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Direct evidence that tumor cells soften when navigating confined spaces

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ABSTRACT The mechanical properties of cells strongly regulate many physiological and pathological processes. For example, in cancer, invasive and metastatic tumor cells have often been reported to be softer than nontumor cells, raising speculation that cancer cells might adaptively soften to facilitate migration through narrow tissue spaces. Despite growing interest in targeting cell softening to impede invasion and metastasis, it remains to be directly demonstrated that tumor cells soften as they migrate through confined spaces. Here, we address this open question by combining topographically patterned substrates with atomic force microscopy (AFM). Using a polydimethylsiloxane open-roof microdevice featuring tapered, fibronectin-coated channels, we followed the migration of U2OS cells through various stages of confinement while simultaneously performing AFM indentation. As cells progress from unconfined migration to fully confined migration, cells soften and exclude Yes-associated protein from the nucleus. Superresolution imaging reveals that confinement induces remodeling of actomyosin stress fiber architecture. Companion studies with flat one-dimensional microlines indicate that the changes in cytoarchitecture and mechanics are intrinsically driven by topographical confinement rather than changes in cellular aspect ratio. Our studies represent among the most direct evidence to date that tumor cells soften during confined migration and support cell softening as a mechanoadaptive mechanism during invasion.

INTRODUCTION

The mechanical properties of cells deeply underlie many physiological functions, such as adhesion, migration, and differentiation. Alterations in cellular mechanics are strongly associated with a number of disease processes; for example, many of the classically described hallmarks of cancer (Hanahan and Weinberg, 2000, 2011) are predicated on dramatic changes in cellular mechanical properties. In particular, the ability of tumor cells to invade tissues and metastasize to distant sites requires tumor cells to overcome mechanical barriers posed by migration through confined spaces

within the extracellular matrix and between cells (Fidler, 2003; Talmadge and Fidler, 2010; Lambert *et al.*, 2017). The growing appreciation that tumor cells may need to adaptively change their mechanics to drive disease has motivated significant efforts to explore altered cellular mechanics as a physical biomarker and the mechanotransductive machinery as a therapeutic target. Many of these studies have focused on identifying differences in mechanical properties between tumor cells and their nontumorigenic counterparts (Lekka *et al.*, 1999; Zhang *et al.*, 2002; Cross *et al.*, 2008; Faria *et al.*, 2008; Li *et al.*, 2008; Kumar and Weaver, 2009; Swaminathan *et al.*, 2011; Rebelo *et al.*, 2013; Alibert *et al.*, 2017; Rianna and Radmacher, 2017a). Tumor cells are frequently found to be softer than normal cells, which has raised the attractive idea that tumor cells adaptively soften during invasion and metastasis (Guck *et al.*, 2005; Xu *et al.*, 2012).

Despite its intuitive appeal, the adaptive-softening hypothesis is primarily based on population-level comparisons between normal and tumor cells (Guck *et al.*, 2001, 2005; Hou *et al.*, 2009; Lekka and Laidler, 2009; Byun *et al.*, 2013; Lee and Liu, 2014; Pachenari *et al.*, 2014). For example, atomic force microscopy (AFM) studies have indicated that cancerous human bladder cells are an order of

magnitude softer than healthy bladder cells (Lekka *et al.*, 1999). High-throughput optical stretching measurements similarly indicate that human breast cancer cells deform more than mammary epithelial cells and that deformability increases with metastatic potential (Guck *et al.*, 2005). However, these population-based studies paint a somewhat mixed picture. For example, AFM measurements have revealed that leukemia cells are stiffer than leukocytes (Lam *et al.*, 2008), and micropipette aspiration studies show that hepatocellular carcinoma cells are stiffer than normal hepatocytes (Zhang *et al.*, 2002). A variety of other studies employing a range of mechanical modalities provide evidence that tumor cells are either stiffer than their normal counterpart or stiffen during tumor progression, complicating the adaptive-softening hypothesis (Baker *et al.*, 2010; Lopez *et al.*, 2011; Staunton *et al.*, 2016). The lack of clarity in the evolution of cell stiffness during invasion fundamentally derives from a lack of direct measurements of cellular mechanics during the invasion process. This gap has undoubtedly arisen in part from the technical challenge of longitudinally measuring the mechanical properties of a cell during confined migration.

To address this challenge, we engineered a microscale culture platform that allows us to measure changes in cellular mechanical properties during confined migration. Our system is based on an open-roof polydimethylsiloxane (PDMS) microdevice with tapered, extracellular matrix (ECM)–functionalized channels. We observed tumor cells spontaneously migrating from flat to confined spaces while simultaneously measuring elastic properties by AFM. We find that cells soften substantially during channel traversal (Young's modulus values reduce from 5.6 to 2.1 kPa), which is accompanied by the presence of increased actomyosin density at the side walls as well as exclusion of Yes-associated protein (YAP) from the nucleus. Notably, when cells were confined through culture on narrow, one-dimensional (1D) ECM microlines, cells elongated but did not soften, indicating that softening is an intrinsic feature of three-dimensional (3D) geometries.

RESULTS

To determine how tumor cells adapt their mechanics as they migrate through confined spaces, we sought to design a culture paradigm that places cells in confined migration while allowing longitudinal live-cell optical imaging and AFM indentation measurements. As a starting point, we employed open PDMS microchannels, which have been used to model confined migration (Cha *et al.*, 2015; Smith *et al.*, 2016). Our first challenge was to identify geometric conditions under which cells experience lateral cell confinement and begin to physically engage the channel walls. We therefore fabricated individual devices featuring linear, fibronectin-coated PDMS channels with widths ranging from 5 to 100 µm (Figure 1A). When we cultured U2OS human osteosarcoma cells within these devices, we found that the morphology varied strongly with channel width (Figure 1B). On flat PDMS surfaces, cells adopted a highly spread morphology with random orientations, as expected. As the channel width was reduced to 50–100 µm, cells tended to adhere to a single channel wall and assemble stress fibers (SFs) along that wall. When the channel width was further reduced to 20 µm, cells were fully confined within the channel, with elongated morphologies and extensive contacts with both walls (see Supplemental Figure S1 for additional confocal images). Further reduction of channel width to 5–10 µm did not produce confinement, with cells spanning multiple channels and the adjoining mesas (plateaus). AFM imaging confirmed that cells occupy a 3D morphology within the channels, with engagement of the walls through their entire height (Figure 1C; see Supplemental Figure S2 for AFM images of the other channel

widths). We also quantified focal adhesion (FA) and nuclear morphology (Figure 1D). While FA length was indistinguishable across conditions, FA orientation varied significantly relative to the channel axis (calculated as $\theta_{FAs}-\theta_{pattern}$). Specifically, increasingly narrow channels increased the alignment of FAs with the channel. Nuclear area was indistinguishable from flat PDMS across all channel widths, save for a moderate decrease in the narrowest channel (5 µm). Nuclear aspect ratio began to increase relative to flat PDMS for 20-µm-wide channels and progressively increased with further channel narrowing.

To determine whether channel confinement influenced cellular mechanics, we obtained arrays of AFM indentation measurements on cells within 100, 50, and 20 µm channels, performing force maps on the supranuclear regions of each cell (Figure 2, A and B). Each map was made of 256 (16 × 16) force curves and each force curve (Figure 2C) was fitted with the Hertzian model for pyramidal probes to yield a Young's modulus. Young's modulus values were highest on the flat, unconfined surfaces and fell with decreasing channel width (Figure 2D and Supplemental Figure S3). To explore the generality of this relationship in a distinct tumor cell–matrix system, we repeated these measurements with U87 human glioblastoma cells on patterned hyaluronic surfaces and found the same trend, with cells getting softer when laterally confined and squeezed within channels (Supplemental Figure S4).

Having established 20 µm as a critical channel width to induce confinement in U2OS cells, we introduced a new PDMS device with a tapered, Y-shaped channel to allow observation of cell migration from an unconfined to a confined geometry (Figure 3A). An important advantage of the Y-shaped channel design is that it allowed us to capture, on a single surface, "snapshots" of many cells at different stages of confined migration, ranging from fully unconfined to partially confined to fully confined. To better restrict cells to the channels we passivated the mesas between the channels by microcontact-printing polyethylene glycol (PEG) before fibronectin-coating the device. Time-lapse imaging of U2OS cells within these devices revealed successful confined migration, with cells entering the channels from one end and traversing the entire length of the channel (Supplemental Movie S1).

We then used AFM indentation to follow changes in cellular mechanics before and after confinement (Figure 3, B–D). To simplify interpretation, we sorted each cell into one of four categories: 1) unconfined; 2) associated with one wall; 3) associated with two walls but not fully confined; and 4) fully confined. Comparison of cells in each category revealed that U2OS cells progressively soften as they transition from unconfined migration to partially confined migration to fully confined migration (Young's modulus value decreases from 5.6 to 2.1 kPa). These findings are consistent with our measurements in straight channels (Figure 2) and indicate that cells adaptively soften as they encounter increasingly confined geometries. Interestingly, time-lapse imaging failed to yield significant differences in migration speed or directionality as a function of confinement stage, at least over this limited time window (Supplemental Figure S5 and Movie S2).

To gain additional insight into cytoarchitecture during the transition to confined migration, we performed confocal imaging of the actin cytoskeleton within this device. As expected, cells in unconfined geometries primarily assembled SFs at their basal surface, with the expected distribution of dorsal fibers, transverse arcs, and ventral fibers (Lee *et al.*, 2018). However, cells in confined geometries mainly positioned SFs adjacent to the channel walls, creating two parallel arrays of actin bundles and few SFs adjacent to the device floor (Supplemental Movies S3 and S4). To determine whether these changes in mechanics and cytoarchitecure were accompanied by functionally
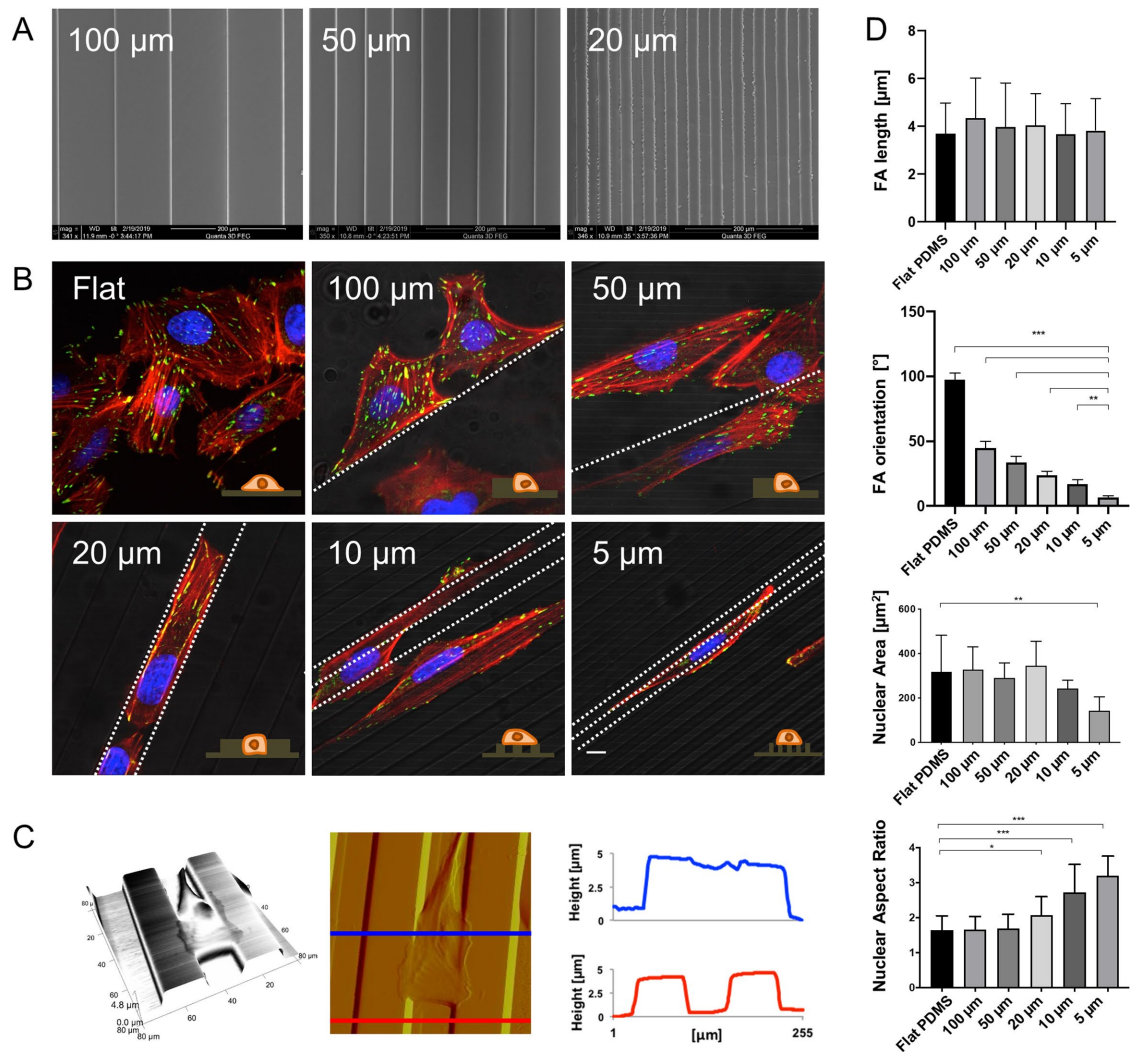
FIGURE 1: U2OS cell morphology varies strongly with PDMS channel width, achieving cell confinement within 20-μm spaced channels. (A) Scanning electron microscopy (SEM) images of linear PDMS channels with widths of 20 μm, 50 μm, and 100 μm, and lateral wall heights of ~5 μm. (B) Confocal immunofluorescence images of U2OS cells on flat surfaces and in 5–100-μm-wide channels showing actin (red), vinculin (green), and nuclei (blue). The fluorescence images are superimposed on grayscale phase contrast images to show the underlying substrate; white dotted lines indicate the pattern direction. Scale bar is 10 μm. (C) Left to right, AFM 3D projection, deflection image, and cross-sectional profiles of U2OS cell confined within 20-μm-spaced PDMS channels. As displayed in the deflection image, the cross-section images are measured on the top of the cell (blue) and on bare channels (red). (D) Quantification of FA length and orientation (top two plots) and nuclear area and aspect ratio (bottom two plots) on flat PDMS and channels of 5–100 μm width. All error bars depict standard deviations ($n = 10$ across three independent experiments; ***, $p < 0.0001$; **, $p < 0.001$; *, $p < 0.01$, as calculated by the Mann-Whitney test between each pair of categories).

relevant mechanotransductive signaling events, we examined localization of YAP, which has been observed to traffic to the nucleus under conditions of high cellular contractility (Dupont *et al.*, 2011; Elosegui-Artola *et al.*, 2017). Confocal imaging revealed strong YAP nuclear localization for cells in unconfined geometries and strong cytoplasmatic localization for cells confined within channels (Figure 4). However, when cells were either associated with one channel wall or incompletely confined (categories 2 and 3 in Figure 3B), YAP mainly localized to the nuclei. This result aligns with our AFM measurements, which show that cells soften as they traverse narrow channels (Figure 3D). In other words, cells on unconfined regions of the substrate are stiff, have well-developed basal SF networks, and show strong YAP nuclear localization. As cells experience increasing

confinement, they soften, increase actin density at the channel walls, and partition YAP from the nucleus into the cytoplasm.

The confocal images illustrate that confinement leads to dramatic changes in cell shape, in addition to changes in mechanics and YAP localization. Importantly, cellular elongation has been reported to alter cytoarchitecture and YAP localization independent of topographical confinement (Bao *et al.*, 2017; Gegenfurtner *et al.*, 2018). This raises the possibility that narrow channels might induce softening and YAP localization by altering cell shape rather than by introducing mechanical constrictions (Bao *et al.*, 2017; Gegenfurtner *et al.*, 2018). To experimentally separate elongation and mechanical confinement, we conducted a parallel set of studies in which we cultured cells on fibronectin (FN) 1D microlines of widths ranging
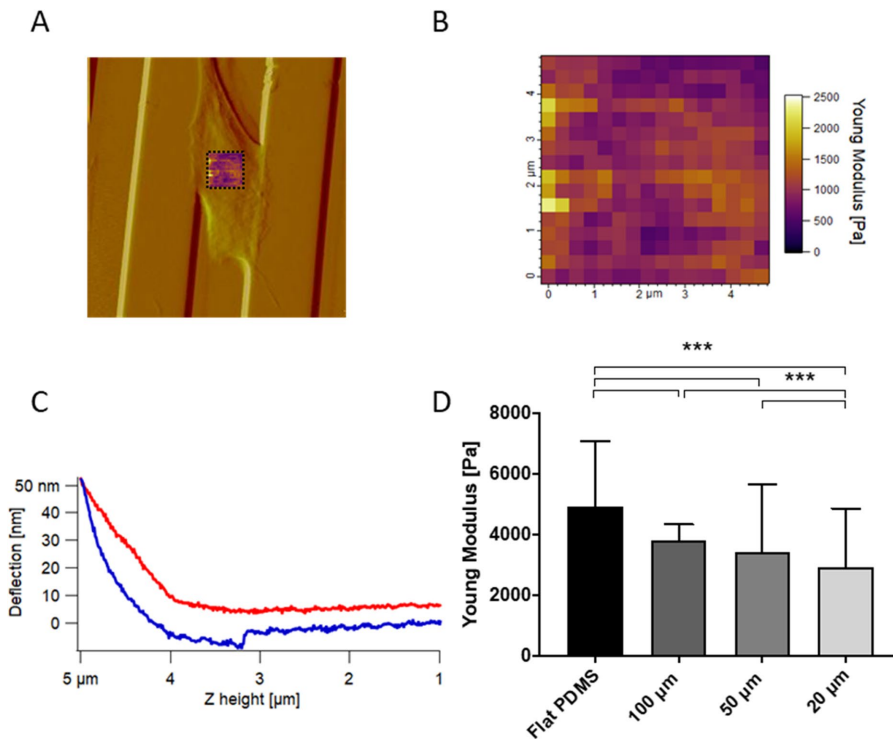
FIGURE 2: Microchannel confinement regulates cell mechanics. (A) AFM deflection image showing that force maps were measured over the central, supranuclear portion of the cell. (B) AFM force map made of 16 × 16 force curves over a 5 μm × 5 μm area; each pixel of the force map represents a force curve. (C) Representative force curve with approach in red and retraction in blue. Each curve within the force volume was fit with the Hertz model to extract the Young's modulus, with the median value across all curves reported in the bar plot. (D) Young's modulus values of U2OS cells seeded on flat PDMS and 20–100 μm channels. As above, each bar represents the median value of all the Young's modulus values collected on all the cells for each category, plotted with interquartile range as error bars. Statistical significance was determined using the Mann-Whitney test between each pair of categories (***, $p < 0.0001$; $n = 2560$, 2304, 3072, and 3328 force curves on 10, 9, 12, and 13 cells across three independent experiments. For each cell, one 16 × 16 force map was recorded on the supranuclear region of the cell body).

have often focused on mechanical comparisons between cancer and normal cells (Lekka *et al.*, 1999; Cross *et al.*, 2008; Faria *et al.*, 2008; Li *et al.*, 2008; Rebelo *et al.*, 2013). These studies have yielded tremendous insight into the role of cellular mechanics during invasion and metastasis and raised the notion that tumor cells must be highly deformable to navigate tight tissue spaces (Guck *et al.*, 2005; Xu *et al.*, 2012). While many studies have indeed shown that cancer cells are softer than normal cells (Lekka *et al.*, 1999; Cross *et al.*, 2008; Faria *et al.*, 2008; Li *et al.*, 2008; Rebelo *et al.*, 2013), the result is far from universal, with some studies indicating that tumor cells are stiffer than normal cells (Zhang *et al.*, 2002; Lam *et al.*, 2008), especially when they are in contact with an ECM (Pickup *et al.*, 2014; Acerbi *et al.*, 2015; Rianna and Radmacher, 2017a,b; Abidine *et al.*, 2018). Moreover, the interpretation of many of these studies is complicated by the differences in genetic background between the normal and tumorigenic culture models, such that it is difficult to know whether measured differences in cell stiffness are intrinsically associated with differences in tumorigenic potential. In fact, when mechanical comparisons have been made within an isogenic tumor progression series, cell stiffness has been observed to increase with tumorigenic potential (Baker *et al.*, 2010). Our study brings additional insights to this question by characterizing tumor cells in various stages of confinement and invasion. As tumor cells experience increasing confinement, they markedly soften, consistent with the idea that deformability supports invasion and metastasis. Our results broadly agree with two recent studies. The first investigated cell traction forces in confined microenvironments and found that 3D confinement reduces cell traction forces (Raman *et al.*, 2013). The second examined migration of breast tumor cells within collagen "microtracks" and found that this confined geometry reduces basal FAs and SFs. Notably, the latter study found that confinement also increases matrix strain, which both complements and contrasts with our AFM stiffness measurements and implies that stiffness and contractility are not fully coupled (Mosier *et al.*, 2019).

A central enabling technology in our work is our open microchannel-based paradigm that allows AFM access for mechanical measurements during confined migration. This innovation addresses the technical challenge of obtaining contact-based mechanical measurements in a 3D-like environment. Our open-roof Y-shaped PDMS device combines the features of confined migration in 3D matrices, with the accessibility of two-dimensional supports to perform longitudinal mechanical measurements. By combining this device with AFM, we were able to follow cell migration from unconfined to confined spaces, while simultaneously measuring mechanical properties on the supernuclear region of the cell. The design of the platform also proved highly amenable to high-resolution optical imaging, including super-resolution imaging. Through these studies, we found that cells in unconfined geometries adopt a highly spread morphology with classical SF networks at the cell base and above the nucleus. As cells

from 2 to 100 μm (Supplemental Figures S6 and S7). Structured illumination microscopy (SIM) imaging revealed that cells closely adhere to the geometry of the microline, with greater aspect ratios observed on narrower microlines (Figure 5).

However, unlike in 3D-like topographical channels, AFM measurements indicated that cells do not appreciably soften on increasingly narrow microlines (Figure 6). Instead, cell stiffness reached a peak of 30-μm-wide lines (Young's modulus values increase from 4.1 to 7.2 kPa), where SIM images revealed prominent thick and parallel bundles of SF (Figure 5). Consistent with this result, YAP localized robustly to the nucleus for all microline widths (Figure 7A). The YAP nuclear/cytoplasmic (N/C) ratio fell only during topographical confinement, with no statistically significant width-dependent variation of the YAP N/C ratio for cells cultured on 1D microlines (Figure 7, B and C). This effect did not strictly correlate with nuclear elongation (Supplemental Figure S8). Thus, confinement-induced softening is an intrinsic effect of a 3D-like geometry and cannot be solely attributed to changes in cell or nuclear shape.
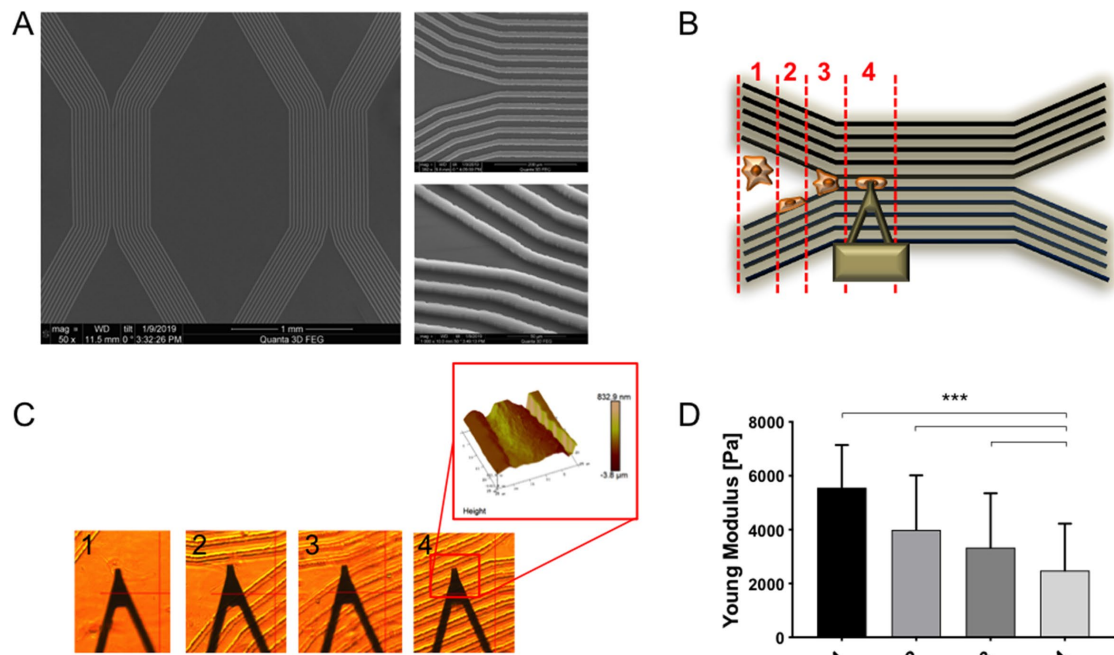
DISCUSSION

We have explored how tumor cells adapt their mechanical properties as they migrate through confined spaces, processes key to invasion and metastasis. Previous efforts to address this question

FIGURE 3: U2OS cells progressively soften as they progress from unconfined to fully confined migration. (A) SEM images of Y-shaped PDMS devices with different magnification and tilted angle. (B) Schematic of cells traversing Y-shaped channels, showing different cell categories: 1) unconfined, 2) associated with one wall, 3) associated with two walls but not fully confined, and 4) fully confined. (C) Optical images of AFM cantilever in contact with different cells and (D) AFM Young's modulus values of U2OS cells sorted in the four different categories. Each bar represents the median value of all the Young's modulus values collected on all the cells for each category, plotted with interquartile range as error bars. Statistical significance was determined using the Mann-Whitney test between each pair of categories (***, $p < 0.0001$; $n = 2560, 3072, 3072$, and 3840 force curves on 10, 12, 12, and 15 cells across three independent experiments. For each cell, a 16×16 force map was recorded on the supranuclear region of the cell body).

experience increasing confinement, they become increasingly elongated and exhibit increased actin density at the lateral device walls. Such cytoskeletal rearrangements have been observed in past microchannel studies (Balzer *et al.*, 2012) but without a clear function. Our study suggests that confinement induces the assembly of SFs at the channel walls, which could facilitate confined migration by exploiting the walls as "handholds" to build traction force and migrate, while simultaneously reducing their cortical stiffness. As a result, our work may eventually help reconcile conflicting views of how stiffness changes in tumorigenesis—namely, cells may become more deformable *and* generate greater contractile forces during confinement, thus simultaneously appearing to be both more and less stiff, depending on the measurement. Nonetheless, higher-resolution imaging and/or a modified channel system is still needed to definitively demonstrate that confinement induces SF assembly along the channel walls. It is also important to note that measures of cortical stiffness (e.g., AFM) are not equivalent to measures of traction force or contractility, even though the two are often strongly correlated (Wang *et al.*, 2002; Schierbaum *et al.*, 2019). Coupling our platform with finer intracellular mechanical measurements should help refine our conceptual model.

Another important outcome from our work is the direct correlation of mechanical changes during confined migration with functionally significant signaling events. In particular, for cells on flat, unconfined regions, cell stiffening is accompanied by strong nuclear YAP localization. With increasing confinement, cells soften, and YAP localizes to the cytoplasm. These results are consistent with previous work in which YAP nuclear localization closely tracks actomyosin assembly and contractility (Elosegui-Artola *et al.*, 2017) and confirm the role of YAP in mechanotransduction (Dupont *et al.*, 2011;

Raghunathan *et al.*, 2014; Nardone *et al.*, 2017). Previous studies observed that cellular elongation has a role in altering cytoarchitecture and YAP localization independently of topographical confinement (Bao *et al.*, 2017; Gegenfurtner *et al.*, 2018). Notably, YAP remained in the nucleus for cells on 1D microlines, indicating that cell softening and YAP function intrinsically depends on topographical confinement and is not simply being driven by changes in cellular elongation. In the future, it will be informative to quantify traction forces at the channel walls and investigate how SF distribution is related to YAP localization. Another interesting open question raised by our studies is which YAP-dependent genes are most strongly affected by confinement geometry and the extent to which the resulting gene products contribute to confined motility.

Our study also suggests a number of other interesting future directions. For example, it would be valuable to quantify changes in cell viscous (loss) modulus during confinement, in addition to the elastic properties measured here. It would also be informative to learn whether confinement-induced softening tracks in some interesting way with malignant potential, for example, using a progression series with well-matched genetic backgrounds. Finally, it will be important to ask whether our observations also hold in more tissue-like systems, including both organotypic culture models and animal models.
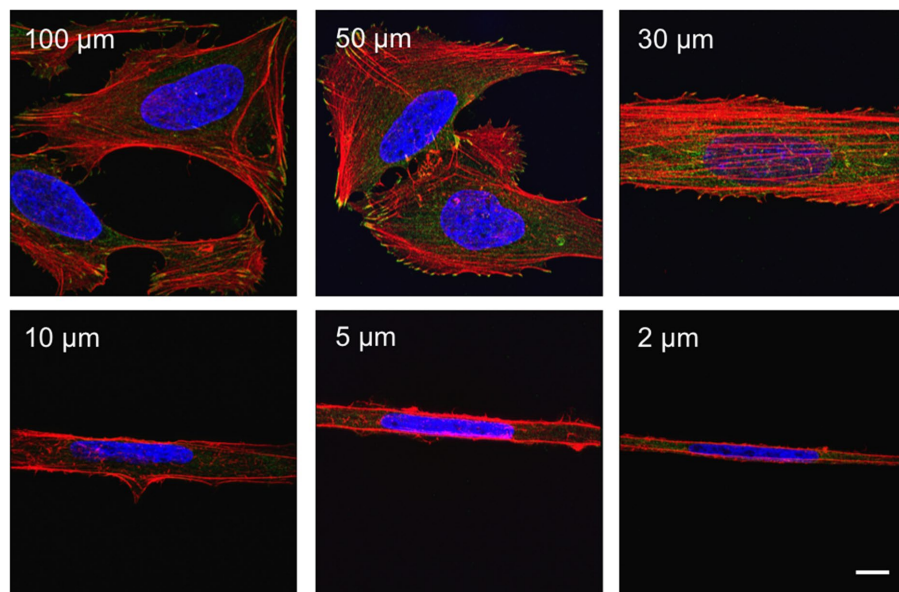
MATERIALS AND METHODS
Fabrication of patterned PDMS substrates
PDMS channels were created with photolithography and replica molding techniques (Xia *et al.*, 1997; Gates *et al.*, 2005). Photolithography masks were designed with AutoCAD and produced from Artnet Pro (San Jose, CA). A clean silicon wafer (WaferNet, CA) was
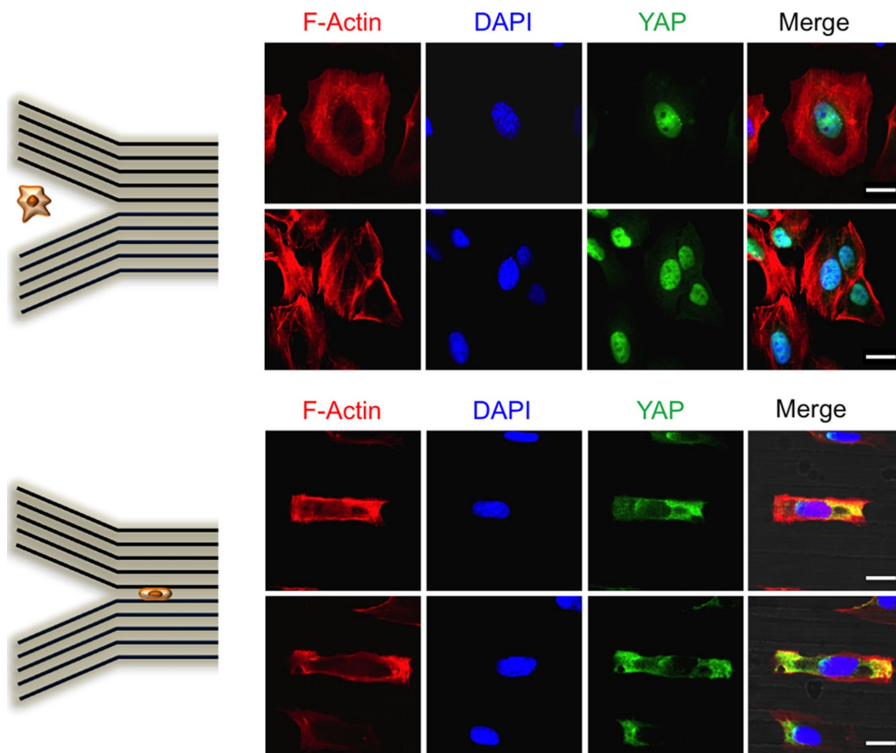
FIGURE 4: YAP localization is influenced by topographical cell confinement. Confocal images of U2OS cells fixed and stained for actin (red), nuclei (blue), and YAP (green). The top panel shows cells seeded on unconfined regions and the bottom panel shows cells in confined channels. Scale bars are 20 µm.

Midland, MI) and the prepolymer solution was poured onto the wafer. The wafer was degassed in vacuum and then prepolymer was cured for 2 h at 80°C (or overnight at room temperature [RT]). Finally, the PDMS was gently peeled off the wafer. Before proceeding with cell culture, PDMS substrates were extensively washed with ethanol, dried, and subjected to plasma treatment for 3 min to reduce hydrophobicity. Finally, substrates were incubated with fibronectin (FN) for 3 h before proceeding with cell seeding. U2OS cells were then seeded on PDMS substrates for 24 h, after which cells were either maintained as live cultures for AFM measurements or fixed and processed for fluorescence imaging.
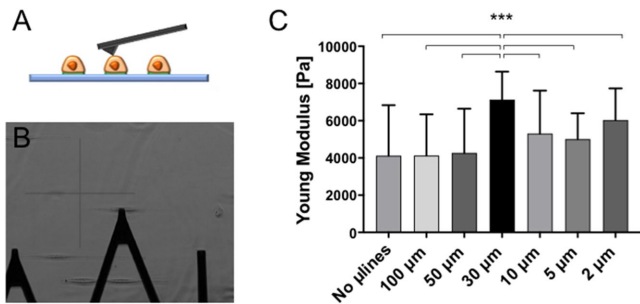
PEG passivation

To restrict cell adhesion to the channels, we passivated the surrounding mesas with PEG. A cover glass was coated with poly-L-lysine grafted to polyethylene glycol (PLL-g-PEG, Surface Solutions) for 1 h and then brought into close contact with the patterned PDMS for 15 min to allow PEG adsorption onto the mesas. The PEG-coated PDMS substrates were incubated with FN for 3 h, during which time FN adsorption was expected to be restricted to the channels due to the PEG passivation of the mesas.

spin-coated with SU-8 3005 photoresist (MicroChem, MA) to create a 5-µm-thick film. After soft baking at 95°C for 3 min, the film was exposed to UV light for 15 s through the mask, postbaked at 65°C (1 min) and 95°C (2 min). Finally, the wafer was developed and hardbaked at 200°C for 15 min. PDMS monomer and cross-linker were mixed (ratio 10:1; Sylgard 184 Silicone Elastomer Kit; Dow Corning,

Deep UV-based fabrication of FN microlines

Microlines were produced as previously reported (Lee *et al.*, 2018). Briefly, 18 × 18 mm cover glasses (Fisher Scientific) were incubated with 0.01 mg/ml PLL-g-PEG (Surface Solutions) in 10 mM HEPES for 1 h. After washing twice with phosphate-buffered saline (PBS) and deionized (DI) water (2 min each), PLL-g-PEG–coated glasses were illuminated for 15 min with deep UV light (UVO cleaner; Jelight) through a chrome–quartz photomask encoding multiple microlines ranging from 2 to 100 µm width. After illumination, cover glasses were left in DI water until use in cell culture. Before cell culture, cover glasses were incubated in fibronectin for 3 h at 37°C and then extensively washed with PBS.

Cell culture

U2OS human osteosarcoma cells and U87 human glioblastoma cells were cultured in DMEM supplemented with 10% fetal bovine serum (FBS) and 1% penicillin–streptomycin. Cells were incubated at 37°C in a humidified atmosphere of 95% air and 5% CO_2. All substrates used in this study were incubated with fibronectin for 3 h and then with DMEM medium for 20 min before proceeding with cell seeding. Cells were tested for mycoplasma every 3 mo and authenticated via short tandem repeat profiling.

FIGURE 5: 1D microlines strongly affect U2OS morphology. SIM images of U2OS cells seeded on FN microlines ranging from 2 to 100 µm width. Cells were fixed and stained for actin (red), vinculin (green), and nuclei (blue). Scale bar is 10 µm.

FIGURE 6: U2OS mechanics are only modestly affected by 1D microline width. (A) Schematic representation of an AFM cantilever engaging a cell seeded on 1D microlines of fibronectin. (B) MLCT-Bio cantilever (lever C) indenting living cells. (C) Young's modulus values of U2OS cells seeded on microlines with widths ranging from 5 to 100 μm. Each bar represents the median value of all the Young's modulus values collected on all the cells for each category, plotted with interquartile range as error bars. Statistical significance was determined using the Mann-Whitney test between each pair of categories (***, $p < 0.0001$; $n = 2048$, 2560, 2048, 2816, 3072, and 3072 force curves on 8, 10, 8, 11, 12, and 12 cells across six independent experiments. For each cell, a 16 × 16 force map was recorded on the supranuclear region of the cell body).

Immunofluorescence staining

Cell were fixed in 4% (vol/vol) paraformaldehyde (Alfa-Aeser) for 15 min at RT, permeabilized with 0.1% (vol/vol) Triton X-100 (EMD Millipore) for 3 min, and blocked with 1% (vol/vol) bovine serum albumin (BSA; Thermo Fisher) for 30 min in a humid chamber. Cells were incubated with primary antibodies for 2 h, rinsed with 1% BSA in PBS, and then incubated with secondary antibodies and phalloidin (3:500; Thermo Fisher Scientific) for 30 min in a humid chamber. The following primary antibodies were used for immunostaining: mouse anti-vinculin hVin-1 (1:1000; Sigma) and YAP (D8H1X) XP Rabbit mAb (1:1000; Cell Signaling). As secondary antibodies we used Alexa Fluor 488 goat anti-mouse (1:500) and Alexa Fluor 488 goat anti-rabbit (1:500), both from Life Technologies. Finally, cells were incubated with NucBlue Live ReadyProbes Reagent (Thermo Fisher Scientific) for 20 min in PBS to stain nuclei. Samples were rinsed in PBS and stored in PBS at 4°C in the dark.

Confocal and structured illumination microscopy

Samples were imaged with a Prairie Technologies swept field confocal microscope using an Olympus LUMPlanFL N 60×/1.0 water dipping objective and a Zeiss Elyra PS.1 structured illumination microscope (Zeiss) with a DIC M27 63×/1.4 oil objective (Zeiss).

Image analysis

Quantification analysis of FAs, nuclei, and YAP nuclear/cytoplasmic ratio was performed using the open source software Fiji (Schindelin *et al.*, 2012). Morphometric analysis of FAs was performed after processing images with a brightness/contrast tool, followed by the threshold command to obtain binarized images, and finally subjected to particle analysis. The orientation of FAs was calculated with ($\theta_{FAs}-\theta_{pattern}$), such that a smaller value indicates a stronger orientation to the pattern axis. Nuclear area and aspect ratio were calculated after processing the images with the threshold and particle analyzed, measuring area and shape descriptors. Finally, YAP
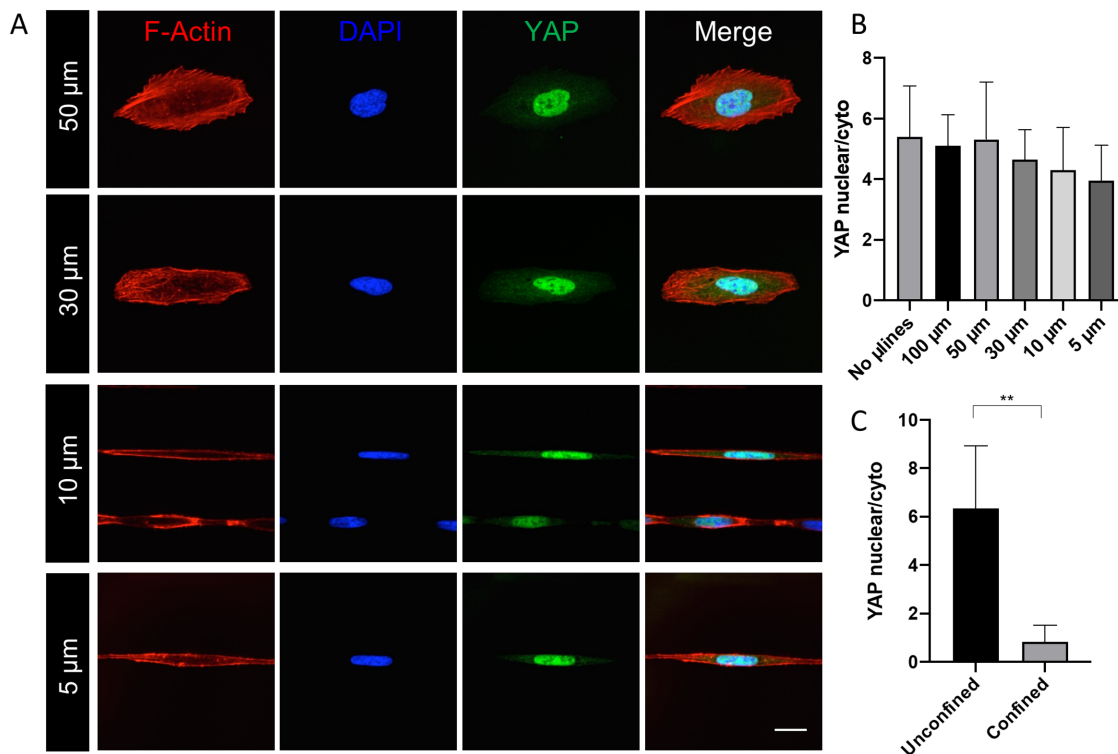
FIGURE 7: YAP localization is not affected by 1D microline width. (A) Confocal images of U2OS cells seeded on FN microlines with different widths (5–50 μm). Cells were fixed and stained for actin (red), nuclei (blue), and YAP (green); merged colors are shown on the rightmost micrographs. Scale bar is 30 μm. (B) Quantification of YAP nuclear/cytoplasmic ratio in cells cultured on FN microlines and (C) on unconfined and confined regions of a Y-shaped PDMS device. Error bars are SD of mean values ($n = 10$ across two independent experiments; **, $p < 0.001$, calculated with the Mann-Whitney test).

nuclear/cytoplasmic ratio was calculated as the ratio between the intensity values for the pixels in the nuclear and in the cytoplasmatic regions of each cell. Both intensity values were divided by the area of the corresponding regions.

Scanning electron microscopy (SEM)
Samples were sputter-coated with Au-Pa for 50 s (85 Å thickness). A FIB-SEM (Zeiss) was used to image surface topography of patterned PDMS substrates.

AFM measurements
AFM measurements on living cells were performed with a Bruker Catalyst (CA, USA) with Bruker MLCT-Bio cantilevers (lever C). To acquire AFM images, MLCT-Bio tips were used in contact mode with a scan rate of 0.5–1 Hz. For cell mechanics measurements, cantilevers were calibrated with the thermal tune method (Hutter and Bechhoefer, 1993) before each experiment, and force volumes were obtained using a 50 nm (0.5 nN) trigger threshold to limit indentations. Arrays of force curves (force maps) were recorded at the central and highest part of the cell body to avoid any thin peripheral regions, which are more susceptible to contributions from the underlying substrate (Domke *et al.*, 2000). Force maps were recorded on 5 µm × 5 µm areas and obtained in a 16 × 16 grid (256 force curves). More than 4000 force curves were obtained for each cell category.

AFM data analysis
The data analysis package IGOR (Wavemetrics, Lake Oswego, OR) was used to analyze AFM data and to extract the Young's or elastic modulus (E). Force-indentation curves were fit to the Hertzian model for pyramidal probes for every force curve within a force map. For each 16 × 16 force map, 256 Young's modulus values were calculated. Finally, the median of E values, calculated from all the force curves, were considered representative of each cell category. Every graph reporting AFM data is presented as a bar graph, reporting the median values of all force curves with interquartile range as error bars.

Statistical analysis
Statistical analysis and graphing were performed in Graph-Pad Prism (v 7.00). The Mann-Whitney test, a nonparametric test for non-Gaussian distributions, was used to assess statistical differences. For AFM mechanical data, two categories at a time were compared with each other, to study statistical differences between all of them. For each category, a range of 8–15 cells were studied, for each of which one force map (made of 256 force curves) was recorded, yielding 2048–3840 force curves per category. Experiments were independently repeated three to six times, with measurements obtained across multiple samples. Additional statistical details are specified in the figure captions.

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