

Imaging and manipulating the structural machinery of living cells on the micro- and nanoscale

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Abstract: The structure, physiology, and fate of living cells are all highly sensitive to mechanical forces in the cellular microenvironment, including stresses and strains that originate from encounters with the extracellular matrix (ECM), blood and other flowing materials, and neighbouring cells. This relationship between context and physiology bears tremendous implications for the design of cellular micro-or nanotechnologies, since any attempt to control cell behavior in a device must provide the appropriate physical microenvironment for the desired cell behavior. Cells sense, process, and respond to biophysical cues in their environment through a set of integrated, multi-scale structural complexes that span length scales from single molecules to tens of microns, including small clusters of force-sensing molecules at the cell surface, micron-sized cell-ECM focal adhesion complexes, and the cytoskeleton that permeates and defines the entire cell. This review focuses on several key technologies that have recently been developed or adapted for the study of the dynamics of structural micro-and nanosystems in living cells and how these systems contribute to spatially-and temporally-controlled changes in cellular structure and mechanics. We begin by discussing subcellular laser ablation, which permits the precise incision of nanoscale structural elements in living cells in order to discern their mechanical properties and contributions to cell structure. We then discuss fluorescence recovery after photobleaching and fluorescent speckle microscopy, two live-cell fluorescence imaging methods that enable quantitative measurement of the binding and transport properties of specific proteins in the cell. Finally, we discuss methods to manipulate cellular structural networks by engineering the extracellular environment, including microfabrication of ECM distributions of defined geometry and microdevices designed to measure cellular traction forces at micron-scale resolution. Together, these methods form a powerful arsenal that is already adding significantly to our understanding of the nanoscale architecture and mechanics of living cells and may contribute to the rational design of new cellular micro-and nanotechnologies.

Keywords: imaging, living cells, micro-and nanoscale, laser ablation, fluorescence recovery after photobleaching, fluorescent speckle microscopy, micropatterning, cytoskeleton, extracellular matrix.

Introduction

Normal tissue function depends on the coordinated action of a large number and diverse variety of cell types that must proliferate, function, and ultimately die at precisely defined times and places, and this intricate choreography relies heavily on a rich and continuous dialogue between individual cells and between cells and their environment. Cell biologists have long assumed that the words of this conversation are largely biochemical in nature, ie, that cells base their actions primarily on the concentrations of soluble signals to which they are exposed. However, one of the most exciting breakthroughs in cell biology in the past decade is the recognition that the *physical* microenvironment of a cell, including applied mechanical forces and the geometric arrangement, protein density, and mechanical compliance of the extracellular matrix (ECM), can regulate cell behavior in equally profound ways (Discher et al 2005). For

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example, individual mesenchymal stem cells differentiate into adipocytes (fat cells) when they are prevented from completely spreading, even when they are saturated with growth factors that would normally induce them to form bone cells (McBeath et al 2004). Similarly, mammary epithelial cells assemble into normal tubular structures when cultured in relatively compliant (soft) ECMs; when they are cultured in ECMs that are slightly more stiff, these cells form abnormal structures, detach from their neighbors, and proliferate more rapidly – all hallmarks of a developing breast tumor (Paszek and Weaver 2004). For this reason, any attempt to control or exploit cell behavior in a technological context must pay attention to how the physical microenvironment influences cell physiology. Viewed another way, physical crosstalk between the cell and its surroundings represents a design parameter that may be modulated to achieve a desired physiological outcome.

The crosstalk between cells and their surroundings is mediated by an intimately connected and interdependent set of structures that range in size from single molecules to networks that span the whole cell over tens of microns. The coordinated action of these nano- and microstructures lies at the heart of many of the characteristic processes of living cells, including adhesion, motility, and maintenance of shape. For example, cells adhere to specific extracellular matrix (ECM) components through transmembrane integrin proteins, which spatially cluster upon engagement and serve as nucleation sites for focal adhesion complexes between the cell and ECM. These micron-scale complexes contain literally dozens of distinct proteins that physically connect integrins to the cellular cytoskeleton, sense and biochemically transduce mechanical forces, and regulate the presentation and activity of various transmembrane receptors that initiate signaling (Ingber 2003; Vogel and Sheetz 2006). As cells spread and grip the ECM, they must assemble strong focal adhesion complexes that are capable of supporting increasing tractional forces generated by cellular contractile elements. Equally importantly, directional cell motility on a planar substrate depends on the ability to assemble focal adhesion complexes at the leading edge of the cell and disassemble them at the trailing edge (Lauffenburger and Horwitz 1996).

Thus, to dissect the molecular mechanisms through which cells sense and respond to their physical microenvironment, it is critical to understand how all elements in the cytoskeleton, focal adhesions, and the ECM come together in an organized fashion to allow the cell to generate and sense mechanical forces. This requires the ability

to precisely and quantitatively measure a wide variety of biophysical properties, including binding and polymerization kinetics, dynamics, and mechanical properties. Traditionally, the vast majority of these measurements have been obtained in purified systems, ie, systems in which molecular components have been purified and removed from their cellular contexts. While these approaches enable precise, quantitative measurements that have contributed greatly to our understanding of cell structure and function, they cannot recapture the spatial and temporal architecture that is essential to transmission of mechanical information in living systems and enables such life-defining processes as locomotion. Over the past decade, however, a remarkable set of technologies has emerged that seeks to satisfy both goals by obtaining quantitative measurements in living cells. Some of these technologies represent increasingly sensitive ways to image and measure the dynamics and mechanics of structures on the molecular scale, and others represent platforms with which to directly manipulate micro- and nanoscale cell structural elements. Here, we review three of these technologies: subcellular laser ablation, advanced live-cell fluorescence imaging methods (including fluorescence recovery after photobleaching and fluorescent speckle microscopy), and cellular micropatterning. These technologies promise both to dramatically advance our ability to understand the conversation between cells and their physical environment and to build our own vocabulary of commands as we attempt to exploit this knowledge in the design of technologies that interface with living systems.

Subcellular laser ablation

Mechanical inputs from the ECM represent one of the most important and well-studied classes of cues in the cellular microenvironment. Because the cellular cytoskeleton confers shape and mechanical properties to the whole cell, it is absolutely critical to a cell's ability to detect, process, and respond to these mechanical cues. More specifically, by setting its composition, three-dimensional architecture, and contractile state, the cytoskeleton serves as both a passive structural network through which environmental forces are distributed and an active generator of forces that deform the ECM and permit cell spreading and motility. While it is clear that the cytoskeleton is key to cell structure and mechanics in a global sense, considerably less is known about the loads borne by specific cytoskeletal filaments in living cells, or how individual filaments contribute to the structure and

mechanics of the whole cell. Answering these questions requires methods capable of selectively probing and disrupting individual cytoskeletal structures in living cells. Ideally, these methods should be capable of interrogating living cells on the submicron length scale in a minimally invasive fashion without killing them in the process.

Subcellular laser ablation has emerged as a powerful tool to accomplish these goals. Pioneered in the late 1970's by Michael Berns and colleagues (Strahs and Berns 1979; Koonce et al 1982; Berns et al 1991; Berns 1998; Botvinick et al 2004), and then refined over the next two decades by the laboratories of Karsten König (König et al 1999; Tirlapur et al 2001; Tirlapur and König 2002), Conly Rieder (Khodjakov and Rieder 2001; Faruki et al 2002; Maiato et al 2005), and others, this technique utilizes a focused laser beam to selectively ablate nano-to microscale structures in living cells. While the first applications of this method used continuous-wave visible-range or ultraviolet lasers, more recently, pulsed lasers have become the norm. In particular, ultrashort laser pulses (eg, nano-to femtosecond) are focused through a high-numerical aperture objective lens onto an intracellular target that may be visualized by phase contrast or fluorescence. Material at the focus undergoes nonlinear multiphoton absorption of laser energy, leading to optical breakdown and photoablation. Ideally, the laser energy is delivered quickly enough to prevent energy dissipation by heat transfer, but at a sufficiently low energy and pulse frequency to limit the zone of damage. For this reason, pulse energy, pulse width, and repetition rate have all been shown to be important parameters in limiting the precision of this method. For example, in chemically fixed cells, delivery of femtosecond laser pulses at kilohertz repetition rates and at pulse energies ranging from 1.4 nJ to 2.3 nJ can produce zones of photodamage between 0 and 1 μm . Importantly, the spatial values were determined by transmission electron microscopy (TEM) imaging of cells that had been fixed and fluorescently stained; regions of cells irradiated at the lowest energies (~ 1.4 nJ) were rendered nonfluorescent even though subsequent TEM failed to reveal damage, demonstrating that those regions had been photobleached rather than damaged (Heisterkamp et al 2005). Thus, in interpreting these experiments, it is important to distinguish between photobleaching and photoablation.

Subcellular laser ablation has emerged as a powerful tool for the measurement of mechanical properties of load-bearing cytoskeletal elements in their living, intracellular context. Perhaps the most closely studied cytoskeletal structures by this method are actomyosin stress fiber

bundles, which are contractile structures that connect to the cell-ECM interface and enable cells to exert tractional forces on their surroundings. In one of the first applications of laser ablation in living cells, Berns and coworkers visualized stress fibers by phase contrast microscopy, irradiated them with a laser microbeam, and observed their retraction and subsequent repair on a time scale of several hours (Strahs and Berns 1979). In later work, they showed that the rate of stress fiber repair depended on the integrity of the microtubule and fine actin cytoskeleton and not, surprisingly, cellular protein synthesis (Koonce et al 1982). In recent work, laser ablation has been used to study the mechanics of single stress fibers in living cells and their contributions to cell shape and ECM strain (Figure 1). Here, the authors visualized the actin cytoskeleton in living endothelial cells using yellow fluorescent protein (YFP)-tagged actin and severed stress fibers at the cell base

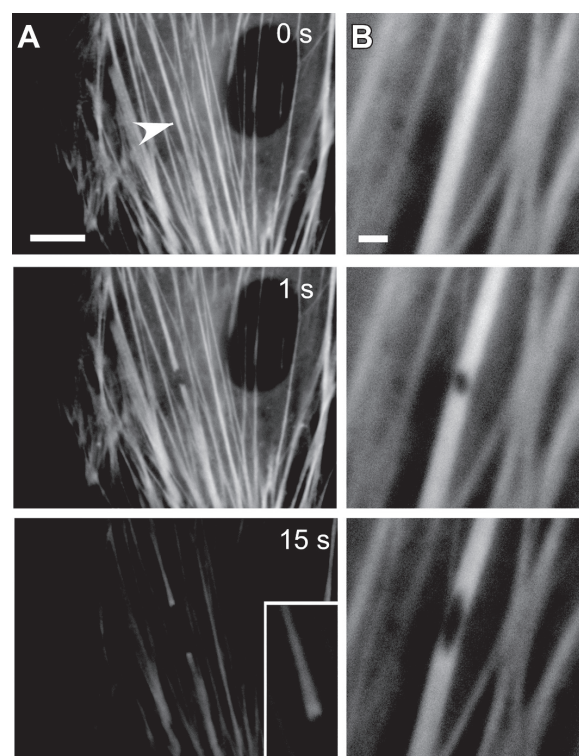


Figure 1 Subcellular laser ablation. In subcellular laser ablation (SLA), a series of high-intensity, ultrashort laser pulses are focused on an intracellular target, resulting in nanoscale destruction of material with minimal damage to surrounding structures and without killing the cell. Here, SLA is used to probe the micromechanics of yellow fluorescent protein-tagged actomyosin stress fiber bundles in an endothelial cell. Laser irradiation results in complete severing (A, arrow) or puncturing (B) of selected stress fibers, leading to profound remodeling on a time scale of 15 s; the ends of severed stress fibers retract (A) and splay apart (inset), and the irradiated region of punctured stress fibers deform into a progressively elongated ellipse (B) (Bar = 2 μm). Reproduced with permission from: Kumar S, et al. 2006. Viscoelastic retraction of single stress fibers and its impact on cell shape, cytoskeletal organization, and extracellular matrix mechanics. *Biophysical Journal*, 90:3762–73. Copyright © 2006, Biophysical Society, <http://www.biophys.org>.

with femtosecond laser pulses (Kumar et al 2006). Following irradiation, the ends of the stress fibers retracted in a straight line (ie, in parallel with the axis of the fiber) with viscoelastic recoil dynamics with characteristic times of 15–20 sec; in some cases, it was possible to puncture, rather than incise, stress fibers, yielding holes with diameters as small as 300 nm that distended into ellipses as the weakened stress fiber retracted. Importantly, the degree to which incision of one stress fiber influenced the cytoskeletal architecture and shape of the rest of the cell depended strongly on the compliance of the ECM on which cells were cultured. For cells cultured on rigid substrates (eg, glass), incision of a single stress fiber, or even multiple parallel stress fibers, produced essentially no rearrangements in cell structure. Conversely, incision of even one stress fiber in cells cultured on compliant (~4 kPa) polyacrylamide substrates produced cellular elongation and widespread rearrangements of cytoskeletal structures many microns from the site of incision. In the latter case, traction force microscopy measurements revealed that dissipation of tractional stresses into the ECM substrate were concentrated near sites of stress fiber insertion into the ECM; ie, focal adhesion complexes. In related studies, subcellular laser ablation was used to locally dissipate tension on focal adhesions by severing adjacent stress fibers, resulting in accelerated turnover of specific focal adhesion proteins. This result was consistent with results obtained upon global dissipation of cellular tension through the use of compliant culture substrates or through treating the cell with contractility-inhibiting drugs (Lele et al 2006a).

Subcellular laser ablation has also been used to sever cellular microtubules in various contexts. For example, laser ablation has been used to disrupt various parts of the mitotic spindle in dividing yeast in order to determine the distribution of tensile and compressive loads among astral and kinetochore microtubules (Khodjakov et al 2004; Tolic-Norrelykke et al 2004). More recently, this method has been applied to cortical microtubules, which play mechanical roles in establishing and stabilizing cellular structure. Botvinick et al. (Botvinick et al 2004) used a picosecond laser to irradiate and sever cytoplasmic microtubules tagged with various green fluorescent protein variants, including YFP and cyan fluorescent protein (CFP); as expected, following microtubule incision, one of the severed ends rapidly depolymerized, corresponding to the exposure of a bare minus end. Interestingly, based on subsequent TEM imaging of the ablated cells, the degree of damage differed significantly depending on whether CFP or YFP was used as a fluorescent tag, and

both cases differed from green fluorescent protein (GFP). This strongly suggested that the mechanism of photodamage is different for each fluorophore, which in turn results from differences in their respective absorption spectra associated with differences in amino acid composition. Heisterkamp et al. (Heisterkamp et al 2005) used a femtosecond laser to sever a curved GFP-tagged microtubule beneath the nucleus of living cells. Following laser severing, the bent microtubule rapidly straightened and then depolymerized, reflecting the release of stored elastic bending energy.

Fluorescence recovery after photobleaching

An important step in elucidating the influence of the physical microenvironment on cell function is to understand both the dynamics of structural elements in living cells, and how mechanical loads borne by these structures translate into specific biochemical changes at precise cellular times and locations. There are at least three challenges in obtaining these types of data. First, traditional biochemical assays rely on average measurements of populations of cells at fixed points in time, which, by definition, fail to capture information about events in specific portions of specific cells. Second, structures may appear relatively static macroscopically despite the presence of active and rapid molecular-scale remodeling events, such as subunit binding and unbinding; thus, even prolonged imaging of subcellular structures may fail to capture these subtle yet potentially important dynamics. Third, when fluorescently labeled, many cellular structural elements such as microtubules and intermediate filaments fluoresce along their entire length, thereby offering little to no negative contrast to track internal rearrangements in the body of the filament.

Fluorescence recovery after photobleaching (FRAP) has proved a powerful means to address all of these issues. In FRAP, regions of a fluorescently-tagged structure are selectively rendered nonfluorescent through irradiation with a pulse of high-intensity light at the excitation wavelength of the fluorophore, thereby photobleaching the population of fluorescent subunits in the region and rendering them optically invisible. Unlike subcellular laser ablation, FRAP does not destroy material or compromise protein function and only affects fluorescently tagged molecules whose excitation wavelength corresponds to the wavelength of irradiation. Because the loss of fluorescence in a bleached fluorophore is permanent on the time scale of the experiment, any recovery of fluorescence in the targeted structure must be due to a

combination of the departure of the bleached subunits and the arrival of new, unbleached fluorophores into the bleached region. Both of these processes may result from either a combination of transport of material within the structure (eg, diffusion) or exchange of subunits between the material and the surrounding medium (ie, binding and unbinding). Thus, given a physical model that describes the dynamics of subunit binding and dynamics, FRAP data (fluorescence vs. time) may be fitted to the model to extract kinetic parameters that describe the binding and transport properties of the labeled molecule.

FRAP has been extensively applied to interrogate the dynamics of complex and heterogeneous cytoskeletal systems, particularly intermediate filaments and actin-based structures. In one study (Yoon et al 2001), FRAP was used to probe the network dynamics of keratin intermediate filaments within cultured epithelial cells. Here, cells were transfected with fluorescently-tagged keratin subunits which then incorporated into the tonofibril network (parallel bundles of keratin-rich cables) of cultured epithelial cells, thereby labeling the tonofibrils. When a line was bleached across many tonofibrils, the bleached spots recovered very slowly (half-times of 1–2 hours), thus yielding a relatively stable fiduciary marker to track internal tonofibril dynamics. Surprisingly, bleached zones on adjacent tonofibrils often moved in opposite directions and at different rates, thereby revealing for the first time the complexities of assembly of keratin transport and turnover in these cells. Similarly-inspired studies with the nuclear envelope proteins lamin A and lamin B1 provided insights into the molecular-scale dynamics of envelope assembly and disassembly during cell division (Moir et al 2000). FRAP has also yielded tremendous insight into the internal dynamics of actin-based structures relevant to cell adhesion, mechanics, and motility, including actomyosin stress fiber bundles. Peterson et al (Peterson et al 2004) expressed fluorescently-tagged α -actinin and myosin, two actin-binding proteins found in stress fibers, in cultured fibroblasts and used FRAP to investigate the binding kinetics of these proteins to various regions of the stress fibers. Interestingly, they found that both proteins exchange more rapidly at the center of stress fibers than near their peripheral attachments to focal adhesions; together with high-resolution imaging of myosin distributions during cell contraction, the FRAP data enabled the authors to demonstrate that stress fibers are capable of simultaneously stretching and contracting along their length. In another study, FRAP was used to examine the exchange kinetics of α -smooth-muscle-actin (SMA) in the presence of an SMA peptide fragment (Clement et al 2005). By showing that a peptide derived

from the amino terminal sequence of SMA slowed recovery of SMA fluorescence and therefore retarded SMA exchange, the authors were able to develop a model in which the amino terminus of SMA facilitates incorporation of the protein into stress fibers.

More recently, FRAP measurements have helped to elucidate the influence of the mechanical microenvironment, including mechanical force and ECM rigidity, on the unbinding kinetics of specific focal adhesion proteins at the cell-ECM interface. These studies were made possible by new innovations in the acquisition and analysis of FRAP data that permit separation of contributions from molecular binding and diffusion to the recovery curve (Lele et al 2004). In one study, the authors measured dissociation rates of zyxin and vinculin from focal adhesions before and after dissipating intracellular tension by a variety of methods, including adding contractility inhibitors, culturing the cells on compliant ECMs, and severing attached stress fibers by laser ablation (Lele et al 2006a). These studies revealed that the unbinding rate of zyxin, but not vinculin, increased when tension on the stress fiber was released. Thus, zyxin was postulated to serve as a molecular mechanosensor whose binding properties, and presumably conformational properties, are highly sensitive to the tensile state of the cell (Figure 2). This approach has since been extended to the measurement of binding interactions between nuclear histone proteins and chromatin, thereby lending new insight into the biophysical basis of transcriptional control (Lele et al 2006b).

Fluorescent speckle microscopy

Fluorescent speckle microscopy (FSM) represents another powerful tool for the characterization of the microscale internal dynamics of cellular structural elements (Danuser and Waterman-Storer 2006) (Figure 3). In contrast to FRAP, which derives its ability to probe molecular dynamics from the selective inactivation of fluorophores within a continuously fluorescent structure, FSM follows the motions of fluorescent subunits within an otherwise nonfluorescent structure; this discontinuous fluorescence gives the structure a “speckled” appearance from which the name of the method is derived. Intracellular structures are typically made fluorescent in FSM experiments by either microinjecting a very low concentration of fluorescently-tagged proteins into a cell, or by expressing the protein as a GFP fusion construct at very low levels. While FSM may be performed with a wide-field epifluorescence microscope, it is increasingly being used in the context of total internal reflection

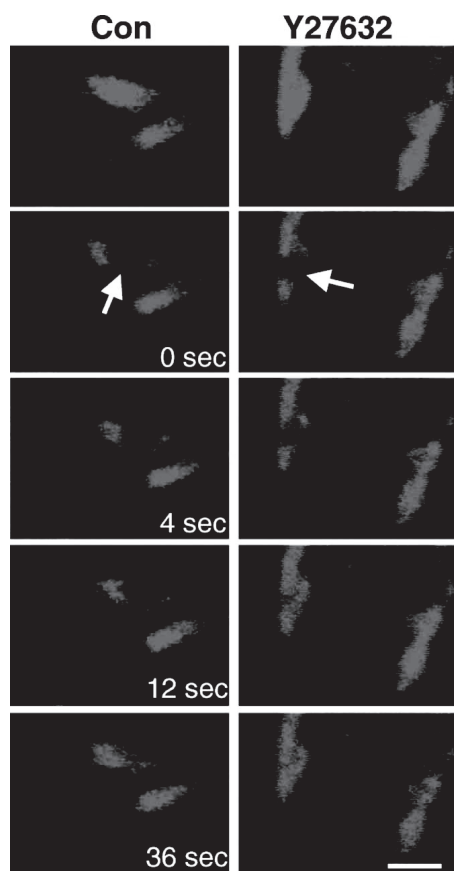


Figure 2 Fluorescence recovery after photobleaching. In fluorescence recovery after photobleaching (FRAP), a fluorescent intracellular structure is irradiated at high intensity and at the excitation wavelength, photobleaching the structure and rendering it optically invisible without destroying it. The kinetics with which the photobleached subunits are replaced by fluorescent subunits yields information about subunit transport and binding. Here, green fluorescent protein-tagged zyxin in focal adhesions is photobleached (arrow) in the absence (left column) or presence (right column) of Y27632, a pharmacologic inhibitor of intracellular contractility. Inhibition of contractility dramatically accelerates the turnover of zyxin, demonstrating that the unbinding kinetics of this protein are sensitive to the mechanical properties of the cell. (Bar = 2 μm). Reproduced with permission from: Lele TP, et al. 2006. Mechanical forces alter zyxin unbinding kinetics within focal adhesions of living cells, *Journal of Cellular Physiology*, 207:187–94. Copyright © 2006 Wiley-Liss, Inc.

fluorescence microscopy, which permits high-resolution imaging of structures at the cell-substrate interface such as the cortical cytoskeleton and focal adhesions. FSM has been used with great success to follow the internal dynamics of microtubules (Waterman-Storer and Salmon 1998), actin filaments within cellular networks (Ponti et al 2004), and a wide variety of focal adhesion proteins, including vinculin, paxillin, and talin (Geiger et al 2001). Whereas early FSM experiments were analyzed by manually tracking the motions of fluorescent particles, advanced automated tracking algorithms have emerged which allow rapid and highly quantitative interpretation of FSM data, leading to the direct estimation of local cellular viscoelastic properties and

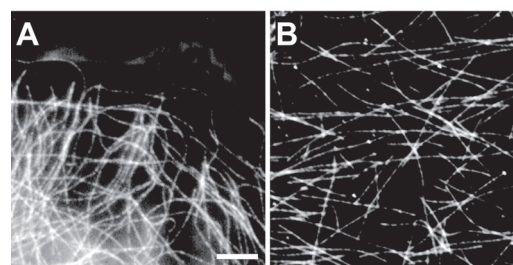


Figure 3 Fluorescent speckle microscopy. In fluorescent speckle microscopy (FSM), an intracellular structure is fluorescently labeled in a sparse fashion, such that the structure takes on a “speckled” appearance with discontinuous regions of fluorescence. The relative dynamics of these speckles reveals quantitative insight into the internal dynamics of the structure. Here, microtubules are labeled for an FSM study in a living epithelial cell (A) and in a purified preparation (B). (Bar = 5 μm). Reproduced with permission from: Waterman-Storer CM, and Salmon ED. 1998. How microtubules get fluorescent speckles. *Biophysical Journal*, 75:2059–69. Copyright © 1998, Biophysical Society, <http://www.biophysj.org>.

measurements of the degree of kinematic coupling between different cellular structural complexes (Waterman-Storer and Salmon 1998).

FSM has uncovered particularly exciting details of the dynamics of the actin cytoskeleton in migrating cells. In a seminal FSM study (Ponti et al 2004), rhodamine-actin was microinjected into cultured epithelial cells, and the resulting actin speckles were utilized to track zones of actin polymerization and depolymerization near the leading edges of cells during migration. By carefully tracking the growth, disappearance, and trajectories of actin speckles, the authors constructed high-resolution spatial maps of actin flow which defined two distinct dynamic networks: the lamellipodium, a zone of fast, retrograde actin flow within 1–3 μm from the leading edge of the cell, and the lamella, an adjacent and much broader zone of slower retrograde flow. Subsequent immunofluorescence imaging revealed that the lamellipodium was enriched in the actin branch-promoting factors ADF/cofilin and Arp 2/3, and the lamella was enriched with the contractile proteins myosin II and tropomyosin. These observations, together with pharmacological inhibition data, supported a model in which the lamellipodium gives rise to protrusion of the leading edge, and the lamella drives advancement of the whole cell through actomyosin contraction. Thus, detailed analysis of a simple yet elegant FSM experiment provided rich and novel insight into the contributions of different portions of the actin cytoskeleton to coordinated cell movement.

All of the above methods represent ways of studying the molecular aspects of crosstalk between cells and their physical microenvironment by observing and manipulating structures inside living cells. We now turn to technolo-

gies that have been created to tap into this crosstalk by engineering the geometry of the extracellular matrix of cells. In some cases, these approaches have been used to study the role of cellular structure in driving cell function, and in others they have been used to measure physical interactions between cells and their surroundings at the micro- and nanoscale.

Microcontact printing approaches to controlling cell geometry

Direct patterning of extracellular matrix proteins by microcontact printing offers a powerful way to control the geometry of single cells and small groups of cells in two-dimensional culture. Analogous to conventional ink-stamp printing, microcontact printing is based on the controlled deposition of ECM proteins by apposition of a protein-coated stamp with a cell culture substrate (eg, glass) and subsequent passivation of the non-stamped regions of the substrate. By controlling the topographical relief pattern on the stamp, one may create micron-sized islands of defined location, size, and shape that in turn dictate the distribution and geometry of cells cultured on the substrate (Chen et al 1997; Lauer et al 2001; Brock et al 2003; McBeath et al 2004). Typically, photolithography is used to create the inverse of the desired relief pattern on a silicon wafer that will serve as a “master.” A layer of photoresist (eg, SU-8) is spun onto the wafer at a controlled thickness. The photoresist layer is then selectively exposed to UV light via a photomask of the desired pattern. Finally, the resist is developed by application of a developer solution that, in the case of negative resist, removes the unexposed regions of the photoresist, leaving behind the desired microtopographical pattern on the wafer. An elastomer, frequently poly(dimethylsiloxane) (PDMS), is then cast and crosslinked atop the master; separation of the polymer and master yields a flexible stamp with the desired microtopographical pattern on its surface. Importantly, many PDMS stamps can be cast from the same master making this a highly cost-effective, reproducible, and high-throughput means of patterning proteins. Often, in order to facilitate adsorption of ECM proteins, the surface of the PDMS stamp is modified from a methyl-terminated surface to a hydrophilic hydroxyl-terminated surface with either air plasma or ultraviolet ozone (Ye et al 2006). Following passive adsorption of ECM protein, the coated surface is placed into conformal contact with a substrate, stamping the ECM protein into the desired pattern. Non-stamped regions of the substrate are

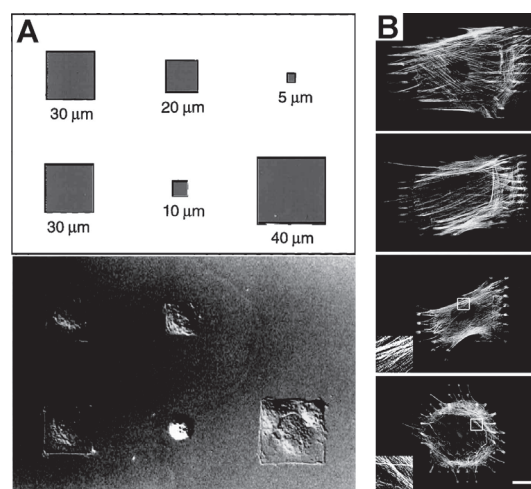


Figure 4 Control of cellular structure with micropatterning. Nomarski differential interference contrast image of endothelial cells cultured on square fibronectin islands (**A**). The cell shapes conform to the geometry of the adhesive islands as defined in the schematic. Reprinted with permission from: Chen CS, et al. 1997. Geometric control of cell life and death. *Science*, 276:1425–8. Copyright © 1997, AAAS. <http://www.sciencemag.org>. Microcontact printing of focal adhesion-sized islands of fibronectin onto glass (**B**). Islands of a variety of sizes (20, 10, 6, and 2 μm , from top panel to bottom panel) were used to support the culture of myofibroblasts, which were then stained for F-actin. While all of the myofibroblasts cultured on the substrate developed actin-based stress fibers, stress fibers in cells cultured on islands larger than 6 μm preferentially incorporated α -smooth muscle actin from the cytosol, indicating higher contractility (not shown). Inset shows a 4 \times magnification of F-actin stress fibers. (Bar = 20 μm). Reproduced with permission from: Goffin JM, et al. 2006. Focal adhesion size controls tension-dependent recruitment of α -smooth muscle actin to stress fibers. *The Journal of Cell Biology*, 172:259–68. Copyright © 2006, The Rockefeller University Press.

often passivated with an amphiphilic block copolymer (eg, Pluronic) to inhibit protein adsorption and cell binding.

Ingber, Whitesides and coworkers have used microcontact printing to create two-dimensional fibronectin islands of defined geometry for the culture of single cells, a technique that has enabled them to study the effect of cell size on cell fate independent of ECM protein density, cell-cell contacts, or media conditions (Singhvi et al 1994; Chen et al 1997) (Figure 4A). Endothelial cells cultured on the adhesive islands conformed to the size and geometry of the stamped ECM, and cell spreading could be promoted by patterning either a single, cell-sized ECM island (eg, 900 μm^2) or multiple small focal adhesion-sized islands distributed over a cell-sized area. Importantly, spreading area strongly influenced whether cells proliferated or underwent apoptosis; specifically, as cells were allowed to spread to larger and larger areas, the frequency of proliferating cells increased whether the ECM was presented as a large, continuous island or multiple, discontinuous islands distributed over a large area (Chen et al 1997).

McBeath and coworkers (McBeath et al 2004) leveraged this approach to investigate the role of cell area in

driving differentiation of human mesenchymal stem cells (hMSCs). Remarkably, they found that hMSCs confined to small ($1024\ \mu\text{m}^2$) islands appeared round and differentiated exclusively into adipocytes, whereas hMSCs cultured on large ($10,000\ \mu\text{m}^2$) islands spread and differentiated exclusively into osteoblasts. Cells cultured on islands of intermediate size ($2025\ \mu\text{m}^2$) yielded mixed populations of adipocytes and osteoblasts. The authors hypothesized that these cell area-dependent effects were due to changes in intracellular contractility; ie, cells allowed to spread to larger areas generated more contractile forces than those confined to small areas. To test this hypothesis directly, the authors manipulated hMSC contractility by controlling the activity of the small GTPase RhoA, which strongly promotes actomyosin activation, by transfecting hMSCs with either a constitutively active or a dominant negative version of RhoA. Cells made to express constitutively active RhoA predominantly differentiated into osteoblasts, while cells made to express dominant negative RhoA predominantly differentiated into adipocytes. Remarkably, this result held even when the hMSCs were cultured in media normally expected to promote proliferation or differentiation into the other cell type; eg, hMSCs with abrogated RhoA activity differentiated into adipocytes even when cultured in osteogenic media. Thus, the degree to which hMSCs can generate tensile forces against the ECM figures centrally in the fate decision between adipogenesis and osteogenesis. These studies, which provided fundamental insight into our understanding of hMSC physiology, would not have been possible without microcontact printing technology.

More recent studies have taken advantage of microcontact printing to control the geometry and distribution of individual focal adhesion complexes between the cell and ECM (Goffin et al 2006) (Figure 4B). Here, arrays of fibronectin islets spaced several microns apart with a constant width ($1.25\ \mu\text{m}$) and lengths ranging from $2\text{--}20\ \mu\text{m}$ were printed on a rigid, planar substrate. Cells cultured on these substrates formed focal adhesion complexes that assumed the size and shape of the patterned islands. Varying the length of the focal adhesion sites influenced the biochemical composition of the associated intracellular stress fiber bundles; in particular, stress fibers associated with long ($>8\ \mu\text{m}$) islands, and hence long focal adhesion complexes, preferentially incorporated α -smooth muscle actin (α -SMA), a marker of high contractility. The robustness of this system allowed for the simultaneous alteration of several mechanical cues including ECM elasticity and density as well as islet spacing and size. By systematically varying each parameter, the authors

demonstrated that islet size dominantly influenced α -SMA localization to stress fibers.

Recent advances in soft lithography: Measuring cell-ECM traction forces

While the earliest applications of soft lithography to cell biology focused almost exclusively on the control of cell size and shape, more recent approaches have sought to measure functional properties, including the spatial distribution of cell-ECM traction forces. For example, Tan and coworkers (Tan et al 2003) developed a microfabricated post array detector (mPAD) system to measure cell traction forces and contractility. For this technology, the authors fabricated arrays of deformable, cell-adhesive microposts, schematically similar to a “bed of needles” (Figure 5A). As cells spread and exerted tractional forces on these microposts, they elastically deformed them; from the magnitude and direction of the deformation, the tractional force on a given micropost could be directly determined. The mechanical properties of each mPAD system, and therefore the range of measurable forces, could be tuned through the geometry of the constituent microposts (eg, $2\text{--}10\ \mu\text{m}$ in diameter and $3\text{--}50\ \mu\text{m}$ in height). Furthermore, by using modified microcontact printing methods to selectively stamp fibronectin islands onto each mPAD, the researchers added another dimension of control by controlling of area available for cell spreading. By varying the area of the stamped islands, the authors demonstrated that the total area of focal adhesions regulated the magnitude of traction force across the whole cell. Moreover, as spreading area increased, total tractional force increased; as expected, tractional force could be altered by manipulating intracellular RhoA activity.

A similar technique was subsequently used to study traction forces exerted by an epithelial monolayer (du Roure et al 2005) (Figure 5B). Here, the authors fabricated a substrate of narrow, densely packed PDMS micropillars ($2\ \mu\text{m}$ diameter, $1\ \mu\text{m}$ post-post spacing) in an effort to improve the spatial resolution of the force measurements. In an important extension of the earlier technology, the authors examined cell-ECM tractional mechanics in a monolayer patch of epithelial cells, rather than only those of isolated cells. The authors found that the greatest force was exerted at the edge of the monolayer, and that tractional forces exerted by these cells exceeded those generated by isolated cells, suggesting cooperative mechanical behavior within the monolayer. This finding was later confirmed and significantly extended in a multicellular study with the mPAD technology (Nelson et al 2005).

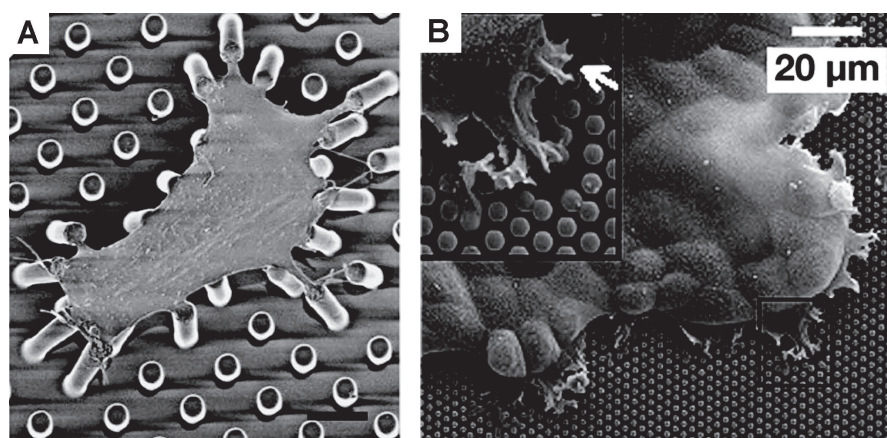


Figure 5 Measurement of cellular traction forces with micropost systems. Scanning electron micrograph of a smooth muscle cell cultured atop an array of flexible PDMS pillars with fibronectin-coated tips (**A**). The degree to which each post deflects directly reveals the contractile force exerted by the cell at that position on the extracellular matrix. (Bar = 10 μ m). Reproduced with permission from: Tan JL, et al. 2003. Cells lying on a bed of microneedles: an approach to isolate mechanical force, *Proceedings of the National Academy of Science of the United States of America*, 100:1484–9. Copyright © 2003, PNAS. <http://www.pnas.org>. Scanning electron micrograph of an epithelial monolayer on an array of closely packed PDMS pillars (**B**). These measurements revealed that the highest contractile force is produced by the leading edge (inset) of the monolayer; moreover, cells within monolayers exerted greater tractional forces than isolated cells, suggesting cooperative mechanical behavior. (Bar = 20 μ m). Reproduced with permission from: du Roure O, et al. 2005. Force mapping in epithelial cell migration, *Proceedings of the National Academy of Science of the United States of America*, 102:2390–5. Copyright © 2005, PNAS. <http://www.pnas.org>.

Nanoscale patterning of cell adhesion receptors

Surface patterning can also be a useful tool on a scale unachievable with microcontact printing. The mechanics of cell adhesion have been controlled on the molecular level using nanopatterning. Arnold and colleagues (Arnold et al 2004) utilized a diblock copolymer micelle self-assembly system to pattern 8 nm diameter gold nanodots in a rigid hexagonal pattern with precise interparticle spacings (28, 53, 73, 85 nm). The dots were then functionalized with cyclic arginine-glycine-aspartate (RGD)-containing peptides that enabled ligation of transmembrane integrins. Importantly, the small size of each dot permitted conjugation of a single peptide, which in turn bound only a single integrin receptor. Thus, by controlling the spacing of the dots, the authors were able to precisely manipulate spacing between adjacent integrins. As expected, small spacing between dots permitted enhanced integrin clustering, which in turn promoted cell adhesion and spreading as well as formation of robust focal adhesion complexes and stress fiber bundles. Conversely, greater separation distances between the dots (≥ 73 nm) precluded integrin clustering and severely limited cell attachment and spreading.

Conclusions and future prospects

Traditional approaches to studying cell physiology have emphasized treating cultured cells with defined concentrations of soluble cues and characterizing their responses.

While these studies have established the foundation for our modern understanding of receptor-mediated signal transduction, hormonal physiology, and other fundamental aspects of cell biology, it is becoming increasingly clear that these kinds of inputs do not tell the whole story. In particular, the physical microenvironment in which cells exist can provide signals that strongly modulate or even oppose the effects of these soluble cues, suggesting that elements of the physical microenvironment may provide an important but largely underappreciated role in regulating cell behavior and consequently tissue assembly, development, and pathology. This growing recognition is spawning an entirely new generation of cell biological experiments in which parameters such as ECM stiffness, geometry, and topography have been included as experimental variables, which has in turn created a need for entirely new platforms and tools to create the appropriate experimental conditions. Because much of the cellular machinery that senses, processes, and responds to these physical inputs range in size from single molecules to tens of microns, methods capable of probing nano- and microscale phenomena in living cells have proven tremendously useful in elucidating these signaling mechanisms and serving as a handle with which to engineer cell behavior. In this review, we have discussed three foundational technologies that are beginning to make particularly significant impact: subcellular laser ablation, advanced live-cell fluorescence methods including FRAP and FSM, and cellular micropatterning. The first two methods have enabled examination of subcellular mechanics and dynamics at

unprecedented spatial and temporal resolution, and the third method has permitted both the precise engineering of cell structure and quantitative measurement of the mechanics of cell-ECM adhesion and traction.

Several challenges remain to be tackled in the future. In the case of subcellular laser ablation, efforts are ongoing to increase the spatial and temporal resolution of material destruction by using pulsed lasers that offer optimal combinations of pulse width, repetition rate, and energy (Vogel et al 2005). Moreover, because lasers may be used for capturing particles (eg, through an optical trap), cell stretching, and a variety of other processing modalities, efforts are beginning to emerge to incorporate laser ablation into multimodal laser processing systems, enabling microdissection and capturing capabilities in a single platform (Stuhrmann et al 2006). Similarly, because laser ablation is a multiphoton process that may be induced with the same sorts of near-infrared lasers that are typically used for multiphoton imaging, it has become attractive to develop combined systems for imaging and ablation. For example, the same femtosecond laser system was recently used to image *Drosophila melanogaster* embryos by both third harmonic generation and two-photon modalities as well as to ablate multicellular regions of the embryo (Supatto et al 2005). In this way, the authors were able to correlate expression of a fluorescent, mechano-sensitive reporter gene with local disruptions of the multicellular force balance between cells in different portions of the embryo. Finally, one of the key limitations of subcellular laser ablation is the lack of intrinsic molecular specificity; in other words, all material at the laser focus is destroyed rather than specifically-tagged macromolecules. Here, a complementary technique known as chromophore assisted laser inactivation (CALI) is beginning to address this need; CALI is based on irradiation of a fluorophore-tagged intracellular target at the fluorophore's excitation wavelength, but at a much higher intensity than would be required for imaging (Rajfur et al 2002; Tour et al 2003; Tanabe et al 2005; Tour 2005). The resulting laser-induced photochemical reaction leads to highly localized production of short-lived reactive oxygen species, which then inactivate the tagged protein.

Several new advances are also in progress for FRAP and FSM. First, beyond tracking the dynamics of single structural elements, FRAP and FSM are increasingly being used to study correlations between the dynamics of different structural complexes in relation to one another. For example, Danuser, Waterman-Storer and colleagues have begun to fluorescently label and simultaneously obtain FSM data for both actin and focal adhesion proteins

in an effort to understand how focal adhesions mediate mechanical coupling between the cytoskeleton and ECM (Danuser and Waterman-Storer 2006). Moreover, FRAP has been combined with various mechanical measurements or perturbations to gain insight into the connection between intracellular molecular dynamics and communication with the physical microenvironment. Earlier, we discussed the combination of FRAP with SLA and contractility-inhibiting drugs to study mechanical regulation of focal adhesion turnover (Lele et al 2006a). FRAP has also recently been combined with osmotic pressure manipulations to probe the effect of mechanical force on lipid mobility in outer hair cells (de Monvel et al 2006), and FRAP has been used in parallel with optical trapping methods to correlate focal adhesion protein turnover with force-induced cytoskeletal remodeling (von Wichert et al 2003). With the expanding battery of methods to measure and deliver force to cells at ever-smaller length scales, these types of studies will increase and promise to revolutionize our understanding of how cells couple mechanics and biochemistry on the nanoscale.

Finally, there are several future opportunities for micropatterning approaches. One important challenge that remains is to develop patterning strategies that realistically capture the architecture of multicellular tissues. For example, Nelson and coworkers have cultured endothelial cells on micron-to-millimeter-sized ECM islands, thereby permitting adhesion of many cells in a confined area (Nelson et al 2005). Careful examination of proliferation rates in different portions of the island revealed that cells proliferate most rapidly at the edges, even when N-cadherin-mediated cell-cell contacts were broken. Finite-element calculations and mPAD force measurements demonstrated that cells at the edges of the islands also experienced the highest net mechanical forces, suggesting an intrinsic connection between mechanical force and cell fate determination. Micropatterning strategies are also increasingly being used to support co-culture of heterogeneous collections of cells, in an effort to work towards "bottom-up" engineering of functional tissues (Bhatia et al 1998; Bhatia et al 1999; Tourovskaia et al 2003). Perhaps the greatest unmet challenge in this area is to develop three-dimensional patterning strategies for single cells with precise control of ECM geometry and biochemistry. The creation of microwell-like structures that simulate a three-dimensional environment but still allow controlled ECM biochemistry while isolating cells appears to be one promising approach (Ostuni et al 2001); the culture of cells on substrates with topographically-defined three-dimensional

structures is certainly another (Norman and Desai 2005; Norman and Desai 2006).

The intersection between cell biology and the micro- and nanosciences has never been greater. As cell biologists and bioengineers seek to understand the molecular-scale mechanisms that facilitate crosstalk between cells and their physical microenvironment with greater precision and sophistication, tools designed to interface with cells at mesoscopic length scales will become indispensable. While the technologies described in this review all began as highly specialized methods and in many cases were first developed by physicists and engineers, they are all rapidly becoming accessible to traditional cell biologists and are being closely integrated with the standard tools of cell and molecular biology. We are headed toward a future in which the well-rounded practice of cell biology will require comfort and facility with the micro- and nanotechnologies; if the nascent efforts described in this review are any indication, this will be a bright future indeed for our understanding of cellular structure and function.

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