

Molecular Bases of Cell–Cell Junctions Stability and Dynamics

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Epithelial cell–cell junctions are formed by apical adherens junctions (AJs), which are composed of cadherin adhesion molecules interacting in a dynamic way with the cortical actin cytoskeleton. Regulation of cell–cell junction stability and dynamics is crucial to maintain tissue integrity and allow tissue remodeling throughout development. Actin filament turnover and organization are tightly controlled together with myosin-II activity to produce mechanical forces that drive the assembly, maintenance, and remodeling of AJs. In this review, we will discuss these three distinct stages in the lifespan of cell–cell junctions, using several developmental contexts, which illustrate how mechanical forces are generated and transmitted at junctions, and how they impact on the integrity and the remodeling of cell–cell junctions.

Cell–cell junction formation and remodeling occur repeatedly throughout development. Epithelial cells are linked by apical adherens junctions (AJs) that rely on the cadherin–catenin–actin module. Cadherins, of which epithelial E-cadherin (E-cad) is the most studied, are Ca^{2+} -dependent transmembrane adhesion proteins forming homophilic and heterophilic bonds in *trans* between adjacent cells. Cadherins and the actin cytoskeleton are mutually interdependent (Jaffe et al. 1990; Matsuzaki et al. 1990; Hirano et al. 1992; Oyama et al. 1994; Angres et al. 1996; Orsulic and Peifer 1996; Adams et al. 1998; Zhang et al. 2005; Pilot et al. 2006). This has long been attributed to direct physical interaction of E-cad with β -catenin (β -cat) and of α -catenin

(α -cat) with actin filaments (for reviews, see Gumbiner 2005; Leckband and Prakasam 2006; Pokutta and Weis 2007). Recently, biochemical and protein dynamics analyses have shown that such a link may not exist and that instead, a constant shuttling of α -cat between cadherin/ β -cat complexes and actin may be key to explain the dynamic aspect of cell–cell adhesion (Drees et al. 2005; Yamada et al. 2005). Regardless of the exact nature of this link, several studies show that AJs are indeed physically attached to actin and that cadherins transmit cortical forces exerted by junctional acto-myosin networks (Costa et al. 1998; Sako et al. 1998; Pettitt et al. 2003; Dawes-Hoang et al. 2005; Cavey et al. 2008; Martin et al. 2008; Rauzi et al. 2008). In addition, physical

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association depends in part on α -cat (Cavey et al. 2008) and additional intermediates have been proposed to represent alternative missing links (Abe and Takeichi 2008) (reviewed in Gates and Peifer 2005; Weis and Nelson 2006). Although further work is needed to address the molecular nature of cadherin/actin dynamic interactions, association with actin is crucial all throughout the lifespan of AJs. In this article, we will review our current understanding of the molecular mechanisms at work during three different developmental stages of AJs biology: assembly, stabilization, and remodeling, with special emphasis on the mechanical forces controlling AJs integrity and development.

CELL JUNCTIONS FORMATION

Cell junctions form in two contexts during development. (1) Migrating cells undergo mesenchymal-epithelial transitions (MET) during which they establish membrane contacts with neighbors and initiate assembly of AJs at these sites. Subsequently, cell–cell contacts expand and newly formed AJs serve as landmarks for establishing tissue polarity. (2) In primary embryonic epithelia, cell junctions are formed in a subregion of pre-existing cell contacts, which is defined by upstream polarity cues.

Cell–cell junction formation during MET has been extensively studied in cell cultures: (1) formation of junctions after membrane contacts or after Ca^{2+} switch (activating cadherin adhesive function by raising the extracellular Ca^{2+} concentration); (2) adhesion of cells on cadherin-coated substrata. These studies have provided considerable insights into in vivo epitheliogenesis and epithelial sheet sealing processes, which occur during embryogenesis and wound healing. In this section, we first review how cell junctions assemble in cell cultures, focusing on the role of forces generated by actin polymerization and acto-myosin tension. We then examine how these forces are used during epithelial sheet sealing in whole organisms. Finally, we summarize how cell junctions form in a primary embryonic epithelium, i.e., without MET.

Mesenchymal-Epithelial Transitions (MET): Cell Culture Studies

Cell–cell junction formation during MET suggests the following steps:

1. Membrane protrusions explore the environment to generate initial cell contacts. Cadherin molecules diffusing in the plasma membrane engage in homophilic interactions and form clusters.
2. Homophilic ligation of cadherins triggers actin cytoskeleton rearrangement by directly controlling the recruitment and activity of several actin regulators.
3. Actin reorganization drives contact expansion (an increase of the surface of contact between two cells), and is also linked to the stabilization of adhesive interfaces.

Formation of Initial Cell–Cell Contacts

The first step is an opportunistic event resulting from the exploratory behavior of cells extending actin-based protrusions (lamellipodia and membrane ruffles) (Fig. 1A) (reviewed in Vasioukhin and Fuchs 2001). Immunofluorescence and the use of GFP-fusions show that bright cadherin puncta rapidly form where protrusions touch (Fig. 1B). These puncta are thought to represent clusters of homophilic cadherin dimers, a view supported by the fact that they contain cadherin molecules from both contacting cells (Kametani and Takeichi 2007). However, cadherin clustering mechanisms are still not fully understood and remain debated. They may include lateral interactions in the cadherin extracellular domain and intracellular interactions including interactions with actin (reviewed in Leckband and Prakasam 2006; Pokutta and Weis 2007). Upon contact formation, cadherin molecules previously diffusing “freely” in the membrane become immobilized (Adams et al. 1998), presumably by anchoring to actin (Sako et al. 1998; Iino et al. 2001; Lambert et al. 2002). Within a few tens of minutes, cadherin puncta grow, suggesting a continuous addition of new molecules

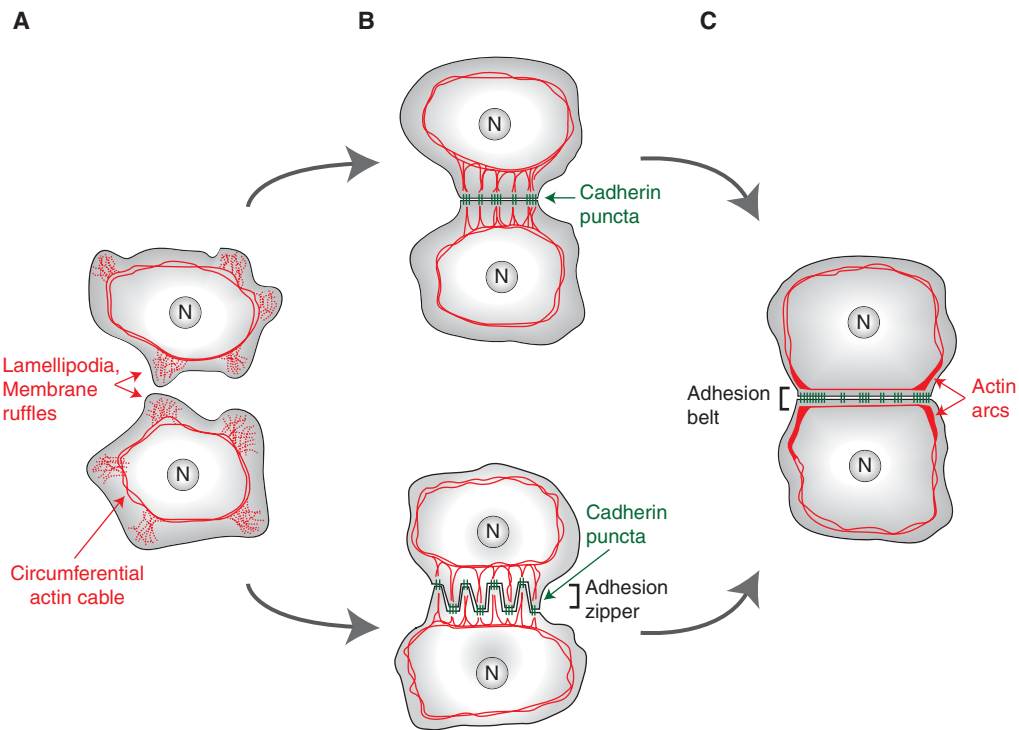


Figure 1. Actin reorganization during cell–cell junction formation in cell culture. Actin is shown in red, cadherin in green. (A) Before cell–cell contact, epithelial cells extend protrusions (lamellipodia and membrane ruffles). (B) Cadherin puncta form at the tips of these projections and are connected to the circumferential actin cable via radial actin bundles. (B, bottom) Contacting keratinocytes adopt an intermediate configuration known as the “adhesion zipper” as a consequence of myosin-mediated tension pulling inward on cadherin puncta. (C) As cell–cell contacts expand and mature, actin arcs focus on the edges of the contact. Actin remodeling along the contact results in formation of the adhesion belt.

(Yonemura et al. 1995; Adams et al. 1996; Adams et al. 1998).

In the *Drosophila* embryo, a similar punctate organization of cadherins has been reported (Muller and Wieschaus 1996; Harris and Peifer 2004). Moreover, puncta were recently shown to represent *bona fide* sites of immobilized E-cad clusters with slower dynamics than outside puncta (Cavey et al. 2008), suggesting that they are equivalent to the spot adherens junctions (SAJs) structures observed at the electron microscopy level (Tepass and Hartenstein 1994; Oda et al. 1998). Therefore the puncta observed in cell cultures and in several epithelia *in vivo* likely represent the same structures, namely sites of homophilic cadherin dimers enrichment.

Actin Reorganization

Before cell contacts, actin forms concentric ring(s) (the “circumferential actin cable/ring”) and a dense meshwork between the ring and the plasma membrane (Fig. 1A) (Yonemura et al. 1995; Adams et al. 1996; Gloushankova et al. 1997; Adams et al. 1998; Krendel and Bonder 1999; Ehrlich et al. 2002; Vaezi et al. 2002; Ivanov et al. 2005a). As contacts form, cadherin puncta are connected to the actin ring via radial actin bundles (Fig. 1B). Subsequently, actin bundles are replaced by finer ones under the region of contact (the “perijunctional actin belt/adhesion belt”) resembling actin organization in epithelia *in vivo* (Hirokawa et al. 1983), and thick bundles (“actin arcs”) focus on the

contact edges (Fig. 1C). This cytoskeletal reorganization is triggered and controlled by cadherins whose homophilic ligation can directly recruit and activate actin regulators including Rac1, Cdc42 (Nakagawa et al. 2001; Noren et al. 2001; Kovacs et al. 2002a), Abl kinase (Zandy et al. 2007), Arp2/3 (Kovacs et al. 2002b; Verma et al. 2004), Cortactin (Helwani et al. 2004), N-WASP (Ivanov et al. 2005a), Formin1 (Kobiela et al. 2004), and Ena/VASP (Vasioukhin et al. 2000) (for reviews, see Bershadsky 2004; Braga and Yap 2005). Cell–cell contact formation is dependent on actin polymerization (Braga et al. 1997; Adams et al. 1998; Vasioukhin et al. 2000; Ivanov et al. 2005a; Zhang et al. 2005; Yamada and Nelson 2007) and adhesion strength (measured as the force required to detach cell doublets) increases in an F-actin- (Angres et al. 1996), Cdc42-, and Rac1-dependent manner (Chu et al. 2004). Cytoskeletal reorganization likely serves two related purposes examined below: expansion and stabilization of the adhesive interface.

Cell–Cell Contact Expansion

Two forces are coordinated to expand cell contacts: (1) Actin polymerization produces membrane protrusions to generate new sites of contact, and (2) acto-myosin tension focusing on contact edges generates a pulling force to facilitate contact expansion.

(1) Lamellipodia and membrane ruffles initiating cell–cell contacts (Gloushankova et al. 1997; Krendel and Bonder 1999) are generated by branched actin networks (Vaezi et al. 2002; Bershadsky 2004). New sites of contact are then generated in adjacent regions. In keratinocytes, the adhesive interface develops in an “adhesion zipper” structure, because of myosin-mediated tension pulling inward on cadherin puncta (Fig. 1B, *bottom*). Actin polymerization from the tip of radial actin bundles could provide the force necessary to resolve the two rows of cadherin puncta into a single belt of mature junctions, but the underlying mechanism is not understood (Vasioukhin et al. 2000; Kobiela et al. 2004). In other

cell types, lamellipodial activity is initially distributed evenly around the cell periphery but becomes restricted to the region of contact and subsequently propagates to adjacent regions in waves generating new sites of membrane apposition (Ehrlich et al. 2002) (Fig. 2A). Supporting this idea, factors promoting branched actin polymerization are enriched at the leading edge of cells spreading on cadherin-coated substrata (Kovacs et al. 2002a; Helwani et al. 2004) and Rac1 is specifically activated in regions of contact expansion (Yamada and Nelson 2007). Moreover, interfering with actin branching impairs cell spreading or contact expansion (Ehrlich et al. 2002; Kovacs et al. 2002a; Helwani et al. 2004; Verma et al. 2004; Ivanov et al. 2005a; Zandy et al. 2007). Conversely, Rac1 constitutive activation increases cell contact expansion rate and adhesion strength (Ehrlich et al. 2002). Together, these studies show that actin polymerization in lamellipodia is specifically concentrated in regions adjacent to the site of initial contact and provides a pushing force required for membrane apposition. These fluctuating contacts are then ligated and rectified by cadherin homophilic dimers.

(2) Myo-II-mediated tension is also important for contact expansion. Myo-II localizes to peripheral actin bundles in epithelial cells (Krendel et al. 1999; Krendel and Bonder 1999; Bertet et al. 2004; Zallen and Wieschaus 2004; Shewan et al. 2005; Zhang et al. 2005; Yamada and Nelson 2007) and is required to bundle actin filaments (Vaezi et al. 2002; Ivanov et al. 2005a; Shewan et al. 2005; Zhang et al. 2005). Cell adhesion induces Myo-II activation (Ivanov et al. 2005a), which can be directly triggered by cadherin homophilic ligation (Shewan et al. 2005). In keratinocytes, Myo-II appears to act negatively on contact expansion by generating the adhesion zipper structure (Vaezi et al. 2002). However, in other cell types Myo-II acts positively on contact expansion (Yamada and Nelson 2007). How can Myo-II activity participate in contact expansion?

As contacts expand, cadherin puncta (Adams et al. 1998) and ConA-coated beads

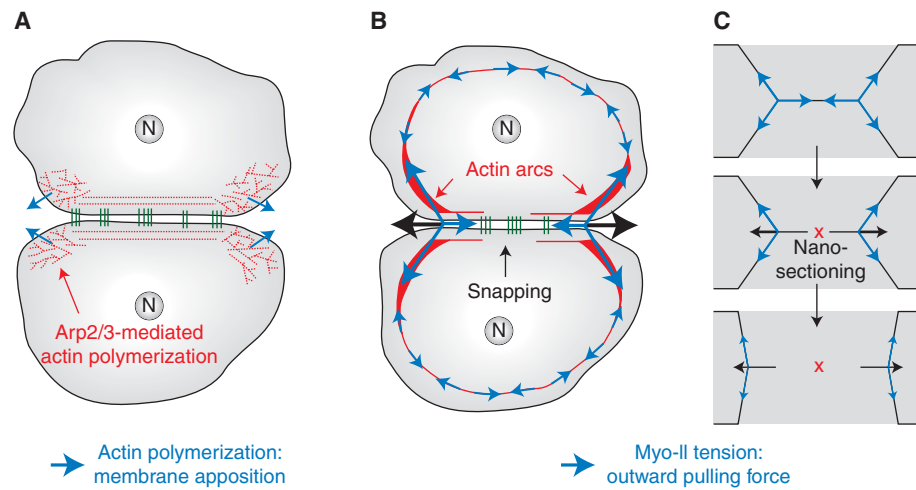


Figure 2. Forces driving cell–cell contact expansion in cell culture. (A) Polymerization of branched actin networks at the edges of a contact generates a pushing force (blue arrows) necessary to create new sites of membrane apposition, which are then ligated by homophilic cadherin dimers. (B,C) Analogy for the role of myosin-II-mediated tension (blue arrows) during contact expansion in cell culture (B) and after laser nano-sectioning in an epithelium (C). In both cases, the sum of tension forces applied at the contact edges is initially null, but once actin bundles are sectioned by snapping along the cell–cell contact (B) or laser nano-sectioning (C), these forces are not balanced anymore. This produces a net outward pulling force (black arrows), which drives contact expansion.

that mark plasma membrane proteins (Gloshankova et al. 1997) move toward the contact edges (i.e., tangentially) at similar speeds, revealing a flow of cortical material toward the edges. The cortical forces responsible for this flow likely result from acto-myosin contractility. Activated Myo-II and activated Rho are enriched on the actin arcs, which contract at the edges of cell contacts (Krendel and Bonder 1999; Yamada and Nelson 2007) (Fig. 2B). In addition, during contact expansion, actin bundles along the contact zone break and retract toward the edges of the contact (i.e., tangentially), in a process termed “actin bundle snapping” (Krendel and Bonder 1999) (Fig. 2B). This suggests a model whereby the local unbalance of cortical acto-myosin forces at contact edges drives expansion. Actin bundles along the contact have to resist the outward pulling forces generated on the actin arcs. However, actin bundle snapping along the contact alleviates such a resistance, resulting in a net outward force pulling on the contact edges that drives contact expansion (Fig. 2B). Snapping along cell contacts could be a

consequence of increased tension generated on actin arcs, combined with actin remodeling triggered by cell–cell adhesion. Moreover, retraction of these bundles after snapping could drag cadherin puncta toward the edges (Adams et al. 1998). Such a mechanism would be analogous to laser nano-sectioning experiments performed in live epithelial cells of *Drosophila* embryos (Cavey et al. 2008; Rauzi et al. 2008). Indeed, nano-sectioning of a tensile acto-myosin cortical network along a cell–cell contact causes the actin network to retract, resulting in a net expansion of the contact (Fig. 2C). Actin retraction induces E-cad puncta redistribution away from the region of sectioning by tethering E-cad puncta to the contractile acto-myosin network (Cavey et al. 2008; Rauzi et al. 2008).

Junction Formation by MET in Embryos

Interfering with AJs components in embryonic tissues results in tissue collapse at varying developmental stages (Larue et al. 1994; Riethmacher et al. 1995; Tepass et al.

1996; Uemura et al. 1996; Torres et al. 1997; Carmeliet et al. 1999; Vasioukhin et al. 2001; De Vries et al. 2004). One interesting phenomenon is when cell–cell contacts are increased during compaction of late 8-cell mouse embryos. Interfering with cell adhesion using anti-E-cad antibodies and Ca^{2+} depletion both block compaction (Kemler et al. 1977; Hyafil et al. 1980; Pratt et al. 1982; Shirayoshi et al. 1983; Johnson et al. 1986). Compaction may be driven by similar forces as contact expansion in cell culture. Compaction requires actin polymerization (Pratt et al. 1982; Fleming et al. 1986; Clayton et al. 1999) and involves a redistribution of E-cad to the baso-lateral domain where cell contacts expand (Vestweber et al. 1987). Interestingly, numerous membrane protrusions (microvilli) form bridges between cells along regions of contacts (Calarco and Epstein 1973; Fleming et al. 1986), suggesting a role analogous to that of lamellipodia in cell cultures. In later stages, microvilli are excluded from regions of cell contacts (Fleming et al. 1986), as membrane protrusive activity may not be compatible with stabilization of the interface (see below). This may be controlled

in part by the ERM (Ezrin-Radixin-Moesin) protein Ezrin, which organizes actin networks to form microvilli and has to be excluded from regions of cell–cell contacts for compaction to be completed (Dard et al. 2001; Dard et al. 2004). A role for acto-myosin tension in driving compaction has not been directly investigated yet, but Rho inhibition (using C3-transferase) disrupts actin and E-cad organization, blocking compaction (Clayton et al. 1999).

Cell–cell contact establishment also occurs during the sealing of epithelial sheets during embryogenesis and wound healing. At the end of embryogenesis, morphogenetic rearrangements leave holes in the dorsal epidermis of *Drosophila* embryos and in the ventral hypodermis of *C. elegans* embryos. Similar holes are created when embryonic or adult tissues are wounded. In all cases, migrating epithelial sheets cover and eventually seal these holes (Fig. 3) (reviewed in Jacinto et al. 2001; Martin and Parkhurst 2004). As for the establishment of cell–cell contacts in cell culture, actin protrusions and acto-myosin cables are both involved in epithelial sheet sealing. An acto-myosin cable

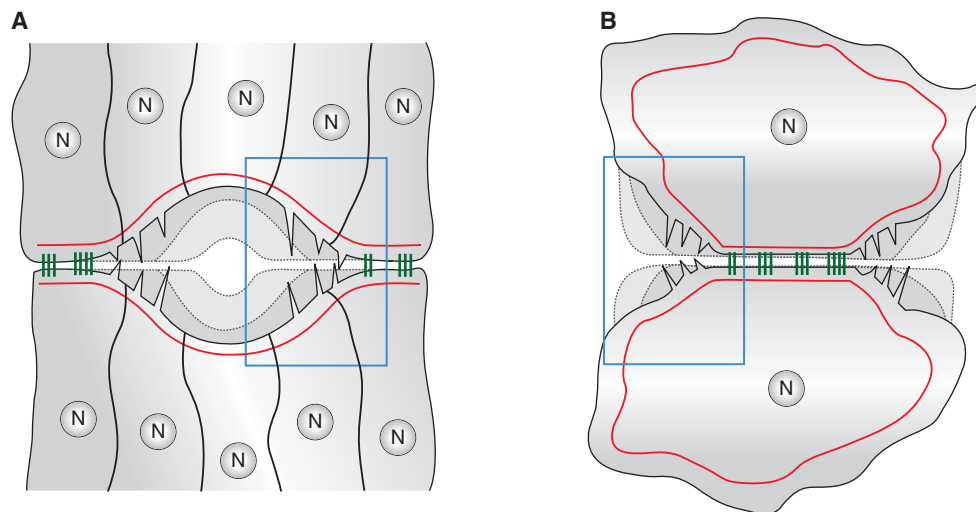


Figure 3. Parallel between epithelial sheet sealing and contact expansion in cell culture. (A) Epithelial sheet sealing at the end of embryogenesis and during wound healing and (B) cell–cell contact expansion in cell culture. More advanced stages are shown by dashed lines. (A) A contracting acto-myosin cable (red) closes the hole and the final sealing step is facilitated by actin-based protrusions. Note the similarity of structures involved in the two systems (blue rectangles).

assembled at the periphery of the hole/wound provides contractile force to progressively close the hole in flies (Harden et al. 1999; Magie et al. 1999; Kiehart et al. 2000; Bloor and Kiehart 2002; Jacinto et al. 2002) during wound healing (Wood et al. 2002)(reviewed in Jacinto et al. 2001; Redd et al. 2004) and possibly in worms as well (Williams-Masson et al. 1997; Raich et al. 1999). Filopodia establish connections over the holes and are required during the final phase to seal the two sheets together. They may exert pulling forces to help sealing (Williams-Masson et al. 1997; Raich et al. 1999; Jacinto et al. 2000; Bloor and Kiehart 2002; Jacinto et al. 2002; Soto et al. 2002; Wood et al. 2002; Gates et al. 2007; Sheffield et al. 2007; Millard and Martin 2008). Electron microscopy studies have revealed interdigitated filopodia during sealing (Redd et al. 2004) harboring AJs puncta at their tips (Raich et al. 1999; Vaezi et al. 2002). This structure appears analogous to the adhesion zipper observed in keratinocytes (Vasioukhin et al. 2000).

Cell–Cell Junctions Formation by Non-MET Processes

In the examples mentioned so far, AJs formation defines a spatial cue for organizing the cytoskeleton and recruiting apical–basal polarity complexes. In turn, these complexes stabilize and maintain AJs, possibly via regulation of the junctional actin cytoskeleton (reviewed in Knust and Bossinger 2002; Nelson 2003; Ebnet 2008). However, in the *Drosophila* embryonic primary epithelium, AJs formation occurs in a small region of the surface of contact between cells, which is defined by already-present apical–basal polarity cues. The polarity protein Par3/Bazooka (Baz) is the first component recruited to the apical region, in a microtubule- and Dynein-dependent process (Harris and Peifer 2004; Harris and Peifer 2005). Par3/Baz then initiates AJs assembly and recruits further polarity components, which maintain AJs in later stages (Muller and Wieschaus 1996; Bilder et al. 2003; Tanentzapf and Tepass

2003; Harris and Peifer 2004; Hutterer et al. 2004; Harris and Peifer 2005). Par3/Baz oligomers could serve as structural adaptors for AJs integrity (Benton and St Johnston 2003) or could link AJs to actin via the nectin/afadin system (Wei et al. 2005). Polarity complexes are also probably implicated in regulating actin polymerization via the small GTPases. In mammals, Par3 serves as a platform connecting Rho signaling to Rac1 regulation to control front/rear polarity in migrating cells. Rac1 is activated by its GEF Tiam1/2, which is in a complex with Par3/aPKC/Cdc42 (Nishimura et al. 2005). Phosphorylation of Par3 by the RhoA-ROCK pathway separates Rac1 from Tiam1/2, resulting in Rac1 inactivation (Nakayama et al. 2008). In flies, Par3 has also been implicated in organizing the cortical actin cytoskeleton by acting on the recruitment of Moesin, an ERM protein (Pilot et al. 2006) and in the control of phosphoinositides levels (von Stein et al. 2005), which can impact on actin dynamics and organization in many ways (for reviews, see Zheng 2001; Yin and Janney 2003).

MAINTENANCE OF CELL–CELL JUNCTIONS

Once cell contacts have been established, actin polymerization and acto-myosin tension are required for the stabilization and maturation of adhesive interfaces. Actin network organization and Myo-II-mediated tension control the organization of adhesion molecules at the cell surface and the dynamics of the plasma membrane, which are both crucial for maintaining stable cell interfaces and tissue architecture.

Immobilization of E-cadherin Clusters

The organization of cadherins at cell interfaces depends on the integrity of the actin cytoskeleton (Pilot et al. 2006). In *Drosophila* embryos, apical–basal polarity cues (Par3/Baz and the phosphoinositide derivative PIP2) recruit the synaptotagmin-like protein Bitesize, which in turn recruits Moesin to the apical region.

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These factors organize the cortical actin network and ensure its integrity, which is crucial for a homogeneous distribution of adhesion foci at cell interfaces (Pilot et al. 2006). In fact, cortical actin is composed of two intermixed populations of filaments, which control E-cad distribution at two levels (Cavey et al. 2008). (1) Locally, actin filaments with a low turnover control the stability of E-cad molecules within puncta. (2) A peripheral contractile network of acto-myosin controls the mobility (displacement) of these puncta along cell–cell contacts. Myo-II-mediated tension is crucial for effectively tethering E-cad puncta in the plasma membrane and thus for controlling their spatial distribution along cell–cell contacts. This suggests that stabilization of adhesion requires the regulated immobilization of homophilic E-cad clusters by a tensile cortical actin network, independently of cluster stabilization per se (Cavey et al. 2008). These results shed light on previous reports, which implicated Myo-II in the spatial organization of cadherin clusters in various cell types (Gloushankova et al. 1998; Krendel et al. 1999; Vaezi et al. 2002; Conti et al. 2004; Shewan et al. 2005; Zhang et al. 2005).

E-cadherin Clusters Stability

Actin depolymerization studies suggest that actin turnover is reduced in mature cell junctions compared to younger ones (Adams et al. 1998; Braga et al. 1999; Ivanov et al. 2005a). Actin turnover is also specifically lower at cadherin puncta compared to neighboring regions in *Drosophila* epithelial cells (Wood et al. 2002; Cavey et al. 2008). In mammalian cells, cadherin stability and actin stability may be directly coupled by Eplin, a protein recruited by E-cad/ β -cat/ α -cat complexes and which is required for stabilizing actin filaments associated with adhesion complexes (Abe and Takeichi 2008). However, recycling of actin filaments associated with adhesion complexes is likely to occur because small GTPases are required to maintain adhesion. Alternatively, GTPase activity may reflect the

need to remodel junctions in dynamic epithelia (reviewed in Braga and Yap 2005; Kooistra et al. 2007; Yamazaki et al. 2007).

Role of Junctional Actin Architecture

Adhesion strengthening seems to involve regulation of actin cytoskeleton organization: Branched networks associated with lamellipodial protrusions are replaced by parallel contractile bundles. This transition probably takes place very shortly after the initial clustering of cadherins and appears to be controlled by AJs components. Factors promoting branched actin polymerization (Rac1, Arp2/3, and cortactin) are relatively depleted from “older” regions of cell contacts (Helwani et al. 2004; Verma et al. 2004; Yamada and Nelson 2007). α -cat was proposed to directly control this transition by repressing Arp2/3 activity (Drees et al. 2005). As AJs mature, progressive enrichment of α -cat would result in local inhibition of Arp2/3 (Perez-Moreno and Fuchs 2006; Pokutta and Weis 2007). In addition, factors promoting unbranched F-actin elongation can be recruited to AJs, such as Formin1 and Ena/VASP in keratinocytes (Vasioukhin et al. 2000; Kobiela et al. 2004). Another formin, Diaphanous (Dia), is recruited to cell contacts and required for junction maintenance (Sahai and Marshall 2002; Carramusa et al. 2007; Homem and Peifer 2008). Dia may be directly recruited at junctions by α -cat and p120, which recruit its activator Rho1 (Magie et al. 2002). Bundling factors at nascent AJs may also contribute to the transition of actin organization (for reviews, see Adams 2004; Broderick and Winder 2005; Mege et al. 2006). In this context, direct observation of actin network architecture at AJs is an important, albeit challenging, avenue.

Tension and Rigidity of the Interface

Linear arrays of unbranched actin filaments favor Myo-II tension, which can stabilize cell–cell junctions in two ways.

(1) The protrusive activity of cell membranes is thought to destabilize adhesive



interfaces (Gloushankova et al. 1997; Sahai and Marshall 2002; Zhang et al. 2005). Myo-II-mediated tension inhibits the formation of protrusions by its ability to align actin filaments parallel to the cell membrane (Gloushankova et al. 1997). In cell cultures, protrusive activity decreases after cell–cell contact formation (Gloushankova et al. 1997; Ehrlich et al. 2002) and increases upon Myo-II inhibition in keratinocytes (Vaezi et al. 2002; Zhang et al. 2005). Similar results were observed in *Drosophila* (Bloor and Kiehart 2002; Jacinto et al. 2002). Interfering with Dia function (polymerization of unbranched actin filaments) increases the membrane protrusive activity and creates gaps in cell contacts (Sahai and Marshall 2002). The formation of a continuous belt of adhesion is compromised upon Myo-II inhibition in keratinocytes and MCF-7 cells but whether this is a direct consequence of increased protrusions is unclear (Shewan et al. 2005; Zhang et al. 2005).

(2) Myo-II-mediated tension may also affect the distribution of cadherin molecules at junctions. For instance, applying tension on the cell membrane is sufficient to drive the clustering of cadherins independently of actin. The current model proposes that membrane tension along a cell contact brings the two cell membranes in close proximity, thereby favoring new homophilic cadherin interactions to form and thus trapping these molecules in the region under tension (Delanoe-Ayari et al. 2004).

DYNAMICS AND REMODELING OF CELL–CELL JUNCTIONS

Throughout embryonic development and in adults, morphogenetic processes, which shape tissues and organisms, require constant remodeling of cell junctions. What underlies the dynamics of cell–cell junctions? They first require a constant turnover of AJs components at the cell surface. Second, they result from the regulated balance of two forces at the cell surface: adhesion and cortical tension.

Adhesion tends to increase the surface of contacts with neighbors, whereas cortical tension tends to decrease it (for a more in-depth discussion, see Lecuit and Lenne 2007).

Regulation of Adhesion

Cadherin Endocytosis and Recycling

The turnover of AJs components is achieved by endocytosis and recycling of cadherins to the cell surface (reviewed in D'souza-Schorey 2005; Ivanov et al. 2005b; Yap et al. 2007). Indeed, several morphogenetic processes involving junction remodeling are blocked upon inhibition of endocytosis (Jarrett et al. 2002; Classen et al. 2005; Ulrich et al. 2005; Shaye et al. 2008). Several studies point to a role for actin in mediating cadherin endocytosis (Le et al. 2002; Ivanov et al. 2004) (reviewed in Kaksonen et al. 2006). Myosin may facilitate endocytosis in Ca^{2+} -depleted cells by assembling a contractile acto-myosin ring, which could provide mechanical force for vesicle formation at AJs (Ivanov et al. 2004). On the other hand, association with actin has been proposed to protect E-cad from endocytosis based on experiments suggesting that homophilically engaged E-cad is not endocytosed, whereas free E-cad is (Izumi et al. 2004). However, this issue remains controversial because cadherin homophilic dimers—which are presumably associated to actin—can be dissociated by endocytosis (Trojanovsky et al. 2006). Recently, the Par polarity proteins Cdc42, aPKC, and Par6 (but not Par3/Baz) have been implicated in regulating AJs stability in *Drosophila* by controlling E-cad endocytosis. Cdc42 is thought to couple actin polymerization (via WASp and Arp2/3) with vesicle scission (via Cip4 and Dynamin) to promote E-cad endocytosis and hence turnover (Georgiou et al. 2008; Leibfried et al. 2008; Harris and Tepass 2008).

E-cadherin recycling is equally important for the regulation of AJs dynamics. In *Drosophila* thoracic epithelial cells, members of the exocyst complex (Sec5, Sec6, and Sec15) directly control the recycling of E-cad to AJs

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via interactions with β -cat (Langevin et al. 2005). In the *Drosophila* pupal wing epithelium, irregularly arranged cells remodel their contacts to pack into a highly regular hexagonal array (Classen et al. 2005) (Fig. 4A). This process requires E-cad endocytosis (Rab5), recycling

(Rab11), and exocytosis (Sec5). E-cad exocytosis may be under the control of the planar cell polarity pathway, which may bias E-cad recycling to specific cortical locations to promote regular hexagonal packing (Fig. 4A) (Classen et al. 2005).

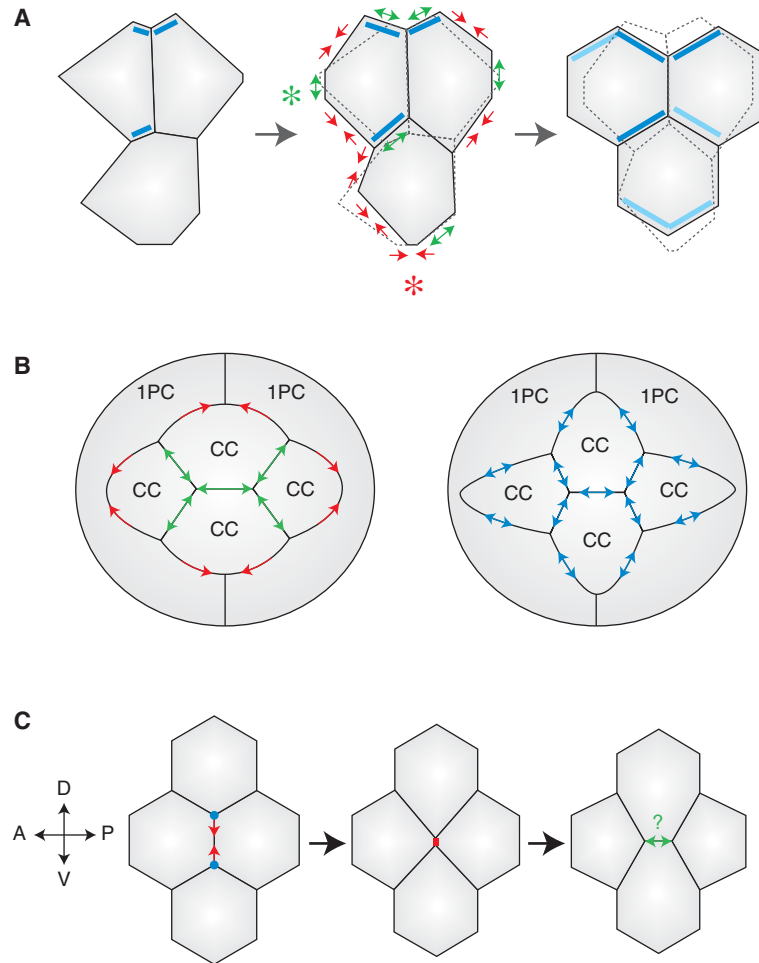


Figure 4. Remodeling cell–cell junctions during morphogenesis. (A) Hexagonal packing in fly wing epithelial cells. Remodeling involves cell–cell contact shrinking (red arrows), expansion (green arrows), loss of some contacts (red asterisk), and creation of new contacts (green asterisk). The previous shape of cells is indicated by dashed lines in the *middle* and *right* panels. Flamingo enrichment (dark blue rectangles) and emergent polarity (light blue rectangles) may spatially bias exocytic delivery of E-cad to promote hexagonal packing. (B) Pattern formation in fly retina. (*Left* panel) Wild type ommatidial cluster with four cone cells (CC) surrounded by two primary pigment cells (1PC). Strong adhesion (N-cad + E-cad) increases inter-CC contacts (green arrows) at the expense of weaker adhesion (E-cad only) with 1PCs (red arrows). (*Right* panel) Uniform adhesion in a mutant ommatidium (blue arrows) distorts the cell pattern. (C) Cell intercalation during germ band elongation in fly embryos. The T1 transition involves shrinking of junctions between A/P neighbors (vertical junctions) and creation of new junctions between D/V neighbors (horizontal junctions). Shrinking is triggered by increased acto-myosin tension along A/P junctions (red arrows; vertices are indicated by blue dots). Expansion of the new D/V interface could rely on adhesion (green arrows), as in cell culture systems.

Generation of Tissue Patterns by Differential Adhesion

The concept of differential adhesion states that differences in adhesion strength between different cell types induce cell sorting, that is, the segregation of groups of cells based on their relative affinity for each other (Steinberg 1963). The strength of interaction is hypothesized to be different between homophilic and heterophilic cadherin dimers. However, cell sorting can happen with similar adhesion strengths (Niessen and Gumbiner 2002; Prakasam et al. 2006) and likely depends on additional parameters including the kinetics of the interactions and the actin cytoskeleton (discussed in Leckband and Prakasam 2006; Lecuit and Lenne 2007). Regardless of the exact mechanisms underlying differential adhesion, the idea that it can drive cell sorting has been largely confirmed in vivo (Godt and Tepass 1998; Gonzalez-Reyes and St Johnston 1998). This concept can be extended to the generation of cell patterns, which do not involve the complete separation of different cell types but subtle cell shape changes. One striking example comes from the morphogenesis of the *Drosophila* ommatidia in the developing retina (Fig. 4B). Ommatidia are composed of four cone cells (CCs) located in the center of a cluster and expressing both N-cad and E-cad, surrounded by two primary pigment cells (1PCs) expressing only E-cad. Genetic analyses have shown that increased adhesion strength in CCs because of N-cad expression maximizes inter-CCs contacts at the expense of CC-1PCs contacts and is responsible for establishing the specific geometry of CCs (Hayashi and Carthew 2004). Another aspect of ommatidial geometry is dictated by differential adhesion, relying on the immunoglobulin (Ig)-domain cell adhesion molecules Hibris and Roughest (Bao and Cagan 2005).

An increase in contact surface is hypothesized to result from increased adhesion strength along the interface. Adhesion strength depends on the surface levels of cadherins (Angres et al. 1996; Duguay et al. 2003; Chu et al. 2004), on the inherent properties

(strength, kinetics) of homophilic versus heterophilic interactions (reviewed in Prakasam et al. 2006), as well as on the interaction of cadherins with actin and on actin dynamics (Angres et al. 1996; Imamura et al. 1999; Vasioukhin et al. 2000; Drees et al. 2005; Yamada et al. 2005; Zhang et al. 2005; Cavey et al. 2008). How does an increase in adhesion strength lead to an increase in contact surface? In an ideal system in which no forces resist the deformations induced by changes in contact surface, a zipping-like mechanism could induce contact surface expansion. However, living cells represent a much more complex system in which the deformation of cell shape produces a restoring force—cortical elasticity—which resists to contact expansion. Stronger adhesion alone is thus unlikely to be sufficient to expand the contact surface. As we have seen in the first section, the expansion of cell–cell contacts relies on actin-based forces (protrusions mediated by actin polymerization, and acto-myosin tension), which cooperate to bring cell membranes in close apposition and thereby counteract deformation-induced resistance to contact expansion. Such forces are very likely to be required to increase contact surfaces in the examples mentioned above. One could even imagine that increased adhesion strength only acts via actin remodeling to promote contact surface expansion. The magnitude of adhesion strength (i.e., amount and/or adhesive strength of cadherin dimers) may thus not be directly relevant to contact expansion but may simply reflect the degree of mobilization of actin-based forces. Alternatively, adhesion may regulate the kinetics of contact expansion but not the equilibrium geometry of cell contacts.

Cortical Tension and Junction Dynamics

Recent studies have characterized some important features of junctional mechanics underlying morphogenetic processes (Farhadifar et al. 2007; Kafer et al. 2007; Rauzi et al. 2008). The models were inspired from work done with soap films in which bubbles assemble in geometries almost identical to that of cells

in an epithelium and which can be explained as resulting from progressive exploration of local surface energy minima (for details, see Carthew 2005; Lecuit and Lenne 2007). These studies suggest that the extent of cell–cell contacts is dictated by (1) local forces contributed by adhesion and acto-myosin tension, and (2) global forces generated by acto-myosin tension and dependent on the state of cell deformation.

Explaining Steady-State Cell Patterns

This paradigm has been used to explain pattern formation in the fly retina. A simple model that only incorporates the action of adhesive forces cannot recapitulate the patterns observed in wild type ommatidia and mutants in which the expression levels of cadherins are altered. In contrast, these different patterns are faithfully reproduced when including cortical tension as a force resisting cell adhesion (Kafer et al. 2007). Similarly, the steady state geometry of the wing imaginal disc epithelium before hexagonal packing can be modeled in terms of adhesive and elastic tension forces (Farhadifar et al. 2007). Comparing computer simulations to in vivo data yielded estimates of the relative contributions of elastic forces and line tensions at work at cell contacts and suggested that in this epithelium, local acto-myosin contractility predominates over adhesion (Farhadifar et al. 2007).

Tensile Networks Regulating Junction Dynamics

Following similar physical working hypotheses, a recent study probed the spatial distribution of junctional forces underlying the dynamics of tissue elongation (Rauzi et al. 2008). In gastrulating *Drosophila* embryos, epithelial cells of the ventro-lateral tissue (the germ band) intercalate to promote germ band elongation (GBE). This process involves the regulated disassembly of cell contacts between antero-posterior (A/P) neighbors and the creation of new contacts between dorso-ventral (D/V) neighbors (Bertet et al. 2004) (Fig. 4C). An anisotropy in Myo-II localization, more specifically, Myo-II

enrichment along shrinking A/P junctions, is required for intercalation (Bertet et al. 2004), suggesting that polarized Myo-II distribution could generate anisotropic tension that would drive cell intercalation. Computer simulations and comparisons with in vivo data indicated that anisotropic tension could be sufficient to drive GBE. Laser-nanodissection of cortical actin measured the anisotropy at AJs and its dependence on Myo-II (Rauzi et al. 2008).

It is worth noting that tensile forces, which shape cells, need not be generated directly at the cell cortex. This was recently illustrated by the observation that apical cell constriction, which underlies tissue bending and invagination in several developmental contexts (reviewed in Lecuit and Lenne 2007), is driven by contraction of a centrally located acto-myosin network in the plane of AJs (Martin et al. 2008). This contrasts with other models stating that an acto-myosin purse string physically attached to AJs is responsible for constriction. During *Drosophila* mesoderm invagination, acto-myosin aggregates located in the medial part of cells undergo pulses and pull on the cell cortex to drive constriction as a ratchet (Martin et al. 2008). These recent results add another level of complexity to the description of junctional mechanics controlling cell contacts and cell shape.

Interplay of Adhesion and Cortical Tension

As illustrated throughout this review, actin plays a lead role in the regulation of cell contacts, by sustaining adhesion and cortical tension, whose balance dictates the extent of cell contacts. The cadherin/catenin system provides one among several other mechanical links between the actin skeletons of adjacent cells and thereby integrates intra- and inter-cellular forces to the scale of the whole tissue during morphogenesis. How such coupling impacts on the amplitude and the spatial range of force transmission at the cell cortex is one of the most important and open fields of investigation now, in which the contributions of different candidate linkers will have to be evaluated in detail. The extent of frictions

between cadherins and contacting tensile actin networks is also likely to define the kinetics of cell shape changes. There is evidence that actin filaments are subdivided in functionally distinct populations, which are dedicated to specific purposes (Zhang et al. 2005; Cavey et al. 2008). Future work will have to focus on the mechanisms controlling the functional subdivision of the actin cytoskeleton in sub-cellular domains. It will be equally important to understand how the balance of forces is fine-tuned to reach equilibrium and obtain stable cell patterns in a tissue. Many experimental and pathological conditions show that misregulating the central players of adhesion and cortical tension leads to disequilibrium and has dramatic consequences on tissue integrity. For instance, excess tension can rupture cell–cell contacts (Sahai and Marshall 2002; Diogon et al. 2007), whereas the loss of cell–cell adhesion triggers epithelia–mesenchymal transitions (EMT) and is associated with cancer (reviewed in Thiery and Sleeman 2006; Baum et al. 2008).

Vertices as Central Regulatory Units of Epithelial Remodeling

Vertices are geometrical points in a tissue where three or more cells meet and can define both cell shape and cell dynamics. The number of neighbors a cell has simply reflects the number of vertices it is part of and the length of cell–cell contacts reflects the inter-vertices distances. Thus, epithelial remodeling can be explained as the displacement of vertices and most importantly the subsequent exchange of neighbors at vertices. Understanding how these two processes are regulated is thus key to understanding epithelial remodeling.

A drastic change in inter-vertices distance occurs during GBE in *Drosophila* embryos, when A/P junctions shrink during T1 transitions (Fig. 4C) (Bertet et al. 2004; Rauzi et al. 2008). Theoretically, anchoring of acto-myosin contractile bundles to the cell membrane specifically at vertices could be sufficient to remodel cell contacts. However, acto-myosin bundles are anchored to AJs puncta all along

cell–cell junctions and not simply at vertices (Cavey et al. 2008). Moreover, vertices impose structural limits on the movement of stable E-cad clusters that cannot “cross” a vertex. Consistent with this, nano-ablation experiments indicate that vertices represent physical barriers to the lateral displacement of retracting actin bundles and E-cad clusters (Cavey et al. 2008; Rauzi et al. 2008), supporting the idea that vertices represent special sites of attachment of acto-myosin bundles to the cortex.

The regulation of AJs stability specifically at vertices also requires further understanding. Anchoring of acto-myosin bundles to the cortex depends on AJs (Dawes-Hoang et al. 2005; Martin et al. 2008) suggesting that the stability of AJs complexes, namely homophilic E-cad clusters, at vertices is essential to drive junction shrinkage. However, T1 transitions, the process whereby four cells meeting at a vertex after junction shrinkage produce two new three-way vertices (Fig. 4C), requires local remodeling of homophilic E-cad clusters. How E-cad stability is regulated during this step is currently unknown. There may be specific mechanisms targeted to vertices to regulate AJs integrity, which remain to be characterized.

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