Dynamics of the $\alpha 6\beta 4$ Integrin in Keratinocytes ∇

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> The integrin $\alpha 6\beta 4$ has been implicated in two apparently contrasting processes, i.e., the formation of stable adhesions, and cell migration and invasion. To study the dynamic properties of $\alpha 6\beta 4$ in live cells two different β 4-chimeras were stably expressed in β 4-deficient PA-JEB keratinocytes. One chimera consisted of full-length β 4 fused to EGFP at its carboxy terminus (β 4-EGFP). In a second chimera the extracellular part of $\beta 4$ was replaced by EGFP (EGFP- $\beta 4$), thereby rendering it incapable of associating with $\alpha 6$ and thus of binding to laminin-5. Both chimeras induce the formation of hemidesmosome-like structures, which contain plectin and often also BP180 and BP230. During cell migration and division, the β 4-EGFP and EGFP- β 4 hemidesmosomes disappear, and a proportion of the β 4-EGFP, but not of the EGFP- β 4 molecules, become part of retraction fibers, which are occasionally ripped from the cell membrane, thereby leaving "footprints" of the migrating cell. PA-JEB cells expressing β 4-EGFP migrate considerably more slowly than those that express EGFP- β 4. Studies with a β 4-EGFP mutant that is unable to interact with plectin and thus with the cytoskeleton (β4^{R1281W}-EGFP) suggest that the stabilization of the interaction between $\alpha 6\beta 4$ and LN-5, rather than the increased adhesion to LN-5, is responsible for the inhibition of migration. Consistent with this, photobleaching and recovery experiments revealed that the interaction of β 4 with plectin renders the bond between α 6 β 4 and laminin-5 more stable, i.e., β 4-EGFP is less dynamic than β 4^{R1281W}-EGFP. On the other hand, when α 6 β 4 is bound to laminin-5, the binding dynamics of β 4 to plectin are increased, i.e., β 4-EGFP is more dynamic than EGFP- β 4. We suggest that the stability of the interaction between $\alpha \delta \beta$ 4 and laminin-5 is influenced by the clustering of $\alpha 6\beta 4$ through the deposition of laminin-5 underneath the cells. This clustering ultimately determines whether $\alpha 6\beta 4$ will inhibit cell migration or not.

INTRODUCTION

Keratinocytes adhere to the basement membrane by hemidesmosomes that serve as anchoring sites for the intermediate filament system and play a critical role in stabilizing the association of the dermis with the epidermis. The transmembrane components of hemidesmosomes comprise the laminin-5 (LN-5) binding integrin $\alpha 6\beta 4$ and the bullous pemphigoid antigen (BP)180. These proteins are connected via the hemidesmosomal proteins plectin and BP230 to the keratin intermediate filament system (reviewed by Jones *et al.*, 1998; Borradori and Sonnenberg, 1999).

Based on their structural constituents, two subtypes of hemidesmosomes are distinguished. Type I hemides-

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* Corresponding author. E-mail address: a.sonnenberg@nki.nl. Abbreviations used: LN-5, laminin-5; PA-JEB, pyloric atresia associated with junctional epidermolysis bullosa; EGFP, enhanced green fluorescent protein; BP, bullous pemhigoid antigen; IL2R, interleukin-2 receptor; FRAP, fluorescence recovery after photobleaching mosomes contain $\alpha 6\beta 4$, plectin, BP180, and BP230 (Green and Jones, 1996), whereas type II hemidesmosomes contain only $\alpha 6\beta 4$ and plectin (Uematsu *et al.*, 1994). Recently, the tetraspanin CD151 was identified as another component of both type I and II hemidesmosomes (Sterk *et al.*, 2000). Type I or classical hemidesmosomes are present in basal keratinocytes of squamous and complex epithelia (Nievers *et al.*, 1999). Type II hemidesmosomes are found in intestinal epithelial cells and some other cultured epithelial cell types (Uematsu *et al.*, 1994; Orian-Rousseau *et al.*, 1996; Fontao *et al.*, 1997). The association of type II hemidesmosomes with intermediate filaments is less robust than that of type I hemidesmosomes, which may imply a more dynamic regulation of their assembly.

The stability of type I hemidesmosomes is illustrated by their continued presence during mitosis (Riddelle *et al.*, 1992; Baker and Garrod, 1993), thereby ensuring that cells with a strong proliferative potential remain present in the basal compartment of the skin. However, during wound healing, hemidesmosomes are disassembled to allow keratinocytes to migrate on a newly deposited LN-5 matrix (Martin, 1997; Decline and Rousselle, 2001). Several growth factors have been implicated in the regulation of the disassembly of hemidesmosomes, including the epidermal and hepatocyte growth factors (Mainiero et al., 1996; Trusolino et al., 2001). As a result of interaction of these growth factors with their cognate receptors, the β 4 subunit is tyrosine phosphorylated and recruits the signaling adaptor protein Shc (Mainiero et al., 1996; Mariotti et al., 2001; Trusolino et al., 2001). Conceivably, the phosphorylation of β 4 on tyrosine residues may prevent its incorporation into hemidesmosomes. Studies by Rabinovitz et al. (1999), however, have revealed that EGF receptor-mediated disruption of hemidesmosomes depends on the ability of this receptor to activate protein kinase C and may involve the direct phosphorylation of the β 4 cytoplasmic domain on serine residues. In addition, there is evidence suggesting that $\alpha 6\beta 4$ activates phosphoinositide 3-OH (PI-3) kinase (Shaw et al., 1997; Shaw, 2001) and interacts with the actin cytoskeleton in filopodia and lamellipodia (Rabinovitz *et al.*, 1997, 1999). This ability of α 6 β 4 to activate PI-3-kinase signaling has been connected to the promotion by this integrin of the migration and invasion of carcinoma cells (Rabinovitz and Mercurio, 1997; Shaw et al., 1997; Gambaletta et al., 2000; Hintermann et al., 2001). Activation of PI-3 kinase by $\alpha 6\beta 4$ may also contribute to adhesion and spreading of keratinocytes via binding of $\alpha 3\beta 1$ to LN-5 (Nguyen *et al.*, 2000). Finally, it has been shown that proteolytic processing of the LN-5- α 3 and - γ 2 chains may determine whether this matrix protein supports stable adhesion or instead migration (Giannelli et al., 1997; Goldfinger et al., 1998). Taken together these data show that cell migration is regulated by both extrinsic (proteases) and intrinsic (signaling molecules) factors.

New insights into the dynamic properties of focal contacts have been provided by studying green fluorescent (GFP)tagged integrin α or β subunits in live cells (Smilenov *et al.*, 1999; Ballestrem et al., 2001; Laukaitis et al., 2001). To gain more information about the dynamics of hemidesmosomes, we expressed two different enhanced GFP (EGFP)-tagged β 4 chimeras in B4-deficient PA-JEB keratinocytes (Schaapveld et al., 1998). In one, EGFP was fused to the carboxy-terminal end of the β 4 cytoplasmic domain (β 4-EGFP), and in the other the extracellular domain of β 4 was replaced by EGFP (EGFP- β 4), which made this chimera incapable of associating with the α 6 subunit and thus of binding LN-5. However, independently of binding to ligand, it can still induce hemidesmosome formation through its interaction with the hemidesmosomal component plectin (Nievers et al., 1998, 2000).

We studied the formation of hemidesmosomes by these two chimeras by using time-lapse video microscopy. Our data indicate that hemidesmosomes containing either β 4-EGFP or EGFP- β 4 are both dynamic structures that are assembled and redistributed during cell migration and division. Furthermore, we show that as a result of binding of $\alpha 6\beta 4$ to plectin, the binding of $\alpha 6\beta 4$ LN-5 is stabilized, which inhibits cell migration and reduces the dynamics of $\alpha 6\beta 4$. We thus provide novel insights into the role of the $\alpha 6\beta 4$ integrin in migration.

MATERIALS AND METHODS

Cell Lines

Immortalized β 4-deficient keratinocytes have been derived from a patient with pyloric atresia associated with junctional epidermolysis

bullosa (PA-JEB; Niessen *et al.*, 1996; Schaapveld *et al.*, 1998). Fulllength β4 cDNA and a cDNA encoding a chimeric protein consisting of the extracellular and transmembrane domains of the interleukin 2 receptor (IL2R) fused to the cytoplasmic domain of β4 were stably expressed in PA-JEB cells by retroviral infection to generate PA-JEB/β4 (Sterk *et al.* 2001) and PA-JEB/IL2R-β4 (Nievers *et al.*, 1998), respectively. PA-JEB/β4-EGFP, PA-JEB/EGFP-β4, and PA-JEB/β4^{R1281W}-EGFP cells were generated as described below. All cells were maintained in keratinocyte serum-free medium (SFK; Life Technologies-BRL; Rockville, MD) supplemented with 50 µg/ml bovine pituitary extract, 5 ng/ml epidermal growth factor, 100 U/ml penicillin, and 100 U/ml streptomycin. To stimulate hemidesmosome formation, cells were grown for 24 h in calciumrich medium, consisting of HAM-F12 Nutrient Mixture (Life Technologies-BRL) and DMEM (Life Technologies-BRL) in a ratio of 1:3.

Plasmid Constructs and Generation of PA-JEB/β4-EGFP and PA-JEB/EGFP-β4 Keratinocytes

The chimeric β4-EGFP and EGFP-β4 constructs were produced in pcDNA3. The β4-EGFP chimera consisted of the full-length human β 4A sequence, a sequence encoding six glycine residues followed by EGFP (Invitrogen, Carlsbad, CA). The EGFP-β4 chimera consisted of the amino acid signal sequence of the β 4 subunit, EGFP, and transmembrane and cytoplasmic domains of human β 4A integrin. Both constructs were generated by overlap PCR using the following sequences: β4 c-terminus/6*glycine, 5'-catgccgccgccgccgccgccgctttggaagaactgttggtc-3'; 6*glycine-EGFP, 5'-ggcggcggcggcggcggcggcatggtgagcaagggcgaggag-3'; β4 signal sequence-EGFP, 5'-ctcctcgcccttgctcaccatgcggtttgccaaggtcccaga-3'; EGFP-β4 transmembrane, catgccatggtcttgtacagctcgtccatgcc-3'. The two PCR products were digested with EcoRI and cloned into the corresponding site of the pcDNA3 vector. The $\beta 4^{R1281W}$ -EGFP construct was generated by replacing a 5-kb EcoRI/EcoRV fragment from β4-EGFP cDNA with a corresponding fragment containing a CGG to TGG mutation in codon 1281 of β 4 (Geerts *et al.*, 1999). Subsequently, the EGFP- β 4, β 4-EGFP, and β 4^{R1281W}-EGFP cDNAs were released from pcDNA3 by digestion with EcoRI, and the resulting fragments were ligated into the corresponding site of the retroviral LZRS-IRES-zeo expression vector, a modified LZRS retroviral vector conferring resistance to zeocin (Kinsella and Nolan, 1996; van Leeuwen et al., 1997) to result in the LZRS-EGFP-β4-IRES-zeo, LZRS-β4-EGFP-IRES-zeo, and LZRS-B4R1281W-EGFP-IRES-zeo constructs, respectively. Correctness of the DNA constructs was verified by sequence analysis The retroviral constructs were transfected into the Phoenix packaging cell line (Kinsella and Nolan, 1996) by the calcium phosphate precipitation procedure, and after 2 days supernatants containing recombinant viruses were collected. Transduction of the recombinant viruses in PA-JEB cells was performed for 10 h at 37°C. PA-JEB cells expressing β 4-EGFP, β 4^{R1281W}-EGFP, or EGFP- β 4 were isolated by FACS, expanded, and analyzed.

Antibodies

The rat mAb GoH3 is a blocking antibody directed against the extracellular part of the integrin α 6 subunit (Sonnenberg *et al.*, 1987). The mouse mAb K20, anti- β 1, was purchased from Biomeda (Foster City, CA). The mouse mAb 450–11A, directed against the cytoplasmic domain of β 4 was from PharMingen International (San Diego, CA). The rabbit polyclonal antibody against the LN-5 α 3 chain was a kind gift of Dr. R. Timpl (Max Planck Institut für Biochemie, Martinsried, Germany). The mouse mAbs 121, anti-plectin/HD1 (Hieda *et al.*, 1992), and 233, anti-BP180 (Nishizawa *et al.*, 1993), were generously provided by Dr. K. Owaribe (Nagoya University, Nagoya, Japan). The human mAbs 5E and 10D against BP230 (Hashimoto *et al.*, 1993) were a kind gift of Dr. T. Hashimoto (Kurume University, Kurume, Fukuoka, Japan). The mouse mAb TB30, against the extracellular domain of the interleukin-2-receptor (IL2R), was purchased from the Central Laboratory of the Red Cross

Blood Transfusion Service (Amsterdam, The Netherlands). Mouse mAb 3C12, anti-ezrin, was from Lab Vision Corp. (Fremont, CA), mouse mAb RV202, anti-vimentin, was kindly provided by Dr. F. Ramaekers (University of Limburg, Maastricht, The Netherlands), mouse mAbs against α and β tubulins were from Sigma Chemical (St. Louis, MO; clones B-5-1-2 and 2-28-33), mouse mAb KL-1, anti-keratin, was from Immunotech (Marseille, France), rabbit antiserum against human LN-5 (Marinkovich *et al.*, 1992) was a kind gift of Dr. R. Burgeson (Cutanous Biology Research Center, Charlestown, MA). The mouse mAb P48, also known as 11B1.G4, was clustered as CD151 in the VI International Leukocyte Typing Workshop (Ashman *et al.*, 1997). The mouse B34 mAb directed against GFP was purchased from BabCO (Richmond, CA).

The sheep anti-mouse horseradish peroxidase–coupled antibodies were purchased from Amersham Corp. (Arlington Heights, IL), Texas Red–conjugated goat anti-mouse rat or rabbit antibodies were obtained from Molecular Probes (Eugene, OR), Texas Red– conjugated donkey anti-human antibodies and Cy-5–conjugated goat anti-rabbit antibodies and were from Jackson ImmunoResearch Laboratories (West Grove, PA).

Time-lapse Observations and Fluorescence Recovery after Photobleaching Experiments

Time-lapse observations were made in a tissue culture device at 37°C and viewed under a Leica TCS-NT confocal microscope (Deerfield, IL) equipped with argon/krypton laser. The krypton/argon laser was used to excite the EGFP-tagged proteins at 488 nm, and emissions above 515 nm were collected. Images of β 4-EGFP and EGFP- β 4 were collected every 2–15 min for periods up to 4 h. Phase-contrast images of cells were taken during time-lapse observations to obtain the corresponding cell shape image.

Fluorescence recovery after photobleaching (FRAP) experiments were performed by selecting a region of β 4-EGFP or EGFP- β 4 hemidesmosomes located at the cell periphery, and oval-shaped regions were bleached using the krypton/argon laser for 1 s at 100% power, resulting in a bleached spot of 1 μ m diameter. Images were collected after bleaching every 15 s for 10 min. The fluorescence intensity in the bleached region of the β 4-EGFP or EGFP- β 4 hemidesmosome during 10 min of recovery was normalized to the fluorescence intensity measured in a nonbleached region. This procedure allowed us to account for the decreased fluorescence due to overall bleaching of the entire field as a result of image collection. Phase-contrast images of cells were taken during FRAP analysis to ensure that there was no significant change in cell shape and position during periods of observation. Imaging from live cells on our confocal system prohibits the collection of large numbers of images, so that reliable fitting of more than one component is not possible. In the inhibitor studies, antibodies (GoH3) were added at a concentration of 25 μ g/ml 24 h before FRAP analysis.

Preparation of Laminin-5 Matrices

PA-JEB/ β 4-EGFP and PA-JEB/EGFP- β 4 keratinocytes were grown to confluency in six-well tissue culture plates, washed three times with PBS, and incubated overnight at 4°C in PBS containing 20 mM EDTA and a cocktail of protease inhibitors (Sigma). After incubation the cells were removed by forceful pipetting, and the remaining matrices were dissolved in SDS sample buffer. For Western analysis a fraction (1/4) of the matrices in the well was loaded.

Immunofluorescence Microscopy

PA-JEB/ β 4, PA-JEB/ β 4-EGFP, PA-JEB/EGFP- β 4, and PA-JEB/ IL2R- β 4 keratinocytes grown on glass coverslips were washed and fixed with 1% (wt/vol) formaldehyde for 10 min. Fixed cells were washed twice with PBS and permeabilized in 0.5% (vol/vol) Triton X-100 in PBS for 5 min. Cells were rinsed with PBS and incubated in 2% (wt/vol) BSA in PBS for 1 h, followed by incubation with the primary antibody for 1 h. After washing twice with PBS, Texas Redand Cy-5–conjugated secondary antibodies directed against mouse and rabbit immunoglobulins, respectively, were applied for another 1 h. Actin was labeled with phalloidin Alexa 568 at a 1:40 dilution. After washing twice with PBS, coverslips were mounted on microscope slides with Mowiol (Longin *et al.*, 1993) and viewed under a Leica TCS-NT confocal laser-scanning microscope. All steps were performed at room temperature.

Western Blotting Analysis

PA-JEB/β4, PA-JEB/EGFP-β4, and PA-JEB/β4-EGFP keratinocytes were lysed in 1% (wt/vol) SDS, 10 mM Tris-HCl (pH 7.4), containing the proteinase inhibitors phenylmethylsulfonyl fluoride (1 mM), soybean trypsin inhibitor (10 μ g/ml), and leupeptin (10 μ g/ml). The protein extracts of 2 \times 10⁵ cells were loaded per lane on 10% polyacrylamide gels, separated by electrophoresis, and transferred to Immobilon-PVDF membranes (Millipore, Bedford, MA). The membranes were stained with Coomassie blue to indicate the markers, destained (45% methanol, 5% acetic acid in demineralized water), and blocked by incubation in 5% nonfat dry milk in TBSTbuffer (10 mM Tris-HCl, pH 7.5, 150 mM NaCl, 0.3% Tween-20) for 30 min at room temperature. Then, the membranes were incubated with the mouse primary antibodies B34 (anti-GFP) or 450-11A (anti- β 4), diluted 1:500 in 0.5% dry milk in TBST, for 1 h at room temperature. After washing three times with TBST, the membranes were incubated with secondary sheep anti-mouse Ig-coupled horseradish peroxidase (1:5000 dilution) for an additional hour at room temperature. Immunoreactive bands were visualized using enhanced chemiluminescence, according to the manufacturer's instructions (Amersham Corp.).

Measurement of Cell Motility

PA-JEB/EGFP- β 4, PA-JEB/ β 4-EGFP, and PA-JEB/ β 4^{R1281W}-EGF keratinocytes were grown to confluence on glass coverslips in keratinocyte-SFM. To assess the relative contribution of cell migration in the absence of proliferation, cells were treated with 10 μ g/ml mitomycin C (Sigma Chemical) 2 h before wounding. A cell-free area was introduced by scraping the monolayer with a yellow pipette tip, followed by three washes with PBS to remove cell debris. Scratched areas were photographed at ×200 magnification. Cells were subsequently incubated at 37°C for 48 h in keratinocyte-SFM, and the wounded monolayers were photographed again. In the inhibitor studies, antibodies (GoH3) were added at a concentration of 25 μ g/ml, 1 h before wounding.

RESULTS

β4-EGFP and EGFP-β4 Induce the Assembly of Hemidesmosomes

In previous studies using keratinocytes transfected with an IL2R- β 4 chimera, it was shown that $\alpha 6\beta$ 4 can induce the formation of hemidesmosomes without binding to its ligand (Nievers *et al.*, 1998, 2000). Under these circumstances hemidesmosome formation is driven by the cytoplasmic domain of the β 4 subunit and is dependent on its association with plectin. To study ligand-dependent and -independent hemidesmosome formation in live cells, two different β 4-EGFP chimeras were constructed. The β 4-EGFP chimera consisted of EGFP fused to the carboxy-terminal end of the β 4 cytoplasmic domain, whereas in the EGFP- β 4 chimera the extracellular domain of β 4 was replaced by EGFP (Figure 1A). The chimeras were introduced by retroviral transduction into β 4-deficient PA-JEB keratinocytes to create stable cell lines. Both EGFP chimeras were expressed, and their

molecular masses were as expected (Figure 1B). Introduction of the EGFP chimeras did not affect the expression of the other members of the integrin family at the cell surface, because FACS analysis demonstrated that the levels of $\alpha 2$, $\alpha 3$, $\alpha 5$, and $\beta 1$ subunits were unaffected (unpublished data).

Fluorescence microscopy of PA-JEB cells expressing the β 4-EGFP or EGFP- β 4 chimera showed that both chimeras are able to induce the formation of hemidesmosome-like structures, the distribution pattern of which resembles that produced by the expression of β 4 or IL2R- β 4 in PA-JEB cells, respectively (Figure 2A). Furthermore, the hemidesmosome-like structures contain plectin, BP180, and BP230. β 4-EGFP was always colocalized with patches of LN-5 deposited underneath the cells, whereas sometimes EGFP- β 4 was not colocalized with LN-5 (Figure 2B). These results support the assumption that β 4-EGFP when associated with α 6 interacts with LN-5, whereas the EGFP- β 4 chimera does not.

Analysis of the matrices deposited by β 4-EGFP– and EGFP- β 4–expressing cells by immunoblotting, using polyclonal anti-LN-5 antibodies, confirmed the presence of LN-5 in these matrices and furthermore showed that the α 3 chain of a proportion of laminin-5 had been proteolytically processed. The ratio of unprocessed (190 kDa) and processed α 3 (165 kDa) chain was slightly different between the two cell lines; more α 3 chain being processed in PA-JEB/EGFP- β 4 than in PA-JEB/ β 4-EGFP cells. There was no evidence for extracellular processing of the β 3 chain. Furthermore, the total content of laminin-5 in the matrices deposited by PA-JEB/EGFP- β 4 and PA-JEB/ β 4-EGFP cells was comparable.

In summary, both EGFP chimeras are able to induce the assembly of hemidesmosomes, but although this assembly by EGFP- β 4 is entirely driven from within the cell (Nievers *et al.*, 2000), that by β 4-EGFP is also induced by an interaction of its extracellular domain with LN-5.

Dynamics of Hemidesmosome Formation

Time-lapse videomicroscopy was used to study the distribution of hemidesmosomes during the random movement of keratinocytes. When cultured in high Ca²⁺ medium, PA-JEB/ β 4-EGFP and PA-JEB/EGFP- β 4 keratinocytes hardly migrate but move primarily in situ by continuously extending and retracting their membrane. Changes in the distribution pattern of hemidesmosomes are readily detected in these cells over a period of 3 h (Figure 3A). Existing hemidesmosomes in the central region of the cell disappear, and new hemidesmosomes are formed at the cell margins. Because of the extensions of the membrane of the cell in various directions, the distribution pattern of hemidesmosomes often has the appearance of a cauliflower. Cell retraction is accompanied by the formation of retraction fibers in which β 4-EGFP, but not EGFP- β 4, is present. The β 4-EGFPpositive retraction fibers originate from hemidesmosomes, probably because β 4-EGFP when associated with α 6 cannot be released from its ligand without effort. In migrating cells, retraction fibers are formed at the rear of the cell and occasionally are left behind as "footprints" (Figure 3B).

Time-lapse videomicroscopy also revealed that as the cell further retracts its membrane the β 4-EGFP retraction fibers become thinner and longer, and new ones emerge (Figure 3C, A–F). Fusion of retraction fibers is also observed (Figure 3C, arrow, panels E and F). Eventually, the fibers become smaller

and are pulled back inside the cell. In panels F–J, it can be seen that the empty space left behind by the migrating cell is quickly occupied by another cell that extends its membrane from the leading edge. No β 4-EGFP can be detected in these membrane extensions, which suggests that α 6 β 4 is not an essential component of newly formed filopodia or lamellipodia. Together these results show that both β 4-EGFP and EGFP- β 4 hemidesmosomes are assembled and redistributed in a short period of time. Only β 4 chimeras (β 4-EGFP) that interact with LN-5 are retained in retraction fibers.

Retraction Fibers Mediate the Final Bond of PA-JEB/β4-EGFP Cells with LN-5 during Migration and Mitosis

Hemidesmosomes do not disassemble when keratinocytes divide (Baker *et al.*, 1993). Nevertheless, during the mitotic process, cells have to undergo cellular rounding, which is accompanied by a complete reorganization of the cytoskeleton. To investigate whether the localization of hemidesmosomes changes during mitosis, we recorded the fluorescence of β 4-EGFP– and EGFP- β 4–containing hemidesmosomes during spontaneous cell divisions. Figure 4A shows a series of images of a dividing PA-JEB/ β 4-EGFP cell. The images were taken at 15-min intervals over a period of 90 min and after completion of cell division (+8 h). When mitosis begins, the cell starts to round up and to detach from the matrix. The hemidesmosomes located at the cell borders are converted into retraction fibers (Figure 4A, 30 min). When



Figure 1. (A) Diagram of β 4-EGFP and EGFP- β 4 constructs. In β 4-EGFP the complete β 4 subunit was fused to EGFP and in EGFP- β 4, the extracellular domain of the β 4 subunit was replaced by EGFP. (B) Western blot analysis of β 4-EGFP and EGFP- β 4 protein levels in PA-JEB/β4-EGFP and PA-JEB/EGFP-β4 keratinocytes. Whole cell lysates were resolved by SDS-PAGE and subjected to immunoblotting with monoclonal antibodies against β 4 (450-11A) and GFP (B34). β 4-EGFP and EGFP- β 4 are present as a single band of the expected molecular mass. (C). Western blot analysis of laminin-5 in matrices produced and deposited by PA-JEB/β4-EGFP and PA-JEB/EGFP-B4 keratinocytes. Matrices were solubilized, separated by SDS-PAGE under reducing conditions, and processed for Western blot analysis using anti-laminin-5 antiserum. The four major immunoreactive proteins of 190, 165, 155, and 145 kDa correspond to the unprocessed and processed forms of α 3, and to the γ 2 and β 3 chains of laminin-5, respectively.

Δ PA-JEB/β4 and PA-JEB/β4-EGFP cells



PA-JEB/IL2R-B4 and PA-JEB/EGFP-B4 cells



Β PA-JEB/β4-EGFP cells



the rounding of the cell is completed, it is still attached to the LN-5 matrix via a network of *β*4-EGFP positive retraction fibers (45 min). These fibers keep the cell at its position during the mitotic process (60-75 min) and subsequently will facilitate the spreading of the daughter cells (Cramer and Mitchison, 1993; Mitchison and Cramer, 1996). After completion of cell division, the daughter cells spread on the

components

LN-5 matrix that was already present beneath the dividing cell, and new hemidesmosomes are formed at exactly the same position at which the β 4-EGFP retraction fibers arose from the matrix (Figure 4A, 8 h). Thus, as in migrating cells, in mitotic cells the β 4-EGFP containing hemidesmosomes are converted into retraction fibers. The reverse reaction, the conversion of retraction fibers into *β*4-EGFP containing

Figure 2. (A) Similar localization of β 4-EGFP and β 4, and of EGFP- β 4 and IL2R- β 4. Top: PA- $JEB/\beta4$ and PA-JEB/ $\beta4$ -EGFP keratinocytes in a mixed culture were fixed and incubated with antibodies against the cytoplasmic domain of $\beta 4$ (450–11A), followed by Texas red-conjugated goat antimouse IgG. Both β 4 and β 4-EGFP

hemidesmosomes occur in a cauliflower-like staining pattern, and there is no difference between $\beta 4$

and β 4-EGFP hemidesmosomes. Bottom: PA-JEB/IL2R-B4 and PA-JEB/EGFP-β4 keratinocytes in a mixed culture were fixed, and the IL2R-β4 construct was labeled with antibodies directed against the IL2R part of the chimera (in the IL2R- β 4 chimera, the extracellular part of β 4, is replaced by the extracellular part of the IL2 receptor), followed by Texas red-conjugated secondary antibodies. β 4-EGFP

and IL2R- β 4 that do not interact

with LN-5 are mainly localized at the periphery of cells and the hemidesmosomes formed by them

appeared to be more clustered than those formed by β 4-EGFP or β 4. No differences can be observed between EGFP-B4 and IL2R-B4 hemidesmosomes. Composite images were generated by superimposition of the green and red signals. Areas of overlap appear yellow in the images. (B) Colocalization of *β*4-EGFP and EGFP-*β*4 with the hemidesmosomal components BP180, BP230, plectin, and LN-5. PA-JEB keratinocytes ex-

pressing β 4-EGFP (left panel) or EGFP- β 4 (right panel) were fixed and immunostained for BP180, BP230, plectin, or LN-5. Composite images were generated by superimposition of the green and red signals. Areas of overlap appear yellow in the images. Both chimeras are colocalized with the

BP180, BP230, plectin, and LN-5. Only β 4-EGFP is colocalized with

hemidesmosomal

LN-5. Bar, 10 µm.

A PA-JEB/β4-EGFP



PA-JEB/EGFP-β4

В



 B4-EGFP
 LN-5



Figure 3. Dynamics of hemidesmosome formation. (A) Selected fluorescence micrographs taken from a time-lapse recording at time points 0, 90, and 195 min, depicting the change in distribution of 64-EGFP and EGFP- β 4 hemidesmosomes in PA-JEB/ β 4-EGFP and PA-JEB/ EGFP- β 4 cells, respectively. The left and two middle panels show an overlay of transmission optics and fluorescence. To improve the contrast of the composite, the green fluorescence of the EGFP chimeras is shown in blue. Right panel: a composite of the fluorescence images at time points 0 (in green) and 195 min (pseudocolored in red), with areas of overlap represented in yellow. Note that after 195 min, new β 4-EGFP hemidesmosomes (top panel, red signal) have assembled at the cell borders, whereas preexisting hemidesmosomes (top panel, green signal) in the central part of the cell have disappeared. Bar, 15 μ m. (B) "Footprints" of PA-JEB/ β 4-EGFP keratinocytes. PA-JEB/ 84-EGFP keratinocytes were fixed and immunostained for LN-5 (red). Composite images were generated by superimposition of the green and red signals. Areas of overlap appear yellow in the images. Migrating β 4-EGFP keratinocytes leave footprints containing ripped off β4-EGFP membrane fragments still attached to the underlying matrix. Bar, 10 µm. (C) Time-lapse images of PA-JEB/ β 4-EGFP demonstrating that hemidesmosomes convert into retraction fibers during cell movement. Left panels: fluorescent images; right panels: the corresponding transmission optics. Note that when the cell body moves forward, hemidesmosomes are gradually converted into retraction fibers (A-F). Some fusion of retraction fibers occurs and eventually the retraction fibers become smaller and are pulled back into the cell (arrows, D-G). The arrow in panel A points at the appearance of retraction fibers. The arrow in panel J points to membrane extensions of the neighboring cell. Note that the disassembly of hemidesmosomes and their conversion into retraction fibers lead to withdrawing of the cell in the left corner, thereby allowing the cell below it to migrate into the cleared space (F-J).

PA-JEB/β4-EGFP cells

Α

В



PA-JEB/EGFP-β4 cells



Figure 4. Hemidesmosomes of β 4-EGFP and EGFP- β 4 keratinocytes disappear during mitosis. Time-lapse images of dividing PA-JEB/ β 4-EGFP (A) and PA-JEB/EGFP- β 4 (B) cells. Shown are fluorescent images with the corresponding transmission optics in the right corner of every panel. Note that β 4-EGFP hemidesmosomes convert into retraction fibers, whereas EGFP- β 4 hemidesmosomes do not but are reduced to small structures that no longer appear as hemidesmosomes. Retraction fibers not only form at the rear end during cell contraction, but also promote respreading of cells after mitosis, consequently the hemidesmosomal pattern before and after mitosis is similar.

hemidesmosomes, also occurs and is most frequently seen after completion of cell division, when the rounded cells begin to spread.

 $EGFP-\beta \overline{4}$ hemides mosomes do not convert into retraction fibers but are redistributed during mitosis (Figure 4B). Their number at the sites where the cell is attached to the matrix decreases during the process of mitosis. Finally, when the cell is rounded, a spot-like staining pattern represents the only structure that remains of the EGFP-B4 hemidesmosomes. After cell division, the EGFP-B4 hemidesmosomes reappear in the daughter cells that start to spread. The hemidesmosomal pattern that is finally formed after spreading of the daughter cells is the same as before the onset of mitosis (Figure 4B, 8 h). The transmission images further demonstrate that spreading of the PA-JEB/EGFP-B4 cells is associated with extension of the membranes and the formation of bleb-like structures (Figure 4B, 225-240 min). Together, these data demonstrate that during mitosis \u03b84-EGFP hemidesmosomes are converted into retraction fibers, whereas those containing EGFP-B4 are reduced to small structures that no longer resemble hemidesmosomes. After cell division is completed hemidesmosomes are reassembled.

β4-EGFP Is Not Associated with the Intermediate Filament System in Retraction Fibers

Because retraction fibers that contain β 4-EGFP appeared to originate from hemidesmosomes, we investigated which other components of hemidesmosomes are present in them.

No BP180, BP230, or plectin could be detected in β 4-EGFPpositive retraction fibers. Actually plectin, that mediates the linkage of β 4 to the intermediate filament system, is already dissociated from $\alpha 6\beta$ 4 before the retraction fibers become visible (Figure 5A). This suggests that the integrin must be detached from the intermediate filament system before retraction fibers can be formed. Consistent with the fact that plectin is not present in retraction fibers, we could not detect the filament proteins, keratin or vimentin, or tubulin in these fibers either (Figure 5, B, C and D). On the other hand, the tetraspanin CD151 is colocalized with β 4-EGFP along the retraction fibers and thus is the only other component that is present together with $\alpha 6\beta$ 4 in both hemidesmosomes and retraction fibers (Figure 5E).

The β 1 integrins are not colocalized with β 4 in hemidesmosomes but are present in the focal contacts surrounding them (Schaapveld *et al.*, 1998). Because the retraction fibers of migrating fibroblasts are composed of actin and β 1 (Chen, 1981), we investigated whether these components are also present in the retraction fibers of PA-JEB/ β 4-EGFP cells. As shown in Figure 5, β 1 is detected in retraction fibers, but it is not colocalized with β 4-EGFP (Figure 5F). As expected, Factin is also present in retraction fibers, and it was found to be more prominently present at their base (Figure 5G). No filamin, spectrin, talin, vinculin, or zyxin could be demonstrated in retraction fibers, which would imply that the β 1 integrins are not connected to the actin cytoskeleton. How-



Figure 5. Components of retraction fibers. PA-JEB/ β 4-EGFP were fixed and incubated with mAb 121 against plectin (A) or with mAb KL-1 against keratin (B), a mixture of mAbs B-5-1-2 and 2-28-33 against mouse α and β tubulins (C), mAb RV202 against vimentin (D), mAb P48 against CD151 (E), mAb K20 against β 1 (F), phalloidin Alexa 568 to stain actin filaments (G), mAb 3C12 against ezrin (H) and rabbit polyclonal anti–LN-5 antibodies (*H*). Texas-red– and Cy-5–conjugated secondary antibodies against mouse and rabbit IgG were used to detect bound antibodies. Composite images were generated by superimposition of the green and red signals, with areas of overlap represented in yellow. Where three colors were used (H), composite images were also generated by superimposition of the red, green, and Cy-5 signals, with areas of overlap represented in white. Because β 4-EGFP and keratin did not appear in the same confocal plane, (B) is presented as a maximum projection of XYZ stacks. β 1, actin, and ezrin, are not part of hemidesmosomes, but appear at the base of retraction fibers, and although present in the same structure are not colocalized with β 4-EGFP (F–H). The intermediate filament proteins, keratin and vimentin, and tubulin stain throughout the body of the cell and are not part of hemidesmosomes (B–D). The keratinoxyte remains still attached to its LN-5 matrix by retraction fibers (H). The hemidesmosomal components β 4-EGFP in hemidesmosomes but is dissociated during the conversion of these structures into retraction fibers (A). Bar, 10 μ m.

ever, ezrin, a protein that connects the cortical actin filaments with the plasma membrane, was detected in retraction fibers and, like F-actin, is more prominently localized at their base (Figure 5H). Ezrin is not present in hemidesmosomes. The β 4-EGFP and ezrin-positive fibers are bound to LN-5 left behind after the cells have moved (unpublished data), confirming that they are in fact retraction fibers. Taken together these results suggest that $\alpha 6\beta 4$ and $\beta 1$ integrins present in retraction fibers are not associated with the cellular cytoskeleton.

Stabilization of Membrane Extensions by β 4-EGFP Is Not Required for Cell Migration

In several reports it has been shown that $\alpha 6\beta 4$ associates with actin and is localized at the leading edge of invading carcinoma cells, where it is assumed to contribute to migration by stabilizing filopodia and lamellipodia (Rabinovitz *et al.*, 1997; O'Connor *et al.*, 1998; Goldfinger *et al.*, 1999; Decline and Rousselle, 2001). We, therefore, studied the localization of $\alpha 6\beta 4$ in PA-JEB/ $\beta 4$ -EGFP keratinocytes migrating into a wound bed. In the wound area the migrating cells were of two types. One type contained large lamellipodia with $\beta 4$ -EGFP evenly distributed over the cell (Figure 6), whereas in the other type $\beta 4$ -EGFP appeared at the leading edge. These cell types most likely represent different phases in the migration of the cell, the first type representing a cell that is actually migrating, whereas the migration of the cells with $\beta 4$ containing adhesion sites at their base has slowed down or perhaps even completely stopped. Although in the more stationary cells both $\alpha 3\beta 1$ and $\beta 4$ -EGFP are present at the

PA-JEB/β4-EGFP

PA-JEB/ EGFP-β4



Figure 6. During migration in a wound bed PA-JEB/ β 4-EGFP and PA-JEB/EGFP- β 4 form type I or type II hemidesmosomes. Localization by immunofluorescence of the hemidesmosomal components β 4, plectin, and BP180 in PA-JEB/ β 4-EGFP or PA-JEB/EGFP- β 4 keratinocytes crossing the wounded area after 48 h. Note the different cell types present. Fan-shaped cells with β 4-EGFP and EGFP- β 4 equally distributed over the cells (resulting in PA-JEB/EGFP- β 4 with hardly visible green fluorescence) most likely represent truly migrating cells. The other cells likely just stopped migrating and both β 4-EGFP and EGFP- β 4 appear at the leading edge of the cell, where α 3 β 1 is also present. Plectin is always colocalized with β 4-EGFP and EGFP- β 4 at the leading edge, whereas BP180 is occasionally present. Thus both type I and type II hemidesmosomes are formed by keratinocytes migrating into a wound bed. Bar, 20 μ m, except for (A), where it indicates 10 μ m.

leading edge, they are clearly not colocalized, $\alpha 3\beta 1$ being closer to the leading edge than $\beta 4$ -EGFP. Similar results were obtained with PA-JEB cells expressing EGFP- $\beta 4$. To investigate whether the hemidesmosomal structures formed during migration are type I or type II hemidesmosomes, PA-JEB/ $\beta 4$ -EGFP and PA-JEB/EGFP- $\beta 4$ keratinocytes crossing the wound bed were incubated with antibodies against plectin or BP180. As shown in Figure 6, although plectin was always found together with the $\beta 4$ -EGFP and EGFP- $\beta 4$ clusters at the leading edge of the keratinocytes, BP180 was only occasionally colocalized with them. Thus, PA-JEB/ $\beta 4$ -EGFP and PA-JEB/EGFP- $\beta 4$ keratinocytes form both type I and type II hemidesmosomes during migration.

The Integrin $\alpha 6\beta 4$ Slows Down Migration of Keratinocytes in Response to Wounding

The finding that $\alpha \beta \beta 4$ induces the formation of hemidesmosome-like structures during migration of PA-JEB/ $\beta 4$ -EGFP keratinocytes prompted us to further investigate the role of this integrin in cell migration using in vitro wound healing assays. PA-JEB/ $\beta 4$ -EGFP and PA-JEB/EGFP- $\beta 4$ keratinocytes were grown to confluence after which a scratch wound was introduced in the monolayer. Phase contrast micrographs taken at 48 h demonstrated that the migration of PA-JEB/ $\beta 4$ -EGFP keratinocytes is slower than that of PA-JEB/EGFP- $\beta 4$ keratinocytes, implicating an inhibitory effect of $\beta 4$ on cell migration when it can bind to its ligand LN-5 (Figure 7). Indeed, the mAb GoH3 known to block the adhesion of $\alpha 6\beta 4$ to LN-5 enhances migration of PA-JEB/ $\beta 4$ -EGFP cells. Furthermore, when the interaction of $\beta 4$ -EGFP with LN-5 is inhibited by the mAb GoH3, the $\beta 4$ -EGFP hemidesmosomes adopted the appearance of EGFP- $\beta 4$ hemidesmosomes (see characteristic patterns of $\beta 4$ -EGFP- and EGFP- $\beta 4$ -hemidesmosomes in Figure 2A), the clusters of hemidesmosomes were more clearly stained and defined, and there was no $\beta 4$ -EGFP in retraction fibers.

Because $\alpha 6\beta 4$ can interact with plectin and thereby become connected to the cytoskeleton, it is possible that the inhibition of migration by $\alpha 6\beta 4$ is not due to the increased adhesion to LN-5 but rather to the stabilization of this adhesion as a result of the interaction of $\alpha 6\beta 4$ with the cytoskeleton. To investigate this, we generated an EGFP version of a mutant $\beta 4$ subunit, $\beta 4^{R1281W}$, that can bind to LN-5 but not to plectin (Geerts *et al.*, 1999; Koster *et al.*, 2001). We found that when this mutant is stably expressed in $\beta 4$ negative PA-JEB keratinocytes, migration was hardly affected. In fact, the PA-JEB/ $\beta 4^{R1281W}$ -EGFP keratinocytes move as fast as PA-JEB/EGFP- $\beta 4$ keratinocytes (Figure 6). We conclude that the inhibition of migration by $\alpha 6\beta 4$ is due to the stabilization of the bond of $\alpha 6\beta 4$ and LN-5 through the interaction of $\beta 4$ with plectin.

FRAP Analysis of EGFP-tagged Fusion Proteins in Live Keratinocytes

FRAP was used to determine the dynamics of β 4-EGFP and EGFP- β 4 in hemidesmosomes of live keratinocytes. PA-JEB/



Figure 7. Cell motility is inhibited by binding of $\alpha 6\beta 4$ to plectin. PA-JEB/ $\beta 4$ -EGFP, PA-JEB/EGFP- $\beta 4$ and PA-JEB/ $\beta 4^{R1281W}$ -EGFP keratinocytes were grown to confluence on glass coverslips and treated for 2 h before wounding with 10 μ g/ml mitomycin C to prevent proliferation of cells. The monolayers were then wounded and cell migration was determined after 48 h in the presence or absence of $\alpha 6$ blocking antibodies. Expression of the LN-5 binding $\beta 4$ -EGFP chimera in PA-JEB cells slows down cell migration. Blocking of the $\alpha 6\beta 4$ -LN-5 interaction by using the $\alpha 6$ blocking antibody GoH3 at a concentration of 25 μ g/ml, enhanced migration of PA-JEB/ $\beta 4$ -EGFP keratinocytes and made it comparable to that of PA-JEB/EGFP- $\beta 4$ cells. There was no effect of $\alpha 6\beta 4$ on migration when a mutant of $\beta 4$ ($\beta 4^{R1281W}$), incapable of interacting with the cytoskeleton was used. Blocking of the $\alpha 6\beta 4$ -LN5 interaction did not promote migration. In the bar graph, results are expressed as mean \pm SD (n = 3).

β4-EGFP and PA-JEB/EGFP-β4 cells were examined by time-lapse two-photon excitation microscopy (Figure 8). After photobleaching, fluorescence of β4-EGFP had recovered for 32% (±2%) within the first 4 min and for 56% (±6%) after 10 min. The recovery of EGFP-β4 occurred much more slowly, i.e., 12% (±1%) after 10 min. There was no fast recovery in the first 4 min, and the rate of the EGFP-β4 recovery remained constant throughout time. Thus, in hemidesmosomes of live cells, EGFP-β4 is less dynamic than β4-EGFP. Because the β4-EGFP and EGFP-β4 chimeras only differ in their capacity to bind the ligand LN-5, this suggests that interaction with ligand increases the dynamics of α6β4. To investigate whether indeed the dynamics of β4 are de-

pendent on its interaction with LN-5, binding of β 4-EGFP to LN-5 was blocked by adding the α 6 blocking antibody GoH3 to PA-JEB/B4-EGFP keratinocytes, a treatment that results in a distribution of hemidesmosomes comparable with that in PA-JEB/EGFP- β 4 (see above). When these cells were subjected to FRAP analysis, recovery of fluorescence occurred as slowly as that in the case of EGFP- β 4. Thus, the bond between $\alpha 6\beta 4$ and LN-5 is not stable and can be broken, which makes the integrin a dynamic protein. In PA-JEB/ $\beta 4^{R1281W}$ -EGFP cells, clusters of $\beta 4^{R1281W}$ -EGFP can be observed. However, these clusters cannot be considered to be hemidesmosomes, because plectin, an essential component of hemidesmosomes, cannot bind to this β 4 mutant. FRAP analysis of $\beta 4^{R1281W}$ -EGFP was performed to investigate the contribution of the interaction of β 4 with the cytoskeleton to the dynamics of β 4. The recovery of fluorescence was faster with $\beta 4^{R1281W}$ -EGFP than with $\beta 4$ -EGFP. Within the first 4 min, fluorescence recovers for 58% ($\pm 4\%$), and eventually the recovery reaches 77% (\pm 5%) after 10 min. Diffusion coefficients for the different proteins were not calculated because the fluorescence of EGFP-B4 and B4-EGFP+GoH3 hardly recovered and because the fluorescence recovery curves for β 4-EGFP and β 4^{R1281W}-EGFP do not fit with a single exponential (see MATERIALS AND METHODS). However, the observed differences between the different recovery curves are consistent and striking. Thus, the bond between $\alpha 6\beta 4$ and LN-5 is less easily broken (i.e., less dynamic) when $\alpha 6\beta 4$ is associated with cytoskeleton.

DISCUSSION

The integrin $\alpha 6\beta 4$ is an essential component of hemidesmosomes and is necessary for tightly anchoring keratinocytes to the extracellular matrix (Jones *et al.*, 1998; Borradori and Sonnenberg, 1999). When epithelial cells are induced to migrate in response to wounding, they lose their hemidesmosomes, probably to reduce their strong adhesion to the substratum (Riddelle *et al.*, 1992; Gipson *et al.*, 1993). In carcinoma cells, $\alpha 6\beta 4$ contributes to migration and invasion by activation of PI-3 kinase signaling (Chao *et al.*, 1996; Shaw *et al.*, 1997). Thus $\alpha 6\beta 4$ plays a dual and apparently paradoxical role because it is essential both to keep cells stationary and to promote migration.

In this study, we analyzed the dynamics of $\alpha \beta \beta 4$ in stationary and migrating keratinocytes by expressing a $\beta 4$ subunit tagged with EGFP, in $\beta 4$ -deficient keratinocytes. Furthermore, the involvement of ligand binding in the dynamics of $\alpha \beta \beta 4$ was investigated by using an EGFP- $\beta 4$ chimera that is unable to bind to LN-5. Time-lapse videomicroscopy demonstrated that in moving and dividing keratinocytes, $\beta 4$ -EGFP and EGFP- $\beta 4$ hemidesmosome-like structures are assembled and redistributed within minutes. Unfortunately, we could not determine whether these hemidesmosomal structures are type I or type II hemidesmosomes, because the other hemidesmosomal components plectin, BP180, and BP230 could not be made visible during the time-lapse recordings.

During migration, the leading edge of lamellipodia and filopodia is the site where new adhesions are formed. Both in randomly moving cells and in cells that migrate into cleared areas after the monolayer has been wounded, clustered β 4-EGFP appears at the leading edge of the keratino-



Figure 8. Fluorescence recovery after photobleaching (FRAP) of EGFP chimeras is dependent on LN-5 binding. PA-JEB/EGFP-β4, PA-JEB/β4-EGFP, PA-JEB/β4^{R1281W}-EGFP keratinocytes and PA-JEB/β4-EGFP cells that had been treated with the α6 blocking antibody GoH3 were photobleached for 1 s using 100% laser power at t = 0. Fluorescence recovery was subsequently monitored by acquiring images every 15 s for a period of 10 min. Top panels: fluorescence images of cells before bleaching, immediately after bleaching, and 10 min after bleaching. The bleached area is depicted by a white oval. Bottom panels: representative recovery curves of fluorescence in the bleached area. The fluorescence was related to the initial fluorescence (stat 100%) and corrected for loss of fluorescence due to bleaching and scanning procedure. The presented figures represent a typical experiment of which four that were performed with similar results. After photobleach-ing, fluorescence of β4-EGFP had recovered for 32% (±2%) within the first 4 min and for 56% (±6%) after 10 min. The recovery of EGFP-β4 (±0.5%). The β4^{R1281W}-EGFP shows the fastest recovery of β4^{R1281W}-EGFP. Within the first 4 min the fluorescence recovers for 58% (±4%). Eventually, leading to a recovery of β4^{R1281W}-EGFP fluorescence of 77% (±5%) after 10 min. Thus, the dynamics of the interaction of β4 with LN-5 is suppressed by the association of β4 with the cytoskeleton.

cytes, as has also been shown by others (Goldfinger et al., 1999). The LN-5 binding integrin $\alpha 3\beta 1$ was also concentrated at these sites and was even closer to the leading edge of the cell than β 4-EGFP. Several reports have suggested a role of $\alpha 6\beta 4$ in stabilizing newly formed filopodia and lamellipodia in order to facilitate migration (Rabinovitz and Mercurio, 1997; O'Connor et al., 1998). However, we show that EGFP-B4, which cannot interact with LN-5, also becomes concentrated at these sites. This suggests that β 4 is involved in a different process, i.e., the formation of new hemidesmosomes. This assumption is supported by the colocalization of plectin and, in some cells of BP180 and BP230, with EGFP- β 4 at these sites. In contrast, in highly motile cells displaying a characteristic fan-shaped morphology, neither β 4-EGFP nor EGFP- β 4 are clustered but are diffusely distributed throughout the cell. The mechanism responsible for the localization of EGFP- β 4 at the leading edge is not known, but we assume that it is targeted by a direct or indirect association with $\alpha 3\beta 1$. A role of the latter integrin in hemidesmosome formation has previously been suggested (Nievers *et al.*, 1998, 2000; Sterk *et al.*, 2000) and is supported by the strong reduction in the number of hemidesmosomes in mice that do not express $\beta 1$ in the skin (Brakebusch *et al.*, 2000); Raghavan *et al.*, 2000).

Cell migration not only depends on the formation of cellmatrix adhesion by the moving cell, but also on breaking existing adhesions at the rear end of the cell. (Regen and Horwitz, 1992; Palecek *et al.*, 1998). If these adhesions are strong, they will be less easily broken, which will result in the appearance of retraction fibers, causing migration to be slowed down. In PA-JEB/ β 4-EGFP keratinocytes many retraction fibers can be observed that originate from hemidesmosomes. These retraction fibers contain β 4-EGFP, but no plectin, which suggests that in them the linkage of β 4 with the cytoskeleton is fractured. Indeed, the intermediate filament proteins keratin and vimentin, to which plectin can bind, are not present in these retraction fibers. Incidentally, some of the β 4 is ripped from the membrane and remains attached to the LN-5 matrix, leaving "footprints" of a migrating cell.

The detachment of plectin from B4 during the transition of hemidesmosomes into retraction fibers may be due to either mechanical stress or signaling events. Several studies have shown that phosphorylation of the β 4 subunit by protein kinase C is associated with a redistribution of $\alpha 6\beta 4$ from the hemidesmosome to the cytosol (Alt et al., 2001) and/or to F-actin-rich cell protrusions (Rabinovitz et al., 1999). Furthermore, the link between β 4 and plectin might be cleaved by proteases such as the Ca²⁺-dependent protease calpain. This stimulates rear end release of CHO cells (Palecek et al., 1998) by cleavage of cytoskeletal linkages. In fact, calpain cleavage sites are present in the cytoplasmic domain of the β 4 subunit (Giancotti *et al.*, 1992). The reverse reaction, the reassociation of plectin with $\alpha 6\beta 4$ occurs when retraction fibers convert into hemidesmosomes, which occurs prominently after mitosis (see Figure 4A).

Our study clearly shows that the introduction of β 4-EGFP into a cell that lacks β 4 slows down the migration of that cell. Previous work has shown that the proteolytic processing of the α 3 chain of LN-5 by plasmin produces an LN-5 molecule that induces the assembly of hemidesmosomes and impedes cell migration (Goldfinger et al., 1998). On the contrary, proteolytic processing of the γ 2 chain of LN-5 has been associated with the induction of cell migration (Giannelli et al., 1997). Although the relative amounts of processed $\alpha 3$ chain in the matrices produced by PA-JEB/β4-EGFP and PA-JEB/EGFP- β 4 cells were slightly different, it is unlikely that this accounts for the observed differences in their migration. In fact, the migration of PA-JEB/ β 4-EGFP cells that produce the most unprocessed α 3 is the slowest. Extracellular processing of the γ 2 chain of LN-5 was not observed, and furthermore there was no obvious difference in the amount of LN-5 deposited by the two cell lines into the matrix. Thus, the difference in migration of PA-JEB/B4-EGFP and PA-JEB/EGFP-β4 cells cannot be due to differences in the proteolytic state or the amount of LN-5 molecules produced by them.

It is possible that the increased adhesion of cells to LN-5, due to the expression of β 4-EGFP, by itself might already slow down migration. However, studies with a mutant β 4 subunit ($\beta 4^{R1281W}$ -EGFP) that cannot interact with plectin and therefore also not with the cytoskeleton (Geerts et al., 1999; Koster et al., 2001) showed that it is in fact the stabilization and not the magnitude of the adhesion to LN-5 that causes this effect. Plectin, which is a dimeric protein but also has been shown to form tetramers (Wiche, 1998), might stabilize the adhesion of $\alpha 6\beta 4$ to LN-5 by linking two or more $\alpha 6\beta 4$ molecules together. However, for this linking to occur, the $\alpha 6\beta 4$ molecules must be close together. We have shown previously that when keratinocytes are seeded on a subconfluent LN-5 matrix, hemidesmosomes are distributed in a pattern that mirrors the pattern of LN-5 underneath the cell (Nievers *et al.*, 2000), which implies that β 4 preferentially interacts with LN-5, rather than with plectin. Thus, it is likely the density of LN-5 molecules underneath the cell that ultimately determines whether the interaction of $\alpha 6\beta 4$ with LN-5 will be stabilized. Accordingly, when LN-5 is not concentrated into patches but is distributed diffusely over a surface, $\alpha 6\beta 4$ will not be clustered. In that case, an interaction of $\alpha 6\beta 4$ with plectin will not stabilize the adhesion of $\alpha 6\beta 4$ to LN-5, and migration will not be slowed down. Indeed, we recently found that in $\beta 4$ -deficient PA-JEB cells transfected with $\beta 4$, the $\alpha 6\beta 4$ integrin accelerates migration on surfaces with homogenously distributed LN-5 (G. Smits and A. Sonnenberg, unpublished observations).

Hemidesmosome-like structures that are formed by β 4-EGFP on the patches of LN-5, deposited by the cells themselves, appeared to be less well clustered than the ones that are formed by EGFP- β 4. This may indicate that in these hemidesmosome-like structures, not all of the bindings between $\alpha 6\beta 4$ and LN-5 are stabilized by interactions of $\beta 4$ with plectin. Consistent with this notion, FRAP analysis showed that the recovery of fluorescence of β 4-EGFP was faster than that of EGFP- β 4, or of β 4-EGFP of which the binding to LN-5 was prevented by the mAb GoH3. In fact, hardly any recovery of fluorescence of EGFP-B4 or of B4-EGFP/GoH3 could be detected over a period of 10 min. On the other hand, the mutant β 4 subunit (β 4^{R1281W}-EGFP) that cannot bind to plectin displayed the fastest recovery of fluorescence. Thus, the binding dynamics of $\alpha 6\beta 4$ interaction with LN-5 on the one hand and with plectin on the other hand influence each other in a reciprocal manner.

Why on the LN-5 patches not all of the interactions between $\alpha 6\beta 4$ and LN-5 are stabilized by plectin is not clear. It is possible that the density of the LN-5 molecules in the patches is too low and that therefore not all of the $\alpha 6\beta 4$ molecule are close enough to each other for their interaction with LN-5 to be stabilized by plectin. Alternatively, not all the LN-5 molecules in the matrices may be optimally presented. This might also explain why in tissue culture, hemidesmosomes almost never reach the level of maturation of true hemidesmosomes in the skin, i.e., with discernible inner and outer electron-dense plaques. In vivo, the presentation and the amount of LN-5 that is present under hemidesmosomes may not be a limiting factor and clustering and binding of $\alpha 6\beta 4$ to the cytoskeleton might therefore be more extensive, allowing the formation of well-defined hemidesmosomes. Indeed, it has been shown that LN-5 is much more concentrated under hemidesmosomess than in regions where no hemidesmosomes are present (Rousselle et al., 1991).

Altogether our results suggest that the density of LN-5 underneath the cell ultimately determines to what extent the binding of $\alpha 6\beta 4$ to LN-5 will become stabilized by interaction of $\beta 4$ with plectin and whether $\alpha 6\beta 4$ will inhibit migration or not.

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C.A.W. Geuijen and A. Sonnenberg

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