

1. **Pick up** Name Folder

- Pick up name folder and set it up at seat.

2. **Sit** with your lab group.

- laptops almost closed (avoid distracting)

3. **Clicker** Attendance

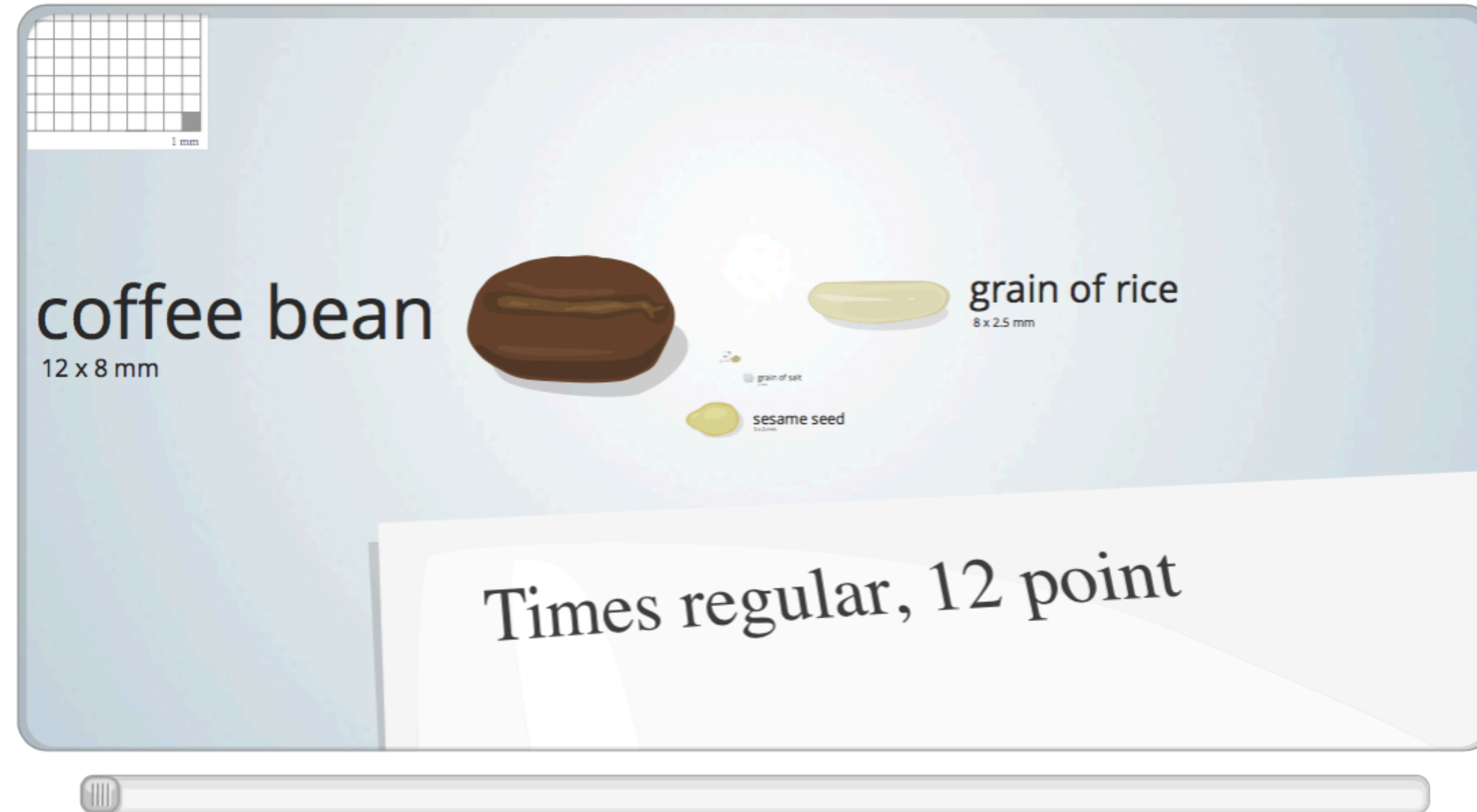
- Launch your Top Hat, and get ready to click.



Budgeting homework time (60 min): Chapter 8, section 8.2 (first half) is 2498 words in length with four data tables and figures that require thinking and notetaking for the Trifecta. Reading at 200 words per minute would mean the section might take 13 minutes to read. Of course, when done properly, when you pause to review figures, try Integrating Questions, and take notes, this assignment will take you more like 60 minutes.

1. _____ **For Thursday's lecture**, read **Chapter 8: Cell Structure and Function**, section 8.2: "Why aren't there giant cells?" and as you read it be sure to take handwritten notes on the first half (**2498 words**). You can stop taking detailed notes when you begin reading the light blue box denoting the second section on "Calculated cell limitations". Just read that short section for deeper understanding.
2. _____ **Try to answer some Integrating Questions and Review Questions**. As you read the ICB textbook always attempt to test yourself a little, answer at least one of each set.
3. _____ (Trifecta): **Prepare to explain (aloud) Figures 8.14, 8.15 and 8.16 in class** (Purpose, Methods, Findings)
4. _____ **Advanced**: Take a peek at some of the published research papers in the Bibliography at the bottom of the page with the goal to find the original figure you studied here in the reading and where it is in the paper?

Cell Size and Scale



Meter	Centimeter	Millimeter	Micrometer	Nanometer	Angstrom	Picometer
10^0 m	10^{-2} m	10^{-3} m	10^{-6} m	10^{-9} m	10^{-10} m	10^{-12} m
1 m	0.01 m	0.001 m	0.000001 m	0.000000001 m	0.0000000001 m	0.000000000001 m
	1/100 m	1/1,000 m	1/1,000,000 m	1/1,000,000,000 m	1/10,000,000,000 m	1/1,000,000,000,000 m
	hundredth of a meter	thousandth of a meter	millionth of a meter	billionth of a meter	ten billionth of a meter	trillionth of a meter

<https://learn.genetics.utah.edu/content/cells/scale/>

trifecta+

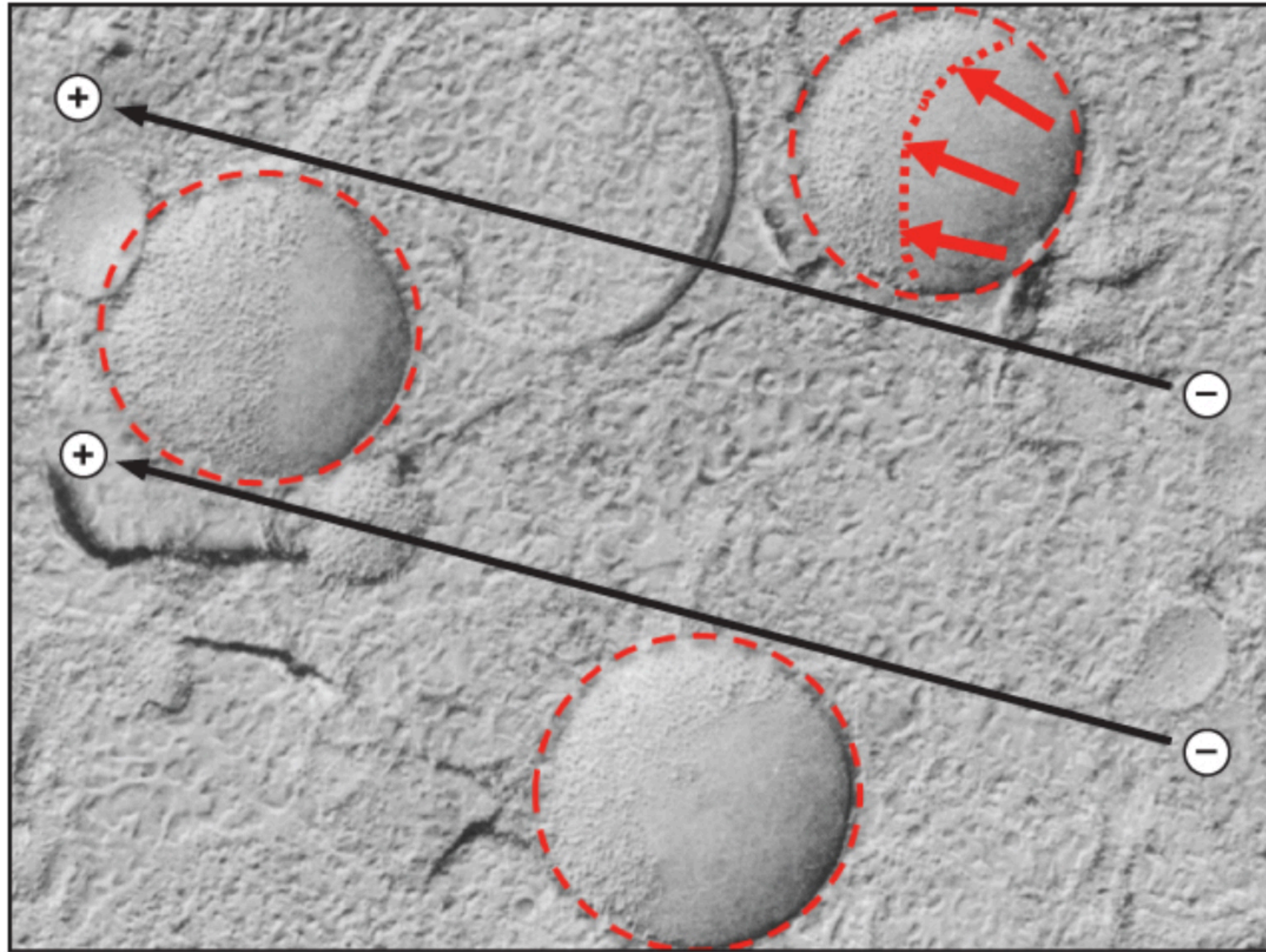


Figure 8.14 Mobility of membrane proteins. Three cells are outlined with red dashed lines. The electrical field is represented by black lines with the indicated polarity. In the top right cell, the movement of proteins is indicated with red arrows, and the edge of the accumulated proteins is indicated with a dotted arc. From Sowers *et al.*, 1981, modified from their figure 6. Sowers, Arthur E. and Charles R. Hackenbrock. 1981. Rate of lateral diffusion of intramembrane particles: Measurement by electrophoretic displacement and rerandomization. PNAS. Vol. 78(10): 6246-6250.

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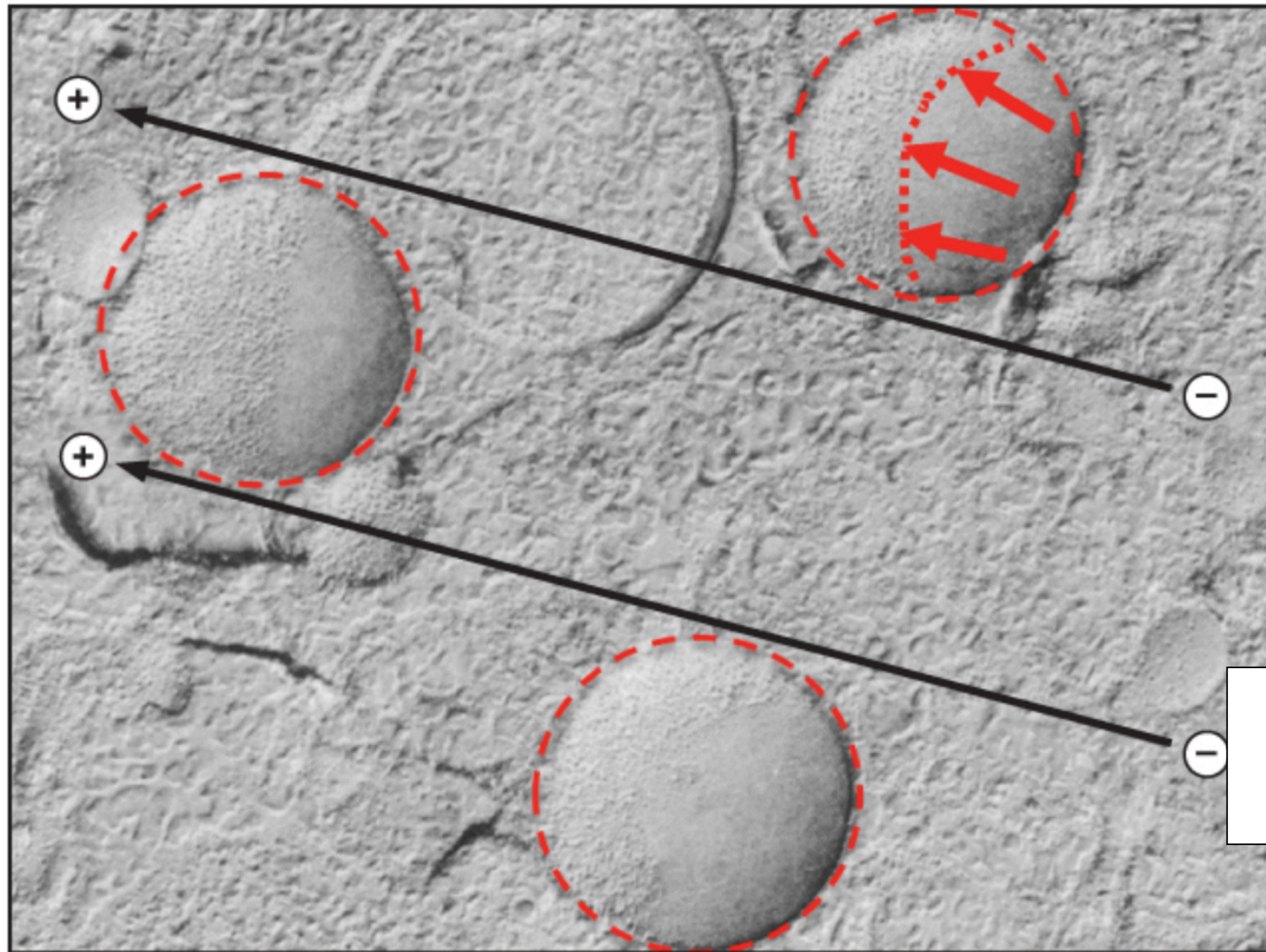
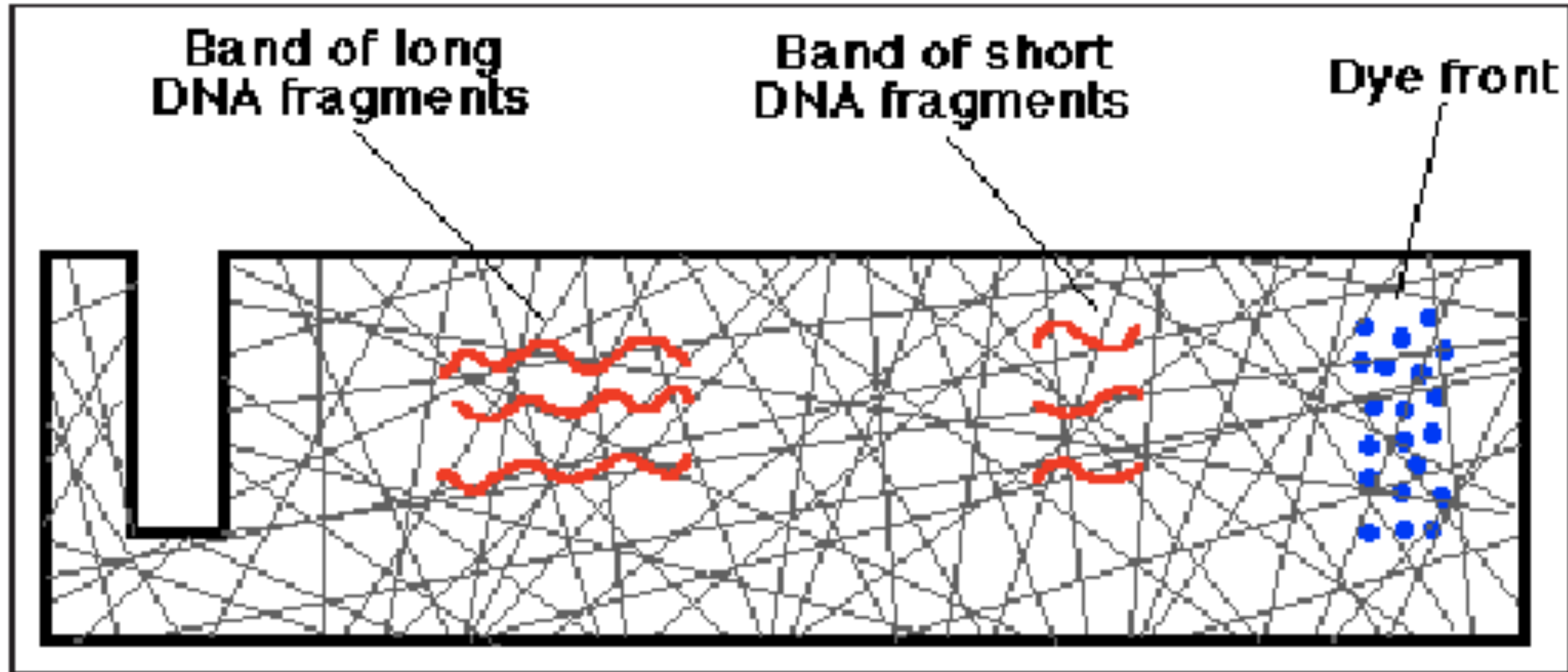


Figure 8.14 Mobility of membrane proteins. Three cells are outlined with red dashed lines. The electrical field is represented by black lines with the indicated polarity. In the top right cell, the movement of proteins is indicated with red arrows, and the edge of the accumulated proteins is indicated with a dotted arc. From Sowers *et al.*, 1981, modified from their figure 6. Sowers, Arthur E. and Charles R. Hackenbrock. 1981. Rate of lateral diffusion of intramembrane particles: Measurement by electrophoretic displacement and rerandomization. PNAS. Vol. 78(10): 6246-6250.

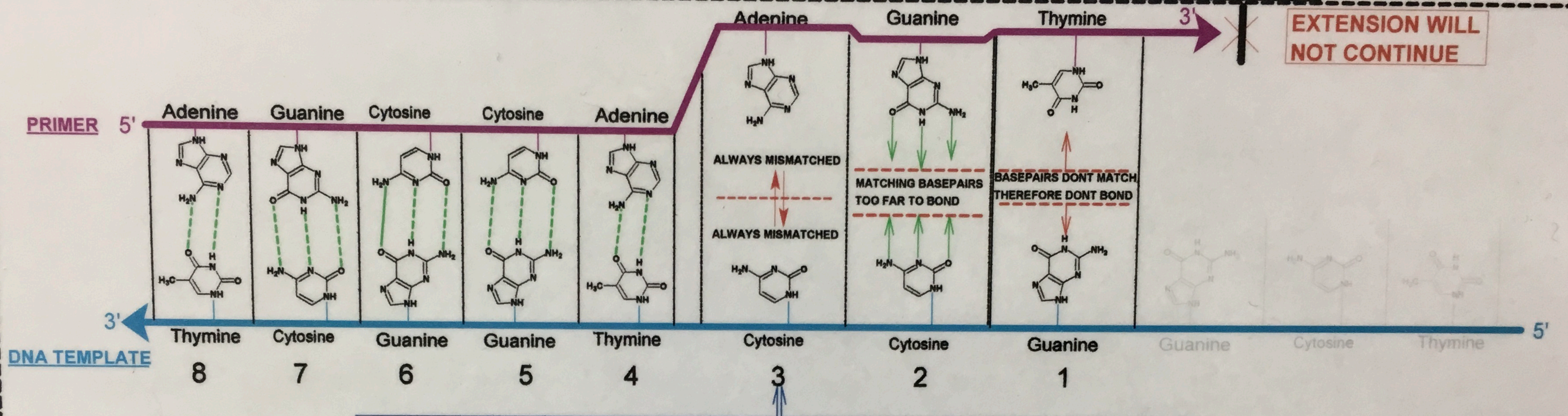
HYPOTHESIS?

Draw what the Hypothesis was
(“Show and Tell” on projector)



HYPOTHESIS (agarose gel electrophoresis)

THE YAKU-BONCZYK PRIMER DESIGN



WHEN USING THE YAKU-BONCZYK PRINCIPLE TO CONSTRUCT PRIMERS, THE POSITION THAT WOULD CORRESPOND TO 3 BASE-PAIRS IN FROM THE 3' POSITION OF THE PRIMER SEQUENCE IS ALWAYS CODED TO MISMATCH!

FIGURE 1: MUTANT-SEEKING PRIMER FOR WILD-TYPE DNA

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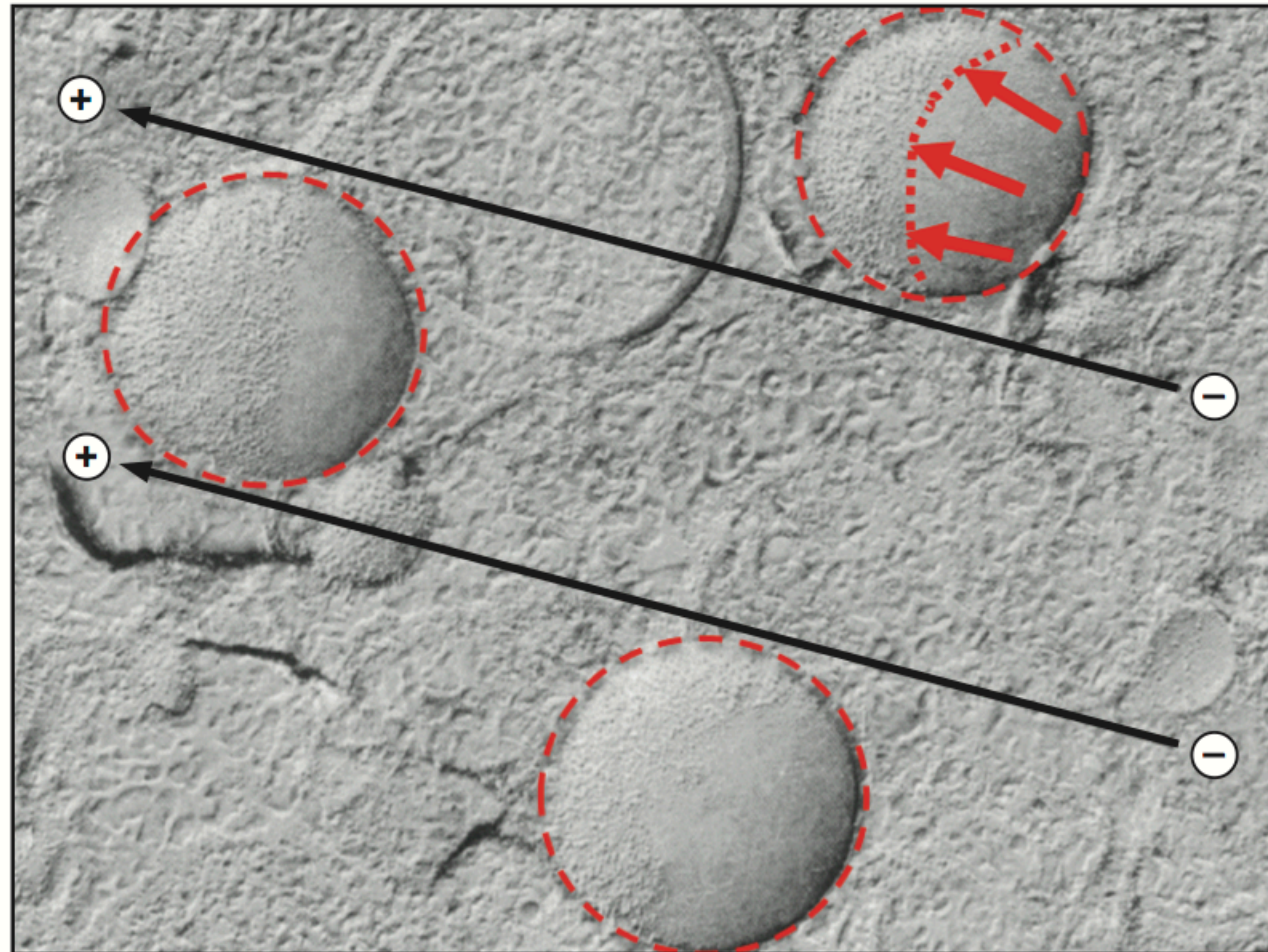


Fig. 8.14

Methods (more details)

Rate of lateral diffusion of intramembrane particles: Measurement by electrophoretic displacement and rerandomization

(lateral diffusion coefficient/integral proteins/freeze-fracture)

ARTHUR E. SOWERS AND CHARLES R. HACKENBROCK

Laboratories for Cell Biology, Department of Anatomy, School of Medicine, U

Communicated by Walther Stoeckenius, June 8, 1981

ABSTRACT A method combining electrophoresis and freeze fracture electron microscopy is described; the method was used to determine the lateral diffusion coefficient of intramembrane

Cell Biology: Sowers and Hackenbrock

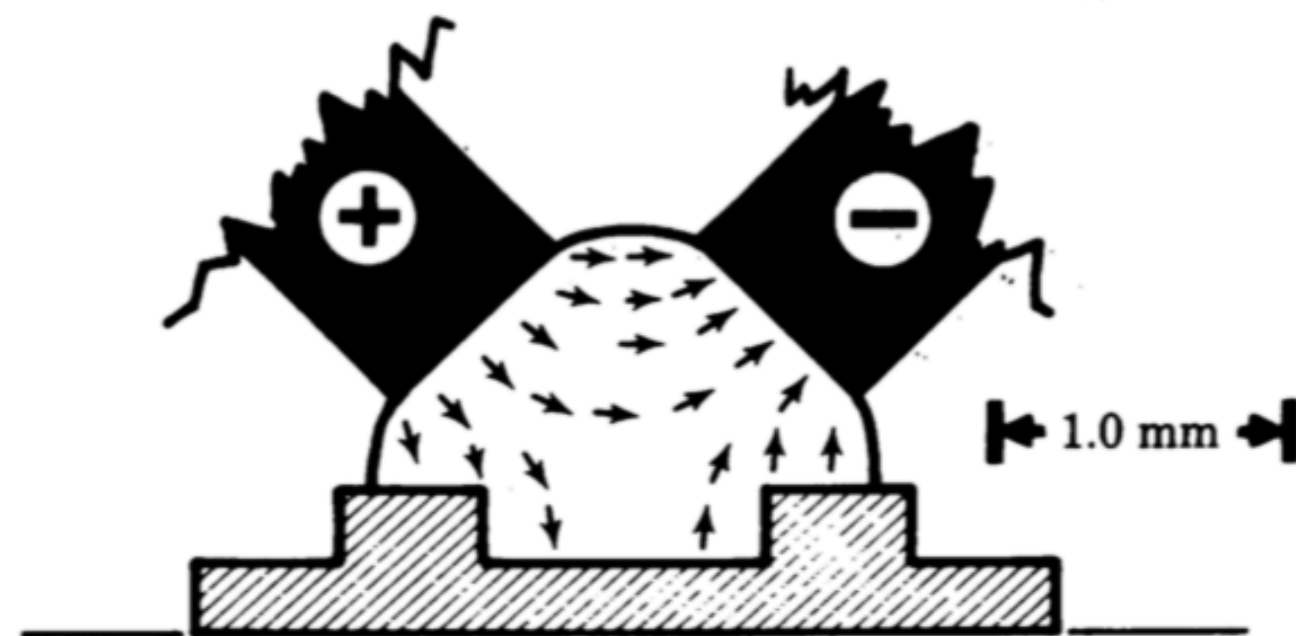


FIG. 1. Relationship of membrane micro-suspension of a standard Balzers gold freeze-fracture specimen holder to Ag/AgCl wire electrodes and the expected electric field as represented by vectors.

mean-square displacement of 57 nm in 10 msec is predicted.

Cell Biology: Sowers and Hackenbrock

Proc. Natl. Acad. Sci. USA 78 (1981) 6249

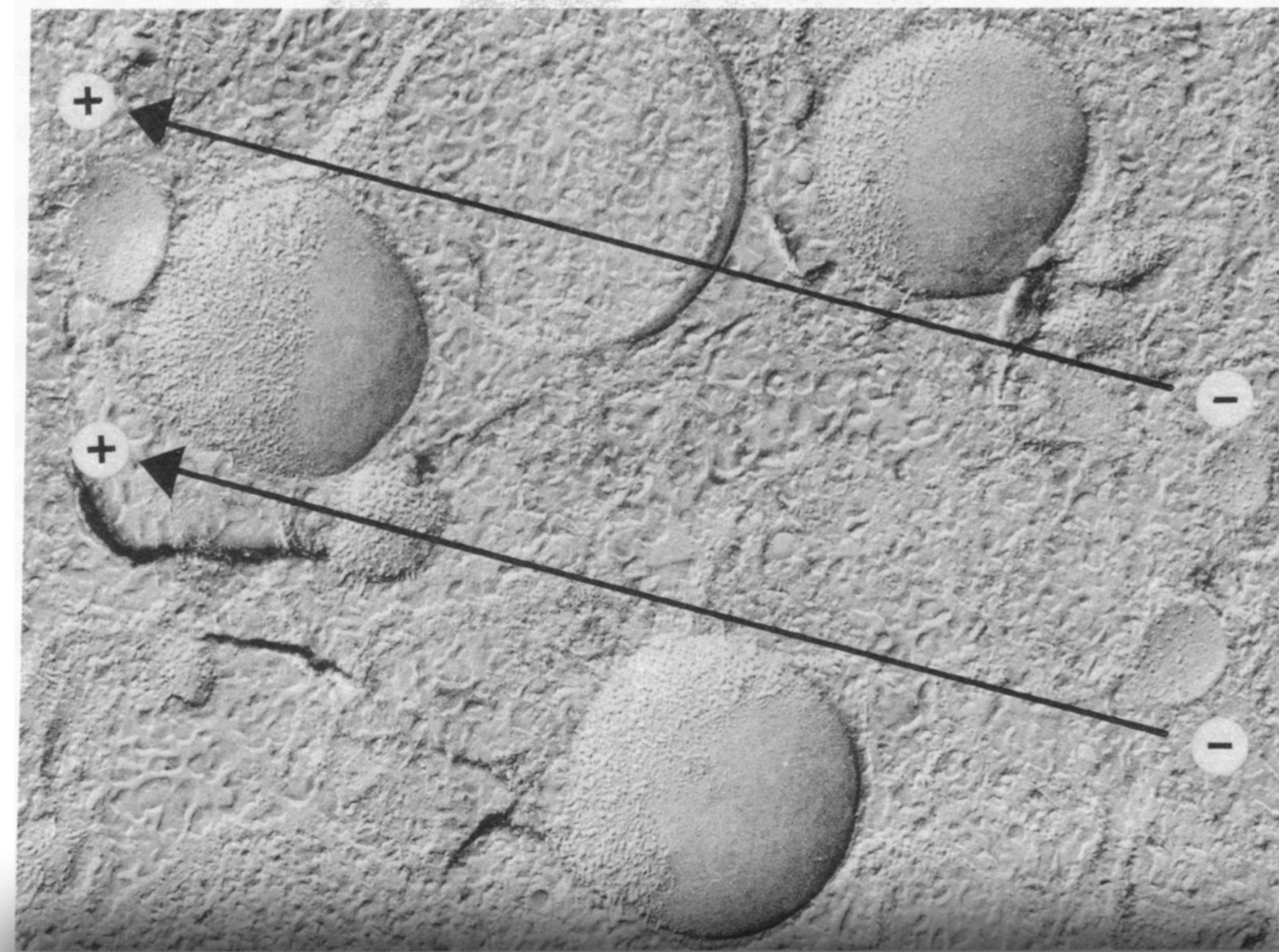
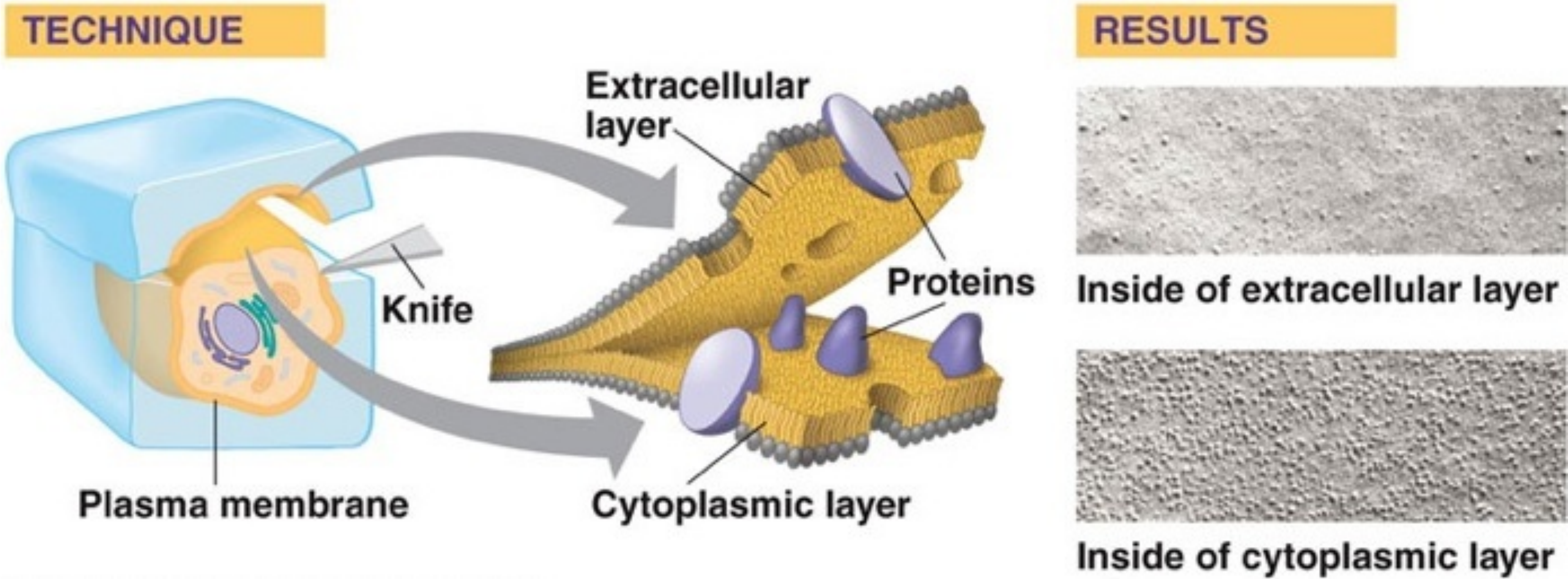


FIG. 6. Membranes frozen after a 3.0-sec exposure to an electric field of about 65 V/mm. Arrows indicate direction of migration of intramembrane particles towards the positive electrode. Convex (*Upper*), and concave and convex (*Lower*) fracture faces reveal migration of intramembrane particles toward the positive electrode. (*Upper*, $\times 55,000$; *Lower*, $\times 62,000$.)

Freeze fracture method (electron microscopy)



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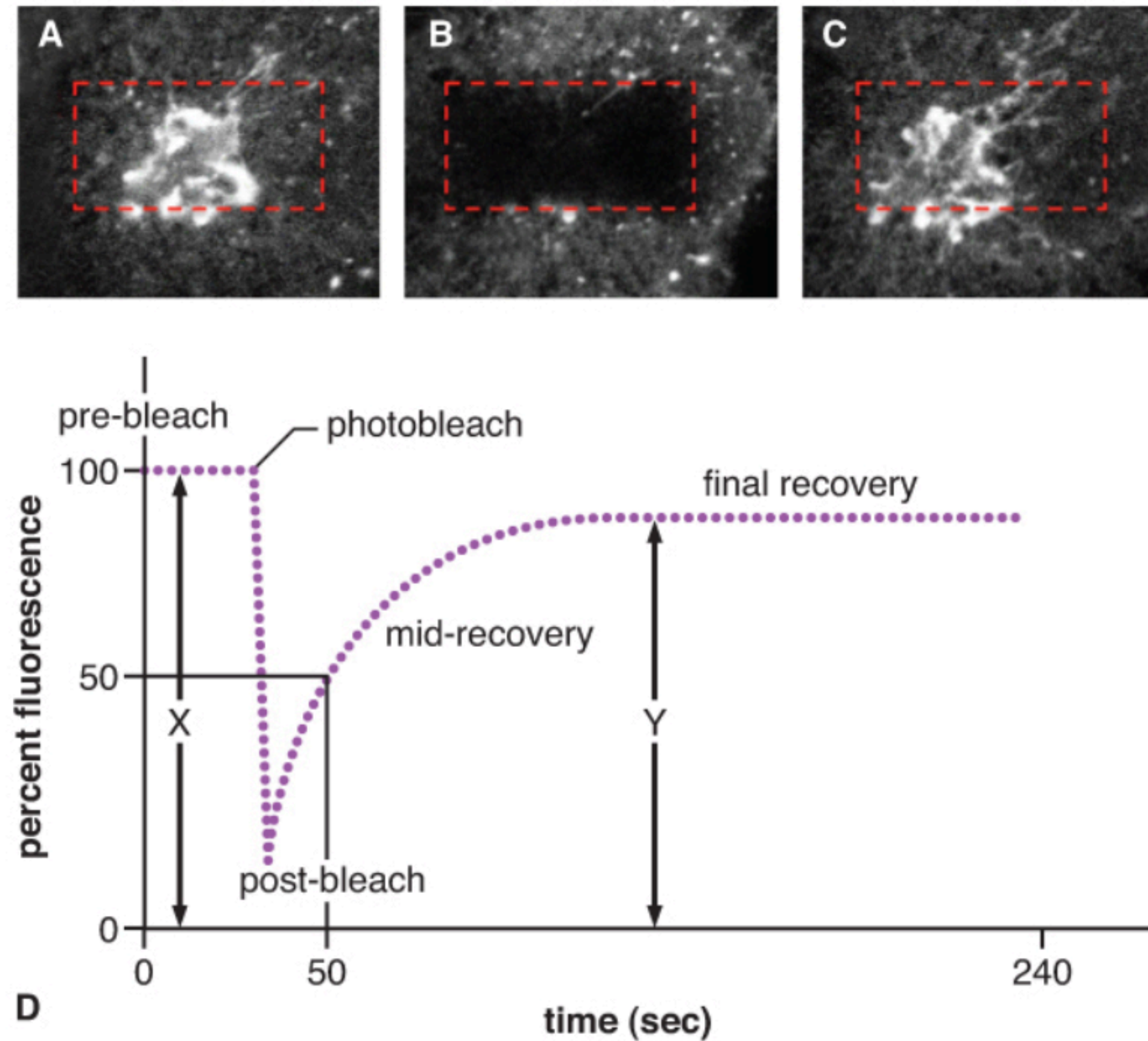


Figure 8.15 FRAP method. Cell immediately before **(A)** and after **(B)** photobleaching inside the boxed area. **C**, After about 2 minutes, the bleached area has recovered its glowing due to mobile proteins. **D**, Diagram of FRAP data showing key steps. [A FRAP movie](#) of the experiment is available online. A-C. Modified from a movie published as part of Presley JF, Cole NB, Schroer TA, Hirschberg K, Zaal KJ, Lippincott-Schwartz J. 1997. ER-to-Golgi transport visualized in living cells. *Nature* 389: 81-85. D. Common Knowledge

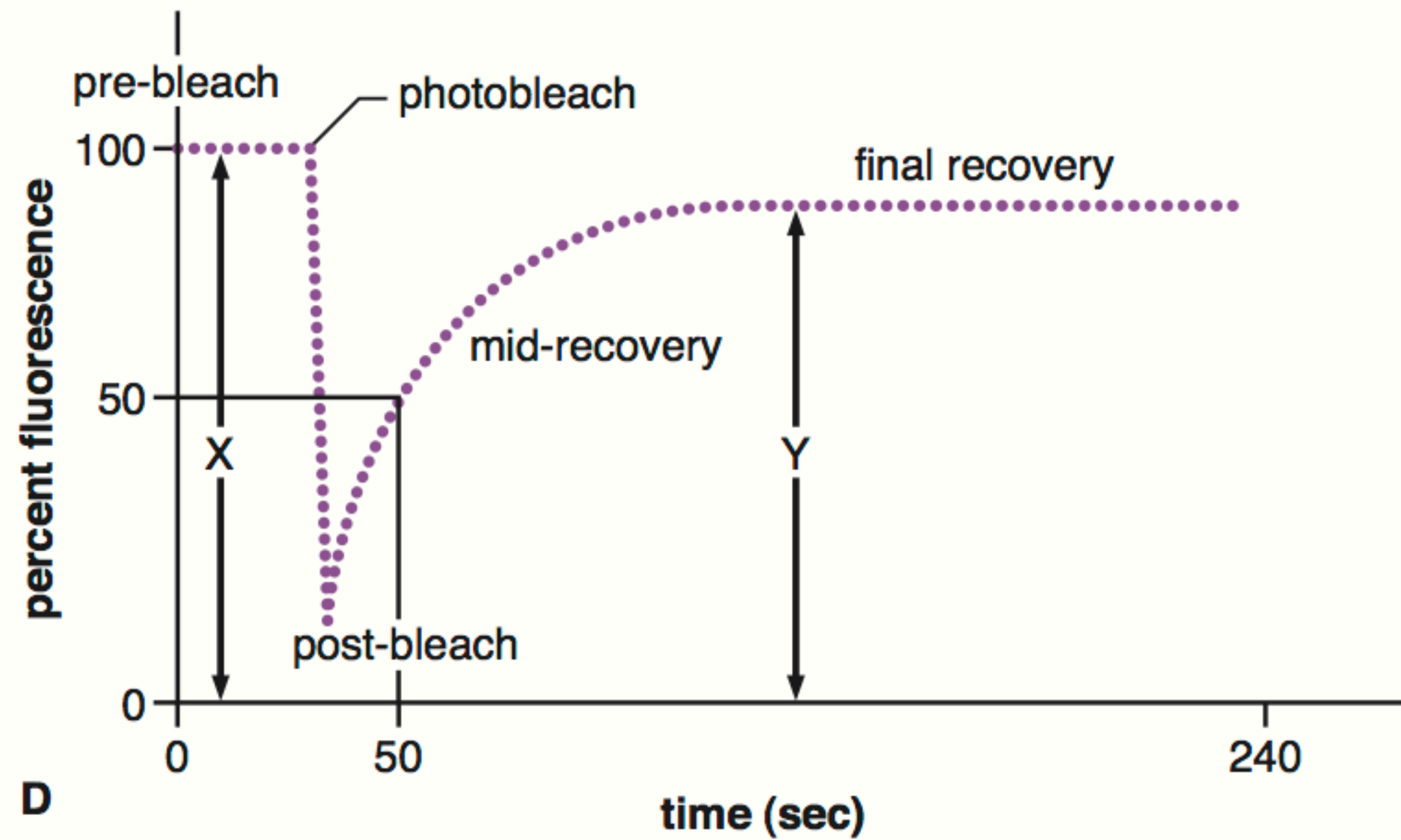
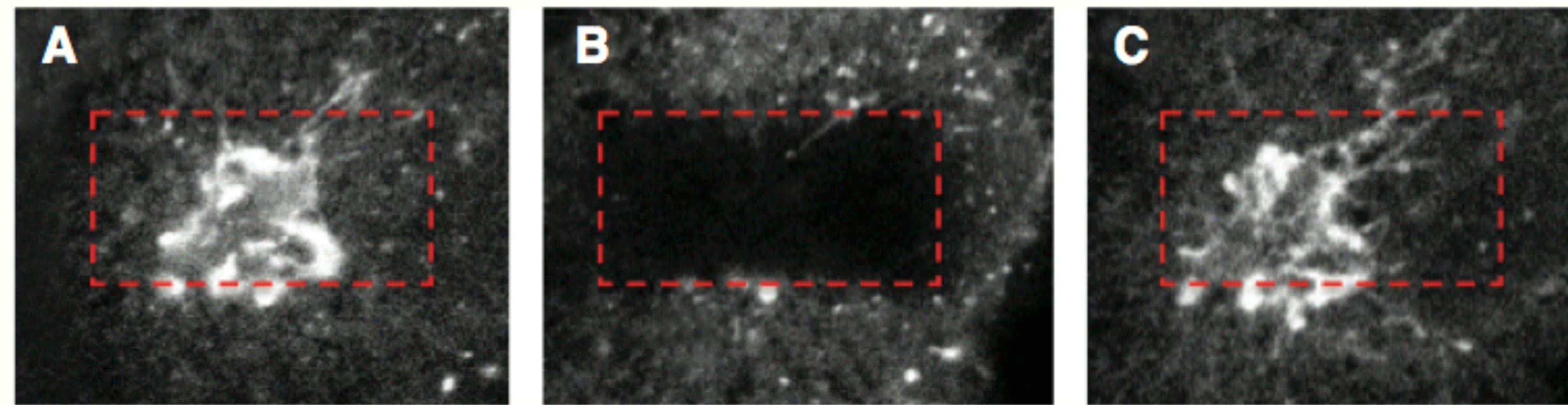


Fig. 8.15

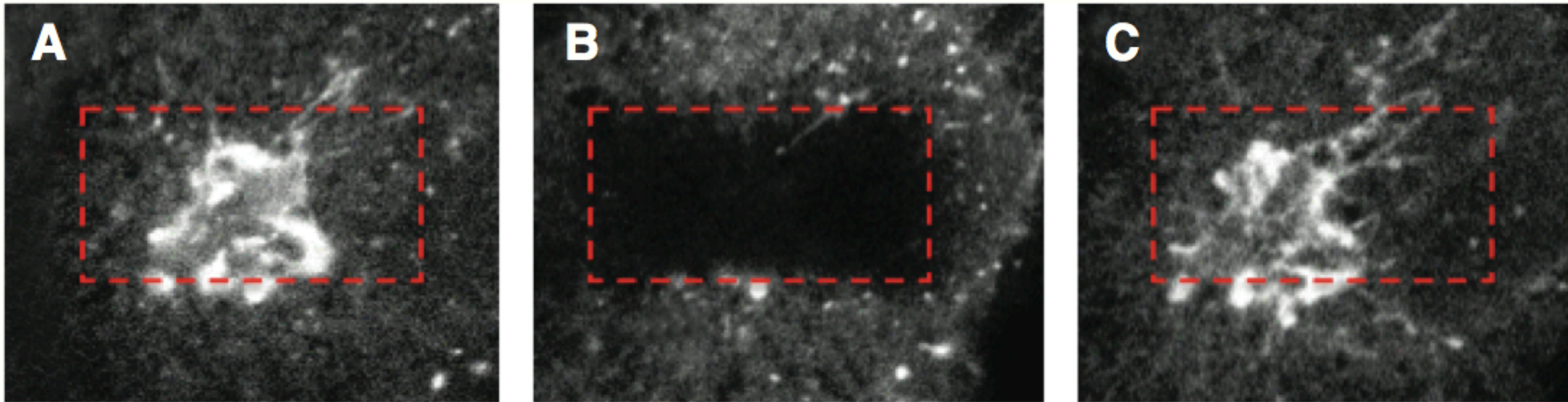
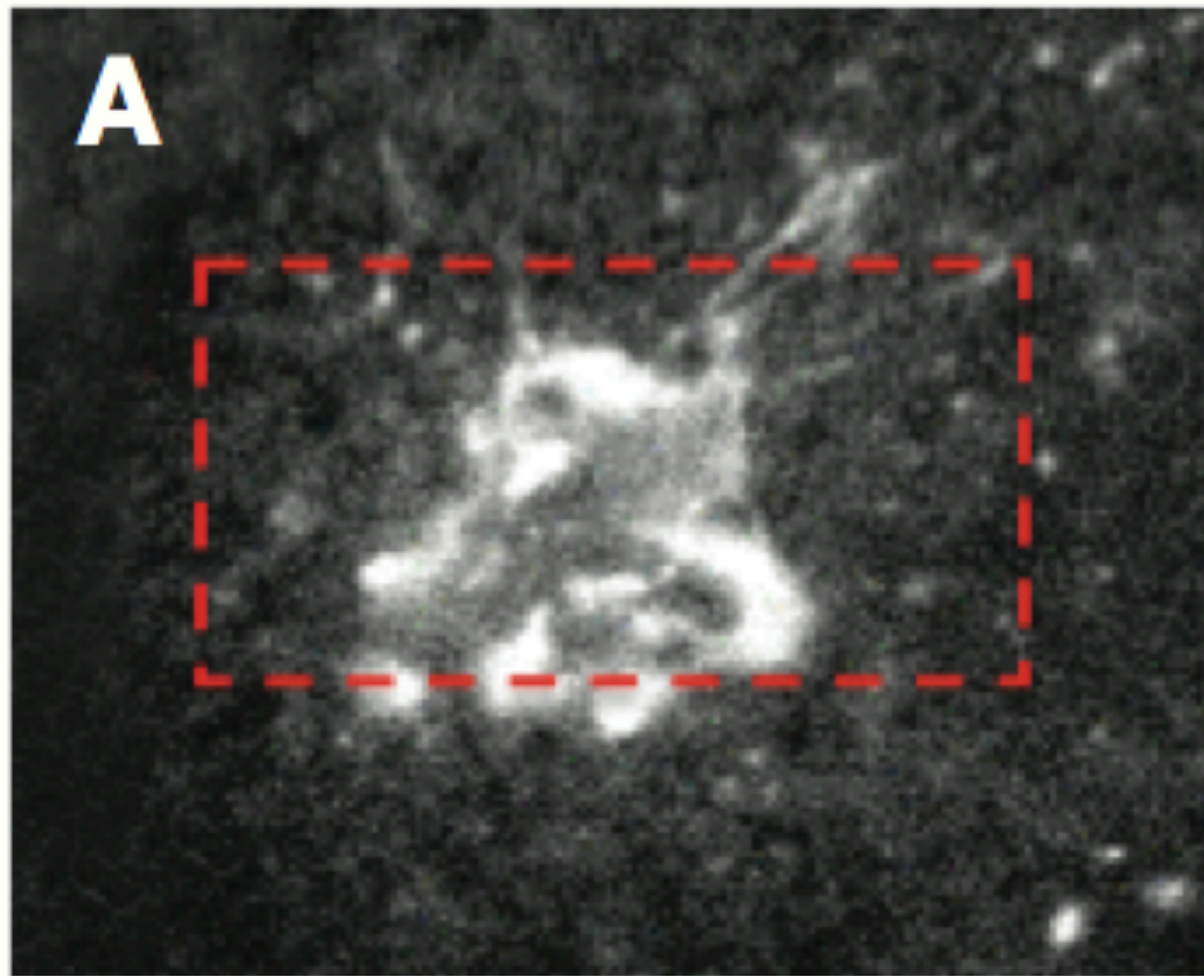


Fig. 8.15

A-C modified from Presley et al., 1997; B: original art
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FRAP Measures Protein Movement



visualize proteins
targeted for photobleaching

Fig. 8.15

FRAP Measures Protein Movement

photobleach proteins

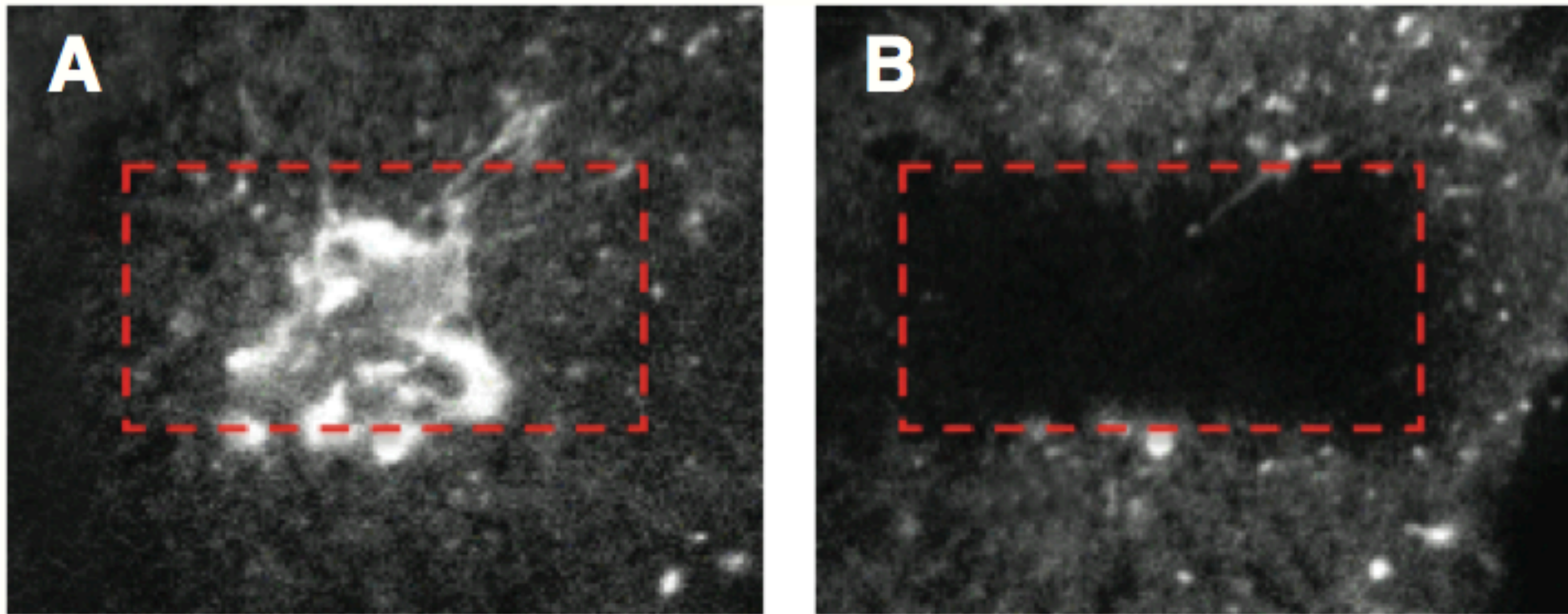
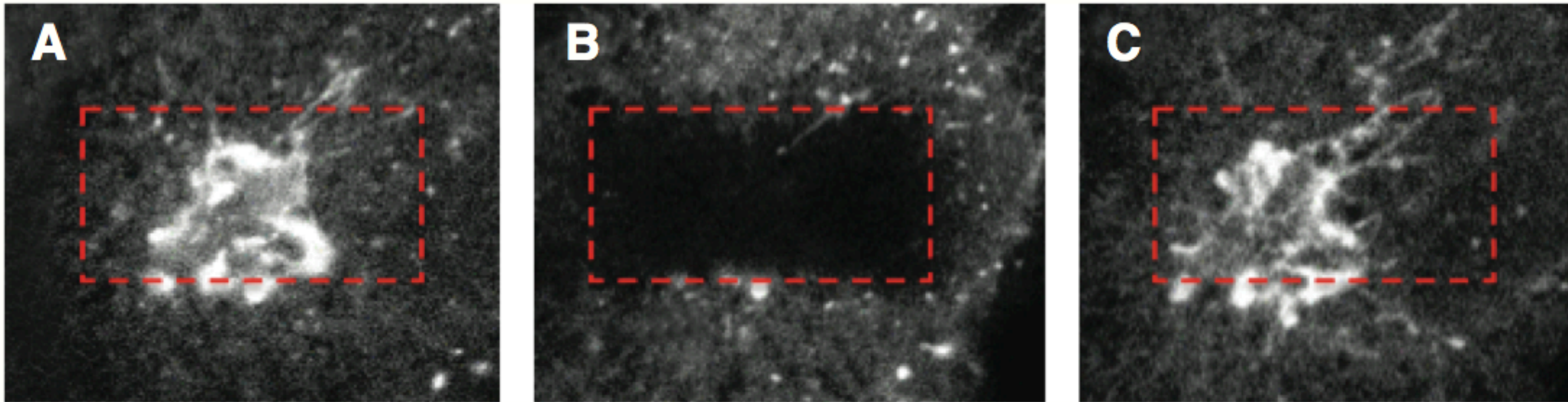


Fig. 8.15

FRAP Measures Protein Movement



watch unbleached
proteins move in

Fig. 8.15

ER-to-Golgi transport visualized in living cells

John F. Presley, Nelson B. Cole, Trina A. Schroer*, Koret Hirschberg, Kristien J. M. Zaal & Jennifer Lippincott-Schwartz

Cell Biology and Metabolism Branch, National Institute of Child Health and Human Development, Building 18T, NICHD, NIH, Bethesda, Maryland 20892, USA

* Department of Biology, The Johns Hopkins University, Baltimore, Maryland 21218, USA

Newly synthesized proteins that are synthesized in the endoplasmic reticulum (ER) are funnelled through the ER for transport to their different destinations. Recent approaches have elucidated the mechanisms of such transport¹⁻³ and have established that transport intermediates are often large and move at high velocity *en route* to the Golgi complex. We have visualized ER-to-Golgi transport of VSVG-GFP in living cells. Upon export from the ER, VSVG-GFP forms many differently shaped, rapidly moving intermediates that translocate inwards to the Golgi complex by using the microtubule motor complex of dynein/dynactin. These intermediates are often greater than 1.5 μm in length and move unidirectionally along microtubule tracks at speeds of up to 1.4 μm s⁻¹. Individual pre-Golgi elements were often greater than 1.5 μm in length and moved unidirectionally along microtubule tracks at speeds of up to 1.4 μm s⁻¹. Staining during this period revealed that VSVG-GFP-containing intermediates move towards the Golgi complex in two pre-Golgi structures. A single long tubule was often observed in the motion of the pre-Golgi motor on the tip of the membranes towards the Golgi complex. Variable times after warm-up, a single element containing VSVG-GFP was plotted in Fig. 2a and b. The fluorescence intensity as they moved towards the Golgi that VSVG-GFP molecules were broken into small vesicle carriers, which were then transported into the Golgi region. Visualized upon shifting to 32 °C (Fig. 3a-d and Quicktime movie).

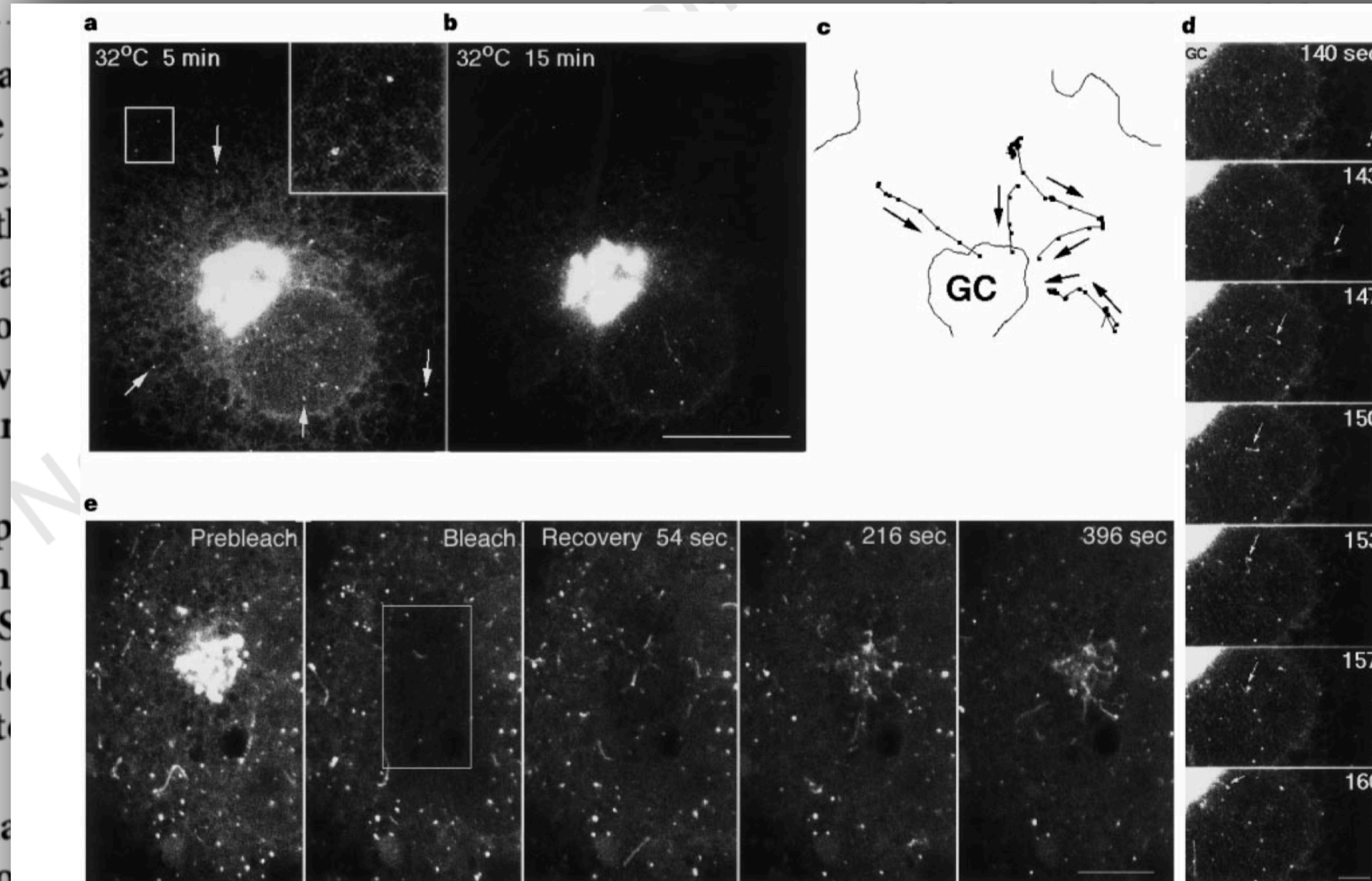


Figure 3 ER to Golgi transport of VSVG-GFP visualized upon shift from 40 °C to 32 °C or in cells whose Golgi area is photobleached. **a-d**, VSVG-GFP-expressing COS cells were incubated for 12 h at 40 °C and then shifted to 32 °C. **a**, Distribution of VSVG-GFP after 5 min at 32 °C. Arrows show examples of pre-Golgi intermediates. **b**, Same cells as in **a** after 15 min at 32 °C. **c**, Schematic diagram of a Golgi complex (GC) with arrows indicating the movement of pre-Golgi intermediates towards it. **d**, Image series showing shape change of a pre-Golgi intermediate as it translocated to the Golgi complex over the time interval shown. Scale bar, 4 μm. **e**, VSVG-GFP-expressing COS cells were incubated for 12 h at 40 °C, shifted to 15 °C for 3 h and then warmed to 32 °C. Fluorescence associated with the Golgi region was photobleached (see Quicktime movie). Within 1 to 5 min after

incubation for 3 h at 15 °C, co-localizing extensively with β-COP (Fig. 1a; 15 °C) and ERGIC53 (data not shown).

To examine how VSVG-GFP is transported from such intermediates into the Golgi region, cells expressing the chimaeric proteins at 15 °C were placed on a microscope stage warmed to 32 °C and fluorescent images were collected at 3.6-s intervals. As shown in Fig. 1b and c (also see Quicktime movie at <http://dir.nichd.nih.gov/CBMB/pb1labob.html>), peripheral pre-Golgi structures containing VSVG-GFP translocated rapidly as units into the centrosomal region where they merged into the large fluorescent area marking the Golgi complex. Figure 1b (centre) maps the movement of several of these structures over the course of 9 min upon warm-up from 15 °C to 32 °C. Such structures moved along straight or curvilinear paths towards the cell centre at speeds of up to 1.4 μm s⁻¹. Individual pre-Golgi elements were often greater than 1.5 μm in

length and moved unidirectionally along microtubule tracks at speeds of up to 1.4 μm s⁻¹. Staining during this period revealed that VSVG-GFP-containing intermediates move towards the Golgi complex in two pre-Golgi structures. A single long tubule was often observed in the motion of the pre-Golgi motor on the tip of the membranes towards the

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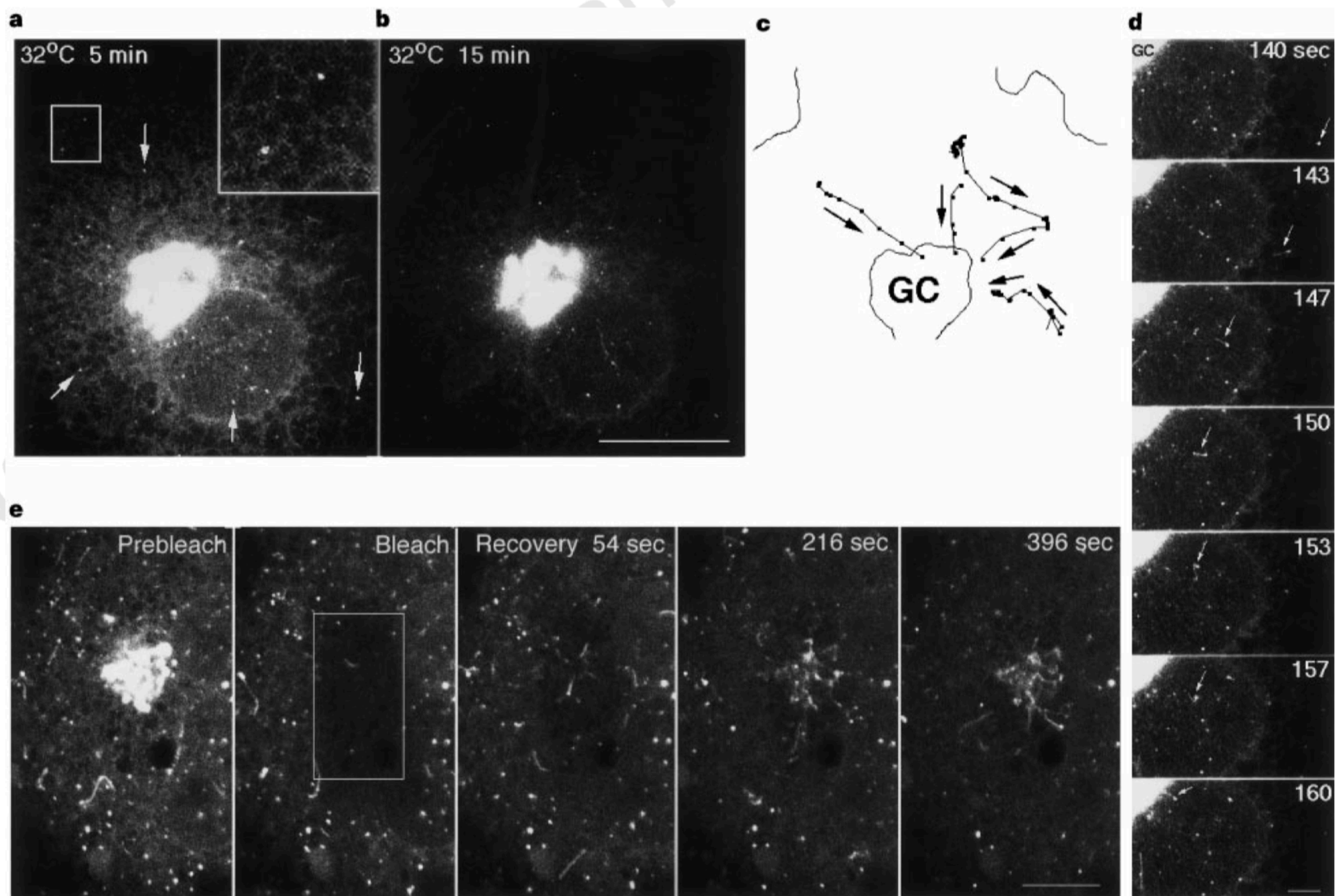
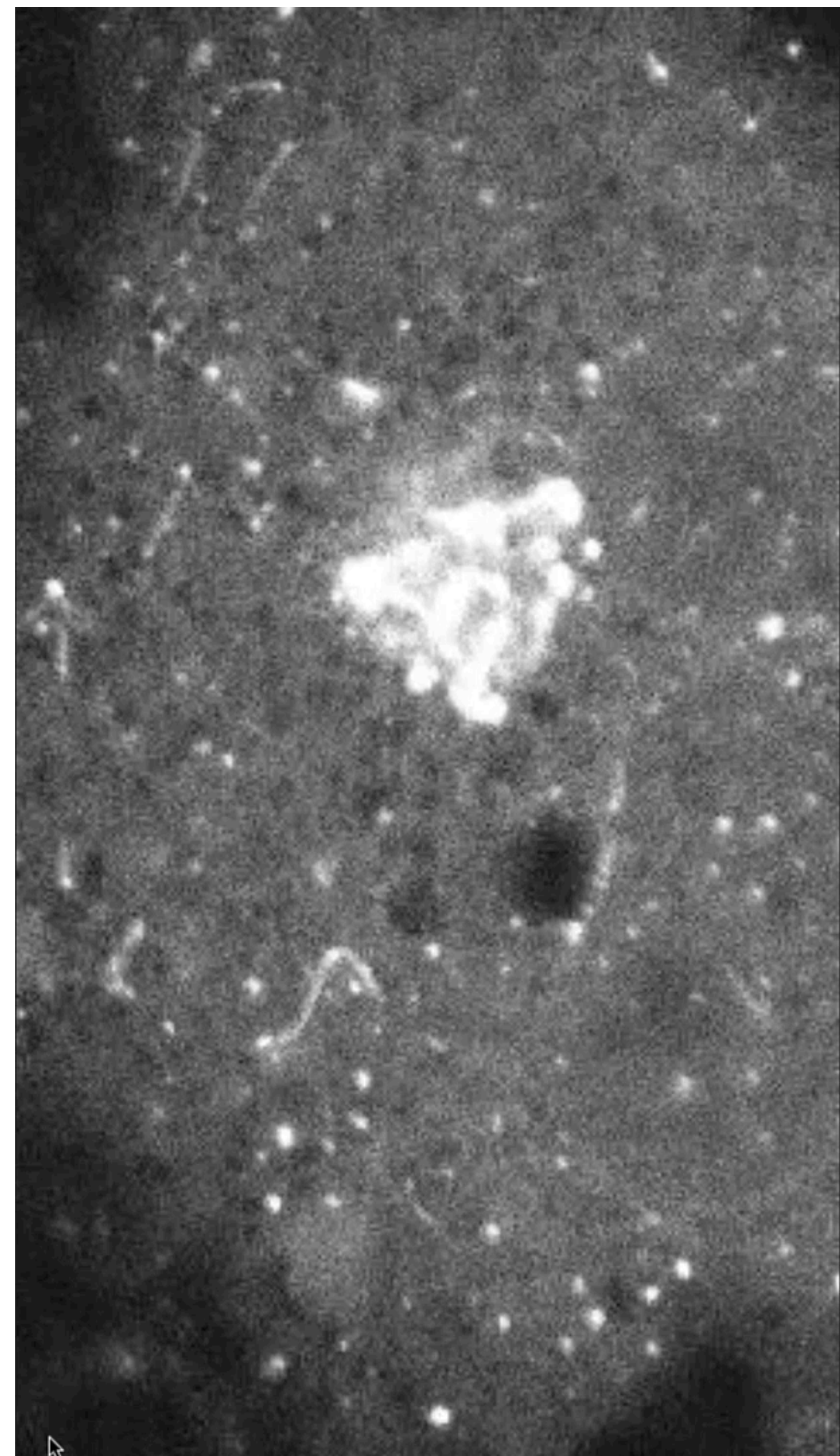


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d, Image series showing shape change of a pre-Golgi intermediate as it translocated to the Golgi complex over the time interval shown. Scale bar, 4 μ m. **e**, VSVG-GFP-expressing COS cells were incubated for 12 h at 40°C, shifted to 15°C for 3 h and then warmed to 32°C. Fluorescence associated with the Golgi

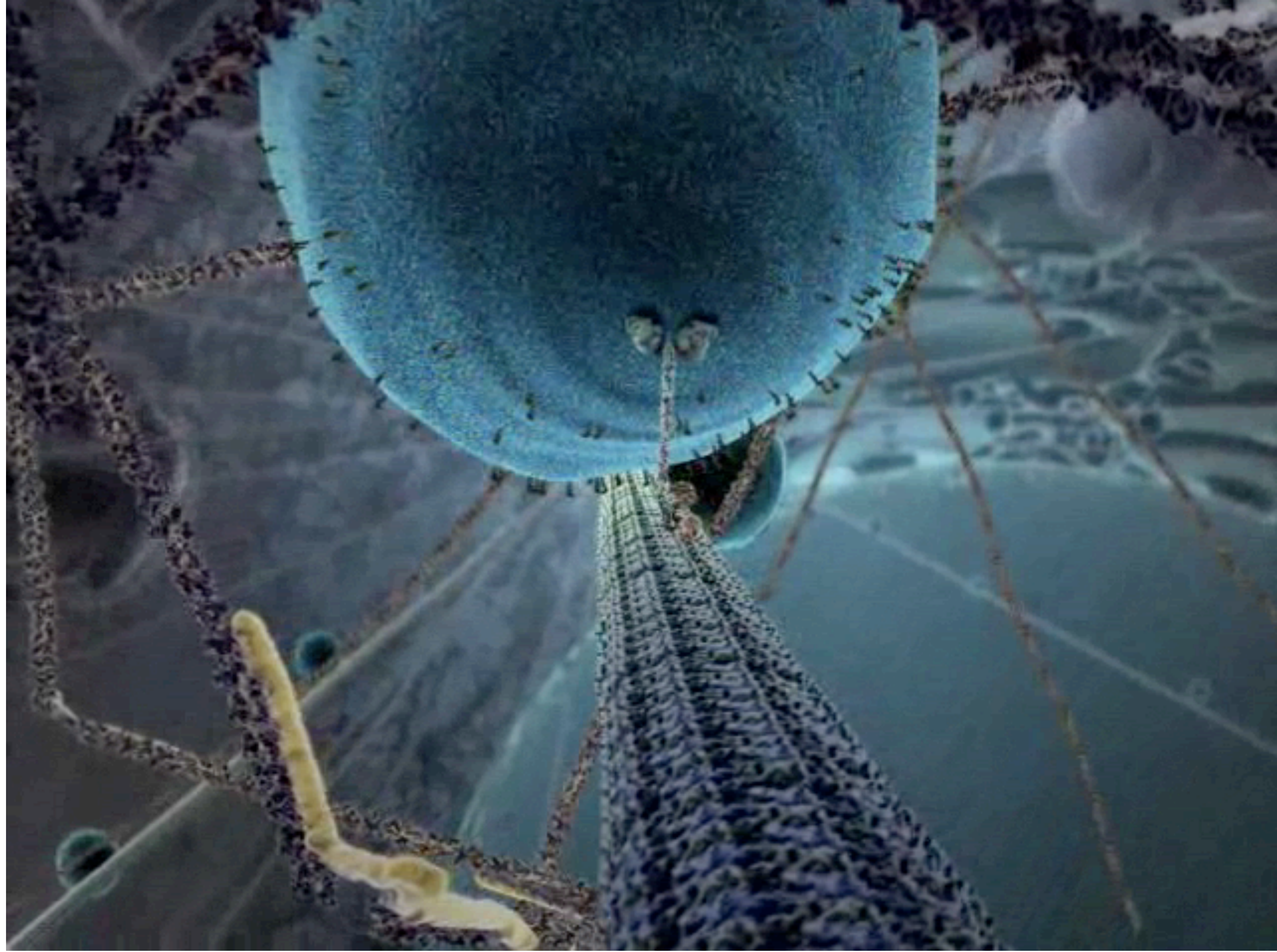
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WHAT YOU WANT TO DO NEXT?

DID CLICKER QUESTIONS

WHAT YOU WANT TO DO NEXT?



trifecta

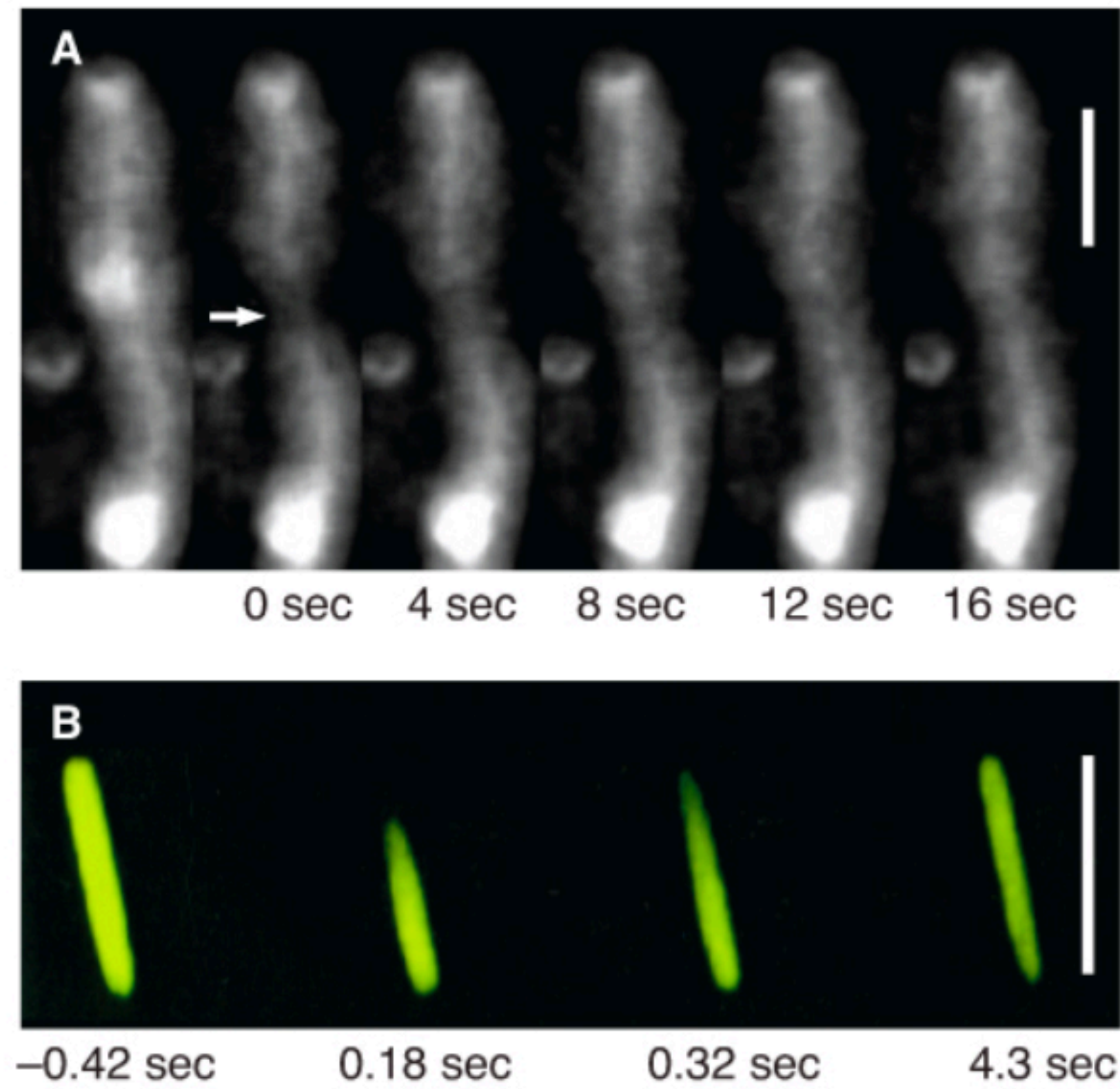


Figure 8.16 A, FRAP experiment with GFP fused with a normal plasma membrane protein; the arrow marks the area that was photobleached. The times are the seconds after bleaching. Bar = 5 μm . **B**, Imaged *E. coli* with GFP in the cytoplasm of the cell at the indicated times in seconds starting 0.42 seconds before photobleaching. Bar = 4 μm . Panel A from Mullineaux *et al.*, 2006, their figure 1. Mullineaux, Conrad W., Anja Nenninger, Nicola Ray, and Colin Robinson. 2006. Diffusion of Green Fluorescent Protein in Three Cell Environments in *Escherichia coli*. *Journal of Bacteriology*. Vol. 188(10): 3442–3448. B. From Elowitz *et al.*, 1999, modified from their figure 1. Elowitz, Michael B., Michael G. Surette, Pierre-Etienne Wolf, Jeffrey B. Stock, and Stanislas Leibler. 1999. Protein Mobility in the Cytoplasm of *Escherichia coli*. *Journal of Bacteriology*. Vol. 181(1):197–203.

Protein Mobility in the Cytoplasm of *Escherichia coli*

MICHAEL B. ELOWITZ,^{1,2*} MICHAEL G. SURETTE,^{2†} PIERRE-ETIENNE WOLF,^{1,2‡}
JEFFRY

Departments of Physics¹ and M

Rece

The rate of protein diffusion in bacteria is a key factor in determining the rates of many biochemical reactions. The apparent diffusion coefficient of green fluorescent protein (GFP) was measured in two wild-type *E. coli* strains. Measurements were made in two wild-type strains in the red-emitting fluorescent state of GFP (M. B. Elowitz, *J. Biol. Chem.* 272:809–812, 1997). The apparent diffusion coefficient is $2.5 \mu\text{m}^2/\text{s}$. A 72-kDa fusion protein containing a GFP domain moves more slowly, with D_a approximately 10% that of GFP. At least two factors: first, D_a is reduced for the larger protein; second, to GFP of a small tag consisting of several amino acids. Cytoplasmic viscosity cannot explain the understanding of intracellular bio-

Response times and reaction rates in *Escherichia coli* depend on the movement of proteins from one location to another in the cell. These proteins may have signaling functions, or they may act as enzymes for cellular reactions. How do such molecules

GFP DIFFUSION IN *E. COLI* 3443

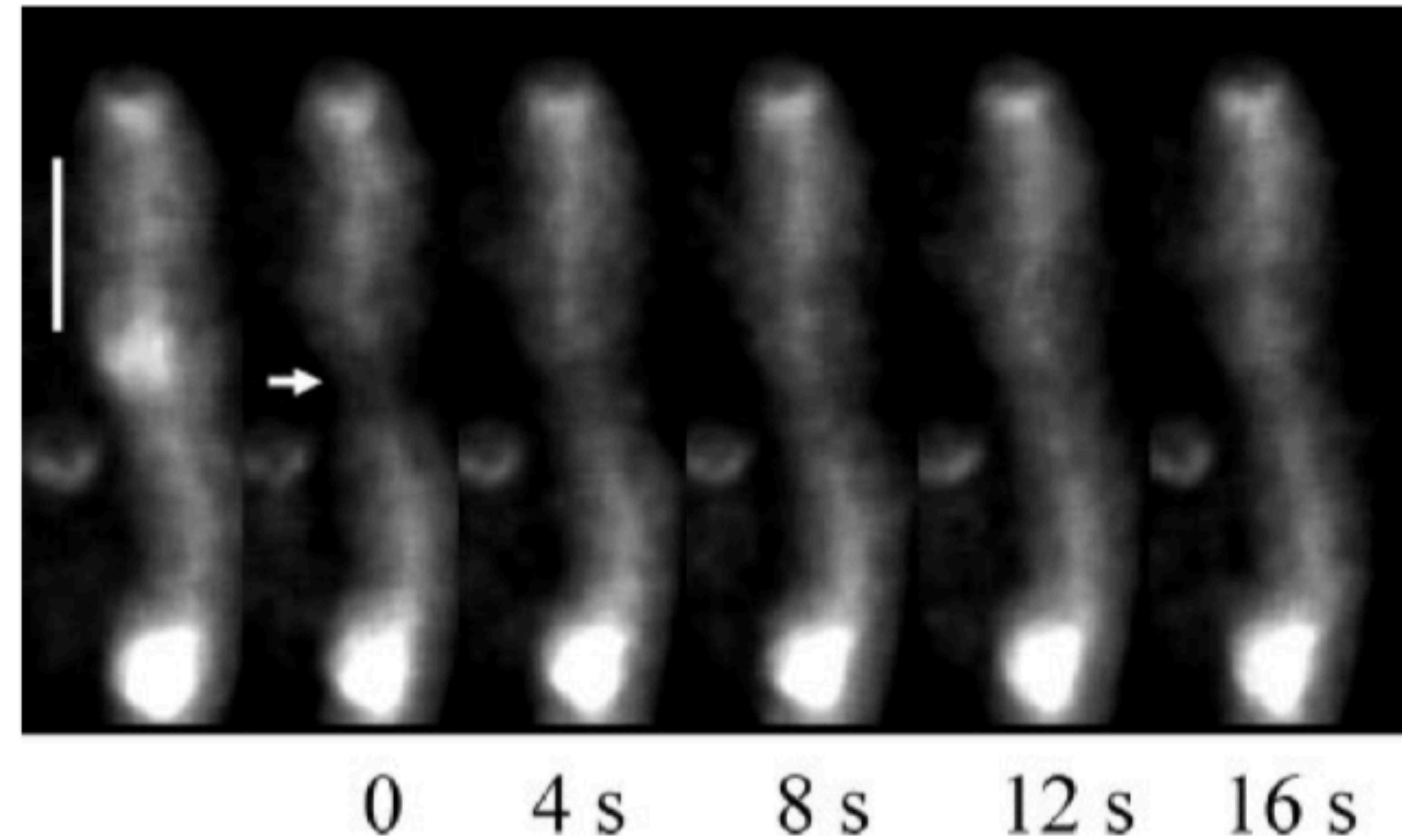


FIG. 1. FRAP image sequence for an elongated *E. coli* cell expressing TatA-GFP localized in the plasma membrane. The far left image is prior to the bleach. The next image is after bleach, and subsequent time in seconds is indicated. The center of the bleach is indicated by the arrow, whose thickness corresponds to the approximate beam width. Scale bar, 5 microns.

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Diffusion of Green Fluorescent Protein in Three Cell Environments in *Escherichia Coli*

Conrad W. Mullineaux,^{1*} Anja Nenninger,² Nicola Ray,² and Colin Robinson²

School of Biological and Chemical Sciences, University of London, United Kingdom,¹ and Department of Microbiology, University of Cambridge, United Kingdom,²

Surprisingly little is known about the rates at which proteins and other macromolecules diffuse in a physical system. There has been much work using fluorescence recovery after photobleaching (FRAP) which is available about diffusion in cells of *Escherichia coli* and other prokaryotes. We have measured the diffusion of GFP in cells of *Escherichia coli* expressing GFP in the cytoplasm, exported into the periplasm, and attached to the integral plasma membrane. We found that a diffusion coefficient comparable to that in water was observed. This showed a very low rate of protein diffusion. The diffusion coefficient was only slightly smaller in the cytoplasm. The diffusion coefficient in the periplasm and on the membrane was only slightly smaller than in the cytoplasm. The diffusion coefficient in the cytoplasm and periplasm are relative to the diffusion coefficient in water.

The diffusion of cell components is important in all living cells. Diffusion may be particularly important in prokaryotes, where systems of active transport appear to be much less developed than those in eukaryotes. We do not know enough about intracellular environments to be able to predict

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VOL. 181, 1999

PROTEIN MOBILITY IN *E. COLI* 199

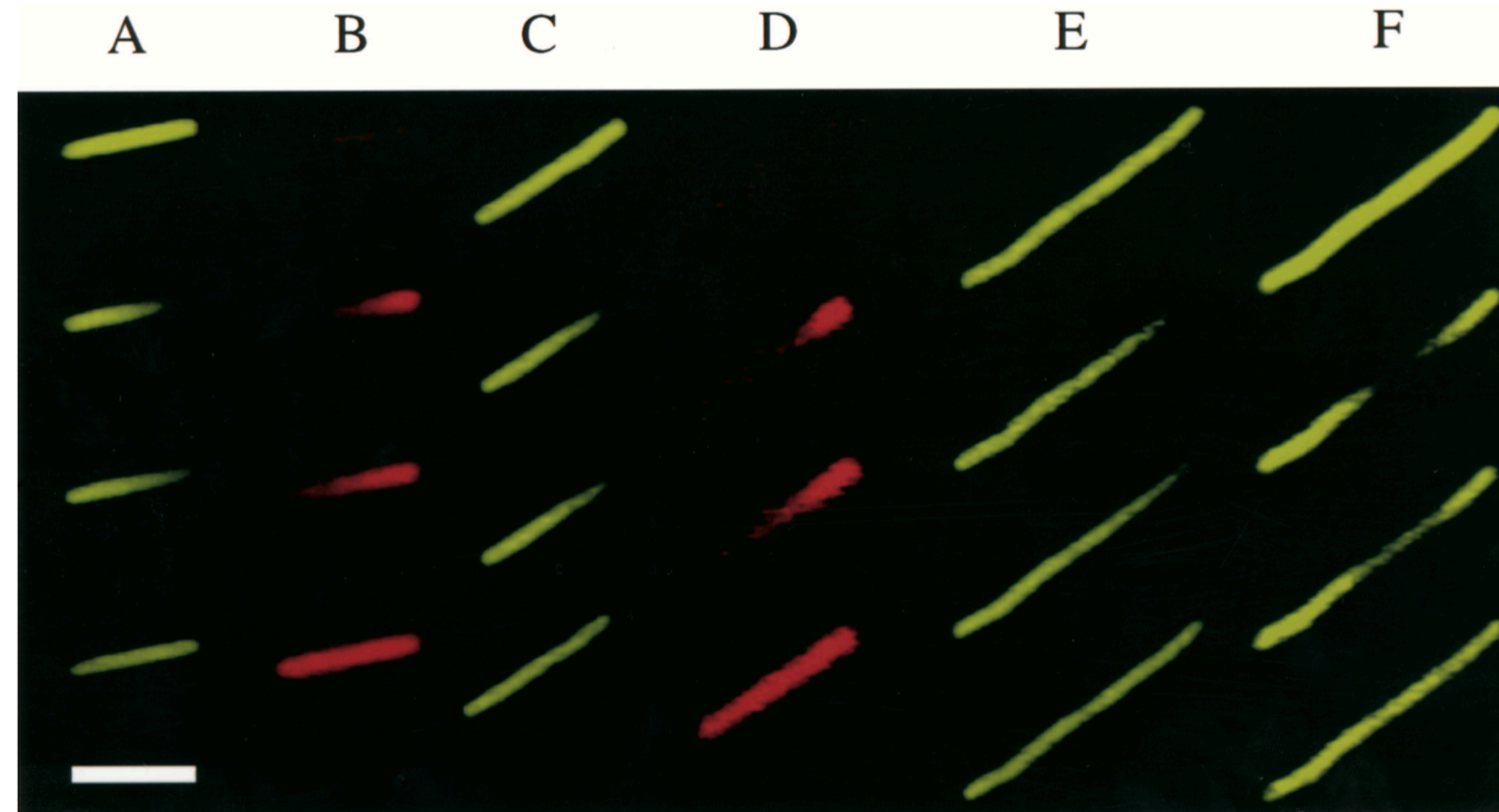


FIG. 1. Snapshots from photobleaching and photoactivation experiments. In each column the first row shows the cell before the laser pulse. The next three images show the cellular fluorescence distribution at subsequent times after the laser pulse. Columns A, C, E, and F show photobleaching (GFP filter set, false color green). Columns B and D show photoactivation (rhodamine filter set, false color red). Columns A to D show two different DH5 α cells expressing GFP (A and B show cell 1; C and D show cell 2). Columns E and F show a cephalaxin-treated DH5 α cell, expressing GFP, being bleached first at the pole (E) and then at the center (F). Time points are as follows ($t = 0$ is set arbitrarily as the end of the laser pulse). (A) $-0.42, 0.05, 0.18, 0.32,$ and 4.3 s. (B) $-0.08, 0.08, 0.35, 0.62,$ and 4.7 s. (C) $-0.5, 0.03, 0.10, 0.23,$ and 0.83 s. (D) $-0.1, 0.03, 0.23, 0.63,$ and 1.7 s. (E) $-0.57, 0.03, 0.43, 0.77,$ and 2.8 s. (F) $-0.57, 0.03, 0.20, 0.37,$ and 1.8 s. Bar = $4 \mu\text{m}$.

of GFP, $1.4 \text{ NS},$
 $1.0 \text{ } \mu\text{m}^2$

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Table 8.3

Table 8.3 Mobility of proteins in and on *E. coli* \pm 1 standard deviation.

protein	location	diffusion rate	fold slower
GFP	water	87.0 \pm 2.0	n.a.
GFP	cytoplasm <i>E. coli</i>	8.0 \pm 2.3	~10 X
GFP over produced	cytoplasm <i>E. coli</i>	3.6 \pm 0.7	~24 X
GFP + protein with sugar	cytoplasm <i>E. coli</i>	2.5 \pm 0.6	~35 X
GFP	periplasm <i>E. coli</i>	2.6 \pm 1.2	~33 X
GFP + membrane protein	membrane <i>E. coli</i>	0.13 \pm 0.03	~669 X

