STRESS-INDUCED ALIGNMENT OF ACTIN FILAMENTS AND THE MECHANICS OF CYTOGEL

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Experimental evidence suggests that anisotropic stress induces alignment of intracellular actin filaments. We develop a model for this phenomenon, which includes a parameter reflecting the sensitivity of the microfilament network to changes in the stress field. When applied to a uniform cell sheet at rest, the model predicts that for sufficiently large values of the sensitivity parameter, all the actin filaments will spontaneously align in a single direction. Stress alignment can also be caused by a change in external conditions, and as an example of this we apply our model to the initial response of embryonic epidermis to wounding. Our solutions in this case are able to reflect the actin cable that has been found at the wound edge in recent experiments; the cable consists of microfilaments aligned with stress at the wound boundary of the epithelium. These applications suggest that stress-induced alignment of actin filaments could play a key role in some biological systems. This is the first attempt to include the alignment phenomenon in a mechanical model of cytogel.

1. Biological background. The mechanical balance of forces in living tissue has formed the basis for many recent models of developmental morphogenesis. This approach was introduced by Odell et al. (1981), who developed a discrete model for the folding of embryonic epithelia. Building on this work, G. F. Oster, J. D. Murray and their colleagues (Oster et al., 1983; Murray et al., 1983; Murray and Oster, 1984a,b; Oster et al., 1985a,b; Murray et al., 1988) developed a series of continuum mechanochemical models for pattern formation in a wide range of embryonic systems. In the Oster–Murray approach, cytogel is treated as an isotropic viscoelastic continuum, and the non-viscous response to applied strain is the same as for, say, a simple rubber sheet, except for the exertion of an isotropic contraction stress. Experimentally,
such traction forces have been observed in a number of cell types (see Harris, 1982; and refs in Stopak et al., 1985). In the model equations, this traction stress is analogous to a "negative pressure", and is crucial to the predictions of the various models.

In reality, of course, there are other differences between the non-viscous response to applied strain of simple non-living materials, such as a rubber sheet, and of cytogel. In this paper, we focus on one such difference, namely the alignment of actin filaments with the stress field, which is illustrated schematically in Fig. 1. Intracellular actin filaments determine many of the mechanical properties of a confluent cell sheet (Pollard, 1990), and the alignment phenomenon has been recognized but neglected in previous applications of the Oster–Murray approach (for example, Oster and Odell, 1984a). Here we develop a new model which reflects stress alignment, and discuss two simple applications, which highlight the importance of the phenomenon in some biological systems.

We begin by briefly reviewing the experimental evidence for stress-induced alignment of actin filaments. The seminal work is by Kolega (1986). He studied the cytoskeleton of fish epidermal cells held under tension either by micromanipulation or by cellular locomotive activity, and found that the actin filaments were aligned parallel to the tension. Moreover, in the case of micromanipulation, filament alignment occurred within a few minutes, as a direct response to applied tension.

Although not as clear as Kolega's (1986) study, previous authors had found a similar response to applied tension. Chen (1981) investigated the retraction of fibroblast tails following detachment from the substratum. In common with previous authors (for example Spooner et al., 1971; Ludueña and Wessells, 1973) he found that a taut elongated tail contained bundles of microfilaments extending parallel to the long axis, and thus aligned with the stress field. Upon detachment, these microfilament bundles lost their orientation and returned to a random mesh-like arrangement; this conversion took about 15 sec. Valberg
and Albertini (1985) used a magnetic particle method to study the effects of applied tension on pulmonary macrophages from the lungs of hamsters. They caused the macrophages to ingest magnetic particles, which then entered phagosome organelles, and they applied external magnetic fields. This resulted in positive staining for actin coinciding with the location of the magnetic particles, which again suggests stress-induced microfilament alignment. Finally, a number of authors have studied the alignment of actin filament bundles in endothelial cells under shear stress. This is an important feature of blood flow, and occurs in both isolated cells and in confluent cell sheets (see, for example, Franke et al., 1984; Wechezak et al., 1989).

2. A Model for Filament Alignment. The detailed biological mechanisms by which an applied stress field induces actin filament alignment are largely unknown. We therefore base our model on simple, intuitively plausible assumptions. Most importantly, we assume that alignment occurs as a direct response to the ratio of the principal components of stress. This implies that multiplying the stress tensor by an isotropic factor will have no effect on the actin filament network; in reality, there may be an increase in the proportion of actin that is polymerized (Pender and McCulloch, 1991), but here for simplicity we take any such increase to be negligible. We will denote the principal axes of the stress tensor \( \sigma \) as the 1- and 2-axes, with corresponding principal values \( \sigma_1 \) and \( \sigma_2 \). These axes are orthogonal since \( \sigma \) will always be symmetric. With this notation, we represent the microfilament network via the density function \( F(\phi; \rho) \), defined by the condition that \( F(\phi; \rho) \delta \phi \) is the continuum average of the proportion of the actin filaments inclined to the 1-axis at angles in the range \( (\phi, \phi + \delta \phi) \), when \( \sigma_1/\sigma_2 = \rho \). Thus \( F \) is also a function of space, but since all our calculations will be at a single point in space, we do not include \( r \) as an argument of \( F \), for notational simplicity. The symmetries of the system are such that \( F(-\phi, \rho) \) and \( F(\phi + \pi, \rho) \) must both equal \( F(\phi, \rho) \), and we can therefore restrict our attention to values of \( \phi \) in \([0, \pi/2]\). We require \( F \) to satisfy the following conditions.

(a) \( F(\phi; \rho) \rightarrow \delta(\pi/2 - \phi) \) as \( \rho \rightarrow 0 \) and \( F(\phi; \rho) \rightarrow \delta(\phi) \) as \( \rho \rightarrow \infty \), where \( \delta \) denotes the Dirac delta function. That is, when the stress field is unidirectional, all filaments are oriented in that direction.

(b) \( \int_{0}^{\pi/2} F(\phi; \rho) \, d\phi = 1 \), since \( F \) is a density function. Thus the total number of microfilaments is unchanged by the process of stress alignment: as discussed above, we assume that changes in the stress field do not affect the extent of actin polymerization.

(c) \( F(\phi; 1) \) is a constant. That is, when the stress field is isotropic the filaments are oriented randomly. Condition (b) implies that the value of this constant is \( 2/\pi \).
(d) \( F(\phi; \rho) = F(\pi/2 - \phi; 1/\rho) \), by symmetry.

These conditions do not determine \( F \), but they are fairly restrictive. In particular, the definition of the Dirac delta function as \( \delta(x) = \lim_{\varepsilon \to 0}(1/\varepsilon \sqrt{\pi}) \exp[-x^2/\varepsilon^2] \) suggests the following functional form:

\[
F(\phi; \rho) = F_0(\rho) \left[ \frac{1}{\psi(\rho)} \exp\left\{- \frac{k(\pi/2 - \phi)^2}{\psi(\rho)^2}\right\} + \psi(\rho) \exp\{- k\psi(\rho)^2 \phi^2\} \right]
\]

(1)

where \( k \) is a positive constant and \( \psi(0) = 0, \psi(1) = 1 \) and \( \psi(\infty) = \infty \). We choose \( F_0(\rho) \) such that condition (b) is satisfied. If we take \( \psi(\rho) = \rho^p \) for some \( p > 0 \) then condition (d) is also satisfied.

It remains to consider condition (c), that \( F(\phi; 1) \) is independent of \( \phi \). This does not hold exactly for any non-zero value of \( k \), but it is approximately true for values of \( k \) close to 1. We choose \( k \) to minimize the variation of \( F(\phi; 1) \) on \([0, \pi/2]\), and we show in the Appendix that this gives \( k \approx 0.988 \), and reduces the variation to about 1.4%. The form of \( F(\phi; \rho) \) with this value of \( k \) is plotted in Fig. 2 for a range of values of \( \rho^p \). The parameter \( p \) remains undetermined, and reflects the sensitivity of the microfilament network to changes in the stress field.

![Figure 2](image-url)

Figure 2. The form of \( F(\phi; \rho) \), defined in (1), plotted as a function of \( \phi \), for \( \rho^p = 0.01, 0.05, 0.1, 0.2, 0.3, 0.5, 0.75, 1.0 \); \( F(0; \rho) \) increases with \( \rho^p \). It is sufficient to consider \( \rho \leq 1 \), since \( F(\phi; 1/\rho) = F(\pi/2 - \phi; \rho) \).

We denote by \( G_0(r) \) the total actin filament density at position \( r \). We assume that each actin filament generates/supports a force along its length; thus the stress exerted by the mesh of actin filaments is determined by the "actin filament density components" along the principal axes of the stress tensor, which are related to \( F \) as follows:
\[ G_1(r) = G_0(r) \int_0^{\pi/2} F(\phi; \rho(r)) \cos \phi \, d\phi \]
\[ G_2(r) = G_0(r) \int_0^{\pi/2} F(\phi; \rho(r)) \sin \phi \, d\phi = G_0(r) \int_0^{\pi/2} F(\phi; 1/\rho(r)) \cos \phi \, d\phi \]

using the symmetry condition (d). Note that \( G_1 \) and \( G_2 \) do not depend on the full form of \( F(\phi; \rho) \); rather, they depend only on the integral:

\[ \int_0^{\pi/2} F(\phi; \rho) \cos \phi \, d\phi \equiv f(\rho), \text{ say.} \] (2)

Condition (a) implies that \( f(0) = 0 \) and \( f(\infty) = \pi/2 \), while condition (b) gives \( f(1) = 1 \). Further, \( f \) is a strictly increasing function of \( \rho_p \). Motivated by these properties, we found that to a very good approximation:

\[ f(\rho) \approx \frac{(\pi/2)\rho_p}{\rho_p^2 + (\pi/2 - 1)} \]

as illustrated in Fig. 3. Thus we represent actin filament alignment by the relations:

\[ G_1(r) = \frac{(\pi/2)\sigma_1^p G_0(r)}{\sigma_1^p + (\pi/2 - 1)\sigma_2^p} \] (3a)
\[ G_2(r) = \frac{(\pi/2)\sigma_2^p G_0(r)}{\sigma_2^p + (\pi/2 - 1)\sigma_1^p}. \] (3b)

Under certain conditions, microfilament alignment can occur spontaneously, as a result of the stress exerted by the filaments themselves; more generally, it can be caused by a change in external conditions, for example at the boundary of a cell sheet. We now consider an example of each of these two types of alignment.

3. Applications

3.1. Auto-alignment of actin filaments. Actin filament alignment occurs in response to the local stress field, and in the absence of external influences, these stress forces are exerted by the actin filaments themselves. We thus expect intuitively that auto-induction of filament alignment will destabilize a random filament mesh-work if alignment depends sufficiently steeply on stress anisotropy. We now use our model (3) to investigate this phenomenon. Our stability calculation assumes that the two regulatory processes that are reflected in (3) have different time scales. The actin-generated stress alters
almost instantaneously with changes in filament density, while the alignment of filaments takes about 10 sec or more to adjust to changes in the stress field (Kolega, 1986; Chen, 1981).

We consider the very simple case of a cell sheet at equilibrium, with its boundaries held fixed and with no strain at any point. The sheet will then be under tension, even though the strain field is zero, because of the traction forces exerted by the cells. This traction stress will depend on the local actin filament density, and following previous authors (for example Oster, 1984; Oster et al., 1985a), we take this dependence to be a simple proportionality. Thus:

\[ \sigma_1 = \tau_0 G_1 \quad \text{and} \quad \sigma_2 = \tau_0 G_2 \]  

where \( \tau_0 \) is a positive constant, and the 1- and 2-axes are in this case in arbitrary orthogonal directions. Writing \( \gamma = G_1 / G_2 \), these imply that \( \rho = \gamma \). As discussed above, this dependence of stress \( \rho \) on changes in alignment \( \gamma \) is essentially instantaneous, while the dependence of \( \gamma \) on changes in \( \rho \) that is implied by our equations (3) occurs on a longer time scale. Together, (3) and (4) give an equation for \( \gamma \), with multiple solutions. We now use the difference in time scale of the two responses to investigate the stability of these various equilibrium configurations of the microfilament network.

Suppose that \( \rho = \rho_e \) and \( \gamma = \gamma_e \) is a solution of (3) and (4), so that:

\[ \gamma_e = \rho_e = \frac{\rho_e^p [1 + (\pi/2 - 1) \rho_e^p]}{\rho_e^p + \pi/2 - 1}. \]
Then our assumption that (4) is satisfied instantaneously implies that if we perturb the alignment ratio to a new value $\gamma_e + \epsilon$, the stress ratio will simultaneously change to $\rho_e + \epsilon$. If this value of the stress ratio $\rho$ were maintained, our equations (3) imply that the actin filament density ratio would move to the value:

$$\gamma_{pert} = \frac{(\rho_e + \epsilon)^p [1 + (\pi/2 - 1)(\rho_e + \epsilon)^p]}{(\rho_e + \epsilon)^p + \pi/2 - 1}.$$ 

The initial direction of change of alignment ratio $\gamma$ is thus increasing if $\gamma_{pert} > \gamma_e + \epsilon$; and since $\rho$ follows $\gamma$ instantaneously, this entails an increase of $\rho$ also, that is an increase in the departure from equilibrium—in other words, an instability. Thus $(\rho_e, \gamma_e)$ will be stable to small perturbations if and only if:

$$\frac{d}{d\rho} \left\{ \frac{\rho^p [1 + (\pi/2 - 1)\rho^p]}{\rho^p + \pi/2 - 1} \right\}_{\rho = \rho_e} < 1.$$ 

since this derivative is clearly non-negative for all $\rho$. We now write

$$H(\rho) \equiv \frac{\rho^p [1 + (\pi/2 - 1)\rho^p]}{\rho^p + \pi/2 - 1} - \rho. \quad (5)$$

Then for a solution of our model we require $H(\rho) = 0$, and this solution is stable if and only if $H'(\rho) < 0$. In Fig. 4 we plot $H(\rho)$ for a number of values of $p$. We restrict attention to $0 \leqslant \rho \leqslant 1$, since our problem is symmetric in the 1- and 2-axes. There is an abrupt change in the form of $H$ as $p$ increases through 1, since:

$$H'(0) = \begin{cases} \infty & \text{for } 0 < p < 1 \\ (4 - \pi)/(\pi - 2) & \text{for } p = 1 \\ -1 & \text{for } p > 1. \end{cases}$$

The figure shows that the number of solutions and their stability depends on $p$:

for $0 < p \leqslant 1$: \quad $\rho = 0$ is an unstable solution \\
\hspace{1cm} $\rho = 1$ is a stable solution

for $1 < p < p_{\text{crit}}$: \quad $\rho = 0$ and $\rho = 1$ are both stable solutions \\
\hspace{1cm} There is also an unstable solution $\rho = \hat{\rho} \in (0, 1)$

for $p > p_{\text{crit}}$: \quad $\rho = 0$ is a stable solution \\
\hspace{1cm} $\rho = 1$ is an unstable solution.

Here $p_{\text{crit}}$ is defined by $H'(1) = 0$, which gives $p_{\text{crit}} = \frac{1}{2} [1 + 1/(\pi/2 - 1)] \approx 1.38$. In the case $1 < p < p_{\text{crit}}$, the two stable solutions $\rho = 0$ and $\rho = 1$ have basins of
attraction \([0, \hat{\rho}]\) and \((\hat{\rho}, 1]\), and as shown in Fig. 5, \(\hat{\rho}\) is extremely small for values of \(\rho\) as high as 1.1, so that for practical purposes \(\rho = 0\) is then unstable.

Recalling that \(\rho = \gamma = G_1/G_2\) at equilibrium, \(\rho = 0\) corresponds to unidirectional actin filaments, while \(\rho = 1\) corresponds to a random filament meshwork. Thus our model predicts that the configuration of the microfilament network in a cell sheet at rest will depend on the sensitivity of actin filament alignment to changes in the stress field. When this sensitivity is low, the network will have a random, mesh-like arrangement, but when it is sufficiently high, this random configuration will be driven unstable, and all the microfilaments will become spontaneously aligned in a single, random direction. Such spontaneous filament alignment could play an important role in morphogenetic processes such as convergent extension (Gerhart and Keller, 1986; Lewis et al., in preparation).

3.2. Boundary-induced filament alignment. Stress-induced alignment of actin will also have important effects in the responses of an epithelial sheet to externally applied forces and boundary conditions. This is well exemplified in
Figure 5. The value of the unstable solution $\hat{\rho} \in (0, 1)$ of $H(\rho) = 0$. This solution exists for $1 < p < p_{\text{crit}}$, but is very small when $p$ is close to 1, so that the anisotropic solution $\rho = 0$ of $H(\rho) = 0$ has a very small basin of attraction.

The case of wound healing, where a surgical cut creates an epidermal free edge, and the cellular response to wounding is to a large extent a response to this free edge. In the epidermis of chick and mouse embryos, Martin and Lewis (1991, 1992) have recently shown that wounding induces marked alignment of actin filaments, resulting in a thick cable of filamentous actin around almost all of the wound margin, localized within the leading row of basal cells. By contrast, wounding in adult epithelia induces increased proliferation and lamellipodial crawling (Winter, 1962, 1972; Clark, 1989; Sherratt and Murray, 1990, 1991, 1992). We now use our model (3) to investigate the initial formation of this actin cable, as an example in vivo microfilament alignment caused by a free boundary. In the epidermal cell sheet, the intracellular actin filaments are linked via cell–cell adherens junctions in a transcellular network, and following wounding this network undergoes rapid changes, forming a new quasi-equilibrium state in which an actin cable runs around the wound margin; this cable appears to act like a purse-string, causing the wound gradually to close. Here we restrict attention to the quasi-equilibrium configuration, with its prominent feature of an actin cable at the wound margin.

We take the microfilament network as two-dimensional, since its thickness is much smaller than the typical wound dimensions of about 500 μm, and for simplicity, we consider the case of a circular wound. The principal axes of the stress tensor $\sigma$ will then be aligned at all points with the radial and tangential coordinate axes. In addition to the traction stress, we also have to take account of strain and elastic contributions to the stress, since the epithelium is displaced. At equilibrium, there will be no viscous contribution to $\sigma$, and we take its principal values to be given by:
\[
\begin{align*}
\sigma_{rr} &= G_r[\varepsilon_{rr} + \Gamma \nabla \cdot \mathbf{u}] + \frac{G_r}{\varepsilon_{rr}} \tau, \\
\sigma_{\theta\theta} &= G_\theta[\varepsilon_{\theta\theta} + \Gamma \nabla \cdot \mathbf{u}] + \frac{G_\theta}{\varepsilon_{\theta\theta}} \tau.
\end{align*}
\]

Here \(G_r(r)\) and \(G_\theta(r)\) are the radial and tangential "components" of actin filament density at the material point initially at \(r\), \(\mathbf{u}(r)\) is the displacement of that point from its pre-wounding position, \(\varepsilon_{rr}\) and \(\varepsilon_{\theta\theta}\) are the principal values of the strain tensor \(\varepsilon = \frac{1}{2}(\nabla \mathbf{u} + \nabla \mathbf{u}^T)\), \(\tau\) is the traction stress per actin filament, and \(E\) and \(\Gamma\) are positive constants. The form (6) is the standard representation of stress at equilibrium in the Oster–Murray approach (see in particular the model of Murray and Oster, 1984a, for epithelial morphogenesis), although in previous applications, \(G_r = G_\theta\). Many authors have taken the traction stress per actin filament, \(\tau\), to be an increasing function of the local compaction, since as the microfilament network is compressed, the degree of filament overlap increases and additional myosin cross-bridges form (Oster, 1984; Oster and Odell, 1984a,b; Oster et al., 1985a). For generality, we therefore take \(\tau = \tau_0/(1 - \beta \Omega)\), where \(\tau_0 > 0\) and \(\beta \geq 0\) are constants, and \(\Omega\) is the fraction of its pre-wounding area by which a small region of cytoplasm contracts in response to wounding. In a radially symmetric circular geometry, \(\Omega = 1 - (1 + u')/(1 + u/r)\), where \(r = |\mathbf{r}|\), \(u(r) = u(r)\mathbf{e}_r\) and prime denotes \(d/dr\).

The models (3) and (6) together give implicit equations for stress as a function of strain:

\[
\begin{align*}
\sigma_{rr} &= \frac{(\pi/2) \sigma_{rr}^P G_0}{\sigma_{rr}^P + (\pi/2 - 1) \sigma_{\theta\theta}^P} \cdot \{Eu' + \Gamma (u' + u/r) + \tau_0/[1 + \beta (u' + u/r + uu'/r)]\}, \\
\sigma_{\theta\theta} &= \frac{(\pi/2) \sigma_{\theta\theta}^P G_0}{\sigma_{\theta\theta}^P + (\pi/2 - 1) \sigma_{rr}^P} \cdot \{Eu/r + \Gamma (u' + u/r) + \tau_0/[1 + \beta (u' + u/r + uu'/r)]\}.
\end{align*}
\]

At equilibrium, these elastic and traction stresses balance the elastic restoring forces due to attachment to the substratum. Following Murray and Oster (1984a), we represent the restoring forces by \(\lambda G_0 \mathbf{u}\), where the positive constant \(\lambda\) reflects the strength of the attachments. The dependence on \(G_0\) arises because we assume that the attachments are fixed in the basal lamina and that this becomes wrinkled (so that the attachments are compressed) in response to wounding, to an extent reflecting the compaction of the cell sheet. Experimental justification for this dependence is given by Hergott et al. (1989), who find that the density of focal contacts increases in parallel to that of actin filament bundles in the response of chick embryonic corneal cells to wounding in an organ culture system.

Thus the equation to be solved for the new equilibrium configuration is:
\[ \nabla \cdot \sigma = \lambda G_0 u \]  
\[ (8) \]

with \( \sigma \) given by (7). For general values of \( p \), (7) cannot be solved analytically to give expressions for \( \sigma_{rr} \) and \( \sigma_{\theta\theta} \) as functions of \( u \). We therefore restrict attention to the case \( p = 1 \). We have shown in the previous section that there will then be no spontaneous microfilament alignment prior to wounding. Therefore, solutions of (7) in which one of the principal values of stress is identically zero are not relevant, since we require both \( G_r \) and \( G_\theta \) to be non-zero at points far from the wound. The solutions for \( \sigma_{rr} \) and \( \sigma_{\theta\theta} \) in the case \( p = 1 \) are thus:

\[ \sigma_{rr} = G_0 \left\{ \frac{E(2u'-(\pi-2)u/r)}{4-\pi} + \Gamma(u'+u/r) + \frac{\tau_0}{1+\beta(u'+u/r+uu'/r)} \right\} \]  
\[ (9a) \]

\[ \sigma_{\theta\theta} = G_0 \left\{ \frac{E(2u/r-(\pi-2)u')}{4-\pi} + \Gamma(u'+u/r) + \frac{\tau_0}{1+\beta(u'+u/r+uu'/r)} \right\}. \]  
\[ (9b) \]

It remains to consider the relationship between \( G_0 \) and \( u \). The results we present here are based on the assumption that the total amount of filamentous actin within each cell remains constant during the initial response to wounding. Since we are treating the cell sheet as a continuum, the mathematical formulation of this assumption is \( G_0(1-\Omega) = \kappa \), a spatially homogeneous constant; recall that \( \Omega = 1 - (1+u') \) is the local compaction. The implications of assuming more localized conservation of filamentous actin are presented elsewhere (Sherratt, 1992). To clarify the roles of the various model parameters, we nondimensionalize (8) by defining:

\[ r^* = r/R \quad u^* = u/R \quad E^* = E/\tau_0 \quad \Gamma^* = \Gamma/\tau_0 \quad \lambda^* = \lambda R^2/\tau_0 \]

where * denotes a dimensionless quantity, and \( R \) is the initial wound radius. Substituting these rescalings into (8) and (9), and dropping the asterisks for notational simplicity, gives a highly nonlinear second order ordinary differential equation for \( u(r) \):

\[ u'' = \frac{1}{[2E/(4-\pi)+\Psi](r+u)+\Gamma(r-u^2/r)} \left\{ \lambda ru \left[ 1+u'+u/r + uu'/r \right] \right\} \]

\[ - \left( u' - \frac{u}{r} \right) \left[ \frac{2E}{4-\pi} \left( 1 + \frac{u}{r} - u^2 + \frac{uu'}{r} \right) + \Gamma(1-u^2) + \Psi(1+u') \right] \]  
\[ (10) \]

where

\[ \Psi = \frac{(\pi-2)Eu}{(4-\pi)r} - \frac{1+\beta+2\beta(u'+u/r+uu'/r)}{[1+\beta(u'+u/r+uu'/r)]^2}. \]
We treat the cell sheet as infinite, which is valid provided the wound area is small compared to the whole sheet. The biologically appropriate boundary conditions are then $\sigma_{rr} = 0$ at $r = 1$ (the wound edge) and $u = 0$ at $r = \infty$.

Within the context of our continuum approximation across the cell sheet, the actin cable observed experimentally should correspond to a sharp increase in the calculated density of tangentially aligned actin filaments near the wound edge. This indeed is our finding in the model solutions for a wide range of parameter values: a typical solution is shown in Fig. 6. Here we plot $G_r$ and $G_\theta$ as functions of $r$; (2.3) and (3.9) together imply that:

\begin{align}
G_r &= G_0 \frac{\Gamma(u' + u/r) + 1/[1 - \beta \Omega] + E[2u' - (\pi - 2)u/r]/(4 - \pi)}{\Gamma(u' + u/r) + 1/[1 - \beta \Omega] + Eu'} \\
G_\theta &= G_0 \frac{\Gamma(u' + u/r) + 1/[1 - \beta \Omega] + E[2u/r - (\pi - 2)u]/(4 - \pi)}{\Gamma(u' + u/r) + 1/[1 - \beta \Omega] + Eu/r}
\end{align}

(11a) (11b)

Figure 6. The effective densities of radially and tangentially aligned actin filaments, $G_r$ and $G_\theta$, defined in (11). The solution for $u(r)$ is obtained by solving the model equation (10) numerically using a finite difference scheme and Newton’s method, with deferred correction and continuation in $\beta$ (Pereyra, 1979). The dimensionless parameter values are $E = 0.5$, $\Gamma = 0.8$, $\lambda = 3$, $\beta = 0.485$. This solution exhibits a sharp increase in the density of tangentially aligned actin near the wound edge, and thus reflects the phenomenon of the actin cable found experimentally by Martin and Lewis (1991, 1992).

The corresponding model without filament alignment can be obtained by setting the parameter $p = 0$ (recall that the above solutions are for $p = 1$). We anticipate that the absence of stress-induced alignment of microfilaments parallel to the wound edge will increase the radial traction stress exerted by aggregated actin near this edge, and thus reduce the ability of the restraining elastic stress to hold the epidermal sheet in mechanical equilibrium. This is
confirmed by the model solutions: for the parameter values used in Fig. 6, a solution of the model does not exist in the case \( p = 0 \) (Sherratt, 1992). Model solutions do exist in this case for larger values of the elasticity parameters \( E \) and \( \Gamma \), although the parameter domain in which a solution exists is considerably smaller than in the case \( p = 1 \). Moreover, for appropriate parameters, these solutions do imply an aggregation of actin density \( G_0 \) near the wound edge; however, they are unable to capture the experimentally observed tangential alignment of this aggregated actin.

Biologically, our results suggest that the initial formation of the actin cable at the edge of embryonic epidermal wounds can be explained simply as a by-product of the post-wounding mechanical equilibrium in the epidermis, without ascribing any special properties, such as elevated levels of regulatory chemical or a phenotype change, to the cells at the wound edge. Such changes may occur (Pender and McCulloch, 1991), but our model suggests that they are not necessary for the formation of an actin cable.

4. Conclusions. Despite considerable biological evidence for stress-induced alignment of the cellular microfilament network, the phenomenon has been neglected in previous mechanical models for tissue deformation. We have developed a model for this effect, in which actin filaments align according to the local anisotropy of the stress field, and our formulation includes a parameter reflecting the sensitivity of this response. We have used the model to investigate the equilibrium state of the microfilament network in a cell sheet at rest, and the response of embryonic epidermis to wounding. In the former system, filament alignment can occur spontaneously, while in the latter it is induced by the free edge at the wound front. These applications suggest that stress-induced actin filament alignment could play a key role in some biological systems.

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APPENDIX

The value of \( k \). The definition (1) of \( F(\phi; \rho) \) does not exactly satisfy condition (c), that \( F(\phi; 1) \) is constant. In this Appendix, we show that the condition can be satisfied to a good approximation for appropriate values of \( k \). Specifically, we look for the value of \( k \), not near zero, that minimizes the variation of:

\[
g(\phi) = \exp[-k\phi^2] + \exp[-k(\pi/2 - \phi)^2] \equiv F(\phi; 1)/F_0(1)
\]

(A1)
on $0 \leq \phi \leq \pi/2$. [Trivially, $g(\phi) \equiv 2$ when $k = 0$, but this case is not biologically relevant.] The form of $g(\phi)$ for a selection of values of $k$ is shown in Fig. A1. Differentiating equation (A1) with respect to $\phi$ gives:

$$
\frac{dg}{d\phi} = 2ke^{-k\phi^2}(\pi/2 - \phi) \left[ \exp\{k\pi(\phi - \pi/4)\} - \frac{\phi}{\pi/2 - \phi} \right]
$$

$$
= 0 \Leftrightarrow \exp\{k\pi(\phi - \pi/4)\} = \frac{\phi}{\pi/2 - \phi}.
$$

(A2)

![Graphs showing the function $g(\phi)$ for different values of $k$.]

Figure A1. The form of the function $g(\phi)$, defined in equation (A1), as the parameter $k$ varies. (a) $k = 0.7$; (b) $k = 0.93$; (c) $k = k_{\text{crit2}} \approx 0.988$, defined in (A3); (d) $k = 1.1$. The dashed line indicates the equal values of $g(0)$ and $g(\pi/2)$.

It is sufficient to consider this equation on $[0, \pi/4]$, by the symmetry of $g(\phi)$ about $\phi = \pi/4$. The two sides of (A2) are plotted in Fig. A2 for a range of values of $k$. From this, it is clear that in addition to the root at $\phi = \pi/4$, equation (A2) has a second root in $(0, \pi/4)$ if $k > k_{\text{crit1}}$, where $k_{\text{crit1}}$ satisfies:

$$
\frac{d}{d\phi} \left[ \exp\{k\pi(\phi - \pi/4)\} - \frac{\phi}{\pi/2 - \phi} \right]_{\phi = \pi/4} = 0.
$$

This uniquely determines $k_{\text{crit1}}$ as $8/\pi^2$, $\approx 0.81$. 
Figure A2. The right-hand (full curve) and left-hand (dashed curves) sides of equation (A2) for $k = 0.2, 0.4, 0.6, \ldots, 2.0$; the value of the left-hand side decreases, for any given $\phi$, as $k$ increases. The number of solutions of the equation on $[0, \pi/4]$ increases from one to two as $k$ increases through the critical value $k_{\text{crit}1}$.

We now write:

$$g_{\text{max}}(k) = \max_{0 \leq \phi \leq \pi/2} g(\phi) \quad \text{and} \quad g_{\text{min}}(k) = \min_{0 \leq \phi \leq \pi/2} g(\phi).$$

By symmetry these are also the maximum and minimum values, respectively, on $[0, \pi/4]$. Then for $k < k_{\text{crit}1}$:

$$g_{\text{max}}(k) = g(\pi/4) = 2e^{-k\pi^2/16}$$

and

$$g_{\text{min}}(k) = g(0) = 1 + e^{-k\pi^2/4}.$$  

For $k > k_{\text{crit}1}$, $g_{\text{max}}(k) = g(\phi_{\text{max}})$, where $\phi_{\text{max}}$ is the unique solution of:

$$\exp\{k\pi(\phi - \pi/4)\} = \frac{\phi}{\pi/2 - \phi}$$

in the open interval $(0, \pi/4)$. This cannot be found analytically, but can easily be calculated numerically for a given value of $k$. Also for $k > k_{\text{crit}1}$, $g_{\text{min}}(k) = \min\{g(0), g(\pi/4)\}$. It is clear from Fig. A1 that there is a critical value $k_{\text{crit}2}$ such that:

for $k_{\text{crit}1} < k < k_{\text{crit}2}$, $g(0) < g(\pi/4)$, so that $g_{\text{min}}(k) = 1 + e^{-k\pi^2/4}$

and for $k > k_{\text{crit}2}$, $g(0) > g(\pi/4)$, so that $g_{\text{min}}(k) = 2e^{-k\pi^2/16}$.

Here $k_{\text{crit}2}$ satisfies $g(0) = g(\pi/4)$, that is:

$$1 + e^{-k\pi^2/4} = 2e^{-k\pi^2/16}.$$  

Thus $k_{\text{crit}2} = -(16/\pi^2)\log x$, where $x$ is a solution of $x^4 - 2x + 1 = 0$. Now:

$$x^4 - 2x + 1 = (x - 1)(x^3 + x^2 + x - 1).$$
The root $x=1$ corresponds to the trivial case of $g(\phi)=2$ when $k=0$. Substituting $x = \frac{1}{3}(2\sqrt{2} \sinh x - 1)$ in the cubic equation shows that this has a unique solution, which gives:

$$k_{crit2} = -\frac{16}{\pi^2} \log \left\{ \frac{2\sqrt{2}}{3} \sinh \left[ \frac{1}{3} \sinh^{-1} \left( \frac{117}{2\sqrt{2}} \right) \right] - \frac{1}{3} \right\} \approx 0.987886. \quad (A3)$$

Figure A3. The maximum and minimum values of $g(\phi)$ on $[0, \pi/2]$, $g_{max}(k)$ and $g_{min}(k)$, plotted as functions of the parameter $k$. The vertical lines $k=k_{crit1}$ and $k=k_{crit2}$ are also drawn. Trivially, $g(\phi)=2$ when $k=0$, but for values of $k$ away from zero, the variation of $g(\phi)$ is minimized in both an absolute and relative sense when $k=k_{crit2}$.

In Fig. A3 we plot $g_{max}(k)$ and $g_{min}(k)$ as functions of $k$. These suggest that the variation of $g(\phi)$ is minimized when $k=k_{crit2}$, and this is confirmed by numerical minimization using quadratic interpolation. Specifically, numerical calculations show that both $[g_{max}(k) - g_{min}(k)]$ and $[g_{max}(k) - g_{min}(k)]/[g_{max}(k) + g_{min}(k)]$ are minimized at $k=k_{crit2}$, for $k$ not near zero. [Trivially, $g_{max}(k)=g_{min}(k)$ when $k=0$, since then $g(\phi)=2$.] Thus the variation of $g(\phi)$ on $0 \leq \phi \leq \pi/2$ is minimized in both an absolute and relative sense when $k=k_{crit2}$. For this value of $k$, the variation of $g$ on $[0, \pi/2]$ is about 1.4%.

**LITERATURE**


Harris, A. K. 1982. Traction and its relations to contraction in tissue cell locomotion. In *Cell*


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