Reconstitution of Amoeboid Motility in vitro Identifies a Motor-Independent Mechanism for Cell Body Retraction

Katsuya Shimabukuro[1], Naoki Noda[2], Murray Stewart[3], and Thomas M. Roberts[1,]*

[1] *Department of Biological Science, Florida State University, Tallahassee, FL 32306*

[2] *Marine Biological Laboratory, Woods Hole, MA 02543*

[3] *MRC Laboratory of Molecular Biology, Hills Road, Cambridge CB2 2QH, UK*

[*] author for correspondence:

     209 Biomedical Research Facility

     Department of Biological Science

     Florida State University

     Tallahassee, FL 32306

     roberts@bio.fsu.edu

     phone: 850-644-3237

     fax: 850-644-0481

Running head: motor-independent retraction in amoeboid movement

Summary

Crawling movement in eukaryotic cells requires coordination of leading edge protrusion with cell body retraction [1-3]. Protrusion is driven by actin polymerization along the leading edge [4]. The mechanism of retraction is less clear; myosin contractility may be involved in some cells [5] but is not essential in others [6-9]. In *Ascaris* sperm, protrusion and retraction are powered by the major sperm protein (MSP) motility system instead of the conventional actin apparatus [10-11]. These cells lack motor proteins [12] and so are well-suited to explore motor-independent mechanisms of retraction. We reconstituted protrusion and retraction simultaneously in MSP filament meshworks, called fibers, that assemble behind plasma membrane-derived vesicles. Retraction is triggered by depolymerization of complete filaments in the rear of the fiber [13]. The surviving filaments reorganize to maintain their packing density. By packing fewer filaments into a smaller volume the depolymerizing network shrinks and thereby generates sufficient force to move an attached load. Thus, this work provides direct evidence for motor-independent retraction in the reconstituted MSP motility system of nematode sperm. This mechanism could also apply to actin-based cells and may explain reports of cells that crawl even when their myosin activity is compromised.

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Results and Discussion

Generation of MSP comet tails in which protrusion and retraction occur

simultaneously.   Previous work demonstrated that the cytoskeletal dynamics associated

with leading edge protrusion and cell body retraction could be reconstituted

independently in a simple cell-free extract obtained as the 100,000 X g supernatant

(S100) from lysed *Ascaris* sperm [14,15]. For example, addition of 1 mM ATP to five-

fold diluted S100 reconstitutes leading edge protrusion by triggering MSP polymerization

behind plasma membrane-derived vesicles to form a fiber that pushes the vesicles

forward as it grows (Figure S1A, Video S1) [14]. Conversely, treatment of these fibers

with YOP, a tyrosine phosphatase, caused them to disassemble and shrink (Figure S1B),

recapitulating the cytoskeletal dynamics involved in cell body retraction [15,16]. To

explore the mechanism of retraction and its relationship to protrusion we identified

conditions in which both processes can be reconstituted simultaneously but at different

locations within the same fiber. By adjusting either the dilution of the extract or the

concentration of added ATP, about 60% of batches of S100 could be induced to form

fibers that exhibited simultaneous but spatially separated protrusion and retraction. For

example, at 80% S100 and 1 mM ATP, filament assembly occurred at the surface of the

membrane vesicle and pushed the vesicle forward (Figure 1A, Video S2). At the same

time, filament disassembly further rearward resulted in a gradual decrease in optical

density along the length of the fiber. Addition of 50 μM ATP to 20% S100 resulted in

similar behavior. These fibers maintained a nearly constant length as they moved along

and exhibited a pattern very similar to that of the actin comet tails that form behind

intracellular pathogens such as *Listeria* or beads coated with agents that activate assembly of dendritic actin filament arrays [17]. Therefore, we refer to fibers exhibiting this pattern as comet tail fibers.

Retraction force generation in comet tail fibers. Often debris in the S100 stuck to comet tails fibers. Examination of the movement of these debris showed that the disassembling part of the fiber generated sufficient force to move a load. For example, Figure 1B shows a comet tail fiber with a large dark object stuck to its trailing end. This object remained attached and was pulled forward at the same rate as growth at the opposite end of the fiber pushed the vesicle. Smaller objects stuck to fibers appeared as dark specks and could be used as fiduciary marks to chart retraction within the fiber itself. Figure 1C (see also Video S3) shows a comet tail fiber with a speck stuck to the retracting rear region. The distance between that speck and another stuck to the substrate (near the growing end of the fiber at the start of the sequence) decreased with time as the fiber-bound speck was pulled forward. Over the same interval, elongation of the fiber pushed the vesicle at the growing end of the fiber away from the substrate–attached speck. The speck on the fiber moved with respect to the substratum while the distance between it and the vesicle at the growing end of the fiber remained nearly constant. This forward movement of specks attached to the rear of comet tail fibers demonstrated that fiber disassembly was able to perform work.

Fluorescent speckle microscopy [18] confirmed that debris attached at the rear of comet tail fibers were being pulled forward and not moving independently over the fiber.

Addition of 40 nM Cy3-labelled MSP to 80% S100 resulted in the formation of comet

tail fibers with fluorescent speckles (Figure 1D, Video S4). Although many speckles at

the rear of the comet tail fiber disappeared quickly, probably due to loss of fluorescent

MSP subunits during filament depolymerization, several persisted long enough to chart

their movement. Those near the front of the fiber remained stationary relative to the

substratum as the vesicle moved away. By contrast, speckles at the rear moved forward

along with the fiber. Thus, the behavior of these speckles showed that the MSP filaments

themselves, like attached objects, are pulled forward by forces generated during

retraction.

The transition from protrusion to retraction. Generating comet tail fibers allowed us to

probe the mechanism of retraction by exploring how the filament network changed as it

switched from growth at the front to retraction at the rear. We used correlative confocal

fluorescence and electron microscopy to measure changes in the mass and packing

density of the filament network along the length of comet tail fibers grown in the

presence of Cy3-MSP (Figure 2A, B). The total fluorescence intensity in cross-sections

generated from z-dimension stacks (Figure 2C-E) provided an estimate of the filament

mass at defined positions along the length of the fiber (Figure 2G). The same fibers

were identified in platinum replicas and used to measure fiber diameter at the same

positions (Figure 2F). This allowed us to estimate filament density as the total Cy3-MSP

fluorescence per unit cross-sectional area (Figure 2G). If the filament network did not

reorganize during retraction the total filament mass and packing density should decrease

together. However, we found that near the rear of the fiber, where retraction was nearly

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complete, on average cross-sections retained only 5% of the of the total fluorescence observed at the front of the fiber but still had 45-50% of their initial fluorescence density (Figure 2G,H). These data suggest that filament packing density is at least partially conserved during retraction.

EM tomography of platinum replicas of comet tail fibers confirmed this conservation of packing density. We generated tomograms at the front, middle, and rear of the fiber (Figure 3A-D) that we then used to track and measure every filament in each tomogram (Figure 3E-G). As expected, the number of filaments per region decreased substantially with distance from the vesicle so that at the rear of the comet tail, where retraction was nearly complete, the number of filaments was only ~10% of that at the front where retraction had not yet started (Figure 3H). Surprisingly, the average lengths of filaments at the front and rear of the fiber were indistinguishable (Figure 3I), suggesting that whole filaments are lost in an all-or-none depolymerization rather than by all filaments gradually shortening. Analogous catastrophic depolymerization of actin filaments has been observed in *Listeria* comet tails and in *Xenopus* egg extracts [19].

We estimated filament packing density in cross-sectional views generated from the tomograms (Figures 3E-G) by measuring the area within the perimeter of the filament-occupied space in each region and counting the number of filaments within that perimeter. This analysis showed that although the numbers of filaments in the middle and back region were only ~40 and ~10%, respectively, of that in the front (Figure 3I), the filament packing density was similar in all three regions (Figure 3J). We were unable

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to identify additional platinum replicas of comet tail fibers suitable EM tomography but did examine several comet tail fibers by conventional EM (Figure 3K-O).   The diameter of each of these fibers narrowed dramatically from the middle to the back, where depolymerization and retraction occurred, and the surviving filaments in the rear of the fiber were packed tightly, features consistent with conservation of packing density.

Filament rearrangement during retraction. Retracting fibers could conserve packing density either by retaining filaments in the fiber core while peripheral filaments depolymerize, or by repacking the filaments that survive depolymerization to compensate for the loss of their neighbors. To discriminate between these possibilities, we examined comet tail fibers by polarization microscopy using an LC-Polscope system. This optical system produces computed images that show the magnitude of retardance at each pixel as shades of gray while the orientation of birefringence is displayed as a series of lines (see Supporting Information; Figure S1) [20-22]. Calibration of the slow axis of birefringence using arrays of MSP filaments of known orientation showed that the orientation of birefringence generated by LC-Polscope microscopy parallels the long axis of the MSP polymer (Figure S2).

To determine if the pattern of birefringence orientation in fibers changed during retraction, we captured successive images of a specific region of a comet tail fiber as it underwent retraction (Figure 4A). The boxed region shown in this sequence was not retracting at the start of this interval but had nearly completed retraction by the end. Initially, much of the birefringence in this region was oriented approximately normal to

the fiber axis with much less oriented axially (Figure 4B). As this region began to retract, its retardance decreased and its birefringence orientation shifted toward the fiber axis (Figure 4A). The fraction of axis-aligned pixels (those within 15° of the long axis) increased over 4-fold (Figure 4C) and we observed orientation parallel to the axis at 110 sec that was not present earlier (Figure 4B). Thus, the change in the birefringence pattern is not due to selective survival of axis-aligned filaments and, instead, fiber retraction involves an active rearrangement of at least some of the surviving of filaments.

Mechanism of motor-independent retraction. Although actomyosin contraction contributes to cell body retraction in some crawling cells [5], several recent reports have shown that cells can still crawl when their myosin activity is compromised [6-9]. These observations suggest that motor-based contraction may not account completely for retraction even in actin-based cells. Sun and colleagues [23] have elaborated a mechanism for generating contraction that does not require the mechanochemical activity of conventional motor proteins. According to this model, the free energy of a filament network is minimized when the entropic contribution based on filament dispersion is balanced by enthalpic energy resulting from attractive interactions between filaments; that is, the network has an optimum packing density. This model envisions that depolymerization of filaments will reduce the packing density of the network so that it becomes suboptimal.   Provided that that the interaction between filaments is transient, the remaining filaments will move closer together (thus increasing the packing density) so that the overall energy of the system decreases until the enthalpic gain and entropic loss balance. Thus, minimizing the energy of the filament network causes it to contract to its optimum packing density.

The relatively constant packing density observed along retracting MSP fibers provides direct experimental evidence to support an optimum packing model and suggest the following mechanism for depolymerization-mediated retraction in comet tail fibers. Retraction starts with catastrophic depolymerization of individual filaments. As filament loss proceeds, the interactions between the filaments decrease allowing them to move more freely. The resulting increase in the movement of the surviving filaments would result in collisions between filaments and the formation of new interactions. As a result, the filament network in the fiber would tend to rearrange spontaneously bringing filaments closer together to restore the packing density to its optimum value. Consistent with this hypothesis, as comet tails fibers retract, >90% of filament mass is lost, but ~50% of their pre-retraction packing is retained. Mathematical modeling of MSP fiber disassembly and retraction as a comparable two-step process involving filament detachment from the network followed by depolymerization [24] predicts a strikingly similar retention of 50% of initial packing density in the fiber. This conservation of filament packing density with progressively fewer filaments would cause the network to shrink and pull the rear part of the network and any attached objects forward. In crawling sperm, disassembly of the cytoskeleton occurs at the base of the lamellipod where it joins the cell body [13]. Although the way in which the cell body of is mechanically coupled to the MSP cytoskeleton has not been defined, shrinkage of the cytoskeletal network like that observed in comet tail fibers could pull the cell body forward. This mechanism is consistent with mathematical modeling of *C. elegans* sperm locomotion, which predicts that disassembly-induced cytoskeletal stress can account for cell body retraction [25].

Although our results were obtained using the specialized *Ascaris* sperm (MSP) motility system, the principles identified here are likely to also be applicable to more general actin-based cell motility. Our results highlight how a pulling force can be generated by filament depolymerization and rearrangement without the necessity for motor proteins, consistent with recent work that has indicated that such mechanisms may also be important in actin-based cells [9]. Moreover, our results provide empirical evidence to support the proposal [23] that balancing enthalpic interaction and entropic dispersion of filaments within a network can power retraction in amoeboid cells.

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Figure Legends

Figure 1. Comet tail fiber dynamics and force generation. **A**. Comet tail fiber in which MSP polymerizes at the front and disassembles toward the rear at about the same rate so that the fiber maintains its length over time. **B.** Time lapse images of a comet tail fiber with a dark piece of debris attached to its trailing end. This "cargo" is pulled forward at about the same rate as the movement of the vesicle at the opposite, growing end of the fiber. **C.** A comet tail fiber moving past a dark speck (arrow) attached to the glass. A second speck (arrowhead) attached to the rear part of the fiber is pulled forward, toward the substrate-attached speck, while the vesicle is pushed away. This pattern illustrates the simultaneous growth and shrinkage that occur in comet tail fibers. **D.** Time lapse images of a comet tail fiber grown in S100 supplemented with a small amount of Cy3-MSP to produce a fluorescent speckle labeling pattern. Two horizontal white lines have been added to assist in following the movement of individual speckles. One speckle (white arrow) in the rear part of the fiber moves forward due to retraction in the region while a second (black arrow) near the vesicle remains stationary as the vesicle moves away. Numerals indicate elapsed time in min (**A-C**) or sec (**D**). Bar, 2 μm in **A-C**, 5μm in **B**. (see also Movies S1-S4; Figure S1)

Figure 2. Filament packing density is partially conserved during retraction. **A**. DIC image of a comet tail fiber grown in the presence of Cy3-MSP. **B** The same fiber viewed by confocal fluorescence microscopy. The fiber is show as a projection of a stack of images through the z axis. **C-E**. Cross sections at the positions indicated by the dashed lines in panel A. The images were generated from the same z stack used to produce panel

B.. Linear contrast adjustment was applied to make all pixels containing fluorescent signal visible   **F.** Changes in the total filament mass (dashed line) and the relative filament density (solid line) at selected points along the length of the comet tail fiber. Total filament mass was calculated as the total fluorescence intensity in arbitrary units in all pixels within each cross section of the fiber. The relative filament density was determined by dividing the total fluorescence intensity by the area of the cross section. That area was determined by using EM images to measure the radius the fiber at each position where a cross section was obtained and then calculate its cross sectional area. Both total filament mass and relative filament density were normalized to the values obtained for a cross section near the front of the fiber and plotted against distance from the vesicle. **G.** Plot of normalized relative filament density versus the percent loss of fluorescence (100 – normalized total fluorescence) for 194 cross sections obtained from 8 fibers. The solid line is a plot of the linear regression. Note that even when the total filament mass in cross sections is almost completely gone the fibers retain ~45% of their filament density.

Figure 3. Platinum-carbon replica and electron tomograms of a comet tail fiber. **A.** Panoramic view of the comet tail fiber. Bar, 1 μm. **B-D.** Higher magnification views of three regions of identical length from the non-retracting front (**B**), actively retracting middle (**C**), and nearly completely retracted rear (**D**) of the comet tail fiber in **A**. Bar, 200 nm. **E-G**. Cross sectional views of 3D models produced by tracking the filaments (yellow) in the regions shown in **B-D**, respectively, in tomographic reconstructions of the fiber. The view shown is a cross section looking down the fiber axis from the forward

face of each region. **H.** Graph showing the average filament length in tomograms in **E**, **F**, and **G**. The number of filaments measured are in parentheses. **I**. The total filament mass within each of the three tomograms determined as the number of filaments in each normalized to number in the front region tomogram. **J**. The filament density in tomograms determined as the total number of filaments divided by the volume, which was estimated as the filament-containing area of the cross section times the length of the tomogram along the comet tail axis.

Figure 4. Rearrangement of filaments during fiber retraction detected by LC-PolScope microscopy (see Figure S1 for an explanation of LC-Polscope computed images). **A**. Time lapse sequences of retardance (upper) and slow axis orientation (lower) images. Numerals indicate elapsed time in sec. Bar = 2 μm. In the retardance images the magnitude of retardance at each pixel is shown as shades of grade according to the scale at the upper left. In the lower set of panels the orientation of the slow axis of birefringence at every fifth pixel is displayed as a line. Note that the lines do not represent individual filaments. **B.** Plot of the relative frequency distribution of orientation lines in the boxed region in **A** at 0 and 110 sec. The plots were generated from 708 and 162 data points for 0 and 110 sec, respectively. **C.** A graph showing the percentage of orientation lines in the boxed region in **A** within ± 15° of the fiber axis over time. (see also Figure S2)

A–G: electron micrographs and filament tracings. Scale bars in A, D, and G.

H

Mean Filament Length (μm)

(535) (230) (51)

Front Middle Back

I

Normalized Filament Mass

Front Middle Back

J

Normalized Filament Density

Front Middle Back

K L M N O

A

0.8 nm

0

0 60 110

B

Frequency

0.3

0.2

0.1

0 s

110 s

-90 -60 -30 0 30 60 90

Orientation of
Birefringence (deg)

C

Degree of
Axis align (%)

50

40

30

20

10

0 20 40 60 80 100 120

Time (sec)