Activation of αvβ3 Integrin Alters Fibronectin Fibril Formation in Human Trabecular Meshwork Cells in a ROCK-Independent Manner

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PURPOSE. Fibronectin fibrillogenesis is an integrin-mediated process that may contribute to the pathogenesis of primary open-angle glaucoma (POAG). Here, we examined the effects of αvβ3 integrins on fibrillogenesis in immortalized TM-1 cells and human trabecular meshwork (HTM) cells.

METHODS. TM-1 cells overexpressing wild-type β3 (WTβ3) or constitutively active β3 (CAβ3) integrin subunits were generated. Control cells were transduced with an empty vector (EV). Deoxycholic acid (DOC) extraction of monolayers, immunofluorescence microscopy, and Oncell western analyses were used to determine levels of fibronectin fibrillogenesis and fibronectin fibril composition (EDA+ and EDB+ fibronectins) and conformation. αvβ3 and α5β1 Integrin levels were determined using fluorescence-activated cell sorting (FACS). Cilengitide and an adenovirus vector expressing WTβ3 or CAβ3 integrin subunits were used to examine the role of αvβ3 integrin in HTM cells. The role of the canonical α5β1 integrin-mediated pathway in fibrillogenesis was determined using the fibronectin-binding peptide FUD, the β1 integrin function-blocking antibody 13, and the Rho kinase (ROCK) inhibitor Y27632.

RESULTS. Activation of αvβ3 integrin enhanced the assembly of fibronectin into DOC-insoluble fibrils in both TM-1 and HTM cells. The formation of fibronectin fibrils was dependent on α5β1 integrin and could be inhibited by FUD. However, fibrillogenesis was unaffected by Y27632. Fibrils assembled by CAβ3 cells also contained high levels of EDA+ and EDB+ fibronectin and fibronectin that was stretched.

CONCLUSIONS. αvβ3 Integrin signaling altered the deposition and structure of fibronectin fibrils using a β1 integrin/ROCK-independent mechanism. Thus, αvβ3 integrins could play a significant role in altering the function of fibronectin matrices in POAG.

Keywords: fibronectin, trabecular meshwork, glaucoma, integrins

Primary open-angle glaucoma (POAG) is the most common type of glaucoma in the United States, and a major risk factor for POAG is an elevation in intraocular pressure (IOP). There is increasing evidence that higher than normal resistance to aqueous humor outflow from the trabecular meshwork (TM) is responsible for this elevation in IOP. Recent studies suggest that excess deposition of extracellular matrix (ECM) proteins, especially fibronectin, could be a key factor in the development of restricted aqueous humor outflow from the anterior segment. In addition, both POAG and steroid-induced glaucoma (SIG) are associated with an increased accumulation of various ECM proteins, including fibronectin.

This excess deposition of fibronectin has been attributed to elevated levels of TGFβ2 in aqueous humor or in response to treatment with glucocorticoids such as dexamethasone. Recent studies suggest that the glucocorticoid- and TGFβ2-induced increases in fibronectin may be related since higher levels of TGFβ2 follow glucocorticoid treatments. TGFβ1- and TGFβ2-induced increases in IOP in a rat model have also been shown to correlate with increased fibronectin labeling within the TM.

Fibronectin fibrils are a major component of the ECM in the TM and recent in vitro studies have shown that TM cells produce both the EDA+ and EDB+ fibronectin isoforms. Fibronectin is one of the earliest ECM fibrils to be assembled in vivo, and it has been found to act as a scaffold upon which other ECM protein matrices can be assembled. Within the TM it appears to be involved in the assembly of nascent matrices of type IV collagen, laminin, and fibrillin into the ECM. Fibronectin fibrils also act as bioreservoirs for growth factors such as TGFβ2 and enzymes like LOX1, both of which have been implicated in glaucoma. The EDA domain of fibronectin has been implicated in the formation of myofibroblasts and fibrosis and is a factor that may play a role in POAG and SIG. Fibronectin fibrils therefore would be expected to play a critical role in maintaining ECM homeostasis in the normal TM and in the development of POAG and SIG.

Unlike some ECM proteins such as type I collagen, the incorporation of fibronectin into fibrils is a multistep process...
that involves integrin signaling and the contractile properties of the tissue. Fibronecin fibril formation is initiated when the secreted soluble protein dimer binds to cell surface integrins. Once bound, the fibronecin dimer is unfolded and stretched, resulting in a conformational change that exposes specific fibronecin–fibronecin binding sites within the molecule that promote the assembly of a larger, insoluble fibril. The unfolding and stretching of the soluble fibronecin dimers is believed to be mediated by the guanosine triphosphatase (GTPase) RhoA and the contractile properties of the actomyosin network.

Integrins are a family of cell surface receptors consisting of a noncovalently bound heterodimer of α and β subunits. At least six fibronecin binding integrins have been identified in the TM, including α5β1, which is the major integrin that mediates fibronecin fibril assembly. Several other integrins, however, are capable of initiating fibril assembly including αβ3 integrin. Expression of the α5, αv, and β3 integrin subunits in TM cells is affected by glucocorticoids, and studies from our lab have shown that the overall levels and activity of αβ3 integrin in human TM (HTM) cells are increased in response to glucocorticoids such as dexamethasone. Although α5β1 and αβ3 integrin expression is affected by TGFβ1 in other cell types, to date only one study has reported that TM cells increased expression of αv and β3 integrin subunits in response to TGFβ2.

Given the important role that fibronecin appears to play in regulating the assembly of the ECM within the TM, we sought to examine the relationship between αβ3 integrin signaling and fibronecin matrix assembly using an immortalized TM cell line, TM-1, previously engineered to stably overexpress either a wild-type (WTβ3) or constitutively active (CAβ3) αβ3 integrin, and HTM cells expressing an activated αβ3 integrin. We specifically looked to see if the activated conformation of αβ3 integrin was involved. Our previous studies have suggested that this conformation caused HTM cells to exhibit phenotypes associated with POAG and SIG, including actin cytoskeletal rearrangements into cross-linked actin networks (CLANs) and inhibition of phagocytosis. We report here that this activated conformation of αβ3 integrin also increases fibronecin fibril assembly in TM cells. This increase in fibrils includes multiple fibronecin isoforms, a stretched conformation of fibronecin, and appears to be independent of Rho kinase (ROCK) activity.

**Materials and Methods**

**Materials**

Rabbit polyclonal fibronecin antiserum was produced in our lab and validated by ELISA analysis and immunofluorescence microscopy. Normal rabbit serum was purchased from Vector Laboratories (cat. #S-5000; Burlingame, CA, USA). The mouse anti-EDA+ fibronecin clone IST-9 (cat. #ab65328), mouse anti-EDB+ fibronecin clone BC-1 (cat. #ab154210), mouse anti-β3 integrin clone CRC54 (#ab34409), mouse anti-β5 integrin clone SNAKA51 (cat. #MABT201) were purchased from EMD Millipore. Recombinant FUD (functional upstream domain) peptide, derived from the Streptococcus pyogenes F1 adhesin protein, was expressed and prepared as previously described.

**Adenovirus 5 (Ad5) WTβ3-mCherry/CAβ3-mCherry Construction**

The wild-type cDNA for the human β3 integrin subunit was obtained from Thermo Fisher Scientific and cloned into the pLVX-IRES-Puro vector (Takara Bio USA, Mountain View, CA, USA) as previously described. A DNA fragment containing a Kozak sequence was then cloned onto the amino terminus of the β3 integrin cDNA along with an mCherry tag at the carboxyl terminus. This WTβ3 integrin-mCherry transgene was then cloned into the Xho1/Xba1 site of the pacAd5CMV/mcsSV40 shuttle vector (Ad5-WTβ3). Site-directed mutagenesis was used to create the pacAd5CMV-β3 integrin T562N-mCherry-SV40PA vector (Ad5-CAβ3). Cloning and site-directed mutagenesis of the Ad5-WTβ3-mCherry and Ad5-CAβ3-mCherry vectors were done by GenScript (Piscataway, NJ, USA) and validated by cDNA sequencing. The engineered vectors, along with the pacAd5CMV/mcsSV40PAAd5 empty vector (Ad5-EV), were each packaged at the University of Iowa Viral Vector Core.

**Cell Culture**

Immortalized TM-1 cells overexpressing either a wild-type β3 integrin subunit (WTβ3) or a constitutively active β3 T562N integrin subunit (CAβ3) were generated as previously described. A cell line transfected with an empty vector (EV) was used as a control. All TM-1–derived cell lines were cultured in routine growth medium consisting of low-glucose Dulbecco’s modified Eagle’s medium (DMEM) (Sigma Aldrich Corp.), 10% fetal bovine serum (FBS) (Atlanta Biologicals, Atlanta, GA, USA), 2 mM L-glutamine (Sigma Aldrich Corp.), 0.2% Promicin (InvivoGen, San Diego, CA, USA), and 0.05% gentamicin (Mediatech, Manassas, VA, USA). Cells were kept under selection in 2 μg/mL puromycin. The N27TM-6 strain of normal HTM cells was isolated from a 27-year-old female donor and characterized as previously described. HTM cells were routinely grown in the same growth medium used for TM-1 cell lines except for the use of 15% FBS and 1 ng/mL FGF-2 (PeproTech, Rocky Hill, NJ, USA).

In experiments in which HTM cells were treated with or without cilengitide (CGT) or dexamethasone (DEX) to activate the αβ3 integrin, HTM cells were plated at a density of 3 × 10^4 cells/well in growth medium into 96-well plates. Upon reaching confluence, cells were fed daily with growth medium for 7 days. Cells were then switched to low serum (1% FBS) and treated for 12 to 14 days with control medium, medium plus 0.1% ethanol (vehicle), or 500 nM DEX or medium containing 50, 100, or 200 μM CGT plus either vehicle or DEX. At the end of the treatment period cells were processed for On-cell western (OCW) analysis as described below.

For experiments in which HTM cells were transduced with Ad5 viral vectors expressing mCherry-β3 integrin transgenes, cells were plated at 4 × 10^5 cells/well in normal growth medium in 24-well plates. Just prior to reaching confluence, cells were transduced with either Ad5-EV, Ad5-WTβ3 integrin-mCherry, or Ad5-β3 T562N integrin-mCherry for 4 hours at a multiplicity of infection (MOI) of 100. Twenty-four hours post
transduction, cells were refeed with normal growth medium. Upon reaching confluence, cells were refeed daily with normal growth medium for 7 days. Cells were then refeed with 10% FBS-containing medium for 48 hours followed by 1% FBS-containing medium for another 48 hours prior to processing for OCW analysis as described below or immunofluorescence microscopy.

**Fluorescence-Activated Cell Sorting (FACS) Analysis**
Cells in growth medium were detached from plates using Cell Dissociation Buffer (Sigma Aldrich Corp.) and blocked in PBS plus 5% BSA on ice. They were then incubated with IgG only, mAb LM609, mAb PID6, or mAb 12G10 at 5 μg/mL for 1 hour on ice. Cells were washed and labeled with Alexa Fluor 488 goat anti-mouse IgG (cat. #A11029; Thermofisher Fisher Scientific) diluted 1:4000 for 30 minutes on ice. The cells were then fixed with PBS + 1% paraformaldehyde (PFA). Data were collected using either a FACS Calibur (Becton Dickinson, Franklin Lakes, NJ, USA) or an Attune NxT (Thermo Fisher Scientific) flow cytometer and analyzed using FlowJo software (FlowJo, LLC, Ashland, OR, USA). The FlowJo software was used to calculate the normal geometric mean fluorescence (NGMF).

**On-Cell Western Analysis (OCW)**
TM-1-derived cell lines were plated at a density of 5 × 10⁴ cells/well in low-serum (1% FBS) medium into 96-well plates for 3 to 4 hours to allow the cells to attach and spread. The cells were then treated with or without 500 nM FUD, 10 μg/mL mAb 13 or control rat IgG, or 5 μg/mL Y27632 for 24 hours. To determine the level of deoxycholic acid (DOC)-insoluble fibronectin fibrils, cell layers were extracted with 1% DOC-containing buffer and processed for OCW analysis as previously described.12 To determine the level of fibronectin in either the unextracted or DOC-extracted wells, cell layers or DOC-extracted matrices in the wells were fixed with 4% PFA and labeled with anti-fibronectin antibodies followed by an IRDye 800CW-conjugated secondary antibody. The fibronectin-labeled wells were blanked against wells labeled with normal rabbit serum or an irrelevant mouse mAb. The level of fibronectin labeling was normalized to the total protein content/well as determined by labeling with IRDye 680RD NHS ester. Quantification of the signals at each wavelength was performed using Li-COR Image Studio v. 5.0.21 software (Li-Cor Biosciences, Lincoln, NE, USA). Except where noted in the legend, the results represent the mean ± SE of data pooled from two to six experiments depending upon the exact assay.

**MTT Cell Viability Assay**
Cells were plated into 96-well plates and allowed to attach for 3 to 4 hours. Cells were refeed with media containing 1% FBS with or without 500 nM or 2 μM FUD for 24 hours. As a control, some cells were treated for 24 hours with 0.1% saponin. Viability was determined using a CellQuanti-MTT assay kit (BioAssay Systems, Hayward, CA, USA) as previously described.12 The results represent the mean ± SE of data pooled from two to six experiments depending upon the exact assay.

**Immunofluorescence Microscopy**
Cells were fixed with 4% PFA in PBS at room temperature for 20 minutes and blocked in PBS plus 1% BSA prior to labeling with the various primary antibodies. For immunolabeling studies, cells were labeled with primary antibodies diluted in block. Primary antibodies were detected with Alexa Fluor 488 goat anti-rabbit IgG (cat. #A11034), Alexa Fluor 546 goat anti-mouse IgG (cat. #A1050), Alexa Fluor 546 goat anti-rabbit IgG (cat. #A11035), or Alexa Fluor 488 goat anti-mouse IgG. All secondary antibodies were diluted in block. Alexa Fluor secondary antibodies were purchased from Thermo Fisher Scientific. Nuclei were labeled using Hoechst 33342 (cat. #H1399, Thermo Fisher Scientific). Fluorescence was observed using a Zeiss Imager.Z2 epifluorescence microscope (Carl Zeiss AG, Oberkochen, Germany) equipped with a digital camera (AxioCam 702 mono) and image acquisition software (Zen v. 2.3). All experiments were independently performed two to four times. For the experiments in which cells were treated with or without the ROCK inhibitor Y27632, Z-stacks were acquired from three fields of view/coverslip. Using the image acquisition software, these Z-stacks were then deconvoluted and reconstructed into three-dimensional images or compressed into single images. Cells transduced with Ad5-viral vectors were not labeled with primary or secondary antibodies prior to observation using fluorescence microscopy.

**Quantitative PCR**
RNA from confluent cultures was isolated using the QIAshredder and RNeasy Plus Mini Kits (Qiagen, Inc., Valencia, CA, USA) and cDNA was generated using the High Capacity cDNA Reverse Transcription Kit (Applied Biosystems, Foster City, CA, USA) according to the manufacturers’ instructions. Quantitative (q)PCR was performed using a QuantStudio 7 Flex Real-Time PCR system and SYBR Green PCR Master Mix (both Thermo Fisher Scientific). Results were normalized to the housekeeping gene succinate dehydrogenase (SDHA). The primers (Integrated DNA Technologies, Coralville, IA, USA) used for the qPCR reactions are listed in the Table.

**G-LISA RhoA Activation Assay**
Confluent cultures of EV or CAβ3 cells were serum starved for ~16 hours prior to being left untreated or stimulated with 10% FBS for 15 or 30 minutes, respectively. Cells were then processed for analysis using the RhoA G-LISA activation assay kit (cat. #BK124; Cytoskeleton, Denver, CO, USA) according to the manufacturer’s instructions. Briefly, cell lysates were incubated for 30 minutes at 4°C in microtiter wells that had been precoated with the Rho binding domain from a Rho effector protein, rhotekin, that specifically binds active RhoA. Wells were washed and incubated for 45 minutes at room temperature with a mouse anti-RhoA antibody. Wells were washed again and incubated with horseradish peroxidase (HRP)-conjugated goat anti-mouse IgG secondary antibody. The plates were then washed and the level of label in each well was detected using a HRP detection reagent provided by the manufacturer for 10 to 15 minutes at 37°C. The reaction was stopped with HRP stop buffer. Wells were read at 490 nm to determine the content of GTP-bound RhoA present in each well.

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**TABLE. Primes Used for Real-Time RT-PCR**

<table>
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<tr>
<th>Type</th>
<th>Sequence</th>
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<tr>
<td>Human fibronectin</td>
<td>Reverse: GCCAGTGCAAGCATACACAGTG</td>
</tr>
<tr>
<td>Human EDA-fibronectin</td>
<td>Reverse: GTCAACCCTGACCTGGAAACTTG</td>
</tr>
<tr>
<td>Human EDB-fibronectin</td>
<td>Reverse: CGTGGACCACCGCTAAACCTC</td>
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Statistical Analysis

Statistical analysis was performed using ANOVA. Where pairs of treatment groups were compared, ANOVA analysis was used in conjunction with Tukey’s honestly significant difference test.

RESULTS

Effect of Activation of \(\beta3\) Integrin Signaling on Fibronectin Matrix Assembly in Immortalized TM Cells

To determine if \(\alpha v \beta 3\) integrin signaling influenced fibronectin fibrillogenesis, confluent monolayers of the EV cell line or cell lines overexpressing the WT\(\beta 3\) or CA\(\beta 3\) integrin subunits were fixed and labeled for fibronectin (Fig. 1A). By immunofluorescence microscopy, no obvious differences in labeling patterns or intensity were observed in the two cell lines overexpressing the \(\beta 3\) integrin subunits compared to control EV cells. Quantification of the cell surface-labeled fibronectin fibrils in intact confluent monolayers of WT\(\beta 3\) and CA\(\beta 3\) cells showed a 1.7-fold increase (\(P < 0.002\)) in fibronectin in cells overexpressing a CA\(\beta 3\) integrin subunit compared to EV cells (Fig. 1B). CA\(\beta 3\) cultures also expressed a 1.4-fold increase (\(P < 0.02\)) in fibronectin compared to WT\(\beta 3\) cultures, suggesting that the activation state of the integrin, rather than just overexpression, was affecting levels of fibronectin in the intact cell layer.

In order to differentiate between cell surface-bound soluble fibronectin and fibronectin assembled into an insoluble fibril, cell cultures were extracted with DOC. The DOC extraction
procedure removes soluble, cell surface-bound and intracellular fibronectin but not fibronectin assembled into insoluble fibrils. It is these insoluble fibrils that are the functional form of fibronectin that serve to support a variety of cell behaviors under both normal and pathological conditions in vivo. As shown in Figure 1C, DOC-extracted monolayers immunolabeled for fibronectin showed a clear increase in fibronectin matrices formed by CAβ3 cultures compared to both EV and WTβ3 cultures (Fig. 1C). OCW analysis of the DOC-extracted monolayers (Fig. 1D) confirmed the immunofluorescence microscopy results and showed a ~2-fold increase (P < 0.01) in DOC-insoluble fibrils in CAβ3 cultures compared to EV cultures. When compared to WTβ3 cultures, the CAβ3 cultures showed a ~3-fold increase in DOC-insoluble fibrils (P < 0.0002). Thus the increase in insoluble fibrils was not due to the mere overexpression of the β3 integrin subunit since a similar increase was not observed in the levels of insoluble fibrils expressed in the WTβ3 cultures compared to the EV cultures.

The increase in fibronectin fibrils was also not due to changes in the level of fibronectin expression, since Western blot analysis of whole cell lysates (Fig. 1E) did not show any consistent differences in fibronectin levels between the three cell lines. Interestingly, both WTβ3 and CAβ3 cells demonstrated a 1.5-fold increase in fibronectin mRNA levels (P < 0.01) compared to EV cells (Supplementary Fig. S1). However, there was no difference between fibronectin mRNA levels in WTβ3 and CAβ3 cells, indicating that differences in mRNA levels were not a factor in the observed differences in fibronectin fibril levels. In order to determine if differences in fibronectin isoform expression were a factor in this phenomenon, we also analyzed the mRNA levels of the EDA− and EDB+ isoforms of fibronectin. As shown in Supplementary Figure S1, although the EDA+ fibronectin isoform appeared to be expressed at higher levels in both WTβ3 and CAβ3 cultures compared to EV cultures, the differences were not statistically significant. EDB+ fibronectin mRNA levels were comparable between the three cell lines.

We then wanted to verify that the WTβ3 and CAβ3 cells expressed similar levels of αvβ3 integrins on their cell surface given the differences in the levels of their assembled fibronectin matrices. FACS analysis using the mAb LM609 to αvβ3 integrin (Supplementary Fig. S2) confirmed our earlier study and showed that EV cells express very low levels of αvβ3 integrin on the cell surface while WTβ3 and CAβ3 cells showed significantly higher levels of αvβ3 integrin. The cell surface levels of αvβ3 integrin in WTβ3 and CAβ3 cells, however, were comparable.

**Effect of β3 Integrin Activation on Fibronectin Matrix Assembly in Normal HTM Cells**

We next sought to confirm that activation of αvβ3 integrin signaling increased fibronectin matrix assembly in primary HTM cells. For this, HTM cells were cultured in the presence of DEX, which we previously demonstrated led to the activation of αvβ3 integrin and increased fibronectin matrix assembly, with or without increasing concentrations of the cyclic RGD peptide CGT, which has been reported to activate αvβ3 signaling when used at high concentrations. As shown in Figure 2A, HTM cells treated with vehicle together with 100 μM CGT demonstrated 2.7-fold more DOC-insoluble fibronectin matrix (P < 0.005) compared to cells treated with vehicle alone. Higher levels of DOC-insoluble matrix were also observed in the presence of 50 and 200 μM CGT; however, these levels were not significantly greater than with vehicle alone. In contrast, cells treated with DEX and all three CGT concentrations demonstrated significantly more DOC-insoluble fibronectin matrix (P < 0.01) compared to cells treated with DEX alone. Consistent with our earlier study, DEX treatment by itself significantly increased DOC-insoluble fibronectin matrix compared to vehicle-treated cells (P < 0.002).

In a second set of experiments, HTM cells were transduced with Ad5 viral vectors expressing either WTβ3 or CAβ3 integrin-mCherry transgenes (Figs. 2B, 2C). An empty Ad5 viral vector (EV) was used as a control. Twelve days post transduction, OCW analysis showed that HTM cells transduced with Ad5-CAβ3 assembled 1.35-fold more DOC-insoluble fibronectin matrix compared to control cells transduced with Ad5-EV (P < 0.003). Ad5-CAβ3–transduced cells also assembled 1.2-fold more DOC-insoluble fibronectin matrix (P < 0.03) than cells transduced with Ad5-WTβ3. Together these data confirmed our findings using immortalized TM-1–derived cell lines that overexpressing an activated αvβ3 integrin enhances fibronectin fibril formation.

**Effect of Overexpressing αvβ3 Integrins in TM Cells on α5β1 Integrin-Mediated Fibronectin Fibril Assembly**

Because α5β1 integrin plays a significant role in fibronectin fibrillogenesis in most cells, we also wanted to determine if α5β1 integrin levels were different in the three cell lines. FACS analysis using mAb P1D6 against the α5 integrin subunit (Fig. 3A) or mAb 12G10 against the β1 subunit (Fig. 3B) indicated that there were minor (≤1.2-fold) but statistically significant differences (P < 0.001) in α5 and β1 integrin subunit levels between the three cell lines. These differences did not, however, correlate with the observed differences in levels of fibronectin matrices formed by the three cell lines.

We next wanted to determine if the insoluble fibronectin fibrils formed in CAβ3 cultures involved the canonical α5β1 integrin–mediated processes commonly thought to regulate fibril formation (Fig. 4A). In particular, we wanted to determine if antibodies to α5β1 integrin disrupted binding of soluble fibronectin to the cell surface and whether these fibrils were sensitive to disruption by the peptide FUD, which was previously found to disrupt fibronectin fibrillogenesis in HTM cultures. For this, we treated EV, WTβ3, and CAβ3 cultures for 24 hours with either the β1 integrin function-blocking mAb 13, previously shown to block the early cell surface binding step in fibronectin matrix assembly, or FUD. All three cell lines were then processed for OCW analysis and immunofluorescence microscopy.

In intact, unextracted cell layers (Fig. 4B), mAb 13 behaved as expected and treatment significantly decreased cell surface binding of fibronectin in all three cell lines. Decreases of 54%, 53%, and 32% (all P < 0.01) were seen in EV, WTβ3, and CAβ3 cell lines, respectively. Immunofluorescence microscopy verified the OCW analysis. As shown in Figure 4C, confluent monolayers of untreated EV, WTβ3, and CAβ3 cells form a well-developed fibronectin matrix. However, a 24-hour exposure to mAb 13 caused a marked reduction in fibronectin levels in all three cell lines, consistent with previous studies. Thus, all three cell lines were using the α5β1 integrin to bind fibronectin and mediate the early cell surface steps in fibril formation. With respect to DOC-insoluble fibronectin fibrils (Fig. 4E), mAb 13 had little or no effect on the incorporation of the remaining cell surface–bound fibronectin into the insoluble ECM in all three cell lines.

Responses to FUD treatment in intact monolayers were similar. In both WTβ3 and CAβ3 cultures, FUD caused significant decreases of 39% and 42% in the level of fibronectin in intact cell layers (both P < 0.01), respectively, compared to untreated WTβ3 and CAβ3 cultures (Fig. 4B). In FUD-treated EV cultures the 25% reduction in fibrils was not quite
statistically significant ($P = 0.08$) compared to untreated EV cultures. These results were confirmed by immunofluorescence microscopy (Fig. 4D), which showed that treatment with FUD for 24 hours clearly reduced fibril formation in confluent monolayers of EV, WTβ3, and CAβ3. These results are consistent with our earlier study. That earlier study also confirmed that the effects of the FUD were specific, as a mutated version of the peptide had little or no effect on fibronectin fibril assembly by TM-1 cells.

FUD also appeared to have an inhibitory effect on insoluble fibronectin fibril formation in EV and WTβ3 cultures (54% and 23%, respectively), but neither of these decreases was
adhesions (Fig. 4A). As shown in Figure 5, all three cell types had strong fibronectin matrix assembly called fibrillar adhesions and focal adhesions in TM cells. We next sought to determine if activated αvβ3 integrins also localized to fibrillar adhesions in CAβ3 cells where fibronectin fibrillogenesis was occurring and whether this could explain the enhanced fibronectin fibril assembly observed in these cells. For this we used the mAb CRC54, which we previously used to detect activated αvβ3 integrins in TM cells. Consistent with our earlier studies, little or no activated αvβ3 integrin was detected in EV cells. Neither focal adhesions (sites of cell attachment) nor fibrillar adhesions in EV cells contained activated αvβ3 integrin, although there were abundant fibronectin fibrils present (Figs. 6A–C). We also did not see any activated αvβ3 integrins in fibrillar adhesions in either WTβ3 or CAβ3 cells. However, activated αvβ3 integrins were found in focal adhesion complexes in both WTβ3 and CAβ3 cultures (Figs. 6D, 6G). Not surprisingly, these αvβ3 integrin-positive focal adhesion complexes were more prominent in CAβ3 cells compared to WTβ3 cells. In both cell lines, however, few if any of the αvβ3-integrin–positive focal adhesion complexes colocalized with fibronectin fibrils (Figs. 6E–6I). Thus, activated αvβ3 integrins were not involved in the sites of fibronectin fibril assembly. CRC54-positive structures were confirmed to be focal adhesions by double-labeling cultures of all three cell lines with phalloidin, to visualize actin stress fibers and either mAb LM609 (Supplementary Fig. S4) or mAb CRC54 (Supplementary Fig. S5).

**Effect of Inhibiting ROCK Activity on the Incorporation of Fibronection Fibrils Into the DOC-Insoluble Matrix**

We next sought to determine whether or not RhoA was involved in the enhanced incorporation of fibronectin into the DOC-insoluble fibrils exhibited by CAβ3 cells, since it is well established that Rho GTPase signaling plays a role in regulating fibronectin fibrillogenesis in TM cells and other cell types. We first compared the level of RhoA activity in EV cells and CAβ3 cells in response to serum stimulation (Fig. 7). Since EV and WTβ3 cells behaved the same in earlier
FIGURE 4. Both mAb 13 and FUD inhibit fibronectin fibril formation. (A) Schematic showing the canonical $\alpha_5\beta_1$ integrin-mediated pathway involved in fibronectin fibrillogenesis.\textsuperscript{58} As previously reported,\textsuperscript{46} both $\alpha_5\beta_1$ and $\alpha_v\beta_3$ integrins can colocalize to focal adhesions in TM cells. During the initial stages of fibril formation, soluble fibronectin dimers bound to $\alpha_5\beta_1$ integrins are unfolded within focal adhesions together with $\alpha_v\beta_3$ integrins. The fibronectin–$\alpha_5\beta_1$ integrin complex, but not $\alpha_v\beta_3$ integrin, is then translocated to regions of fibril formation called fibrillar adhesions.\textsuperscript{58} While in fibrillar adhesions, fibronectin-fibronectin binding interactions promote the assembly of DOC-insoluble fibrils. Both the $\beta_1$ integrin function-blocking mAb 13 and FUD inhibit this process as indicated in the schematic. (B) OCW analysis of unextracted confluent monolayers of EV, WT$\beta_3$, and CA$\beta_3$ cells. Cells were left untreated (NT) or treated for 24 hours with 500 nM FUD, 10 $\mu$g/mL mAb 13, or control IgG. Monoclonal Ab 13 significantly reduced fibronectin fibril formation in all three cell lines ($P < 0.01$) while FUD significantly reduced fibril formation in the WT$\beta_3$ and CA$\beta_3$ cells, respectively ($P < 0.01$). The $\sim 25\%$ reduction in response to FUD seen in the EV cultures was not statistically significant ($P = 0.08$). Data are pooled from two separate assays with triplicate determinations ($n = 6$) and represent the mean ± SE. (C) Immunofluorescence images of intact, unextracted monolayers of EV, WT$\beta_3$, and CA$\beta_3$ cells that were left untreated or treated with 10 $\mu$g/mL mAb 13 for 24 hours prior to labeling with rabbit fibronectin antiserum. Monoclonal Ab 13 disrupted fibronectin fibril formation in all three cell lines. The labeling is a combination of both soluble and insoluble fibronectin fibrils. Scale bar: 50 $\mu$m. (D) Immunofluorescence images of intact, unextracted confluent monolayers of EV, WT$\beta_3$, and CA$\beta_3$ cells that were left untreated or treated with 10 $\mu$g/mL mAb 13 for 24 hours prior to labeling with rabbit fibronectin antiserum. FUD disrupted fibronectin fibril formation in all three cell lines. The labeling is a combination of both soluble and insoluble fibronectin fibrils. Scale bar: 50 $\mu$m. (E) OCW analysis of DOC-extracted monolayers of EV, WT$\beta_3$, and CA$\beta_3$ cells. Cells were treated as in (B) prior to DOC extraction; mAb 13 did not have an effect on DOC-insoluble fibronectin levels in any of the three cell lines. FUD only statistically significantly decreased DOC-insoluble fibronectin levels in the CA$\beta_3$ cell cultures relative to both untreated cells and cells treated with mAb 13 (both $P < 0.01$). The minor decreases in the insoluble fibronectin matrices seen in both EV and WT$\beta_3$ cells were not statistically significant. Comparing untreated groups, CA$\beta_3$ cells exhibited significantly more DOC-insoluble fibronectin matrix than either EV or WT$\beta_3$ cells, respectively ($P < 0.01$). Data are pooled from two separate assays with triplicate determinations ($n = 6$) and represent the mean ± SE.
Experiments, we used only EV cells in this set of experiments. EV cells exhibited a weak, transient, and statistically insignificant ($P = 0.28$) 23% increase in RhoA activity in response to serum that peaked at 15 minutes post stimulation. CA$\beta$3 cells also exhibited a weak increase in RhoA activity 15 minutes post serum stimulation; however, by 30 minutes, RhoA activity had significantly increased 37% over control levels ($P < 0.02$) and 30% over the levels seen at 15 minutes ($P < 0.045$).

Given that CA$\beta$3 cells appeared to exhibit a higher level of RhoA activity compared to EV cells, we then examined whether or not the enhanced fibronectin fibrillogenesis in CA$\beta$3 cells was mediated by RhoA signaling. For this study, we treated confluent monolayers of EV, WT$\beta$3, and CA$\beta$3 cells with or without the Y27632 inhibitor, which targets the downstream effector of RhoA called ROCK. Phase microscopy of inhibitor-treated cells (Supplementary Fig. 8G) showed that all three cell lines responded to Y27632 and exhibited retraction and/or cell rounding compared to untreated cells. The enhanced level of fibronectin in the insoluble matrix of CA$\beta$3 cultures, however, was unaffected by the presence of the ROCK inhibitor (Fig. 8A). In fact, Y27632 had no effect on the levels of fibronectin insoluble matrices in all three cell lines. In addition to there being no significant loss of DOC-insoluble fibronectin fibrils, immunofluorescence microscopy analysis of the three cell lines treated with or without Y27632 showed that the only obvious change in the inhibitor-treated cells was an increase in punctate labeling for fibronectin (Fig. 8B). Z-stack and three-dimensional reconstruction analysis of images acquired from control and inhibitor-treated EV, WT$\beta$3, and CA$\beta$3 cultures showed that the punctate labeling was predominantly intracellular (Supplementary Fig. S7).

**Effect of Overexpressing zv$\beta$3 Integrins on the Incorporation of Structurally Different Fibronectin Fibrils Into the Insoluble ECM of TM Cells**

We then considered the possibility that constitutively active zv$\beta$3 integrin signaling changed the structure of the fibronectin in fibrils assembled by CA$\beta$3 cells. Thus, we first used the mAb L8, which recognizes a conformation-sensitive epitope
that is exposed within fibronectin fibrils that have had their quaternary structure unfolded and their tertiary and/or secondary structure stretched in response to cell-derived mechanical forces (Fig. 9A). Analysis of intact monolayers by immunofluorescence microscopy (Fig. 9B) indicated no obvious differences in L8 labeling between the three cell lines. OCW analysis of intact monolayers (Fig. 9C), however, found that WTβ3 cultures demonstrated L8 labeling that was 1.3- to 1.45-fold lower than in EV (P < 0.02) or CAβ3 (P < 0.005) cultures, respectively. In contrast, CAβ3 cultures had significantly higher levels of L8 labeling in DOC-insoluble fibronectin fibrils than the other two cell lines (Fig. 9D). The difference in L8 labeling was 5.7-fold (P < 0.005) comparing CAβ3 and EV cultures while the difference between CAβ3 and WTβ3 cultures was 2.6-fold (P < 0.02). This indicates that fibronectin in the fibrils assembled by CAβ3 cells was more unfolded or stretched compared to the fibronectin in fibrils assembled by the other cell lines.

We then used mAbs IST-9 and BC-1 that detect the alternatively spliced EDA+ and EDB+ domains in fibronectin (Fig. 10A), respectively, to determine if fibrils assembled by CAβ3 cells contained different fibronectin isoforms. As shown in Figure 10B, immunofluorescence microscopy of intact monolayers labeled with mAb IST-9 suggested that CAβ3 cell layers contained slightly more EDA+ fibronectin than the other two cell lines. OCW analysis of intact monolayers (Fig. 10C) confirmed this, although the increase in labeling was not statistically significant. However, there was at least 4-fold more DOC-insoluble EDA+ fibronectin (Fig. 10D) in CAβ3 cultures compared to the other two cell lines (both P < 0.002). Together these data suggest that although the three cell lines may be expressing the same total levels of EDA+ fibronectin, CAβ3 cell are assembling more EDA+ fibronectin into DOC-insoluble fibrils.

Using mAb BC-1 to detect EDB+ fibronectin in fibrils, both immunofluorescence microscopy (Fig. 10B) and OCW analysis (Fig. 10C) showed that intact monolayers of EV cells labeled significantly stronger for this isoform than WTβ3 cells (P < 0.002) and CAβ3 cells (P < 0.02). Labeling for EDB+ fibronectin in fibrils in intact cultures of EV cells was 2.5- and 1.5-fold greater than that seen in WTβ3 and CAβ3 cultures, respectively. In contrast, when we examined the levels of...
EDB+ fibronectin fibrils in cultures extracted with DOC (Fig. 1D), we found that CAβ3 cells incorporated at least 17-fold more DOC-insoluble EDB+ fibronectin fibrils compared to both EV and WTβ3 cultures (both $P < 0.03$). Thus, both the composition (EDA+/EDB+ isoforms) and stretched conformation of fibronectin in fibrils made by CAβ3 cells appear to be different compared to the fibrils assembled by EV and WTβ3 cells.

**DISCUSSION**

In this study we have shown that activation of αvβ3 integrins in TM cells enhances fibronectin fibrillogenesis. This is consistent with our earlier studies that found that HTM cells treated with glucocorticoids expressed higher levels of activated αvβ3 integrins38,40,41 and these cells assembled higher levels of DOC-insoluble fibronectin matrices.12 These fibrils are formed using an alternative ROCK-independent pathway and appear to have different structural properties and composition. Such differences are likely to alter the signaling properties of the ECM and behavior of TM cells.66 Theoretically these differences could contribute to the development of pathological changes observed in glaucoma, especially SIG, where the αvβ3 integrin is likely to be activated.38,67

This is not the first time an integrin other than α5β1, the main integrin responsible for mediating fibronectin fibril assembly,35,36 has been found to be involved in fibronectin fibril formation. There are a limited number of other integrin heterodimers that can promote fibronectin fibrillogenesis, including αvβ3 integrins,68–70 which in some instances appear to play a more significant role in matrix assembly than α5β1 integrins.71 These earlier studies are consistent with the results presented here demonstrating a pronounced enhancement in fibronectin fibrillogenesis in TM cells overexpressing constitutively active αvβ3 integrins.

The increase in fibril formation was not due to a disruption in the canonical α5β1 integrin-mediated fibronectin fibril assembly mechanism that appeared to be functioning in TM cells, including those overexpressing a constitutively active β3 integrin. These studies show that mAb 13, which blocks binding of fibronectin to α5β1 integrins in the early stages of fibril formation, was able to block binding of fibronectin in CAβ3 cultures, suggesting that this early integrin-mediated step was still involved. In addition, fibrillar adhesions, which are the sites where fibronectin fibrils are assembled, only contained α5β1 integrins, further suggesting that an α5β1 integrin step was involved. In contrast, αvβ3 integrins were not found in fibrillar adhesions, implying that αvβ3 integrins were not involved in the initial binding of fibronectin during fibril formation. This suggests that the enhanced deposition of fibronectin fibrils in CAβ3 cells may be due to αvβ3 integrin playing a role in the later stages of fibril formation when the transition of cell surface-soluble fibronectin into insoluble fibrils occurs72 and/or it is altering the signaling pathways that govern fibronectin fibril formation.

How αvβ3 integrin signaling is affecting fibril formation is still unclear. The increased assembly of these fibronectin fibrils could be due to the enhanced RhoA activity of the CAβ3 cells in response to serum (see Fig. 7). Early studies demonstrated that fibronectin fibrillogenesis is a RhoA-dependent process,34,64 and RhoA signaling has also been suggested to play a role in regulating fibronectin matrix formation in TM cell cultures.52 RhoA activity has also been reported to be upregulated when the β3 integrin subunit is overexpressed.73 The changes in RhoA activity would influence the contractile state and traction force generated by a cell through sites of focal adhesions, which in turn can significantly impact a cell’s ability to assemble a fibronectin matrix.34,59,74 The implication here is that traction and contractile forces on fibronectin promote unfolding and stretching of the fibronectin dimer, which in turn initiates a cascade of intermolecular interactions between multiple fibronectin dimers that propagate the formation of a DOC-insoluble fibril. Such an enhanced unfolding and stretching of fibronectin’s conformation was seen in CAβ3 cells when we labeled the matrices with the L8 antibody that recognizes a conformation-sensitive epitope.

Interestingly, when we tried to use the ROCK inhibitor Y27632 to correlate the increased incorporation of fibronectin into the DOC-insoluble matrix with RhoA signaling, we failed to see a decrease in fibronectin fibril formation. Earlier studies

![Figure 7.](image-url)
have suggested that other pathways, independent of RhoA/ROCK, can play a role in fibronectin fibrillogenesis. Studies by Hill et al.17 showed that when living rat eyes were treated with TGF-\( \beta \)1 there was an increase in fibronectin that was only modestly decreased by inhibiting RhoA activity using siRNA. A similar result has also been reported in endothelial cells that appear to be capable of assembling a fibronectin matrix in the absence of a classical RhoA-dependent pathway.63 Additional studies in \textit{Xenopus} embryos \( ^{75} \) and human foreskin fibroblasts \( ^{58} \) also suggested that RhoA/ROCK-independent pathways could play a significant role in promoting fibronectin fibrillogenesis.

Together these observations support our data that a RhoA/ROCK signaling pathway does not appear to play a significant role in the enhanced fibronectin matrix deposition by CA\( \beta \)3 cells. Instead, there appears to be a ROCK-independent pathway that is mediating enhanced fibril formation in these cells. Studies by Shiller et al.\( ^{57} \) have recently proposed that an alternative RhoA-mediated pathway can control contractility and hence may be involved in fibril formation. In this alternative pathway (Fig. 11), the RhoA/ROCK pathway is mediated to a large extent by \( \alpha \)v\( \beta \)1 integrin signaling while signaling via \( \alpha \)v-class integrins such as \( \alpha \)v\( \beta \)3 mediates RhoA signaling via the guanine nucleotide exchange factor (GEF) GEF-H1 and the formin mDia. Other studies support this idea. Zamir et al.\( ^{56} \) found that fibrillar adhesion formation (hence fibrillogenesis) occurred via a RhoA/ROCK-independent mechanism while another study found that inhibiting mDia blocked fibrillar adhesion formation and impaired fibronectin remodeling.\( ^{77} \) Other alternative pathways have also been proposed. Fernandez-Sauze et al.\( ^{55} \) have proposed an alternative Rho-independent mechanism involving ILK and Rac1 that could be responsible for the phosphorylation of myosin light chain and the generation of the contractile forces needed for matrix assembly. Thus, it is possible that the CA\( \beta \)3 cells are utilizing one or both of these pathways to control the contractile forces that enhance fibronectin matrix assembly.

These studies also showed that the composition and stretched state of fibronectin in fibrils appeared to be altered in CA\( \beta \)3 cells. Our studies indicate that the DOC-insoluble fibrils in cultures of CA\( \beta \)3 cells contain substantially more EDA\( ^{+} \) and/or EDB\( ^{+} \) fibronectin than those assembled by EV or WT\( \beta \)3 cells. This was despite the fact that intact EV and CA\( \beta \)3 monolayers demonstrated comparable levels of EDA\( ^{+} \) fibronectin and EV monolayers exhibited significantly more EDB\( ^{+} \) fibronectin labeling than either WT\( \beta \)3 or CA\( \beta \)3 cultures. Yet

**FIGURE 8.** DOC-insoluble fibronectin matrix assembly is independent of ROCK activity. (A) OCW analysis of EV, WT\( \beta \)3, and CA\( \beta \)3 cells treated with or without 5 \( \mu \)M Y27632 ROCK inhibitor for 24 hours prior to extraction with 1% DOC and subsequent fixation. CA\( \beta \)3 cells exhibited significantly enhanced matrix assembly relative to both EV and WT\( \beta \)3 cells in the absence and presence of the ROCK inhibitor (all \( P < 0.01 \)). Data are pooled from two separate assays with triplicate determinations (\( n = 6 \)) and represent the mean ± SE. (B) Compressed Z-stack images of intact, confluent monolayers of EV, WT\( \beta \)3, and CA\( \beta \)3 cells treated as in (A) prior to DOC extraction and labeled with rabbit anti-fibronectin antiserum. Although the fibronectin fibrils are relatively unchanged in the inhibitor-treated cells, there was an apparent increase in intracellular punctate fibronectin labeling (see Supplementary Fig. S7) in the inhibitor-treated cells. Scale bar: 50 \( \mu \)m.
very little EDA+ and/or EDB+ fibronectin could be detected in the insoluble matrix of EV cells. This suggests that activation of αvβ3 integrin is involved in promoting the assembly of fibronectin fibrils that include one or both of these domains.

Interestingly, cultured HTM cells treated with either glucocorticoids, which can cause αvβ3 integrin activity to be upregulated, or TGFβ2 can both upregulate the EDA+ and EDB+ fibronectin isoforms.11,12 In addition, EDA+ fibronectin has been reported to be elevated in glaucomatous TM tissues.11 This suggests, especially in SIG, that αvβ3 integrin activity may be involved in ECM deposition in these circumstances.

In addition to this change in fibril composition, our studies indicate that there is a higher percentage of L8 labeling in the DOC-insoluble fibrils found in CAβ3 cultures. Since the L8 antibody detects an epitope that is exposed when fibronectin is stretched or unfolded in response to cell-derived mechanical forces,54,65,78,79 this suggests that fibronectin in fibrils in CAβ3 cultures may be under more tension or mechanical forces.

This enhanced unfolding and/or stretching of fibronectin fibrils is associated with more rigid fibrils in aging matrices and would be expected to have different biochemical properties that would affect cell behavior.78 The observation that EDB+ fibronectin is more prevalent in fibrils assembled by CAβ3 cells further substantiates this idea that fibrils assembled when αvβ3 integrin is activated have different biological properties. Together these studies show that activation of αvβ3 integrins,
especially by glucocorticoids, may result in the assembly of a fibronectin matrix with biological properties that differs from a matrix assembled under conditions where a\textsuperscript{v}b\textsuperscript{3} integrin is either expressed at very low levels such as in EV cells or not active such as in WT\textsuperscript{b}3 cells. Whether these changes could contribute to the enhanced rigidity of the ECM associated with glaucoma is unknown.

In summary, in addition to the b\textsuperscript{3} integrin–induced alteration in fibronectin fibrillogenesis shown here, earlier work from our lab also found that b\textsuperscript{3} integrin signaling played a role in altering the TM cytoskeleton to form CLANs\textsuperscript{41,46,47} and served as a negative regulator of TM phagocytosis\textsuperscript{40,80}. Collectively, alterations in these biological processes are associated with the pathogenesis of certain forms of glaucoma such as POAG and SIG.\textsuperscript{57,81–83} This suggests that dysregulation of b\textsuperscript{3} integrin signaling may play a significant role in the pathogenesis of these glaucomas and possibly other forms of the disease. Future studies using the tamoxifen-inducible Cre\textsuperscript{+}/\textsuperscript{−}b\textsuperscript{3} integrin\textsuperscript{flox/flox} mouse model we recently developed to knock down expression of a\textsuperscript{v}b\textsuperscript{3} integrin in the trabecular meshwork\textsuperscript{84} should help determine the role of this integrin in POAG.

\textbf{Figure 10.} Constitutively active a\textsuperscript{v}b\textsuperscript{3} integrin increases deposition of specific fibronectin isoforms into the DOC-insoluble matrix. (A) Fibronectin schematic showing the locations of the EDA and EDB alternatively spliced domains. The mAb BC-1 recognizes a conformation-specific epitope formed in the seventh type III repeat when the EDB domain is present.\textsuperscript{85} The IST-9 mAb detects a sequence found within the EDA domain. (B) Intact, confluent monolayers of EV, WT\textsuperscript{b}3, and CA\textsuperscript{b}3 cells were immunolabeled with mAbs IST-9 and BC-1, respectively. No clear differences in labeling intensity were observed between the three cell lines for mAb IST-9. EV cells, however, demonstrated more intense labeling with mAb BC-1 than both WT\textsuperscript{b}3 and CA\textsuperscript{b}3 cells. Negative control cells were labeled with mAb GAL-13 against β-galactosidase and showed no significant labeling (not shown). This experiment was performed twice with identical results. Scale bar: 50 μm. (C) OCW analysis of intact, unextracted monolayers of EV, WT\textsuperscript{b}3, and CA\textsuperscript{b}3 cells plated for 24 hours. No significant difference was found between any of the three cell lines with respect to EDA\textsuperscript{+} fibronectin. Both WT\textsuperscript{b}3 cells (P < 0.002) and CA\textsuperscript{b}3 cells (P < 0.02) demonstrated less EDB\textsuperscript{+} fibronectin in intact monolayers relative to EV cells. Results represent the mean ± SE and consist of data pooled from two assays using quadruplicate and triplicate determinations, respectively (n = 7). (D) OCW analysis of DOC-insoluble fibronectin fibrils in EV, WT\textsuperscript{b}3, and CA\textsuperscript{b}3 cultures. CA\textsuperscript{b}3 cells demonstrated significantly higher levels of DOC-insoluble EDA\textsuperscript{+} fibronectin fibrils than EV and WT\textsuperscript{b}3 cells (P < 0.002). CA\textsuperscript{b}3 cells demonstrated significantly higher levels of DOC-insoluble EDB\textsuperscript{+} fibronectin fibrils than EV and WT\textsuperscript{b}3 cells (P < 0.03). Results represent the mean ± SE and consist of data pooled from two assays using triplicate determinations (n = 6).
Schiller et al. suggest that there are alternative ways in which stress fibers can be assembled by integrin signaling. Signaling from fibrillogenesis. Together these two processes lead to stronger contractile forces that could ultimately lead to the enhanced fibronectin matrix assembly demonstrated by CAβ3 integrins. In addition, these actin filaments are necessary for the formation of fibrillar adhesions that are the sites of fibronectin deposition. None

References


