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Force response and actin remodeling (agglomeration) in fibroblasts due to lateral indentation

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Abstract

We report the loading and unloading force response of single living adherent fibroblasts due to large lateral indentation obtained by a two-component microelectromechanical systems force sensor. Strong hysteretic force response is observed for all the tested cells. For the loading process, the force response is linear (often with small initial non-linearity) to a deformation scale comparable to the undeformed cell size, followed by plastic yielding. In situ visualization of actin fibers by tagging with green fluorescent protein indicates that during the indentation process, actin network possibly decomposes irreversibly at discrete locations where well-defined circular actin agglomerates appear all over the cell, which explains the irreversibility of the force response. Similar agglomeration is observed when the cell is compressed laterally by a micro plate. The distribution pattern of the agglomerates strongly correlates with the arrangement of the actin fibers of the pre-indented cell. The size of the agglomerates increases with time as t^α , initially with $\alpha = 2-3$ followed by $\alpha = 0.5-1$. The higher growth rate suggests influx of actin into the agglomerates. The slower rate suggests a diffusive spreading, but the diffusion constant is two orders of magnitude lower than that of an actin monomer through the cytoplasm. Actin agglomeration has previously been observed due to biochemical treatment, gamma-radiation, and ischemic injury, and has been identified as a precursor to cell death. We believe this is the first evidence of actin agglomeration due to mechanical indentation/compression. The study demonstrates that living cells may initiate similar functionalities in response to dissimilar mechanical and biochemical stimuli.

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1. Introduction

Study of the mechanical behavior of cells has gained considerable attention during the past 10 years [1–3]. The study is motivated by the observation that a wide range of cell functionalities (e.g., motility, growth, and differentiation) are related to mechanical stimuli. Cells undergo mechanical deformations that originate from various sources such as during systolic and diastolic cycles. Large deformations, on the order of cell size, are also common in soft biological tissues during normal functioning [4–6], injuries [7], and during growth on scaffolds [8,9].

Force response of single living fibroblasts subject to large stretches have been measured by microelectromechanical systems (MEMS) force sensors, and the response was found to be strongly linear, reversible and repeatable [10]. Cell response and cytoskeletal reorganization under indentation and compression yet remain mostly unexplored. Indentation may occur during mechanical injuries. Endothelial cells of blood vessels are subjected to cyclic radial compression which may play a role in the damage of endothelium during atherosclerosis.

For this paper, we used a two-component MEMS force sensor to measure the loading and unloading force response of single living fibroblasts subject to lateral indentation where the maximum cell deformation is not limited by cell thickness or substrate stiffness. In order to explore possible mechanisms of cell force response,

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we employed the green fluorescent protein (GFP) technique to visualize the evolution of the actin network during indentation. We found a direct correlation between the measured force response and the adaptive transformation of actin network. These results shed new light on how cells sense and respond to mechanical stimuli, particularly injuries.

2. Materials and methods

2.1. Force sensors

Fig. 1A shows a two-component force sensor, which consists of a flexible sensor beam, a rigid bar, a sensor

probe attached to the free end of the rigid bar, and a measurement reference bar attached to the sensor base. The probe can be designed according to the desired contact configuration between the probe and cells. Fig. 1C shows the scanning electron microscopy (SEM) micrograph of a two-component force sensor. Here, the lengths of the sensor beam and rigid bar were $L_1 = 1$ mm and $L_2 = 0.429$ mm, respectively. The cross section of the sensor beam was $2.0 \mu\text{m}$ (in plane of paper) $\times 13.1 \mu\text{m}$ (normal to the plane). The rectangular shape and dimensions (about $2 \mu\text{m} \times 5 \mu\text{m}$) of the probe are illustrated in the upper right box of Fig. 1A. The depth of the probe was $13.1 \mu\text{m}$. The sensor was made of single crystal silicon by the process described in Ref. [11]. An $x - y - z$ piezo actuator was

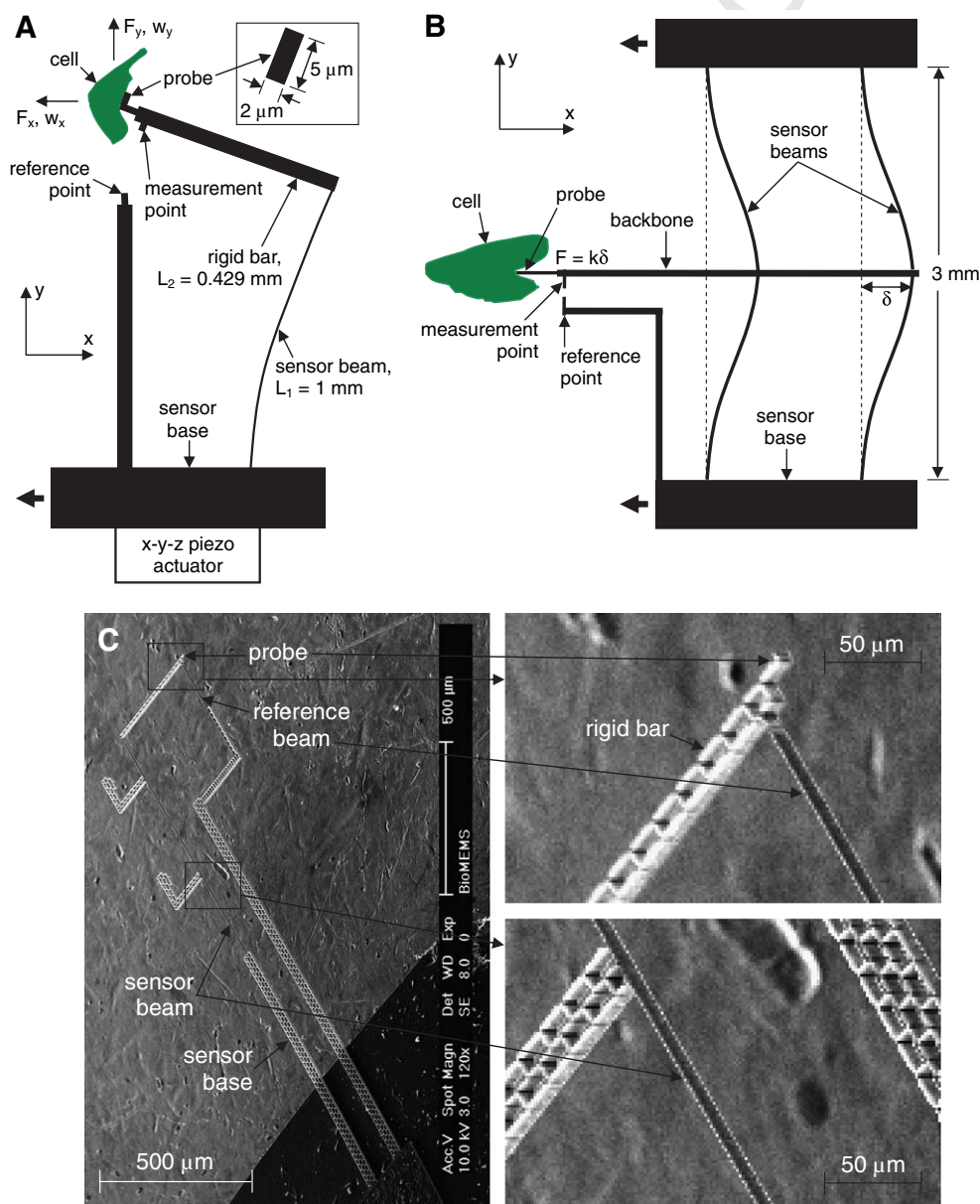


Fig. 1. (A) Schematic drawing of a two-component MEMS force sensor. (B) Schematic drawing of a single-component MEMS force sensor. In (A) and (B) the sensor probe is about 1–2 μm over the surface of the substrate on which the cells are cultured. (C) Scanning electron microscope images of the two-component force sensor and the zoom in views for the probe and sensor beam, respectively.

77 used to move the sensor base along the $-x$ direction to lat-
78 erally indent the cells (Fig. 1A).

79 Two sets of data were recorded during indentation:
80 deformation of the cell and the corresponding force
81 response. Cell deformations in the x and y directions,
82 denoted by D_x and D_y , respectively, at the point of contact
83 between the cell and the probe were measured from the
84 movement of the sensor probe (since the sensor probe
85 was always in contact with the cell). The force exerted by
86 the cell on the probe is estimated from the sensor deflec-
87 tions, w_x and w_y , measured from the relative displacements
88 between the measurement point and the reference point.
89 The two components of the force, F_x and F_y , were obtained
90 from
91

$$93 \begin{bmatrix} F_x \\ F_y \end{bmatrix} = \frac{2EI}{L_1^3} \begin{bmatrix} 6 & 3\frac{L_1}{L_2} \\ 3\frac{L_1}{L_2} & 2\left(\frac{L_1}{L_2}\right)^2 \end{bmatrix} \begin{bmatrix} w_x \\ w_y \end{bmatrix}, \quad (1)$$

94 where $E = 170$ GPa is the Young's modulus and I is the
95 moment of inertia of the sensor beam. If, for example,
96 $w_x = 1$ μm , $w_y = 0.5$ μm , from Eq. (1), we get $F_x =$
97 28.2 nN, $F_y = 36.9$ nN. The magnitudes of the cell
98 deformation and force response are $\sqrt{D_x^2 + D_y^2}$ and
99 $\sqrt{F_x^2 + F_y^2}$, respectively. The cell deformation in the x
100 direction is considered negative when the cell is under
101 indentation. The cell deformation in the y direction is posi-
102 tive when the cell is deformed upward (see for example
103 Fig. 1A). Cell deformation, sensor deflection and force re-
104 sponse are measured with respect to the initial indentation
105 state (after initial observation of the cell for 20 min under a
106 small indentation; see Section 3).

107 A single-component force sensor was used for indenta-
108 tion while visualizing the actin network by tagging with
109 GFP. The schematic of such a sensor is shown in
110 Fig. 1B. It has two parallel beams that measure the force
111 response. The sensor applies indentation and measures cell
112 force response only along the x direction. The sensor
113 beams were 3 mm long with 2.1 $\mu\text{m} \times 7.7$ μm in cross sec-
114 tion, which results in a sensor stiffness of 14.4 nN/ μm .
115 The width and depth of the sensor probe were also
116 2.1 μm and 7.7 μm , respectively.

117 2.2. Cell culture and force response measurement

118 The cells were cultured from a monkey kidney fibroblast
119 (MKF) cell line (CV-1, ATCC, Manassas, VA). They were
120 cultured in a medium with 90% DMEM (ATCC, Manas-
121 sas, VA) and 10% FBS (ATCC, Manassas, VA) in an envi-
122 ronment with 37 $^\circ\text{C}$ temperature and 5% CO_2 . For in situ
123 visualization of actin during force response measurement,
124 the cells were transfected with the pEGFP-Actin vector
125 (BD Biosciences Clontech, Palo Alto, CA) by using
126 CLONfectin (BD Biosciences Clontech, Palo Alto, CA).
127 The force response measurement was conducted in air at
128 room temperature after 60 h of transfection. An inverted

optical microscope (Olympus IX81, Olympus America, 129
Melville, NY) with an objective of $20\times$ was used to image 130
the cell deformations and sensor deflections. A cooled 131
charge-coupled device camera (MagnaFire S99806, Olym- 132
pus America, Melville, NY) and its corresponding data 133
acquisition software were used with the microscope to col- 134
lect the images and measure the data. The entire experi- 135
mental system has been described in detail in Ref. [11]. 136
The force resolution of the measurement system using the 137
two-component force sensor was estimated to be 0.6 nN 138
by multiplying the displacement measurement resolution, 139
 0.14 μm , with the spring constant of the sensor beam in 140
the x direction. Each indentation step was taken within 141
 1 s by moving the sensor base with the piezo actuator. 142
The cell deformation and force response were recorded 143
 15 s after each indentation step. The time delay between 144
two consecutive indentation steps was kept at 50 s unless 145
otherwise stated. In order to contact a cell, the sensor 146
probe was first lowered to gently contact the bottom of 147
the Petri dish on which the cells were cultured. The probe 148
was then lifted up by about 1 – 2 μm , and moved laterally 149
to contact and indent a well-spread and attached cell by 150
a few micrometers. The initial indentation established a 151
stable (non-sliding) contact configuration between the 152
front surface of the probe and the cells. The cell was then 153
observed for the next 20 min for any shape change and 154
movement. The cells that did not undergo any apparent 155
morphological change/migration were considered for force 156
response study. 157

158 3. Results and discussion

159 3.1. Loading and unloading force response

160 Fig. 2 shows a typical force response of an MKF subject 161
to large lateral indentation by the two-component force 162
sensor and three phase contrast images during indentation. 163
Fig. 2A shows the magnitude of the cell force response 164
resultant, $\sqrt{F_x^2 + F_y^2}$, versus the magnitude of the cell 165
deformation resultant, $\sqrt{D_x^2 + D_y^2}$; Fig. 2B shows the cell 166
deformation components, D_x and D_y ; Fig. 2C shows the 167
cell force components, F_x and F_y . The rest of the panels 168
show the phase contrast images at different cell deforma- 169
tion states.

170 Fig. 2A shows that the force response is linear until the 171
cell indentation reaches about 35 μm (a–c), which is compa- 172
rable to the cell size, followed by plastic yielding which is 173
possibly triggered by significant damage of the cell cytoskel- 174
etal structure (from c to d). We will show later that remodel- 175
ing of the actin network occurs during this yielding 176
process, which is an adaptive biological response, different 177
from the mechanical one. Such plastic yielding was not 178
observed under large stretches [10]. At the beginning of the 179
unloading process, the force response dropped sharply (d– 180
e). Two linear responses (e–f and f–g) followed with further 181
unloading. Thus the force response was strongly hysteretic.

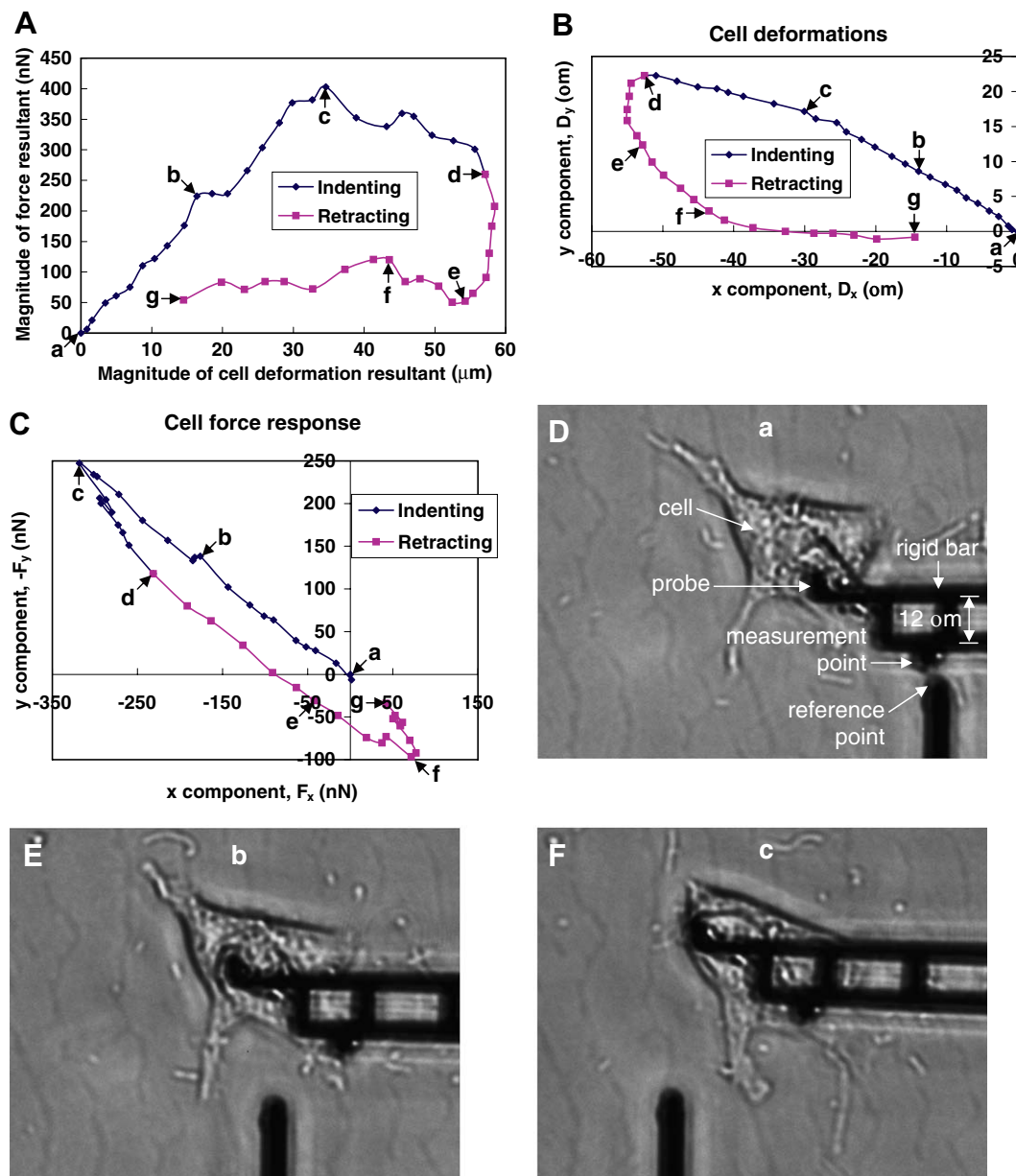


Fig. 2. Results for a widely spread monkey kidney fibroblast obtained by the two-component force sensor. (A) Magnitude of cell force response versus magnitude of cell deformation. (B) Cell deformations in the x and y directions. (C) Cell force response in the x and y directions. (D)–(F) Phase contrast images at the cell deformation states specified in (A)–(C).

182 Fig. 2B and C show that as the cell was deformed
 183 between a and c, the directions of the deformation and
 184 the force response remained the same (straight line between
 185 a and c). But the resistance of the cell against deformation
 186 (i.e., the cell stiffness) in the y direction was higher than
 187 that in the x direction. Hence the angle between the deformation
 188 vector (a–c) and $-x$ axis was 30.8° , whereas the
 189 angle between the force response vector (a–c) and $-x$ axis
 190 was 37.1° . This result shows that the structural anisotropy
 191 revealed by the magnetic twisting cytometry for human airway
 192 smooth muscle cells [12], where the induced surface
 193 deformation of the cells was $\sim 0.2\text{--}0.6\ \mu\text{m}$, may be applica-
 194 ble to a deformation range of about two orders of magni-
 195 tude higher. Fig. 2B also shows that cell deformation is

irreversible, as the loading and unloading paths were
 196 different. 197

The linear response during the first phase of indentation
 198 (Fig. 2A) is surprising, as it differs from some earlier
 199 reports. For example, non-linear (stiffening) cell force
 200 response was observed under indentation by cell poker
 201 technique [13] and AFM [14–16], as well as under compres-
 202 sion by a pair of microplates [17]. Note that, in all these
 203 experiments, cell deformations were orthogonal to the sub-
 204 strate, and were small, limited by cell thickness. The tenseg-
 205 rity cell structural model predicts a cell stiffening force
 206 response [2,18,19]. Non-linear mechanical behavior of bio-
 207 logical soft tissues is also well-established [4,20,21]. We
 208 believe that the linear force response observed in this study
 209

210 under large lateral indentations may arise due to biological
211 adaptation of the cell, i.e., the cell response is both mechan-
212 ical and biological. The observed remodeling of the actin
213 network (see next section) reveals a dramatic biological
214 response under sustained indentation. In order to explore
215 the time dependence of such a biological response, we car-
216 ried out an indentation experiment over a time span of 1.7 s
217 (Fig. 3), in contrast to 37.5 min (Fig. 2). We again found a
218 linear force response, with small initial non-linearity. Thus,
219 if the linear response is due to biological adaptation, it
220 must take place within a time scale much shorter than a
221 second.

222 We also note that, during indentation, although the cells
223 adhere on the substrate, local actin network may be par-
224 tially damaged and some of the cell attachments with the
225 substrate may unbind due to the invasion of the sensor
226 probe. Either of these mechanisms will cause mechanical
227 softening, which, together with expected mechanical stiffen-
228 ing during indentation, may result in an apparent linear
229 response. However, Figs. 2 and 4 suggest that local damage
230 of actin network compared to the overall network and local
231 unbindings compared to the overall span of attachments
232 are small. Hence their influence on the overall force
233 response is expected to be small.

234 3.2. Remodeling of actin network

235 In order to explore the underlying mechanism of the
236 force response, we observed the deformation of the actin
237 filaments in transfected MKFs during indentation force
238 response measurement by the single-component force sen-
239 sor (Fig. 1B). Fig. 4A shows the force response of an
240 MKF. The response shows initial non-linearity, an inter-
241 mediate linear regime, and a long plastic regime.
242 Fig. 4C–H shows the representative fluorescent images at
243 times indicated. Due to photobleaching, the relative bright-
244 ness corresponding to the actin in these images decreased
245 with increasing observation time, but the distribution of

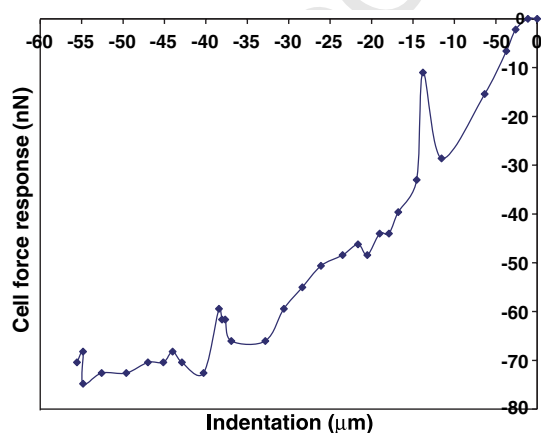


Fig. 3. Fast indentation force response of a monkey kidney fibroblast obtained by the single-component force sensor. The time span for the entire experiment is 1.7 s, compared to 37.5 min in Fig. 2.

246 the actin can still be clearly seen. Panels C and D are in
247 the initial non-linear/linear force response regime where
248 deformation of the actin is highly localized, i.e., only the
249 actins on the way of the probe and its nearby region are
250 deformed, but there is no significant deformation far away
251 from the probe. Panels E–H are in the plastic response
252 regime where part of the actin network far away from
253 the probe lose distinct features and form well-defined cir-
254 cular structures. The higher brightness of these circular struc-
255 tures indicates that these regions are actin agglomerates.
256 With increasing time, the agglomerates grow in size but
257 not in numbers, and a slight relaxation in the force
258 response is observed.

259 Fig. 5 shows agglomeration in three MKFs during and
260 after indentation with a probe and a plate. In Fig. 5A,
261 the agglomerates continued to increase in size even after
262 the probe is withdrawn, and the actin network did not
263 reappear implying that the process of actin remodeling is
264 irreversible. These observations suggest that large sustained
265 indentation or compression induce decomposition of the
266 stress fibers and reorganization of the decomposed stress
267 fibers into well-defined agglomerates in the cell. In
268 Fig. 5B, a plate was used to compress a cell instead of
269 indenting it by a probe. Such compression avoids possible
270 local damage of cell membrane by a probe. The observa-
271 tion of actin agglomeration implies that such remodeling
272 is possibly a general response of a cell under compression
273 or indentation. Fig. 5B also shows local unbinding of the
274 actin network may occur even far away from the region
275 of compression. Fig. 5C shows agglomeration in a cell that
276 is indented and the cell membrane is punctured.

277 3.3. Observations and interpretations

278 3.3.1. Size and distribution pattern of agglomerates

279 Although agglomerates appear all over the cell, they are
280 located at discrete locations only. These agglomerates are
281 mostly circular and hence are likely to occupy spherical
282 or penny shaped domains (constrained by cell membrane,
283 Fig. 5A, middle panel). The fluorescent intensity over most
284 of their size is similar to or higher than the pre-existing
285 actin network (where the agglomerates are formed). This
286 suggests that the agglomerate is formed by actin contrib-
287 uted not only by the actin network it might have replaced
288 but also by other resources as well.

289 3.3.2. Spatial pattern of agglomerates

290 Figs. 4 and 5 show that the spatial distribution of actin
291 agglomerates follow the actin fiber distribution pattern in
292 the pre-indented cell. These agglomerates appear and grow
293 simultaneously. During the initial phase of growth, they
294 appear to be away from the basal cell membrane (on the
295 substrate) and closer to the apical (top) cell membrane,
296 which is noticed by focusing at the plane of the substrate
297 and above. As the agglomerates increase their size, they
298 become bounded by the top and bottom membranes as
299 seen from the phase contrast image (Fig. 5A, middle

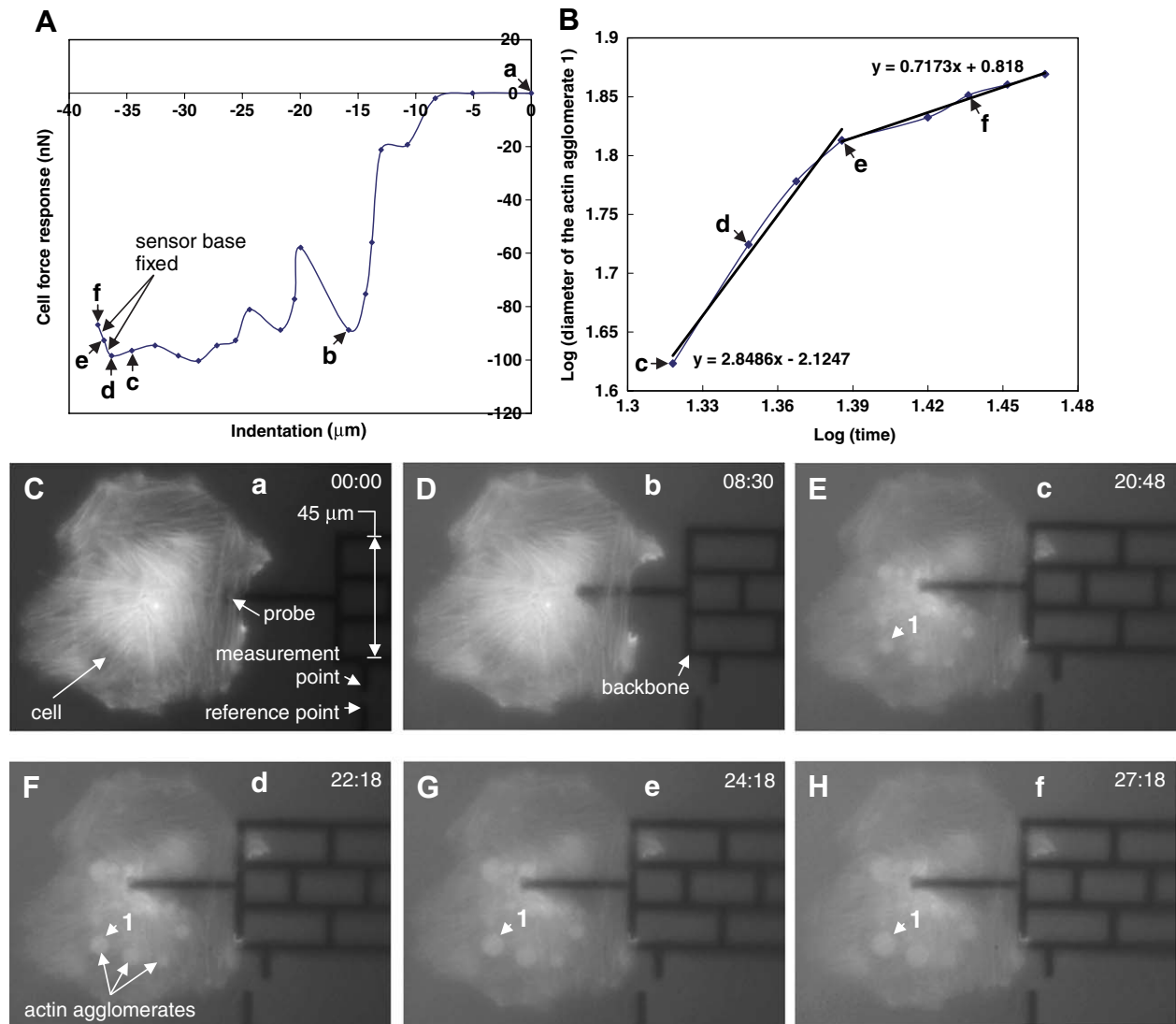


Fig. 4. Results for a GFP-actin transfected monkey kidney fibroblast (MKF). (A) Indentation force response obtained by the single-component force sensor. (B) Logarithm of the diameter of the circular actin agglomerate 1 (formed in the cell and indicated in (E)–(H)) versus logarithm of the time. The diameter of the agglomerate is in the units of image pixel and the time is in units of minutes. (C)–(H) Representative fluorescent images during the measurement at the cell deformation states specified in (A) (min:s). (C) The MKF before the indentation. (G) and (H) were recorded 2 and 5 min after (F), respectively, with the sensor base fixed.

300 panel). In Fig. 5C, where the cell membrane is punctured,
 301 agglomeration is neither simultaneous nor are the agglom-
 302 erates distributed all over the cell. Their distribution is ini-
 303 tially localized near the injured region. With time, they
 304 begin to appear in regions away from the injury. We will
 305 return to this observation to develop a hypothesis on the
 306 agglomeration mechanism.

307 Prior to indentation, the fibers were straight, stable (no
 308 apparent thermal fluctuation) and long, on the order of
 309 30 μm or higher. Since the persistence length of single actin
 310 fibers is about 18 μm [22], the long straight in vivo fibers
 311 suggest that they were under tension and/or each fiber con-
 312 sisted of many single fibers. Absence of fibers along orthog-
 313 onal directions (i.e., fibers along one direction only, Figs. 4
 314 and 5) suggests that most of the regional intracellular
 315 forces were along the direction of the fiber alignment.

The alignment of agglomerates along the direction of fibers 316
 and their simultaneous growth (in cells where cell mem- 317
 brane is not punctured, Figs. 4 and 5A and B) suggest that 318
 they were initiated at points where stress change was high 319
 under indentation. Also, Fig. 5C shows voids and cracks 320
 in the actin network that developed during indentation 321
 between parallel fibers suggesting that forces orthogonal 322
 to the fiber direction develop which separate them from 323
 one another laterally. 324

3.3.3. Cell viability 325

In order to test whether the cells were alive after forma- 326
 tion of agglomerates, we applied 0.4% trypan blue solution 327
 (Sigma–Aldrich, St. Louis, MO) in the cell medium after 328
 the indentation experiment. We found that the cells with 329
 agglomerates became blue indicating that they were dead. 330

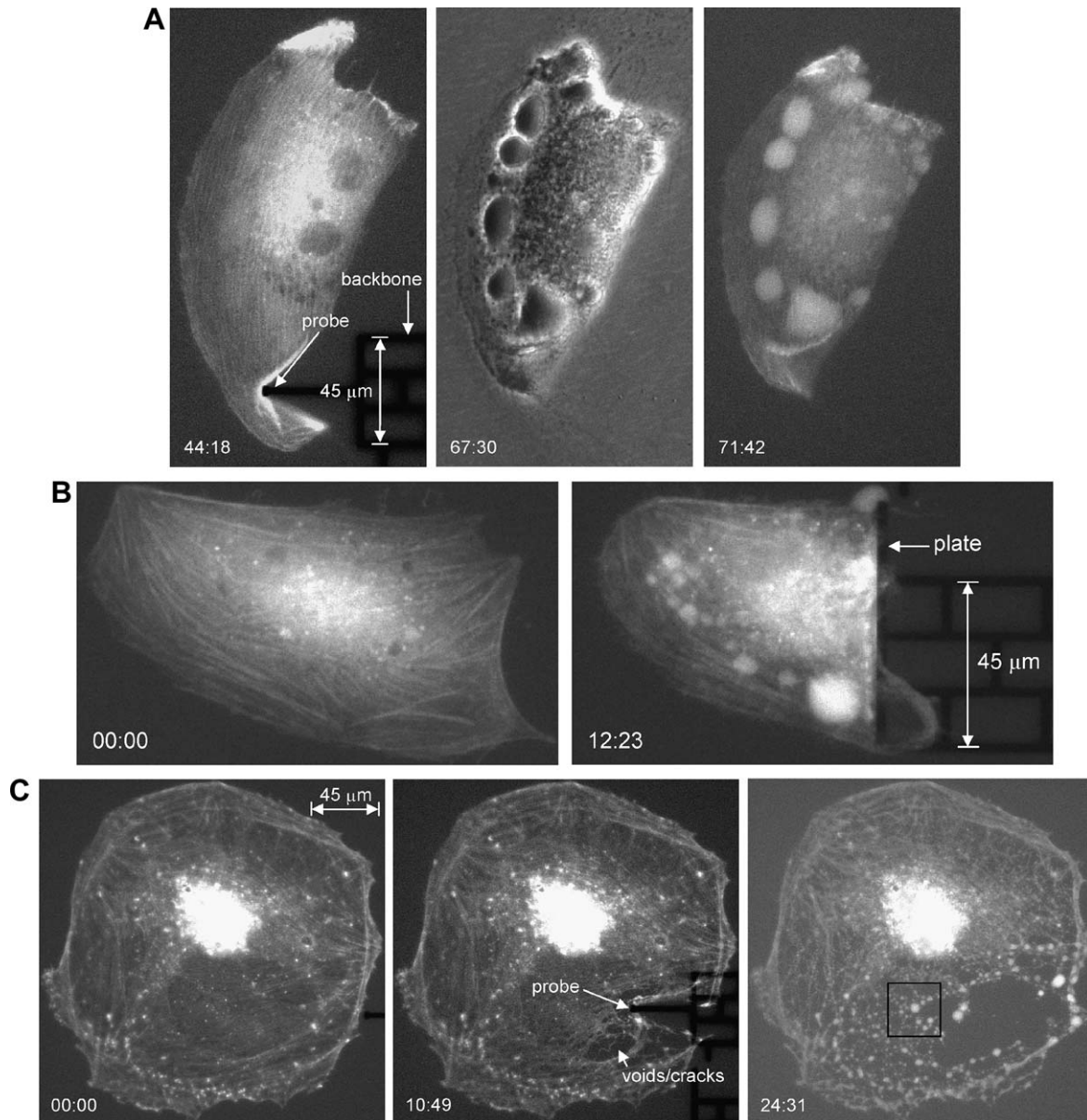


Fig. 5. (A) Response of the actin network in a GFP-actin transfected monkey kidney fibroblast due to the probe indentation and retraction (the middle panel is a phase contrast image). (B) Response of the actin network in a GFP-actin transfected monkey kidney fibroblast due to a plate compression. (C) Response of the actin network in a GFP-actin transfected monkey kidney fibroblast due to the probe indentation and retraction, where the cell membrane is punctured (min:s).

331 Thus, agglomeration might be the pre-apoptotic state of
332 the cell induced by mechanical indentation.

333 3.3.4. Time to agglomeration

334 We indented a cell by about $32\ \mu\text{m}$, and held the inden-
335 tation fixed until agglomeration could be detected. We
336 found that the time to agglomeration was about 1 min. It
337 thus appears that sustained indentation beyond a threshold
338 value initiates agglomeration.

339 3.3.5. Agglomeration in untagged cells

340 Actin agglomerates appeared as blebs in the phase con-
341 trast image of the cell (Fig. 5A, middle panel). In order to

test whether agglomeration dynamics under indentation is
an artifact of GFP tagging of actin, we indented cells with-
out fluorescent tags. Blebs appeared in these cells as well
under indentation, implying that agglomeration is a gener-
al response to mechanical probing unaffected by tagging.

342 3.4. Agglomeration kinetics

343 Fig. 4B shows the time dependence of the diameter of
the actin agglomerate 1, indicated in Fig. 4E-H, plotted
on a log-log scale. Here, the diameter of the agglomerate
is in the units of pixel (of the images) and the time is in
the units of minutes. The dependence shows two distinct

353 regions, i.e., the initial region with a slope of 2.85 and the
 354 remaining region with a slope of 0.72, i.e., initially the
 355 agglomerates grow at a fast rate with time as $t^{2.85}$, followed
 356 by a slow rate of $t^{0.72}$, where t is the time. Note that, due to
 357 photobleaching, the growth can be measured only for a
 358 limited time. Hence, such transition of growth rates may
 359 not be traced unless it occurs before photobleaching.

360 In order to form a statistics of growth rates, we deter-
 361 mined the exponents (α) for 18 such agglomerates from
 362 cells that did not undergo any observable shrinkage or
 363 expansion while the agglomerates were growing. Such size
 364 change would influence the diameter change of the agglom-
 365 erates. Fig. 6A shows the histogram of growth exponents.
 366 Note that a few agglomerates contributed two exponents,
 367 a high and a low one representing two growth rates if they
 368 appeared before photo-bleaching. The statistics show two
 369 peaks in the vicinity of $\alpha = 0.5$ and 2.5.

370 3.4.1. Interpretation of the faster growth rates

371 The higher exponents ($\alpha = 2-3$) suggest a non-diffusive
 372 mechanism. The fast growth cannot be explained by
 373 decomposition of the pre-existing actin network spanned
 374 by the domain of the agglomerate alone. For then the
 375 intensity of the agglomerate would be significantly lower
 376 than the initial actin fibers contributing to the agglomerate.

The actin, initially constrained within the fibers, would
 have to occupy a much bigger volume thus lowering the
 concentration and the corresponding intensity. Experimentally,
 we see the reverse. The increased amount of actin
 within the agglomerate is probably contributed by actin
 monomers available in the cytoplasm generated by uncou-
 pling of sequestering molecules. A possible mechanism of
 agglomerate dynamics is discussed later.

385 3.4.2. Interpretation of the slower growth rates

386 The lower growth rate ($\alpha = 0.5-1$) suggests a diffusive
 387 mechanism, i.e., once the actin fibers decompose at a
 388 region, and if there is no addition of actin, then the accu-
 389 mulated actin diffuses outward. If the diffusing species sat-
 390 isfies the Fick's law of diffusion, then its concentration, C ,
 391 is governed by
 392

$$\frac{\partial C}{\partial t} = D\nabla^2 C, \text{ and } C \text{ evolves with time as } C = \frac{C_0}{(Dt)^{\frac{3}{2}}} e^{-\frac{r^2}{4Dt}} \quad (2) \quad 394$$

where D is the diffusion constant, r is the distance from the
 center of the diffusing agglomerate, and C_0 is a constant. If
 the radius of the agglomerate is characterized by R where
 $C(R) = C^* \ll$ peak concentration at any $t > 0$, then from
 Eq. (2), $R \sim t^{\frac{1}{2}}$, i.e., $\alpha = 0.5$, similar to the exponent ob-
 tained experimentally. Hence the slow growth rate is most
 likely determined by diffusion. We further note that the
 brightness at any point of the two-dimensional (2D) image
 of an agglomerate is contributed by the accumulated flores-
 cence over the depth of the sphere (if the agglomerate is
 spherical). Thus the brightness intensity, $I(x)$, at any point
 x from the center is given by

$$\begin{aligned} I(x) &= 2 \int_0^{\sqrt{R^2-x^2}} C(r) dy \\ &= 2 \int_0^{\cos^{-1} \frac{x}{R}} C\left(\frac{x}{\cos \theta}\right) x(1 + \tan^2 \theta) d\theta \\ &= \frac{2C_0 x}{(Dt)^{\frac{3}{2}}} \int_0^{\cos^{-1} \frac{x}{R}} (1 + \tan^2 \theta) e^{-\frac{x^2}{4Dt \cos^2 \theta}} d\theta, \end{aligned} \quad (3) \quad 409$$

where x is the distance from the center of the agglomerate
 (2D image).

Fig. 6B shows the 2D image (inset) of an agglomerate
 that grows with time as t^α with $\alpha = 0.93$, and its intensity
 profile (the experimental curve in Fig. 6B is obtained by
 the image analysis software Scion Image [23]) in gray scale
 units along the diameter. Also plotted in the figure is the
 predicted intensity profile $I(x)$ along the diameter using
 Eq. (3). Here, C_0 and Dt are chosen such that the predicted
 and experimental peaks match. The close match between
 the experimental and theoretical intensity profiles point
 to a diffusive spreading of the agglomerate.

Note, however, that the diffusion time of the actin
 monomer to travel across the cell (tens of micrometers) is
 on the order of a second [24]. Since the mean square dis-

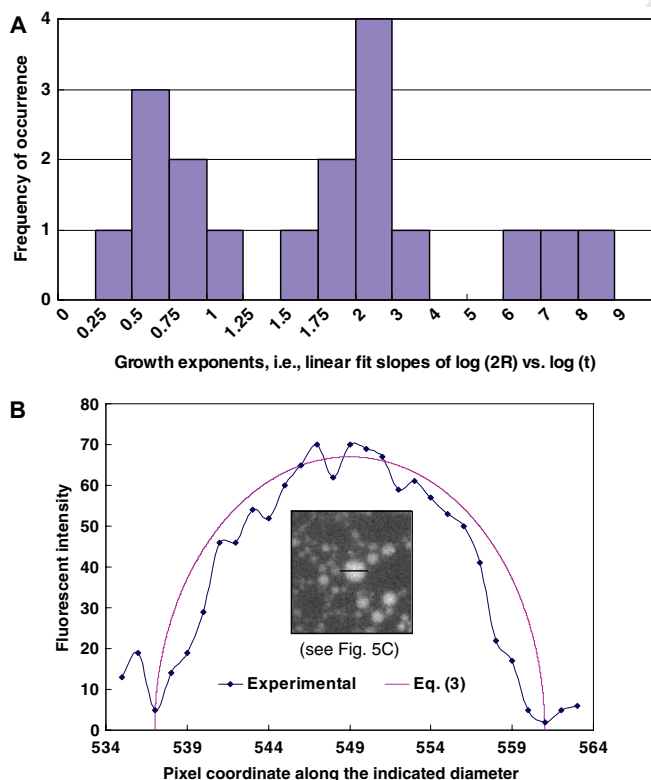


Fig. 6. (A) Histogram of growth exponents of 18 agglomerates formed in the actin network in GFP-actin transfected monkey kidney fibroblasts. (B) Fluorescent intensity profile (the experimental curve) in gray scale units along the diameter of an actin agglomerate shown in Fig. 5C. The predicted intensity profile by Eq. (3) is also shown, where C_0 and Dt are chosen such that the predicted and experimental peaks match.

425 placement, $\langle r^2(t) \rangle$, of a diffusing species is given by,
 426 $\langle r^2(t) \rangle = 2Dt$, we get $D \sim 50 \mu\text{m}^2/\text{s}$ for $\langle r^2 \rangle \sim 100 \mu\text{m}^2$ and
 427 $t \sim 1$ s. If the species within the agglomerate were actin
 428 monomers with $D \sim 50 \mu\text{m}^2/\text{s}$, then they would diffuse
 429 out within 10 s, and the fluorescent intensity would
 430 decrease by two orders of magnitude by then. Since the
 431 agglomerates, some of which appear to be isolated from
 432 the rest of the actin network, grow over a time span of sev-
 433 eral minutes, it implies that the diffusion coefficient (if dif-
 434 fusion is the mechanism of spreading) of actin species
 435 within the agglomerates is a few orders of magnitude less
 436 than that of actin monomer. This suggests that the agglom-
 437 erates consist of species that are much larger than mono-
 438 mers (e.g., segments of actin fibers), or they may have
 439 mutually attractive interaction between them which retards
 440 spreading.

441 3.5. Hypothesis on actin agglomeration

442 3.5.1. Agglomeration is initiated by indentation/compression 443 and triggered by the increase of the intracellular Ca^{2+} 444 content

445 Actin agglomeration observed here is possibly triggered
 446 by the increase of the intracellular Ca^{2+} content. Ca^{2+}
 447 could be released from intracellular storage vesicles, a pro-
 448 cess stimulated by IP₃, and IP₃ could be generated by
 449 mechanically (here it is the mechanical indentation/com-
 450 pression) induced activation of phospholipase C [25]. The
 451 time for intracellular Ca^{2+} to increase may be only 60 s
 452 [26], similar to the time to initiate actin agglomeration
 453 (about 1 min, see above) under sustained indentation.
 454 Another possibility for the increase of the intracellular
 455 Ca^{2+} content is the influx of Ca^{2+} through mechanical
 456 force-induced pores on the cell membrane from extracellu-
 457 lar space.

458 Thus, under sustained indentation/compression, Ca^{2+} is
 459 released from intracellular sources or introduced from
 460 extracellular space through pores at stress high points.
 461 The excessive Ca^{2+} may trigger uncoupling of sequestering
 462 proteins (such as thymosin) from actin monomers in the
 463 cytoplasm [26]. This could generate an abundant supply
 464 of actin monomers ready to be polymerized to form actin
 465 filaments. The excessive Ca^{2+} also activates gelsolin which
 466 severs actin filaments (in the existing actin network and
 467 newly formed actin filaments) into actin fragments and
 468 caps their plus ends [24]. Thus, the combined effect of acti-
 469 vated actin monomers and presence of gelsolin may lead to
 470 a local rapid accumulation of actin fragments, which may
 471 explain the fast growth rate of agglomerates. The process
 472 continues until the local Ca^{2+} concentration decreases to
 473 the normal intracellular level. Further spreading of accu-
 474 mulated actin segments is then achieved by diffusion, with
 475 a diffusion constant much smaller than that of single actin
 476 monomer due to the larger size of actin fragments. Sus-
 477 tained indentation causes sustained release (instead of typi-
 478 cal short temporal bursts) of Ca^{2+} which may lead to
 479 apoptosis [27].

If the cell membrane is fractured (Fig. 5C), then Ca^{2+} 480
 can largely enter the cell from the external medium. The 481
 locally increased Ca^{2+} level near the ruptured region trig- 482
 gers the above process causing actin agglomeration. Such 483
 agglomeration spreads towards the rest of the cell with time 484
 due to the diffusion of extracellular Ca^{2+} towards the inter- 485
 ior of the cell, in contrast to the simultaneous formation of 486
 agglomerates at the stress high points all over the cell. The 487
 overall spatial distribution of the agglomerates due to the 488
 fracture of cell membrane appears random, whereas that 489
 due to intracellular high stress follows the stress fiber 490
 pattern. 491

492 4. Remarks 493

1. Agglomeration of actin is not unusual and has been 494
 observed in mice neurons induced by the treatment of 495
 cytochalasin D [28], in chicken muscle actin induced 496
 by gamma-radiation [29], in the reorganization of depo- 497
 lymerized actin in *Dictyostelium* cells by the treatment of 498
 latrunculin A [30], and in proximal tubule cells by ische- 499
 mic injury [31]. The actin network is found to depoly- 500
 merize and form agglomerates in the late feeding stage 501
 of working larvae, which coincide with the apoptotic 502
 wave [32]. We have also observed actin agglomeration 503
 in MKFs due to the treatment of trypsin-EDTA (images 504
 not shown). To the best of our knowledge, this is the 505
 first evidence of actin agglomeration followed by cell 506
 death due to local mechanical indentation and damage. 507
 It is interesting to note that cells may signal similar pro- 508
 cesses due to such diverse stimuli as mechanical and bio- 509
 chemical ones. 510
 It was reported that eupodia, being F-actin-containing 511
 cortical structures, are formed locally in invasive loco- 512
 motion in *Dictyostelium* at the base of a lamellipodium 513
 advancing to invade a tight space, and it appears that 514
 the mechanical stress at the leading edge modulates 515
 the structural integrity of actin and its binding proteins, 516
 such that eupodia are formed when anchorage is needed 517
 to boost for invasive protrusion of the leading edge [33]. 518
 It was also reported that podosomes, being invasive 519
 adhesion structures and an F-actin-rich core surrounded 520
 by a ring structure containing proteins such as vinculin 521
 and talin, are formed locally in transformed fibroblasts 522
 by oncogenic Src causing disruption of actin stress fibers 523
 [34,35]. Thus, the *Dictyostelium* eupodia and the podo- 524
 somes are formed to strengthen the adhesion between 525
 the cells and substrate and to bear the mechanical force 526
 exerted by the cell migrating movement, and of course 527
 they should be subjected to this mechanical force during 528
 the cell migrating movement. The actin agglomerates we 529
 observed here are different. As mentioned in the above, 530
 here the depolymerization of stress fibers and agglomera- 531
 tion of actin are initiated at high stress points existing 532
 in the actin network induced by sustained indentation, 533
 i.e., the mechanical force induces the actin agglomerates. 534
 Also, the actin agglomerates do not strengthen the cells 535

since the increase in the cell force response was not observed after the agglomerates were formed. There are two other significant differences between the actin agglomeration observed here and the eupodia and podosomes. First, the actin agglomerates grow significantly with time (as shown in the above), whereas the significant growing of eupodia and podosomes was not observed after they were formed. Second, the actin agglomerates do not seem to be attached to or even located close to the intracellular side of the adhering cell surface, whereas eupodia and podosomes are ventral structures. But, depending on the cell types, all these actin assemblies show different types of mechanobiological response of the cells to their mechanical force/stress environments, although maybe initiated and formed by different mechanisms.

Mechanical modifications of the actin network can be mediated also by hyperosmotic stresses. Accumulation of actin filaments along the margin of the *Dictyostelium* cells was observed after the contraction of these cells in response to hyperosmotic stress [36], but actin agglomeration well inside the cells was not observed.

- The study also reveals a fundamental difference between a living specimen and a non-living one. The former responds both mechanically and biologically when subjected to a mechanical stimulus, whereas the latter responds only mechanically. The biological response is a result of a cascade of events and signals that we have very little knowledge about. When we measure the response, we measure both the mechanical and the biological contributions (such as actin agglomeration far from the point of indentation) with very little detail on how to decouple one from the other. Nature, most likely, engages both. If the mechanical stimulus is small and is over a short time, the biological response may be negligible. The challenge is in reducing the large parameter space of the biological response to a few important ones so that one can formulate a quantitative approach in interpreting the measured combined response.

5. Conclusions

In this paper, the loading and unloading force response of single living fibroblasts due to large lateral indentation has been measured by MEMS force sensors. The actin network of cell cytoskeleton has been observed in situ by using the GFP technique, which reveals how cells adapt to mechanical indentation and injuries by cytoskeletal reorganization, and provides mechanistic interpretation for the measured force response.

We found: (1) The indentation force response is highly irreversible (i.e., hysteretic) in contrast to the observed reversible and repeatable linear cell force response for the case of large local stretch [10]. (2) All the loading processes show linear force response (sometimes with small initial non-linearity) for indentations as large as the size

of the cells. Plastic yielding follows on further indentation. (3) Cell mechanical response is anisotropic. (4) Upon sustained indentation, the actin network possibly decomposes irreversibly at discrete locations all over the cell and forms circular agglomerates. (5) The spatial distribution of the agglomerates follow the stress fiber orientation of the pre-indented cell. (6) The size of the agglomerates grows with time initially at a fast rate as t^α with $\alpha = 2-3$ followed by a slow rate with $\alpha = 0.5-1$. The transition from fast to slow rates happens abruptly. Actin agglomeration has been observed in living cells subject to ischemic injury, toxic treatment, and at certain developmental stages of larvae, and is believed to be the pre-apoptotic phase of such cells. To the best of our knowledge, this is the first evidence of actin agglomeration due to local mechanical indentation. The finding suggests that living cells, when mechanically indented, injured or compressed, may also initiate actin reorganization and agglomeration with consequent loss of stiffness. In other words, two distinct stimuli, one mechanical and the other biochemical, may result in similar cell signalings. The study may have implications in understanding the role of radial compression on the damage of the endothelium in blood vessels during formation of plaque and progression of atherosclerosis.

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