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The key word ‘Integrin segregation’ was spelled incorrectly in the print version of this article. The online version is correct.

We apologise for this mistake.

The mechanical integrin cycle

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Summary

Cells govern tissue shape by exerting highly regulated forces at sites of matrix adhesion. As the major force-bearing adhesion-receptor protein, integrins have a central role in how cells sense and respond to the mechanics of their surroundings. Recent studies have shown that a key aspect of mechanotransduction is the cycle by which integrins bind to the matrix at the leading cell edge, attach to the cytoskeleton, transduce mechanical force, aggregate in the plasma membrane as part of increasingly strengthened adhesion complexes, unbind and, ultimately, are recycled. This mechanical cycle enables the transition from early complexes to larger, more stable adhesions that can then rapidly release. Within this mechanical cycle, integrins themselves exhibit intramolecular conformational change that regulates

their binding affinity and may also be dependent upon force. How the cell integrates these dynamic elements into a rigidity response is not clear. Here, we focus on the steps in the integrin mechanical cycle that are sensitive to force and closely linked to integrin function, such as the lateral alignment of integrin aggregates and related adhesion components.

Supplementary material available online at
<http://jcs.biologists.org/cgi/content/full/122/2/179/DC1>

Key words: Cell-matrix adhesion, Integrin dynamics, Mechanotransduction, Vinculin, Talin, Integrin seregation, Adhesion remodelling

Introduction

Cells shape tissues by pulling on neighboring cells and extracellular matrices (ECMs), creating specific levels of tension. In turn, cells are finely attuned to the forces and rigidity of their surroundings. As rigidity is defined by the force per unit displacement, rigidity-sensing cells must measure both force and displacement. Across different tissue types, rigidities are in the range of 1–100 kPa, from the softness in which fat cells or neurons thrive, to the relative stiffness that is home for chondrocytes (Discher et al., 2005). Yet, despite differences in tissue and cell types, force and position are crucial aspects of many, if not most, cell-matrix interactions. For example, fibroblasts and endothelial cells periodically contract fibronectin to test its rigidity (Giannone et al., 2004). When that process is analyzed in detail, it appears to be controlled by a series of mechanical steps that result in periodic rows of $\alpha V\beta 3$ -integrin aggregates together with early adhesion components (Giannone et al., 2007). Similarly, to produce a uniform displacement on substrates of increasing rigidity, epithelial cells recruit additional motor proteins to generate higher forces (Saez et al., 2005). This implies a rapid feedback mechanism at sites of integrin-mediated attachment, between the rigidity-sensing system and the force-producing machinery.

A cell, probing its environment, initiates matrix adhesion through actin-dependent protrusions that bring integrins at the leading edge in contact with the matrix where they can bind (Fig. 1A). The binding of integrin to the ECM is rapidly followed by integrin binding to the actin cytoskeleton, which is typically moving inwards from the site of assembly at the leading edge towards the cell center. Thus, a pulling force is quickly generated across nascent integrin-matrix linkages (Fig. 1B). Within seconds, these initial sites of integrin-ECM linkage begin to strengthen as additional components are recruited under force (Fig. 1C) (Galbraith et al., 2002; von Wichert et al., 2003b). On a larger scale, the pulling force also acts to either pull the matrix over the cell or pull the cell over the matrix. Following this matrix movement or, in the case of cell migration, following this cell movement, the integrins then release

from the matrix. When forces are sufficiently high – that is, when the substrate is sufficiently rigid – sites of integrin-mediated adhesion undergo further maturation, extending anisotropically several μm in length as additional proteins are recruited. These centripetally polarized supramolecular structures have been termed ‘focal adhesions’. In supplementary material Fig. S1, a single integrin is highlighted during each step of this mechanical cycle. The mechanism of cell-derived tension that drives this mechanical cycle is described in Box 1.

Force-dependent strengthening of adhesion sites is remarkable, because all receptor-ligand bonds eventually break under high forces. For example, the lifetime of the notoriously strong avidin-biotin bond is reduced from more than a day to ~1 minute under a force of 5 pN (Merkel et al., 1999). Two logical explanations for force-dependent adhesion strengthening are increased recruitment of integrin receptors, which is known to occur when integrin-mediated adhesions exhibit physical growth under force, and/or catch bonds, receptor-ligand complexes that exhibit an increased lifetime under mechanical load. Moreover, as the proteins in these force-regulated adhesion sites turn over rapidly to enable cell spreading and migration, their strength and position are also remodeled under force. Cells clearly control this cycle of integrin-dependent attachment, force production and release to generate precise tissue morphologies. A guiding question at the forefront of current cell science is how these sites of integrin-mediated adhesion participate in this process of cell-tissue morphodynamics.

In an effort to understand how integrin-mediated force regulates protein recruitment and the strengthening of focal-adhesions, physical models have been derived. For example, elastic strain is proposed to induce anisotropic protein aggregation under force because of the resulting asymmetric extension and compression of the focal-adhesion entity (Besser and Safran, 2006; Nicolas et al., 2004). Alternatively, protein aggregation under force has been proposed to be the result of a purely thermodynamic process, whereby stress-induced changes in the chemical potentials of focal-adhesion proteins are compensated for by the binding of additional focal-adhesion proteins (Shemesh

Box 1. Cells steer adhesion-site maturation by forces that are generated through actin assembly and actomyosin contractions

In the most peripheral, 1–4- μm region of a spreading cell, termed the lamellipodium, relatively fast rates of actin assembly and disassembly generate force on the cell membrane. This force is comparatively small (Dubin-Thaler et al., 2008; Raucher and Sheetz, 2000) and has been linked to the initiation of integrin-mediated adhesion (Alexandrova et al., 2008; DeMali et al., 2002).

Just behind the lamellipodium, in a region termed the lamellum, the retrograde actin flow is slower. The boundary between the lamellipodium and the lamellum appears to be regulated by the development of nascent adhesion sites into mature focal adhesions (Alexandrova et al., 2008). The contraction of actomyosin filament bundles, called stress fibers, drives the growth of focal complexes into focal adhesions. Myosin II A and B are the isoforms responsible for generating this force (Cai et al., 2006). Whereas myosin II inhibition blocks the conversion from focal complexes to focal adhesions, the application of a local, external pulling force can replace the role of actomyosin contractility and restore focal-adhesion formation (Riveline et al., 2001).

Force generated from actomyosin contractions is also thought to contribute to early adhesion-site formation in the lamellipodium (Galbraith et al., 2002). In support of this notion, adhesion-site formation at the cell edge has been found to have the same periodicity as myosin-mediated contractions (Giannone et al., 2004). Recently, lamellipodium actin has been shown to form an adaptable, mechanical link between the site of an actin polymerization at the leading edge and the myosin motor activity in the lamellum (Giannone et al., 2007). Thus, the interplay between forces generated by actin polymerization and by myosin-mediated contractility work cooperatively to drive the integrin mechanical cycle (Dubin-Thaler et al., 2008).

et al., 2005). Although these models describe general physical mechanisms of how force governs focal-adhesion morphodynamics, recent experimental and computational findings have revealed how mechanical stress directly regulates the function of integrins and several other associated molecules, such as talin and vinculin. Further, force-dependent tyrosine phosphorylation can result in dramatic changes in signaling pathways that alter integrin-ligand binding (Tamada et al., 2004; Sawada et al., 2006). Together, an understanding of how different signals affect the distinct molecular dynamics of the mechanical integrin adhesion cycle can illuminate the basis of changes in cell and tissue shape.

In this Commentary, we will focus on the integrin-dependent motility that has been studied extensively in primarily mammalian fibroblastic, immune and endothelial cells, with many corresponding features also evident in stem cells and cancer cells. First, we will discuss intramolecular integrin dynamics, which govern integrin activation. Second, we will consider focal-adhesion assembly in terms of integrin aggregation under force, which exhibits a linear directionality that may regulate intracellular signaling pathways.

Integrin activation under force

An understanding of integrin-mediated mechanosensing begins with integrin activation, which governs integrin-binding kinetics and clustering (Cluzel et al., 2005; Kim et al., 2004). Integrin activation occurs allosterically, involving long-range intramolecular conformational changes that can originate from the extracellular or

cytoplasmic end of the integrin heterodimer. Integrin heterodimers comprise non-covalently bound α - and β -subunits, which associate to form the extracellular ligand-binding head, two multi-domain ‘legs’, two single-pass transmembrane helices and two short cytoplasmic tails. All known integrin heterodimers contain the βA domain (also called the I-like or βI domain), which is located at the extracellular end of the β -subunit. Mutational and monoclonal-antibody experiments have shown that the switch from low- to high-binding affinity in the ECM-binding integrin headpiece involves an increase in the hinge angle between the βA - and hybrid-domains (Luo et al., 2003; Luo et al., 2004; Mould et al., 2003). X-ray crystallographic structures provide the stationary endpoints of this conformational switch in the unliganded closed-hinge and the ligand-bound open-hinge β3 -integrin headpiece domains (Xiao et al., 2004; Xiong et al., 2001). Molecular dynamics (MD) simulations of the β3 -integrin headpiece domains have illustrated the Ångstrom-level structural pathway of ligand-induced hinge-angle opening (Puklin-Faucher et al., 2006).

One hallmark of allosteric proteins such as integrins is their bi-directionality, which means that the same activating structural pathway can be induced by extracellular (‘outside in’) or intracellular (‘inside out’) factors (Hynes, 2002). In vivo events that are known to activate integrins are the ligand binding by the extracellular head (Takagi et al., 2002) or the talin binding by the intracellular tail of the β -subunit (Tadokoro et al., 2003). In the absence of force, integrin activation occurs within seconds. However, there are clearly mechanical signals that can induce events downstream of activation in seconds, such as integrin aggregation (Giannone et al., 2004) and adhesion-protein assembly (Galbraith et al., 2002; Riveline et al., 2001; von Wichert et al., 2003b). In the case of T cells, firm integrin adhesiveness was shown to be tightly regulated by mechanical signals that involve the combination of force from shear-fluid flow and immobilized chemokines (Woolf et al., 2007). Consistent with these observations, when the application of ligand-mediated mechanical force was simulated in steered MD (SMD) investigations, it was shown to accelerate the allosteric pathway to activation in the integrin headpiece to the sub-microsecond timeframe (Puklin-Faucher et al., 2006). As binding to ECM ligands is known to activate integrins under equilibrium conditions (Takagi et al., 2002) and binding is needed for force to be transduced across integrins, the major effect of force on integrin activation may be in accelerating the allosteric activation pathway and, thereby, in stabilizing bonds that would otherwise dissociate within sub-seconds.

It is logical to postulate that the application of a force vertical to the membrane would induce an activating conformational change in integrins (as shown in Fig. 1). Such a force could be generated even though the force vector is often almost parallel to the membrane, rather than perpendicular. $\alpha\text{V}\beta\text{3}$ integrins can stably bind fibronectin with only a modest ($\sim 11^\circ$) increase in the angle of their headpiece hinge and with a severe bend (of $\sim 135^\circ$, based on crystallographic data) in their extracellular legs (Adair et al., 2005). After matrix is bound, this bond could potentially be stabilized by force-induced conformational change. For example, with only the β - and not the α -cytoplasmic tail linked to the cytoskeleton, force could vary the interdomain headpiece hinge via separation of the heterodimer legs as the β -subunit becomes aligned along the force vector.

Intracellularly, binding of the talin head to the β -subunit of the integrin tail has been shown to activate integrins by disrupting membrane-proximal and transmembrane associations with the

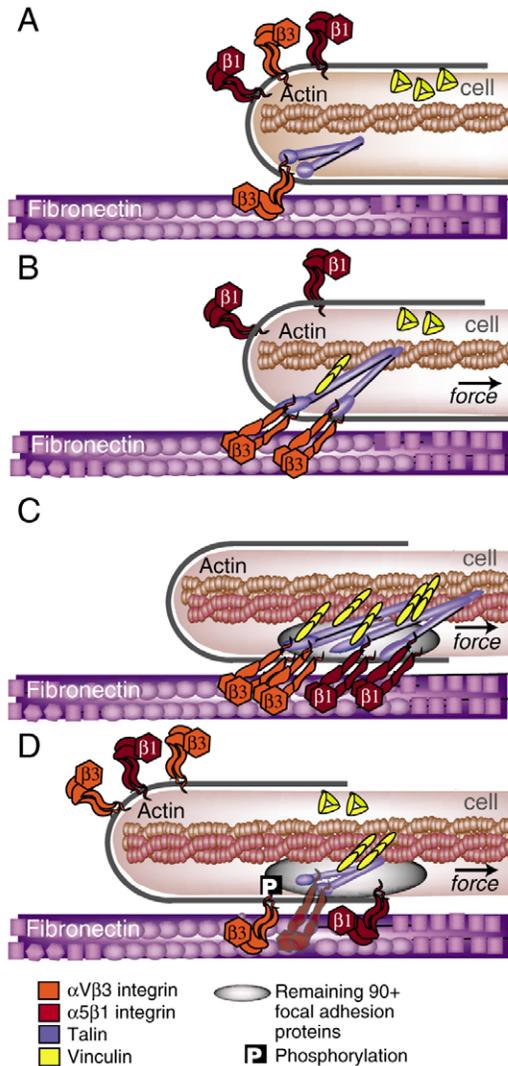


Fig. 1. The mechanical integrin cycle. (A) Cell-ECM adhesion occurs when actin-dependent protrusions bring integrins at the leading edge (orange) in contact with the matrix (purple) where they can bind. (B) Next, the integrins link to the actin cytoskeleton through adaptor proteins, such as talin (blue), Shp2, filamin or α -actinin. Integrins bind to these adaptor proteins through their β -tails. Rearward actin flow, generated by actin polymerization and actomyosin contractions (see Box 1) induces a pulling force on the integrin-ECM linkage. On sufficiently rigid substrates, this may serve to accelerate an integrin-activating conformational change, as well as a talin stretch, which may expose buried vinculin-binding sites (yellow). Although the bent conformation of the ligand-bound α V β 3-integrin crystal structure produced much controversy in the integrin field (Liddington and Ginsberg, 2002; Mould et al., 2003), it has subsequently been shown in electron microscopy experiments to stably bind fibronectin (Adair et al., 2005). Force might accelerate the switch to high-binding affinity by freeing the ligand-bound integrin head from the constraints of neighboring domains, which would essentially accelerate the allosteric pathway to the activated state (Puklin-Faucher et al., 2006). (C) The cell begins to pull itself over the site of adhesion. Intramolecular conformational changes in α 5 β 1 integrins facilitate their inward translocation, whereas α V β 3 integrins remain anchored at the edge. This segregation of integrins may further facilitate the talin stretch. At this stage of adhesion, a wide variety of intracellular focal-adhesion proteins are accumulated in the adhesive plaque (grey oval). (D) Ultimately, highly clustered integrins switch from high- to low-binding affinity, possibly catalyzed by the phosphorylation of β 3-integrin tails. Membrane exocytosis places recycled, low-affinity integrins at the end of microtubules, often 2–4 μ m away from the leading edge. The integrin turnover in focal adhesions (from C to D) is ~1–3 minutes (Hu et al., 2007). For the description of a single integrin see supplementary material Fig. S1.

neighboring α -subunit domains (Tadokoro et al., 2003; Wegener et al., 2007). Recently, the structurally homologous kindlin family of proteins has been shown to interact directly with β 3- and β 1-integrin tails and to catalyze (kindlin-2, also known as FERMT2) or even supersede (kindlin-3, also known as FERMT3) integrin activation by talin (Ma et al., 2008; Moser et al., 2008). To influence the ECM-binding affinity of the integrin head, the structural change induced by kindlin and talin at the integrin tails must propagate across the multiple leg domains of the ~28-nm-long integrin molecule. As described above, ligand-mediated force may accelerate this allosteric structural change (Alon and Dustin, 2007; Puklin-Faucher et al., 2006).

As the highly flexible integrin β -tails provide a scaffold for a wide range of cytoskeletal proteins (Calderwood et al., 2003) and are extremely flexible, there is also the possibility that ligand-mediated force could accelerate binding to kindlin and talin by making the binding sites more accessible through disruption of the membrane-proximal and transmembrane integrin-heterodimer associations. In support of this, the presence of the head part of talin – but not the rod – appears to stabilize integrin binding to fibronectin even in the absence of actin binding (Zhang et al., 2008). The binding of single fibronectin trimers is highly dependent upon talin, as is a weak slip bond with the actin cytoskeleton (Jiang et al., 2003). Also, swapping α - and β -tails blocked lateral integrin aggregation, but moving the β 1 tail further from the membrane by lengthening the membrane-proximal domain of the α 5-chimera with a spacer restored the lateral aggregation that was dependent upon the β 1 tail (Partridge et al., 2006). This result implies that allowing the distal β -cytoplasmic domain to adopt a distinctive conformation by freeing it from the proximal α -cytoplasmic domain is the structural event that drives aspects of ligand-dependent integrin signalling such as lateral aggregation. Together, these findings imply that the physical unmasking of kindlin- and talin-binding sites on the integrin β -tail can stabilize their structural and functional state (Ulmer et al., 2003). Although there is considerable evidence that the early linkages between integrins and the cytoskeleton depend upon kindlin-2, kindlin-3 and talin, there are other integrin-tail-binding partners that can also link to the contractile actin cytoskeleton in other adhesion processes. These include filamin, α -actinin, melusin, SH2-domain-containing protein-tyrosine phosphatase (Shp2), skelemin, integrin-linked kinase and, possibly, myosin (Phillips et al., 2001; Critchley and Ginggras, 2008; Kiema et al., 2006; Pavalko et al., 1991; von Wichert et al., 2003a).

Force generation, integrin segregation and adhesion growth

Nascent focal adhesions, termed ‘focal complexes’, mediate high forces relative to mature focal adhesions (Beningo et al., 2001) and grow in size in linear proportion to the traction forces exerted upon them (Balaban et al., 2001). These traction forces are, in turn, directly proportional to the rigidity of the extracellular substrate (Saez et al., 2005). During mechanosensing, the density of integrins that surround these sites of attachment increases in a directional fashion. As shown schematically in Fig. 1C and with antibody staining in Fig. 2, α V β 3 integrins remain anchored at the distal end, closer to the leading edge, and α 5 β 1 integrins translocate to the proximal end, closer to the center of the cell. The spatial segregation of α 5 β 1- from α V β 3-integrins has been well established during the transition from focal to fibrillar adhesions, when cells create extracellular fibrils from plasma fibronectin (Pankov et al., 2000; Zamir et al., 2000). During this process, termed ‘fibrillogenesis’, α 5 β 1 integrins translocate

inwards by a distance of $\sim 10 \mu\text{m}$. This movement along the actin cytoskeleton serves to elongate and organize newly formed fibrils into the ECM. In the absence of fibrillogenesis, for example, when the fibronectin matrix is non-deformable (Katz et al., 2000), $\alpha 5\beta 1$ integrins also aggregate in a linear fashion inside focal adhesions. In this fashion, mature focal adhesions exhibit a linear, centripetal spatial segregation of $\alpha V\beta 3$ - and $\alpha 5\beta 1$ -integrins that extends ~ 5 - $6 \mu\text{m}$ along contractile actomyosin bundles (Felsenfeld et al., 1999). As this highly regulated integrin activity is governed by a force-dependent structural change and follows the primary force vector, it is logical to propose that force and rigidity choreograph this directionality in integrin dynamics.

At the leading edge of mechanosensing cells, $\alpha V\beta 3$ integrins form the nascent cell-ECM contacts and mediate initial force-accelerated adhesion-strengthening events (von Wichert et al., 2003b). For example, during the early stages of cell-adhesion-contact formation, activation of the Src family kinase Fyn requires that surface-bound $\alpha V\beta 3$ integrins interact with receptor-like protein tyrosine phosphatase α (RPTP α) (Su et al., 1999; von Wichert et al., 2003b). This RPTP α -induced activation of Fyn at the leading edge of rigidity-sensing cells is necessary for the force-dependent strengthening of $\alpha V\beta 3$ -integrin-cytoskeleton connections (Jiang et al., 2006; von Wichert et al., 2003b).

Similar to $\alpha V\beta 3$ integrins, $\alpha 5\beta 1$ integrins have also been shown to bind to the ECM at the leading edge of the cell and translocate inwards (Nishizaka et al., 2000). Release of $\alpha 5\beta 1$ integrins from fibronectin-coated beads at the back of the lamellipodium, where adhesions often end, was observed in experiments using optical tweezers (Nishizaka et al., 2000). Humphries and co-workers have now shown that the inwards translocation of $\alpha 5\beta 1$ integrins within focal adhesions corresponds to a series of distinctive conformational changes in the extracellular domains of $\alpha 5\beta 1$ integrins, namely from bent to straight and then to separated (Clark et al., 2005). Similarly, a recent study shows that the inwards translocation of $\alpha 5\beta 1$ integrins causes centripetal focal-adhesion orientation but relies upon the extracellular binding interactions and the subsequent conformational changes of $\alpha 5\beta 1$ - but not $\alpha V\beta 3$ -integrins (Huvencers et al., 2008). Alternative explanations for the spatial segregation of $\alpha V\beta 3$ - and $\alpha 5\beta 1$ -integrins include integrin-recycling pathways (White et al., 2007) and retrograde flux of actin filaments (Guo and Wang, 2007).

Talin

Talin binds directly to $\beta 1$ -, $\beta 2$ - and $\beta 3$ -integrins. Recently, differences in the way that talin interacts with integrins have come to light. For instance, although a small fragment of the N-terminal talin head domain, the F3 sub-domain, is sufficient to activate $\beta 3$ integrins (Calderwood et al., 2003; Garcia-Alvarez et al., 2003), this sub-domain is not sufficient to activate $\beta 1$ integrins. To produce detectable $\beta 1$ -integrin activation, the entire ~ 50 kDa talin head is required (Bouaouina et al., 2008).

Similar to activation, clustering has structural origins. Clustering of $\alpha V\beta 3$ - and $\alpha 5\beta 1$ -integrins, which follows activation (Kim et al., 2004), is driven by interactions with the talin head domain (Cluzel et al., 2005; Zhang et al., 2008). Talin also contains a second integrin-binding site in its C-terminal rod domain, which directly interacts with both $\beta 3$ - and $\beta 1$ -integrin tails (Parsons et al., 2008; Tremuth et al., 2004; Xing et al., 2001). Interaction with the talin rod, however, does not lead to integrin activation; rather, it provides the link to the cytoskeleton, thus enabling the substrate traction forces that are necessary for sustained cell spreading (Moes et al., 2007; Zhang et al., 2008). It has been suggested that the

integrin-activating conformational change that is induced by binding of the talin head may result in the exposure of a de novo high-affinity integrin-binding site for the talin rod (Moes et al., 2007). In line with this idea, a recent fluorescence resonance energy transfer (FRET) analysis identified specific interactions between $\beta 1$ integrins and the talin rod domain but not the talin head domain, leading the authors to suggest that interactions between the $\beta 1$ -integrin and the talin head domain may be transient and/or restricted to early adhesion complexes (Parsons et al., 2008).

The initial binding of $\beta 1$ integrins to talin is linked to their synchronous sideways movement when clustered at the leading edge by polymerizing actin fibers (Galbraith et al., 2007). Clustering of $\alpha 5\beta 1$ integrins has been linked with the transition from focal adhesions into fibrillar adhesions (Clark et al., 2005). When ligand-dependent translocation of $\alpha 5\beta 1$ integrins in human fibroblasts was blocked and integrin clustering was then induced with monoclonal antibodies, movement of $\alpha 5\beta 1$ integrins out of focal adhesions into fibrillar adhesions was observed. Thus, clustering of $\alpha 5\beta 1$ integrins can independently drive their directional, centripetal translocation into fibrillar adhesions (Clark et al., 2005), where talin is replaced by the adaptor protein tensin (Pankov et al., 2000). Together, these findings point to a model in which talin, which is ~ 60 nm long, may ultimately be oriented inside focal adhesions by head and rod contacts with centripetally organized $\alpha V\beta 3$ - and $\alpha 5\beta 1$ -integrins. Force-induced deformation of the integrin-talin linkage may then facilitate exchange to the adapter protein tensin.

Although we are focused here on mammalian integrin interactions, it is interesting to note the similarities and differences relative to *Drosophila melanogaster* that have recently come to light. Although the integrin link to the cytoskeleton via the talin rod is conserved in both species (Tanentzapf and Brown, 2006), talin is not sufficient to activate the $\beta 1$ -integrin orthologue in *Drosophila* (termed βPS) (Helsten, 2008). Whereas the integrin link to the cytoskeleton through the talin rod is conserved in both species (Tanentzapf and Brown, 2006), talin is not sufficient to activate *Drosophila* βPS integrins (Helsten et al., 2008). The authors of this recent finding speculate that the regulation of integrin activation by talin in mammalian cells may have developed "...later in vertebrate evolution to provide exquisite regulation of integrin affinity in highly motile cells" (Helsten et al., 2008).

Vinculin

The talin rod contains multiple binding sites for the adaptor protein vinculin, which is known to be involved in focal-adhesion dynamics (Coll et al., 1995; Volberg et al., 1995) and is recruited to sites of talin-integrin adhesion under tension (Galbraith et al., 2002; Zaidel-Bar et al., 2003). Importantly, talin's binding sites for vinculin are buried inside α -helix bundles under equilibrium conditions and have been shown in silico to be exposed by stretching force in recent SMD simulations (Hytonen and Vogel, 2008). Indeed, traction forces on integrin-mediated adhesions have been shown to result in recruitment of vinculin within tens of seconds (Galbraith et al., 2002) and vinculin activity, in turn, regulates both paxillin recruitment and integrin turnover (Humphries et al., 2007). Currently, direct experimental verification of vinculin binding-site exposure by talin mechanical stretch, as illustrated in Fig. 1A-C, is lacking.

Notably, a decisive factor in adhesion strengthening is not simply the increase in integrin density, but rather the distance between individual integrin molecules. As revealed in a recent study of integrin adhesion strengthening, cyclic Arg-Gly-Asp

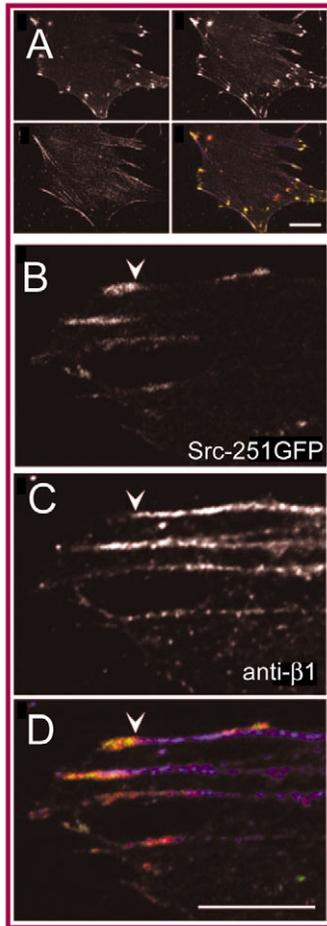


Fig. 2. Segregation of α V β 3- and α 5 β 1-integrins in focal adhesions. The segregation of α V β 3- and α 5 β 1-integrins in focal adhesions (depicted schematically in Fig. 1D) is shown here by antibody staining (Felsenfeld et al., 1999). (A) Src-deficient cells that expressing a truncated form of Src tagged with GFP (Src-251GFP) were fixed and stained with antibodies recognizing β 1 integrins, α V β 3 integrins and vinculin. (B) In higher-magnification views, the α V β 3 subunit can be seen to colocalize at the periphery of focal adhesions, with vinculin and Src-251GFP. Vinculin staining was co-distributed with that of Src-251GFP in all cases. (C) By contrast, β 1 integrins distributed in longer peripheral stripes (consistent with the distribution of stress fibers) that did not overlap with the distribution of α V β 3-Src-251GFP. Arrowhead indicates the boundary of staining. (D) Overlap of GFP and vinculin staining without β 1 integrins is indicated by yellow pixels. Scale bars: 10 μ m (A) and 5 μ m (D). Images reproduced with permission (Felsenfeld et al., 1999).

(RGD) ligands, which preferentially bind α V β 3 integrins, need to be spaced at a distance of 55 nm or less for adhesions to be reinforced (Selhuber-Unkel et al., 2008) and vinculin molecules to bind (Cavalcanti-Adam et al., 2006). Vinculin also assists α 5 β 1-integrin clustering through a tight association with talin (Humphries et al., 2007). Similar to mammalian talin, the N-terminal head and C-terminal tail domains of mammalian vinculin have been pinpointed as being the locations that support clustering and mechanical linkage to the actin cytoskeleton, respectively (Humphries et al., 2007). Together, these findings portray a model of focal-adhesion dynamics in which clustered cytoskeletal-integrin-ECM linkages are crosslinked by talin homodimers, which then regulate integrin dynamics by recruiting vinculin when stretched in a directional fashion (Fig. 1C).

Force acceleration of biochemical pathways

In addition to spatial arrangements, temporal changes are a crucial component of integrin-mediated mechanotransduction. This is illustrated on the intramolecular scale in MD and SMD simulations of the integrin headpiece, which show that mechanical force accelerates the same allosteric pathway to hinge-angle opening that is induced by ligand binding under equilibrium conditions (Puklin-Faucher et al., 2006). Currently, direct experimental verification of integrin activation by mechanical force is lacking, owing in part to the tendency of integrins to cluster upon adhesion and thus obscure measurements of monovalent substrate interactions. On the intermolecular scale, vinculin and talin can drive integrin clustering in focal adhesions independently of tensile force (Humphries et al., 2007). Similarly, in mature focal complexes, α V β 3 integrins interact with Src family kinases directly (Arias-Salgado et al., 2003), versus interacting through RPTP α in nascent integrin-ECM linkages under force (von Wichert et al., 2003b).

In RPTP α (RPTPA)-knockout cells, as well as talin 1 (*TLN1*)-knockout cells, integrin-ECM adhesion sites eventually form and become strengthened in a manner similar to that observed on a faster timescale under force (Kostic et al., 2007; Priddle et al., 1998). This indicates that other proteins (probably talin 2) can substitute for talin 1 in building integrin-cytoskeleton connections (Giannone et al., 2003; Zhang et al., 2008). Correspondingly, a model of force-accelerated integrin activation has recently been proposed in which integrins that are not anchored to the ECM are more mobile under force and are thus readily recruited to sites that have already been stabilized by anchored integrins (Rose et al., 2007).

Together, these observations point to a common mechanosensing mechanism: force-induced acceleration of biochemical events that are likely to occur, albeit on slower timescales, in a diffusion-controlled fashion in the absence of force. In this scenario, force strengthens adhesions by accelerating reactions and governing the spatial pattern of the cell-ECM interface, whereas relaxation of force inhibits strengthening and accelerates dissociation.

Force, integrin detachment and adhesion remodeling

Although detachment has received less attention, it is an extremely important, tension-dependent step in the integrin mechanical cycle, because cells must release to move and restructure their environment. The lifetime of focal adhesions as distinct entities is in the order of 5-10 minutes (Ren et al., 2000). Within focal adhesions, integrins that are directly linked to the ECM exhibit the slowest dynamics relative to other focal-adhesion proteins, with exchange rates in the order of 1-3 minutes (Ballestrem et al., 2001; Hu et al., 2007). Vinculin, focal adhesion kinase (FAK) and talin have much faster exchange rates than integrins (Hu et al., 2007; von Wichert et al., 2003a), and integrin turnover is preceded by the switch from high to low integrin-binding affinity (Cluzel et al., 2005). What drives this hierarchy in protein exchange rates, and how does that govern integrin-binding affinity under force? More generally, how is adhesion-site remodeling controlled? To address this larger question, which is crucial for understanding tissue homeostasis, we will consider the role of integrin segregation in adhesion remodeling in terms of syndecans, phosphorylation and recycling.

Syndecans

Syndecans, similar to integrins, are a family of transmembrane ECM-adhesion receptors. Whereas integrins bind to peptide motifs in their ECM ligand (e.g. the RGD loop on the tenth fibronectin

type III module), syndecans bind to heparin-binding motifs in their ECM ligand (e.g. the heparin-binding motif on the thirteenth type III fibronectin module). Together, syndecans and integrins have been shown to facilitate the transduction of multiple signaling pathways (for a review, see Morgan et al., 2007). In particular, syndecan 1 has been shown to modulate $\alpha V\beta 3$ -integrin-binding affinity (Beauvais et al., 2004), whereas syndecan 4 interacts, albeit indirectly (Zimmermann et al., 2005), with $\alpha 5\beta 1$ integrin in focal adhesions. The engagement of the ECM with syndecan 4 is linked with activation of FAK and Src, two non-receptor kinases that have important roles in the weakening of integrin-cytoskeletal linkages under force (Felsenfeld et al., 1999; Galbraith et al., 2002; von Wichert et al., 2003a). In vivo, FAK inhibition occurs in a rigidity-dependent fashion as its expression is required for durotaxis on collagen (Wang et al., 2001). It also occurs in a temporal fashion, as the prolonged association of FAK within focal adhesions is linked with increased FAK activity and increased focal-adhesion disassembly (Giannone et al., 2004). When activated, FAK and Src have been shown to form a complex with one another, which extends the lifetime of their active states (Lietha et al., 2007). Whereas Src has been shown to colocalize with $\alpha V\beta 3$ - but not $\alpha 5\beta 1$ -integrins (Felsenfeld et al., 1999), evidence of regulation of syndecan 4 by FAK is robust (reviewed by Morgan et al., 2007). Together, these findings suggest that, similar to focal-adhesion strengthening through talin crosslinking across linearly arranged $\alpha V\beta 3$ and $\alpha 5\beta 1$ integrins, focal-adhesion turnover can be similarly regulated by the linear arrangement of $\alpha V\beta 3$ - and $\alpha 5\beta 1$ -integrin, in synergy with syndecan 1 and syndecan 4, and crosslinked inside and outside the cell by the FAK-Src complex and fibronectin module, respectively.

Integrin phosphorylation

The dissociation of talin from $\beta 3$ -integrin tails is probably an important event along the pathway to force-induced focal-adhesion turnover. When the localization of talin to adhesion sites is altered by the injection of antibody (Nuckolls et al., 1992) or sequestration of phosphoinositides (Martel et al., 2001), there is no simultaneous disruption of mature adhesion sites. However, antibody injection disrupts newly formed adhesion sites or prevents their formation, indicating the crucial role of talin in early rather than mature adhesion sites. As talin cannot bind to phosphorylated $\beta 3$ -integrin tails, one possible mechanism of force-accelerated integrin turnover comes from the tension-induced increase in kinase and phosphatase activity that has been shown to occur in the vicinity of the $\beta 3$ -integrin tail under force (Giannone and Sheetz, 2006; Tamada et al., 2004).

Talin activates $\beta 3$ integrins by influencing specific interactions in the membrane-proximal region of the integrin tail (Wegener et al., 2007). In contrast to talin, binding of the signaling adaptor Shc (Cowan et al., 2000) requires phosphorylation of the $\beta 3$ -integrin tails. Interestingly, Shc requires phosphorylation of only the membrane-distal and not the membrane-proximal tyrosine of the $\beta 3$ tail (Cowan et al., 2000), thus leaving the membrane-proximal region free to renew its association with the α -subunit, which could switch the integrin from the high-affinity to the low-affinity state. Perhaps the binding of Shc to the $\beta 3$ -integrin tail promotes the switch from high to low integrin-binding affinity by promoting renewed interactions between integrin transmembrane domains and membrane-proximal segments of the cytoplasmic tails. In support of this hypothesis, Shc association with $\alpha V\beta 3$ integrins has been shown to be induced under shear-fluid flow in endothelial cells, in which dynamic remodeling of the adhesion plaques to allow strategic positioning for sustaining force requires

constant integrin association with and dissociation from the ECM (Chen et al., 1999).

It has been previously proposed that tyrosine phosphorylation functions as a 'molecular switch' for the binding interactions of $\beta 3$ -integrin tails following adhesion, activation and clustering (Calderwood et al., 2003). Although a recent *in vivo* study has revealed that cytoplasmic regulation of $\beta 1$ -integrin function is phosphorylation independent – with the hydrophobic interactions that the tyrosines support having the key role instead (Chen et al., 2006) – phosphorylation of the cytoplasmic tail of $\beta 3$ integrins is crucial for platelet aggregation (Blystone et al., 1997; Jenkins et al., 1998; Law et al., 1999). Knock-in mice, in which each of the tyrosines (Y) in the two NPxY motifs on the $\beta 3$ -integrin tail was replaced by phenylalanine, displayed platelet clotting deficiencies (Law et al., 1999). These mutational studies showed that disruption of $\beta 3$ -tail phosphorylation does not disrupt initial aggregation rates but, rather, disrupts the ability of $\beta 3$ integrins to maintain aggregation. Interestingly, fluorescence recovery after photobleaching (FRAP) experiments have shown that exchange rates are faster in high-density $\beta 3$ -integrin clusters and slower in low-density $\beta 3$ -integrin clusters (Ballestrem et al., 2001). Intracellular tension that is induced by RhoA or blocked by the protein-kinase inhibitor staurosporine correlates with the formation and maintenance of high-density or low-density $\alpha V\beta 3$ -integrin focal adhesions, respectively (Ballestrem et al., 2001). Increased integrin density under force has thus been proposed to lead to lower-affinity integrin-ECM binding (Ballestrem et al., 2001). Together, these findings suggest that clustering of $\alpha V\beta 3$ integrins, and the subsequent competition for $\beta 3$ -integrin tails by force-increased kinase activity, may be a key event in focal-adhesion turnover.

Recycling of integrins to the leading edge

Although it is possible for the integrins to diffuse back to the leading edges of active cells, observations of the diffusion of unliganded integrins have revealed that they often undergo active movements towards the leading edge (Schmidt et al., 1993). Similar to the sites of integrin release from adhesions, the major sites of membrane exocytosis are located at the ends of microtubules, often 2–4 μm back from the leading edge, and it takes integrins nearly 10 minutes to diffuse to the leading edge at the observed diffusion rates ($D = (0.1 \text{ micrometer})^2/\text{second}$) (Schmidt et al., 1995; Schmidt et al., 1993). Recently, recycling of $\alpha V\beta 3$ integrins to the leading edge has been proposed to resensitize the integrins to ligand occupation, by acting to return them to the membrane that is competent to bind and promote migration (White et al., 2007). Thus, recycling of integrins to the leading edge through the active transport mechanism is likely to be an important component of adhesion remodeling during rapid cell motility.

Conclusions and perspectives

The integrin mechanical cycle (supplementary material Fig. S1) is crucial for cellular function and depends upon the biophysical and biochemical changes in integrin structure and post-translational modifications of both integrins and associated proteins. In the initial binding to matrix, both intracellular and extracellular factors can alter the level of integrin binding, in terms of affinity (e.g. talin-head activation or matrix binding) as well as avidity (e.g. edge localization and lateral interactions with externally activated proteins such as syndecans). Stabilization of integrin-matrix binding and the formation of an adhesion complex by lateral recruitment of more integrins is probably catalyzed at early times by force-dependent alterations in talin and, possibly, in the integrin itself. As the lifetimes of components

in the adhesion complexes are much shorter than the lifetimes of focal adhesions themselves, there must be a self-renewal process that supports tension while enabling turnover. Recruitment of new actin filaments and actin-binding components in overlapping arrays through radial or simply lateral aggregation could facilitate this.

The local and directional growth of focal adhesions under force can be explained in terms of a network of mechanotransduction pathways. Force probably accelerates integrin activation, both by extracellular and intracellular rearrangements, induces protein recruitment through protein stretching and accelerates integrin clustering, leading to deactivation. Dispersal of adhesion contacts is favored by the loss of tension. The recycling of integrins to the leading edge of moving cells allows the integrin cycle to begin again. Force clearly alters the biochemical steps in the integrin cycle, which enables cellular signaling pathways to affect the strength of the adhesions indirectly by altering cell-force generation, or directly by altering adhesion protein dynamics.

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