

Cell differentiation through tissue elasticity-coupled, myosin-driven remodeling

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Cells may lack eyes to see and ears to hear, but cells do seem to have a sense of ‘touch’ that allows them to feel their microenvironment. This is achieved in part through contractility coupled adhesion to physically flexible ‘soft’ tissue. Here we summarize some of the known variations in elasticity of solid tissue and review some of the long-term effects of cells ‘feeling’ this elasticity, focusing on differentiation processes of both committed cell types and stem cells. We then highlight what is known of molecular remodeling in cells under stress on short time scales. Key roles for forces generated by ubiquitous and essential myosin-II motors in feedback remodeling are emphasized throughout.

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Introduction

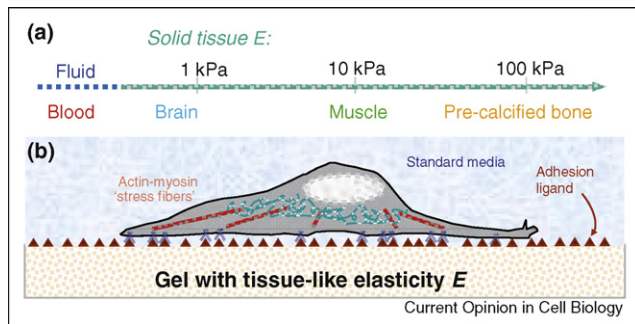
Solid tissues are soft, with the exception of bone, and they possess an elasticity E that not only varies between many different tissues (Figure 1a) but also exerts a considerable influence on how resident tissue cells behave. Indeed, cells that are dissociated from a solid tissue are generally not viable in a fluid suspension — they must adhere to a ‘solid’ which, by definition, recovers its shape after pushing and pulling, even at the scale of a cell. Solid tissues such as skin, muscle, and brain, are all relatively elastic, with the macroscale elasticity evident in their recovery of shape within seconds after mild poking and pinching or even after sustained compression, such as sitting. This is in contrast to fluid tissues such as blood and lymph which flow readily on a similar time scale and contain distinct cells, such as red and white blood cells, that are functional without sustained attachment. The impact of solid tissue elasticity on adherent cells is the focus here, with recent insights from stem cells and structural proteomics adding

to past reviews (e.g. [1]) of findings that indicate tissue elasticity E is felt by cells, affecting cell structure and function.

Both matrix composition and cell activity contribute to tissue elasticity or stiffness at a scale that cells can actively probe and sense. With collagen as an example: collagen type, amount, diameter of fibers, crosslinking (e.g. cellular lysyl oxidase activity), plus noncovalent interactions with other matrix proteins will all contribute to the matrix elasticity. Recent measurements of the elasticity of zebrafish embryos [2**] that were treated with the nonmuscle myosin-II (NMM II)-specific inhibitor blebbistatin also document a dramatic decrease in the effective elasticity, illustrating the contribution of myosin-derived tension (like tension in a guitar string) to the elasticity of the entire organism. While adherent tissue cells and extracellular matrix contribute to a characteristic if not strictly tissue-specific elastic microenvironment, cells generally anchor and pull on their surroundings through myosin-II-based contractility and transcellular adhesions of integrins plus other adhesion molecules [3]. The resistance felt by a given cell derives from tissue matrix, an adjacent cell, or perhaps — in culture — a synthetic substrate intended to model soft tissue (Figure 1b). Disease can bring significant changes in tissue elasticity: indeed, ‘sclerosis’ — as in atherosclerosis, otosclerosis, scleroderma, and more — is greek for hardening of tissue.

Contractile forces generated by ubiquitous crossbridging interactions of actin and myosin-II filaments in stress fibers are transmitted to the substrate as ‘traction’ forces that cause visible wrinkles in a thin film or lateral displacements of markers at the surface of a soft gel [4–7]. On gels with collagen-I covalently attached, epithelial cells and fibroblasts [8**] were the first cells reported to detect and respond distinctly to soft versus stiff substrates; differences were suggested to depend on myosin-II as they were inhibited by BDM (2,3-butanedione monoxime) — although this drug is now known to have multiple effects beyond myosin inhibition. Since then, neurons [9,10], muscle cells of various types [11–13], mesenchymal stem cells (MSCs) [14**], plus many other tissue cell types [15–18] have been shown to sense substrate stiffness, and at least some of the results have confirmed the importance of nonmuscle myosin-II through the inhibition of elasticity-dependent behavior changes with blebbistatin. Most cell types are found to respond to the elasticity E of the substrate within hours by spreading and assembling both adhesions and

Figure 1



Tissue elasticity scale (a) and model elastic culture systems (b).

cytoskeleton in proportion to E up to some saturating value beyond which changes in E exert no influence. Given that an isoform of myosin-II is also responsible for the work done by skeletal muscle, an analogy to lifting weights and exercise seems appropriate: to your bicep, a load of 1 kg undoubtedly feels very different from a load of 10 kg, whereas pushing or pulling on an immovable object like the handle of a locked door is a very distinct isometric exercise. Similar sensitivity to E seems to apply to most anchored cell types with similar implications for growth and remodeling within individual cells.

Soft tissue E measurements and model systems

The intrinsic resistance of a solid to a stress, regardless of topography and thickness (e.g. basement membrane), is measured by the solid's elastic modulus E , which is most simply obtained by applying a force — such as poking with an atomic force microscope (AFM) [19] — to a section of tissue or other substrate and then measuring the relative displacement. Tissues with small E show larger indentations or displacements under a given force.

E appears to adequately characterize many tissues not only at a macroscopic scale but also at the microscale of cell-exerted tractions — despite heterogeneity within tissue. Subtleties include matrix fiber effects and length scales of greatest relevance to cell sensing. Measurement issues include sample preparation; for example, E of whole brain in macroscopic measurements can vary by a factor of two or more depending on specifics of preparation, tissue perfusion, etc. [20]. In addition, many probing methods involve high frequency stressing [21] whereas relevant time scales for cell-exerted strains seem likely to range from seconds to hours with signal integration likely for longer time scales. Nonetheless, comparisons of three diverse tissues that contain a number of different and illustrative cell types show that brain tissue with $E \sim 0.1$ –1 kPa [20] is softer than striated muscle with $E \sim 10$ kPa [12,22], which is softer than osteoid with $E \sim 20$ –40 kPa [14^{••}]. Even though bone is about as rigid as

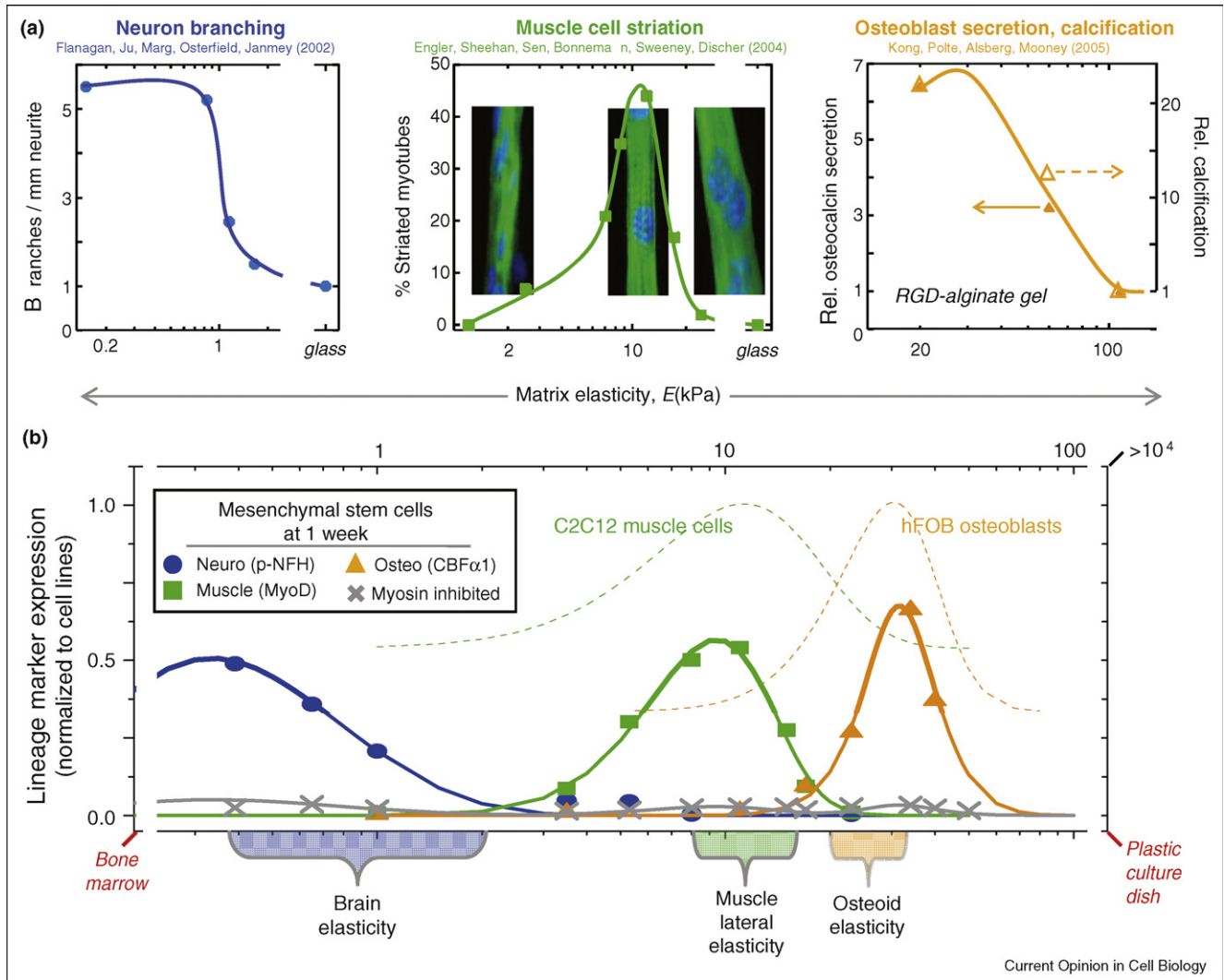
glass and plastic, it begins as compliant, precalcified, and heavily crosslinked collagenous 'osteoid' [14^{••}]. Fibrotic tissues and wounds have been measured to have E overlapping with this range [23,24]. Although the mapping of soft tissue microelasticity at a resolution typical in histology seems important and likely to be revealing, the implication from current studies is that there are distinct elastic microenvironments for neurons in brain, for myotubes in fiber bundles, and for osteoblasts on osteoid.

Tactile sensing of tissue stiffness requires a matrix ligand such as collagen for attachment, but once the ligand density is no longer limiting on a substrate [11], this cell-driven sensing feeds back on adhesion and cytoskeleton as well as on net contractile forces. Inert polyacrylamide gels with covalently attached collagen-I [8^{••}] have emerged as the most common system for controlling E in cell biology. By controlling the extent of polymer cross-linking in the gels, E can be adjusted over several orders of magnitude from extremely soft to stiff. Whether the collagen is fibrillar or not in such systems does not appear to have been scrutinized, but the collagen film does not contribute significantly to microelasticity [11] based on AFM measurements that also show E does not change even after many days of exposure to serum proteins — contrary to recent speculations [25].

Focal adhesion proteins paxillin [11,14^{••}] and vinculin [8^{••}] are found in small, diffuse, and dynamic adhesion complexes in cells on soft, lightly crosslinked gels ($E \sim 1$ kPa). In contrast, stiff, highly crosslinked gels ($E \gg 20$ kPa) show cells with stable focal adhesions, typical of those seen in cells attached to glass. Similarly, rigidification of cell-derived fibrillar matrices with glutaraldehyde shifts adhesions from fibrillar to large, nonfibrillar focal adhesions similar to those found on fixed substrates of fibronectin [26[•]]. Cytoskeletal assembly and cell tension [14^{••}] follow the same trends as adhesions. Thus the stiffer the matrix, the larger the adhesions and the stronger the cell pulls.

On a series of gel substrates, cell types that are committed to lineages such as neurons, muscle, and bone will modulate their phenotype in accordance with tissue elasticity (Figure 2a). Branching morphogenesis of neurons has been found to be maximal on soft matrices that mimic E_{brain} and minimal on both stiffer gels and glass [9,10]. Likewise, the striation of skeletal muscle myosin in muscle cells was maximal on stiffer matrices that mimic E_{muscle} , although myoblast fusion and expression levels of markers such as skeletal muscle myosin appeared otherwise unaffected [12]. Similar results were also obtained with multilayer cultures in which a first layer of cells grown on rigid glass showed no striation whereas an upper layer of cells that feel the soft bottom layer of cells showed strong striation. Lineage-specific secretions of osteoblasts [27[•]] also showed a relative

Figure 2



Matrix elasticity effects on differentiation of committed cells (a) and mesenchymal stem cells (b).

maximum near E_{osteoid} , suggesting that rigid bone is derived from remodeling of a suitably nonrigid matrix, although only three values of E were examined. The results of the osteoblast study were obtained with the noncollagenous ligand RGD attached to alginate gels rather than polyacrylamide. In addition, the elasticity dependence of the fibroblast–myofibroblast transition was demonstrated using a nonhydrous rubber rather than a hydrated gel-like polyacrylamide [28]. These latter two studies reporting elasticity dependence on nonpolyacrylamide substrates support the premise that the collective property E of a substrate rather than its detailed chemistry influences cell structure and function. It should be emphasized that the differentiating factor in all of these studies and those reviewed below was a property of the insoluble microenvironment rather than the soluble milieu.

Stem cells are particularly E -sensitive

MSCs appear especially sensitive to tissue elasticity. These cells reside in the bone marrow and are believed to enter the circulation and contribute to tissue regeneration and repair after injury, such as a muscle tear. Bone marrow aspirates are either fluid, with an ‘intercellular substance’ measured decades ago to be about 100-fold more viscous than water [29], or else have a very small E [30], and the rare MSCs in marrow are generally separated from the many other marrow cell types by their differential adhesion to rigid plastic. Adhesion occurs within hours [31] and is likely mediated by serum matrix proteins such as fibronectin that commonly adsorb rapidly to plastic and are also prominent in bone marrow. When replated on collagen-I-coated gels that mimic the soft tissue elasticities, a significant fraction of the slowly proliferating MSCs (in 10% serum) began within

4–24 hours to take on cell shapes that resembled a branched neuronal lineage on E_{brain} gels, a spindle-shaped myoblast on E_{muscle} gels, or a well-spread osteoblast on E_{osteoid} gels [14**]. Collagen-I is not reported to be a matrix molecule within marrow, and while this particular ligand might be necessary, collagen-I on glass was not sufficient to induce lineage-specific morphogenesis. Elasticity was clearly the differentiating factor, and the pluripotency of MSCs makes these cells especially sensitive to microenvironmental factors.

Expression of lineage-specific proteins such as MyoD occurred several days after plating, and a broader confirmation of lineage induction by one week was documented through both lineage-specific protein and transcript profiling (Figure 2b). When calibrated against established differentiated cell lines, both protein and transcript levels for the three lineages proved consistent in showing that E -induction is about half that of the established cell lines. Profiling also showed that MSCs downregulated Collagen-I, contrary to recent speculations based on highly synthetic fibroblastic cultures [25] and indicating that these multi-potent stem cells respond to their given matrix rather than remodel it. Indeed, not all stem cells are likely to be equivalent: Embryonic stem cells (ESCs) are expected to synthesize matrix for de novo tissue, and ESCs in vivo frequently generate teratomas, unlike MSCs. The MSCs would nonetheless commit to their E -induced lineages after several weeks in culture, based on the fact that induction media composed of potent synthetic glucocorticoids (e.g. dexamethasone) could redirect lineages at early but not late time points. Moreover, when myogenic or osteogenic induction media cocktails were added to the various cell on gel systems, the soluble and insoluble effects proved additive in inducing expression levels similar to committed cell lines.

One prediction from the cell biology studies of human-derived MSCs above follows from the fact that osteoid elasticity [14**] possesses an elasticity E similar to that of a fibrotic infarct scar [23]. Osteogenesis is therefore predicted to occur if MSCs are injected into an infarct scar — this is exactly what was found in mouse studies in which MSCs and hematopoietic stem cells (HSCs) were injected into infarcts, with only the MSCs causing ossification [32]. Neither the potential risks for human trials nor the need for basic studies of the mechanism of elasticity-induced signals in stem cell biology should be neglected.

Matrix-coupled, myosin-driven remodeling

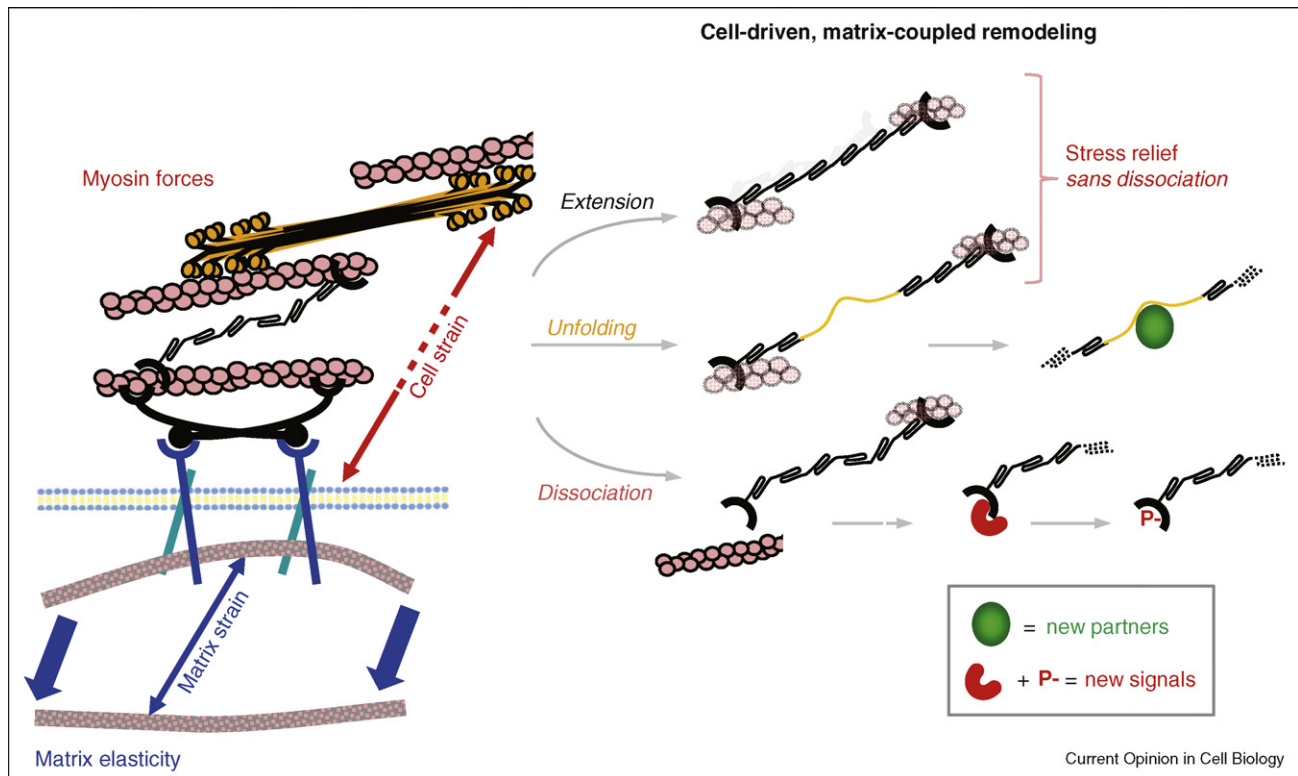
Molecular mechanisms of elasticity sensing by cells seem likely to be collective and dependent on many interacting components of the cyto-adhesion apparatus. Cell tension is expected to be important, and differentiation of MSCs was indeed blocked by myosin-II inhibition with blebbistatin [14**]. In addition, lineage specification was

associated with significant changes in the levels NMM IIB and C, with considerable downregulation on soft matrices (0.1–1 kPa) and modest upregulation on stiffer matrices (>10 kPa); NMM IIA showed the smallest variation with E among a dozen myosin motors studied, suggestive of a constitutive role in cell biological processes. The findings are consistent with the lack of differentiation in NMM IIA null mice, which are embryonic lethal at day ~6 with no signs of heart or vasculature development despite normal levels of expression of many key factors such as the transcription factor GATA-4 [33]. Embryoid bodies of the NMM IIA null cells showed no proliferation defect but appeared flaccid rather than as the typical tensed spheroids.

In the earliest studies of stiffness sensing, tyrosine phosphorylation on multiple proteins (including paxillin) appeared broadly enhanced in cells on stiffer gel substrates [8**]. Pharmacologically induced, nonspecific hyperphosphorylation could drive focal adhesion formation on soft materials, and key roles for membrane localized phosphatases have also emerged [34]. Likewise on soft substrates, overexpression of GFP-actin was found to over-ride the limited spreading, whereas overexpression of GFP-paxillin had no effect [11]. α 5-Integrin was reportedly downregulated on soft gels but overexpression had no effect on cell spreading [18]; in contrast, engineered clustering of integrins could strongly influence stiffness sensing [35*]. Microtubules have been proposed to act as ‘struts’ in cells, but quantification of their contributions to cells on gels had shown that they provide only a minor fraction of the resistance (14%) to cell-generated contractile tension [36]; the majority of a cell’s tension or stress is thus resisted by the strain of the matrix and microenvironment.

The list of molecular players in E -sensing will undoubtedly grow as will a myriad of interacting binding partners, but understanding of the molecular dynamics of cell-driven matrix-coupled remodeling seems likely to depend as much on new approaches that can clarify how forces within cells remodel proteins and their assemblies through extension, conformation, and/or dissociation (Figure 3). Extension of proteins should, under small forces, resemble stretching of a spring, but tension forces exerted on folded domains can in principle relieve the stress either by unfolding one of many domains or by dissociating one end of the protein from binding partners that anchor it down. Many ubiquitous cytoskeletal proteins including myosin’s rod domain [37], filamin [38], spectrin [39], and ankyrin [40] have been studied by AFM as purified single molecules and found to undergo forced unfolding; likewise, many adhesion proteins including VCAM [41] and integrins [42] unfold or dissociate from their ligands under force. Transitions clearly depend on the levels of force and the duration of stress, with typical transition time scales of <1 s at forces

Figure 3



Schematic representation of how myosin forces strain the cell and also strain the matrix as resisted by matrix elasticity. The strain within the cell is manifested as protein extension, domain unfolding, and/or protein-protein dissociation. These processes might simply relieve stress in the cell, or they might create sites for new binding partners or new signals.

of <100 pN that 1–20 myosin molecules can generate. Cyto-adhesion structures are therefore prone to force- f driven kinetics (i.e. $rate \approx rate_0 \exp(f/f_0)$ where f_0 is a molecule-specific constant), but identifying which proteins or complexes remodel or reorganize has of necessity motivated new methods.

Cys shotgun and other methods begin clarifying molecular dynamics

Cysteine is a reactive but relatively hydrophobic amino acid that is often buried within tertiary or quaternary structures. Reactivity of cysteine's thiol group had been exploited in solution to a limited extent to probe protein interactions [43] and folding [44], and *in situ* Cys labeling of membrane proteins such as GPCRs had yielded insights into accessibility and ligand-induced changes in individual proteins [45]. Proteomic-scale Cys shotgun labeling of intact cells has now been shown to be feasible with mass spectrometry (MS) [46••]. The basic premise is to label Cys with thiol-reactive, membrane-permeable fluorescent probes in both stressed and unstressed cells, and then search for differential labeling of proteins by multiple methods, including MS, to identify the proteins and the position of the cysteines that, under stress, have

shifted from buried and inaccessible to exposed and accessible to a fluorescent reactant.

Cys shotgun methods were first developed with the simplest possible mammalian cell, the red blood cell, which was sheared in a flow device while suspended with an encapsulated dye. The membrane-localized cytoskeleton of this cell is a simple but ubiquitous spectrin-actin network, and it showed shear-enhanced labeling within minutes as visualized by fluorescence microscopy. Both α and β spectrin showed shear-enhanced labeling in SDS-PAGE and MS analyses — although no differences were seen with actin, ankyrin, and other peripheral membrane proteins. Some of the Cys with increased labeling were clearly hidden in crystal structures or homology models of domains, and recombinant proteins studied with temperature-dependent labeling kinetics in native versus denaturing conditions proved consistent with domain unfolding as assessed by circular dichroism. Cys shotgun methods were subsequently applied to MSCs on polyacrylamide gel substrates in tensed versus blebbistatin-relaxed states [46••] and to cardiomyocytes exerting their rhythmic contractions on soft versus stiff substrates [47]. Among the most prominent differences induced by matrix elasticity were Cys sites that implicate differential

unfolding in myosin-II, differential unfolding and dissociation of filamin, and shifts in the polymerization state of vimentin.

New binding partners and new phosphorylation-based signals (Figure 3, right) might arise directly or indirectly from molecular extension, unfolding, or dissociation. Initial progress in finding such force-sensitive signaling proteins has identified p130-Cas. Within spread and fixed cells, binding of a conformation-sensitive antibody to p130-Cas in regions of the cells expected to generate the highest traction forces has suggested force-induced extension of a central proline-rich region in p130-Cas making it available for phosphorylation by Src family kinases [48**]. A role for p130-Cas in stiffness sensing by cells has yet to be demonstrated. Nonetheless, with the growing capabilities of MS for detecting functional modifications that are either natural (phosphorylation) or synthetic (modified Cys), proteomic-scale insights into the various pathways involved in how cells feel the elasticity of their microenvironment seem within reach.

Acknowledgements

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