Integration of contractile forces during tissue invagination

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Introduction

Throughout development, epithelial sheets of cells undergo dramatic rearrangements, such as tissue folding and invagination, to generate complex three-dimensional organs in a process called epithelial morphogenesis (Leptin, 2005; Lecuit and Lenne, 2007; Quintin et al., 2008). Epithelial morphogenesis results from coordinated cell shape changes and movements that collectively deform tissues (Holtfreter, 1943; Gustafson and Wolpert, 1962; Sweeton et al., 1991; Shih and Keller, 1992). Individual cell shape changes and movements require that forces are generated within cells. In addition, for cellular forces to result in coherent changes in tissue architecture, forces must be transmitted between epithelial cells. The mechanical coupling of epithelial cells implies that the integration of cellular forces can in turn influence individual cell dynamics (Peralta et al., 2007; Gorfinkel et al., 2009; Pouille et al., 2009; Solon et al., 2009). Therefore, understanding tissue morphogenesis requires determining how cellular forces are integrated across tissues and whether tissue-level forces feed back to control individual cell behaviors.

A common cell shape change that results in epithelial morphogenesis is apical constriction. Apical constriction reduces the apical cross-sectional area of an epithelial cell, which transforms a columnar cell to a wedge or cone shape. Coordinated apical constriction of mechanically coupled cells promotes epithelial folding and tissue invagination (Odell et al., 1981). During Drosophila gastrulation, apical constriction appears to be required for the invagination of the prospective mesoderm, a strip of ~1,200 cells along the ventral midline of the embryo (Leptin and Grunewald, 1990; Parks and Wieschaus, 1991; Sweeton et al., 1991). Apical constriction of mesoderm cells is polarized, or anisotropic, such that cells constrict predominantly in the ventral–lateral (v–l) direction, remaining longer along the anterior–posterior (a–p) axis (Fig. 1 A; Sweeton et al., 1991). The mechanism responsible for anisotropic constriction is not known, but one consequence is that whereas the tissue contracts in the v–l direction, a–p tissue length remains relatively constant, resulting in a long, narrow ventral furrow.

Epithelial tears allow cells normally elongated along the a–p axis to constrict isotropically, which suggests that apical constriction generates anisotropic epithelial tension that feeds back to control cell shape. Epithelial tension requires the transcription factor Twist, which stabilizes apical myosin II, promoting the formation of a supracellular actomyosin meshwork in which radial actomyosin fibers are joined end-to-end at spot AJs. Thus, pulsed actomyosin contractions require a supracellular, tensile meshwork to transmit cellular forces to the tissue level during morphogenesis.

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Abbreviations used in this paper: AJ, adherens junction; a–p, anterior–posterior; Arm, Armadillo; dsRNA, double-stranded RNA; v–l, ventral–lateral.
Results

Anisotropic apical constriction possibly reflects tissue mechanics

Polarized constriction could result from an individual cell behavior, such as directional actomyosin-based force generation along the v-l axis. Alternatively, a mechanical property of the tissue, such as greater tension along the a-p axis, could bias the direction in which cells constrict. Because ventral furrow cell apical constriction is pulsed (Martin et al., 2009), we decided to distinguish between these possibilities by determining whether the increase in cell polarization correlates with pulses of actomyosin contraction in individual cells. To visualize apical cell shape, we imaged ventral furrow cells in live embryos expressing the membrane markers Spider-GFP and Resille-GFP (Morin et al., 2001). We quantified the polarization of apical constriction by fitting apical cell outlines with ellipses and measuring two parameters (Fig. 1 B). Aspect ratio and anisotropy had similar values in elongated cells during furrow formation because cells were aligned along the a-p axis such that $\theta \approx 0$ (unpublished data).
Mean anisotropy and aspect ratio gradually increased to ~2 before cells invaginated into the embryo interior (Fig. 1, C and D). Analysis of individual cells demonstrated that transient increases in anisotropy did sometimes coincide with contraction pulses (Fig. 1, E and F, i). However, anisotropy also increased between contractions (Fig. 1, E and F, ii). Overall, there was not a strong correlation between the contriction rate and the rate of change in anisotropy (r = 0.1 ± 0.24, n = 125 cells, three embryos), which suggests that directional contraction of individual cells is not solely responsible for the anisotropy. Furthermore, anisotropic constriction still occurred in bicoid nanos torso-like (bnt) embryos that lack a-p polarity, which demonstrates that directional information for individual cell constriction is not provided by the a-p patterning system (Fig. 1 D). These results suggest that rather than being caused by discrete cellular behaviors, anisotropic apical constriction could result from the global mechanics of the tissue.

**AJs integrate contractile forces to generate epithelial tension**

AJs, which contain components of the transmembrane adhesion molecule E-cadherin and the adaptors β-catenin and α-catenin, link neighboring cells to each other and link the cell surface to the actin cortex (Gumbiner, 2005). Partial loss-of-function mutants in the Drosophila β-catenin (Armadillo [Arm]), such as armΔ4301, result in lower levels of Arm, E-cadherin, and α-catenin at the cell surface (Cox et al., 1996; Sokac and Wieschaus, 2008; Sawyer et al., 2009). Therefore, to determine whether there are mechanical properties of the ventral furrow that might explain cell anisotropy, we examined the consequences of reducing AJ levels using maternal and zygotic armΔ4301 mutants (armΔ2). Because contractile forces for apical constriction are generated by myosin II on the apical surface, we first visualized apical myosin-GFP in force for apical constriction are generated by myosin II on Video 1; Martin et al., 2009). In some meshwork on the ventral surface of the embryo (Fig. 2 A and Video 2). Laser ablation has served as an excellent method to measure relative tensile forces in epithelial tissues undergoing morphogenesis (Kiehart et al., 2000). Targeted UV laser irradiation severs actin- and myosin II–containing structures, resulting in the recoil of the surrounding tissue whose initial velocity (displacement/time) immediately after ablation is proportional to the magnitude of the resting tension (Hutson et al., 2003; Fernandez-Gonzalez et al., 2009). Because tissue retraction associated with local ablations could be influenced by the orientation of cellular actomyosin structures, we first made 20-µm-long multicellular laser incisions along either the a-p or v-l axes and measured the initial myosin II displacement perpendicular to the incision at various distances from the incision (Fig. 3 A). Incisions across the width of the ventral furrow resulted in dramatic myosin displacement all along the a-p axis of the embryo (Dw), which indicates tension along this axis (Fig. 3 B and Video 2). Laser incisions parallel to the furrow resulted in significantly less displacement along the v-l axis (Dv; Fig. 3 B and Video 2), which is consistent with there being lower v-l tension. To more accurately measure the initial velocity of tissue retraction, we also performed 1-µm point ablations in which we imaged myosin II displacement within 3 s of initiating ablation (compared with 9 s for line ablation, Fig. S2). Ablation of isometric-looking myosin structures resulted in retraction of the surrounding tissue (Fig. 3 D). To determine if this displacement was directional, we measured the initial radial displacement of myosin structures (Dr) from the site of ablation at different angles (θ) representing different directions relative to the a-p and v-l axes (Fig. 3 E). The initial radial displacement in wild-type embryos was 2–3-fold higher along the a-p axis of the embryo (Fig. 3 F). These results confirm the conclusions from our analysis of armΔ2 mutants and demonstrate that epithelial tension is predominantly directed along the length of the ventral furrow.

**Epithelial tension along the a-p axis causes anisotropic apical constriction**

The higher tension along the a-p axis suggests the possibility that ventral furrow cells attempt to constrict isotropically, but encounter more resistance and thus constrict less along the a-p axis. To determine whether epithelial tension biases constriction to occur in the v-l direction, we examined how the disruption of epithelial tension that results from AJ depletion affects cell shape change in both live and fixed embryos.
epithelial tearing events (Fig. 4, D, E, and F). These results strongly suggest that individual cells are not hard-wired to undergo polarized contraction, but that apical constriction generates higher epithelial tension along the a-p axis, providing a mechanical constraint that feeds back to control the resulting individual cell shape.

Actomyosin fibers connected by spot AJs form a supracellular meshwork

We next examined how contractile forces are transmitted across the tissue. The epithelial tears in \( \text{arm}^{M/Z} \) embryos suggested that apical actomyosin meshworks are coupled to each other through AJs. Therefore, we examined how cytoskeletal structures are linked between cells during ventral furrow formation.
using fixed embryos labeled with myosin II, actin, and E-cadherin. Before furrow formation, the majority of myosin II was present as isolated spots on the apical surface of individual cells, the largest of which likely resulted from pulses of myosin II coalescence (Fig. 5 A). As myosin II levels increased and a furrow developed, myosin II and actin formed supracellular...
fibers that extended across multiple cells and were incorporated into the tissue-wide meshwork (Fig. 5 A and B). Supracellular actomyosin fibers bridged the apical surface of the cell, such that the ends of myosin II fibers met at cell–cell interfaces (Fig. 5 B, i). Multiple radial actomyosin fibers often originated from a central myosin II spot or ring, which formed branches in the tissue-wide meshwork (Fig. 5 B, i). In addition to radial actomyosin fibers, some supracellular actomyosin fibers ran along cell boundaries (Fig. 5 B, ii). In ventral furrow cells, subapical AJs ~4 µm below the apical cortex are disassembled and reappear at the apical surface as spotlike structures between cell vertices (Fig. 5 C; Dawes-Hoang et al., 2005; Kölsch et al., 2007; Martin et al., 2009). Spot AJs appeared to connect radial actomyosin fibers end-to-end to generate the supracellular fibers (Fig. 5 C). Interestingly, spot AJs linking actomyosin fibers were often stretched along the fiber, possibly indicating tension being exerted on the junction (Fig. 5 C). Thus, spot AJs appear to integrate cellular actomyosin networks into supracellular actomyosin fibers, resulting in a tissue-wide actomyosin meshwork.

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Figure 4. Anisotropic apical constriction results from α-p epithelial tension. (A and B) Apical myosin II (green) and subapical F-actin used to visualize cell outlines (white) in fixed armRNAi embryos. (A) A tear similar to those observed in live embryos. Red arrows indicate cells that have constricted isotropically. (B) A later-staged embryo where myosin II has rounded up into rings. (C) Disruption of AJs results in loss of cell–cell adhesion. Time-lapse images of cell outlines [Spider-GFP] during a tear in an armRNAi embryo. Red cell–cell contacts are lost and different cells come into contact [blue]. (D) Isotropic apical constriction occurs upon tearing. Time-lapse images of cell outlines [Spider-GFP] in an E-CadherinRNAi embryo. Blue dots indicate initially anisotropic cells that constrict isotropically upon loss of epithelial integrity. Apical area (E) and anisotropy (F) were quantified for the cells labeled in D. Dotted lines indicate the time of the tear. Bars: (A and B) 10 µm; (C) 5 µm; (D) 5 µm.
Spot AJs transmit tension from actomyosin fibers across the plasma membrane

To test whether spot AJs transmit tension between cellular actomyosin fibers, we examined more closely the cellular organization and dynamics of myosin II in embryos with reduced AJs. Supracellular actomyosin fibers initially formed in arm<sup>WZ</sup> embryos, which is consistent with this allele retaining partial function (Fig. 2 A). However, upon epithelial tearing, myosin II fibers rapidly contracted, demonstrating that myosin II fibers generate contractile force in a manner similar to stress fibers (Fig. 2 C, i; and Video 4; Kumar et al., 2006). Loss of cell adhesion resulted in myosin II contraction into rings or balls across the entire epithelium (Fig. 2, A and C, ii and iii; and Fig. 4, A and B). Myosin II spots continuously flowed into myosin II rings, suggesting that unrestrained actomyosin contraction occurs in the absence of opposing contractile forces from surrounding cells (Fig. 2 C, iii; and Video 5). These results suggest that spot AJs balance tensile forces generated by cellular actomyosin networks to allow the formation of stable actomyosin fibers that transmit tension across the tissue.

The retraction of actomyosin fibers in embryos with reduced AJs could reflect a disruption of the extracellular linkage between adjacent cell plasma membranes or the intracellular connection between the plasma membrane and the actomyosin cortex. To distinguish between these models, we examined whether cell membranes completely separate during tearing events using scanning EM. In the ventral furrow of wild-type embryos, boundaries between cells cannot be discerned because of tight adhesion of these cells through apical AJs (Fig. 6 A). In contrast, individual
ventral cells can easily be identified in arm\textsuperscript{MZ} mutants because cells lose tight adhesion to one another, specifically in the ventral furrow region (Fig. 6 B). In addition, we identified tearing events in which adjacent rows of cells became separated (Fig. 6 C). Scanning EM demonstrated that the cell expansion observed using confocal microscopy represented cells rounding up upon loss of adhesion. Consistent with the global contraction of the surrounding tissue in live embryos, the isotropically constricting cells adjacent to the tear contained dramatic membrane blebs, which suggests that the apical cortex rapidly contracts upon loss of epithelial tension across the tissue.

Despite the dramatic loss of adhesion, cells in arm\textsuperscript{MZ} embryos maintained thin membrane connections, or tethers, to their neighbors (Fig. 6, B and C). Live imaging of the apical surface during tearing events demonstrated that membrane tethers form as cell apices contract away from each other (Video 6). Membrane tethers remained connected to clusters of membrane blebs, which colocalized with myosin II rings and some remaining cortical E-cadherin (Figs. 6 D and S1 C). Tethers often contained detectable levels of F-actin, but not myosin II, which suggests that tethers fail to transmit tension (Fig. S1 D). We propose that membrane tethers form when the actomyosin cortex detaches from the one side of a fiber–AJ–fiber connection (Fig. 6 E). This detachment did not indicate a specific function for β-catenin in coupling the actomyosin cortex to AJs because membrane tethers were also observed in both \textit{α-cateninRNAi} and ECad2RNAi embryos (Fig. S1 E). Thus, reducing the number of AJs possibly focuses the large amount of contractile force generated by ventral furrow cells onto the few remaining AJ–cytoskeleton connections, exceeding the forces of adhesion and causing junctions to detach from the tensile meshwork. However, because we have been unable to completely disrupt AJs, we cannot rule out the possibility that membrane tethers result from cell–cell adhesion that is E-cadherin and Arm independent. Regardless of the exact nature of the membrane connection, tether formation upon disruption of AJs further argues that spot AJs balance contractile forces across the plasma membrane, allowing tension to be transmitted between cells.

**Epithelial tears and loss of cell-cell adhesion fail to occur in the absence of Snail-mediated actomyosin contraction**

We have hypothesized that the epithelial tears observed in embryos with disrupted AJs result from contractility of the supracellular actomyosin meshwork. We previously showed that the transcription factor Snail is required for pulsed contractions and apical actomyosin meshwork assembly (Martin et al., 2009). Therefore, we determined whether Snail is required for the epithelial tears and loss of cell adhesion observed in arm mutant embryos by performing double RNAi experiments in which the expression of sna was reduced in combination with arm. A control double-stranded RNA (dsRNA; CG3651RNAi) had no
To occur in the absence of Snail-mediated actomyosin assembly, supporting the model that tension generated by the actomyosin contraction is responsible for these phenotypes.

Twist-mediated stabilization is required to transmit epithelial tension

The transcription factor Twist affects actomyosin contractility in the ventral furrow differently than Snail. Pulses of actomyosin contractility in tissues • Martin et al.

Figure 7. Snail and Twist are required for epithelial tearing. (A) Images of myosin-GFP in live embryos injected with the indicated dsRNA. snaRNAi inhibits the formation of myosin II rings and the loss of cell–cell adhesion normally observed in armRNAi embryos. twiRNAi inhibits tearing, but myosin contractions still occur. (B) Images of cell outlines (Spider-GFP) in armRNAi and arm-twiRNAi embryos. Knockdown of Twist suppresses the formation of discrete epithelial tears. (C) Epithelial tears and loss of cell–cell adhesion occur within the Snail expression domain. Images of a fixed armM/Z embryo stained for Snail (green) and neurotactin (purple, cell outlines). An apical image of neurotactin is included to illustrate the position of the tear. The ventral midline is offset to visualize the lateral border of the mesoderm where two Snail-expressing nuclei do not initially lose adhesion [white bracket]. Bars, 10 µm.

effect on its own or with armRNAi (Fig. 7 A and Video 7), which demonstrates that multiple dsRNAs can be combined without affecting the potency of RNAi. Knockdown of sna and arm suppressed the formation of epithelial tears and the loss of adhesion normally observed in armRNAi embryos (n = 5 embryos; Fig. 7 A and Video 8). In addition, snaRNAi suppressed apical myosin II recruitment and contraction into rings in armRNAi embryos. Thus, epithelial tearing and loss of cell adhesion fail to occur in the absence of Snail-mediated actomyosin assembly, supporting the model that tension generated by the actomyosin contraction is responsible for these phenotypes.
contraction still occur in *twi* mutant and *twiRNAi* embryos. However, the resulting constriction fails to be stabilized, possibly because of a defect in maintaining cortical tension (Martin et al., 2009). Therefore, we examined whether Twist is required for the epithelial tears in *armRNAi* embryos to distinguish the roles of pulsed actomyosin contraction and Twist-dependent stabilization in generating tissue-wide epithelial tension. Pulses of myosin II coalescence occurred in *twi-armRNAi* embryos, similar to *twiRNAi* alone (Fig. 7 A and Video 9). Similar to *armRNAi* embryos, cells lost adhesion and formed tethers and dramatic blebs in *twi-armRNAi* embryos (Figs. 7 B and S1 F). Thus, pulsed contractions generate local tension that can disrupt cell–cell adhesion. Pulses continued even after cells lost adhesion, which suggests that the pulsed contraction does not require mechanical interactions between cells. Importantly, cell dissociation occurred without the formation of tissue-wide tears along the a-p axis of the embryo (n = 10 embryos; Fig. 7, A and B; and Videos 9 and 10), which suggests that pulsed actomyosin contractions are insufficient to generate global epithelial tension. To further examine Twist’s role in generating tension, we performed laser incisions across the width of the furrow and measured a-p myosin displacement (D_a-p) in control (H_2O-injected) and *twiRNAi* embryos. *twiRNAi* embryos exhibited significantly lower levels of recoil at all distances from the incision, demonstrating that Twist is indeed required to transmit tension between ventral furrow cells (Fig. 3 C). Because Twist enhances Snail expression, it was possible that the disruption of epithelial tension was caused by lower levels of Snail activity (Kosman et al., 1991; Leptin, 1991). However, knockdown of the two Twist targets, *fog* and *t48* (Costa et al., 1994; Kölsch et al., 2007), also suppressed the formation of tissue-wide epithelial tears in *fog-t48-armRNAi* embryos, demonstrating that Twist activity is independent of Snail (unpublished data). Consistent with the importance of Twist in generating tissue-wide tears, epithelial tears in *arm^mc2* mutants are restricted to ~14 cells within the 18 cell–wide Snail expression domain (Fig. 7 C), corresponding to the domain of highest Twist expression, where transcriptional targets such as folded gastrulation are activated (Leptin, 1991; Costa et al., 1994). Thus, Twist-mediated stabilization is required to transmit tension across the tissue, which is required for epithelial tears in *arm* mutant embryos.

**Twist stabilizes apical myosin II fibers between contraction pulses to form a supracellular actomyosin meshwork**

Why do pulsed myosin II contractions in *twiRNAi* embryos fail to transmit tension across the tissue? To address this question, we used myosin-GFP and membrane-mCherry to compare cellular myosin II dynamics in wild-type and *twiRNAi* embryos. In wild-type cells, myosin II fibers remain on the apical cortex after contraction pulses during the stabilization phase of apical constriction (Fig. 8 A; Martin et al., 2009). This results in a gradual increase in cortical myosin II levels over a series of contractions (Fig. 8 B). In contrast, myosin II seemed to disappear from the apical cortex after contraction pulses in *twiRNAi* embryos, with myosin II levels falling back to near the same level as before the pulse (Fig. 8, A and B). To determine whether *twiRNAi* affects cortical myosin II levels after contraction, we averaged normalized myosin II intensity profiles for individual pulses in control (CG3651RNAi) and *twiRNAi* embryos. We found that mean cortical myosin II intensity in *twiRNAi* embryos dropped to a significantly lower level after a contraction pulse than control embryos (Fig. 8 C; P < 0.0001). This Twist-dependent stabilization of cortical myosin II was even more dramatic in *arm-twRNAi* embryos. In contrast to *armRNAi* embryos, where myosin II contracts into a stable cortical ring in each cell, myosin contractions in *arm-twRNAi* embryos were pulsed, with myosin II disappearing from the cortex after coalescing (Fig. 8 D). Myosin II dynamics in *arm-twRNAi* embryos were indistinguishable from *fog-t48-armRNAi* (n = 5 embryos), demonstrating that destabilization of cortical myosin II resulted directly from loss of function in the Twist pathway and not from lower Snail activity (Fig. 8 D). Given that contraction pulses are asynchronous in adjacent cells (Martin et al., 2009), we hypothesized that the failure to retain cortical myosin II between pulses would disrupt supracellular meshwork formation because only a subset of cells would have cortical myosin II at a given time. To test this, we segmented myosin-GFP images from control and *twiRNAi* embryos and measured the area of the largest myosin-GFP structure at each time point (Fig. 8 E). With our segmentation threshold, control embryos assembled large myosin II networks on the order of 200–300 µm², or the size of several cells (Fig. 8 F). However, *twiRNAi* embryos rarely exhibited myosin II networks larger than 50 µm², which corresponds to the apical area of an individual cell. We propose that Twist-dependent stabilization of apical myosin II fibers is required to form the supracellular actomyosin meshwork, which allows tension to be stably transmitted across the epithelium during ventral furrow formation.

**Discussion**

**AJs integrate tensile force along the length of the ventral furrow**

Recent studies have demonstrated the importance of actomyosin and AJs for individual cell shape changes and morphogenesis (Bertet et al., 2004; Dawes-Hoang et al., 2005; Franke et al., 2005; Blankenship et al., 2006; Gorfinkel and Arias, 2007; Martin et al., 2009; Sawyer et al., 2009). However, less is known about the underlying mechanisms by which dynamic contractile forces are transmitted between cells of an epithelium and the resulting tissue mechanics. In this study, we examined the mechanics of ventral furrow formation both by lowering AJ levels and by ablating the apical actomyosin meshwork. Reducing AJ levels sensitized cell adhesion to stress, resulting in discrete, tissue-wide epithelial tears. We propose that tears in AJ-depleted embryos result from the loss of mechanical connections between cells when contractility exceeds the forces of adhesion. Interestingly, tears resulted from tissue separation along the a-p axis, which suggests that tension transmitted by AJs is mostly oriented along the length of the ventral furrow (Fig. 9 A). We confirmed this anisotropy in epithelial tension by showing that laser ablations in the ventral furrow also result in preferential...
Epithelial tension controls individual cell shape

Ventral furrow cells undergo polarized apical constriction, constricting predominantly in the v-l direction and remaining longer along the a-p axis (Sweeton et al., 1991). Because we failed to find a correlation between contraction pulses and cell anisotropy and because epithelial tension itself is anisotropic, we tested whether polarized apical constriction results from the mechanical constraints of the tissue. Indeed, epithelial tears allowed adjacent anisotropic cells to constrict isotropically, which suggests that ventral furrow cells attempt to constrict isotropically, but encounter greater resistance to constriction along the a-p axis (Fig. 9 B). During germ band extension, tension generated by supracellular actomyosin cables influences myosin actomyosin retraction along the a-p axis. The global a-p retraction of the surrounding tissue after epithelial tearing suggests that the ventral furrow is a mechanically coherent epithelium, in which high levels of tensile force are integrated along the length of the furrow.

It is surprising that the vast majority of the epithelial tension generated by apical constriction is oriented along the length of the furrow, the axis with the least tissue movement. Indeed, one might have expected greater levels of tension along the v-l axis given that apical constriction is thought to pull lateral cells toward the ventral midline around the circumference of the embryo (Fig. 9 A). Future experiments are required to determine how anisotropic tension in the ventral furrow contributes to the bending and invagination of this epithelium.

**Figure 8.** Twist is required for myosin II stabilization and supracellular meshwork formation. (A) Twist is required to stabilize apical myosin II fibers after contraction. Time-lapse images of myosin-GFP (green) and membrane-mCherry (purple). Bar, 5 µm. (B) Quantification of apical area (red) and myosin intensity (blue) in individual cells from wild-type and twiRNAi embryos. (C) Twist is required to maintain cortical myosin II after a contraction pulse. Normalized myosin intensity for individual contraction pulses was averaged. Data are means ± SD, which is indicated by the error bars (control: n = 329 pulses, 120 cells, two embryos; twiRNAi: n = 524 pulses, 106 cells, two embryos). (D) Twist is required to stabilize myosin II rings. Time-lapse images of myosin-GFP in embryos treated with arm dsRNA together with either twi or log/148 dsRNA. Bar, 5 µm. (E) Myosin-GFP (top and bottom, green) in live control or twiRNAi embryos. Images were segmented using intensity and area thresholds to identify myosin II structures (bottom, red). Bar, 10 µm. (F) Twist is required to form the supracellular actomyosin meshwork. The area of the largest myosin II structure was quantified for each time point in videos of control-injected or twiRNAi embryos.

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junctions and form supracellular actin meshworks across the apical surface of epithelia (Yonemura et al., 1995; Adams et al., 1996; Vaezi et al., 2002). Furthermore, actin meshworks consisting of radial actin fibers emanating from central actin spots are also observed in *Xenopus* cells undergoing convergence and extension (Skoglund et al., 2008; Rolo et al., 2009). Actomyosin organization as a two-dimensional branched meshwork could represent a different mechanical paradigm for epithelial morphogenesis that enables cells to form a mechanically coherent sheet in which tensile forces are not restricted to cell interfaces, but bridge the surface of cells across the entire tissue.

**Figure 9.** Causes and consequences of contractile force integration. (A) Distribution of apical tensile forces that accompany ventral furrow formation. Black arrows indicate the movement of lateral cells toward the ventral midline. Red arrows indicate epithelial tension, which is predominantly directed along the length of the furrow (a-p axis). (B) Tissue-level forces influence individual cell shape changes. Ventral furrow cells attempt to contract isotropically (black arrows), but a-p tension (red arrows) resists constriction in this direction, resulting in anisotropic constriction. (C) Model for how contractile forces are integrated to generate epithelial tension in the ventral furrow. In wild-type embryos, pulses of actin (red) and myosin II (green) contraction constrict cell apices, and actomyosin fibers that remain on the apical surface between pulses maintain cortical tension (black arrows above cells), thus stabilizing cell shape. Actomyosin fibers linked by spot AJs (blue) form a supracellular meshwork that allows forces to be stably transmitted between cells, generating global epithelial tension (red arrows). In the absence of Twist, cells lack actomyosin fibers between contractions and fail to assemble a supracellular actomyosin meshwork. Thus, contraction pulses in cells stretch their immediate neighbors, and tension fails to be propagated across the epithelium.

**Spot AJs transmit tensile forces oriented perpendicular to cell interfaces**

Epithelial tension observed during ventral furrow formation likely resulted from contractile forces generated by apical actomyosin meshworks. We found that cellular actomyosin fibers were coupled end-to-end at spot AJs to form supracellular fibers and a tissue-wide meshwork. Disruption of AJs caused an imbalance of forces across the plasma membrane, resulting in membrane tethers. This supports a model in which tensile forces are integrated through discrete AJ anchors that link actomyosin networks in adjacent cells (Fig. 9 C). Supracellular actomyosin cables that transmit tension parallel to cell interfaces have been observed during germ band extension, dorsal closure, and compartment boundary formation (Rauzi et al., 2008; Rodriguez-Diaz et al., 2008; Fernandez-Gonzalez et al., 2009; Landsberg et al., 2009). However, in the ventral furrow, supracellular actomyosin fibers are mostly oriented perpendicular to cell interfaces (Fig. 9 C). This orientation is similar to that of actin in forming cultured epithelia, where radial actin fibers terminate in spot junctions and form supracellular actin meshworks across the apical surface of epithelia (Yonemura et al., 1995; Adams et al., 1996; Vaezi et al., 2002). Furthermore, actin meshworks consisting of radial actin fibers emanating from central actin spots are also observed in *Xenopus* cells undergoing convergence and extension (Skoglund et al., 2008; Rolo et al., 2009). Actomyosin organization as a two-dimensional branched meshwork could represent a different mechanical paradigm for epithelial morphogenesis that enables cells to form a mechanically coherent sheet in which tensile forces are not restricted to cell interfaces, but bridge the surface of cells across the entire tissue.

**Pulsatile and stable contractile networks cooperate to change cell shape and transmit tension across the tissue**

Knockdown of the transcription factor Twist using RNAi separates the activities of pulsatile actomyosin contraction and cell shape stabilization between contractions (Martin et al., 2009). Pulses of actomyosin contraction continued to occur in both *twi*RNAi and *arm-twিRNAi* embryos. In *arm-twিRNAi* embryos, pulsed contractions disrupted cell–cell adhesion and formed membrane tethers, which suggests that pulsed contractions generate cellular tension that can separate the actomyosin network from AJs. A recent study of amnioserosa apical constriction proposed that contraction pulses are activated by mechanical interactions (i.e., stretching) between cells (Solon et al., 2009). However, in ventral furrow cells, pulsatile actomyosin contractions occurred even after cells lost adhesion, which suggests that pulsed contractions are fundamentally cell autonomous, possibly resulting from dynamic behaviors of cytoskeletal networks.

**Figure 9.** Causes and consequences of contractile force integration. (A) Distribution of apical tensile forces that accompany ventral furrow formation. Black arrows indicate the movement of lateral cells toward the ventral midline. Red arrows indicate epithelial tension, which is predominantly directed along the length of the furrow (a-p axis). (B) Tissue-level forces influence individual cell shape changes. Ventral furrow cells attempt to contract isotropically (black arrows), but a-p tension (red arrows) resists constriction in this direction, resulting in anisotropic constriction. (C) Model for how contractile forces are integrated to generate epithelial tension in the ventral furrow. In wild-type embryos, pulses of actin (red) and myosin II (green) contraction constrict cell apices, and actomyosin fibers that remain on the apical surface between pulses maintain cortical tension (black arrows above cells), thus stabilizing cell shape. Actomyosin fibers linked by spot AJs (blue) form a supracellular meshwork that allows forces to be stably transmitted between cells, generating global epithelial tension (red arrows). In the absence of Twist, cells lack actomyosin fibers between contractions and fail to assemble a supracellular actomyosin meshwork. Thus, contraction pulses in cells stretch their immediate neighbors, and tension fails to be propagated across the epithelium.
(Munro et al., 2004; Paluch et al., 2005). Importantly, our data also showed that pulsed contractions were insufficient to generate epithelial tears. Thus, an additional mechanism is required to translate pulsatile contractile forces and cell shape changes into tissue-wide epithelial tension.

Twist-mediated stabilization is critical to transmit cellular forces to the tissue level. We show here that Twist was required for apical myosin II fibers to remain on the apical cortex between contraction pulses. These actomyosins fibers bridging the apical cortex likely stabilize the cell apex, preventing previously observed stretching events that would dissipate force transmission across the tissue (Martin et al., 2009). Furthermore, Twist-dependent myosin II stabilization allowed the formation of the supracellular actomyosin meshwork, which was critical for generating tissue-wide tension. We propose that this actomyosin meshwork results in a mechanically integrated tissue, allowing propagation of both pulsatile and stable tensile forces (Fig. 9 C). Thus, while pulsed contractions generate dramatic changes in individual cell shape, stabilizing tension produced by the supracellular actomyosin meshwork transmits contractile forces to the tissue level, generating epithelial tension.

Pulsatile and stable contractile forces also cooperate to contract the amnioserosa during dorsal closure (Solon et al., 2009). During dorsal closure, amnioserosa contraction is stabilized by actomyosin cable assembly in the surrounding epidermis. However, in the ventral furrow, the machinery responsible for pulsation and stabilization are both present in the mesoderm. Future studies are needed to determine how actomyosin networks with unique properties are coordinated within the same cell. Interestingly, Twist is thought to activate G protein–coupled receptor signaling, which activates stress fiber formation in cultured cells (Buhl et al., 1995; Lin et al., 2005). Thus, Twist possibly induces actomyosin fiber assembly independently of pulsed contractions. Alternatively, pulsed contractions could bundle actomyosin meshworks to form fibers that become stabilized by Twist activity. Regardless of these different models, the cooperation of pulsatile and stable networks in morphogenetic events as diverse as epithelial invagination and sheet migration suggests that coordinated pulsation and stabilization represents a conserved mode of tissue morphogenesis.

### Materials and methods

#### Fly stocks and genetics

The following stocks containing fluorescent fusion proteins were used: Spider-GFP (95–1) and Resilie-GFP (117–2) (Morin et al., 2001), membrane-mCherry [this paper], myosin-mCherry [sqh-GFP; Royo et al., 2002], and myosin-mCherry [sqh-GFP; Martin et al., 2009]. To examine cell shape in embryos devoid of α-p polarity, we used the stock w; Resilie-GFP; bicoid
d0 nanos2 torso-like44/43/TM3; Sb. We analyzed embryos from mothers that were homozygous for bicoid
d0 nanos2 torso-like44/43. To generate armV2 mutants, we created armE3401 germ-line clones using the FLP/DFS system (Chou and Perrimon, 1992). We visualized myosin II in armV2 mutants by generating a stock that was armV2 and low GFP (Royo et al., 2002). We crossed females of this genotype to w ovo FRT101/Y; flip-138 males to obtain armE3401 FRT101/w ovo FRT101; flip-138/+ females. These females were heat shocked as larvae for 2 h at 37°C each day to induce mitotic recombination in the germ line. We imaged embryos from the following cross: armE3401 FRT101/w ovo FRT101; flip-138/+ females x FM7+/++; flip-138/+ males. Half of these embryos showed loss of cell–cell adhesion, which is consistent with half being rescued zygotically.

**Construction of membrane-mCherry**

A membrane-mCherry marker was created by fusing the N-terminal 20 amino acids of the rat Gap43 gene, which contains a myristoylation sequence, to mCherry. This membrane-mCherry fusion was cloned into a pBluescript vector containing the sqh promoter and 3′ untranslated region (Martin et al., 2009). The 3-kilobase KpnI/XbaI sqh-mCherry::mCherry-sqh3′ fragment was then inserted into a transformation vector containing the attB site (pTiger; courtesy of S. Ferguson, State University of New York at Fredonia, Fredonia, NY). The resulting construct was sent to Genetic Services, Inc., for integration into the attP2 site using the phiC31 integrase system (Groth et al., 2004).

#### Live cell imaging

To prepare embryos for live imaging, embryos were dechorionated with 50% bleach, washed with water, and mounted ventral side up on a slide covered with embryo glue (Scotch tape resuspended in heptane) between two No. 1.5 coverslip spacers. A coverslip was placed over the embryo and Halocarbon 27 oil was added to the resulting chamber. The spacers prevent the overlying coverslip from contacting the ventral surface of the embryo such that the embryo was not compressed. Embryos imaged under these conditions developed normally and subsequently hatched, demonstrating that our imaging conditions had minimal impact on development. All imaging was performed in Halocarbon 27 oil at room temperature (−23°C).

**Spider-GFP, Resilie-GFP, sqh-mCherry;Spider-GFP, and sqh-GFP; membrane-mCherry videos were obtained with a confocal microscope (SP5; Leica), a 63×/1.3 NA glycine-immersion objective (Leica), an argon ion laser, and a 561-nm diode laser. Images were acquired using a pinhole setting from 1–2 airy units. For multichannel imaging, we set the excitation bandpass to 495–550 nm to detect GFP and 578–650 nm to detect mCherry. sqh-GFP single-channel videos were obtained using the aforementioned imaging system or with a spinning disk confocal microscope (Ultraview; PerkinElmer) controlled with Volocity Acquisition software (Improvision), a 60×/1.4 NA oil-immersion objective (Nikon), an argon/ krypton laser, and an electron-multiplying charge-coupled device camera (Cy100-13; Hamamatsu).

**Laser ablation**

Embryos were prepared for live imaging and were imaged using an Ultraview spinning disk confocal microscope equipped with a 63×/1.4 NA oil-immersion lens controlled with MetaMorph software (Universal Imaging). An N2 Micropoint laser (Photronics Instruments) tuned to 365 nm was focused on sqh-GFP structures on the ventral surface of the embryo to ablate actomyosin structures. Point ablations were performed by ablating isotropic sqh-GFP spots with a point ~1 µm in diameter, which took ~670 ms. Line ablations were performed by making nine sequential point ablations to make a 20-µm incision, which took ~6.4 s. Z stacks were acquired immediately before and after ablation in order to measure displacement of myosin II structures upon release of tension.

#### Image processing and analysis

Images presented were processed using Image (http://rsb.info.nih.gov/ij/) and Photoshop CS (Adobe Systems, Inc.). A Gaussian smoothing filter with a radius of one pixel was used to reduce noise in published images. Myosin II images presented in all figures are maximum intensity z projections of the apical ~5 µm of cells in the middle of the image. Because myosin II is almost entirely present on the apical surface of cells, these images represent a surface projection of the embryo. Images of cell outlines are z slices ~2 µm below the apical surface.

**Image segmentation** was performed using custom MATLAB (MathWorks) software. Raw images were bandpass filtered with effective cutoff wavelengths of ~1.4 µm (low) and ~6.5 µm (high) for individual cells. We measured the intensity of cortical myosin II in individual cells using three-dimensional time-lapse videos that were ~5 µm in depth. To separate cortical myosin II structures from the diffuse cytoplasmic staining, we smoothed sqh-GFP images using a Gaussian smoothing filter with a three-pixel kernel size, σ = 0.5 pixels, and clipped intensity values to three standard deviations above the mean. We then made maximum-intensity z projections of myosin II (averaging the two highest-intensity values) and integrated the intensity

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of all the pixels in a given cell. Data for apical area, myosin intensity, and anisotropy were smoothed using a Gaussian smoothing filter (σ = 18–24 s, three point times) to remove noise.

Recoil of sqh-GFP structures after laser ablation was quantified using custom software in which myosin spots were hand selected and tracked from the pre-ablation frame to the post-ablation frame (Fernandez-Gonzalez et al., 2009).

To measure the continuity of the supracellular meshwork in control-injected and twiRNAi embryos, we thresholded maximum-intensity projections of sqh-GFP images using the mean pixel intensity as a cutoff. We then used the “bwmorph” function in MATLAB to identify connected objects and identified the largest object at each time point in a time-lapse video.

Embryo fixation and staining

To visualize cytoskeletal structures, embryos were fixed with 8% paraformaldehyde/heptane for 30 min, manually devitellinized, stained, and mounted in AquapolyMount (Polysciences, Inc.). Endogenous sqh-GFP fluorescence was used to visualize myosin II, and Alexa Fluor 568 phalloidin (Invitrogen) was used to visualize F-actin. Drosophila E-Cadherin (DCad2; Developmental Studies Hybridoma Bank) and rabbit anti-Snail (a gift from M. Biggin, Lawrence Berkeley National Laboratory, Berkeley, CA) at dilutions of 1:100 and 1:1,000, respectively. M. Biggin, Lawrence Berkeley National Laboratory, Berkeley, CA) at dilution buffer (5 mM KCl and 0.1 mM sodium phosphate, pH 7.0). Newly heated to >90°C, which was allowed to cool for several hours. The dsRNA was purified by phenol/chloroform extraction and resuspended in injection buffer (5 mM KC1 and 0.1 mM sodium phosphate, pH 7.0). Newly laid embryos were injected laterally and incubated for 2.5–3 h at room temperature (23°C) before imaging gastrulation. We injected A2 dsRNA at a concentration of ~1 mg/ml. However, tears in the myosin II network could be observed even after diluting dsRNA to a lower concentration. For most experiments, Arm dsRNA was used at 0.2 mg/ml. For double RNAi experiments, Arm dsRNA was used at 0.2 mg/ml, and Twist and Snail dsRNA was used at 2 mg/ml. For triple RNAi experiments, Arm dsRNA was used at 0.2 mg/ml, and Fos and T48 were used at 0.8 mg/ml. For all experiments, Arm dsRNA was used at 0.2 mg/ml, and Twist and Snail dsRNA was used at 2 mg/ml. For triple RNAi experiments, Arm dsRNA was used at 0.2 mg/ml, and Fos and T48 were used at 0.8 mg/ml and 0.65 mg/ml, respectively.

Online supplemental material

Fig. S1 shows that knockdown of E-cadherin, Arm, or α-catenin disrupts the myosin II network and results in membrane tethers. Fig. S2 shows myosin-GFP displacement as a function of time for the two types of laser ablation experiments shown in Fig. 3. Video 1 shows myosin-GFP in wild-type and arm-MY22 mutant embryos. Video 2 shows laser ablations in myosin-GFP embryos. Video 3 shows myosin-ncCherry and membrane-GFP in an armRNAi embryo. Video 4 shows the contraction of a myosin II fiber in an arm-MY22 mutant embryo. Video 5 shows the unrestrained myosin II contraction that occurs after tearing in an armRNAi embryo. Video 6 shows membrane tether formation in a membrane-GFP embryo. Video 7 shows myosin-GFP in a double RNAi control arm-CG3651RNAi embryo. Video 8 shows the suppression of tears and loss of cell–cell adhesion in a myosin-GFP arm-snaRNAi embryo. Video 9 shows that contraction pulses continue despite loss of cell–cell adhesion in a myosin-GFP arm-twiRNAi embryo. Video 10 shows the suppression of epithelial tears in a membrane-GFP arm-twiRNAi embryo. Online supplemental materials is available at http://www.jcb.org/cgi/content/full/jcb.200910099/DC1.

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References


Figure S1. Knockdown of E-cadherin or α-catenin disrupts the myosin II network and results in membrane tethers. (A) RT-PCR of ECad2, arm, and α-catenin transcripts after RNAi. The percentage of knockdown of the specific transcript was measured to be 60% for ECad2RNAi and 90–95% for armRNAi and α-cateninRNAi. (B) Myosin-GFP in live ECad2RNAi and α-cateninRNAi embryos. Bottom images are 5 min later than top images. Note the dramatic tears along the α-p axis and the contraction of myosin II into rings. Bar, 10 µm. (C) Myosin II rings in armRNAi embryos are enriched in the remaining cortical E-cadherin. Images represent fixed myosin-GFP armRNAi embryos stained for E-cadherin. Bar, 5 µm. (D) Membrane tethers contain F-actin but not myosin II. Fixed myosin-GFP (top) or membrane-GFP (Spider-GFP, bottom) armRNAi embryos were stained for F-actin. Arrowheads indicate membrane tethers that appear to contain F-actin. Bar, 5 µm. (E) Apical membrane (Spider-GFP) in live ECad2RNAi and α-cateninRNAi embryos. Membrane tethers connect dense membrane accumulations, similar to armRNAi embryos. Bar, 5 µm. (F) Scanning EM of armRNAi and arm-twiRNAi embryos. In arm-twiRNAi embryos, cells dissociate and continue to form numerous blebs and tethers. Bars: (B) 10 µm; [C–F] 5 µm.
Figure S2. **Myosin-GFP displacement as a function of time for different laser ablation experiments.** Individual curves represent the displacement of individual myosin II structures that retracted various distances after laser ablation. Gray box indicates the duration of the ablation. Arrows indicate the time when displacements were measured for the data in Fig. 3. Line ablations (left) lasted ~6.4 s, and thus the tissue relaxed farther before the first post-ablation image. Point ablations (right) lasted 0.67 s, providing a better estimate for the initial retraction velocity.

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**Video 1.** Maximum intensity z projections of myosin-GFP during ventral furrow formation in wild-type and arm\(^{M/Z}\) mutant *Drosophila* embryos. Images were analyzed by time-lapse confocal microscopy using a laser-scanning confocal microscope (SP5; Leica). Image stacks were taken every 12 s. The video speed is 120-fold faster than real time (10 frames/second). Related still images are shown in Fig. 2 A.

**Video 2.** Maximum intensity z projections of myosin-GFP in *Drosophila* embryos where a laser incision was made in the myosin II meshwork. The white bars in the first frame illustrate the location of the incision made in each embryo. Images were analyzed by time-lapse confocal microscopy using a spinning-disk confocal microscope (Ultraview; PerkinElmer). Image stacks were taken every 3 s. The video speed is 30-fold faster than real time (10 frames/second). Related still images are shown in Fig. 3 A.

**Video 3.** Maximum intensity z projection of myosin-mCherry (top) and single Z-slice of membrane-GFP below the apical surface (bottom, to illustrate cell outlines) in an armRNAi *Drosophila* embryo. Images were analyzed by time-lapse confocal microscopy using a laser-scanning confocal microscope (SP5; Leica). Image stacks were taken every 19.5 s. The video speed is 120-fold faster than real time (6.15 frames/second).

**Video 4.** Maximum intensity z projection of myosin-GFP in an arm\(^{M/Z}\) mutant *Drosophila* embryo. Myosin fiber contracts upon separation of the actomyosin meshwork. Images were analyzed by time-lapse confocal microscopy using a spinning-disk confocal microscope (Ultraview; PerkinElmer). Image stacks were taken every 2.4 s. The video speed is 10-fold faster than real time (4.18 frames/second). Related still images are shown in Fig. 2 C (i).
Video 5. **Maximum intensity z projection of myosin-GFP in an armRNAi Drosophila embryo.** Tears in the meshwork are associated with myosin contraction into rings or balls and constitutive flow of myosin spots into these rings. Images were analyzed by time-lapse confocal microscopy using a spinning-disk confocal microscope (Ultraview; PerkinElmer). Image stacks were taken every 2.4 s. The video speed is 30-fold faster than real time (12.66 frames/second). Related still images are shown in Fig. 2 C (iii).

Video 6. **Membrane tether formation in a membrane-GFP (Spider) Drosophila embryo.** Time-lapse images were taken of a single z slice on the apical surface of an armRNAi embryo. Images were analyzed by time-lapse confocal microscopy using a laser-scanning confocal microscope (SP5; Leica). Image stacks were taken every 6 s. The video speed is 120-fold faster than real time (20 frames/second).

Video 7. **Maximum intensity z projection of myosin-GFP in an arm-CG3651RNAi Drosophila embryo.** Images were analyzed by time-lapse confocal microscopy using a spinning-disk confocal microscope (Ultraview; PerkinElmer). Image stacks were taken every 2.75 s. The video speed is 60-fold faster than real time (21.82 frames/second). A related still image is shown in Fig. 7 A.

Video 8. **Maximum intensity z projection of myosin-GFP in an arm-snaRNAi Drosophila embryo.** snaRNAi inhibits loss of cell adhesion and epithelial tears. Cells only round up when they eventually enter mitosis. Images were analyzed by time-lapse confocal microscopy using a spinning-disk confocal microscope (Ultraview; PerkinElmer). Image stacks were taken every 2.34 s. The video speed is 60-fold faster than real time (25.64 frames/second). A related still image is shown in Fig. 7 A.

Video 9. **Maximum intensity z projection of myosin-GFP in an arm-twiRNAi Drosophila embryo.** twiRNAi does not prevent loss of adhesion, but inhibits the formation of discrete epithelial tears. Note that contraction pulses occur even when cells appear to lose adhesion. Images were analyzed by time-lapse confocal microscopy using a spinning-disk confocal microscope (Ultraview; PerkinElmer). Image stacks were taken every 2.75 s. The video speed is 60-fold faster than real time (21.82 frames/second). Related still image is shown in Fig. 7 A.

Video 10. **Single z slice of membrane-GFP (Spider) in arm-twiRNAi Drosophila embryo.** Note that twiRNAi inhibits the formation of discrete tears as cells lose adhesion. Images were analyzed by time-lapse confocal microscopy using a laser-scanning confocal microscope (SP5; Leica). Image stacks were taken every 11 s. The video speed is 60-fold faster than real time (5.5 frames/second). A related still image is shown in Fig. 7 B.