

Short Communication

Altered Expression of Laminins in Crohn's Disease Small Intestinal Mucosa

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Laminins are a large family of heterotrimeric basement membrane molecules that mediate crucial cell functions such as adhesion, proliferation, migration, and differentiation. Up to now, three distinct laminins have been identified in the normal human small intestinal epithelium. Laminin-1 ($\alpha1\beta1\gamma1$) and laminin-5 ($\alpha3\beta3\gamma2$) are mainly expressed at the base of villus cells, whereas laminin-2 ($\alpha2\beta1\gamma1$) is restricted to the bottom of the crypts. The expression of these molecules has not yet been studied in Crohn's disease (CD), but it could be altered, in light of the important changes occurring in the architecture of the crypt-villus axis under the active state of the disease. To test this hypothesis, the expression of laminin $\alpha1$, $\alpha2$, and $\alpha3$ subunits was analyzed in control, inflamed, and corresponding uninflamed CD small intestinal specimens by indirect immunofluorescence and reverse transcriptase-polymerase chain reaction. Surprisingly, $\alpha1$ and $\alpha3$ remained strongly expressed by all villus cells, whereas $\alpha2$, normally expressed in the bottom of the crypts in control and uninflamed CD specimens, was lacking in inflamed CD specimens. However, this loss of $\alpha2$ expression was associated with a significant up-regulation of both $\alpha1$ and $\alpha3$ expression in the crypts of inflamed CD specimens. A significant up-regulation of the $\alpha1$ subunit was also observed in the crypts of uninflamed CD specimens. At the transcript levels, $\alpha1$ was found significantly higher in inflamed than uninflamed CD specimens. Taken together, these observations identify important alterations in laminin expression in the small intestine with CD and suggest that compositional changes in the epithelial basement membrane may

play a role in this disease. (Am J Pathol 2000, 156:45-50)

Crohn's disease is a chronic inflammatory bowel condition with a relatively high incidence among Caucasians and with a peak age of onset of between 15 and 25 years.^{1,2} Whereas the disease may affect any segment of the digestive tract, it most frequently involves the small intestine (small intestine only, 30%; ileum and cecum, 40%).^{2,3} Pathological features of Crohn's disease include transmural and granulomatous inflammation, fibrosis, alteration of the crypt-villus architecture, and epithelial cell injury.^{1,2,4}

The etiology of the disease remains incompletely understood but it is becoming more evident that interacting environmental and genetic factors are both of great importance.^{2,5} Another concept that has begun to receive considerable attention pertaining to intestinal inflammation is the involvement of essentially all cellular and acellular components of the mucosa, including immune cells and epithelial, mesenchymal, and endothelial cells, as well as the extracellular matrix (ECM), and that the dysfunction of any component of this intricate system may lead to a disruption in communication resulting in pathological inflammation.⁵

The importance of ECM molecules in the regulation of inflammation, including the recruitment and state of activation of leukocytes, has been clearly demonstrated in various systems.⁶⁻¹⁰ However, the information is still extremely limited for intestinal inflammation.¹¹ Indeed, in contrast to the normal human intestine where the expression, location, and possible function of most ECM molecules, namely laminins, type IV collagens, fibronectins, and tenascin, as well as their receptors in the integrin family, have been determined,^{12,13} almost nothing is

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known about ECM composition and function in inflammatory bowel diseases.¹⁴

As a first step in investigating this question, we have chosen to focus on laminins. Laminins represent a major component of all basement membranes both quantitatively and functionally.¹⁵ The members of this multigene family have been shown to mediate several cellular activities, namely the promotion of adhesion, migration, growth, and tissue-specific gene expression, depending on the laminin and cell type studied.^{13,15,16} In the human small intestine, three distinct laminins have been identified: laminin-1 ($\alpha1\beta1\gamma1$) and laminin-5 ($\alpha3\beta3\gamma2$), which are mainly expressed at the base of villus cells, and laminin-2 ($\alpha2\beta1\gamma1$), which is restricted to the bottom of the crypts. In this study, the expression and distribution of the three α chains specific to these laminins have been analyzed in controls as well as in inflamed and uninfamed mucosa from patients with Crohn's disease, by indirect immunofluorescence and reverse transcriptase (RT)-polymerase chain reaction (PCR). Our observations identify major alterations in the pattern of laminin expression in crypts of the small intestine with Crohn's disease and therefore support the hypothesis that ECM molecules can participate in the chronic intestinal inflammation cascade.

Materials and Methods

Tissues

Eleven specimens of distal ileum with Crohn's disease were available for the study. For each patient (five males and six females; ages 22–40), samples from both inflamed and uninfamed (resection margin) areas were obtained, and the diagnosis was confirmed by a pathologist. Three of these patients received no treatment before surgery. Six control specimens obtained from the nondiseased part (at least 10 cm distant from lesions) of resected ileum for pathologies other than Crohn's disease (bowel obstruction, primary lymphoma, or tumor) were also included in the study. All tissues were processed within 1 hour after surgery. The project was in accordance with a protocol approved by the Institutional Human Research Review Committee of the Université de Sherbrooke for the use of human material. The preparation and embedding of tissues for cryosectioning were performed as described previously.^{17,18}

Indirect Immunofluorescence

Cryosections, 2–3 μm thick, were cut on a Jung Frigocut 2800N cryostat (Leica Canada, Saint-Laurent, Canada) and fixed in methanol (10 minutes, -20°C) for the detection of $\alpha1$ (4C7¹⁹) and $\alpha2$ (5H2 and 2G9^{19,20}) or in freshly prepared 2% paraformaldehyde in phosphate-buffered saline (60 minutes, 4°C) for the detection of $\alpha3$ (BM2²¹). The secondary antibody was Cy2-conjugated goat anti-mouse IgG (Amersham-Pharmacia, Baie d'Urff, Canada). Sections were stained with Evan's blue (0.01% in phosphate-buffered saline), mounted in glycerol:phosphate-

buffered saline (9:1) containing 0.1% paraphenylene diamine, and viewed with a Reichert Polyvar 2 microscope (Leica Canada) equipped for epifluorescence. In all cases, no immunofluorescent staining was observed when primary antibodies were replaced by mouse non-immune serum.

RNA Extraction and RT-PCR

Total RNAs from the six available series of paired, inflamed and uninfamed CD specimens were prepared by Clontech's Atlas Total RNA Isolation protocol (Clontech, Palo Alto, CA) and stored in RNasecure (Ambion, Austin, TX) at -80°C . First, strand complementary DNA synthesis with Superscript II (Life Technologies, Inc., Burlington, Canada) was performed on 5 μg total RNA by using oligo(dT)^{12–18} (Amersham-Pharmacia) as primer. Conditions for amplification of laminin $\alpha1$ and $\alpha2$ chains and S14, used as endogenous control, have been described previously.^{22,23} For laminin $\alpha3$ chain, we used the sense primer 5'-GGACCTCAACGTCGGTCA-3' and the anti-sense primer 5'-CAGGGATCCTCAGTGTCTGC-3', which amplified a 209-bp amplicon at nucleotides 4814–5023 of laminin $\alpha3$.²⁴ Single-stranded complementary DNA was amplified for 30 cycles of denaturation (1 minute at 94°C) and annealing/extension (1 minute at 53°C and 1 minute at 72°C) in a thermal cycler (Perkin-Elmer DNA thermal cycler, Model 480, Foster City, CA) in the presence of 250 $\mu\text{mol/L}$ deoxyribonucleoside triphosphates and 2.5 U of *Taq* polymerase (Qiagen, Mississauga, Canada).

Results

Distribution of Laminin α Chains along the Crypt-Villus Axis

The expression of the $\alpha1$, $\alpha2$, and $\alpha3$ chains of laminin and their distribution at the epithelial basement membrane along the crypt-villus axis were determined by indirect immunofluorescence with chain-specific monoclonal antibodies. In control specimens not affected by CD, the $\alpha1$ chain was uniformly detected in the villus, but it was below the detection level in most crypts (Figure 1a) as also noted for the $\alpha3$ chain, whereas the $\alpha2$ chain was found to be confined to the lower half of the crypts as previously observed.^{18,25} In uninfamed CD specimens (resection margins), the normal histology of the ileal mucosa was relatively well preserved (Figure 1, b–e), and the distribution of the $\alpha2$ and $\alpha3$ chains was comparable to controls, $\alpha2$ being restricted to the bottom of the crypts (Figure 1d) and $\alpha3$ to the villi (Figure 1e). However, in addition to its normal distribution in the villus, the $\alpha1$ chain was consistently found also in a large proportion of the crypts, in most studied specimens (10 of 11), according to a staining pattern ranging from the lower half of the glands (Figure 1b) to the entire gland (Figure 1c). In inflamed CD specimens, the villi were short and wide, whereas the crypts were irregular and generally exhibited a larger diameter (Figure 1, f–h). The $\alpha1$ and $\alpha3$ chains

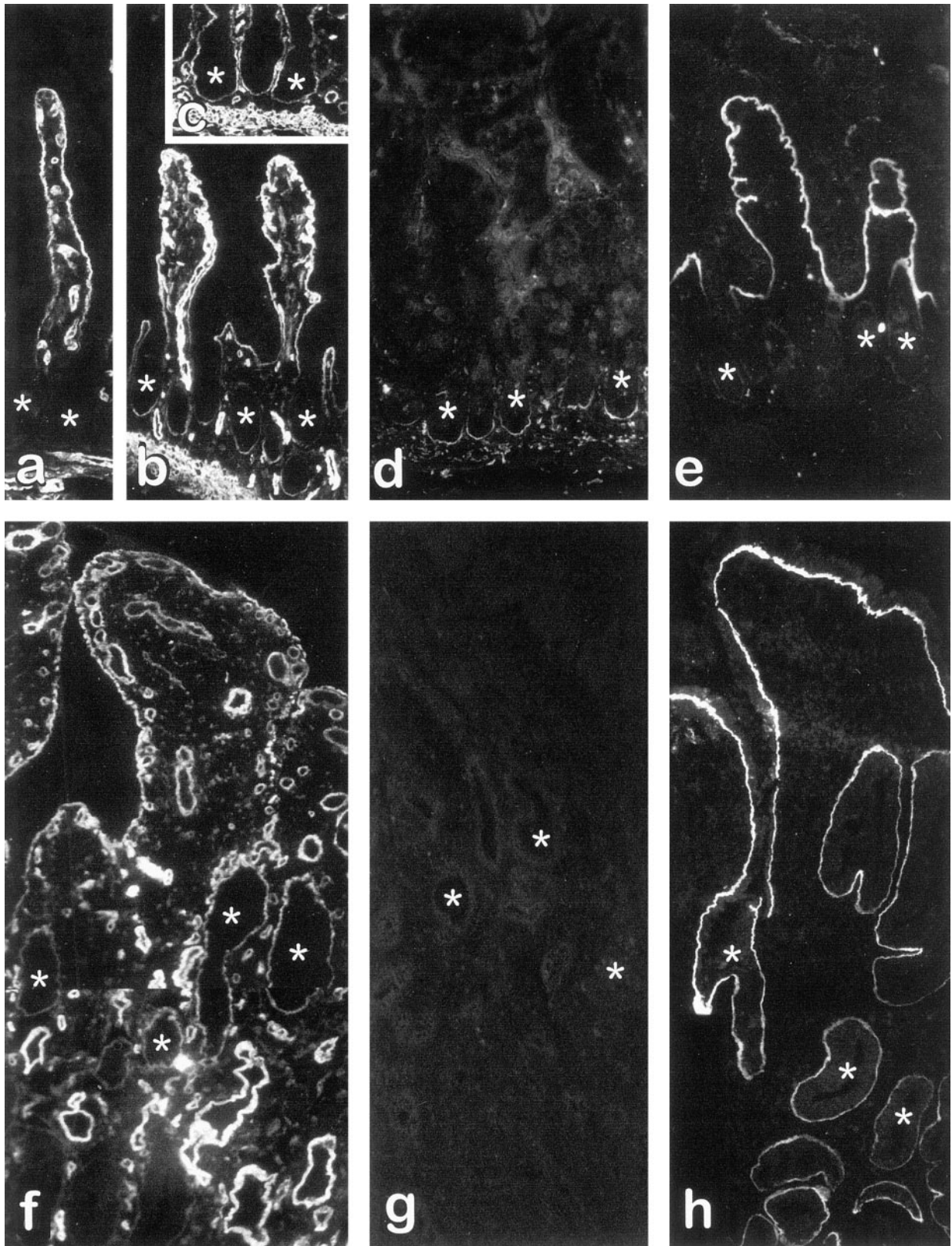


Figure 1. Expression and distribution of laminins in the small intestine from patients with CD. Indirect immunofluorescence micrographs from representative fields of cryosections from control specimens (**a**) as well as uninfamed (**b-e**) and inflamed (**f-h**) paired regions of CD specimens stained for the detection of laminin $\alpha 1$ (**a-c**, **f**), $\alpha 2$ (**d**, **g**), and $\alpha 3$ (**e**, **h**) chains. The $\alpha 1$ chain was found to be expressed in the basement membrane of villus cells under all conditions (**a-c**, **f**) as well as in crypts (**asterisks**) of uninfamed (**b**, **c**) and inflamed (**f**) CD specimens, but not in crypts of control specimens (**a**). The $\alpha 2$ chain remained normally expressed in uninfamed CD specimens, being restricted to the lower half of the crypts (**d**), whereas it was consistently undetectable in most inflamed CD specimens (**g**). The $\alpha 3$ chain was detected only in the basement membrane of the villus in uninfamed tissues (**e**), whereas it was found to stain both villus and crypt in their inflamed counterparts (**h**). Original magnification, $\times 128$. **Asterisks** denote crypts.

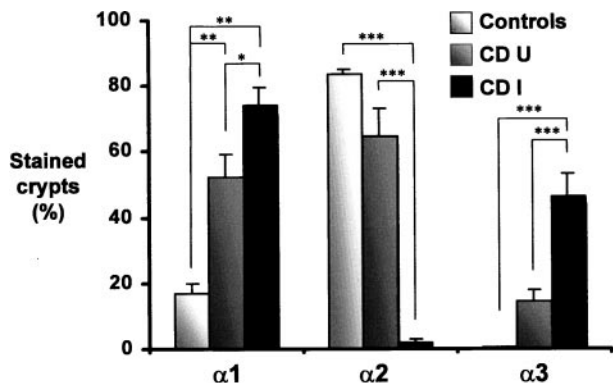


Figure 2. Expression of laminins in the crypts of control and CD small intestine. Proportions (in percentages) of crypts expressing laminin $\alpha 1, 2$, and 3 chains were determined by indirect immunofluorescent staining in control as well as in paired uninflamed (CDU) and inflamed (CDI) CD small intestinal sections by counting a minimum of 100 crypts on each section. Partially stained crypts (see Figure 1b as an example of minimally labeled crypts) were scored as positive. Six distinct specimens were used as controls, whereas 11 paired specimens were analyzed in the CD group. Asterisks over bars indicate a statistical difference: *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.005$.

were found in the basement membrane of all villi and also in a large proportion of crypts (Figure 1, f and h). However, the $\alpha 2$ chain was not detected in these specimens (Figure 1g). As summarized in Figure 2, a significant up-regulation of the $\alpha 1$ chain was observed in the crypts of both uninflamed and inflamed CD specimens, whereas alterations in the other laminin α chains were restricted to the crypts of inflamed CD specimens, the $\alpha 3$ chain being significantly increased and the $\alpha 2$ chain being mostly undetectable (Figure 2).

Expression of Laminin α Chain Transcripts in CD

Laminin α chain expression in CD was further investigated at the transcript level by RT-PCR on 6 of the 11 specimens studied by indirect immunofluorescence. As shown in Figure 3a, the three laminin α chains were found to be expressed in all CD specimens, including $\alpha 2$. Furthermore, the relative amount of $\alpha 1$ mRNA was found to be significantly higher in inflamed than uninflamed CD specimens, although the $\alpha 2$ and $\alpha 3$ transcript levels remained comparable, statistically (Figure 3b).

Discussion

These results showing alterations of laminin expression in the CD small-intestinal mucosa represent clear evidence that ECM molecules can be involved in chronic intestinal inflammation, as postulated recently.⁵ Our data pointed out three particular phenomena occurring in the inflamed and uninflamed CD small intestine, respectively. First, there is a major reorganization of the basement membrane of the crypts in inflamed intestinal segments, which consists of the disappearance of laminin-2 and its replacement by laminin-1 and laminin-5, as indicated by the distribution patterns of their respective α chain ex-

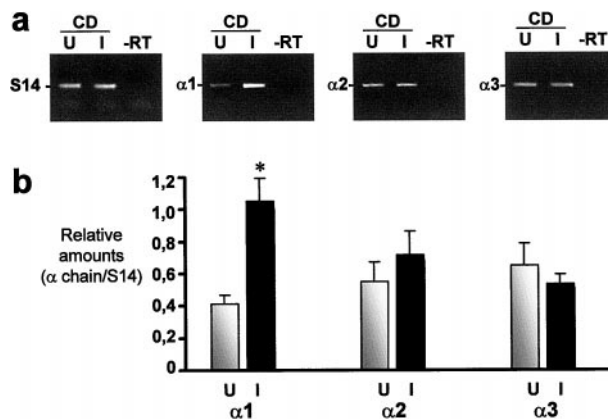


Figure 3. Expression of laminin α chain transcripts in small intestine from patients with CD. **a:** Representative RT-PCR analysis of $\alpha 1, \alpha 2$, and $\alpha 3$ chain mRNA in uninflamed (U) and corresponding inflamed (I) CD specimens. -RT, RT omitted. S14 transcript was determined to ensure complementary DNA integrity and to compare amounts of starting RNA material in the various samples. **b:** Amounts of $\alpha 1, \alpha 2$, and $\alpha 3$ transcripts in paired uninflamed (U) and inflamed (I) specimens ($n = 6$) were determined relative to S14. Statistical differences between U and I were noted for $\alpha 1$ (asterisk: $P < 0.05$).

pression. Indeed, the $\alpha 1$ and $\gamma 1$ chains, which are shared by both laminin-1 and -2, have been found constitutively expressed along the crypt-villus axis in the human small intestine.^{18,26,27} A switch from laminin $\alpha 2$ to $\alpha 1$ chain expression would thus result in the expression of laminin-1 instead of laminin-2. Alternatively, because 4C7 (which is the only antibody available for immunodetection of $\alpha 1$) may also recognize the $\alpha 5$ laminin chain,²⁸ the expression of laminin-10 ($\alpha 5\beta 1\gamma 1$) in the human small intestine remains a possibility. The expression of heterotrimeric $\alpha 3\beta 3\gamma 2$ laminin-5 in inflamed CD crypts is supported by the extracellular deposition of the $\alpha 3$ chain in the basement membrane (see Figure 1h), as well as of $\beta 3$ and $\gamma 2$ chains (data not shown). However, because the $\alpha 3$ chain can also complex with $\beta 1, \beta 2$, and $\gamma 1$, it cannot be excluded, at this time, that a proportion of $\alpha 3$ is expressed as laminin-6 ($\alpha 3\beta 1\gamma 1$) and/or laminin-7 ($\alpha 3\beta 2\gamma 1$).²⁹

The second particular feature observed in the small intestine of CD patients is the expression of the laminin $\alpha 1$ chain in the crypts of uninflamed segments, a phenomenon rarely observed in control specimens from non-CD patients.^{18,26} It is interesting that, in most of these specimens, the crypt staining for laminin-1 invariably included the bottom of the glands (see Figure 1, b and c), the site of laminin-2 expression. This situation is reminiscent of that of the fetus, in which laminin-1 and -2 are coexpressed in the developing crypts.²⁷

A third feature of potential pathological interest is the up-regulation in the expression of the $\alpha 1$ transcript in the small-intestinal mucosa of inflamed CD specimens. Increased production of ECM molecules in inflammatory bowel diseases has been previously observed for type III collagen³⁰ and tenascin,³¹ a phenomenon that seems to be related to an increased proliferation of various cellular elements of the lamina propria, such as smooth muscle cells and fibroblasts.^{4,11} The higher expression of the $\alpha 1$

chain in the inflamed CD mucosa observed herein appears consistent with these previous observations because $\alpha 1$, in contrast to $\alpha 2$ and $\alpha 3$, is also expressed by nonepithelial cellular elements of the intestinal mucosa, namely smooth muscle cells and blood vessels.^{18,25}

Taken together, these observations raise two questions. (a) What is the mechanism of alteration of laminin expression in chronic inflammation? (b) What are the functional consequences of laminin redistribution along the crypt-villus axis? For the first question, cytokines and growth factors are likely to play a central role. Indeed, as recently reviewed, alteration in the profile of numerous cytokines, as well as growth factors, is one of the hallmarks of intestinal inflammation.¹¹ As demonstrated with transforming growth factor β ,³² these molecules can affect gene expression in various cell types including epithelial cells and subepithelial myofibroblasts, both responsible for various laminin chain expression.²³ The presence of elevated levels of matrix metalloproteinases in regions of inflamed mucosa in Crohn's disease may also contribute to compositional changes in the basement membrane.³³ This latter possibility should be considered for the laminin $\alpha 2$ chain, which disappears at the protein level as shown by immunofluorescence, while its transcript remains normally expressed. Cytokines could also be responsible for the higher expression of laminin-1 in the crypts of histologically normal CD mucosa, because the production of some, such as tumor necrosis factor- α and interleukin- 1β and -6, also appears to be abnormally elevated in uninvolved CD mucosal biopsies.³⁴

To the second question pertaining to the functional consequences of alterations in laminin expression, it may be hypothesized that replacement of laminin-2 by laminin-1 and -5 in crypts of inflamed CD specimens, as well as coexpression of laminin-1 and -2 in the crypts of uninvolved CD specimens, is of functional relevance, based on the evidence that laminin-1 and -2 exhibit a distinct ability to modulate intestinal cell-specific gene expression *in vitro*.³⁵ However, laminins mediate their effects through membrane receptors, namely integrins. Many of the laminin-binding integrins have been found to be expressed in the normal small intestine, including $\alpha 2\beta 1$, $\alpha 3\beta 1$, $\alpha 7\beta 1$, and $\alpha 6\beta 4$,^{17,22,36} which can bind differentially to distinct laminins and mediate specific cell functions.^{13,16} Further work will thus be required to better delineate, at the cellular level, the functional implications of laminin expression alteration.

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