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Characterizing Cell Adhesion by Using Micropipette Aspiration

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ABSTRACT   We have developed a technique to directly quantify cell-substrate adhesion force using micropipette aspiration. The micropipette is positioned perpendicular to the surface of an adherent cell and a constant-rate aspiration pressure is applied. Since the micropipette diameter and the aspiration pressure are our control parameters, we have direct knowledge of the aspiration force, whereas the cell behavior is monitored either in brightfield or interference reflection microscopy. This setup thus allows us to explore a range of geometric parameters, such as projected cell area, adhesion area, or pipette size, as well as dynamical parameters such as the loading rate. We find that cell detachment is a well-defined event occurring at a critical aspiration pressure, and that the detachment force scales with the cell adhesion area (for a given micropipette diameter and loading rate), which defines a critical stress. Taking into account the cell adhesion area, intrinsic parameters of the adhesion bonds, and the loading rate, a minimal model provides an expression for the critical stress that helps rationalize our experimental results.

INTRODUCTION

Cell adhesion is involved in a plethora of physiological and pathological cellular processes, and cell-substrate adhesion affects several cellular functions, such as migration, proliferation, and differentiation. Molecules involved in cell-substrate adhesion have been thoroughly studied (1), yet quantitative experimental techniques are still needed to provide a better understanding of the mechanics of cell adhesion. In the context of atherosclerosis, the early stage of the disease involves the transmigration of monocytes in abnormally high numbers, which then accumulate in arterial walls and participate in the formation of atherosclerotic plaques (2). Measuring changes in the adhesion of endothelial cells should help shed light on this process, as mechanical changes in endothelial cells after monocyte adhesion might play a crucial role in pathologically elevated transmigration rates. Indeed, results of a study by Kataoka et al. suggest that in vitro, the contact between white blood cells and endothelial cells modifies the strength of endothelial cell adhesion to their substrate (3). The authors found indirect evidence of a reduced adhesion in the presence of white blood cells by measuring a change in the electrical resistance of a population of endothelial cells. They interpreted the observed decrease of electrical resistance in the presence of white blood cells as evidence of a nanometer-scale increase in the distance between the ventral cell surface and the substrate. However, a more direct characterization of the mechanical properties of endothelial cells is still needed to better understand their response to leukocyte adhesion.

Although the behavior of red blood cells detaching from a surface has been experimentally and numerically investigated while monitoring the applied force (4,5), and though several techniques exist to measure the adhesion of cells adhering weakly to a substrate or to another cell, data are scarce on strongly adherent cells such as endothelial cells. Previous techniques at the single-cell level relied on pulling a micro- or nanoneedle inserted into the cell body (6–8), pulling (9,10) or scraping (11,12) the cell of interest with a bending cantilever, and aspirating the cell with a micropipette (13–16). Experiments were also carried out on cell populations in which cell detachment was induced by centrifugation or hydrodynamic stress (17,18), yielding ensemble-averaged values for the quantities of interest. However, some of these studies involve technical difficulties, such as the need for protein scaffolding of the cells to avoid membrane rupture when they are pulled by a nanoneedle (8). Moreover, knowing the detachment force is not always straightforward. For instance, in the case of aspirating micropipettes used as cantilevers (9,10), the detachment force measured from the cantilever deflection may not be independent of the aspiration pressure used to hold the cell and micropipette in contact, as other experiments with static micropipettes indicate that a high aspiration pressure is enough to detach a cell without having to bend the micropipette (13). In turn, in the latter work, the micropipette was not held in direct contact with the cells, so that the flow of medium into the pipette is expected to have contributed to the detachment force; indeed, in a related study, this was taken into account using fluid-flow simulations (16).

This work presents, to our knowledge, a new technique designed to investigate the detachment of endothelial cells

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adhering to a substrate. This method allows for direct measurement of the detachment force by placing an aspirating micropipette in contact with the surface of an endothelial cell so that a seal is formed and fluid flow into the micropette can be neglected. We can perform detachment assays on a number of individual cells in a single experimental run, thus exploring a range of cell sizes, while imposing the micropipette diameter and the aspirating pressure rate (or loading rate). Finally, interference reflection microscopy (IRM) is used to monitor the adhesion areas, as was previously done in the case of adhering blood cells (9) and tumor cells (18). In a nutshell, this technique enables us to fully probe the influence of geometric parameters, such as cell surface, adhesion areas, and micropipette size, as well as dynamical parameters such as the loading rate. We present a minimal model that captures the features of the mechanical behavior observed in our experiments.

MATERIALS AND METHODS

Endothelial cells

Primary bovine aortic endothelial cells were kindly provided by A.-C. Vion and C. Boulanger and used between passages 4 and 11. They were cultured in Dulbecco's modified Eagle's medium (Invitrogen, Carlsbad, CA) supplemented with 10% fetal bovine serum (Invitrogen) and 1% penicillin/streptomycin (Invitrogen) at 37°C with 5% CO_2. The cells were passaged three times a week and resuspended in fresh culture medium. One to two days before each experiment, the cells were trypsinized with trypLE (Invitrogen) and plated at a subconfluent density onto thin-bottom dishes (standard-bottom low-walled 35-mm μ-Dish, IBIDI, Martinsried, Germany). In another set of experiments, the cells were grown on glass microbeads instead of a petri dish. Typically, 50 dextran Cytodex-3 microcarrier beads (Sigma-Aldrich, Taufkirchen, Germany) were first deposited at the bottom of a μ-Dish in phosphate-buffered saline (PBS; Invitrogen). The PBS was then removed and ~10,000 trypsinated bovine aortic endothelial cells were introduced into the μ-Dish.

Before any experiment, a μ-Dish was brought onto the stage of the microscope, where experiments were performed, at room temperature in cultured medium with 20 mM HEPES (Invitrogen) added, on cells that had been plated onto the petri dish a few hours to a few days earlier. Dependence of the detachment force on the length of time after plating was not examined in this study. For experiments in which the cells were exposed to cytochalasin D, the cells were incubated for 30 min in a solution containing 1 μg/mL cytochalasin D from *Zygosporium masonii* (Sigma-Aldrich). The solution containing the cytochalasin D was then aspirated, and fresh medium was introduced. When treated with nocodazole, cells were incubated for 1 h in culture medium containing 10 μg/mL of nocodazole (Sigma-Aldrich). We verified using live-cell microtubule staining (tubulin tracker green, Life Technologies, Saint Aubin, France) that in these conditions, microtubules were properly destabilized.

Micropatterns

Micropatterned coverslips were prepared as described by Azioune et al. (19). The coverslips were first sonicated in ethanol and plasma-treated. They were then incubated in 10 mM HEPES buffer in the presence of 0.1 mg/mL of polylysine-grafted polyethyleneglycol (PLL-g-PEG; Surface Solutions, Dübendorf, Switzerland). After washing with PBS and water, the coverslips were illuminated with deep ultraviolet (UV) light (UVO Cleaner,

342 Series, Jelight, CA) through a photomask and then incubated for 1 h in 100 mM NaHCO$_3$ in the presence of 50 μg/mL fibronectin. The cells were then plated onto the coverslips, which were mounted on magnetic chambers (Chamlide, Live Cell Instrument, Seoul, South Korea) before each experiment.

Micropipette fabrication

Borosilicate glass capillaries (1 mm outside diameter, 0.78 mm inside diameter, Harvard Apparatus, Holliston, MA) were pulled on a P-97 micropipette puller (Sutter Instruments, Novato, CA). To fabricate the micropettes used in cell aspiration, a MF-900 microforge (Narishige, Tokyo, Japan) was used to cut the extremity of pulled capillaries to the desired diameter, ranging from 4.6 to 30 μm. The body of the micropipettes was then bent at a 45° angle so that their extremity could be held perpendicular to the plane of cell adherence. The micropipettes used to hold the Cytodex-3 beads were fabricated in the same fashion but with a 50-μm diameter at the tip.

Microscope setup

The setup was mounted on a TE300 inverted microscope (Nikon, Tokyo, Japan) located on an air suspension table (CVI Melles Griot, Didam, Netherlands). It was equipped with a 100× oil immersion, 1.3 NA objective for experiment monitoring (Nikon), and lower magnification objectives (40×, 20×, 10×, 4×, and 2×, Nikon Instruments) for micropipette positioning. Images were acquired with a Flash4.0 CMOS camera (Hamamatsu Photonics, Hamamatsu City, Japan) controlled using the software Micromanager (20). The experiments were performed using either brightfield microscopy or IRM. Timelapse movies were acquired at a rate of 1 frame/s under a 100 ms exposure time for IRM.

Interference reflection microscopy

In performing IRM, we used a filter cube equipped with a dual-edge dichroic mirror and a dual-band bandpass emission filter (505/606-nm BrightLine and 524/628-nm BrightLine, respectively, Semrock, Rochester, NY). As an illumination source, we used an Intensilight (Nikon) with an ND8 neutral density filter, and no excitation filter. The polarization of the incoming light was assured by the dichroic mirror, but our illumination was not strictly monochromatic due to the dual-band emission filter. Raw images were processed to remove their background using the software ImageJ (21). See the Supporting Material for details on image processing. To validate this configuration, we tested the setup using 2.2-mm glass beads (Dominique Dutscher, Brumath, France) placed at the bottom of a petri dish in the presence of cell culture medium (see the raw picture obtained in Fig. 1 A). The distance between the surface of the glass bead and the surface of the petri dish is given by $h(d) = R - \sqrt{R^2 - d^2}$, where $R = 1.1$ mm is the radius of the bead and d is the distance from the bead center projected on the dish plane, $0 \le d \le R$ (Fig. 1 A, *inset*). We measured the positions of the local intensity extrema as a function of h and verified that they follow the established equation (18,22)

$$I = \frac{I_M + I_m}{2} - \frac{I_M - I_m}{2} \cos\left(4\pi \frac{\delta}{\lambda}\right), \qquad (1)$$

where I_M and I_m are the maximum and minimum intensity levels, respectively, and $\delta = h(d)$ (Fig. 1 B). The best fit was obtained for a characteristic wavelength of $\lambda = 340$ nm.

When performing experiments with continuous IRM imaging, we qualitatively observed that the illumination seemed to affect the cell mechanics. Indeed, a primary detachment event occurred during which a significant part of the cell detached, but the pipette tip then had to be slightly translated in the xy plane to detach the remaining part of the cell (Movie S2). During
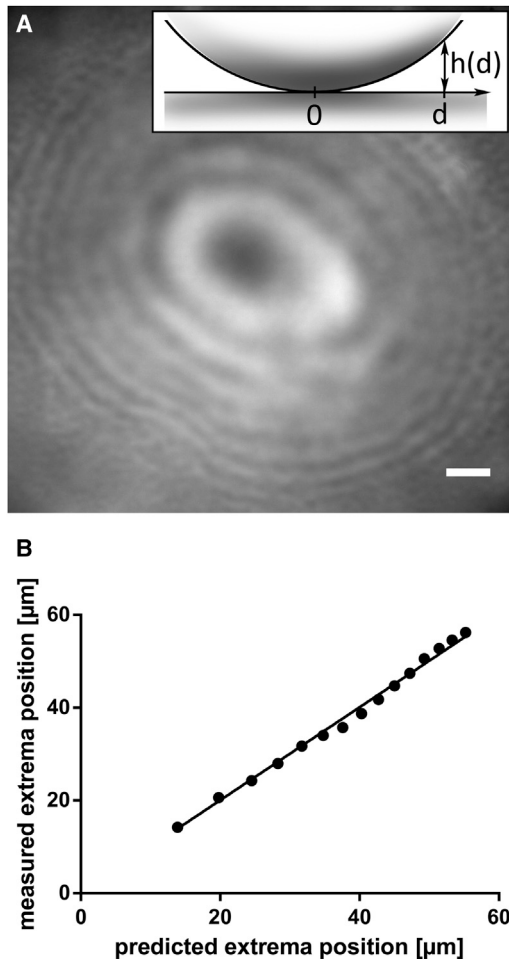
FIGURE 1 Determination of the IRM wavelength. (*A*) IRM image showing the intensity minima and maxima (from destructive and constructive interference, respectively) of the light refracted from a 2.2-mm glass bead. Scale bar, 10 μm. (*Inset*) For each point on the bead surface, the distance to the petri dish is $h(d) = R - \sqrt{R^2 - d^2}$, where $R = 1.1$ mm is the bead radius and d is the projected distance from the center of the bead. (*B*) Plot of the experimental intensity extrema positions versus the predictions from Eq. 1 (18,22), for $\lambda = 340$ nm. The good fit for this value of λ is evidenced by the slope of the linear regression: 1.003 ± 0.007.

our experiments under brightfield illumination, we also observed cell detachment, but no additional micropipette motion was necessary to fully detach the cells. We hypothesize that this change in cell mechanics is due to the UV light used in our IRM setup. Under continuous illumination, the cells are exposed to a very large amount of UV light, which likely causes phototoxic damage. In most experiments, however, we only used IRM to take a snapshot of the adhesive areas in the initial state and thus expect cell damage to be minimal.

Micromanipulators

The microscope was equipped with a motorized micromanipulator carrying a first micropipette holder at a 45° angle, and a manual three-axis stage linked to a UT-2 joint to orient a second micropipette holder (MP285 micromanipulator, Sutter Instruments, Novato, CA; IM-H1 micropipette holders and UT-2 joint, Narishige, Tokyo, Japan; three-axis stage, Thorlabs, Newton, NJ). The first micropipette was used to aspirate adherent endothelial cells, whereas the other was used to hold Cytodex-3 beads.

RESULTS AND DISCUSSION

A constant-rate aspiration technique for cell-detachment assays

We have developed, to our knowledge, a new technique to apply a well-controlled aspiration force to a single endothelial cell adhering to a substrate while quantitatively monitoring the detachment mechanics. We impose an aspiration pressure, ΔP, at a constant rate, r_P, via a micropipette held in contact with the cell body, its tip perpendicular to the substrate (Fig. 2). The aspiration force applied to the cell, $F = \Delta P S_{pipette}$, is readily known, since we control both the aspiration pressure, ΔP, and the pipette section, $S_{pipette}$. Two different setup configurations allow us to visualize and follow the detachment mechanics either in bottom or side view, which we refer to as in-plane and profile modes, respectively.

In the first configuration, or in-plane mode, we designed our setup to be able to work directly with cells cultured on the bottom of a thin-bottom petri dish: this allows us to test a significant number of cells in a single run of experiments (up to 100/day). The petri dish is placed above the microscope objective, parallel to the observation plane. An aspiration micropipette fabricated as described above is micromanipulated so that its tip is brought into contact perpendicular to the luminal surface of the adherent endothelial cells (see Fig. 2 *A* and Movie S1). Monitoring the detachment in the substrate plane allows us to measure the projected cell area over time (see Fig. 2 *B*). Using IRM also gives access to evolution of adhesion areas (see Fig. 4 *A*).

To better understand the detachment process, we performed complementary experiments in a second configuration, the profile mode. In this case, endothelial cells adhering to a microsphere held by a second micropipette are aspirated in a direction parallel to the observation plane, enabling us to observe the detachment from the side (see Fig. 2 *D*). More precisely, the cells were grown on 200-μm dextran Cytodex-3 microspheres initially designed to culture adherent cells in agitation. We used a 50-μm-diameter micropipette to firmly hold a Cytodex-3 bead. We positioned the bead to place endothelial cells perpendicular to the equatorial plane of the microsphere (Fig. 2 *D*, *inset*), which allowed us to visualize a single endothelial cell in profile. We then brought the aspiration micropipette into contact with the endothelial cell, perpendicular to its surface (Fig. 2, *D* and *E*). This visualization method does not give access to the projected cell area, but it provides additional insight into the cell-detachment dynamics.

Detachment force and geometric parameters

The time evolution of the projected cell area is shown in Fig. 2 *B* for three different cells. Although the initial area covered by the cells varies, each case follows a qualitatively similar scenario: the projected area is constant over time
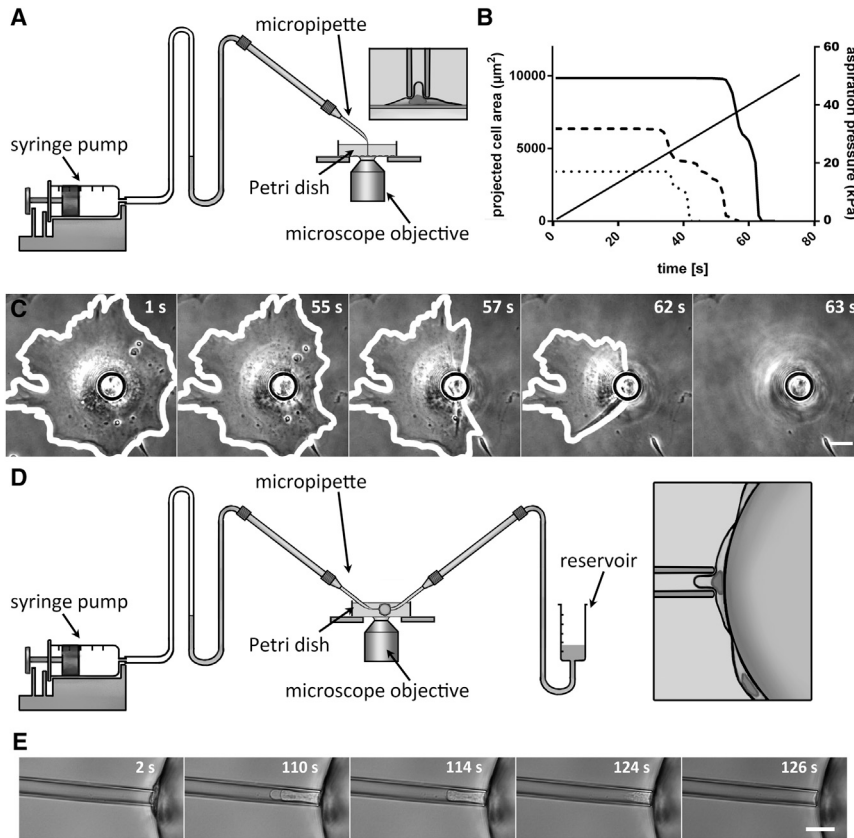
FIGURE 2 Setup: in-plane (*A–C*) and profile modes (*D* and *E*). (*A*) Experimental setup for in-plane aspiration experiments. In this case, the micropipette is positioned perpendicular to the surface of adherent endothelial cells cultured on the bottom of a petri dish. The syringe pump on the left creates a constant-rate aspiration pressure increase. (*B*) Plot of the projected cell area versus time for three different cells. The projected cell area is measured from pictures taken throughout the detachment. (*C*) Time-lapse of a cell throughout a detachment assay (corresponding to the solid line in *B*, and to the cell on the right in Movie S1). Scale bar, 10 μm. (*D*) Experimental setup for profile aspiration experiments. In this case, the aspiration micropipette is still positioned perpendicular to the surface of the endothelial cells, but they adhere to a Cytodex-3 bead, held in position by a second, larger micropipette (on the right). (*E*) Time lapse of the detachment assay of an endothelial cell adhering to a Cytodex-3 bead. Scale bar, 10 μm.

until a breaking point when the projected area rapidly collapses until the cell is fully detached, at a critical aspiration pressure, ΔP^*. This indicates that the detachment of cells from their substrate is a well-defined event in our assay. Moreover, for a constant micropipette diameter, we find that larger cells require a higher aspiration pressure, i.e., a higher aspiration force, $F^* = \Delta P^* S_{\text{pipette}}$, to detach from the substrate.

To further quantify the relation between projected area and detachment force, we performed in-plane detachment experiments on 325 cells (over several dishes and days). We found the detachment force, F^*, to be, at a first approximation, proportional to the initial projected cell area, S_{cell} (Fig. 3 *A*). We can then write

$$F^* = \sigma^* S_{\text{cell}}. \tag{2}$$

A linear fit of the experimental data yields a critical stress of $\sigma^* = 1300 \pm 50$ Pa. All points in this data set collapse on the master curve from Eq. 2, regardless of cell size or pipette diameter. Indeed, both S_{cell} and S_{pipette} vary widely here: S_{cell} ranges from 775 μm^2 to 22,000 μm^2 and S_{pipette} from 16.6 μm^2 to 707 μm^2 (which corresponds to pipette radii of 2.3–15 μm). Conversely, we found ΔP^* to depend on S_{pipette}, as shown in Fig. 3 *B*. We thus conclude that F^*, not ΔP^*, is the relevant observable quantity that characterizes cell detachment.

Since cells resist an aspiration force via cell-substrate adhesions, we used IRM to investigate the relationship between detachment force and adhesive areas at the cell-substrate interface. Indeed, these adhesive areas appear as dark patches in IRM images taken in the in-plane mode (18,22,23). By monitoring the evolution of these adhesive areas over the duration of a complete detachment event, we found that the detachment does not occur by peeling from the periphery but rather appears to be initiated in the region directly under the micropipette, see Movie S2 and Fig. 4 *A*. These experiments lead us to assume that the adhesive areas share the pulling force throughout the aspiration assay and remain stable until rupture. However, a quantification of these experiments during the detachment process is difficult, as the cell membrane areas that are pulled further than half a wavelength (i.e., 170 nm) away from the substrate become as dark as the adhering areas. In what follows, we consider the initial adhesive area, S_{adh}.

We relate the rupture force, F^*, to the initial adhesive area, S_{adh}, instead of the initial projected cell area, S_{cell}, by expressing F^* as

$$F^* = \sigma^{**} S_{\text{adh}}, \tag{3}$$

which defines a second critical stress, $\sigma^{**} = \frac{\sigma^*}{S_{\text{adh}}/S_{\text{cell}}} = 16700 \pm 5600$ Pa ($N = 16$) (see Fig. 4 *B*). The rupture force, F^*, scales with both S_{cell} and S_{adh}, and consistently,
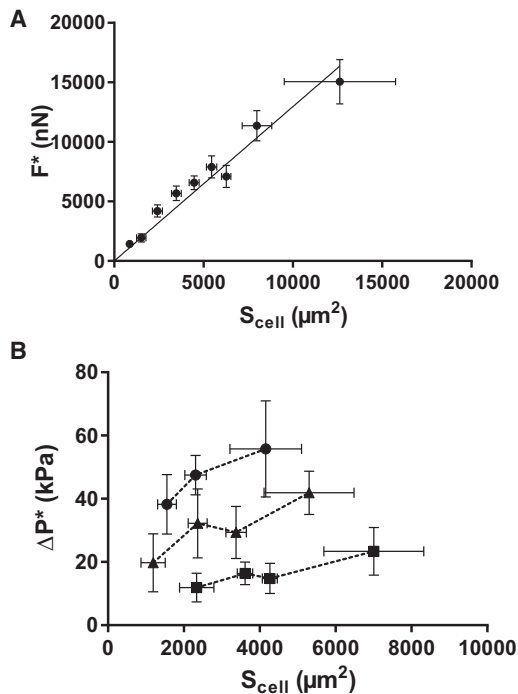
FIGURE 3 Critical stress. (A) Plot of binned experimental data showing that the detachment force scales linearly with the initial projected cell area, S_{cell}, implying that a critical stress, $\sigma^* = 1300$ Pa ± 50 Pa, induces cell detachment. Error bars show the standard deviation. $N = 335$ experimental data points. (B) Critical aspiration pressure, ΔP^*, versus projected cell area for different micropipette diameters. ΔP^* depends on the micropipette section, $S_{pipette}$, used in the experiments ($\bullet$: $S_{pipette} \in [17\ \mu m^2, 72\ \mu m^2]$, $\blacktriangle$: $S_{pipette} \in [78\ \mu m^2, 275\ \mu m^2]$, $\blacksquare$: $S_{pipette} \in [314\ \mu m^2, 707\ \mu m^2]$). Error bars show the standard deviation.

S_{cell} and S_{adh} are correlated (Spearman's correlation $r = 0.54$, $N = 16$ cells), with $S_{adh}/S_{cell} = 0.09 \pm 0.04$ ($N = 16$ cells, mean $\pm$ SD).

The scaling of F^* with S_{adh} is easier to interpret, since the resistance to the aspiration force comes from these adhesion areas. However, the correlation between S_{adh} and S_{cell} does not allow us to conclude whether S_{adh} is truly a more relevant parameter than S_{cell}. To constrain S_{cell}, we cultured cells on circular micropatterns of different sizes, so that S_{cell} was imposed by the disk shape ($S_{cell} = 520 \pm 30\ \mu m^2$ on 25-μm

patterns, $N = 12$; $S_{cell} = 760 \pm 210\ \mu m^2$ on 30-μm patterns, $N = 10$). We took advantage of the fact that S_{adh} still varied, $S_{adh}/S_{cell} = 0.16 \pm 0.05$ (mean $\pm$ SD), to study how the detachment force, F^*, relates to S_{adh}. Interestingly, when detaching cells from the micropatterns, F^* still scaled linearly with S_{adh} as in Eq. 3, and the values obtained for the critical stress, σ^{**}, were comparable to those obtained without micropatterns: $\sigma^{**} = 16,700 \pm 5600$ Pa without a micropattern ($N = 16$) versus $\sigma^{**} = 20,300 \pm 5500$ Pa on micropatterns ($N = 12$). Note that to account for the fibronectin thickness of the micropatterns, we considered a 60 nm threshold to define the adhesion areas on the IRM pictures—instead of 50 nm on petri dishes. Consequently, a single IRM picture of the cell-substrate interface should be sufficient to compute the aspiration force necessary to detach the cell at a given loading rate.

Membrane rupture does not entail cell detachment

The complete detachment process was also monitored in profile mode, in which the aspirated endothelial cell was visualized from the side under brightfield illumination (Fig. 2, D and E). We found that the cell membrane appeared to break systematically before detachment (Movie S3). To assess whether the membrane rupture was a triggering event for cell detachment, we performed in-plane experiments in the presence of propidium iodide in the surrounding medium. Propidium iodide is an intercalating agent that becomes fluorescent when it gains access to intracellular nucleic acids: membrane rupture is thus indicated by a sharp increase in fluorescence (see Movie S4). We measured the force at which the membrane broke for several pipette radii and compared it to the detachment force predicted by Eq. 2 (with $\sigma^* = 1300$ Pa). This computed detachment force was systematically higher than the force at which the membrane was observed to break, i.e., membrane rupture always occurred before cell detachment within this range of pipette radii (Fig. 5). In profile mode, we ran a second series of experiments in which we stopped aspirating and waited for 2 min immediately after membrane rupture. None of the
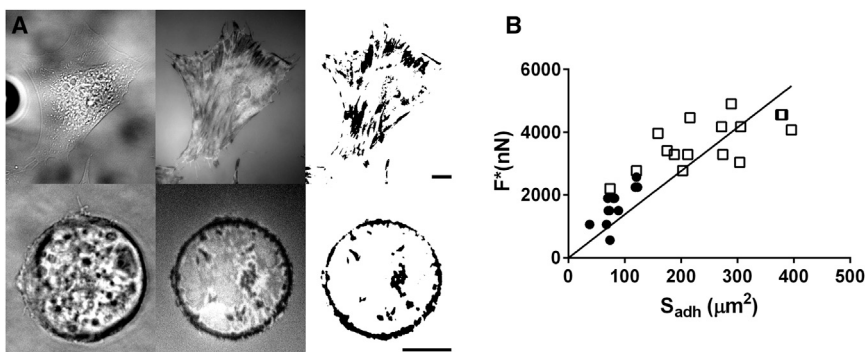

FIGURE 4 Detachment force scales with the adhesion area obtained from IRM images. (A) Each row shows, from left to right, the brightfield picture, the IRM picture, and the binary picture, with the adhesion patches shown in black. (Upper) Cell adhering to an unpatterned petri dish. (Lower) Cell constrained on a circular micropattern. Scale bars, 10 μm. (B) Detachment force versus adhesion area. The detachment forces obtained with ($\bullet$) and without ($\square$) micropatterns scale with the adhesion area (the solid line shows the linear fit of both data sets).
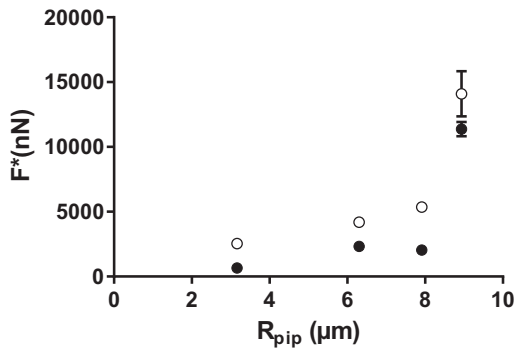
FIGURE 5 Aspiration force at membrane rupture and detachment force. The force at rupture (●), based on experimental data, and the detachment force (○) are predicted using Eq. 2. The membrane systematically breaks before the cell detaches from the substrate. Error bars represent the standard error.

10 cells we tested detached during that period of time. At the end of this waiting period, we aspirated these cells again and verified that they could still resist a significant amount of aspiration before detaching. We ran a set of complementary experiments in in-plane mode. We applied an aspiration pressure of 5.5 kPa on cells in the presence of propidium iodide, until the rapid increase in fluorescence showed that the membrane was ruptured. As soon as rupture was detected, we set the aspiration pressure back to zero while leaving the micropipette in place, and we waited for up to 300 s. During the whole process, we imaged the cell-substrate interface with IRM at a rate of one image every 4 s. The quantification of these experiments showed that S_{adh} stayed relatively constant even in the presence of a broken membrane (see the Supporting Material, Fig. S3, and Movie S5).

To test whether detachment force would still scale with S_{cell} in the absence of membrane rupture, we grew endothelial cells on a low-adhesion surface consisting of a glass coated with a low concentration of PLL-g-PEG molecules (see the Supporting Material). When left for 2 h to adhere, cells developed a limited S_{cell}. Detachment force in these conditions still scaled with both S_{adh} and S_{cell} (Fig. S4). The critical stress, $\sigma^* = 1500 \pm 500$ ($N = 9$ cells), was very consistent with the one obtained on glass substrates, although $\sigma^{**} = 4300 \pm 2300$ Pa was significantly lower (see the Supporting Material for further discussion). We ran another set of detachment experiments on cells grown on PLL-g-PEG in the presence of propidium iodide, and they showed that for the majority of cells (17 of 20 cells tested), the cell membrane did not break before detachment. These sets of experiments strongly suggest that membrane rupture does not necessarily lead to cell detachment, and that it has no influence on the scaling of detachment force with cell projected area. This suggests that membrane integrity is not essential for transmitting forces to the cell-substrate interface. The consideration that membrane rupture does not seem to influence the rupture force, together with

our IRM observation of cells detaching first in the region directly beneath the micropipette instead of peeling from the outer rim toward the center, led us to conclude that the aspiration force is not transmitted to the substrate through membrane tension but rather through the whole cell body. In what follows, we then make the simplifying assumption that the adhesive bonds share the aspiration force in parallel.

Existence of intact adhesion area after cell detachment

As can be observed in Movie S2, some traces that appear dark in the IRM images can be seen after cell detachment. We examined whether the presence of what we will henceforth call dark traces was systematic. Indeed, after the detachment of each of nine cells, although these traces were hard to observe in brightfield illumination, they were systematically present in IRM (see Fig. S7). To investigate the nature of the dark traces, we performed immunostaining of the substrate after cells were detached and found that the dark traces were rich in vinculin (see the Supporting Material and Fig. S8). We quantified the area, S_{adh}^{after}, of these dark traces relative to S_{adh} measured before detachment. By using the same thresholding procedure of IRM images (see the Supporting Material) before and after detachment on $N = 13$ cells, we obtained $S_{adh}^{after}/S_{adh} = 0.06 \pm 0.07$. Thus, on average, 94% of all adhesive areas detach during cell detachment, so we made the simplifying assumption that we could neglect the potential contribution of the undetached adhesion area. However, this raised an interesting question about the process occurring locally to these remaining adhesion areas during cell detachment. One possible scenario is that at these locations, the membrane broke instead of adhesive bonds. We tried to test this hypothesis by running detachment experiments in the presence of propidium iodide while acquiring time lapses. We could not detect such local membrane rupture, and we only observed a major rupture close to the micropipette tip (as shown in Movies S2 and S5). These nanoscale membrane ruptures could still exist, but get repaired too quickly (24), thus avoiding diffusion of propidium inside the cell.

Dynamic force spectroscopy of cell adhesion: cell detachment force depends on the loading rate

Since the seminal work of Evans et al. and further developments by other groups in the field of dynamic force spectroscopy, it has been established that when submitted to a pulling force, f, increasing linearly in time, a single adhesive bond breaks at a force level that is stochastic due to thermal fluctuations and that the most probable value of this rupture force, f^*, depends on the loading rate, i.e., the rate of force increase, df/dt (25–29). All of the aspiration experiments described above were performed with an aspirating pressure

that increased linearly with time, so that the aspirating force applied to the cell at any time is given by

$$F = S_{\text{pipette}}\Delta P(t) = S_{\text{pipette}}r_{\text{P}}t, \qquad (4)$$

where $\Delta P(t)$ is the applied aspiration pressure at time t, and the aspiration rate, $r_{\text{P}} = (d/dt)\Delta P(t)$, is the constant rate at which the aspiration pressure increases. As shown in Fig. 6 A, we find that the critical stress, σ^*, increases with the aspiration rate, r_{P}.

Making a number of simplifying assumptions, we can write a minimal model to rationalize our experimental results for the detachment dynamics, notably the relationship between detachment force, aspiration rate, and adhesive areas. As mentioned above, we first assume that the adhesive bonds share the aspiration force in parallel. Taking a mean-field approach, we assume that each adhesive bond breaks at the same rupture force, f^*, and approximate the number of bonds, N, as $N = S_{\text{adh}}/s_0$, where s_0 is the average area occupied by a single biological adhesive bond. Assuming a negligible transmission of the aspiration force
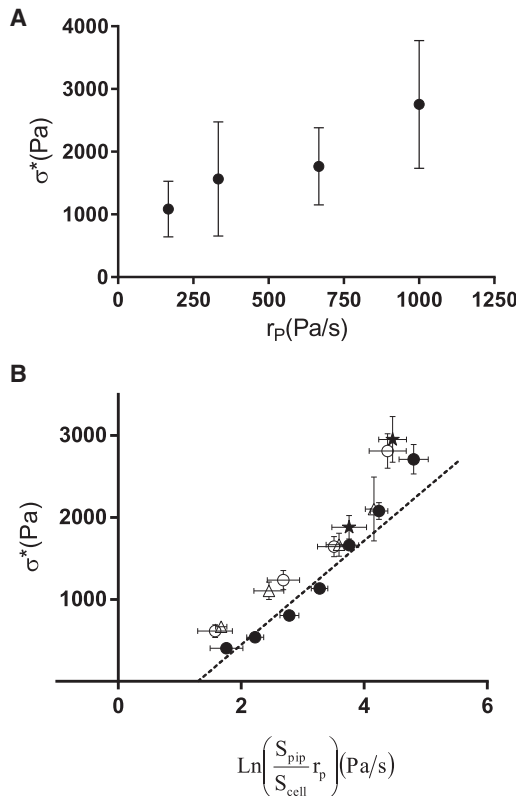

A

B

FIGURE 6 Critical stress as a function of the aspiration rate and master equation. (A) Critical stress versus aspiration rate, from experimental data. Error bars represent the standard deviation. (B) Critical stress versus $\ln((S_{\text{pipette}}/S_{\text{cell}})r_{\text{P}})$ for the conditions where r_{P} varies while $S_{\text{pipette}}/S_{\text{cell}}$ is constant (data points shown in A) (○), $S_{\text{pipette}}/S_{\text{cell}}$ varies while r_{P} is constant (●), cells are exposed to cytochalasin D (△), and cells are treated with nocodazole (★). All data collapse along the master curve, in good agreement with the prediction from Eq. 8. Error bars represent the standard error.

through the membrane, a quasistatic equilibrium between the aspiration force, $F = S_{\text{pipette}}\Delta P(t)$, and the force shared by the N bonds adhering to the substrate is expressed as $F^* = Nf^* = (S_{\text{adh}}/s_0)f^*$. Note that most experiments were conducted under brightfield illumination so that we measured S_{cell} instead of S_{adh}, but as the ratio $S_{\text{adh}}/S_{\text{cell}}$ is well-defined, we can write

$$\sigma^* = \frac{F^*}{S_{\text{cell}}} = \frac{S_{\text{adh}}}{S_{\text{cell}}}\frac{f^*}{s_0}, \qquad (5)$$

which relates cell-scale stresses to molecular-scale forces. We further assume that for a single adhesive bond, the rupture force, f^*, follows a slip-bond behavior with a single energy barrier dominating its dynamics (28,30,31),

$$f^* = \frac{k_{\text{B}}T}{x_{\text{b}}}\ln\left(\frac{\frac{df}{dt}}{\frac{k_{\text{B}}T}{x_{\text{b}}}k_0}\right), \qquad (6)$$

where k_{B} is the Boltzmann constant, x_{b} the microscopic distance characteristic of the bond, and k_0 an off-rate (equivalently the inverse of a lifetime) under no applied force.

The loading rate of the bond is then

$$\frac{df}{dt} = \frac{d}{dt}\left(\frac{F}{N}\right) = \frac{d}{dt}\frac{(r_{\text{P}}t)S_{\text{pipette}}}{\frac{S_{\text{adh}}}{s_0}} = r_{\text{P}}s_0\frac{S_{\text{pipette}}}{S_{\text{adh}}},$$

and hence,

$$\frac{df}{dt} = r_{\text{P}}s_0\frac{S_{\text{pipette}}}{S_{\text{cell}}}\frac{1}{\frac{S_{\text{adh}}}{S_{\text{cell}}}}. \qquad (7)$$

Combining Eqs. 5–7, we obtain the master equation,

$$\sigma^* = \frac{F^*}{S_{\text{cell}}} = \frac{S_{\text{adh}}}{S_{\text{cell}}}\frac{1}{s_0}\frac{k_{\text{B}}T}{x_{\text{b}}}\ln\left(\frac{r_{\text{P}}\frac{S_{\text{pipette}}}{S_{\text{cell}}}}{\frac{S_{\text{adh}}}{S_{\text{cell}}}\frac{1}{s_0}\frac{k_{\text{B}}T}{x_{\text{b}}}k_0}\right), \qquad (8)$$

which predicts the dependence of σ^* on S_{cell}, S_{adh}, S_{pipette}, r_{P}, and the microscopic parameters x_{b}, s_0, and k_0. This new expression for σ^* now captures the dependence on both the geometric parameters and r_{P}.

Since r_{P} and the pipette radius were our control parameters, we readily tested the dependency of σ^* on $S_{\text{pipette}}/S_{\text{cell}}$ and r_{P} independently. We maintained $r_{\text{P}} = 667$ Pa/s constant while varying the pipette surface area from 16.6 μm^2 to 707 μm^2 (over $N = 335$ cells), or kept $S_{\text{pipette}} = 165\ \mu m^2$ constant while varying r_{P} over four different trials, $r_{\text{P}} = 167$ Pa/s, 333 Pa/s, 667 Pa/s, and 1000 Pa/s ($N = 55$ cells). Based on data from the literature (32,33), we took $s_0 = (80\ \text{nm})^2 = 6.4 \times 10^3\ nm^2$ for the average surface area per adhesive bond, and verified that the data satisfactorily collapsed on the master curve given by Eq. 8 (Fig. 6). A linear fit of the data in Fig. 6 B yields a slope of 634 Pa and an intercept with the y axis, at $\ln(r_{\text{P}}(S_{\text{pipette}}/S_{\text{cell}})) = 0$, of -821 Pa. Taking the average value $S_{\text{adh}}/S_{\text{cell}} = 0.09$ and $k_{\text{B}}T = 4.10^{-21}$ at

room temperature, we obtain a characteristic lengthscale of $x_b = 0.11$ nm and an off rate of $k_0 = 0.006$ s^{-1}.

This off rate is consistent with values obtained in single-molecule force spectroscopy; for instance, Evans et al. measured zero-force off rates ranging from 10^{-4} to 10^{-1} s^{-1}. However, although a microscopic distance, x_b, of only 0.1 nm is not unrealistic compared to distances measured in other bonds (34), larger microscopic distances x_b ranging from 0.4 to 0.6 nm were measured for ICAM1-β_2 integrin bonds (35,36).

Note that to keep our model simple, we neglected the stochastic nature of the bond rupture by assuming that all bonds would all break at the most probable rupture force, f^*, and implicitly considered that once broken, adhesive bonds could not reform. However, we are applying very weak loading rates on each bond ($\sim 4.10^{-2} - 4$ pN/s), and rebinding might play a role. Litvinov et al. (32) found much larger off rates for fibrinogen-integrin interactions than we did in this study, on the order of $k_0 \sim 1$ s^{-1}. This large discrepancy with our value for k_0 could be a signature of the rebinding that we neglected in our model. Another major assumption we made was to consider slip bonds, although evidence of both slip and catch bonds was reported for integrins (see Zhang et al. (37,38), Li et al. (39), and Jiang et al. (40) for slip bonds and a more recent work by Jiang et al. (41) for catch bonds). One technical advantage of considering slip bonds is that the relationship between the rupture force, f^*, of a single slip bond submitted to a loading rate is simple and involves only two microscopic parameters. On the other hand, there are multiple catch-bond models, and they involve more parameters (30,42).

The master equation (Eq. 8) underestimates the detachment stress, σ^*, at large values of $(S_{pipette}/S_{cell})r_P$, i.e., at large loading rates, according to Eq. 7: this might be due to our model for the dynamics of adhesive bonds, as it has been shown that the dynamics of some adhesive bonds are dictated by more than one major energy barrier (26,28). To take this into account, Eq. 6 could be refined to include multiple linear terms in the relationship between f^* and $\ln(df/dt)$, with coefficients depending on the energy barrier governing the dissociation dynamics (26,28,29,43). Nevertheless, our simple model captures well the dependence on geometric parameters (S_{cell}, $S_{pipette}$, S_{adh}/S_{cell}), on intrinsic characteristics of adhesive bonds (x_b and k_0), and notably on the rate r_P at which the aspiration pressure is applied. This validates our global picture of detachment dynamics relying on adhesive areas sharing the aspiration stress in parallel.

Effect of cytoskeletal inhibitors

To assess the role of the cytoskeleton, we also investigated the effect of actin depolymerization. To do so, we incubated cells for 30 min with a high concentration of cytochalasin D (1 μg/mL). The cells affected by the drug displayed a star-like form instead of the convex polygonal shapes of untreated cells (see Fig. S9), but showed only a slight decrease in their S_{adh}/S_{cell} ratio in the remaining cell area ($S_{adh}/S_{cell} = 0.07 \pm 0.02$, $N = 30$, in the presence of cytochalasin D, versus 0.09 ± 0.04, $N = 16$, without cytochalasin D). We could roughly estimate S_{cell} before treatment by measuring the surface of the convex polygon in which the cell surface is inscribed. Such a rough estimate made on five cells led us to assume that S_{cell} was reduced by a factor of 0.7 ± 0.1 by cytochalasin D. The fact that S_{adh}/S_{cell} is comparable to untreated cells means that both S_{adh} and S_{cell} were affected by cytochalasin D. Interestingly, even though we again varied the aspiration rate, the measured σ^* collapsed on the same master curve defined by Eq. 8 (see Fig. S10). We further investigated the potential role of cytoskeleton by destabilizing the microtubule network with nocodazole. We obtained an adhesion area, S_{adh}, that was larger relative to S_{cell} than in the control case without nocodazole ($S_{adh}/S_{cell} = 0.17 \pm 0.09\%$, $N = 15$), consistent with the slightly higher values for σ^* than would have been predicted by the master curve for a comparable aspiration rate and pipette diameter (Fig. 6 B). These experiments with cytoskeletal inhibitors show that to be transmitted to the substrate, the aspiration force does not require an intact actin or microtubule cytoskeleton. Although this might be surprising, it is worth noting that force (stress) can be transmitted through a pure liquid, in the same way that aspiration force (pressure) is transmitted through the water inside the body of the micropipette. Hence, one can envision that a solid-like cytoskeletal meshwork is not necessarily required to transmit the forces that detach the cell.

Comparison with other studies

Although direct comparison is not straightforward, we can place our results in the context of the literature on cell detachment. Experiments on biotinylated red blood cells adhering to streptavidin-coated surfaces (4) also led to a detachment process in two phases. During the first phase, the contact radius slowly decreased, and then the adhesive contact underwent a "catastrophe-like" diminution until separation (4). We do not see a slow reduction of adhesion area in our experiments, but the catastrophic diminution of adhesive area found in that study quite matches what we observe. Those authors also introduced a critical force, $f_c = \pi R_c W$, over which the adhesive contact cannot withstand a force in static conditions, where W is the adhesion energy per unit area and R_c the contact radius. From their critical force, we can estimate a critical stress, σ^*, as $f_c/\pi R_c^2 = W/R_c \sim 10^{-4}$J·m^{-2}/$10^{-6}$m ~ 100 Pa. Shortly before the Pierrat et al. study, Prechtel et al. (44) ran similar experiments but with vesicles decorated with lipopeptides and adhering to endothelial cells via integrins. Those authors also observed very rapid detachment of the vesicle (within ~40 ms). They performed experiments at larger

loading rates than ours, but extrapolating their rupture force versus loading-rate relationship leads to detachment forces as low as 100 pN for adhesive patches of typical radius 1 μm, hence a critical stress of only tens of pascals. Francis et al. (9) used flexible aspirating micropipettes as cantilevers to detach red blood cells and *Dictyostelium discoideum* amoeba from hydrophobic or hydrophilic substrates while monitoring the adhesion areas with IRM. For red blood cells, they obtained detachment forces of ~1 nN for an adhesion area of S_{adh} ~ 0.3 μm^2, based on which we can estimate a critical stress of σ^{**} ~ 3000 Pa. For the detachment of *Dictyostelium discoideum*, the authors found a detachment force of ~10 nN for an adhesion area of ~2 μm^2, leading to a critical-stress value of σ^{**} ~ 5000 Pa, close to the one they obtained with red blood cells. We have information neither on the loading rate used nor on the ratio S_{adh}/S_{cell} in that case, but based on the figures by Francis et al. (9), we can roughly estimate that S_{adh}/S_{cell} ~ 0.3, so that σ^* ~ 0.3 σ^{**} ~ 1000 Pa. These values are larger than the one estimated above based on studies by Pierrat et al. and Prechtel et al., but one major reason might be that the latter groups used specific bonds, whereas Francis et al. used a hydrophilic or hydrophobic glass substrate with no serum in the medium (hence no extracellular matrix molecules). We stress that a difference in the red blood cell detachment in the above-mentioned studies is that the force was expected by the authors to act only at the perimeter of the adhesion zone, in a ring of finite width (44), as opposed to our case, where force is shared by adhesive bonds in parallel. Coman used microneedles to pull on cell-cell adhesions (6,7) and obtained typical rupture forces of 10^4 nN (1 mg) for nontumoral cells, but without including information about the cell-cell adhesion area. By (roughly) estimating lateral contact between the epithelial cells used by Coman as ~100 μm^2, we obtain a critical-stress estimate of 10^5 Pa, much larger than our measurements, although with no information about the applied loading rate and allowing for the fact that we may be overestimating the area and cell-cell contacts. Ryu et al. (8) pulled with a nanoneedle on cells covered with a proteic scaffold and obtained detachment forces of ~500 nN. By estimating the area of the cell-substrate interface as ~100 μm^2, we can estimate a critical stress of ~5 kPa. The authors used nanoneedles of spring constant ~1 N/m and a retraction velocity of 5 μm/s, leading to a force increasing at a rate of 5000 nN/s, which, because it is larger than the maximal loading rate (on the whole cell) of $dF/dt = S_{pipette} r_P$ ~ 700 nN/s used in this study, makes their larger critical-stress value consistent with our measurements. In a different experimental configuration, Salánki et al. performed detachment assays using an aspirating micropipette held near (but not in contact with) an adhering cell, with a stepwise increment in aspiration pressure (16). The detachment force was not measured but rather was computed using fluid-flow simulations, and they obtained typical values of 2000 nN. Their cells had a projected area of 500–700 μm^2, yielding a typical stress of 3000–4000 Pa that might be consistent with our measured critical stress σ^*. However, Salánki et al. did not correlate the detachment force with the cell projected area for a given cell type, so it is difficult to speculate whether their data regarding applied loading rate are directly comparable to ours.

Regarding techniques that employ cell detachment by shear, a study using a cantilever of known stiffness to measure the force necessary to scrape cells from their substrate found a constant critical shear stress of 530–750 Pa (11,12). Since their cantilever stiffness was 3.12 N·m^{-1} and its approach velocity 20 μm·s^{-1}, the loading rate applied to the cell was ~ 6×10^4 nN·s^{-1}. At our highest aspiration rate, r_P, and our largest $S_{pipette}$, we impose a loading rate on the whole cell of dF/dt ~ 700 nN·s^{-1}, which corresponds to a critical stress of σ^* ~ 3000 Pa (Fig. 6 *B*), which is larger compared to their critical shear stress (11,12) and would be even larger if we were to interpolate to a higher loading rate. Studies performed in microfluidic channels apply a fluid shear stress to a cell population. Klein et al. (18) increased the shear stress in a stepwise manner and measured a critical fluid shear stress of 3–4 Pa over which 50% of adhered cells would detach. Assuming that this value is representative of a critical stress obtained by shearing cells, this is very small compared to all of the previously mentioned studies, including ours. However, the authors used a model to deduce adhesion forces of ~200 nN and ~300 nN per cell for two different cell types. By dividing by the measured projected cell areas of ~300 and ~80 μm^2, respectively, we can estimate critical stress values of 700 and 400 Pa, respectively, for the two cell types used in that study. It is not clear whether this critical stress is not strongly model-dependent, or whether it can be considered as a critical stress that would be obtained by pulling cells, whereas the critical fluid shear stress of only a few pascals is representative of the critical stress obtained by a dezipping of adhesion molecules. Finally, Christ et al. (17) measured a critical fluid shear stress of 50–100 Pa.

Overall, our measurements are in better agreement with pulling techniques (6–9) than with shearing techniques (11,12,17,18). Scrapping might be more similar to an unzipping of the adhesion molecules on a frontline, as opposed to our situation, where adhesive bonds seem to withstand the pulling force in parallel.

CONCLUSIONS

We have developed, to our knowledge, a new micropipette-based technique to quantify adhesion force at the single-cell level. This technique enables us to fully probe the influence of geometric parameters such as projected cell area, adhesion areas, and micropipette size, as well as dynamical parameters such as the loading rate. We found cell detachment to be a well-defined event and established

a clear correlation between the detachment force and the cell adhesion area. Furthermore, we demonstrated that the detachment force is not a constant but depends on how the force is applied to the cell. Finally, a minimalist model helped us rationalize how the critical stress characterizing cell detachment depends on the cell adhesive area, intrinsic parameters of the adhesive bonds, and the rate at which the force is applied to the cell. We showed that we can predict the force necessary to detach a cell from a substrate by acquiring a single IRM picture of that cell and measuring the area covered by the adhesive bonds. In the context of leukocyte-endothelium interactions, it is therefore reasonable to postulate that the formation of microgaps after monocyte adhesion, described by Kataoka et al. (3), should be detectable directly through IRM. We believe this technique should prove useful for studying in detail the changes at the single-endothelial-cell level induced by monocyte adhesion. It would also be interesting to test how the adhesion of other leukocytes affects the mechanical properties of endothelial cells. Kang et al. (45) have shown that neutrophil adhesion induces a local decrease in endothelial stiffness at the adhesion point that lasts <1 min, but also a stiffness increase in adjacent endothelial cells. Forthcoming results on monocyte and lymphocyte adhesion should help elucidate whether those changes are a universal mechanical response of endothelial cells to the adhesion of leukocytes, or whether these cells can adapt their response to the type of leukocyte adhering to them.

SUPPORTING MATERIAL

Supporting Results and Discussion, ten figures, and five movies and are available at http://www.biophysj.org/biophysj/supplemental/S0006-3495(15)00597-4.

AUTHOR CONTRIBUTIONS

J.H. designed the research; B.H., A.B., and Y.H. performed the research; B.H. and J.H. analyzed the data; and B.H., A.I.B., and J.H. wrote the article.

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Supporting Material for *Characterizing Cell Adhesion using Micropipette Aspiration*

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Calibration of the aspiration system

We applied the aspiration pressure ΔP by dilating a volume of air contained inside a syringe and the tubing connected to it "upstream" of the micropipette immerged in the Petri dish as shown in Figure 2A. A calibration was needed to relate the change in volume ΔV inside the syringe to the aspiration pressure ΔP.

Measuring the ΔP – ΔV relationship with hydrostatic pressure

As a first means of calibration, we replaced the liquid-filled tubing and micropipette shown in figure 2A by a water reservoir open to atmospheric pressure P_{atm} (Fig. S1A-B). By considering the air in the syringe as a perfect gas in isothermal conditions, the product of the air volume by its pressure is constant. Before application of aspiration pressure, the air in the syringe and tubing is contained in a dead volume V_0 at atmospheric pressure P_{atm} (Fig. S1A). After pulling the syringe pump, the new volume $V_0 + \Delta V - V_1$, and the air pressure is $P_{atm} - \Delta P$, where $\Delta V > 0$ is the change in volume inside the syringe, $V_1 = S_{tube} (h_1 - h_0)$ is the air volume lost in the tubing due to the raise of the water level from h_0 to h_1, and S_{tube} is the tube section. The conservation of pressure-volume product writes

$$P_{atm}V_0 = (P_{atm} - \Delta P)(V_0 + \Delta V - V_1), \qquad \text{(S1)}$$

which leads to

$$\Delta P = P_{atm} \frac{\Delta V - V_1}{V_0 + \Delta V - V_1} \qquad \text{(S2)}$$

Equation S2 can be used to measure the dead volume V_0 that depends on tube length, section, and dead volumes inside tube connections. To do so, we moved the syringe pump position, we measured the new water level h_1, from which we deduced $V_1 = S_{tube} (h_1 - h_0)$, as well as the hydrostatic pressure ΔP that is independent of tube section and related to the difference in height $h_1 - h_0$ through $\Delta P = \rho g(h_1 - h_0)$, where ρ is the volumic mass of water and g the gravitational acceleration.

This previous calibration technique is limited by the maximum difference in water level $h_1 - h_0$ achievable, typically 50 cm in our case, hence a maximum hydrostatic pressure difference of 5000 Pa. Importantly, although the term $V_1 = S_{tube}(h_1 - h_0)$ is negligible for large ΔV, it cannot be neglected for small ΔV.

Measuring the ΔP – ΔV with a differential pressure sensor

Another method to calibrate the aspiration pressure system is to use a pressure sensor. We used a home-made pressure sensor using a $\pm$ 25 kPa differential gas pressure sensor chip (Robotshop Inc., Gonesse, France) that we connected to an Arduino Uno board (http://www.arduino.cc) in order to convert the output analog signal of the sensor to a 10-bit signal, which was displayed on a LCD screen device (http://www.dfrobot.com) after a temporal average. In order to convert the output signal to Pascals, we connected the sensor to a syringe (Fig. S1C) and varied the syringe volume $V_0 + \Delta V$ while recording the output signal of the sensor. Importantly, in this case there was a negligible length of tubing connecting the syringe to the sensor. This led us to a measurement of the dead volume V_0 in this particular setting with a relative accuracy better that 1%. Knowing the air pressure by considering it as a perfect gas, we could measure the conversion factor from 10-bit output signal of the sensor to Pascals.

Once calibrated, we could connect the sensor to the actual measurement system (Fig. S1D) and explore much larger aspiration pressures than with the previous calibration technique. Moreover, in the configuration of figure S1D, water level will not change due to the pressure change in the gas, due to the incompressibility of water and the negligible amount of water aspirated by the micropipette tip, so that V_1 in Equation S2 is zero and the equation S2 reduces to

$$\Delta P = P_{atm}\frac{\Delta V}{V_0+\Delta V}, \qquad (S3)$$

which can be rewritten as

$$\Delta V = V_0\frac{\Delta P}{P_{atm}-\Delta P}, \qquad (S4)$$

with the convention here that $\Delta P>0$ for an aspirating pressure. According to equation S3, a linear fit of the ΔV *vs* ΔP plot (Fig. S1E) led us to a measurement of the dead volume $V_0 = 14.2$ mL in our system. Although non-linear, Equation S3 leads to an approximately linear ΔP-ΔV relationship of slope $\frac{P_{atm}}{V_0}$ up to $\Delta V \sim 4$mL (Fig. S1F). For larger volume changes, we corrected for non-linearity using equation S3.
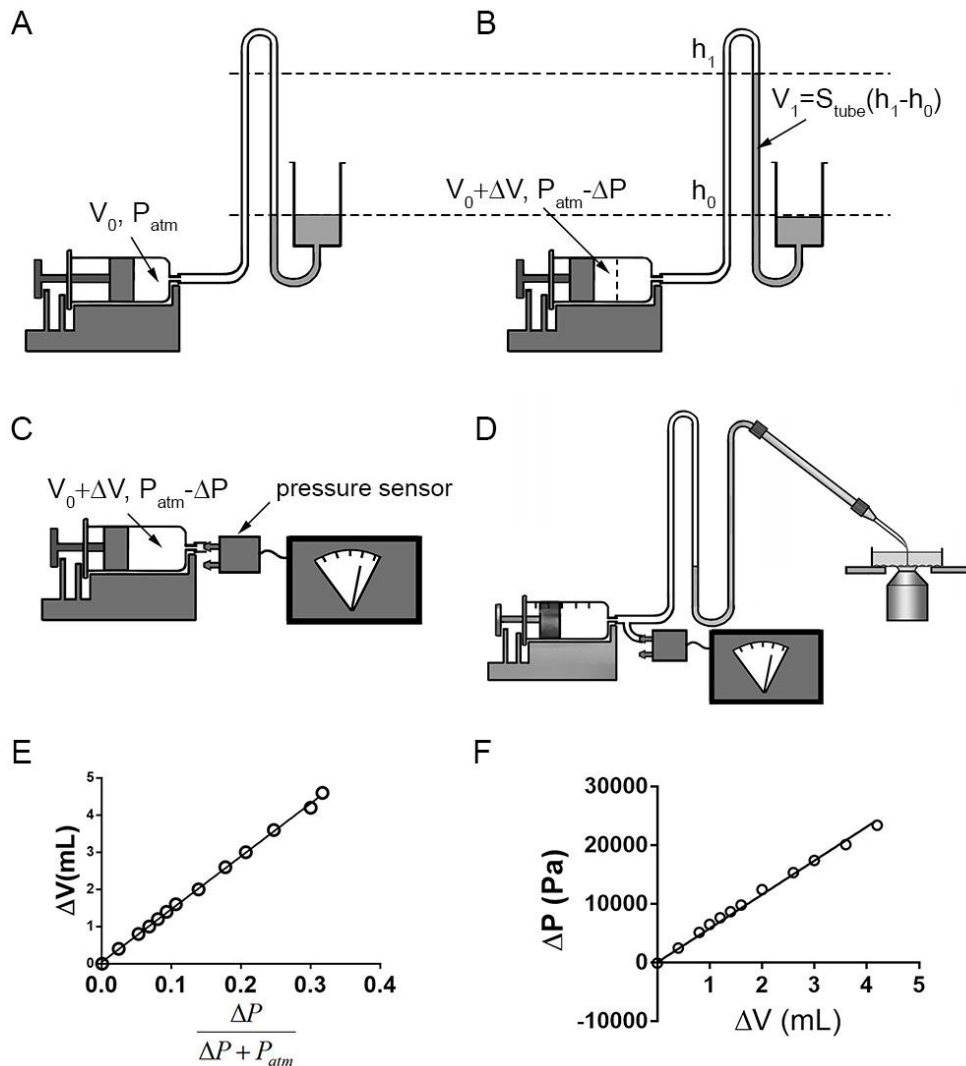
FIGURE S1 Calibration of the aspiration pressure system. (A-B) the micropipette is replaced by a water reservoir open to atmospheric pressure to measure hydrostatic pressure differences by measuring differences in water levels. (C) To calibrate a differential pressure sensor, it was directly connected to a syringe of known inner volume. (D) Once the corresponding value of the output value of the sensor was determined in Pascals, the sensor was connected to the experimental setup. (E) According to Equation S4, the slope of the ΔV *vs* $\frac{\Delta P}{P_{atm}-\Delta P}$ plot is the dead volume of the system V_0. (F) The ΔP-ΔV relationship is nearly linear up to volume changes $\Delta V\sim 4$ mL. Above that value the Equation S4 can be used to correct for non-linearity of the relationship.

Thresholding procedure of IRM images

Images acquired with IRM (Fig. S2A) were processed as follows using the software Image j (ref. [20] in main text). The definition of raw images definition was 16 bits and kept as such until final binarization. An averaging filter was applied ('Filter/mean function', radius 2 pixels), then the background of the image was subtracted in order to correct for inhomogeneous illumination ('Process/Substrack Background' function, with radius 30 pixels, Fig. S2B). Then using an Image j home-made macro, the maximum and minimum gray values of the images where stored in memory. In order to exclude from the process cells neighboring the cell of interest centered in the image, a polygonal shape was defined by the user, and for the sake of helping visual inspection, the outside of the shape was filled with white ('Edit/Clear Outside', Fig. S2C). Then pixels were kept whose value ranges from the minimum up to a value that depended to the height threshold δ determined by Equation 1 in the main text. For $\delta=50$ nm, we obtain $I_{thr} = 0.363*I_m + 0.637*I_M$, where, consistently with Equation 1, I_M and I_m are the maximum and minimum intensity levels, respectively. For $\delta=60$ nm (in the case of the micropipatterns), we obtain $I_{thr} = 0.199*I_m + 0.801*I_M$. The resulting threshold (Fig. 2D) was then applied to obtain a binary image (Fig. 2E) whose area (in black) can be measured in pixel^2 and converted to μm^2 (with our 100X objective 1 pixel represents 0.06 μm).
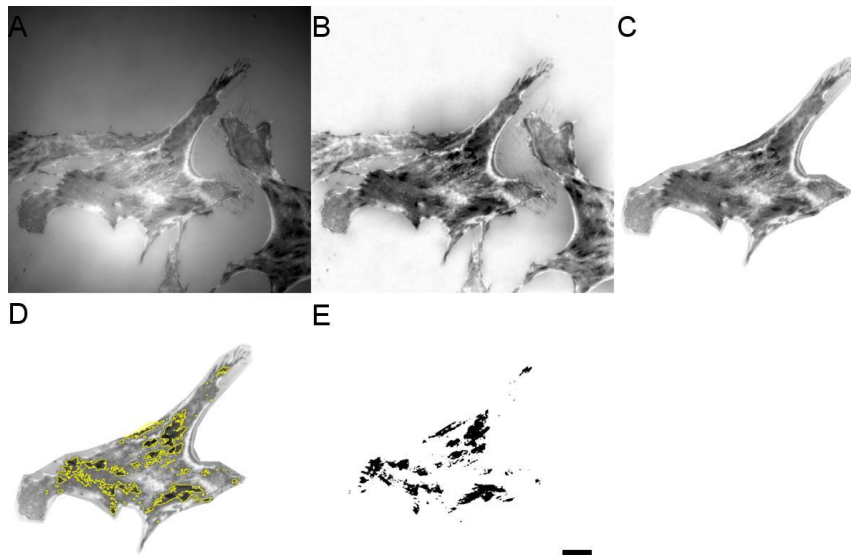

FIGURE S2 Thresholding procedure of IRM images. (A) Raw image. (B) Image after background subtraction . (C) User-defined region to be processed (white outside). (D) Threshold appears in yellow. (E) Binary image obtained after application of the threshold. The bar represents 10 μm.

Adhesion areas are stable after membrane rupture

In order to complement experiments in profile mode to show that breaking cell membrane did not imply cell detachment, we ran a set of complementary experiments in in-plane mode. We applied an aspiration pressure of 5.5 kPa on cells in presence of propidium iodide in the extracellular medium, until the rapid increase in fluorescence showed that the membrane broke. As soon as rupture was detected, we set the aspiration pressure back to zero while leaving the micropipette in place, and we waited up to 300 seconds. During the whole process we imaged the cell-substrate interface with IRM at a rate of one image every 4 seconds. The quantification of these experiments showed that S_{adh} stayed relatively constant even in the presence of a broken membrane (Fig. S3). After typically 200s, the large increase in fluorescence level due to propidium interfered with a proper IRM monitoring in some cells. When the level of propidium fluorescence was very large, we could not quantify S_{adh} properly (over 11 cells, 7 could be analyzed for at least 150 s until propidium fluorescence was too

large) but even in these cases adhesion area did not seem to be strongly impacted. The supplementary movie S5 shows an example of these experiments.
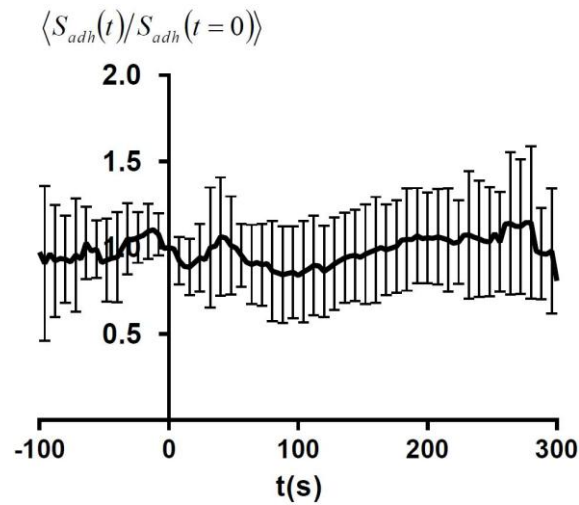

FIGURE S3 Time evolution of the averaged adhesion surface S_{adh} normalized by its value at time t=0 s, which corresponds to the instant at which cell membrane was rupture under a 5.5-kPa aspiration pressure in an in-plane mode experiment, under IRM imaging. As soon as rupture was detected with propidium iodide, the aspiration pressure was set back to zero while leaving the micropipette in place. Average was performed on 7 cells. Error bars represent standard deviation.

Detachment of cells from a low-adhesion surface

As shown before (see Figure 5 in main text), applying an aspiration pressure on cells adhering to glass or Cytodex-3 beads led to a rupture of cell membrane before cell detachment. In order to test the behavior of cells during detachment from a low-adhesion surface, we grew endothelial cells on a surface coated with a low concentration of PLL-g-PEG molecules. We prepared these substrates by incubating plasma-treated μ-Dishes with the PLL-g-PEG solution used to prepare micropatterns but at a 3% concentration (i.e. at 3 μg/mL) in PBS. While PLL assures adhesion to plasma-cleaned glass, the PEG group is anti-adhesive. When left for 2 hours to adhere, cells developed a limited projected area and appeared as more dome-like in brightfield illumination. Interestingly, detachment force in these conditions scaled with S_{cell} (Fig. S4C) with a critical stress $\sigma^* = 1500 \pm 500$ Pa (N=9 cells) very consistent with the one obtained on glass substrates ($\sigma^* = 1300 \pm 50$, N= 335 cells). However, when applying the usual thresholding protocol of IRM images with a 50-nm threshold, we obtained a detachment that still scaled with S_{adh}, but led to a significantly (p<0.0001, unpaired t test) lower critical stress $\sigma^{**} = 4300 \pm 2300$ Pa (N=9 cells) compared to the case without PLL-g-PEG. These values corresponded to a large value of S_{adh}/S_{cell}. One possibility to explain this difference is that applying the same threshold in presence of PEG is not appropriate, since PEG could lead to larger cell-substrate distances. However, setting for instance a threshold of 60 nm instead of 50 nm would select an even larger S_{adh} and go in the other direction, i.e. an even lower value of $\sigma^{**} = \sigma^* / (S_{adh}/S_{cell})$. We ran another set of detachment experiments in presence of propidium iodide in the extracellular medium in order to test whether membrane broke before the forces needed to detach cell in these conditions were reached. Interestingly, for the majority of cells (17 out of 20 cells tested), the cell membrane did not break before detachment. Although the value obtained for σ^{**} is not straightforward to interpret, the consistent value obtained for σ^* suggests that membrane rupture does not influence σ^* for cells grown on more adhesive situations.
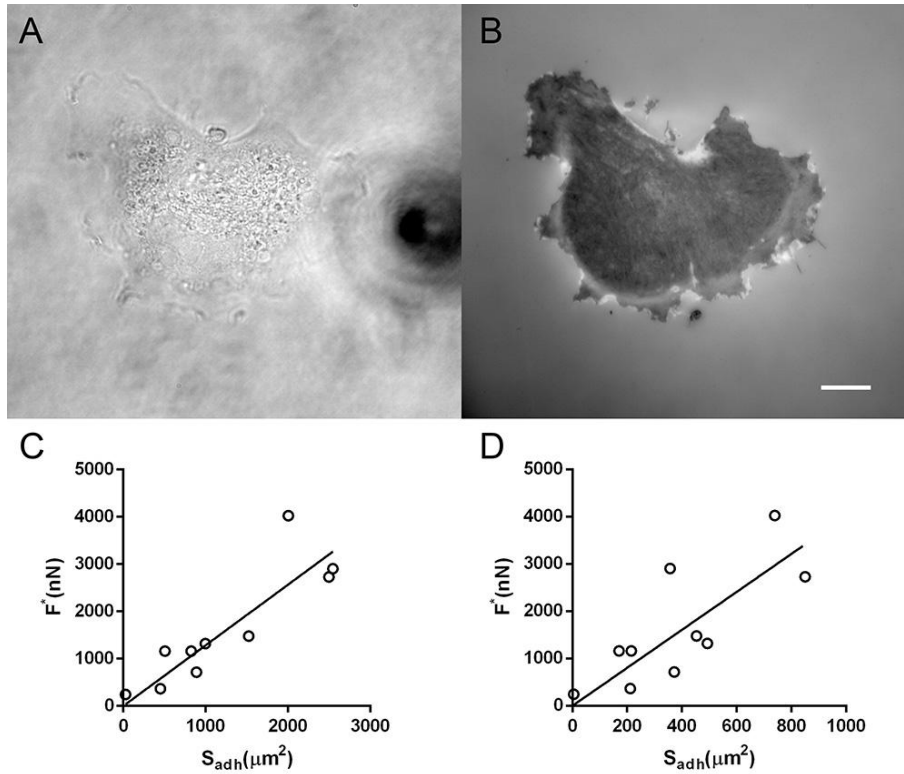
4

FIGURE S4 Brightfield (A) and IRM (B) image of a cell cultured on a glass substrate covered with a low density of PLL-g-PEG. Detachment force F^* vs S_{cell} (C) or S_{adh} (D) for cells cultured in these conditions. The bar in (B) represents 10 μm.

Immunostaining

As a preliminary experiment, we performed immunostaining on cells still attached to their substrate. We stained actin (with Phalloidin Alexa-Fluor-594, Life Technologies), the nucleus (with DAPI, Life Technologies), and most importantly, vinculin (primary antibody: monoclonal Anti-Vinculin antibody produced in mouse, Sigma; secondary antibody: Alexa Fluor 488 AffiniPure Donkey Anti-Mouse IgG (H+L), Jackson). See supplementary figure S5 for an example of stained cell.
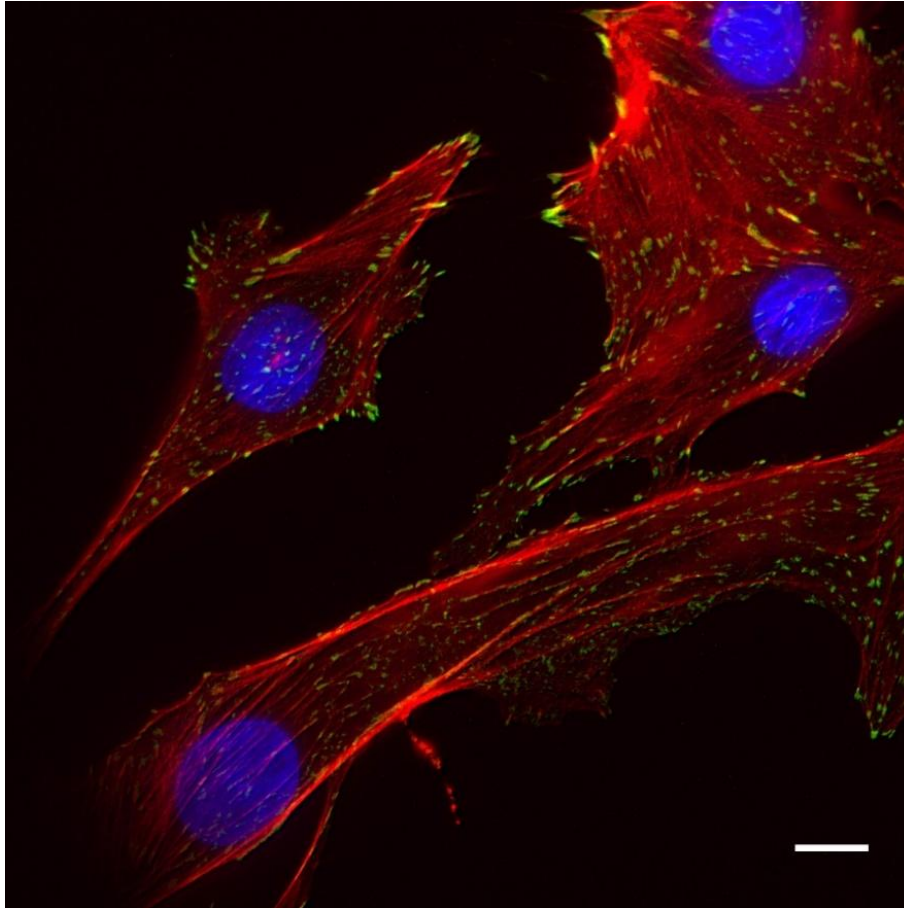
FIGURE S5 Immunostaining. Vinculin staining appears in green, actin staining in red, and nucleus staining in blue. The red channel was processed by applying a "Shadow" filter using Image j software in order to enhance stress fibers. The bar represents 10 μm.

Colocalization of adhesion areas determined with IRM and vinculin

The staining on intact cells described above allowed us to verify that using the 50nm-threshold to define adhesive zones, as used to quantify S_{adh}, we obtained zones that colocalized with vinculin staining (see two examples in figure S6).
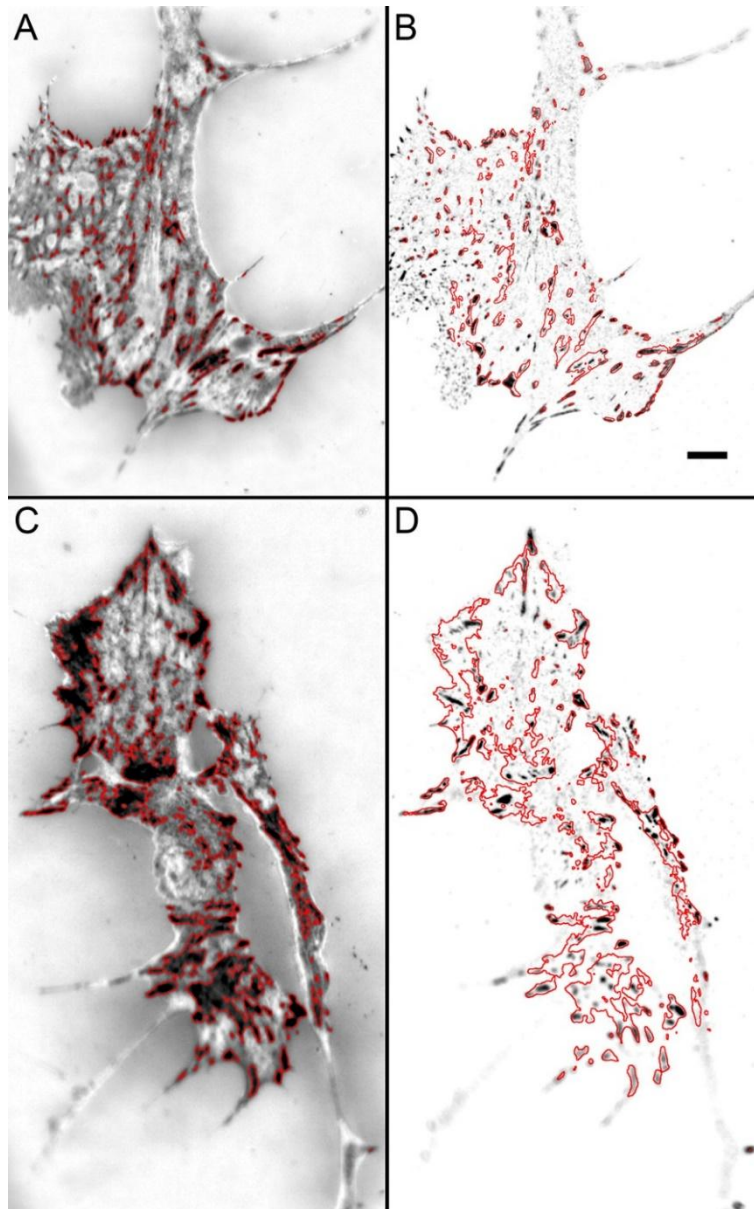
FIGURE S6 Adhesion areas as defined by IRM thresholding match vinculin localization following immunostaining. (A) and (C) Processed IRM images as described previously. The 50-nm threshold is shown in red. (B) and (D) Corresponding image obtained in the GFP channel in epi-fluorescence. The image is inverted, so that vinculin-rich area appear as dark. The threshold obtained from the IRM image is superimposed in red and shows that zones defined as adhesion areas as defined by thresholding IRM images are vinculin-rich.

Presence of traces left after cell detachment

After cell detachment in in-plane mode, the substrate presented some "traces" that were hardly visible in brightfield, but by imaging cells in both brightfield and IRM, we observed that they appeared as "dark traces" in IRM, and that they were systematically present after a cell had been detached (N=9 cells, figure S7).
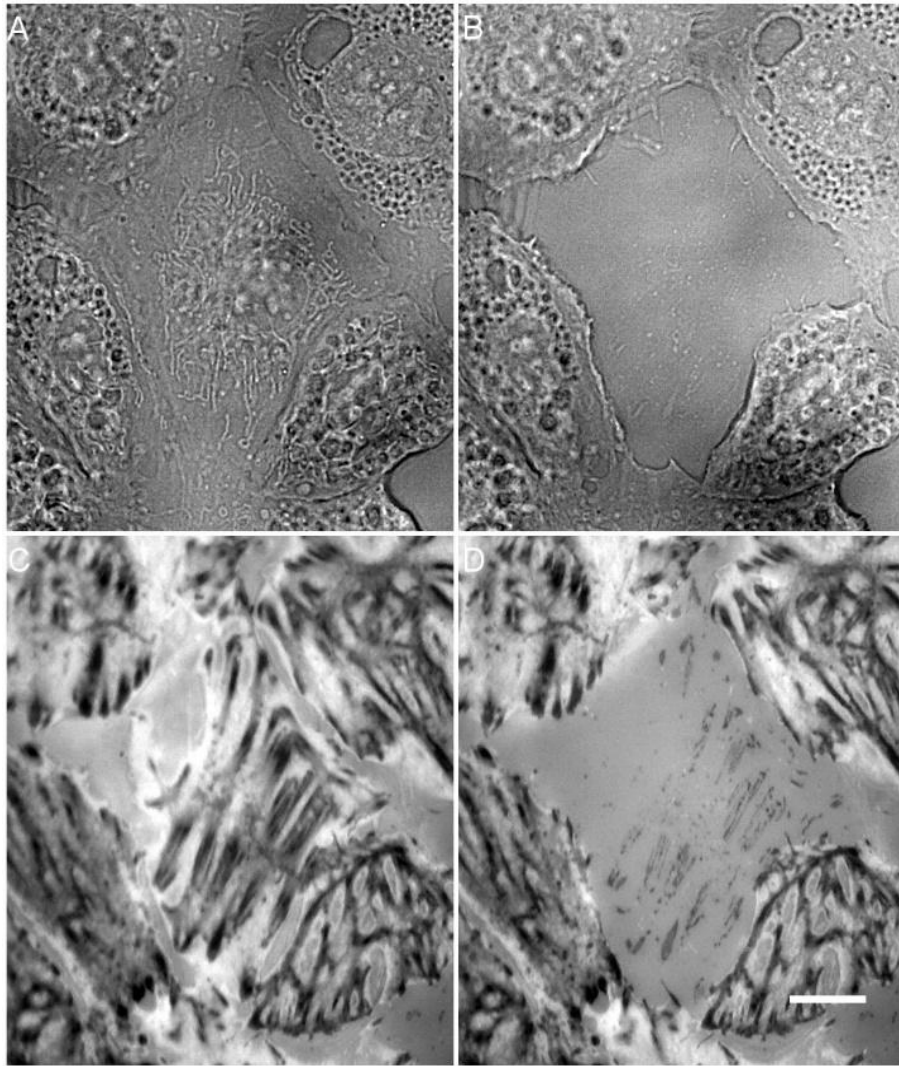
Figure S7 Dark traces are always present after cell detachment. (A) and (B) Images obtained with brightfield illumination before (A) and after (B) cell detachment. (C) IRM image of the same cell as in (A) before detachment. Although hardly detectable in bright field, "dark traces" are clearly distinguishable after the cell has been detached (D). The bar represents 10 μm.

We performed immunostaining in order to investigate whether these dark traces would correspond to adhesion areas. The traces did indeed colocalize with vinculin (Fig. S8), leading us to the conclusion that some adhesion molecules were left on the substrate after cells were detached. The fact that adhesion molecules are left on the substrate contradicts the assumption of our model that all adhesion molecules break during cell detachment. We then sought to quantify this remaining adhesion area S_{adh}^{after} relative to S_{adh} measured before detachment. By using the same thresholding procedure of IRM images before/after detachment on N=13 cells, we obtained $S_{adh}^{after}/S_{adh} = 0.06 \pm 0.07\%$. Thus on average 94% of all adhesive areas break during cell detachment, so that we kept the simplifying assumption of our model that adhesion areas would all detach. However, this raised an interesting question on the process occurring locally to these remaining adhesion areas during cell detachment. One possible scenario is that at these locations, the membrane breaks instead of the adhesive bonds. We tried to test this hypothesis by running detachment experiments in presence of propidium iodide while acquiring timelapses. We could not detect such local membrane rupture and only observe a major rupture close to the micropipette tip (as shown in supplementary movies 4 and 5). These nanoscale membrane ruptures might exist, but heal too quickly, thus avoiding diffusion of propidium inside the cell.
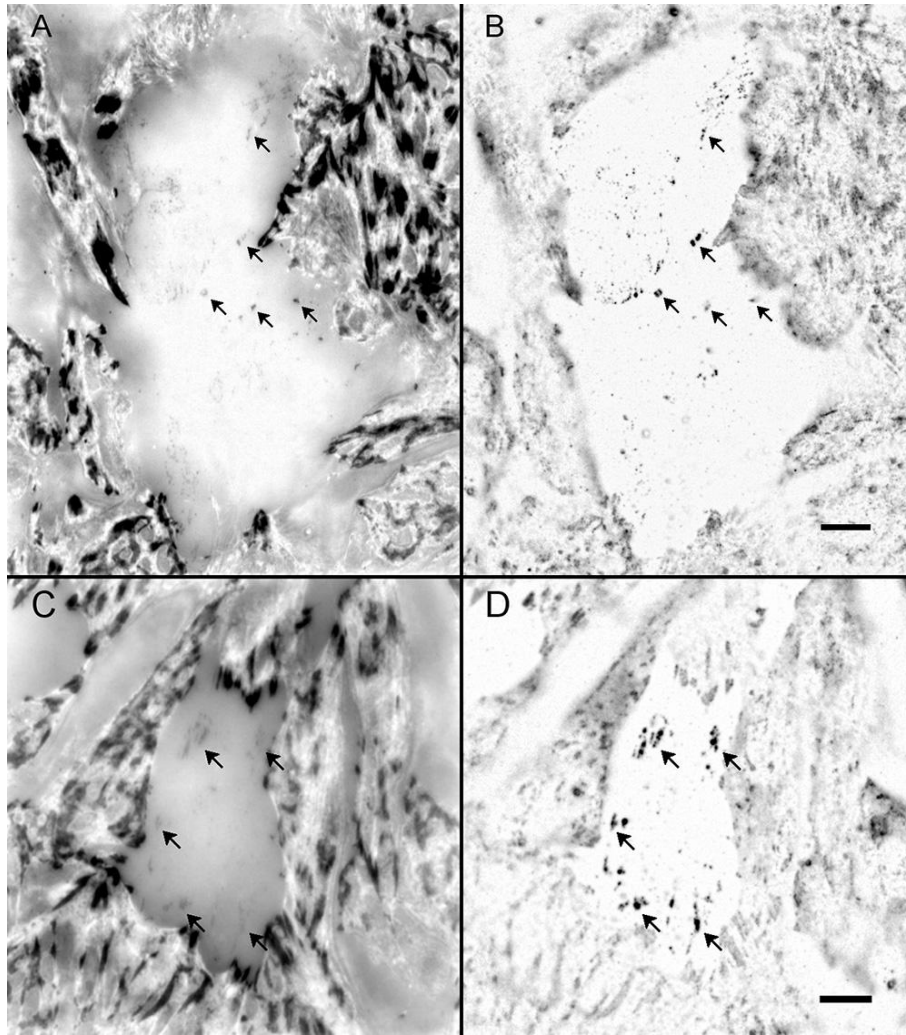
FIGURE S8 "Dark traces" appearing in IRM after cell detachment colocalize with vinculin. (A) and (C) Processed IRM images after two cells were detached. (B) and (D) Corresponding images obtained with epifluorescence in the GFP channel. The images are inverted so that vinculin-rich appear as dark. Arrows show the location of the "dark traces". The bar in (B) and (D) represents 10 μm.

Effect of cytochalasin D

In figure S10A we plotted F* versus Scell for the three different pulling rates r_P that we tested, with 9 or 10 cells for each pulling rate. Although the data are very noisy, individual points collapse well on the same master curve σ^* *vs* $Ln(S_{pip}/S_{cell} \, r_P)$. Furthermore, when the average critical stress σ^* was calculated by averaging F^*/S_{cell}, its value was significantly different (p<0.05, two-tailed t test) between r_P=167 Pa/s and 667 Pa/s, and between 167 Pa/s and 1000 Pa/s (Fig. 10B). Moreover, when compared with the case without cytochalasin D, other parameters being equal, i.e. same pipette diameter and pulling rate, we obtain the same value (p=0.38, Mann Whitney test) for treated and untreated cells (Fig. S10D).
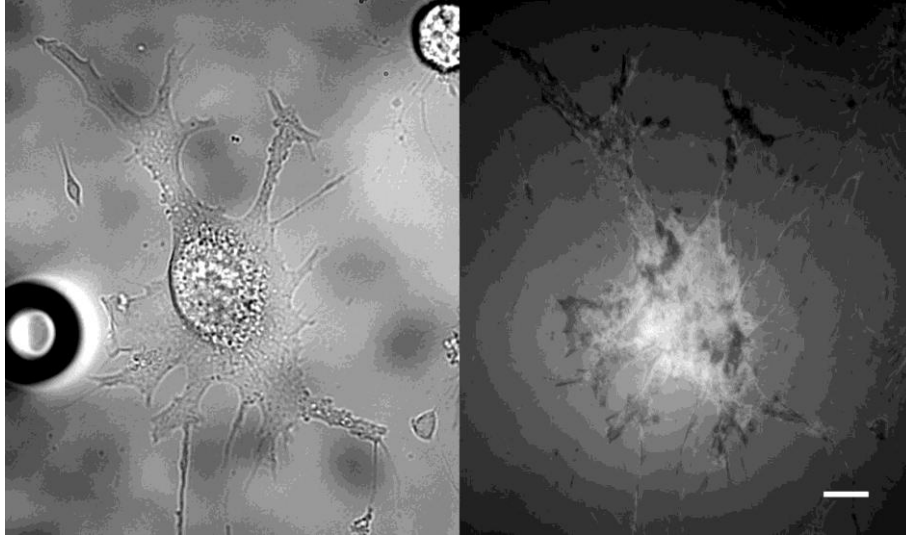
FIGURE S9 Shape of cytochalasin D-treated cells. (A) Image obtained in brightfield illumination. (B) Same cell image in IRM (unprocessed image). The bar represents 10 µm.
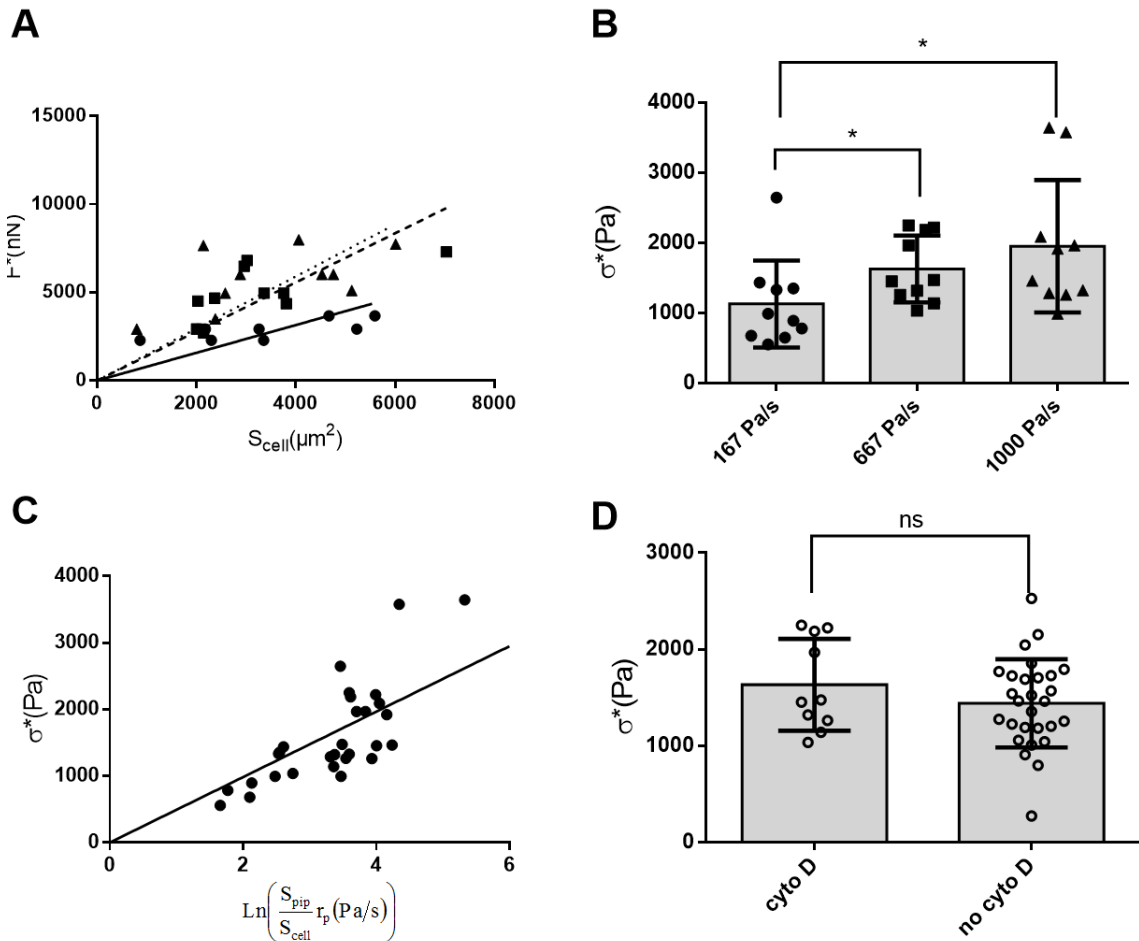

FIGURE S10 (A) Detachment force vs projected cell area for cytochalasin D-treated cells with r_P=167 Pa/s ($\bullet$), r_P=667 Pa/s ($\blacksquare$) and r_P=1000 Pa/s ($\blacktriangle$) (B) Critical stress $\sigma^*=F^*/S_{cell}$ corresponding to (A). (C) Although data are very noisy in (A), they collapse reasonably well on the same master curve as defined in the main text and shown in Figure 6. (D) When compared at same r_P and pipette diameters, critical stress of cytochalasin D-treated and untreated are the same (p=0.38, Mann Whitney test).