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A Specific $\alpha_{5}\beta_{1}$ Integrin Conformation Promotes Directional Integrin Translocation and Fibronectin Matrix Formation

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Summary

Integrin adhesion receptors are structurally dynamic proteins that adopt a number of functionally relevant conformations. We have produced a conformation-dependent anti- α_5 monoclonal antibody (SNAKA51) that converts $\alpha_5\beta_1$ into a ligand-competent form and promotes fibronectin binding. In adherent fibroblasts, SNAKA51 preferentially bound to integrins in fibrillar adhesions. Clustering of integrins expressing this activation epitope induced directional translocation of $\alpha_5\beta_1$, mimicking fibrillar adhesion formation. Priming of $\alpha_5\beta_1$ by SNAKA51 increased the accumulation of detergent-resistant fibronectin in the extracellular matrix, thus identifying an integrin conformation that promotes matrix assembly. The SNAKA51 epitope was mapped to the calf-1/calf-2 domains. We propose that the action of the antibody causes the legs of the integrin to change conformation and thereby primes the integrin to bind ligand. These findings identify SNAKA51 as the first anti-integrin antibody to selectively recognize a subset of adhesion contacts, and they identify an integrin conformation associated with integrin translocation and fibronectin matrix formation.

Keywords

integrin; fibronectin; conformation; monoclonal antibody; matrix assembly

Introduction

Integrins are heterodimeric cell surface receptors that bind to the extracellular matrix (ECM), and provide a physical link to the intracellular cytoskeleton. Integrin function depends on an ability to modulate receptor structure rapidly, and inactive, primed, and ligand-bound conformations with different affinities for ligand-binding have been characterized (Humphries, 2000; Hynes, 2002; Mould, 1996; Shimaoka et al., 2002). Integrin ligand-binding ability can be controlled both by the binding of cytoplasmic factors that induce conformational changes and by regulated positioning on the cell surface to fav3or high-avidity binding. The mechanisms responsible for transferring this signal through the integrin molecule to the extracellular head region, and for regulating ligand-binding, extracellular matrix formation, and remodelling of the cell-matrix interface, are not well

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understood. Several conformational changes have been suggested to underpin integrin priming, and it is possible that a series of events occurs during acquisition of ligand competency. The crystal structure of the $\alpha_v\beta_3$ integrin revealed a bent molecule where the globular head contacted the stalk region (Xiong et al., 2001). Building on this information, a switchblade model for priming was proposed in which divalent cation or ligand occupancy induces a conformational change from the bent to the extended conformation (Takagi et al., 2002). This unbending revealed β subunit activation epitopes and increased ligand-binding affinity (Beglova et al., 2002). Another conformational change associated with integrin priming is the separation of the α and β subunit legs (Kim et al., 2003; Lu et al., 2001; Takagi et al., 2001).

The first integrin crystal structure resolved the atomic details of many of the domains of the heterodimer and confirmed the predicted regions for ligand-binding (Xiong et al., 2001). In addition, conformation-dependent monoclonal antibodies have been valuable for studying the link between receptor shape and activity. The majority of antibodies that modulate the integrin activation state bind to the head region of the integrin (Humphries et al., 2003b). These antibodies allosterically alter the structure of the ligand-binding pocket in the α subunit β propeller and β subunit A-domain through local conformational changes. These local effects can stimulate or inhibit ligand-binding depending on the location of the antibody epitope and the conformation induced. The binding of ligand to the integrin can also affect the expression of certain antibody epitopes. Many of the antibodies that increase ligand-binding or recognize active integrin have ligand-induced binding sites (LIBS) (Bazzoni et al., 1995; Mould et al., 1995b).

Integrins can be localized in different adhesion structures on the cell surface, termed focal complexes, focal adhesions, fibrillar adhesions, and 3D-matrix adhesions. These contacts reflect different stages of interaction of cells with the ECM, and each is formed and disrupted in a dynamic, cyclical manner as cells translocate through sequential recruitment and loss of cytoskeletal and signaling molecules (Geiger et al., 2001; Webb et al., 2004). While focal adhesions provide robust anchorage via transcellular actomyosin-containing stress fibres, fibrillar adhesions are the major sites of fibronectin matrix deposition. Ligated $\alpha_5\beta_1$ integrin molecules translocate centripetally out of focal adhesions generating fibrillar adhesions. This directional movement along the actin cytoskeleton stretches and organizes bound fibronectin into fibrils of the extracellular matrix (Pankov et al., 2000; Zamir et al., 2000).

For integrins to function as vehicles for extracellular matrix deposition, their activity needs to be highly controlled. This control appears to be through conformational modulation (Humphries et al., 2003a; Sims et al., 1991). In this study, we tested the hypothesis that $\alpha_5\beta_1$ integrins associated with fibronectin matrix formation have a particular conformational property. We have identified a unique subpopulation of $\alpha_5\beta_1$ integrins located in fibrillar adhesions that have a specific conformation recognized by a novel anti- α_5 antibody. Integrins in this conformation can undergo directional translocation in fibrillar adhesions and promote fibronectin matrix formation.

Materials and methods

Antibodies

The antibodies used were the mouse anti-human integrin β_1 antibodies TS2/16 (activating; a gift from Francisco Sanchez-Madrid, Universidad Autonoma de Madrid, Spain), 12G10 (activating) (Mould et al., 1995b), and K20 (nonfunction-modulating, Immunotech); rat antihuman integrin β_1 antibodies mAb13 (inhibitory) (Akiyama et al., 1989) and 9EG7 (activating, Pharmingen); rat anti-human integrin α_5 mAb16 (inhibitory) (Akiyama et al.,

1989) and mAb11 (nonfunction-modulating) (Miyamoto et al., 1995); mouse anti-human integrin α_5 antibodies SNAKA52 (inhibitory) and JBS5 (inhibitory, Serotec); mouse anti-human integrin α_V L230 (inhibitory, ATCC); and polyclonal rabbit anti-human fibronectin Rb745 (Cukierman et al., 2001).

Production of hybridomas

 $\alpha_5\beta_1$ integrin was isolated from placenta by two consecutive affinity purifications using mAb13 and mAb16 columns (Mould et al., 1995a). Hybridomas were produced by selection from a fusion of spleen cells from immunized mice and 653 myeloma cells (Mould et al., 1991). Media from the resulting hybridomas was screened by ELISA, and antibody was purified using protein G-Sepharose.

ELISA

 $a_5\beta_1$ or $a_4\beta_1$, diluted to 1 µg/ml in PBS, was used to coat an Immulon-3 assay plate (Dynatech) overnight at 4°C. Wells were blocked with 5% (w/v) BSA in TBS (150 mM NaCl, 25 mM Tris HCl, pH 7.4), and incubated at room temperature for 30 minutes. Culture medium from the hybridomas or antibody was added and incubated at room temperature for 1 hour. Wells were washed 3 times with PBS⁺, then peroxidase-conjugated anti-mouse immunoglobulin (Dako) in PBS⁺ was added. Plates were incubated for 30 minutes at room temperature and washed 3 times with PBS⁺. 0.1% (w/v) 2,2'-azino-bis (3-ethylbenzthiazoline 6-sulphonic acid) (ABTS) in 0.1 M sodium acetate, 0.05 M NaH₂PO₄, pH 5.0, 0.01% (v/v) H₂O₂ was added, and absorbance readings were determined at 405 nm on a Dynatec MR4000 plate reader.

K562 cell adhesion to fibronectin

Fibronectin (Miekka et al., 1982), diluted to 2 μ g/ml in PBS⁺, was coated onto an Immulon-2 U-bottom plate overnight at 4°C. The surface was blocked with 10 mg/ml heatdenatured BSA for 1 hr at room temperature. The plate was washed once with PBS⁺ before addition of DMEM, 25 mM HEPES buffer containing 20 μ g/ml antibody or 100 nM phorbol 12-myristate 13-acetate (PMA). K562 cells were added at 2.5×10⁴ cells/well in DMEM, 25 mM HEPES. The plate was incubated for 30 minutes at 37°C, 10% CO₂. Medium and unbound cells were removed by inversion of the plate and four PBS⁺ washes. 0.2% (w/v) crystal violet, 20% (v/v) methanol solution was added and incubated for 20 minutes. The plate was washed twice by immersion in water, dried before addition of 10% (w/v) SDS, and incubated for 20 minutes before reading absorbance at 600 nm on an Emax plate reader (Molecular Devices).

Production of recombinant Fc-tagged integrins

DNA constructs—All human α_5 -Fc and β_1 -Fc DNA constructs used here have been described previously (Coe et al., 2001). C-terminally truncated versions of α_5 -Fc [residues 1-613 (Δ 613), 1-621 (Δ 621), 1-694 (Δ 694), 1-795 (Δ 795)] and β_1 -Fc [residues 1-455 (Δ 455)] were produced. The β_1 -Fc ADMIDAS mutant (D138A) constructs used were generated as described by Mould *et al.* (Mould et al., 2003a). For production of α_5 -Fc calf domain constructs, DNA was amplified by PCR, inserting a 5' Hind III and a 3' Sal I restriction site, and ligated into pEE12.2hFc. The leader sequence of a murine antibody (Kabat et al., 1987) was also incorporated at the 5' end of each construct to facilitate protein secretion. Calf-1 contained α_5 residues 606-749, calf-2 contained residues 746-951, and calf-1/calf-2 contained residues 606-951.

Expression of $\alpha_5\beta_1$ **-Fc proteins**—CHOL731H cells (Cockett et al., 1991) were grown to ~95% confluence, and 20 µg of α_5 -Fc and β_1 -Fc DNA (or α_5 -Fc calf domain construct

alone) was used to transfect cells using LipofectAMINE 2000TM reagent (Invitrogen) according to the manufacturer's instructions. After 4 days, culture supernatant was harvested by centrifugation at $1000 \times g$ for 5 minutes.

Solid-phase assays

Placental $\alpha_5\beta_1$ integrin, diluted to 1 µg/ml in PBS⁺, was used to coat an Immulon-3 assay plate (Dynatech) overnight at 4°C, or anti- β_1 integrin (K20) 2 µg/ml was coated overnight at 4°C. Wells were blocked with 5% (w/v) BSA in TBS for 2 hours. K20 coated plates were used to tether $0.5 \,\mu$ g/ml placental integrin incubated for 2 hours at room temperature. For capturing recombinant Fc-tagged integrin, goat anti-human γ_1 Fc antibody (Jackson ImmunoResearch Laboratories), diluted to 2.6 µg/ml in PBS⁺, was coated into a immunoassay plate (Costar) overnight at 4°C. Wells were blocked with 5% (w/v) BSA in TBS for 2 hours. Transfection culture supernatants containing integrin-Fc protein were incubated in the wells for 1 hour, and wells were subsequently washed three times with TBS, 1 mM MnCl₂, 1 mg/ml BSA (buffer A). Ligand and antibodies ($10 \mu g/ml$) were added in combination in buffer A. Biotinylated FnIII (6-10) was added at 0.1 μ g/ml and unlabelled FnIII (6-10) at 20 µg/ml (Danen et al., 1995). Cations or EDTA were added at 2 mM. The plate was incubated at 30°C for 3 hours. Wells were washed three times with buffer A, then ExtrAvidin Peroxidase (Sigma) in buffer A was added and incubated for 20 minutes. Alternatively, peroxidase-conjugated anti-mouse or anti-rat immunoglobulin (Dako) was used and incubated for 1 hour. The plate was washed four times with buffer A, ABTS substrate was added, and absorbance was determined at 405 nm.

Indirect immunofluorescence staining

Primary human foreskin fibroblasts (hFF) were a gift from S. Yamada (NIDCR, NIH) and were used at passages 9-18. Cells were plated on 12 mm glass coverslips in DMEM with 10% FCS in a 24-well dish at 5000 cells/well and cultured overnight. The cells were fixed for 20 minutes with 4% (w/v) paraformaldehyde, 5% (w/v) sucrose in PBS. Cells were stained with anti-fibronectin Rb745, Alexa 594-conjugated anti- α_V integrin L230, Alexa 488-conjugated anti- α_5 antibodies SNAKA51, SNAKA52, or mAb11, and Cy2-conjugated anti-rat IgG and AMCA-conjugated anti-rabbit IgG secondary antibodies (Jackson ImmunoResearch Laboratories). Stained samples were mounted in GEL/MOUNTTM (Biomeda Corp.) containing 1 mg/ml 1,4-phenylendiamine (Fluka) to reduce photobleaching. Immunofluorescence images were obtained with a 63x/1.40 oil objective on a Zeiss Axiophot microscope equipped with a Photometrix CH 350 cooled CCD camera. Digital images and image overlays were obtained using MetaMorph 4.6 software (Universal Imaging Corp.).

Epitope mapping (sandwich ELISA)

 $\alpha_5\beta_1$ -Fc was captured in wells from transfection supernatants via anti-human γ_1 Fc antibody as detailed above. Wells were washed three times with buffer A. SNAKA51 was diluted to 10 µg/ml in buffer A and added to the wells for 2 hours. Wells were washed three times, and peroxidase-conjugated anti-mouse antibody (Jackson ImmunoResearch Laboratories), diluted in buffer A, was added for 30 minutes. Wells were washed four times with buffer A, incubated with ABTS substrate, and the absorbance was determined at 405 nm.

Antibody clustering and chasing

hFF were plated on glass coverslips in DMEM, 1% (v/v) fibronectin-depleted FCS containing 25 μ g/ml cycloheximide. After overnight incubation, cells were labelled in vivo by incubation in the same medium containing 10 μ g/ml anti- α_5 antibody for 20 minutes.

After two washes with warm medium, cells were incubated with 2 μ g/ml goat anti-mouse IgG or goat anti-rat IgG in the same medium for 30 minutes (clustering and chasing period). All incubations were at 37°C in a humidified chamber with 10% (v/v) CO₂. At the end of the labelling and chasing periods, samples were fixed and antibody-containing clusters were visualized with CY2-conjugated donkey anti-goat IgG and unclustered antibody with CY2-conjugated donkey anti-mouse IgG (Jackson ImmunoResearch Laboratories). Focal adhesions were stained with Alexa-594-conjugated anti- α_V antibody L230. Samples were mounted and imaged as for immunofluorescence staining.

Fibronectin incorporation into the extracellular matrix

Human salivary gland (HSG) cells were plated into a 6-well dish at 1×10^5 cells/well and cultured overnight. The medium was changed, and cells were incubated a further 24 hours. The cells were then treated with a range of concentrations of biotinylated fibronectin or 10 µg/ml biotinylated fibronectin with or without 25 µg/ml antibody or a range of antibody concentration and incubated overnight or for different time periods. The cells were extracted with 20 mM Tris HCl, pH 8.5, 1% (w/v) deoxycholic acid, 2 mM N-ethylmaleimide, 2 mM iodoacetic acid, 2 mM EDTA, 2 mM PMSF (DOC buffer), extruded through a 23-gauge needle five times and centrifuged at 20,000 g for 20 minutes at 4°C. The pellet was washed once with DOC buffer and prepared for SDS polyacrylamide electrophoresis. Samples were resolved on 4-12% Tris-glycine gradient gels (Novex) and electrotransfered onto nitrocellulose membranes. The membranes were blocked with 5% (w/v) nonfat milk in TBS, 0.05% (w/v) Tween-20 for 30 minutes, immunoblotted with streptavidin peroxidase (Boehringer Mannheim), together with anti-pan cytokeratin (Sigma), using secondary peroxidase-conjugated anti-mouse immunoglobulin (Amersham Pharmacia Biotech). Immunoblots were visualized using the ECL system and Hyperfilm X-ray film (Amersham Pharmacia Biotech).

Results

Identification of a novel antibody that primes $\alpha_5\beta_1$ ligand-binding

Monoclonal antibodies have emerged as a valuable tool for studying the various conformational classes adopted by integrins. To develop additional reagents for probing integrin conformation, we generated antibodies directed against the human placental $\alpha_5\beta_1$ integrin and identified their subunit specificity by ELISA. Two mice were immunised with $\alpha_{5}\beta_{1}$ integrin from which approximately 500 colonies were tested, from the cell fusion products and 12 positive colonies were cloned. An ELISA assay using immobilised $\alpha_4\beta_1$ and $\alpha_5\beta_1$ integrin was used to identify subunit specificity. Antibodies were then tested in a K562 cell adhesion assay to assess their ability to stimulate the $\alpha_5\beta_1$ integrin to bind fibronectin. As previously established, K562-expressed $\alpha_5\beta_1$ integrin was predominantly in an inactive state, and addition of stimulatory anti- β_1 integrin antibodies (TS2/16, 9EG7, and 12G10) or PMA was required to promote cell adhesion (Fig. 1). Nonfunction-modulating anti- α_5 (mAb11) and anti- β_1 (K20) as well as inhibitory anti- α_5 (JBS5 and SNAKA52) antibodies did not support cell adhesion. A new anti-a,5 antibody, SNAKA51 with no immunoreactivity against $\alpha_4\beta_1$, promoted cell adhesion to a similar extent to TS2/16, 9EG7, and 12G10, suggesting that this antibody is also able to stimulate the $\alpha_5\beta_1$ integrin (Fig. 1). From this fusion, SNAKA51 was the only antibody produced that promoted cell adhesion. To confirm the ability of SNAKA51 to stimulate $\alpha_5\beta_1$ integrin function in a biochemical system, solid-phase ligand-binding assays were employed. A biotinylated fibronectin fragment containing type III repeats 6 to 10 [FnIII (6-10)] was used as the ligand. This fragment contains both the RGD and synergy sequence required for $\alpha_5\beta_1$ integrin binding and was used at a concentration at which an antibody effect on ligand binding would be detectable (Mould et al., 1997). SNAKA51 was found to increase ligand-binding to both

placental and recombinant $\alpha_5\beta_1$ integrin above the no-antibody control and the control anti- β_1 antibody (K20), as did the activating anti- β_1 antibody (12G10) (Fig. 2a). This increase in ligand-binding by SNAKA51 was antibody dose-dependent (data not shown).

To determine whether the epitope recognized by SNAKA51 was affected by conformational changes in the integrin induced by ligand-binding or cations, the amount of antibody bound to the $\alpha_5\beta_1$ integrin was measured in the presence and absence of the FnIII (6-10) fragment or in different cation-containing buffers. The addition of FnIII (6-10) increased the amount of SNAKA51 binding to $\alpha_5\beta_1$ over a range of antibody concentrations, for example by nearly 5-fold at an antibody concentration of $0.1 \ \mu g/ml$ (Fig. 2b), indicating that the SNAKA51 epitope is a ligand-induced binding site (LIBS). SNAKA51 binding was increased by the addition of manganese, but not affected by calcium, magnesium or EDTA (data not shown). Using an ELISA measuring the amount of antibody bound to $\alpha_5\beta_1$ integrin, SNAKA51 showed significantly reduced binding to $\alpha_5\beta_1$ integrin, for example 24% at 1 μ g/ml, in comparison to both JBS5 and SNAKA52, which both bound equally well to the integrin, and 77% in comparison to 12G10 (data not shown).

These findings indicated that SNAKA51 increases $\alpha_5\beta_1$ ligand-binding and recognizes a subpopulation of the purified integrin through binding to a cation-independent LIBS epitope.

SNAKA51 specifically stains fibrillar adhesions

Since the SNAKA51 antibody appears to recognize a subpopulation of the $\alpha_5\beta_1$ integrin, we examined its localization on the cell surface. Human foreskin fibroblasts were fixed and stained with a series of anti- α_5 antibodies, anti-fibronectin (Rb745), and anti- α_V integrin antibody (L230), the latter as a marker for focal adhesions. SNAKA51 colocalized with fibronectin in fibrillar adhesions (Fig. 3a, left panels). Inhibitory α_5 antibody SNAKA52 gave diffuse staining across the whole cell surface (Fig. 3a, center panels), which was similar to other inhibitory α_5 or β_1 integrin antibodies (JBS5 and mAb13, data not shown). The nonfunction-modulating α_5 antibody mAb11 stained a combination of these two distributions, which included both integrins that were distributed across the cell surface and those that localized in adhesion complexes (Fig. 3a, right panels). These findings indicate that the subpopulation of integrin recognized by SNAKA51 is specifically localized to fibrillar adhesions and is probably in either a ligand-bound or ligand-competent conformation. This population is distinct from those recognized by both the SNAKA52 and mAb11 antibodies.

Ligand-competent β_1 integrin, detected with the 9EG7 antibody, has been shown to translocate out from focal adhesions into fibrillar adhesions, suggesting that it detects $\alpha_5\beta_1$ (Pankov et al., 2000). Since this integrin localizes to focal adhesions before translocating out into fibrillar adhesions, we compared the localization of SNAKA51-positive with 9EG7positive integrins. 9EG7 stained integrins throughout the focal adhesions and out into the fibrillar adhesions, whereas SNAKA51 labelled integrins that were at the distal edge of the focal adhesion away from the cell perimeter and along the fibrillar adhesions (Fig. 3b). Similar results to the 9EG7 staining were obtained with 12G10 (data not shown). The integrin in the focal adhesion staining was shown to be $\alpha_5\beta_1$ since mAb11 and 9EG7 showed complete colocalization. These findings indicate that the α_5 subunit undergoes a conformational change during the transition from focal adhesions to fibrillar adhesions. The SNAKA51 antibody therefore identifies a unique subpopulation of $\alpha_5\beta_1$ integrin compared to other anti-integrin antibodies.

The $\alpha_5\beta_1$ integrin priming signal is transduced through its legs to the β_1 A-domain

Elucidating the nature of the $\alpha_5\beta_1$ integrin conformation in fibrillar adhesions should help elucidate how the bidirectional control of matrix deposition and intracellular signalling are coordinated. To understand how the SNAKA51 antibody modulates $\alpha_5\beta_1$ function, we initially mapped its epitope. Recombinant truncated $\alpha_5\beta_1$ integrins were expressed as Fctagged proteins (Coe et al., 2001; Ridgway et al., 1996) (Fig. 4a). SNAKA51 recognized recombinant integrin containing full-length extracellular α_5 and β_1 subunits. However, any deletion from the carboxyl-terminus of the a_5 construct, including truncation into the calf-2 domain, abolished binding (Fig. 4b, Δ 795, Δ 694, Δ 621, Δ 613). An integrin containing a truncated $\beta_1 \log (\Delta 455\beta_1)$, together with full-length α_5 , bound SNAKA51 to a similar degree as the full-length protein, confirming the epitope location in the a_5 subunit leg (Fig. 4b). The calf-1 and -2 domains were expressed as Fc-tagged proteins, either together as one construct (residues 606-951) or as individual domains (calf-1 residues 606-749 or calf-2 residues 746-951). The SNAKA51 antibody only recognized the construct containing both calf-1 and calf-2 domains and did not bind to either of the individual domains (Fig. 4b). The epitope for SNAKA51 is therefore within the calf domains; it is conceivable that the calf domains require each other for their correct folding and that the antibody epitope might be in either calf or at the calf-1/calf-2 junction.

Since the SNAKA51 epitope maps to the α_5 leg region of the integrin, the conformational change that it triggers could either be transduced directly up the a_5 subunit or it could cross to the β_1 leg and then up to the ligand-binding site in the integrin head region. The $\alpha_5\beta_1$ Fctagged integrin has a high constitutive level of ligand-binding (Fig. 5), which makes it difficult to observe any further activation. Therefore, we employed a mutant integrin containing the point mutation D138A in the ADMIDAS cation-binding site in the β_1 subunit, which we have shown recently to be constitutively inactive. The ligand-binding ability of this mutant can be rescued with the stimulatory anti- β_1 antibody 12G10, demonstrating that it is competent to bind ligand (Mould et al., 2003a). Mutant integrin was produced in both the full-length β_1 and $\Delta 455\beta_1$ constructs. SNAKA51 and 12G10 promoted ligand-binding by the full-length ADMIDAS mutant, with 12G10 having a more potent effect (Fig. 5, $\alpha_5\beta_1$ D138A). The greater effect of 12G10 is probably due to its epitope being spatially close to the ligand binding pocket and SNAKA51 having to induce a long distance allosteric effect. In the truncated β_1 subunit construct, SNAKA51 stimulation was abolished, while 12G10 was still able to activate the integrin ($\alpha_5 \Delta 455\beta_1$ D138A). These results indicate that the priming signal from the SNAKA51 epitope is transduced between the integrin legs from a_5 to β_1 and then up to the β A-domain, where conformational changes promote the integrin to bind ligand. In agreement with this conclusion, 12G10 and SNAKA51 had a reciprocal effect, with the binding of one increasing the epitope expression of the other (data not shown).

Clustering of active $\alpha_5\beta_1$ leads to integrin translocation

The $\alpha_5\beta_1$ integrin has been shown to translocate from focal adhesions across the cell surface and form fibrillar adhesions (Pankov et al., 2000). Because the SNAKA51 antibody recognizes integrin in a specific conformation that is localized to fibrillar adhesions, we hypothesized that antibody addition might induce formation of these adhesion structures. Human fibroblasts were treated with cycloheximide and cultured in the absence of fibronectin to prevent any ligand-related $\alpha_5\beta_1$ integrin translocation. These cells were treated with anti- α_5 antibodies, and then the antibodies were clustered with secondary antibody. Clustered integrin was then chased to observe integrin movement compared to unclustered antibody, fixing either after the initial labelling or after antibody-chasing without clustering. SNAKA51 initially labelled the entire cell surface, but after clustering it translocated along the cell membrane away from anti- α_V (L230)-labelled focal adhesions and formed linear fibrillar adhesion-like structures along the cell surface (Fig. 6a). In contrast, the inhibitory (SNAKA52) and nonfunction-modulating (mAb11) antibodies remained unorganized on the cell surface even after clustering (Fig. 6b, c). Cells treated with anti- α_5 antibody and chased in the absence of the clustering secondary antibody did not exhibit the translocation away from the focal adhesions observed for clustered SNAKA51. Consequently clustering of the SNAKA51-positive subpopulation, even in the absence of ligand, led to directional integrin translocation. These results indicate a requirement for integrins to adopt a specific activated conformation for translocation to occur.

Integrin priming by SNAKA51 increases the incorporation of fibronectin into the matrix

Translocation of the $\alpha_5\beta_1$ integrin is the driving force for fibronectin fibrillogenesis (Pankov et al., 2000). The translocation ability of the clustered SNAKA51-positive $\alpha_5\beta_1$ integrin conformer into fibrillar adhesion-like structures and its ability to bind fibronectin suggested that the integrin needs to be in this conformation when it drives fibronectin fibrillogenesis. To test this hypothesis, we measured the ability of cells treated with anti- α_5 antibodies to produce fibronectin matrix. Fibronectin incorporated into the extracellular matrix can be characterized separately from soluble and non-matrix associated fibronectin by the use of the detergent deoxycholate. Human salivary gland (HSG) cells (which do not deposit an extensive fibronectin matrix) were treated with a range of concentrations of biotinylated fibronectin. The cells were then extracted with deoxycholate-containing buffer, and the insoluble fraction was analyzed by Western blotting. Biotinylated fibronectin was detected using peroxidase-conjugated avidin and compared to the amount of cytokeratin in the same sample to control for sample loading. Little or no fibronectin was incorporated into the matrix up to a concentration of $5 \,\mu$ g/ml exogenous fibronectin (Figure 7a). A concentration of 10 µg/ml biotinylated fibronectin was used for further assays. The incorporation of fibronectin into the matrix was measured over time, or in the presence of different integrin antibodies. Fibronectin was easily detectable in the insoluble fraction after 3 hours (Figure 7b). SNAKA51 antibody substantially increased the incorporation of biotinylated fibronectin into the insoluble fraction of the HSG cells compared to the no antibody treated control (Fig. 7c). The activating β_1 integrin antibody TS2/16 and inhibitory α_5 integrin antibody SNAKA52 were used as positive and negative controls, respectively. Fibronectin incorporation was suppressed by the inhibitory a_5 antibody (SNAKA52). As little as 1 μ g/ ml of SNAKA51 was capable of increasing fibronectin incorporation into the matrix, and the level detected was not increased by addition of more antibody (Figure 7d). Therefore, SNAKA51 has the ability to activate the $\alpha_5\beta_1$ integrin and promote directional integrin translocation associated with the incorporation of more fibronectin into the matrix.

Discussion

In this study, we have tested the hypothesis that a specific $\alpha_5\beta_1$ integrin conformation is associated with the process of fibronectin matrix formation. We produced monoclonal antibodies directed against the $\alpha_5\beta_1$ integrin and identified an anti- α_5 antibody (SNAKA51) that stimulated $\alpha_5\beta_1$ and promoted cell adhesion and ligand-binding. This antibody had a manganese-sensitive LIBS epitope and bound to a subpopulation of integrin located in fibrillar adhesions. The molecular stimulation initiated by this antibody appears to be transferred from its epitope on the α_5 calf domains, through the β_1 leg, and up to the β Adomain, thereby increasing ligand-binding. By clustering the integrin via this activation epitope, directional translocation of the integrin away from the focal adhesions was stimulated, thereby mimicking normal $\alpha_5\beta_1$ integrin translocation during fibrillar adhesion formation. Induction of the SNAKA51-positive primed integrin conformation increased the incorporation of soluble fibronectin into matrix by cells. Our results demonstrate that a specific $\alpha_5\beta_1$ integrin conformer is required for translocation and formation of fibrillar adhesions and fibronectin matrix.

We generated and characterized an anti- α_5 antibody (SNAKA51) that increased K562 adhesion to fibronectin. SNAKA51 promoted cell adhesion and in vitro ligand-binding to a similar extent as activating β_1 antibodies. Thus, SNAKA51 is a member of a rare group of activating integrin antibodies that bind to the α subunit of non-A domain-containing integrins. Several stimulatory anti- α_{IIb} activating antibodies have been described that bind either to the β -propeller domain (PT25-2 and D33C) (Gulino et al., 1990; Puzon-McLaughlin et al., 2000; Tokuhira et al., 1996) or to a site close to the heavy/light chain border in the membrane proximal region (PMI-1) (Calvete et al., 1991; Loftus et al., 1987). An activating α_L antibody, NKI-L16, binds to the bottom of the thigh domain (Huang and Springer, 1995; Keizer et al., 1988). Another activating α_5 antibody has been identified, but its binding site has not been described in any detail (Chiarugi et al., 2003). The wide range of domains containing the activating antibody epitopes is consistent with dynamic structural changes taking place throughout the molecule during priming.

Using modified ELISA assays, we found that SNAKA51 has a manganese-sensitive LIBS epitope which was mapped to the calf domains of the α_5 subunit leg region (Fig. 8a). This epitope is distant from the ligand-binding site in the integrin head region, suggesting a longdistance conformational transduction of the stimulatory signal from this site to the ligandbinding pocket. SNAKA51 rescued ligand-binding to a deactivated β_1 ADMIDAS mutant, indicating that the antibody is able to induce a conformational change in the β_1 A-domain from its remote epitope. The conformational changes in the β_1 A-domain required for integrin priming have been identified as shifts in the a_1 and a_7 helices (Luo et al., 2003; Mould et al., 2003b). Truncation of the $\beta_1 \log (\Delta 455\beta_1)$ abolished the stimulatory activity of SNAKA51, demonstrating that the signal from the α_5 calf domain is dependent on the β_1 leg being present and suggests a movement between the two subunit legs. We propose that the SNAKA51 signal is transferred through the β_1 leg by movement apart of the two subunits. This separation could be due to a steric effect of the antibody pushing the legs apart and may represent the upstream event normally caused by binding of cytoplasmic factors (Calderwood, 2004; Hughes et al., 1996). The integrin crystal structure reveals contacts between the calf-2 domain of the α subunit and the β subunit EGF-4 and β TD domains (Xiong et al., 2001). The leg movement we propose to be induced by SNAKA51 could break these inter-subunit contacts and cause bending in the flexible region between the EGF-4 and β TD domains of the β subunit, leading to integrin priming. These data suggest that this novel antibody binds only to integrins in a certain conformation/activity state, which occurs via the dynamic equilibrium of integrin conformers, the integrin-associated cytoplasmic complex, or ligation.

The localization of SNAKA51 on fibroblasts demonstrated that its epitope is exposed on a specific population of $\alpha_5\beta_1$ in a ligand-bound or ligand-competent conformation in fibrillar adhesions. This SNAKA51-positive $\alpha_5\beta_1$ was located at the edge of focal adhesions where there is a concentration of fibronectin molecules that marks the initiation point for formation of fibrillar adhesions and fibronectin fibrillogenesis. The SNAKA51 binding in focal adhesions did not fully colocalize with 9EG7 or mAb11, suggesting that several intermediate conformations exist for the $\alpha_5\beta_1$ integrin, and epitope expression for 9EG7 and SNAKA51 binding are indicators of different conformational states.

The clustered, SNAKA51-labelled integrin translocated away from focal adhesions in a directional manner similar to that observed in fibrillar adhesion formation (Pankov et al., 2000). This movement only occurred with clustered SNAKA51, demonstrating that a specific conformation and multiple $\alpha_5\beta_1$ integrins are required for translocation. The ability

to stimulate translocation was not specific to SNAKA51, as 9EG7 triggered the same effect (data not shown). Regardless, we were able to induce and mimic $\alpha_5\beta_1$ integrin translocation even in the absence of ligand, indicating that clustering of a specific integrin conformation is all that is required for this function. These results are consistent with the driving force behind fibrillar adhesion formation being generated from the cytoplasmic protein complex associated with the actin cytoskeleton. This force initiates stretching of the fibronectin molecules required to expose cryptic sites used for fibrillogenesis (Pankov and Yamada, 2002; Zhong et al., 1998). In addition, the fibronectin dimer has four binding sites for $\alpha_5\beta_1$ and so naturally clusters this integrin (see (Pankov and Yamada, 2002)). Clustering may be required to build a large enough cytoplasmic complex to connect to the actin cytoskeleton and drive translocation.

Different functional antibodies affect the activation state of the integrin, making it possible to alter the integrin affinity. We used fibronectin incorporation into the extracellular matrix as a readout for the priming of $\alpha_5\beta_1$ ligand-binding. The HSG cells do not normally deposit an extensive fibronectin matrix. Inducing the SNAKA51-specific $\alpha_5\beta_1$ integrin conformation on HSG cells promoted soluble fibronectin incorporation into the matrix, supporting the conclusion that $\alpha_5\beta_1$ integrin adopts this conformation when driving fibronectin fibrillogenesis.

From current knowledge and the results obtained in this study, we propose a model in which the $\alpha_5\beta_1$ integrin, which is initially diffuse on the cell surface (state 1, Fig. 8b) and perhaps in a bent conformation, is recruited to the focal adhesion where a conformational event occurs, such as unbending, that results in a primed β_1 subunit (state 2). This integrin moves to the distal region of the focal adhesion where either through changes in the cytoplasmic complex or ligation it undergoes a conformational change in the α_5 subunit in which the legs of the integrin are separated and the SNAKA51 epitope is revealed (state 3). This fully primed, or ligand-occupied, and clustered integrin is then actively transported out into fibrillar adhesions and drives fibronectin fibrillogenesis (state 4).

The process of fibronectin matrix formation is important for many cellular functions that involve either cellular sensing of the surrounding environment or using the matrix as a structural support. The $\alpha_5\beta_1$ integrin is the part of the complex molecular motor that crosses the cell membrane barrier to drive fibronectin fibrillogenesis. We have elucidated a conformational path through the integrin structure used for integrin priming. The conformation of both α_5 and β_1 integrin subunits are purposely controlled, and a specific conformation is required for fibronectin matrix formation.

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Figure 1. K562 cell adhesion to fibronectin is promoted by SNAKA51 and other stimulatory anti- β_1 antibodies

K562 cells were allowed to attach to a fibronectin-coated surface $(2 \mu g/ml)$ in the presence of the indicated anti-integrin antibodies $(10 \mu g/ml)$, PMA (100 nM), or no antibody (control). Unattached cells were removed, and remaining cells were fixed and stained with crystal violet. Cell attachment was quantified by absorbance measured at 600 nm.



Figure 2. SNAKA51 promotes ligand-binding to $a_5\beta_1$ integrin, and SNAKA51 binding to $a_5\beta_1$ integrin is increased in the presence of ligand

Using a solid-phase ligand-binding assay, the binding of biotinylated FnIII (6-10) (0.1 μ g/ml) to $\alpha_5\beta_1$ integrin was measured. a) Binding of FnIII (6-10) to directly coated placental $\alpha_5\beta_1$ integrin and anti-Fc captured recombinant integrin in the presence and absence of SNAKA51, 12G10, or K20. Statistical analysis was performed using a 2-tailed t-test in comparison to the no antibody control. * p<0.005, **p<0.0005. b) Dose-dependent binding of biotinylated SNAKA51 to K20 tethered placental $\alpha_5\beta_1$ integrin in the presence or absence of FnIII (6-10) (20 μ g/ml).



Figure 3. SNAKA51 only colocalizes with fibronectin fibers and primed β_1 integrin in fibrillar adhesions

Human fibroblasts were fixed and stained using indirect immunofluorescence for: a) a_5 integrin [SNAKA51 (left panels) or SNAKA52 (center panels) or mAb11 (right panels),

green], fibronectin (blue) and α_V integrin as a focal adhesion marker (L230, red); or b) Top panel β_1 integrin in a primed conformation (9EG7, red) and α_5 integrin (SNAKA51, green), middle panel total β_1 integrin (mAb11 red) and α_5 integrin (SNAKA51, green), bottom panel β_1 integrin in a primed conformation (9EG7, red) and total β_1 integrin (mAb11 green). Bar 20 μ m, insert bar 5 μ m.



Figure 4. The SNAKA51 epitope maps to the calf domains of the a5 subunit

a) Diagrammatic representation of the extracellular domains of the α_5 and β_1 integrin subunits indicating the position of truncations for the various recombinant constructs used in this study. b) The SNAKA51 binding site on the α_5 integrin subunit was mapped using a sandwich ELISA of Fc-tagged truncated or individual domains of recombinant integrin. The Fc-tagged protein was specifically captured using anti-Fc antibodies, and then binding of SNAKA51 to the integrin was measured using enzyme-conjugated anti-mouse secondary antibody.



Figure 5. Induction of priming by SNAKA51 is dependent on the β_1 leg and modulates β_1 A-domain ligand-binding ability

Binding of biotinylated FnIII (6-10) to either wild-type or ADMIDAS mutant (D138A) recombinant $\alpha_5\beta_1$ integrin, containing either a full-length or truncated ($\Delta 455$) β_1 subunit, was measured in the presence or absence of anti-integrin antibodies.



Figure 6. Clustering of SNAKA51-bound α_5 integrin induces integrin translocation out of focal adhesions and across the cell surface

Fibroblasts were plated on glass coverslips and incubated overnight in fibronectin-depleted medium with cycloheximide. Cells were labelled with antibody for 20 minutes, using SNAKA51 (a), SNAKA52 (b), or mAb11 (c). Unbound antibody was rinsed away, and the cells were either fixed ('no chasing'), or the cell-bound antibody was clustered with goat anti-mouse or anti-rat IgG and incubated for a further 30 minutes ('30' chasing') before fixation. For unclustered α_5 integrin, cells were stained with anti-mouse or anti-rat IgG (green). For clustered α_5 integrin, cells were stained with anti-goat IgG (green). In addition, all samples were stained with anti- α_V antibody L230 (red) as a focal adhesion marker. Bar 20 μ m.



Figure 7. SNAKA51 promotes fibronectin incorporation into the deoxycholate-insoluble matrix fraction

Human salivary gland cells (HSG) were treated overnight with a) a range of concentrations of biotinylated fibronectin, or biotinylated fibronectin (10 μ g/ml) and b) harvested at different time points, or c) with anti-integrin antibodies (25 μ g/ml), or d) a range of concentrations of anti- α_5 (SNAKA51). The cells were extracted with deoxycholate buffer, and the insoluble matrix fraction was collected and Western-blotted. Upper panel indicates biotinylated fibronectin incorporation into the insoluble matrix fraction. Lower panel indicates cytokeratin as internal loading controls for the HSG cells.



Figure 8. Model of transitions of $\alpha_5\beta_1$ integrin for fibrillar adhesion formation and fibronectin fibrillogenesis

a) $\alpha_5\beta_1$ integrin predicted domain structure, with a green box (calf domains) indicating the region that contains the SNAKA51 epitope. Blue balls represent cations bound to the MIDAS and ADMIDAS sites in the β A-domain.

b)

- 1. Inactive integrin is diffusely located on the cell surface.
- 2. $\alpha_5\beta_1$ located in focal adhesions expresses epitopes reporting a primed β_1 conformation (e.g. 9EG7). These integrins may or may not be fully bound by ligand.
- **3.** Integrin located at the distal edge of focal adhesions has additional SNAKA51 epitope expression. Clustering of this integrin promotes translocation.
- **4.** Ligated-clustered integrin translocates out of focal adhesions along the actin cytoskeleton, stretching extracellular fibronectin fibrils and driving fibrillogenesis.