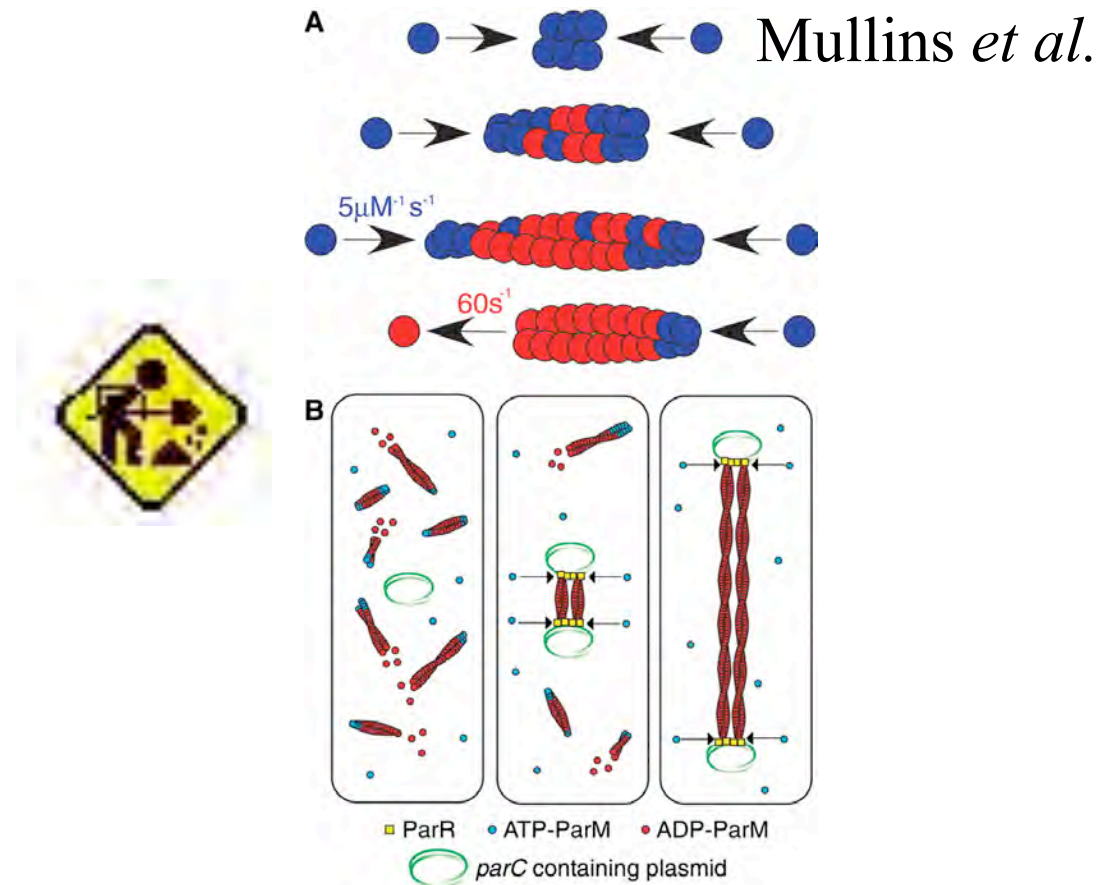


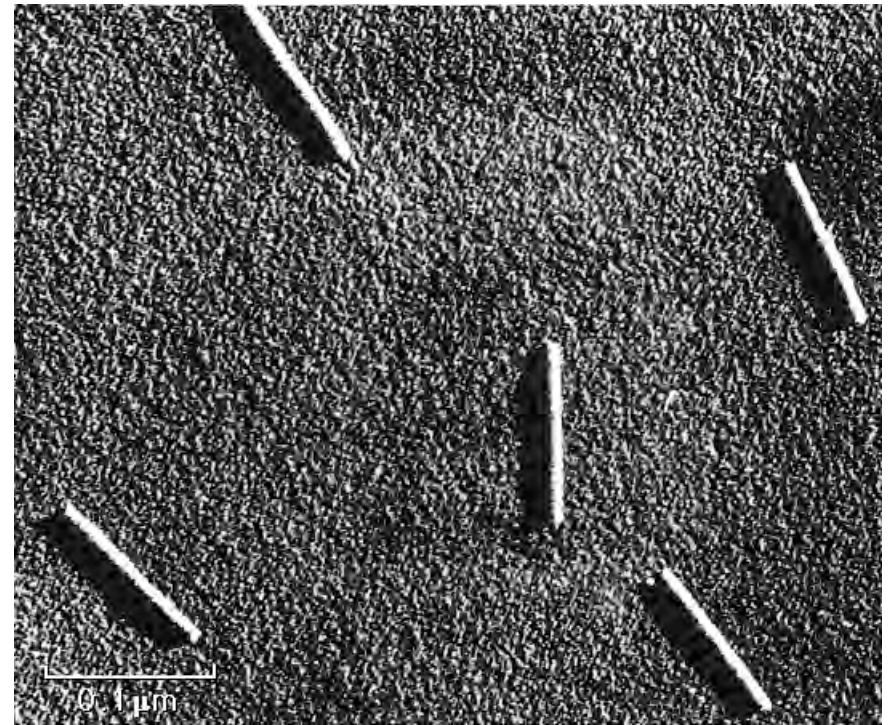
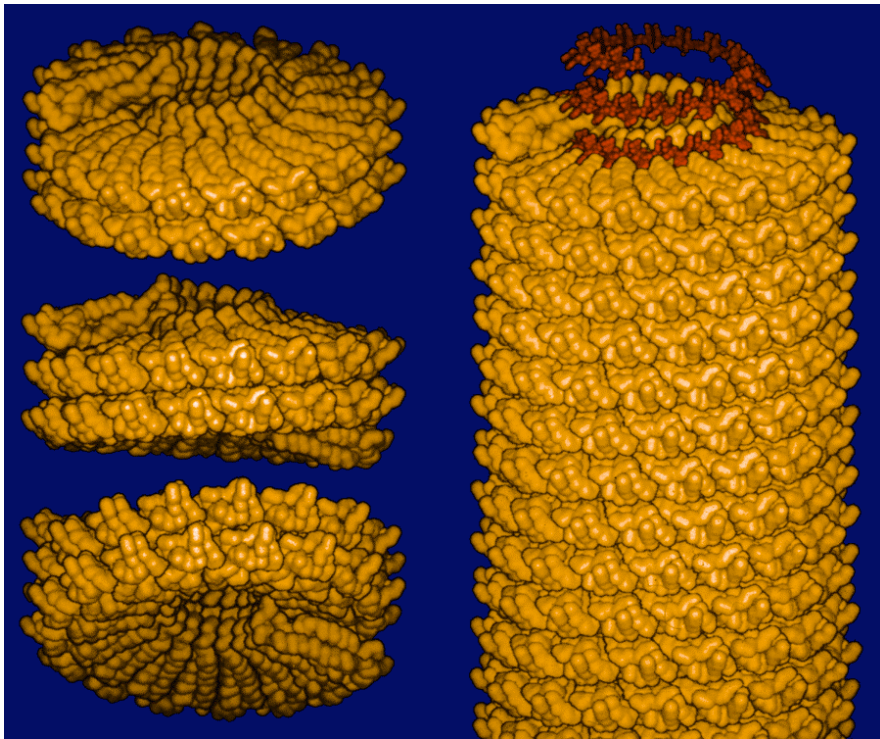
# APh161 - Lecture 11: The Cytoskeleton is Always Under Construction



**Rob Phillips**

**California Institute of Technology**

# *Polymerization Processes 1: Tobacco Mosaic Virus*



Caspar



# A reminder on the cytoskeleton

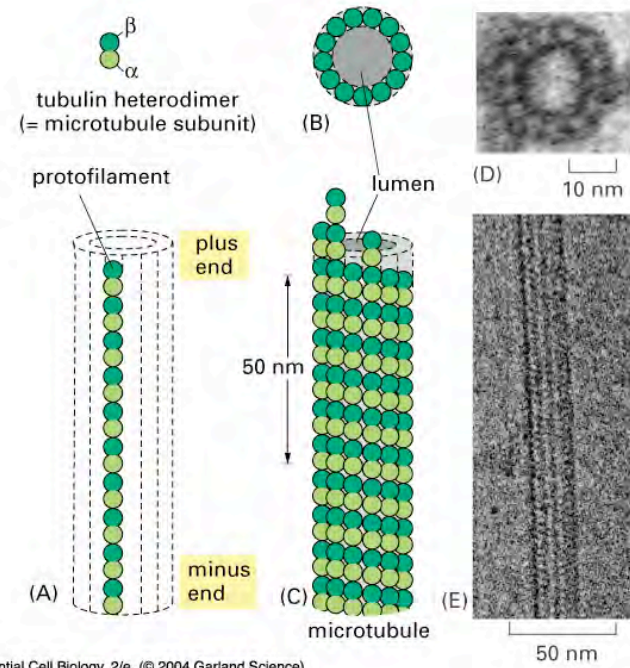
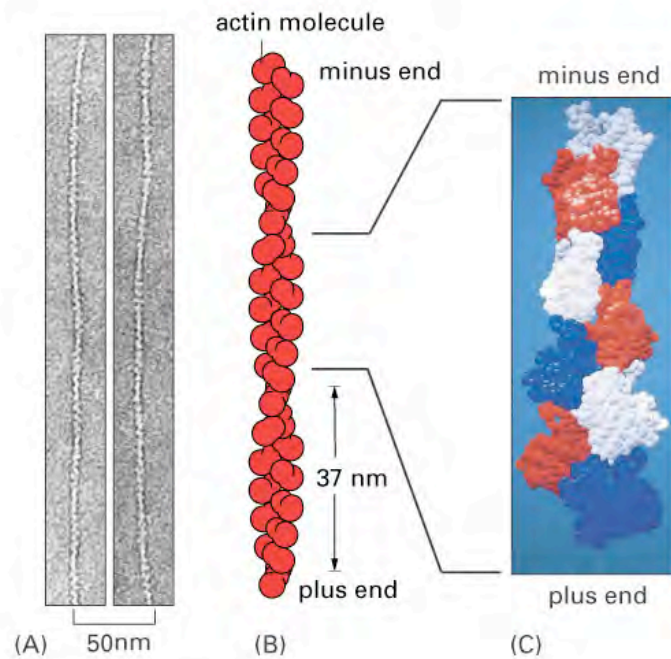
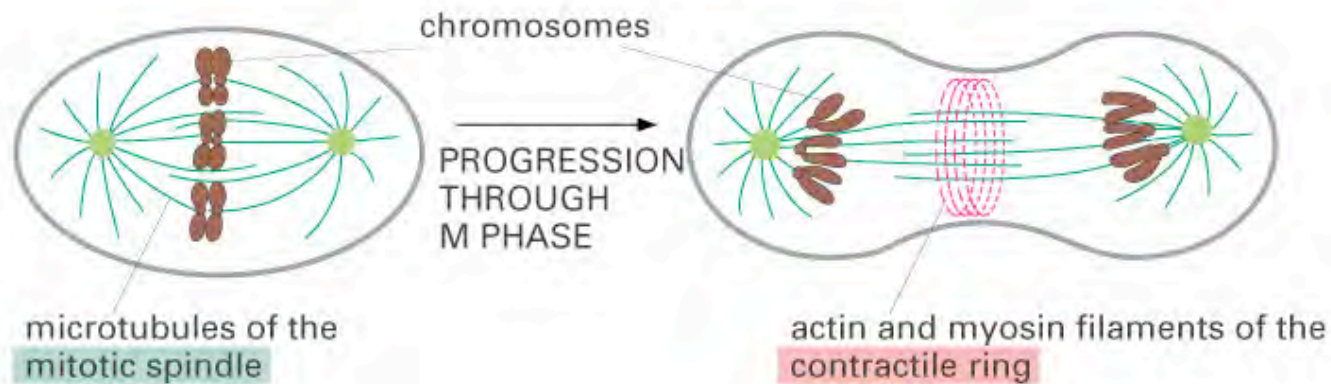


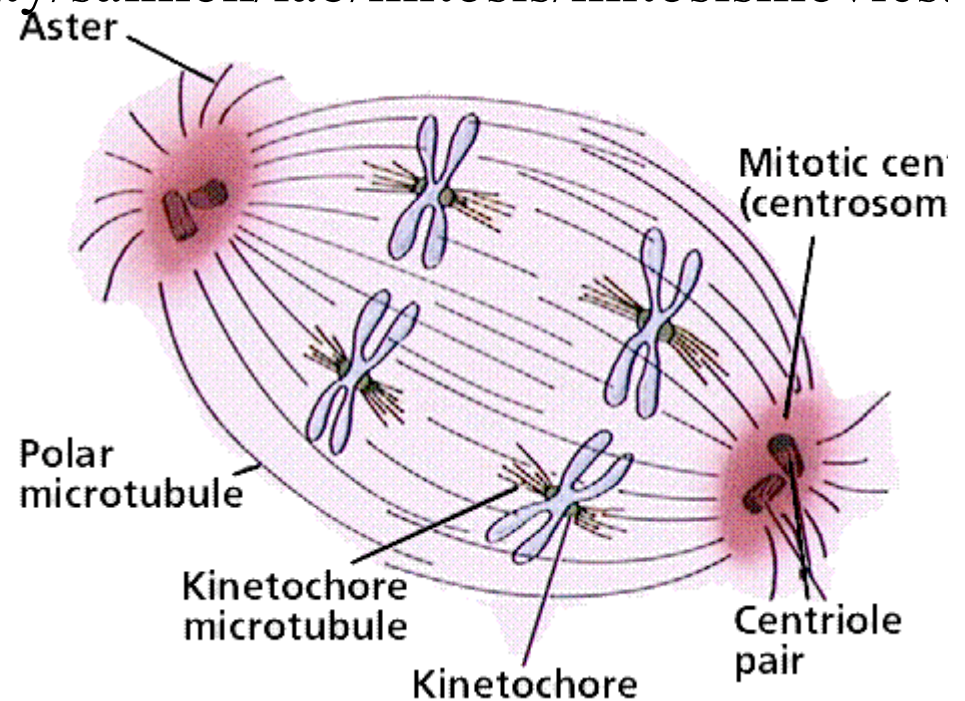
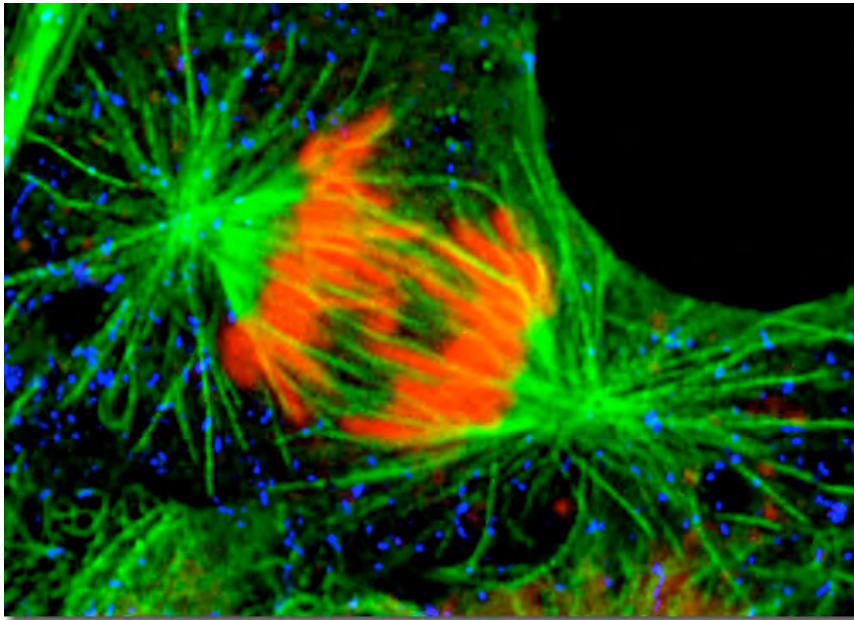
Figure 17-10 Essential Cell Biology, 2/e. (© 2004 Garland Science)

# Cytoskeletal Action During Cell Division

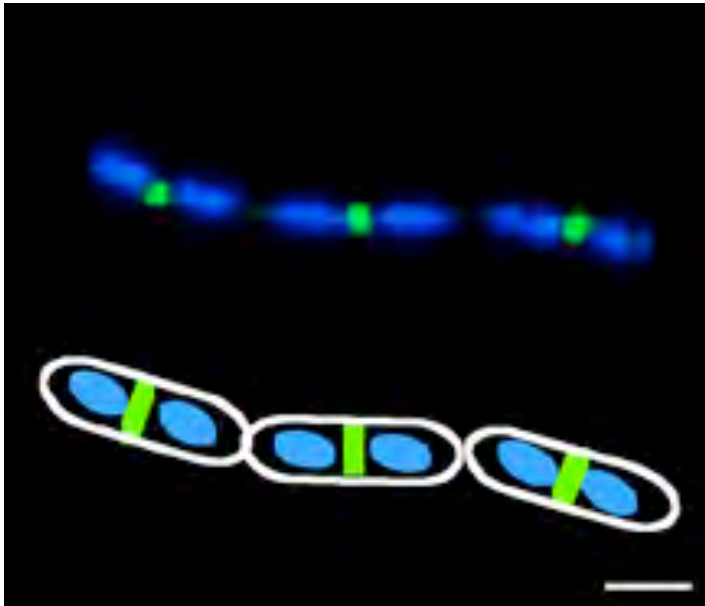


# Cytoskeleton Mediated Mitosis

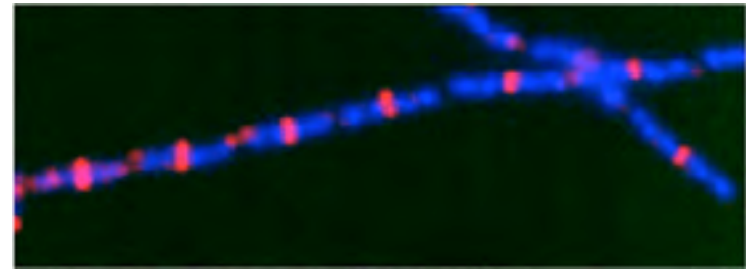
Show the movies in the Lecture 11 directory – see <http://www.bio.unc.edu/faculty/salmon/lab/mitosis/mitosismovies>.



# Cell Division in Bacteria



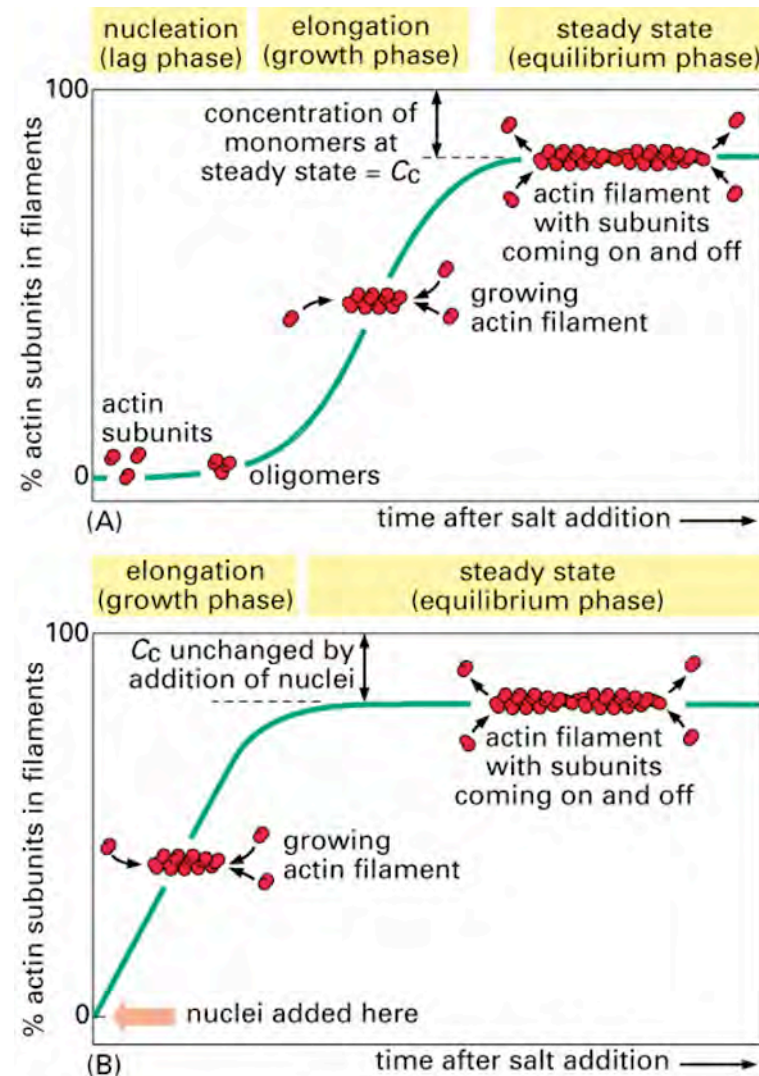
Petra Levin



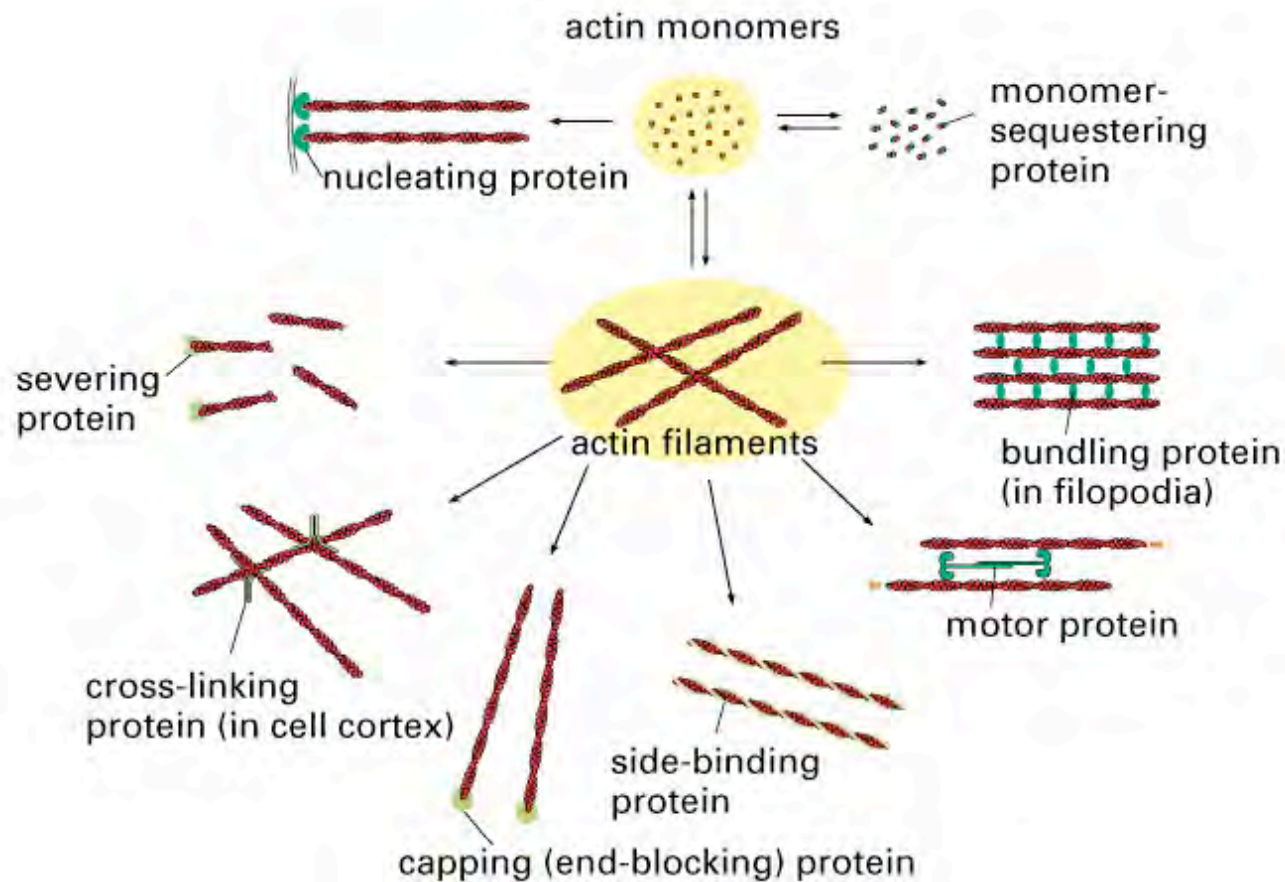
FtsZ rings during division



# Actin Polymerization Rate in vitro

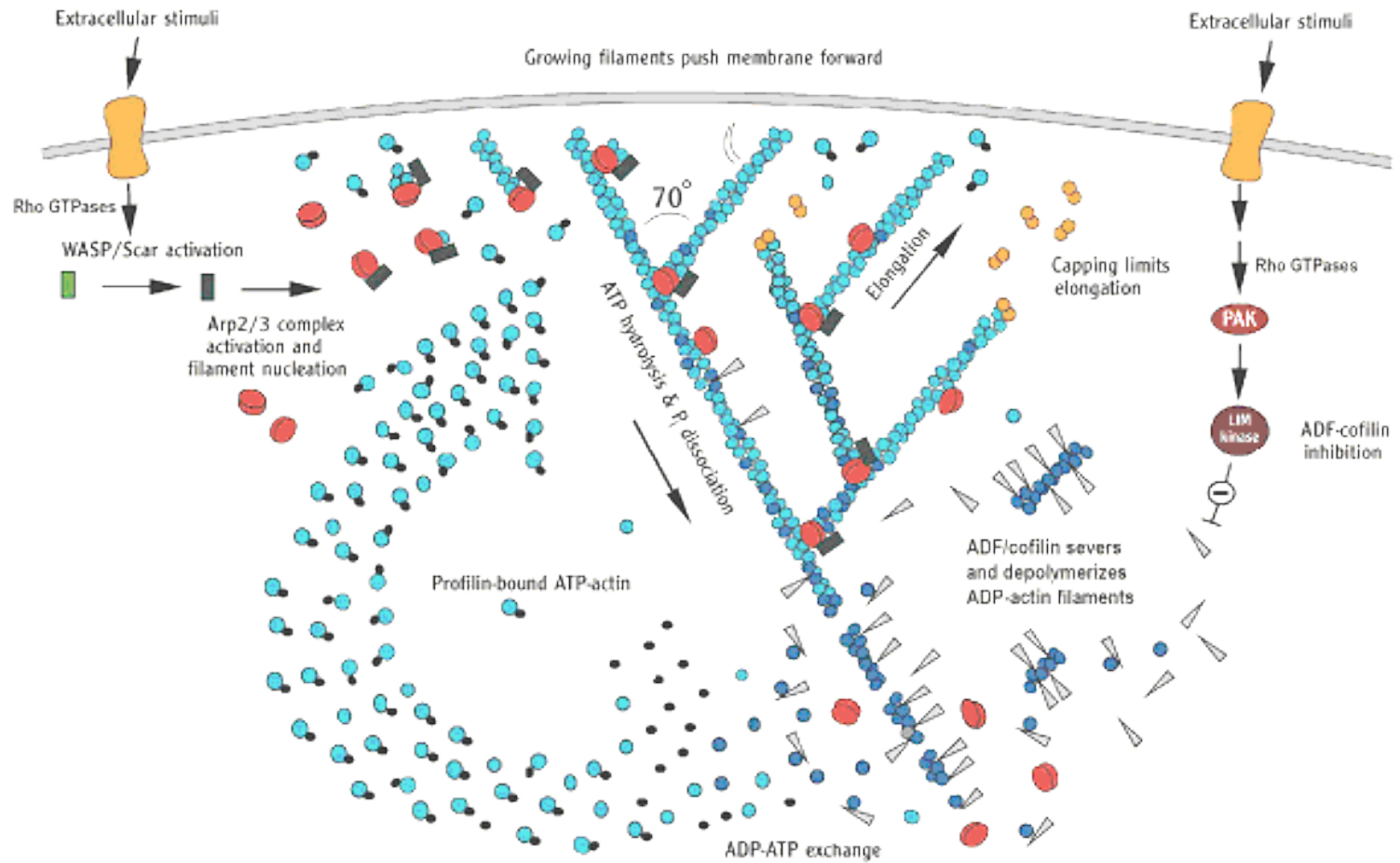


# Actin and its Partners

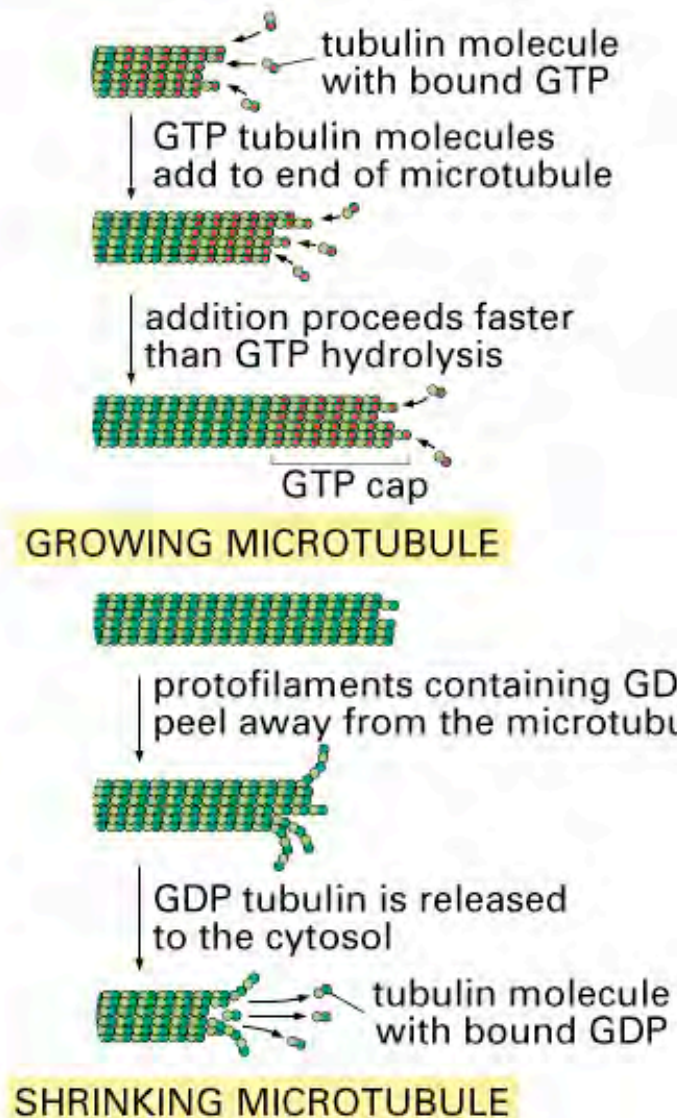




# Life at the Leading Edge



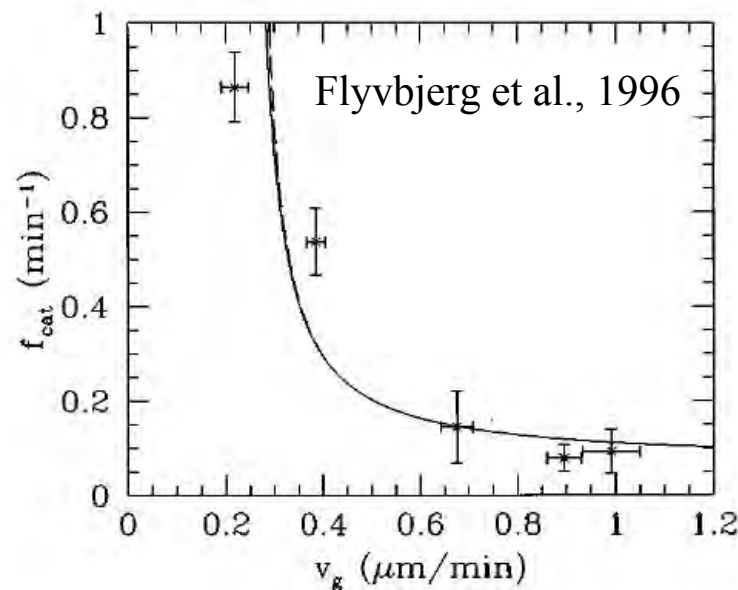
# Microtubule Dynamics: Part of the Polymerization Puzzle



# Microtubule Dynamics In Vitro

Show Borisov movies,  
<http://www.borisylab.nwu.edu/pages/supplemental/treadmill.html>

## Catastrophe rate vs. growth rate



PHYSICAL REVIEW E

VOLUME 50, NUMBER 2

AUGUST 19

### Phase diagram of microtubules

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(Received 4 April 1994)

We map the phase diagram of microtubules as a function of temperature and tubulin concentration. We observe spontaneous and site-nucleated microtubule assembly. At temperatures and concentrations below the onset of spontaneous nucleation, we measure the steady-state proportion of occupied nucleation sites and the distribution of lengths of site-nucleated microtubules. Our observations reveal a transition in the length dynamics of microtubules from bounded to unbounded growth. This transition is also evident in the length dynamics of individual site-nucleated microtubules. The transition to spontaneous microtubule assembly completes the phase diagram. We measure the temperature and concentration dependence of the latent time for spontaneous nucleation of microtubules in the bulk.

PACS number(s): 87.15.-v, 64.60.-i

### I. INTRODUCTION

The protein *tubulin* is found in every cell of every living organism (with the exception of bacteria) [1]. In the presence of guanosine-tri-phosphate (GTP) and magnesium ions ( $\text{Mg}^{2+}$ ), tubulin-GTP complexes form [2] and aggregate [3] into long, hollow cylinders called *microtubules*. Microtubules are about 25 nm in outer diameter, incorporate 1625 tubulin dimers per micrometer of their length [4] and easily grow long enough to span a cell (10–100  $\mu\text{m}$ ). In the cell, microtubules form a network that supports the overall structure and guides internal transport [5]. Fundamental cellular processes, like locomotion, morphogenesis, and reproduction, rely on the ability of microtubules to change their organization. This ability derives from a unique feature of microtubule assembly called *dynamic instability* [6].

The term dynamic instability describes the fact that individual microtubules are dynamic structures that fluctuate erratically between assembling and disassembling. These fluctuations are not microscopic; they often involve most or all of the microtubule. A typical plot of the length of a single microtubule over time is shown in Fig. 1. This so-called dynamic instability depends on the hydrolysis reaction which turns the GTP of the tubulin-GTP complex into guanosine-di-phosphate (GDP) and releases energy ( $\sim 8kT/\text{reaction}$ ) [7].

Dynamic instability has been the subject of research for the past decade. The key element of all models is a competition between growth and the process of hydrolysis, often considered in terms of a growth front and a possible hydrolysis front. Crudely, when the hydrolysis front overtakes the growth front, the microtubule structure becomes unstable. The question of exactly how the hydrolysis and growth reactions are coupled continues to

draw attention [8]. Experimental control parameters include temperature and tubulin concentration, as well as the pH of the solution, the presence of certain other proteins [microtubule-associated proteins (MAP's)] [9], and the concentrations of GTP and  $\text{Mg}^{2+}$  ions [10]. Of the most biologically relevant parameter is perhaps the concentration of tubulin. However, to understand dynamic instability the concentration of tubulin has been the preferred variable, since it affects the growth velocity but not the hydrolysis of GTP.

In the work described here, we vary both temperature and tubulin concentration and map the phase diagram of microtubule growth. We study both spontaneously nucleated assembly and growth from nucleation sites [11]. The resulting phase diagram is shown in Fig. 2.

At low temperatures, microtubules only grow from nucleating sites. For a range of temperatures, a steady state is reached in which only a fraction of the sites have n

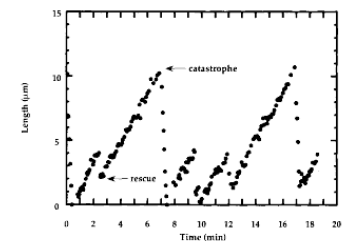


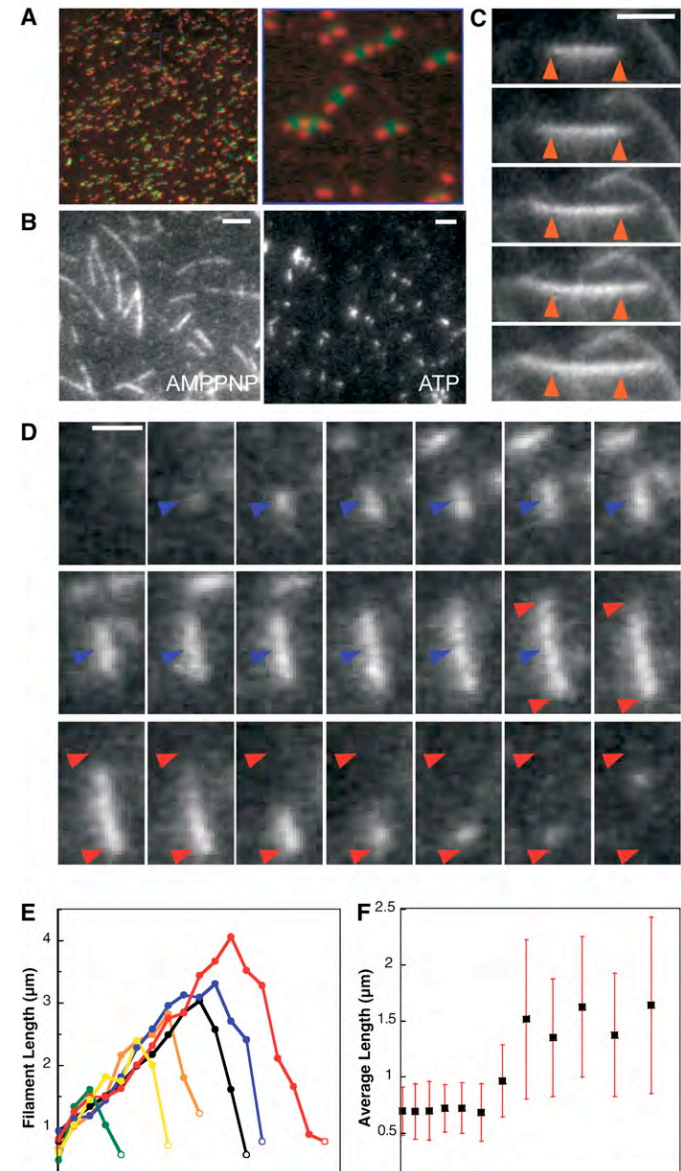
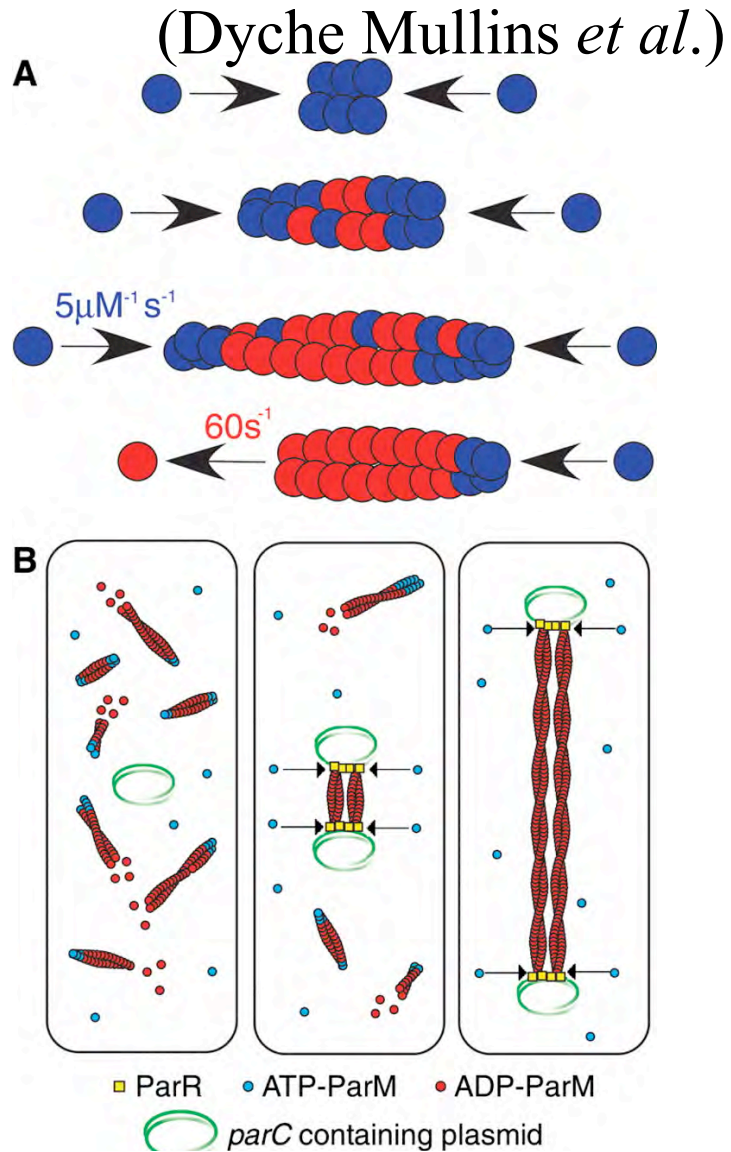
FIG. 1. A typical time series of the length of a single microtubule. The velocities of growth range between 1.6 and 2  $\mu\text{m}/\text{min}$  (40–65 dimers/sec), while the velocities of shortening are 10 times faster. The transitions from growth to shortening are called *catastrophes*; transitions from shortening to growth are called *rescues*.

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<sup>†</sup>Also at NEC Research Institute, 4 Independence Way, Princeton, NJ 08540.

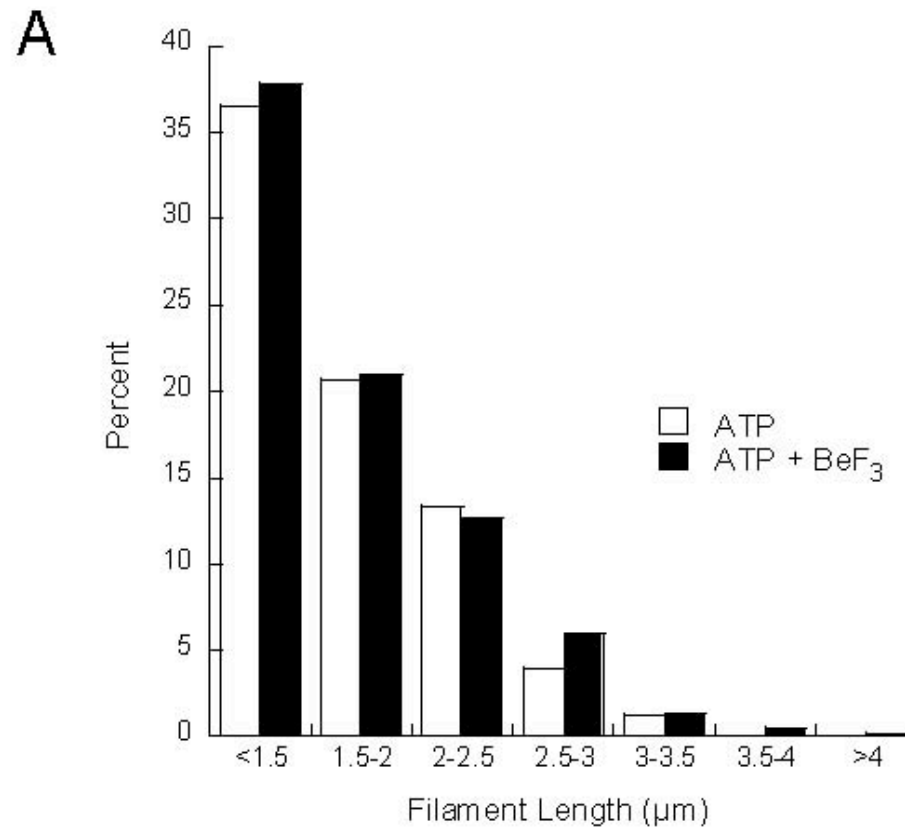


# The Story of ParM



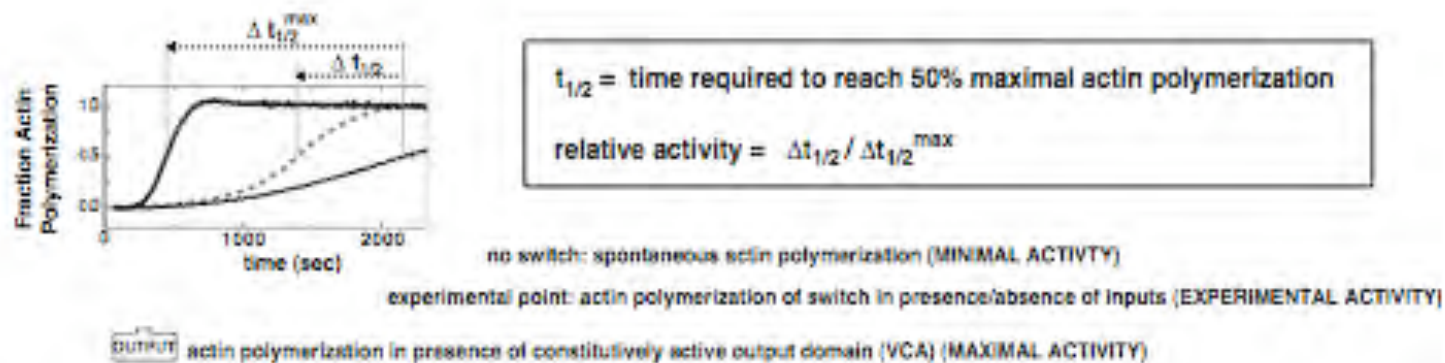
# *ParM Length Distribution*

Mullins et al., Science 2004



# Polymerization of Pyrene Labeled Actin

Lim et al., Science 2004



**Fig. S1. Metric for relative activity of N-WASP switches based on half-time of actin polymerization.** Activity of switch proteins was determined using a fluorescence-based actin polymerization assay (20, and "Methods"). Time required to reach 50% polymerization ( $t_{1/2}$ ) was used as a metric for activity. Minimal activity was defined as the  $t_{1/2}$  observed with spontaneous actin polymerization under these conditions in the presence of Arp2/3 but no nucleation promoting factors. Maximal activity was defined as the  $t_{1/2}$  in the presence of the isolated output domain. Relative activities of individual constructs were scored by measuring the change in  $t_{1/2}$  relative to the difference between maximum and minimum activities.