dilutions were 1:25 and 1:50, respectively. Cells were washed and resuspended in Mg²⁺, Ca²⁺-free PBS. Samples were analyzed on a FAC-Star Plus flow cytometer (Becton-Dickinson, Mississauga, Ontario, Canada) with 488-nm excitation and 530/30-nm band pass filter for FITC. For all flow cytometry analyses, at least 1×10^4 cells were assessed in each sample, and only cells with forward and orthogonal light scatter characteristics similar to whole, intact fibroblasts were included in the analysis by electronic gates previously established for fibroblasts. Mouse monoclonal antibodies for α_2 (1:20 dilution, clone P1E6) and β_1 (undiluted, clone 4B4-RD1) integrins were used for staining.

Northern Analysis

Northern blots were performed on monolayer cultures grown on plastic tissue culture dishes after 3 days of TGF- β_1 treatment or vehicle. As described above for Western blots, cells were subsequently grown on either collagen-coated plates or anchored or floating gels in the presence or absence of TGF- β_1 for 3 days to determine the effect of TGF- β_1 on mRNA levels for α -SMA. The cells were isolated from gels as described for Western blotting, and total RNA was isolated as described.²⁹ RNA samples were separated in denaturing 1.3% formaldehyde-agarose gels, transferred to a nylon membrane (Bio-Rad) and cross-linked by ultraviolet light. The McMolly Tetra program (Soft Gene) was used to design 32-mer oligonucleotides (5'TCCACAGGACATTCA-CAGTTGTGTGCTAGAGA-3' and 5'-CCATGCCAATCT-CATCTTGTCTTCTGCGCAAG-3') complementary to specific sequences of the α -SMA and β -actin mRNA 3' untranslated region, respectively. The oligonucleotides were labeled with [γ ³²P]ATP (Dupont NEN) using 3' end labeling. The blots were washed twice with 0.5% SSC plus 0.5% SDS in dH₂O, each time for 30 minutes at 55°C, and exposed to Kodak X-OMAT films at -70°C between intensifying screens overnight. The blots were stripped and reprobbed with [γ ³²P]ATP-labeled glyceraldehyde phosphate dehydrogenase cDNA.

³⁵S Labeling and Immunoprecipitation

Confluent cell monolayers were serum starved for 24 hours and treated with vehicle or TGF- β_1 for 72 hours. Cells were metabolically labeled for 24 hours with [³⁵S]methionine (100 μ Ci/ml; ICN Biochemicals) in methionine-free α -MEM. Cells were plated on collagen-coated tissue culture plastic, anchored gels, or floating gels in the presence or absence of TGF- β_1 . Cells were isolated from collagen matrices as described above. Cell pellets were solubilized in 200 μ l of lysis buffer. Insoluble material was removed by centrifugation at $10,000 \times g$ for 5 minutes at 4°C. Radioactivity in cell lysates was counted, and equal amounts of radioactivity were used in immunoprecipitation assays. Supernatants were immunoprecipitated with α -SMA antibody overnight at 4°C. Immuno-complexes were recovered by binding to protein A-Sepharose (Zymed Laboratories, South San Francisco, CA) and washed four times with 25 mmol/L Tris-buffered saline (pH 7.4) containing 0.5% Triton X-100 and 1 mg/ml BSA and twice with 0.5 mmol/L NaCl and 25 mmol/L Tris-HCl (pH 7.4). The immunocomplexes were analyzed by electrophoresis on 10% polyacrylamide gels followed by fluorography and scanning for quantification for the density of the band. The data are expressed as percent remaining density for the different sample days, and these data provided an estimate of the percent decay of radioactivity due to nascent α -SMA.

TGF- β Quantification

The levels of active and total TGF- β_1 produced by fibroblasts and the effect of exogenous TGF- β_1 on these

levels were measured with a human TGF- β_1 immunoassay (Quantikine, R&D Systems) with a sensitivity of 5 pg/ml. Cultures of fibroblasts in floating and anchored gels or on collagen plates were prepared as described. Cells were incubated with α -MEM with or without 10 ng/ml TGF- β_1 . After 3 days of culture, supernatants were collected, triplicate samples and activated, and non-activated forms were assayed. Activation was performed by adding 0.1 ml of 1 N HCl to 0.5 ml of cell supernatants for 10 minutes followed by neutralization with 0.1 ml of 1.2 N NaOH. To determine the amount of adherence of TGF- β_1 to collagen, 10 ng/ml TGF- β_1 was incubated in collagen-coated plates without cells, and the medium was assayed after 3 days. The samples containing exogenous TGF- β_1 were diluted 1:5 to adjust the concentration of TGF- β_1 to the linear range of the kit. Samples were analyzed in a 96-well microtiter plate coated with recombinant human TGF- β_1 soluble receptor type II. The optical density of wells was determined with a microtiter plate reader set to 450-nm absorbance. The readings at 570 nm were subtracted from readings at 450 nm to correct for nonspecific absorbance. A standard curve was constructed by plotting the mean absorbance for each standard against the concentration, and a best-fit curve was determined by regression analysis.

To determine the specificity of the effect of TGF- β_1 on α -SMA expression, an anti-TGF- β neutralizing antibody was used to block the effect of TGF- β_1 . Cells were plated on 35-mm collagen-coated dishes and anchored and floating gels. After 24 hours, media were replaced by α -MEM with or without 10 ng/ml TGF- β_1 and with or without anti-TGF- β_1 neutralizing antibody (15 μ /ml; R&D Systems). After 3 days, α -SMA content was quantified by Western blot.

Results

Effect of TGF- β_1 on Collagen Gel Contraction

As several growth factors in serum potentially can influence collagen gel contraction, we have focused specifically on the effect of TGF- β_1 in serum-free conditions. Contraction was determined by measuring gel volume based on the partitioning of $^3\text{H}_2\text{O}$ between the gel phase and the surrounding medium in time course studies.³² If cells were not preincubated with TGF- β_1 and the cytokine was added only from the beginning of the gel contraction assay, only small differences of gel contraction rate were observed between experimentals and controls from 0 to 10 days (data not shown). If, however, cells were preincubated with TGF- β_1 for 3 days before incubation in the gels and the cytokine treatment was continued during the gel contraction phase, large and statistically significant ($P < 0.01$) differences were seen (Figure 1). The floating gels showed a maximal difference of contraction between TGF- β_1 and controls at day 2, which then decreased rapidly. In contrast, anchored gels exhibited a prolonged maximal difference from days 4 to 10. In view of these findings we assessed whether there was a temporal relationship between TGF- β_1 -induced up-regula-

tion of α -SMA and the ability of the cells to contract collagen matrices. Cells on collagen plates were incubated for 1 to 3 days with TGF- β_1 at a concentration of 10 ng/ml. Immunoblotting demonstrated that cells exposed to TGF- β_1 for only 1 day showed little change in α -SMA content whereas incubation for 3 days showed an optimal increase in α -SMA. These data are consistent with the minimal effects of TGF- β_1 on gel contraction without cytokine preincubation and greatly enhanced contraction with preincubation and provided a rationale for conducting all additional experiments with either a 3-day preincubation of cells with 10 ng/ml TGF- β_1 or with vehicle.

Gel Compliance

The floating gels were highly compliant as determined by mechanical testing and offered little resistance to deformation (5.67 ± 0.56 mg/ μ m). As they were physically separated from the culture dish, this type of gel could contract in all three dimensions. In contrast, the anchored gels offered increased resistance to deformation (13.6 ± 2.25 mg/ μ m; $P < 0.01$), and as they were fixed to the sides and the bottom of the dish, these gels could only contract in the z axis. Collagen-coated plates showed high resistance to deformation (57.8 ± 20.4 mg/ μ m; $P < 0.05$) and were not free to contract significantly in any dimension.

TGF- β_1 Enhancement of Stress Fibers Depends on Gel Compliance

In collagen-coated plates (ie, low compliance), compared with controls, TGF- β_1 promoted the development of stress fibers that were prominently stained for α -SMA (Figure 2). As reorganization of stress fibers is promoted by the development of intracellular tension,⁶ we assessed quantitatively whether TGF- β_1 could regulate F-actin content. Affinity labeling of filamentous actin by rhodamine-phalloidin in cultures on collagen-coated plates was measured by single-cell photometry. There was a >3-fold increase in F-actin content compared with untreated controls (TGF- β_1 = 74.3 ± 4.2 intensity units, and controls = 22.7 ± 5.8 intensity units; $P < 0.01$). Confocal imaging of cells stained for filamentous actin in monolayer cultures or anchored or floating gels revealed the development of prominent stress fibers parallel to the long axis of cells in monolayer cultures and to a lesser extent in anchored gels. Stress fibers were not detected in floating gels. As noted above, F-actin staining was significantly enhanced with TGF- β_1 in monolayer (Figure 2, A and B) and to a lesser extent in anchored gels (Figure 2, C and D); however, this enhancement was not detectable in floating gels (Figure 2, E and F), suggesting that stress fiber formation and the development of intracellular tension is dependent on the compliance of the substrate and may regulate TGF- β_1 -induced effects on α -SMA.

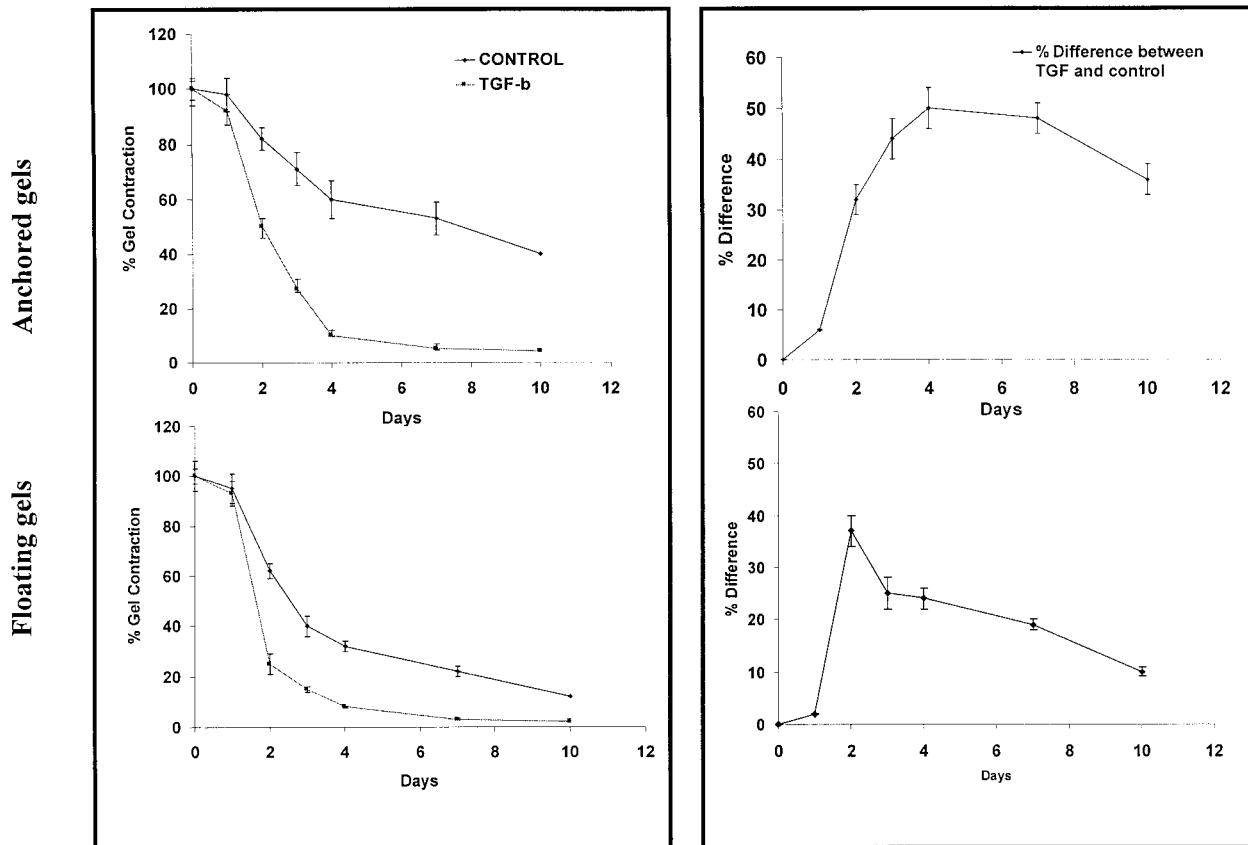


Figure 1. Collagen gel contraction and regulation by TGF- β_1 . Left panels: Cells in anchored (top) and floating (bottom) collagen gels were incubated in α -MEM containing $^3\text{H}_2\text{O}$ ($1 \mu\text{Ci/ml}$) with or without TGF- β_1 (10 ng/ml). At 1, 2, 3, 4, 7, and 10 days, gel volume was determined using scintillation counting of labeled gels. For each time point, $n = 3$ independent samples. Results are presented as mean percent gel volume \pm SEM. Right panels: Data from left panels re-plotted to show percent differences between TGF- β_1 -treated cultures and controls.

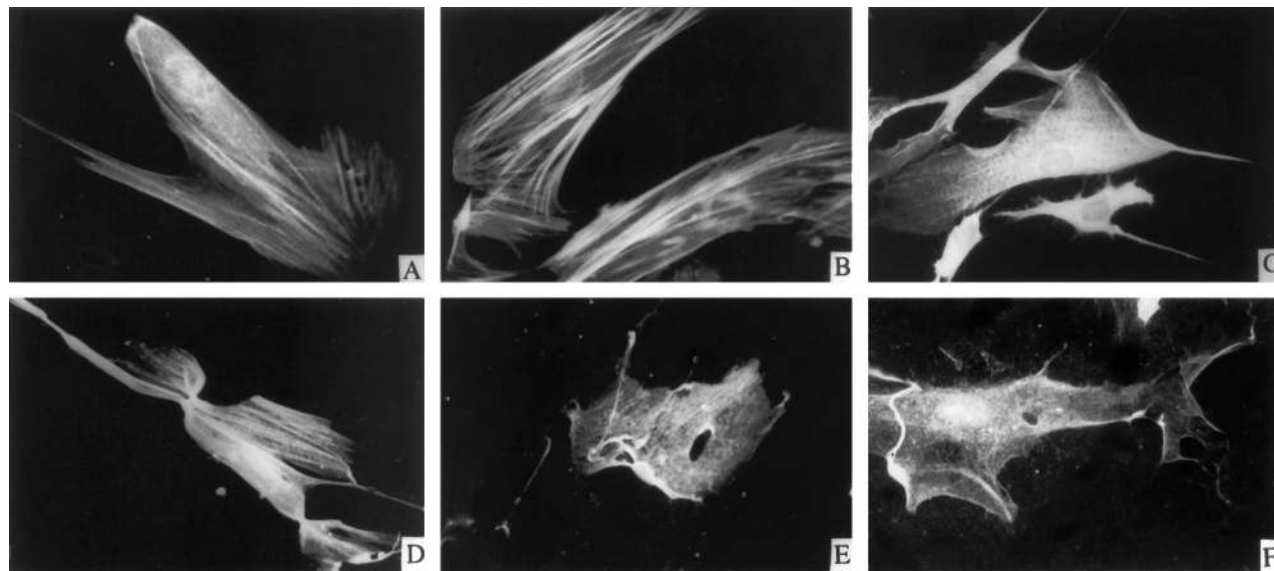


Figure 2. Immunofluorescence of α -SMA in cells on different substrates. Confocal micrographs of cells on collagen-coated plastic (A and B), anchored collagen gels (C and D), and floating collagen gels (E and F) were incubated with vehicle (A, C, and E) or with TGF- β_1 (10 ng/ml ; B, D, and F) for 3 days. α -SMA was immunostained with anti- α -SMA antibody and FITC-conjugated goat anti-mouse antibody.

Effect of TGF- β_1 on α -SMA Protein and mRNA Content

We determined quantitatively whether TGF- β_1 induces changes of α -SMA protein content as a function of β -actin content. Western blotting was performed and β -actin content was used for comparison and for equilibration of protein loading as we found that in preliminary experiments, when lanes were loaded with equal amounts of protein, β -actin did not change appreciably under different experimental conditions. For cells grown on tissue culture plastic but without collagen coating, the amount of α -SMA in serum-free conditions was increased ~three-fold by TGF- β_1 (10 ng/ml; 3 days) compared with controls (Figure 3A). For cells grown on collagen gels, all data were normalized to β -actin content and are expressed as a ratio to overcome any variations in the efficiency of cell extraction and cytoskeletal protein isolation from the different substrates. After adjusting for equal protein loading by BioRad assay, β -actin did not show any change between the different matrices for controls and TGF- β_1 -treated cultures. Cells grown on collagen-coated plates showed ninefold increases in α -SMA: β -actin ratios after a 3-day TGF- β_1 pretreatment and another 3 days of TGF- β_1 treatment compared with controls. When cells were pretreated identically and then grown in anchored collagen gels and treated with TGF- β_1 , the α -SMA: β -actin ratio increased by 2.9-fold compared with controls. For floating gels, TGF- β_1 induced only a 1.4-fold increase in the α -SMA: β -actin ratio (Figure 3B). These findings are consistent with the results of immunofluorescence staining and also indicate that when TGF- β_1 -pretreated cells (with known increases of α -SMA content; Figure 3A) are subsequently grown on different matrices (collagen-coated plastic, anchored gels, and floating gels), they continue to exhibit marked differences in their relative α -SMA content on different matrices. To separate the effect of TGF- β from that of gel compliance on α -SMA levels, we pretreated both experimentals and controls with TGF- β for 3 days, and then on inoculation into collagen gels, cells were treated with either vehicle or TGF- β (treatment protocol 2). Cells grown on collagen-coated plates showed a ~sixfold increase of the α -SMA: β -actin ratio after TGF- β treatment. There were 2.3-fold and 1.2-fold increases for anchored gels and floating gels, respectively, after TGF- β treatment. These ratios were very similar to those observed without TGF- β pretreatment.

Although these experiments were conducted without serum, it was possible that the increases of the α -SMA: β -actin ratios were due to TGF- β -mediated increases of cell numbers. However, cell counts at the end of the treatment periods showed only modest increases between experimentals and controls for all of the collagen gel types (collagen plates: controls, 1.0×10^6 cells/plate, and experimentals, 1.6×10^6 cells/plate; anchored gels: controls, 4.0×10^5 cells/ml of collagen gel, and experimentals, 4.8×10^5 cells/ml of collagen gel; floating gels: controls, 4.0×10^5 cells/ml of collagen gel, and experimentals, 4.2×10^5 cells/ml of collagen gel).

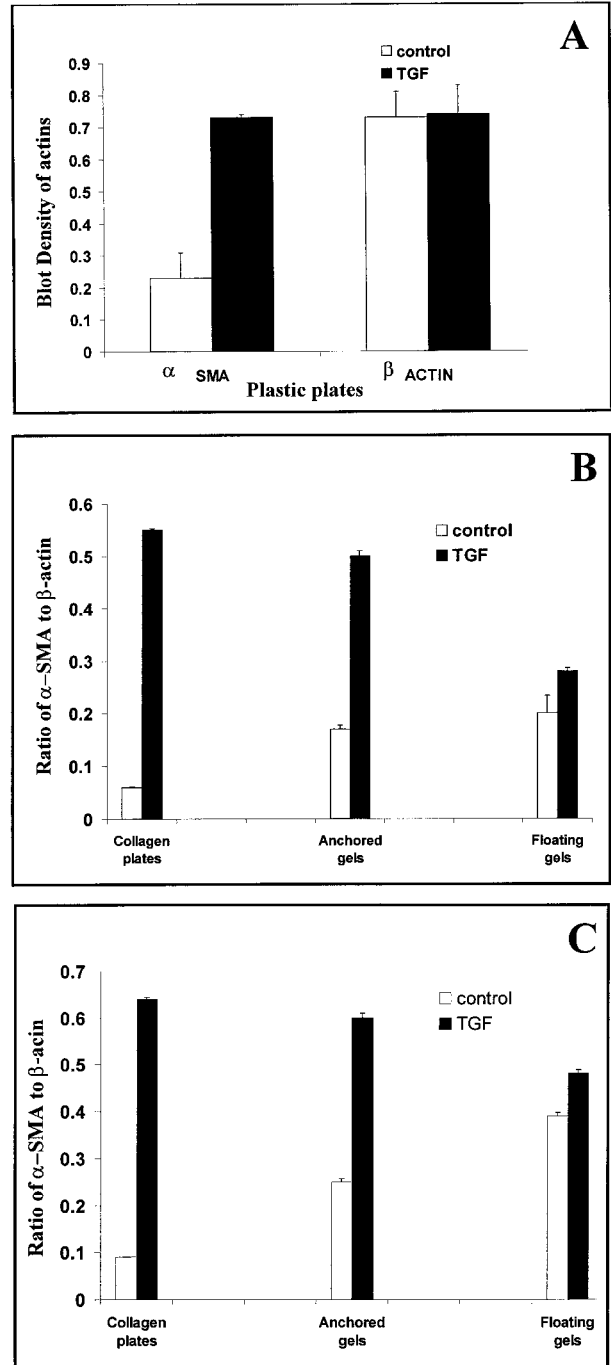


Figure 3. Western blots of α -SMA protein content. **A:** Cells on plastic plates were incubated with vehicle control or TGF- β_1 (10 ng/ml) for 3 days. Immunoblots of α -SMA and β -actin were scanned and the densities plotted (mean density \pm SEM). **B:** Cells were preincubated for 3 days with vehicle or with TGF- β_1 (10 ng/ml) on plastic plates as in **A** and then inoculated on collagen-coated plates or anchored collagen gels or in floating gels and incubated again with or without 10 ng/ml TGF- β_1 for another 3 days. **C:** Both control and experimental cells were preincubated with TGF- β_1 (10 ng/ml) for 3 days and then inoculated on collagen-coated plates or anchored or floating collagen gels for 3 days and treated either with vehicle (controls) or with TGF- β_1 . Western blotting and densitometry were used to assess α -SMA content and β -actin content in three independent samples. Data are expressed as a ratio of the blot densities of α -SMA to β -actin (mean ratio \pm SEM).

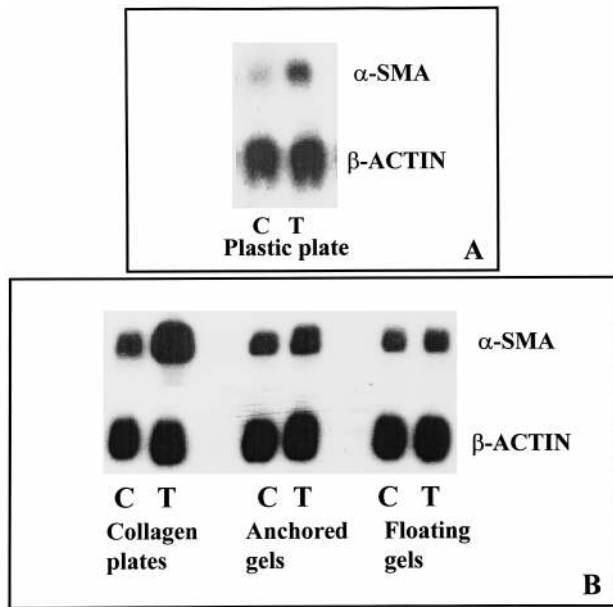


Figure 4. Northern blots of α -SMA and β -actin. Cells were prepared as in Figure 3. **A:** Fibroblasts were incubated with TGF- β_1 or vehicle on tissue culture plastic for 3 days, and mRNA levels were measured by Northern hybridization using oligonucleotide probes specific for either human α -SMA or β -actin. **B:** Cells were preincubated with either vehicle or TGF- β_1 as in **A** and then inoculated in collagen-coated plates, anchored collagen gels, or floating collagen gels and treated with or without TGF- β_1 for 3 days. Cells were assayed for α -SMA and β -actin mRNAs by Northern hybridization. Note the lack of up-regulation of α -SMA by TGF- β_1 in floating gels.

We also assessed whether TGF- β_1 increased α -SMA at the mRNA level. Cells were grown and treated exactly as described for protein analysis. Before plating on collagen substrates, α -SMA mRNA in cells treated with TGF- β_1 for 3 days on plastic plates was increased compared with untreated control cells (Figure 4A) whereas β -actin mRNA levels were unchanged. After a 3-day pretreatment with TGF- β_1 and then 3 days of treatment in collagen matrices, cells on collagen-coated plates exhibited large increases of α -SMA mRNA. Cells in anchored gels showed much smaller increases in TGF- β_1 -induced α -SMA mRNA, whereas cells in floating gels showed no detectable increase in response to TGF- β_1 treatment (Figure 4B). Thus, at both the protein and mRNA levels, the ability of TGF- β_1 to modulate α -SMA was dependent on the compliance of the substrate.

As α -SMA protein may have been either preferentially degraded or lost through transient membrane passages³³ in the different gel types, we measured loss of nascent α -SMA by metabolic labeling of cells for 24 hours on plastic plates with [³⁵S]methionine followed by incubation of cells in gels for up to 4 days. In untreated cultures, cells on collagen-coated plates as well as anchored and floating gels showed similar reductions of radiolabeled α -SMA between 1 and 2 days whereas floating gels showed significantly less decay at days 3 and 4 (Figure 5; $P < 0.01$). In TGF- β_1 -treated cultures, nascent α -SMA content was relatively constant in cells in collagen-coated plates whereas large losses occurred in cells in anchored and particularly floating gels.

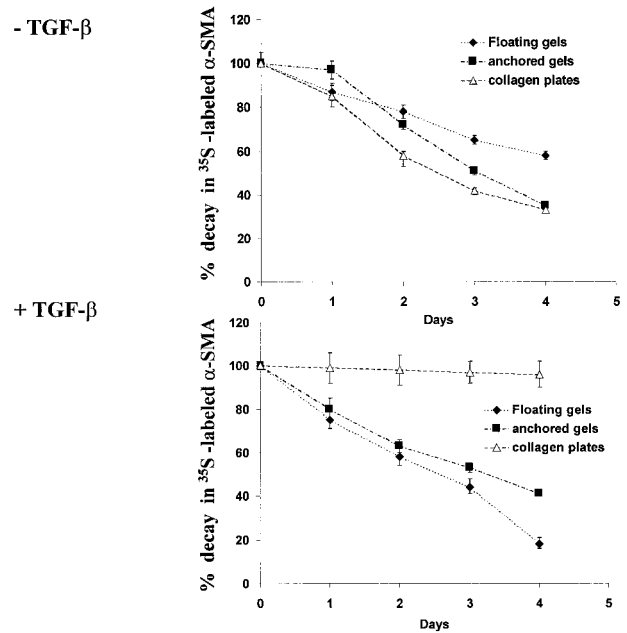


Figure 5. Loss of nascent α -SMA by metabolic labeling analyses. Cells were preincubated with vehicle or TGF- β and labeled for the last 24 hours on plastic plates with [³⁵S]methionine followed by inoculation of cells in different types of collagen gels for up to 4 days with or without TGF- β as indicated. Cells were released from gels and subjected to immunoprecipitation with anti- α -SMA antibody. The relative amounts of nascent α -SMA content were estimated from fluorographic blot densities and expressed as percentage of starting material (ie, day 0 of incubation in gels). Results are expressed as mean \pm SEM percent starting material.

TGF- β_1 Antibody Inhibition of α -Smooth Muscle Actin

Although TGF- β_1 can regulate the synthesis of PDGF in some cell types,³⁴ PDGF does not induce α -SMA expression in dermal fibroblasts³⁵ or in the gingival fibroblasts studied here (data not shown); consequently, it is unlikely that TGF- β_1 acted indirectly by inducing autocrine production of PDGF. To assess the specificity of TGF- β_1 on the up-regulation of α -SMA we blocked TGF- β_1 with a TGF- β neutralizing antibody that reacted with all TGF- β isoforms. Incubation of TGF- β_1 -treated cells on collagen-coated plates and anchored and floating matrices with the inhibiting antibody caused a large and significant reduction in α -SMA levels equivalent to that of untreated controls (Figure 6). These results demonstrated the specificity of TGF- β_1 in up-regulating α -SMA expression but also suggested a possible role for endogenous TGF- β_1 in the induction of α -SMA. Therefore, we measured the actual concentrations of active and total TGF- β_1 by ELISA in the culture conditions to which the cells were exposed. Addition of TGF- β_1 (10 ng/ml) apparently stimulated cells to synthesize TGF- β_1 for all three types of collagen substrates. More than half of the TGF- β_1 produced by cells was in an active form. The levels of active TGF- β_1 produced by cells in collagen-coated plates, anchored gels, and floating gels were 1.05 ± 0.3 ng/ml, 1.09 ± 0.23 ng/ml, and 1.06 ± 0.02 ng/ml, respectively, when exogenous TGF- β_1 (10 ng/ml) was added. In control cultures, the active form of TGF- β_1 was <0.005 ng/ml, and total

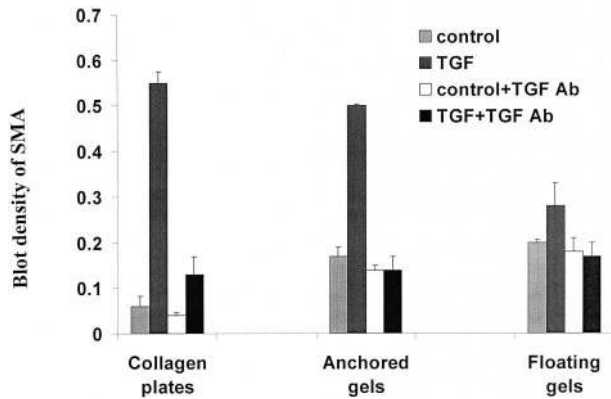


Figure 6. TGF- β antibody blockade of α -SMA induction. Cells were preincubated with vehicle or TGF- β_1 for 3 days on plastic plates as in Figure 3, inoculated into the three different types of collagen gels and incubated with PBS (control) or TGF- β_1 (TGF) with or without 15 μ g/ml anti-TGF- β neutralizing antibody (TGF Ab) for 3 days. Three independent samples were analyzed for the expression of α -SMA by Western blot. Results are expressed as mean density of α -SMA \pm SEM. Values are normalized to β -actin densities for each lane.

TGF- β_1 was 0.7 ng/ml (no significant difference between three different types of gels; $P > 0.2$). These data indicated that the type of gel did not significantly alter the endogenously produced levels of TGF- β_1 , and that cells were exposed to equivalent levels of TGF- β_1 for all types of substrates.

Integrin Content

TGF- β_1 increases $\alpha_2\beta_1$ integrin expression in fibroblastic³⁶ and osteogenic cells,¹⁶ and the $\alpha_2\beta_1$ collagen receptor is important for collagen gel contraction.^{12,13} We asked whether this increase was the result of TGF- β_1 -induced increase of integrin expression as has been reported earlier for certain fibroblastic cell lines cultured on plastic.³⁶ Flow cytometric analyses of the surface expression of α_2 and β_1 integrin subunits of untreated cells grown on collagen-coated plates, anchored gels, and floating gels showed mean staining intensities of 53.5 ± 10.7 , 61.7 ± 4.8 , and 154.9 ± 8.22 , respectively, for α_2 integrin subunits and 105.6 ± 9.8 , 108.6 ± 7.5 , and 121.3 ± 14.5 , respectively, for β_1 integrin subunits. As α_2 integrin extracellular domains are insufficient for collagen gel contraction, it has been suggested that the cytoplasmic domain of the α_2 integrin is involved in force generation required for contraction.¹² Consequently, we extended the flow cytometry results by immunoblotting whole-cell lysates for both α_2 and β_1 subunits in controls and TGF- β_1 -treated cultures. For controls, there was a twofold higher level of α_2 integrin content and a 20% higher β_1 content in cells in floating gels compared with cells grown on collagen-coated plates or anchored gels (Figure 7), consistent with surface expression analyses by flow cytometry. TGF- β_1 induced 1.5-fold and 2-fold increases in α_2 integrin content and 1.5-fold and 1.2-fold increases in β_1 content, respectively, for cells on collagen plates and anchored gels. TGF- β_1 did not induce a significant change of α_2 or β_1 integrin expression in floating gels (Figure 7). These results suggested to us that

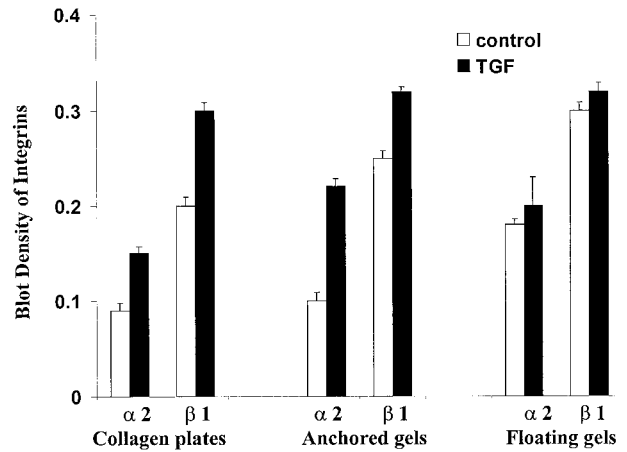


Figure 7. Integrin content of cells in collagen gels. Cells grown on tissue culture plastic were preincubated with vehicle or TGF- β and then inoculated into different types of collagen gels. Cells were incubated with or without TGF- β_1 for another 3 days. Three independent samples were analyzed by Western blot for α_2 -integrin or β_1 -integrin by Western blots. β -actin was used for normalization of protein loading for each lane.

tension is required for TGF- β_1 -induced increases of collagen receptor expression.

Loss of Tension Reverses TGF- β Induction of α -Smooth Muscle Actin

As intracellular tension was evidently required for TGF- β_1 -induced increase of α -SMA, we asked whether release of intracellular tension would abrogate the effect of TGF- β_1 on α -SMA. We used blocking antibodies to induce cell rounding and to release intracellular tension in cells that remained adherent to the collagen substrate. Cells were preincubated with the monoclonal antibodies P1E6 (which blocks the α_2 subunit¹⁴) and/or 4B4 (which blocks the β_1 subunit³⁷) and treated with TGF- β_1 (10 ng/ml) for 3 days. To assess whether the antibodies were relieving intracellular tension as expected, cells on collagen-coated plates were examined by phase contrast microscopy. The cells were initially well spread with or without TGF- β_1 , but after treatment with antibodies for 3 days they exhibited rounding (Figure 8, A–E) and loss of stress fibers. These alterations were most prominent after β_1 integrin antibody incubations and indicated that the anti-integrin antibodies greatly reduced intracellular tension. Quantitative analyses of α -SMA by Western blot demonstrated that preincubation with P1H5 alone reduced the level of α -SMA by 60% in TGF- β_1 cells on collagen-coated plates. Treatment with 4B4 alone decreased TGF- β_1 -induced increases of α -SMA to nearly that of untreated controls without antibody. Preincubation with both antibodies reduced the effect of TGF- β_1 to the level of controls (Figure 8F). For cells on anchored gels and floating gels, the anti-integrin antibodies also reduced the TGF- β_1 -induced increase of α -SMA to control levels, although the amount of reduction was less for anchored gels and much less for floating gels. These results were not caused by cell death as the viability of cells preincubated with antibodies and tested at the end

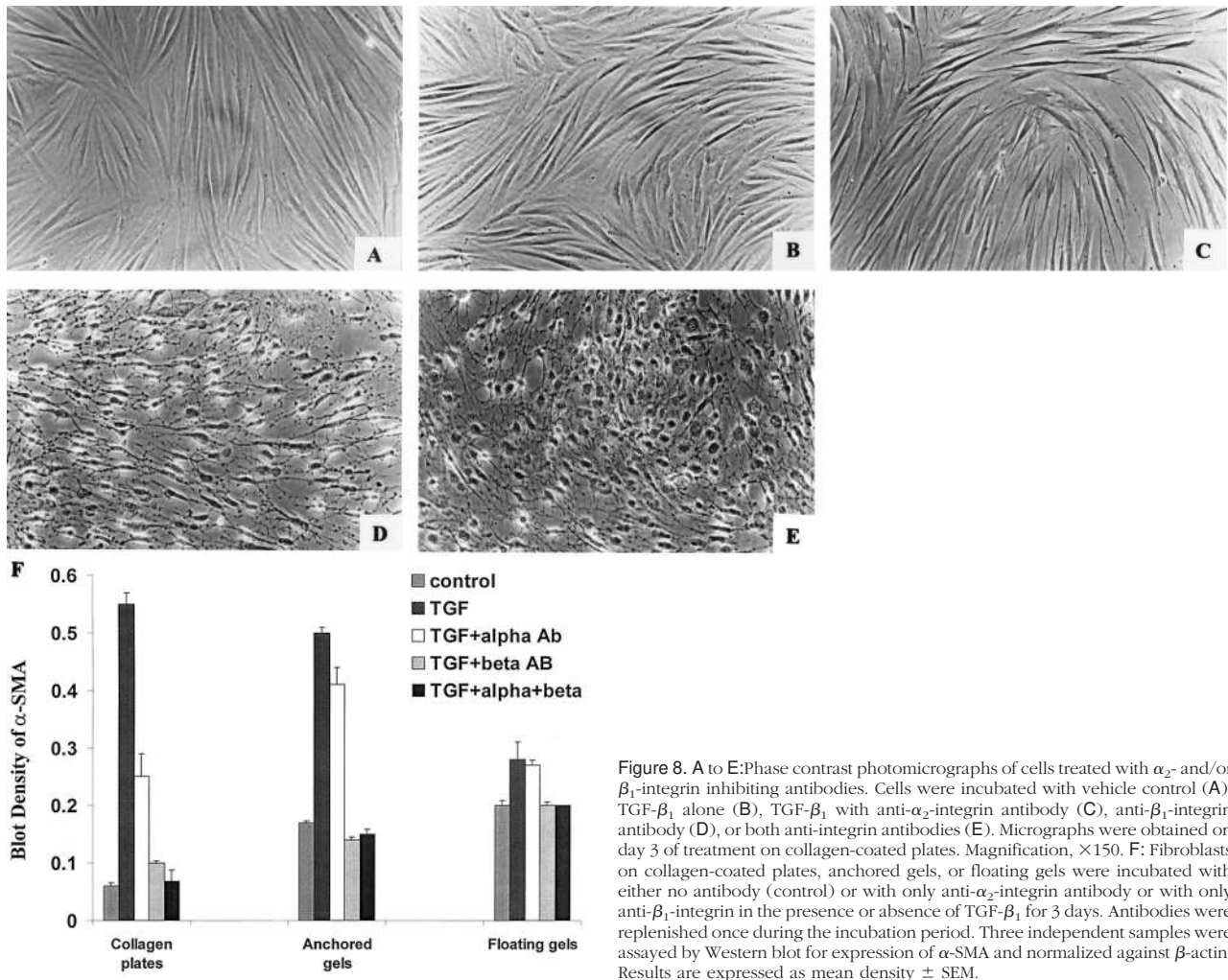


Figure 8. A to E: Phase contrast photomicrographs of cells treated with α_2 - and/or β_1 -integrin inhibiting antibodies. Cells were incubated with vehicle control (A), TGF- β_1 alone (B), TGF- β_1 with anti- α_2 -integrin antibody (C), anti- β_1 -integrin antibody (D), or both anti-integrin antibodies (E). Micrographs were obtained on day 3 of treatment on collagen-coated plates. Magnification, $\times 150$. F: Fibroblasts on collagen-coated plates, anchored gels, or floating gels were incubated with either no antibody (control) or with only anti- α_2 -integrin antibody or with only anti- β_1 -integrin in the presence or absence of TGF- β_1 for 3 days. Antibodies were replenished once during the incubation period. Three independent samples were assayed by Western blot for expression of α -SMA and normalized against β -actin. Results are expressed as mean density \pm SEM.

of the incubations (ie, 4 days) exceeded 95% as measured by trypan blue exclusion. These data indicate that TGF- β_1 requires the development of intracellular tension and $\alpha_2\beta_1$ integrins to induce increased α -SMA expression.

Discussion

The main finding of this study is that the ability of TGF- β_1 to increase α -SMA expression in fibroblasts is determined by the physical resistance of the substrate to cell-generated forces. This finding is consistent with previous *in vivo* data showing a temporal association between the time of appearance of myofibroblasts and the time of increased resistance to wound contraction.² Although the modulatory role of TGF- β_1 on myofibroblast differentiation and expression of α -SMA has been studied previously in monolayer cultures,^{21,27,38} there are no reports on cellular responses to TGF- β_1 under conditions of different tension levels. As healing tissues exhibit significant alterations in their biophysical properties over the time course of wound repair,⁹ we considered that the three collagen substrates used here may provide models for connective tissues at different stages of maturation.

For example, anchored gels resemble granulation tissue in later stages of remodeling with collagen fibers oriented parallel to fibroblasts whereas floating gels are more similar to resting dermis⁸ and to very early stages of wound formation when resistance to intracellular tension is low.⁹ Consistent with these proposed features, the three different types of collagen substrates showed large differences of compliance as measured by mechanical deformation studies, supporting the central notion of this study that the three collagen gel models exhibit large variations of compliance in response to cell-generated force. In turn, in the three culture systems used here, cells exhibited different levels of intracellular tension, as shown by the variation in the abundance and distribution of actin stress fibers in the different systems. Thus, cells in monolayer cultures on collagen-coated plastic developed high levels of intracellular tension, anchored gels developed moderate tension, and floating gels developed low tension. Indeed, fibroblasts grown on low-compliance substrates acquire characteristics of myofibroblasts that have high expression levels of α -SMA.³⁹

We designed experiments to separate the effects of TGF- β induction of α -SMA from the potential effects due to gel compliance. As shown in Figure 3, B and C, if cells

were treated with TGF- β , then compared with controls, the α -SMA content was greatly increased, particularly in collagen plates. This effect was not due simply to the compliance of the gels because if both controls and experimental cultures were both preincubated with TGF- β for 3 days and then inoculated into the different gel types (Figure 3C), the α -SMA content was again greatly increased in TGF- β -treated cultures. Thus, TGF- β not only induced α -SMA but also helped to maintain the cellular levels of this actin isoform. This point is emphasized by examination of [³⁵S]methionine labeling experiments that showed TGF- β -treated cultures on collagen plates had virtually no loss of labeled α -SMA over a 4-day time period if TGF- β was present in the culture medium. If, on the other hand, TGF- β was not present, then the α -SMA content reduced sharply from the time of labeling. Thus, TGF- β both induces and maintains α -SMA content in fibroblasts, particularly in cells on collagen plates in which intracellular tension is high.

Separate experiments showed that TGF- β_1 -induced increases of α -SMA were clearly dependent on the maintenance of intracellular tension during the experimental period. Cell rounding induced by integrin-inhibiting antibodies reversed the effect of TGF- β_1 on α -SMA content to levels of untreated controls for all three types of gel systems. Indeed, the increased decay rates of nascent, [³⁵S]methionine-labeled α -SMA in floating gels indicate that the reduced α -SMA of cells in floating gels is due in part to increased loss of α -SMA, possibly through membrane pores.^{33,40} However, the finding that TGF- β_1 did not increase mRNA for α -SMA in floating gels also indicates that cytokine-mediated induction of α -SMA transcription is regulated in part by intracellular tension. The marked increase of α -SMA mRNA content induced by TGF- β_1 in cells on low-compliance collagen gels is similar to previous observations for smooth muscle cells.⁴¹ To express α -SMA in response to TGF- β_1 , smooth muscle cells require the binding of an undescribed factor to a TGF- β_1 control element along with the binding of serum response factors to two CArG elements in the promoter region. Conceivably, the development of intracellular tension may serve to enhance transcription factor binding to the α -SMA promoter through the production and availability of mechanically coupled enzymes that in turn rely on well developed actin stress fibers traversing the cell. Indeed, mechanotranscriptional processes that regulate enzyme-substrate reactions are believed to be central elements of mechanically dependent transcription machinery⁴² and likely rely on the ability of the cell to generate intracellular tension through the formation of highly adherent attachments to the matrix. In this context, a recent paper by Gabbiani and co-workers⁴³ indicates that a specialized fibronectin isoform (ED-A) is required for the TGF- β induction of the myofibroblastic phenotype. This paper suggests a mechanism by which a specialized extracellular matrix molecule, under the control of TGF- β , is required for outside-inside signaling and possibly for the development of the intracellular tension necessary for α -SMA induction.

Flow cytometric and Western blot analyses of fibroblasts grown on collagen-coated plastic or anchored

gels showed a marked increase of the surface expression and total content of α_2 and β_1 subunits after TGF- β_1 treatment, consistent with earlier data of cells grown on plastic dishes.³⁶ In contrast, cells in floating gels showed only very small increases in protein levels for both subunits. Evidently, TGF- β_1 can increase $\alpha_2\beta_1$ expression, but this effect is dependent on the compliance of the gels. We also showed a central role for $\alpha_2\beta_1$ integrins in TGF- β_1 -induced regulation of α -SMA and maintenance of intracellular tension. These results further support the suggestion that generation of intracellular tension is a requirement for the regulation of α -SMA by TGF- β_1 . Collectively, this study supports the concept that in healing connective tissues, myofibroblastic differentiation is modulated locally by microenvironmental, interactive stimuli, including important interactions between wound-healing cytokines and mechanical forces. Conceivably, therapeutic strategies for enhancement of wound healing and reduction of scar formation could be developed using treatments that reduced generation of intracellular tension.

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