

The integrin $\alpha_v\beta_3$ is a receptor for the latency-associated peptides of transforming growth factors β_1 and β_3

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The integrins $\alpha_v\beta_1$, $\alpha_v\beta_5$, $\alpha_v\beta_6$ and $\alpha_v\beta_8$ have all recently been shown to interact with the RGD motif of the latency-associated peptide (LAP β_1) of transforming growth factor β_1 (TGF β_1), with binding to $\alpha_v\beta_6$ and $\alpha_v\beta_8$ leading to TGF β_1 activation. Previously it has been suggested that the remaining α_v integrin, $\alpha_v\beta_3$, does not interact with LAP β_1 . However, here we show clearly that $\alpha_v\beta_3$ does indeed interact with the LAP β_1 RGD motif. This interaction is similar to other $\alpha_v\beta_3$ ligands in terms of the cations required for adhesion, the concentrations of LAP β_1 required for binding and the ability of a small-molecule inhibitor of $\alpha_v\beta_3$, SB223245, to block the interaction. Using glutathione S-transferase fusion proteins we have mapped a minimal integrin-

binding loop in LAP β_1 and then used this approach to probe the integrin-binding properties of the equivalent loops in LAP β_2 and LAP β_3 . We show that the RGD motif of LAP β_3 also interacts with $\alpha_v\beta_3$, in addition to $\alpha_v\beta_6$, $\alpha_v\beta_1$ and $\alpha_v\beta_5$, whereas the corresponding loop in LAP β_2 does not interact with these integrins. These observations therefore correct a previously reported inaccuracy in the literature. Furthermore, they are important as they link $\alpha_v\beta_3$ and TGF β , which may have implications in cancer and a number of inflammatory and fibrotic diseases where expression of both proteins has been documented.

Key words: adhesion, binding, cell, fibronectin, vitronectin.

INTRODUCTION

The transforming growth factor β (TGF β) family of cytokines affect a variety of cellular processes, including cell proliferation, extracellular-matrix synthesis, integrin expression, immune function and development [1–4]. The mammalian family consists of three members, namely TGF β_1 , TGF β_2 and TGF β_3 , which are secreted as heterotrimeric complexes derived from two genes. Each TGF β gene encodes a protein of 390–414 amino acids which is processed into two polypeptide chains, a 249–282-amino-acid N-terminal subunit and a 112-amino-acid C-terminal subunit. The C-terminal subunit forms the active TGF β cytokine, which is a 25 kDa homodimer. The N-terminal subunit is known as the latency-associated protein (LAP) and forms a homodimer of approx. 90 kDa. Each LAP dimer forms a non-covalent complex with a TGF β dimer termed the small latent complex (SLC), retaining the TGF β cytokine in an inactive conformation. The SLC is usually complexed with a protein called the latent TGF β -binding protein (LTBP), encoded by a separate gene, forming a large latent complex (LLC), predominantly found in the extracellular matrix. The sequestering of TGF β into these inactive complexes provides a mechanism for the fine regulation of the many biological effects mediated by these cytokines [3,4].

There has been much interest in delineating the mechanisms which liberate TGF β activity from the latent complexes. Both proteolytic and non-proteolytic mechanisms for activating latent TGF β_1 have been described. Proteases implicated in TGF β_1 activation include plasmin [5,6], urokinase-type and tissue-type plasminogen activators [7,8], matrix metalloproteases 2 and 9

(MMP-2 and MMP-9) [9] and cathepsin [10], which elicit effects by proteolytic degradation of LAP β_1 . Non-proteolytic activation mechanisms involve interactions with LAP β_1 , inducing a conformational change and exposing the receptor-binding site in TGF β_1 . Both thrombospondin 1 [11–13] and the integrins $\alpha_v\beta_6$ [14], $\alpha_v\beta_1$ [15], $\alpha_v\beta_8$ [16] and, weakly, $\alpha_v\beta_5$ [15] bind to LAP β_1 , and the binding of thrombospondin 1, $\alpha_v\beta_6$, or $\alpha_v\beta_8$ to LAP β_1 has been shown to result in TGF β_1 activation *in vitro*. The thrombospondin and $\alpha_v\beta_6$ mechanisms have been further validated *in vivo* by analysis of thrombospondin 1- [17] and β_6 - [18,19] knockout mice, which show features that may be attributable to a loss of TGF β_1 activity.

Both LAP β_1 and LAP β_3 contain RGD motifs, whereas LAP β_2 does not. As LAP β_1 has been shown to bind to a number of RGD-dependent integrins (all members of the α_v -containing family except $\alpha_v\beta_3$) [14–16], we examined whether it bound to two other RGD-dependent integrin receptors, $\alpha_5\beta_1$ and $\alpha_v\beta_3$. This analysis clearly showed that the remaining α_v -containing integrin, $\alpha_v\beta_3$, bound to LAP β_1 , despite evidence in the literature to the contrary [15]. We demonstrate that the binding of LAP β_1 by $\alpha_v\beta_3$ is RGD-dependent, and has similar characteristics to other $\alpha_v\beta_3$ ligands; namely the cations required for adhesion, the concentrations of LAP β_1 required for binding and the ability of a small-molecule inhibitor of $\alpha_v\beta_3$, SB223245, to block the interaction. We also show that $\alpha_v\beta_3$, $\alpha_v\beta_6$, $\alpha_v\beta_1$ and $\alpha_v\beta_5$ bind to LAP β_3 in an RGD-dependent manner, but not to LAP β_2 , presumably due to the absence of an RGD site. These data are important, particularly given that $\alpha_v\beta_3$ has previously been discounted as a receptor for LAP β_1 [15]. Biologically, the $\alpha_v\beta_3$ interactions with LAP β_1 and LAP β_3 may be clinically important

Abbreviations used: GST, glutathione S-transferase; LAP, latency-associated peptide; MMP, matrix metalloprotease; SLC, small latent complex; TGF β , transforming growth factor β .

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in several diseases including cancer, inflammatory and fibrotic diseases.

EXPERIMENTAL

Materials

Antibodies were obtained from Chemicon (LM609, P1F6, 10D5), Immunotech (4B4, SAM1, AMF7, 69-6-5) and Sigma (MOPC21, UPC10). The $\alpha_v\beta_3/\alpha_v\beta_5$ inhibitor SB223245 [20], the $\alpha_v\beta_1$ inhibitor BIO1211 [21] and the $\alpha_v\beta_6$ inhibitor GW603365A (acetyl-RTDLSLRT-NH₂) [22] were obtained from within GlaxoSmithKline. All oligonucleotides were obtained from Life Technologies. Fibrinogen was purchased from Calbiochem and baculovirus-expressed full-length LAP β_1 was from Sigma. Vitronectin was purified from out-dated plasma as described in [23,24], whereas $\alpha_v\beta_3$ protein was obtained from Chemicon. The K562, A549 and A375M cell lines were originally obtained from the A.T.C.C., whereas DX3 human melanoma cells were available within GlaxoSmithKline [25]. K562-derived cells were cultured in a 1:1 mixture of RPMI 1640 (Hepes modification; Life Technologies)/Dulbecco's modified Eagle's medium (Hepes modification; Sigma), supplemented with L-glutamine (Life Technologies) and 10% fetal calf serum (Gibco). Geneticin (Life Technologies) was added to 1 mg/ml for all transfectants. A375M and A549 cells were cultured in RPMI 1640 (Hepes modification), whereas DX3 cells were cultured in Dulbecco's modified Eagle's medium with high glucose (Life Technologies), all supplemented with L-glutamine and 10% fetal calf serum.

Cloning of integrin subunits

The integrin chains α_v , β_3 , β_5 and β_6 were all cloned by reverse transcriptase PCR of RNA prepared from tissue sources using the Promega RNAagents total RNA isolation system. Oligo(dT) chromatography was subsequently used to purify polyadenylated RNA from the total RNA. α_v , β_3 and β_5 were cloned using RNA from DX3 melanoma cells, and β_6 from HT29 colon carcinoma cells. The primers for the α_v and β_6 cDNAs contained a *Bam*HI restriction site proximal to the initiation codon and an *Eco*RI restriction site distal to the stop codon. For β_3 , a 5' *Hind*III site was included together with a 3' *Xba*I site. For β_5 , *Eco*RI sites were included at both the 5' and 3' ends. For sequence verification, the α_v , β_3 and β_5 cDNAs were cloned into pSP64 (Promega), whereas the β_6 cDNA was inserted into PCR-SCRIPT (Stratagene). All sequences were in agreement with the EMBL/Genbank accession numbers for these cDNAs.

Generation of K562 cell lines stably expressing $\alpha_v\beta_3$, $\alpha_v\beta_5$ or $\alpha_v\beta_6$

Integrin subunit cDNAs were cloned into pCDNA3 (Invitrogen) using the restriction sites incorporated during cloning. The pcDNA3- α_v plasmid was co-transfected with pcDNA3- β_3 , pcDNA3- β_5 or pcDNA3- β_6 into 2×10^6 K562 cells using LIPOFECTAMINETM plus (Life Technologies) according to the manufacturer's instructions. After a 48 h recovery Geneticin was added to the culture medium at 1 mg/ml to select for stably transfected cells. K562- $\alpha_v\beta_3$ and K562- $\alpha_v\beta_5$ transfectants both adhered to the flask plastic during selection, probably due to vitronectin and other matrix components in the serum coating the plastic. Analysis of K562- $\alpha_v\beta_3$ adherent cells by flow cytometry [26] showed high $\alpha_v\beta_3$ expression on this population (Table 1), which was then used for all experiments. K562- $\alpha_v\beta_5$ adherent cells showed biphasic $\alpha_v\beta_5$ expression on flow-cytometric analysis, so were further selected by clonal dilution to yield single clones with reproducible $\alpha_v\beta_5$ expression, of which

Table 1 Analysis of integrin expression on cell lines by flow cytometry

Antibody clones used were 4B4 (β_1), AMF7 (α_v), LM609 ($\alpha_v\beta_3$), P1F6 ($\alpha_v\beta_5$), 10D5 ($\alpha_v\beta_6$) and SAM-1 (α_5), with a goat anti-mouse IgG secondary antibody labelled with Alexa 488, as described in [26]. Data are presented as mean relative fluorescence values. Antibody isotype controls used were MOPC21 (mouse IgG₁) and UPC10 (mouse IgG_{2a}), and were given mean fluorescence values of 1.

Cell line	Mean relative fluorescence					
	β_1	α_v	$\alpha_v\beta_3$	$\alpha_v\beta_5$	$\alpha_v\beta_6$	α_5
K562	13	2.6	1.2	0.95	0.83	11
K562- $\alpha_v\beta_3$	14	106	93	0.96	2.2	13
K562- $\alpha_v\beta_5$	12	6.2	0.91	4.9	0.69	7.3
K562- $\alpha_v\beta_6$	21	8.3	0.71	0.97	8.2	19
A549	352	90	2.2	28	0.68	15
A375M	178	83	85	30	0.74	23
DX3	397	86	45	46	0.75	34

one clone (clone 19) is shown for all experiments. Two further clones showed similar results in adhesion assays [clones 6 and FF (results not shown)]. K562- $\alpha_v\beta_6$ cells in suspension displayed low $\alpha_v\beta_6$ expression by flow-cytometric analysis, so were also selected by clonal dilution with one clone (clone 3) shown for all experiments. A further clone (clone 22) showed similar results in adhesion assays (results not shown). All three cell types were characterized, and displayed expected ligand-binding activities relative to the integrin repertoire expressed at the cell surface (results not shown). Flow cytometry analysis of all cell lines used in this study is shown in Table 1. Further analysis examined $\alpha_v\beta_1$ expression on the transfected lines by immunoprecipitation and Western blot analysis (as no $\alpha_v\beta_1$ -heterodimer-specific antibody is available). This showed the presence of α_v in the β_1 immunoprecipitate from the K562- $\alpha_v\beta_5$, but not the K562- $\alpha_v\beta_3$ or K562- $\alpha_v\beta_6$ cell lines (results not shown), indicating the presence of the $\alpha_v\beta_1$ integrin heterodimer in K562- $\alpha_v\beta_5$ cells only.

Cloning and expression of matrix proteins

Glutathione S-transferase (GST) fusion proteins corresponding to the RGD cell-adhesion domains of tenascin (amino acids 802–891) [27], fibronectin (fibronectin type III repeats 8–10, amino acids 1266–1540), LAP β_1 (amino acids 242–252), LAP β_2 (amino acids 259–269) and LAP β_3 (amino acids 259–269) were generated. Tenascin and fibronectin fragments were generated by reverse-transcriptase PCR from HT29 and DX3 total RNA respectively; other constructs were generated by oligonucleotide insertion. All constructs were cloned into pGEX-2T via the *Bam*HI/*Eco*RI sites except fibronectin, which had a 5' *Xba*I site. Details of all the oligonucleotides used in this study are available from S.B.L. on request. Recombinant proteins were over-expressed in, and purified from, *Escherichia coli* XL1-Blue cells according to standard protocols and as described previously [26], and stored in PBS at -70°C . Protein concentrations were determined using the Coomassie plus protein assay reagent (Pierce). The integrity and purity of each preparation were analysed by SDS/PAGE and shown to be greater than 95% for all proteins except GST-tenascin, which was subject to some apparent proteolytic fragmentation, but was still $>70\%$ pure intact protein. Most proteins were also verified by MS.

Adhesion assays

These assays were performed broadly as described in [26]. Either 3×10^5 cells (all K562-derived cells) or 1×10^5 cells (DX3, A549

and A375M cells) were used per well. Desired cations (either 2 mM $MgCl_2$ or 0.5 mM $MnCl_2$ as indicated in the Figure legends) and other additions were made as indicated. Cells were visualized and quantified using Crystal Violet staining (Sigma). Each Figure is representative of at least three identical experiments and all results are presented as means \pm S.D. from duplicate determinations. Data were fitted to a four-parameter logistic equation for IC_{50} determinations.

ORIGEN® assays

Epoxy beads (Dynal) were pre-coated with $\alpha_v\beta_3$ protein as per the manufacturer's recommendations. Protein ligands were biotinylated with NHS-Biotin (Perbio) as per the manufacturer's protocol. Streptavidin tag was obtained from IGEN. The assays were performed in a buffer containing 25 mM Hepes, pH 7.4, 150 mM NaCl, 1 mM $MnCl_2$, 1 mM $MgCl_2$, 0.1 mM $CaCl_2$, 0.1% Tween-20 and 0.5% BSA in a 150 μ l total well volume. Plates were shaken for 4 h and read on an IGEN M8 Analyser. Each Figure is representative of at least three identical experiments and all results are presented as means \pm S.D. from duplicate determinations. Data were fitted to a four-parameter logistic equation for IC_{50} determinations.

RESULTS

$\alpha_v\beta_3$ interacts with LAP β_1

As LAP β_1 has been shown to bind to a number of RGD-dependent integrins (all members of the α_v -containing family except $\alpha_v\beta_3$) [14–16], we examined whether it bound to two other RGD-dependent integrin receptors, $\alpha_5\beta_1$ and $\alpha_v\beta_3$. The study of specific integrin–ligand interactions is often complicated by the presence of multiple integrins on cell lines. To alleviate this problem we have used K562 cells, which only express the $\alpha_5\beta_1$ integrin endogenously. Wild-type K562 cells were transfected with $\alpha_v\beta_3$ (K562- $\alpha_v\beta_3$), $\alpha_v\beta_5$ (K562- $\alpha_v\beta_5$) and $\alpha_v\beta_6$ (K562- $\alpha_v\beta_6$) to generate stable cell lines. Figure 1 shows that in the presence of magnesium ions K562- $\alpha_v\beta_3$ and K562- $\alpha_v\beta_6$ cells bound well to LAP β_1 , and the adhesion was inhibited by specific function-blocking antibodies to the transfected integrin heterodimer, demonstrating adhesion via $\alpha_v\beta_3$ and $\alpha_v\beta_6$ integrins respectively. K562- $\alpha_v\beta_5$ cells only adhered well to LAP β_1 in the presence of manganese ions, and this binding was mediated by both $\alpha_v\beta_5$ and $\alpha_v\beta_1$ integrins, consistent with small amounts of $\alpha_v\beta_1$ present on the K562- $\alpha_v\beta_5$ cell line (results not shown). Untransfected K562 cells (K562-WT on Figure 1) failed to adhere to LAP β_1 in the presence of either magnesium or manganese ions, confirming that $\alpha_5\beta_1$ does not recognize LAP β_1 , despite it being an RGD-binding integrin. These results indicate that $\alpha_v\beta_3$ is a novel receptor for LAP β_1 , and the inhibition data with antibodies for each cell line suggest that the adhesion event is not the result of the transfected integrin(s) modulating the activity of the endogenous $\alpha_5\beta_1$ integrin ([26,28] and S. B. Ludbrook, unpublished work).

Comparison of LAP β_1 with other $\alpha_v\beta_3$ ligands

$\alpha_v\beta_3$ interacts with a wide variety of ligands, including fibrinogen [29,30], vitronectin [31–33], tenascin [27], fibronectin [34], osteopontin [35], L1 [36], cyr61 [37] and connective tissue growth factor [38]. Ligand recognition by $\alpha_v\beta_3$ can be mediated via both RGD [29,32] and non-RGD [30] motifs; in addition, the β_3 -containing integrins $\alpha_v\beta_3$ and $\alpha_{11b}\beta_3$ contain a high-affinity calcium-binding site on the β_3 subunit that down-regulates ligand

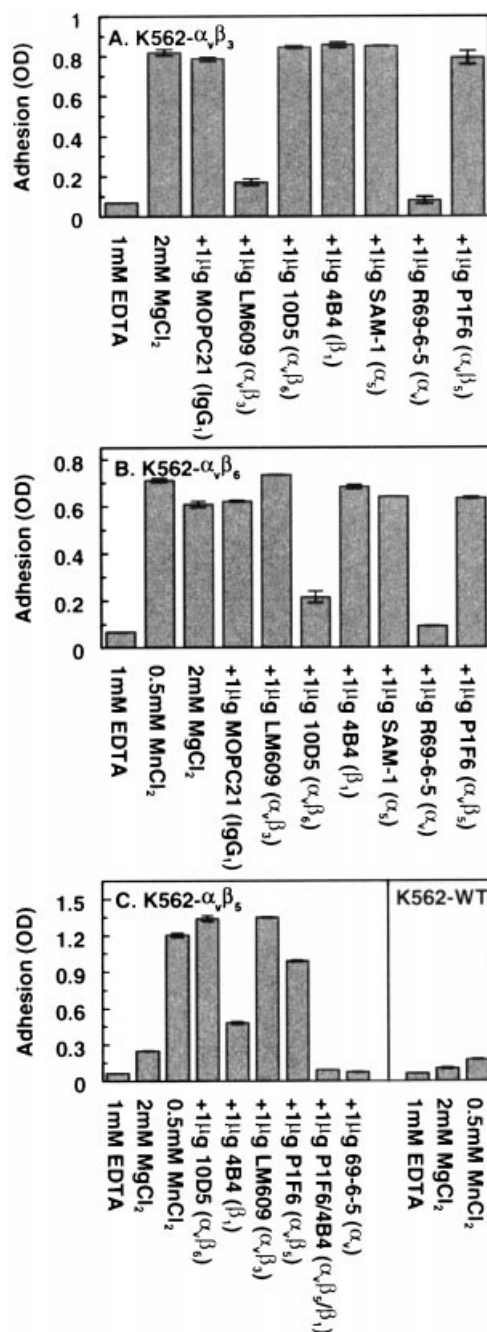


Figure 1 Adhesion of K562- $\alpha_v\beta_3$, K562- $\alpha_v\beta_5$, K562- $\alpha_v\beta_6$ and wild-type K562 cells to LAP β_1

K562- $\alpha_v\beta_3$ (A), K562- $\alpha_v\beta_6$ (B) and K562- $\alpha_v\beta_5$ and wild-type K562-WT (C) cells were allowed to attach to wells coated with 0.5 μ g of LAP- β_1 under conditions as indicated (antibodies added at 1 μ g/well). Antibody-inhibition data were generated using 2 mM $MgCl_2$ (K562- $\alpha_v\beta_3$ and K562- $\alpha_v\beta_6$) or 0.5 mM $MnCl_2$ (K562- $\alpha_v\beta_5$) as the cation.

recognition [39,40]. Therefore, to examine the relevance of the interaction of K562- $\alpha_v\beta_3$ with LAP β_1 , and to determine whether the RGD sequence in LAP β_1 was responsible, three different experiments were performed. Firstly, to determine the amounts of ligands required to support adhesion via $\alpha_v\beta_3$, we compared the adhesion of K562- $\alpha_v\beta_3$ cells to LAP β_1 , vitronectin, fibrinogen, tenascin and a GST-LAP β_1 fragment fusion protein (LAP β_1

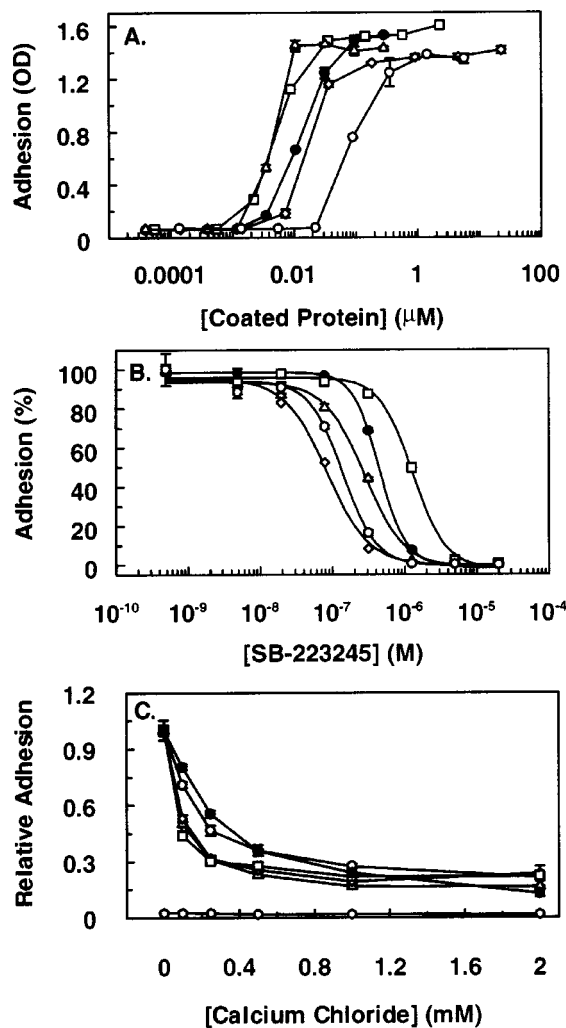


Figure 2 Comparison of LAP β_1 , fibrinogen, vitronectin and tenascin ligands for $\alpha_v\beta_3$

(A) K562- $\alpha_v\beta_3$ cells were allowed to attach to wells coated with a range of concentrations of LAP β_1 (●), fibrinogen (◇), vitronectin (□), tenascin (△) and GST-LAP β_1 (○) in the presence of 2 mM MgCl₂. (B) K562- $\alpha_v\beta_3$ cells were allowed to attach to wells coated with LAP β_1 (0.5 μ g/well), fibrinogen (2 μ g), vitronectin (1 μ g), tenascin (0.5 μ g) or GST-LAP β_1 (0.5 μ g) (symbols as in A) in the presence of 2 mM MgCl₂. The dose–response relationship with SB223245 was tested. Calculated IC₅₀ values were LAP β_1 (442 nM), vitronectin (1283 nM), fibrinogen (83 nM), tenascin (279 nM) and GST-LAP β_1 (139 nM). (C) As panel (B) with the inclusion of GST (0.5 μ g; open hexagons). A dose–response relationship with CaCl₂ was performed. Adhesion levels were normalized, but were similar to those shown in (A), with adhesion to GST-coated wells of 0.074 ± 0.004 absorbance unit (i.e. no adhesion).

amino acids 242–252; ²⁴²GRRGDLATHG²⁵²) in a matrix titration experiment (Figure 2A). This analysis showed that both the full-length and GST fusion protein of LAP β_1 supported K562- $\alpha_v\beta_3$ cell adhesion at similar levels of protein coating to the other ligands used. Secondly, the interaction was analysed by testing the ability of a small-molecule inhibitor of $\alpha_v\beta_3$, SB223245 [20], to block the interaction of K562- $\alpha_v\beta_3$ cells with these ligands. SB223245 blocked $\alpha_v\beta_3$ –ligand interactions with a range of IC₅₀ values. It was most active against the $\alpha_v\beta_3$ –fibrinogen interaction (IC₅₀, 83 nM), relative to GST-LAP β_1 (139 nM), tenascin (279 nM), LAP β_1 (442 nM) and vitronectin (1283 nM; Figure 2B). Thirdly, we assessed the effects on adhesion to each of these ligands of increasing concentrations of CaCl₂. This

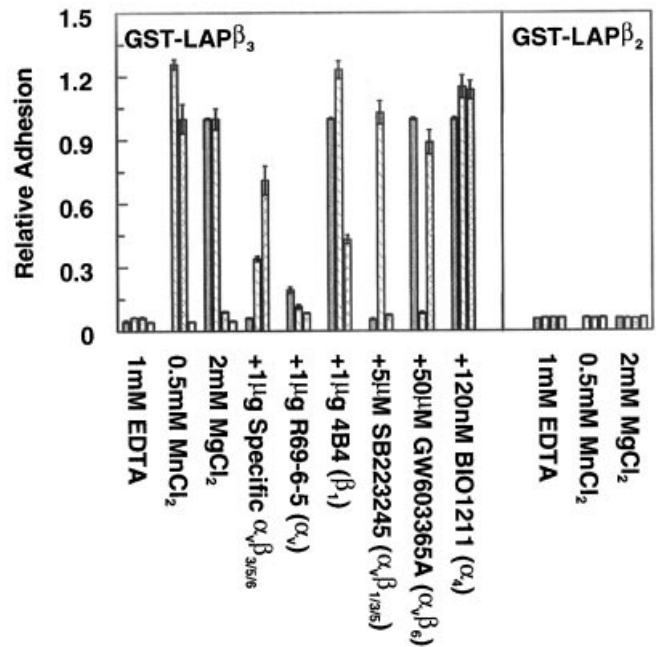


Figure 3 Adhesion of K562- $\alpha_v\beta_3$, K562- $\alpha_v\beta_5$, K562- $\alpha_v\beta_6$ and wild-type K562 cells to GST-LAP β_2 and GST-LAP β_3

K562- $\alpha_v\beta_3$ (shaded bars), K562- $\alpha_v\beta_6$ (hatched bars), K562- $\alpha_v\beta_5$ (cross-hatched bars) and wild-type K562-WT (white bars) cells were allowed to attach to wells coated with 0.5 μ g of GST-LAP β_3 amino acids 259–269 or GST-LAP β_2 amino acids 259–269 under conditions as indicated (antibodies added at 1 μ g/well). Antibody-inhibition data were generated using 2 mM MgCl₂ (K562- $\alpha_v\beta_3$ and K562- $\alpha_v\beta_6$) or 0.5 mM MnCl₂ (K562- $\alpha_v\beta_5$) as the cation. Specific $\alpha_v\beta_3/5/6$ indicates the addition of clone LM609 (anti- $\alpha_v\beta_3$), P1F6 (anti- $\alpha_v\beta_5$) or 10D5 (anti- $\alpha_v\beta_6$) to the specific K562-transfected cell line. SB223245 inhibits $\alpha_v\beta_1$, $\alpha_v\beta_3$ and $\alpha_v\beta_5$ at the concentration used. The complete absence of a bar indicates no test under that condition.

showed that CaCl₂ markedly inhibited K562- $\alpha_v\beta_3$ cell adhesion to all ligands tested (tenascin, fibrinogen, LAP β_1 , GST-LAP β_1 , vitronectin) in a dose-dependent manner in agreement with the literature [39,40]. Collectively these data demonstrate that $\alpha_v\beta_3$ recognizes LAP β_1 with similar characteristics to those observed for other ligands such as fibrinogen, vitronectin and tenascin. By this analysis, of these four ligands, vitronectin is the best ligand for $\alpha_v\beta_3$ followed by LAP β_1 and tenascin, with fibrinogen being the weakest ligand. In addition, as the GST-LAP β_1 protein, which only contains an 11-amino acid fragment of LAP β_1 surrounding the RGD motif, has similar adhesion properties to intact LAP β_1 , it is likely that this RGD-containing region of LAP β_1 is the sole determinant of $\alpha_v\beta_3$ binding.

Integrin binding to LAP β_2 and LAP β_3

The majority of studies analysing the cell-adhesion properties of TGF β LAPs have focused on TGF β_1 , and hence LAP β_1 [14,15]. However, LAP β_3 also contains an RGD motif in the same region, whereas LAP β_2 contains SGD. Therefore we predicted that, like LAP β_1 , LAP β_3 may interact with RGD-dependent integrins whereas LAP β_2 should not. Given the success of our minimized adhesion peptide approach with LAP β_1 , we adopted a similar strategy to test the adhesion properties of LAP β_3 and LAP β_2 . The same regions of LAP β_2 (amino acids 259–269 ²⁵⁹YTS²⁶⁹GDQKTIKS²⁶⁹) and LAP β_3 (amino acids 259–269

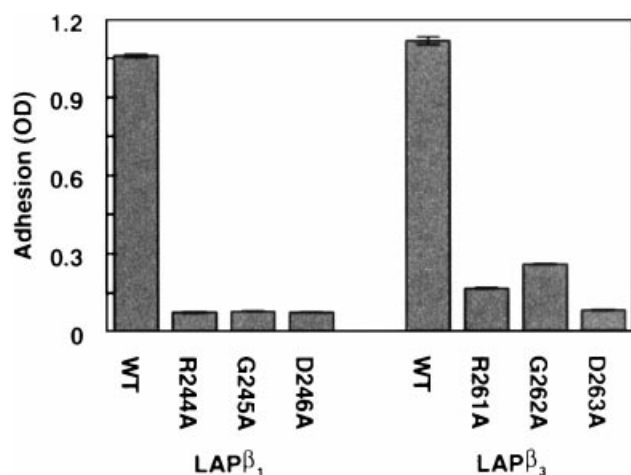


Figure 4 Importance of the LAP β_1 and β_3 RGD sequences for K562- $\alpha_v\beta_3$ binding

K562- $\alpha_v\beta_3$ cells were allowed to attach, in the presence of 2 mM MgCl₂, to wells coated with 0.5 μ g of either wild-type GST-LAP β_1 or GST-LAP β_3 proteins (amino acids 242–252 and 259–269 respectively), or site-specific mutants in which individual residues of the RGD motif had been mutated to alanine.

²⁵⁹HGRGDLGRLKK²⁶⁹) were expressed as GST fusion proteins and tested for their ability to support adhesion of K562- $\alpha_v\beta_3$, K562- $\alpha_v\beta_5$ and K562- $\alpha_v\beta_6$ cells. The GST-LAP β_3 protein (amino acids 259–269) had similar adhesion properties to LAP β_1 in that it supported adhesion of K562- $\alpha_v\beta_3$ and K562- $\alpha_v\beta_6$ cells in the presence of magnesium via $\alpha_v\beta_3$ and $\alpha_v\beta_6$ respectively (Figure 3). K562- $\alpha_v\beta_5$ cells required manganese to further activate the integrins and adhered via a mixture of $\alpha_v\beta_5$ and $\alpha_v\beta_1$ integrins, whereas wild-type K562 cells failed to adhere under either cation condition. As predicted, there was no adhesion of any cell type tested to the GST-LAP β_2 protein (amino acids 259–269; Figure 3).

LAP β_1 and LAP β_3 RGD mutants fail to support $\alpha_v\beta_3$ binding

To confirm the RGD-dependence of integrin binding to LAP β_1 , GST fusion proteins were generated where Arg₂₄₄, Gly₂₄₅ or Asp₂₄₆ were separately exchanged for alanine and the mutant proteins tested for their ability to support adhesion of K562- $\alpha_v\beta_3$ cells (Figure 4). Mutation of any of the three RGD residues abolished binding, clearly demonstrating that $\alpha_v\beta_3$ interacts with LAP β_1 at the RGD motif. Similar results were obtained for LAP β_3 RGD mutant proteins (Figure 4). The LAP β_1 and LAP β_3 mutant proteins also failed to support adhesion of either K562- $\alpha_v\beta_6$ or K562- $\alpha_v\beta_5$ cells (results not shown).

The interactions between $\alpha_v\beta_3$ and LAP β_1/β_3 are direct

Cell-adhesion data strongly suggested that the interactions between $\alpha_v\beta_3$ and LAP β_1/β_3 were direct, as the interaction was cation-dependent, susceptible to inhibition by specific integrin-blocking antibodies, RGD-dependent and inhibited by CaCl₂ addition. To confirm this we assessed the interaction between purified $\alpha_v\beta_3$ protein and LAP β_1 , β_2 and β_3 using ORIGIN[®] assay technology. Firstly we tested whether $\alpha_v\beta_3$ protein could interact with the GST fusion proteins for LAP β_1 , β_2 and β_3 . A cation-dependent signal was observed with both LAP β_1 and

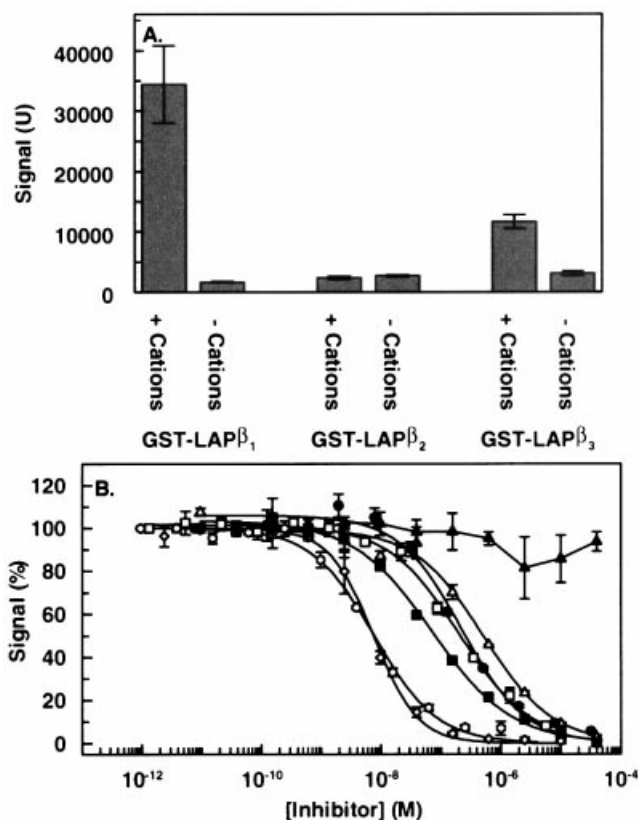


Figure 5 Direct $\alpha_v\beta_3$ protein–protein interaction with GST-LAP β_1/β_3

(A) $\alpha_v\beta_3$ protein (0.4 nM), immobilized to beads, was incubated with 10 nM biotinylated GST-LAP β_1 , GST-LAP β_2 and GST-LAP β_3 proteins, together with 20 nM streptavidin tag, in ORIGIN assay buffer. The assay was performed in either the presence of cations (1 mM MnCl₂, 1 mM MgCl₂ and 0.1 mM CaCl₂) or in their absence (without cations but with 10 mM EDTA). The electrochemiluminescent signal represents the signal obtained in the instrument on binding. (B) Competing protein and small-molecule ligands were tested for inhibitory potencies in the $\alpha_v\beta_3$ –LAP β_1 ORIGIN assay (0.4 nM $\alpha_v\beta_3$, 10 nM biotinylated GST-LAP β_1 , 20 nM streptavidin tag) in a dose–response experiment. The following competitors were used: GST-tenascin (Δ ; IC₅₀, 511 nM), GST-fibronectin (open hexagons; 7.6 nM), vitronectin (\square ; 201 nM), GST-LAP β_1 (\bullet ; 222 nM), GST-LAP β_2 (\blacktriangle ; > 40 μ M), GST-LAP β_3 (\blacksquare ; 72 nM) and SB 223245 (\diamond ; 7.5 nM).

LAP β_3 but not LAP β_2 (Figure 5A), demonstrating specific direct binding between $\alpha_v\beta_3$ and LAP β_1/β_3 . Secondly, as the amplitude of the signal at a given concentration of ligand is a poor indicator of differences in binding affinities (as it is influenced by ligand biotinylation levels and site of biotinylation), and to compare LAP β_1 and β_3 with other $\alpha_v\beta_3$ ligands in this cell-free system, we tested $\alpha_v\beta_3$ ligands as competing proteins in the $\alpha_v\beta_3$ –LAP β_1 ORIGIN[®] assay (Figure 5B). Fibronectin was the most active competitor by this analysis (IC₅₀, 7.6 nM) and was of similar potency to the SB223245 small-molecule inhibitor (7.5 nM). The remaining ligands tested all had IC₅₀ values in a relatively narrow range, where LAP β_1 (222 nM) and LAP β_3 (72 nM) compared well with vitronectin (201 nM) and tenascin (511 nM), and LAP β_2 was inactive (> 40 μ M). These protein-based studies demonstrate firstly that the $\alpha_v\beta_3$ interaction with LAP β_1 and LAP β_3 is direct, secondly that LAP β_1 and LAP β_3 have similar binding affinities for the isolated $\alpha_v\beta_3$ receptor as other $\alpha_v\beta_3$ ligands, and thirdly that $\alpha_v\beta_3$ interacts with soluble LAP β_1 and LAP β_3 , a form that may be encountered *in vivo*.

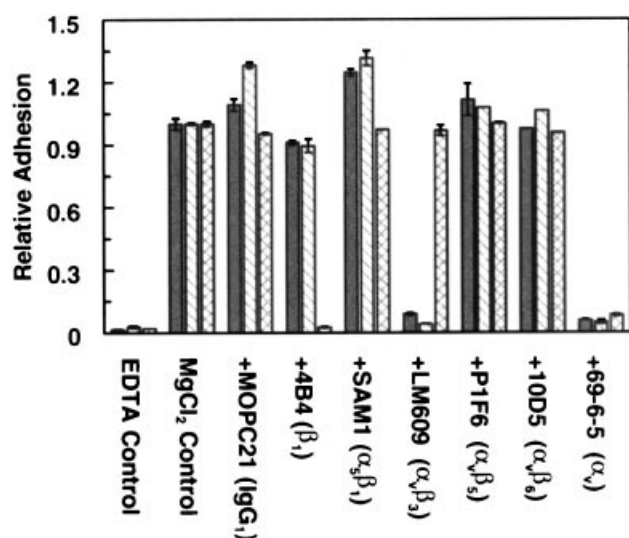


Figure 6 Adhesion characteristics of DX3, A375M and A549 cell lines to LAPβ₁.

DX3 (shaded bars), A375M (hatched bars) or A549 (cross-hatched bars) cells were allowed to attach to wells coated with 0.5 μg of LAPβ₁. Adhesion occurred in 2 mM MgCl₂ or with the antibodies that are indicated under the histogram (1 μg/well).

Native α_vβ₃-expressing cell lines interact with LAPβ₁ via α_vβ₃

The K562-α_vβ₃ cell line used in this study expresses very high levels of α_vβ₃ integrin (see Table 1), relative to the levels of the α_vβ₅ and α_vβ₆ integrins on the K562-α_vβ₅ and K562-α_vβ₆ cell lines respectively. It is possible that the adhesion observed was an artifact of the extremely high levels of α_vβ₃ on the K562-α_vβ₃ cell line. To address this, we investigated the binding profile of cell lines reported to endogenously express the α_vβ₃ integrin to LAPβ₁ and LAPβ₃. The cell lines used were the human melanoma cell lines DX3 [25] and A375M [41], and the epithelial lung carcinoma cell line A549 [42,43]. Analysis of integrin expression on these cell lines showed expression of α_vβ₃ on the DX3 and A375M cells, but negligible levels on the A549 cells (Table 1). Analysis of the adhesion properties of these three cell lines to LAPβ₁, using specific integrin-blocking antibodies, showed that A375M and DX3 cells adhered to LAPβ₁ via α_vβ₃, whereas A549 cells adhered to LAPβ₁ via α_vβ₁ (Figure 6). These results demonstrate firstly that cell lines which endogenously express the α_vβ₃ integrin (DX3 and A375M) do adhere to LAPβ₁ via α_vβ₃. Secondly, the A549-binding data to LAPβ₁ agree with the previous report as being mediated via α_vβ₁ [15], with the absence of an α_vβ₃ component being due to negligible levels of α_vβ₃ being expressed by this cell line. Similar results were obtained with cell adhesion to GST-LAPβ₁ and GST-LAPβ₃ (results not shown).

DISCUSSION

Here we demonstrate that α_vβ₃ interacts with LAPβ₁. This increases the number of integrins known to bind LAPβ₁ to five, the entire α_v family (α_vβ₁, α_vβ₃, α_vβ₅, α_vβ₆ and α_vβ₈). K562-α_vβ₃ cells bind to LAPβ₁ with similar characteristics to other known α_vβ₃ ligands, such as tenascin, fibrinogen and vitronectin, in terms of the ligand concentrations required for adhesion, the cations required for binding and the ability of a small-molecule inhibitor of α_vβ₃ to block the interaction. Like the other α_v

integrin heterodimers, α_vβ₃ interacts with LAPβ₁ via the RGD motif, as mutant proteins in which any of these residues are mutated to alanine fail to support adhesion of K562-α_vβ₃ cells. We have also characterized the integrin-binding properties of the corresponding regions in LAPβ₂ and LAPβ₃. We show that LAPβ₃ is also a ligand for α_vβ₃, α_vβ₆, α_vβ₅ and α_vβ₁, where binding also occurs through the RGD motif. Conversely, LAPβ₂ is not a ligand for any of these integrins, presumably due to the lack of an RGD motif, which has been replaced by SGD. Further analysis of the interaction of α_vβ₃ with LAPβ₁ and LAPβ₃ demonstrated that, in a protein-protein assay, α_vβ₃ interacted with LAPβ₁ and β₃, but not β₂, in a cation-dependent manner. This confirmed not only that the α_vβ₃-LAPβ₁/β₃ interactions are direct, but also that α_vβ₃ can bind to LAPβ₁/β₃ proteins when they are presented in a soluble form, a form that can be encountered *in vivo* in addition to matrix-associated LAPβ₁/β₃. During the preparation of this manuscript, a report demonstrated that α_vβ₆ interacts with LAPβ₃ [44]. Our data confirm this, and also establish that the LAPβ₃ RGD motif possesses similar integrin-recognition properties to the LAPβ₁ RGD sequence. Therefore α_v integrins interact with both LAPβ₁ and LAPβ₃, and have the potential to modulate the localization and possibly activation of TGFβ₁ and TGFβ₃, but not directly TGFβ₂.

Although these data clearly show an interaction between α_vβ₃ and LAPβ₁ and LAPβ₃, a previous study [15] excluded α_vβ₃ as a receptor for LAPβ₁, as A549 cells, which reportedly express α_vβ₃ [42,43], adhered to LAPβ₁ via α_vβ₁. We examined the binding of the DX3, A375M and A549 cell lines, which are all reported to endogenously express α_vβ₃, to LAPβ₁ and LAPβ₃. All three cell lines adhered to LAPβ₁ and LAPβ₃. However, analysis using specific integrin-blocking antibodies showed that DX3 and A375M cells interacted via α_vβ₃, whereas A549 cells interacted via α_vβ₁, as reported in [15]. Flow cytometry analysis of these three cell lines clearly showed high expression levels of the α_vβ₃ integrin on both DX3 and A375M cell lines, but negligible levels on the A549 cells. These data demonstrate that cell lines which endogenously express the α_vβ₃ integrin use α_vβ₃ to interact with LAPβ₁ and LAPβ₃. They also indicate that A549 cells do not use α_vβ₃ to adhere to LAPβ₁ and LAPβ₃ because of very low α_vβ₃ expression levels. The previous report [15] did not show α_vβ₃ expression levels on the A549 cells used and furthermore α_vβ₃ binding to known ligands was not confirmed. However, it is clear in the literature that A549 cells have been shown to adhere to fibrinogen via α_vβ₃ [42,43]. Our A549 cells adhered very poorly to fibrinogen in an α_vβ₁-dependent manner (results not shown), suggestive of either α_vβ₁ being the fibrinogen receptor [45] or the adhesion resulting from the low levels of contaminating fibronectin in the fibrinogen preparation.

α_vβ₃ plays an important role in a variety of physiological and patho-physiological processes, including tumour angiogenesis, rheumatoid arthritis and a number of inflammatory and repair processes [46]. This is reflected by the extensive repertoire of known α_vβ₃ ligands, which include the matrix proteins vitronectin [31–33], fibronectin [34] and tenascin [27], and the cytokines osteopontin [35], cyr61 [37] and connective tissue growth factor [38]. The interactions between α_vβ₃ and distinct ligands may drive different aspects of these biological processes. For example, the α_vβ₃-osteopontin interaction has been shown to be important in mediating the Th1 response via the regulation of interleukin-12 expression from macrophages [47]. In many diseases involving α_vβ₃, up-regulation of TGFβ protein or activity has also been described, such as vascular disorders [48], diabetic retinopathy [49], scleroderma [50] and rheumatoid arthritis [51,52]. It will therefore be interesting to determine what role, if any, the α_vβ₃ interactions with LAPβ₁ and LAPβ₃ play in these diseases.

The interaction between $\alpha_v\beta_3$ and LAP β_1 is important, as it provides a mechanism through which TGF β_1 and TGF β_3 can be localized to the surface of a number of cell types. For example, $\alpha_v\beta_3$ is expressed on angiogenic/activated endothelium, fibroblasts, macrophages, T-cells and smooth-muscle cells [46]. This interaction may be sufficient to activate TGF β_1 and TGF β_3 via a conformational change in LAP, as shown previously for $\alpha_v\beta_6$ [14]. Alternatively, binding to $\alpha_v\beta_3$ may localize LAP β_1 and LAP β_3 to a proteolytically rich environment at the cell surface, resulting in TGF β activation. A similar function has been ascribed to the binding of LAP β_1 to thrombospondin [53] and the mannose 6-phosphate receptor [54]. Moreover, $\alpha_v\beta_8$ has recently been shown to generate TGF β_1 activity by localizing the LAP β_1 -TGF β_1 SLC to the cell surface, thereby permitting membrane-type 1-MMP proteolytic cleavage of LAP β_1 to liberate the TGF β_1 cytokine [16]. Interestingly, $\alpha_v\beta_3$ expression is associated with enhanced cell-surface proteolytic activity by MMP-2 [55], for which LAP β_1 has been shown to be a substrate [9], thereby presenting a potential mechanism to generate TGF β_1 activity from the interaction described here. In addition, the interaction of LAP β_1 and LAP β_3 with $\alpha_v\beta_3$ may directly initiate $\alpha_v\beta_3$ -specific intracellular signalling events commonly associated with integrin-ligand ligation [56]. All of these possibilities are currently the subject of an ongoing study, particularly whether $\alpha_v\beta_3$ either directly activates TGF β by a conformational change in LAP β_1/β_3 , or indirectly via protease activity. For the $\alpha_v\beta_6$ integrin there is clear evidence of the importance of TGF β activity regulation from observations of β_6 -knockout mice [18,19]. There is no such evidence from similar studies in $\alpha_v\beta_3$ -knockout mice [57,58]. However, for the $\alpha_v\beta_3$ -knockout mice the situation is complex as deletion of the β_3 subunit results in the absence of two integrins, $\alpha_v\beta_3$ and the platelet integrin $\alpha_{IIb}\beta_3$. Likewise, deletion of the α_v subunit results in the loss of five integrins, $\alpha_v\beta_1$, $\alpha_v\beta_3$, $\alpha_v\beta_5$, $\alpha_v\beta_6$ and $\alpha_v\beta_8$. Individuals with the α_v deletion fail to reach birth due to a placental defect [59], but the data would clearly fail to suggest $\alpha_v\beta_3$ activation of TGF β activity, in part because $\alpha_v\beta_6$ is also deleted. Research on the β_3 -knockout mice has shown thus far that the mice have a bleeding disorder phenotypically similar to human Glanzmann thrombasthenia, resulting predominantly from the loss of the platelet integrin $\alpha_{IIb}\beta_3$ [57]. In addition it has been shown that $\beta_3^{-/-}$ mice osteoclasts are dysfunctional in bone resorption, resulting from the loss of $\alpha_v\beta_3$ [58]. It would be of interest to determine in these mice if there is a defect in TGF β activation, using appropriate models, for example the bleomycin model of pulmonary fibrosis. Alternatively, to our knowledge investigations of TGF β activity in animal models treated with $\alpha_v\beta_3$ small molecule inhibitors or blocking antibodies have not been performed.

In summary, in contrast with the literature, we have identified the RGD motif of LAP β_1 as a new ligand for $\alpha_v\beta_3$. Furthermore we have shown that the RGD motif of LAP β_3 is recognized by $\alpha_v\beta_1$, $\alpha_v\beta_3$, $\alpha_v\beta_5$ and $\alpha_v\beta_6$, whereas the non-RGD-containing LAP β_2 does not recognize any of these integrins. These interactions may be important in a number of aspects of TGF β and $\alpha_v\beta_3$ biology, particularly the many disease processes associated with both $\alpha_v\beta_3$ and TGF β up-regulation, such as cancer, rheumatoid arthritis and a variety of other diseases involving inflammation and repair.

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