

Spontaneous Waves and Patches of F-Actin in Cells

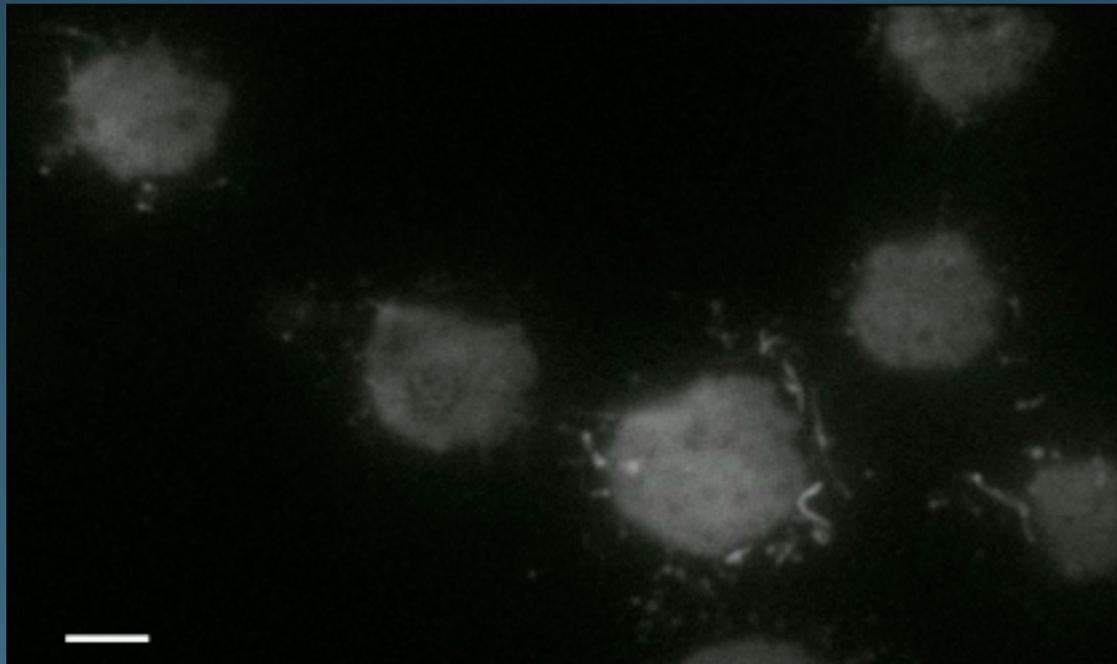
Anders Carlsson, Washington University
in St Louis

Polymerized actin often exhibits spontaneous dynamic behaviors, such as waves, patches, and uniform oscillations

These waves may help cells explore their environment, exert force, and sense tension

- Does known actin biochemistry lead to spontaneous wave formation?
- What types of feedback mechanisms lead to spontaneous actin dynamics?

Actin Waves Appear After Recovery from Latrunculin Treatment



5 μm

(Bretschneider et al 2004)

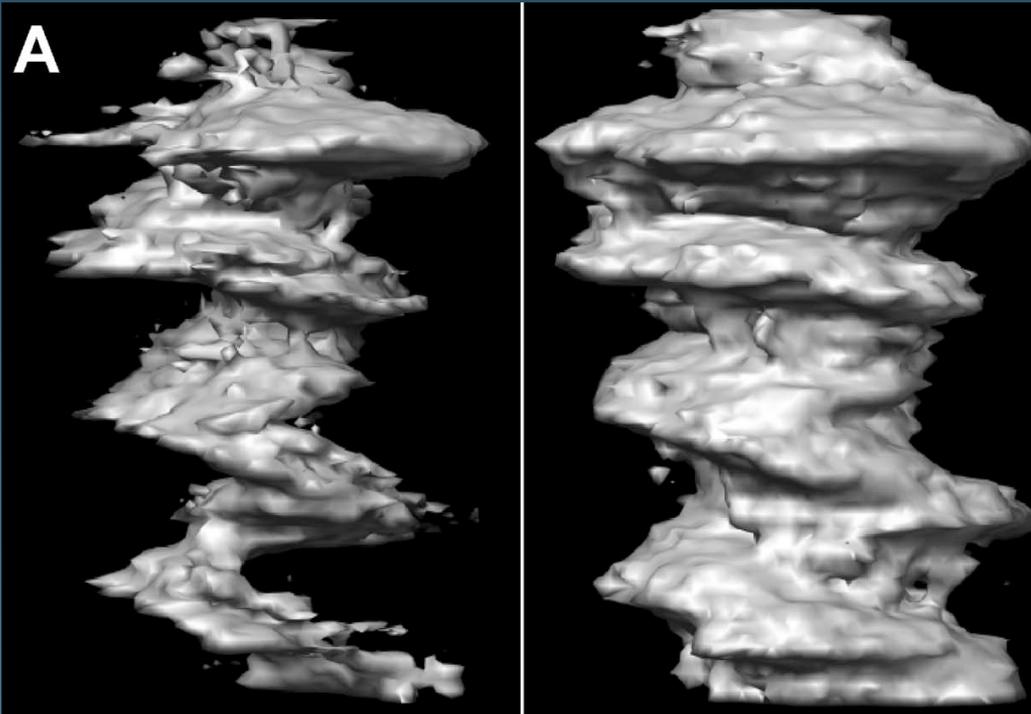
Visualization via total internal reflection
fluorescence microscopy

- Spontaneous waves and patches of actin assembly
- Over time, get patches first, then waves
- Wave speed is 0.1-0.2 $\mu\text{m/s}$

F-actin Waves Are Correlated with Edge Protrusion

F-Actin Intensity

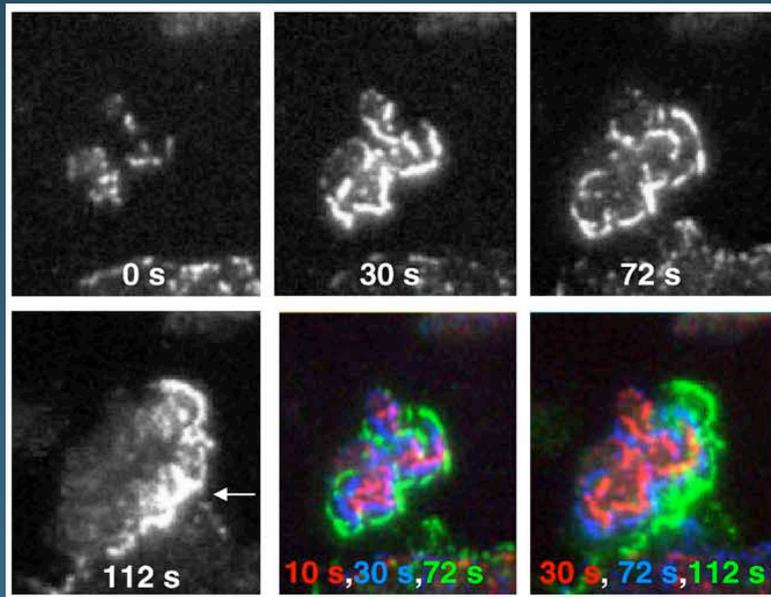
Cell Edge



- Actin waves appear to push membrane out

(Gerisch lab, 2009)

Hem-1 Waves in Chemoattractant-Stimulated Neutrophils



Red, blue, and green are successive times

Waves correlate with edge protrusion

(Weiner et al, 2007)

Actin waves also seen in T cells (Upadhyaya lab)

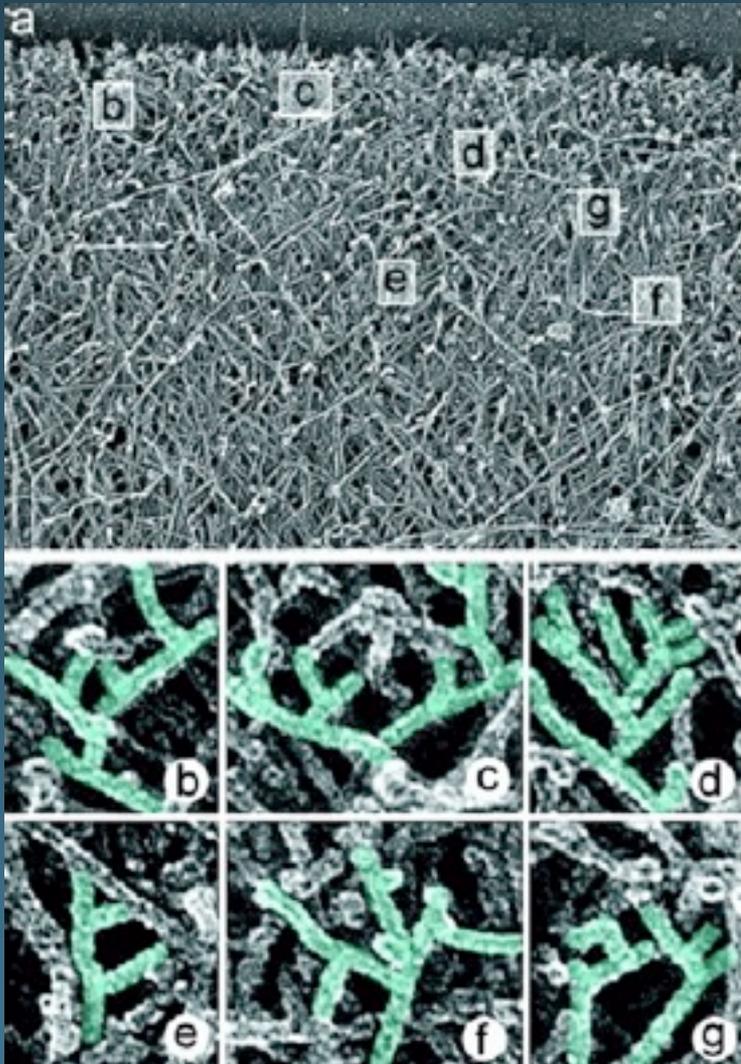
Coarse-Grained Models of Actin Waves

- (Weiner et al 2007, Doubrovinski and Kruse 2008, 2010). Actin filaments plus F-actin nucleators which act cooperatively and are active when at the membrane.
- (Whitelam et al 2009) Nonlinear F-actin field with built-in positive feedback and spontaneously arising orientation anisotropy

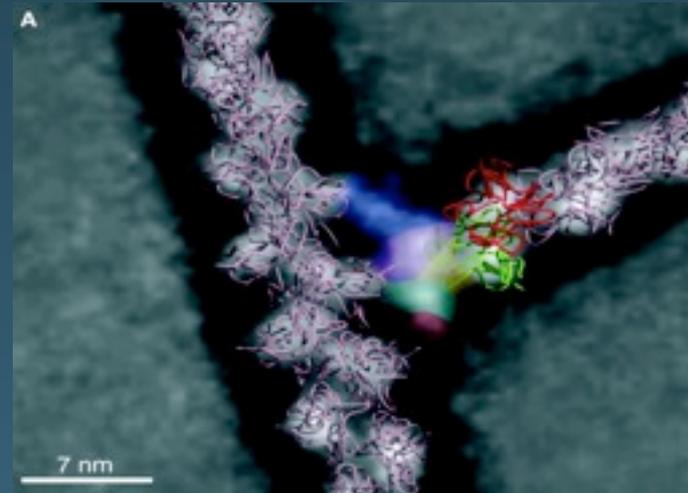
Related work: Falcke, Allard and Mogilner, Vavylonis, Levine and collaborators

Goal here: establish minimal model including known actin biochemistry

Basic Idea: Simulate Dendritic-Nucleation Model of Actin Filament Generation



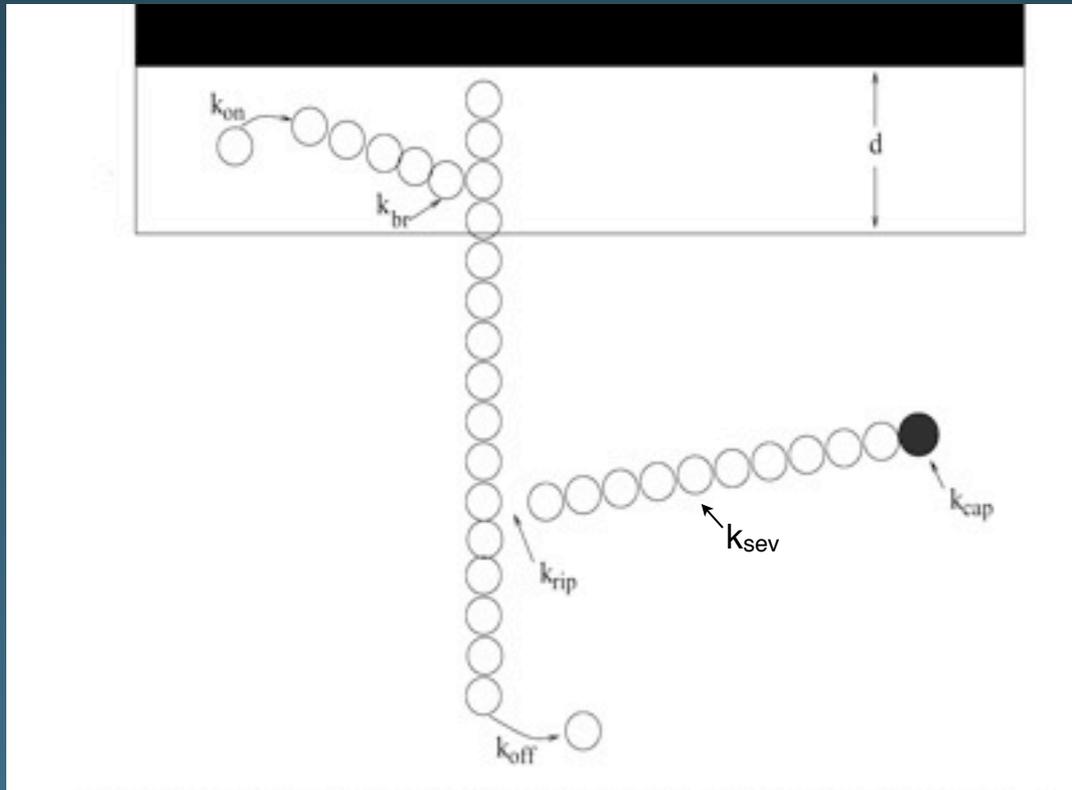
(Svitkina et al 1999)



(Volkman et al, 2004)

- “Arp2/3 complex” is activated by nucleation-promoting factors (NPF) in membrane: “dendritic nucleation”
- In a “Lego-Block” fashion, it starts new filaments on the sides of existing ones

Simulation Approach: Stochastic “Topological” Processes Combined with Brownian Dynamics of Filament Motion



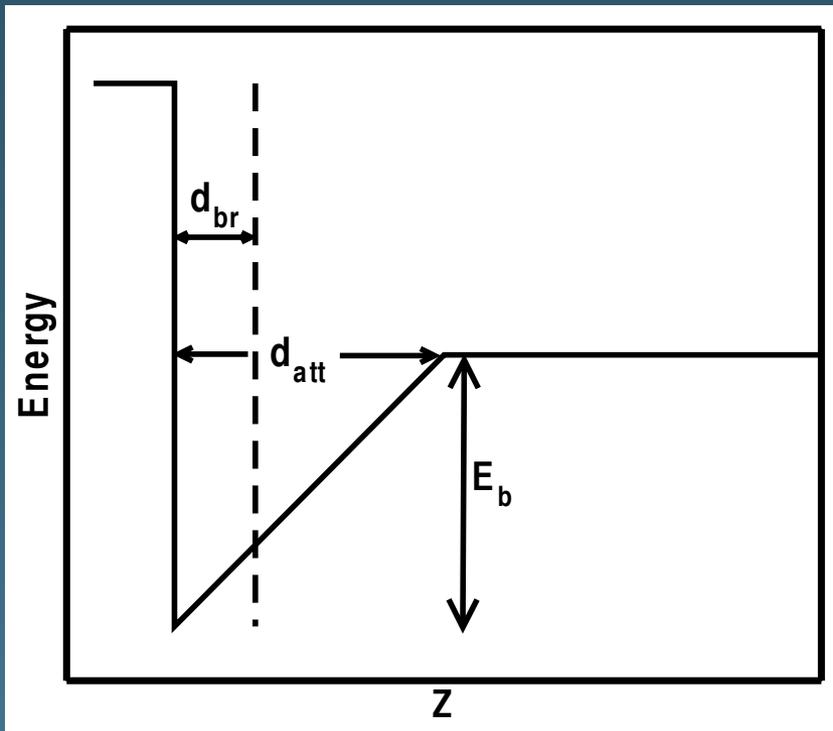
k_{on} : polymerization rate
 k_{off} : depolymerization rate
 k_{br} : branching rate
 k_{cap} : capping rate
 k_{det} : branch detachment rate
 k_{sev} : severing rate

New branches appear within distance d of membrane

Filaments Move via Brownian Dynamics in Membrane's Force Field

Membrane has

- 1) repulsive interactions with all filament tips
- 2) attractive interactions with uncapped barbed ends



$$\Delta R_i = (D/k_B T) F_i \Delta t + \eta \sqrt{2D\Delta t}$$

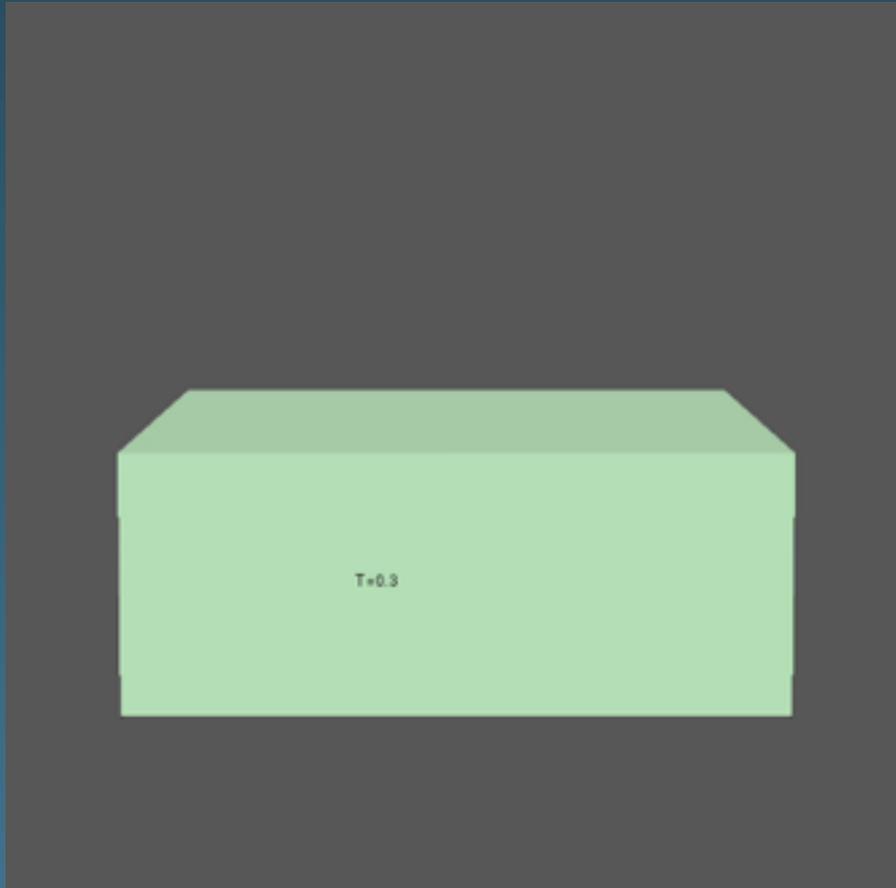
$$d_{br} = 10 \text{ nm}$$

$$d_{att} = 10 \text{ nm}$$

$$E_b = 4.5kT$$

Requires time step of about 10^{-7} s
Treat dendritic clusters as rigidly moving units and ignore cluster-cluster interactions

Dynamics of Actin at Membrane: Initiation of Polymerization



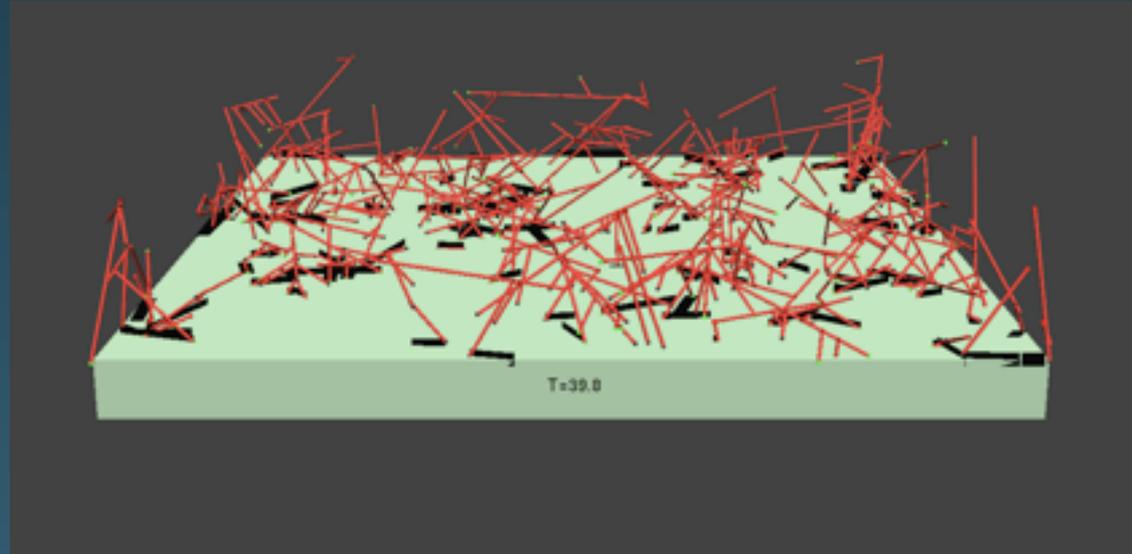
Bright green: uncapped barbed ends

Filaments nucleate at membrane and branch if they stay attached long enough

Most filaments leave the membrane before they branch, giving a large critical cluster size

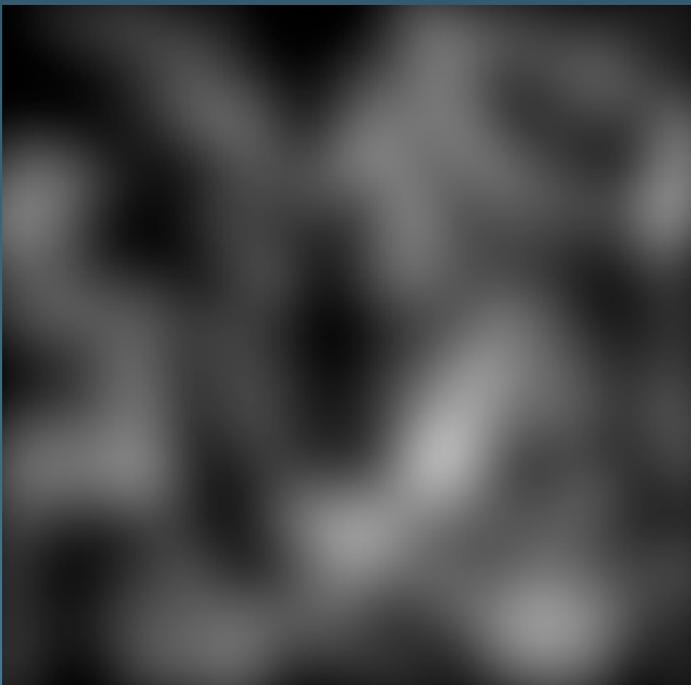
This is similar to the effect of a threshold for positive feedback in continuum models

“Bare” Dendritic
Nucleation Model
Gives no
Identifiable
Waves or
Patches

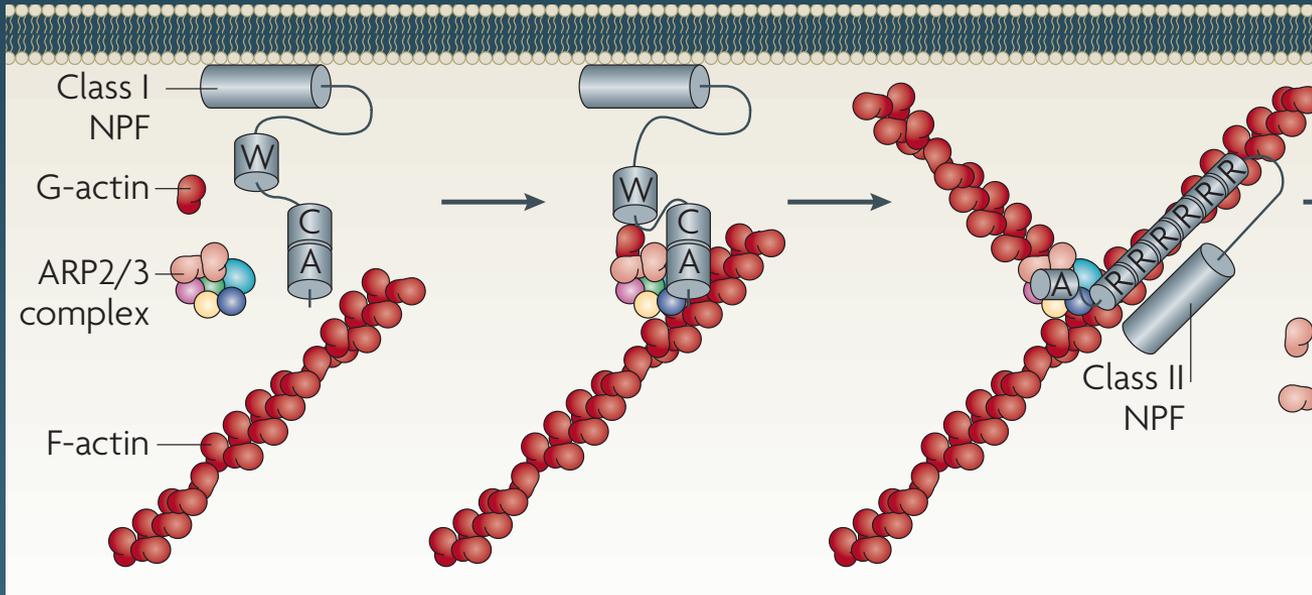


3 μ m

80 μ M actin



Origin of Waves May Lie Upstream of Actin

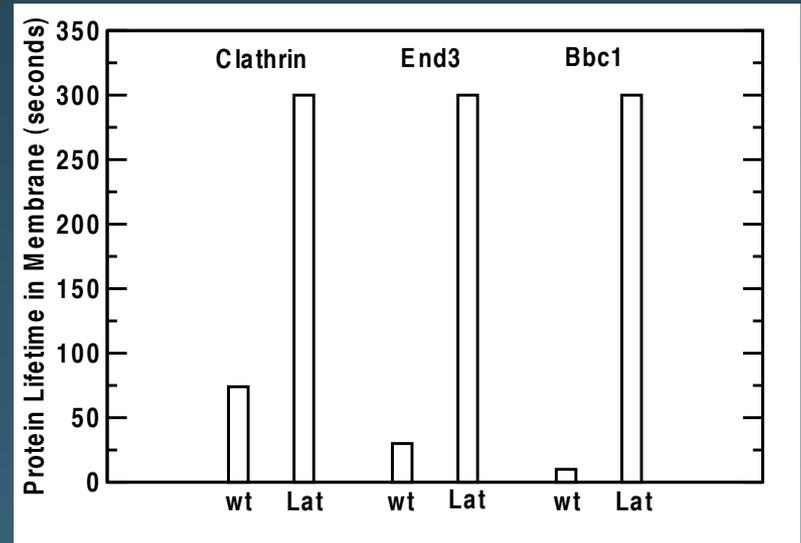


(Welch, 2010)

Nucleation-promoting factors (NPFs) act upstream of actin polymerization and require membrane localization to efficiently activate Arp2/3 complex

Assumption: F-Actin Detaches/ Inactivates Upstream NPFs

Weiner et al (2007)
showed that removal of
the NPF Hem-1 from the
membrane is greatly
slowed by latrunculin
treatment



Data for upstream actin
patch proteins in yeast
(Drubin lab, 2005)

Upstream protein lifetimes
are much longer with
latrunculin treatment¹²

Include F-Actin-Induced NPF Detachment in Model

$$\begin{aligned}\frac{\partial n_a}{\partial t} &= -k_{\text{det}}F(x, y)n_a + k_{\text{att}}\bar{n}_d \\ \frac{d\bar{n}_d}{dt} &= \frac{-1}{A} \int \frac{\partial n_a}{\partial t} dx dy\end{aligned}$$

n_a : 2D density of attached (active) NPFs

n_d : 2D density of detached NPFs (constant)

A : membrane area

F : density of F-actin near membrane

k_{det} , k_{att} : detachment and attachment constants

NPFs are detached by F-actin, then reattach spontaneously

Detachment inactivates the NPFs

Arp2/3 activation $\propto n_a^2$

k_{on}^B	$8.7 \mu M^{-1} s^{-1}$	[3]
k_{on}^P	$1.3 \mu M^{-1} s^{-1}$	[4]
k_{cap}	$8.0 \mu M^{-1} s^{-1}$	[5]
k_{uncap}	$0.0004 s^{-1}$	[5]
k_{nuc}	$0.001 - 0.009 \mu M^{-1} s^{-1}$	
k_{br}	$0.018 \mu M^{-3} s^{-1}$	
k_{dis}	$0.04 s^{-1}$	[6]
k_{sev}	$0.005 s^{-1}$	[7]
k_{att}	$0.025-0.075 s^{-1}$	
k_{det}	$0.005-0.015 \mu M^{-1} s^{-1}$	
A_c^B	$0.07 \mu M$	[5]
A_c^P	$0.69 \mu M$	[5]
[A]	$10-40 \mu M$	
[CP]	$0.15 - 0.5 \mu M$	
[Arp2/3]	$1.0 \mu M$	[8]
E_b	$2 - 5 k_B T$	
D_{memb}	$0.04 \mu m^2 s^{-1}$	[9]
D_{mon}	$4 \mu m^2 s^{-1}$	[10]

Parameter Values

On-and-off rates taken from in vitro measurements, other parameters used to obtain reasonable network structures or varied within reasonable ranges

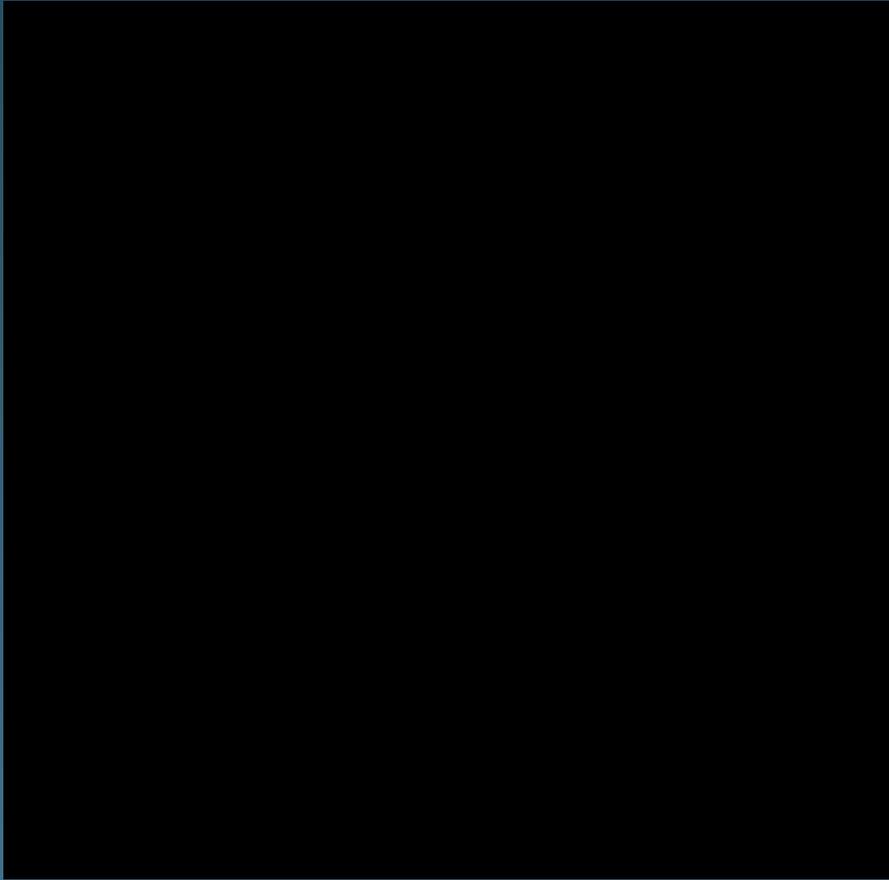
Simulations Show that Branched Actin Networks Form Waves Under Some Conditions



22.5 μM actin 3 μm (PRL, 2010)

Burst of actin polymerization is followed by loss of NPF, which forces actin growth forward

Simulated Fluorescence of Waves



Patchy state can
appear as
interloper

Why Does Dendritic Actin Nucleation Lead to Wave/Patch Formation?

Generic mechanisms leading to wave/patch formation:

- Positive feedback
- Diffusive spreading
- Delayed negative feedback

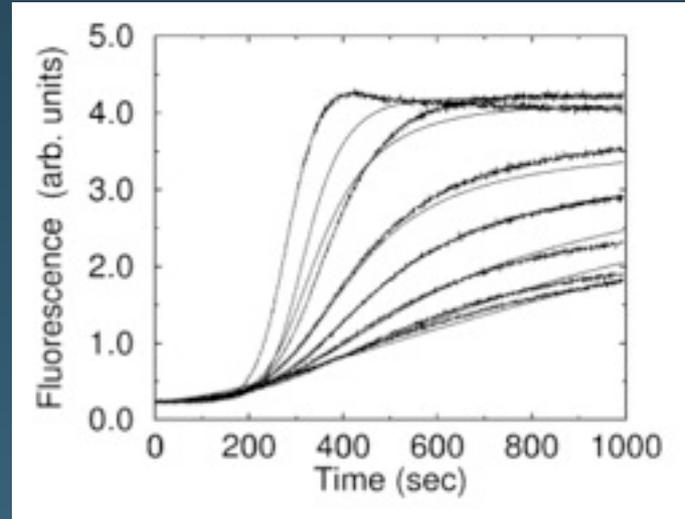
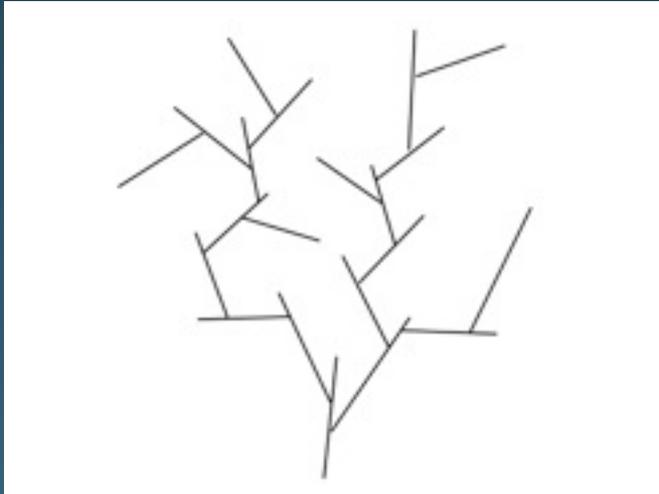
$$\begin{aligned}\dot{u} &= -u + H(u - a) - v + \nabla^2 u \\ \tau \dot{v} &= \mu u - v + L^2 \nabla^2 v,\end{aligned}$$

(Krischer and Mikhailov 1994)

u =activator (F-actin)
 H = positive feedback (step fcn)
 v = inhibitor (absence of NPF)

Branching Causes Positive Feedback

Bulk polymerization with Arp2/3 complex



Martin Wear
(Cooper Lab),
2004

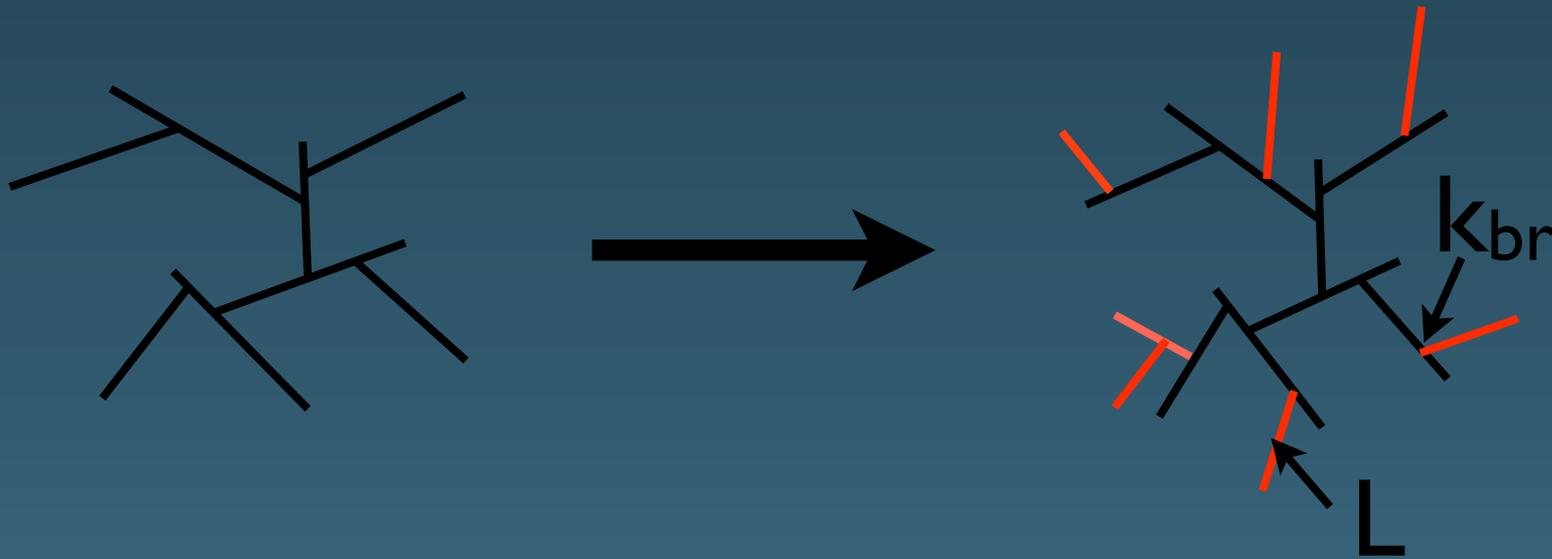
(Fluorescence measures polymerized-actin content)

Number of filaments
↓
$$dN/dt = k_{br} N$$

↑
Branching rate

Polymerization has lag phase followed by exponential growth, characteristic of positive feedback

Diffusionlike Spreading of F-Actin Clusters Results from Branching



$$D_{\text{eff}} \approx k_{\text{br}} L^2 / 12 \approx 0.01 \mu\text{m}^2/\text{s}$$

D_{eff} is probably greater than the physical diffusion coefficient D because of attachment of dendritic clusters to membrane

Wave/Patch Formation is Favored by:

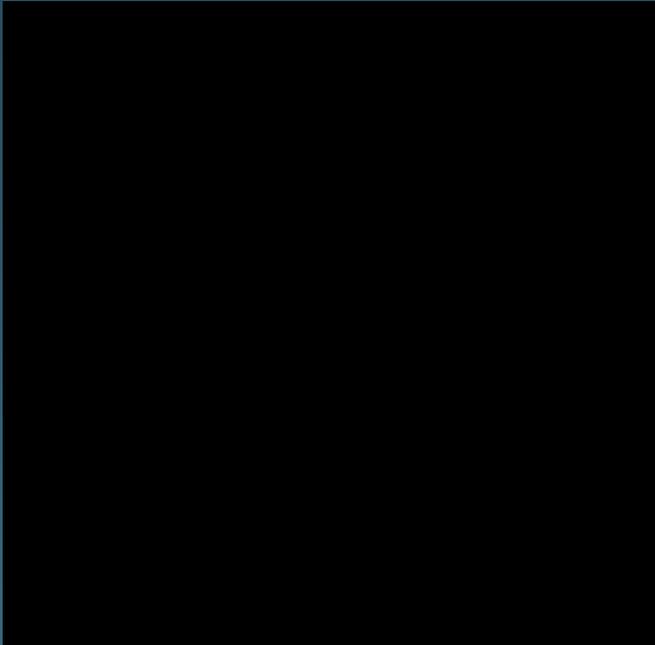
- Slow NPF reattachment
- Slow spontaneous nucleation of actin filaments
- Optimal values of actin polymerization rate and binding strength to membrane

Differences from “Ordinary” Chemical Waves:

- Spreading is not via Brownian motion
- Positive feedback is strongly stochastic

Concrete Predictions:

Reducing actin concentration will cause wave-patch transition

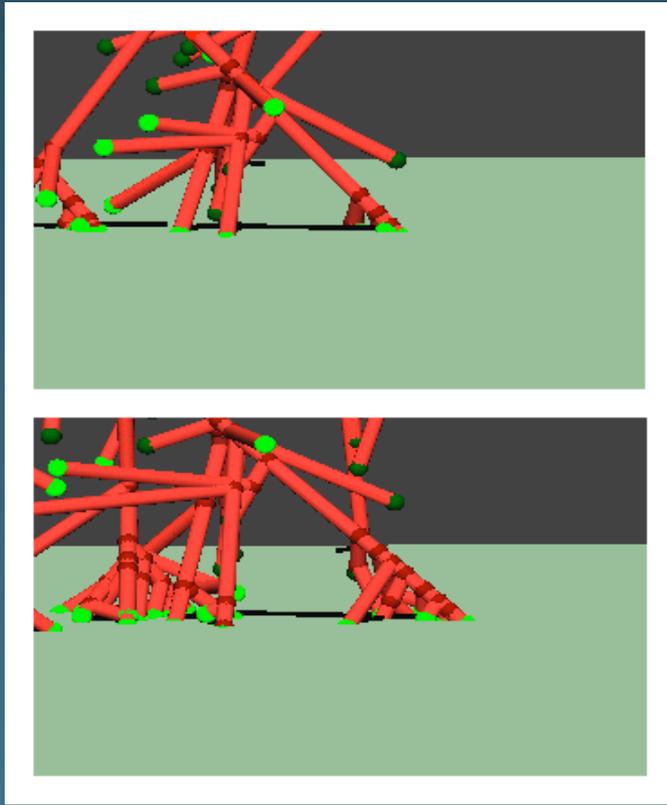


15 μM actin

Patches form because of depletion effects

Consistent with patch-wave transition seen during actin recovery after depletion (Bretschneider et al 2004)

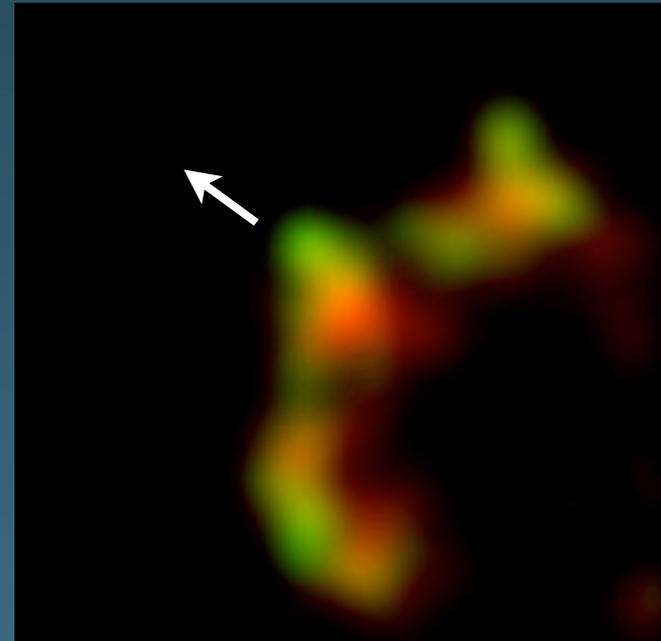
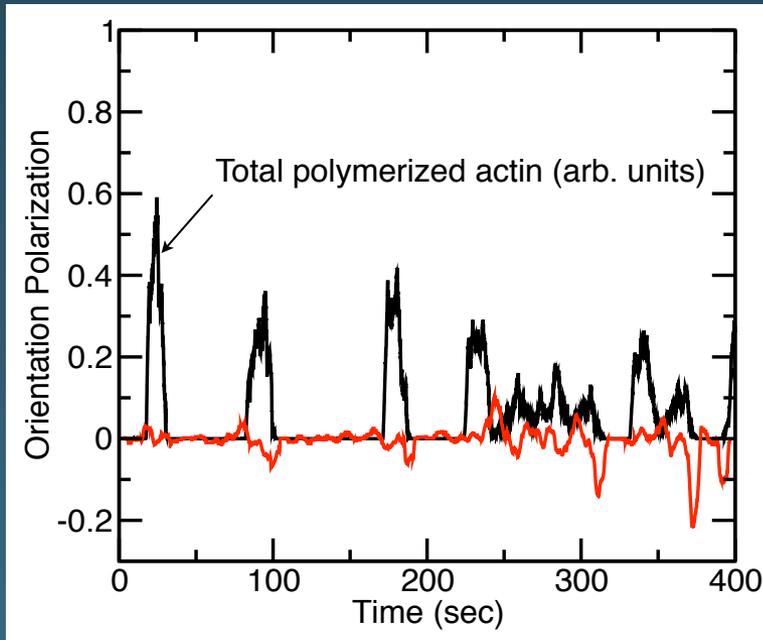
Waves and patches propagate by treadmilling based on branching at edges of waves/patches - not by physical motion of F-actin



0.2 sec between frames

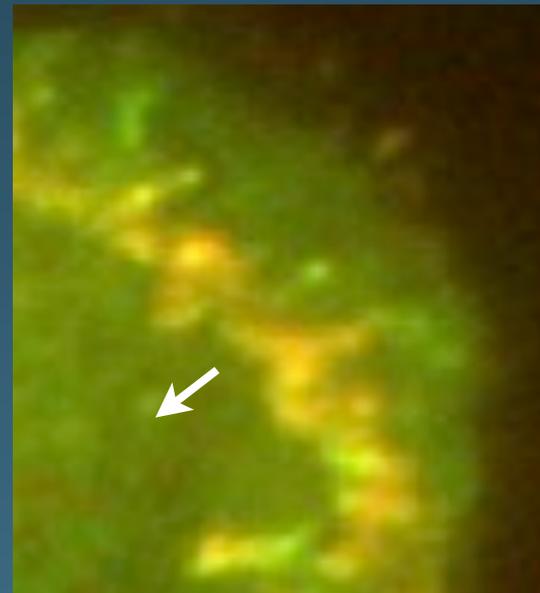
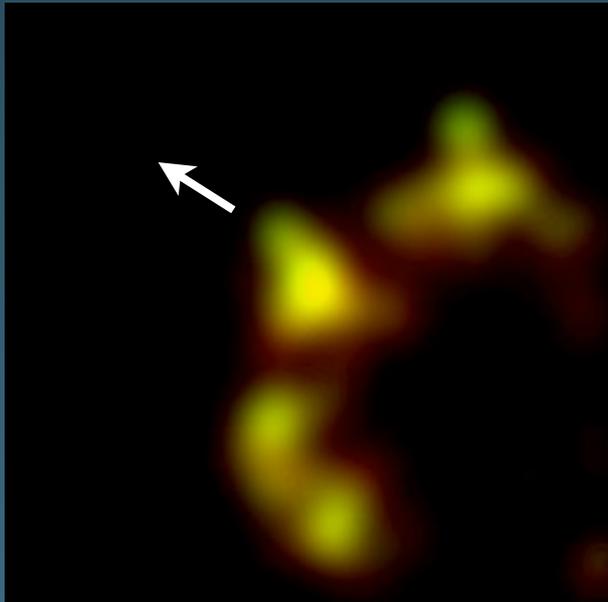
Treadmilling motion has been seen in FRAP experiments (Bretschneider et al 2009)

Filament orientations are not strongly polarized, but the distribution of free barbed ends is:



Green: free barbed ends
Red: F-actin
Arrow: direction of wave motion

Distribution of Arp2/3 complex is broadly similar to that of actin



Green: Arp2/3 complex

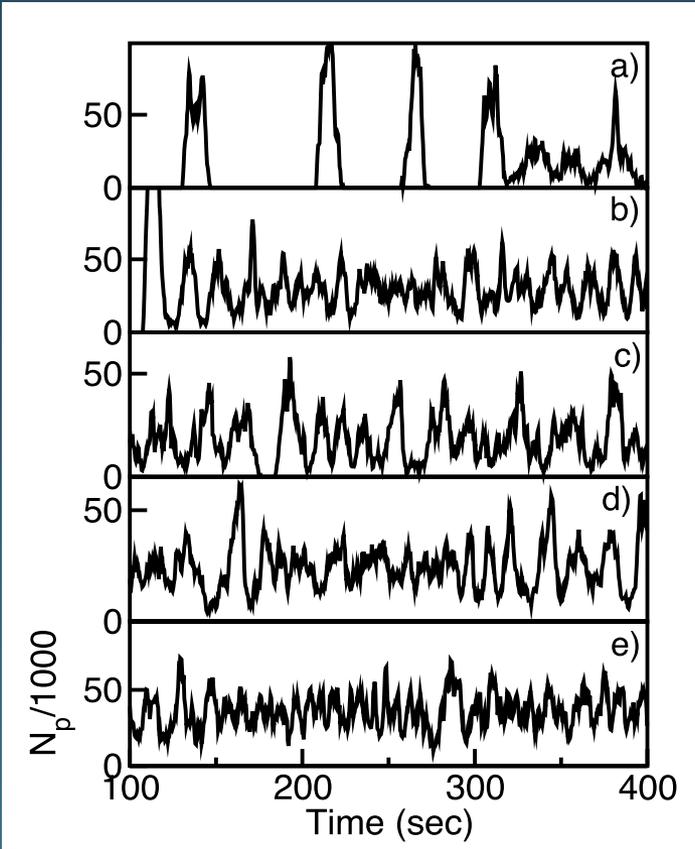
Red: F-actin

Arrow: direction of wave motion

Expt (Bretschneider et al 2009)

Interventions favoring actin polymerization destroy waves

Wavy state has bursts in # polymerized subunits N_p



Baseline (wavy state)

Reduced capping

Faster nucleation

Increased filament binding

Faster NPF reattachment

These effects can be implemented by manipulating key protein concentrations

Diffusion of NPFs slows, and eventually freezes patch/wave motion, while diffusion of filament clusters accelerates wave motion

Speedup due to cluster diffusion is a factor of two if they diffuse freely

Attachment of filaments to substrate tunes the behavior in the same way as the actin concentration:

Weak attachment - patches

Medium-strength attachment - waves

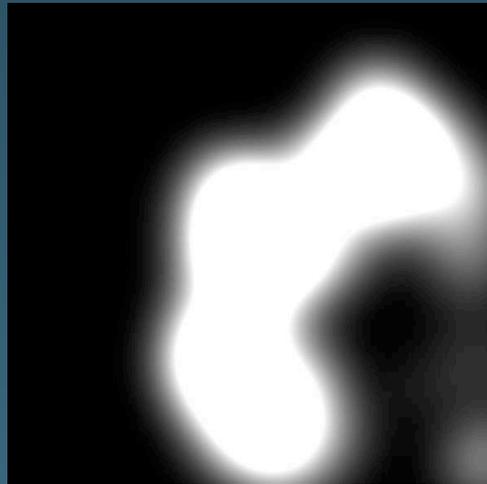
Strong attachment - uniform coverage

Dynamic Phases of F-Actin

Patch



Wave

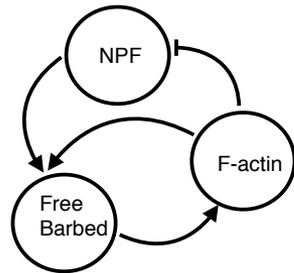


Random

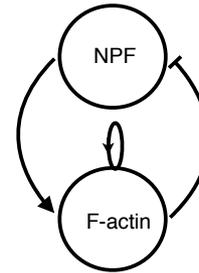


Increasing actin concentration

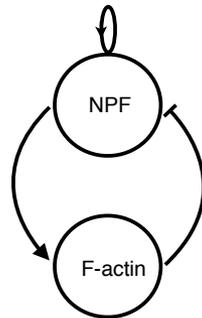
Possible NPF-Actin Feedback Loops



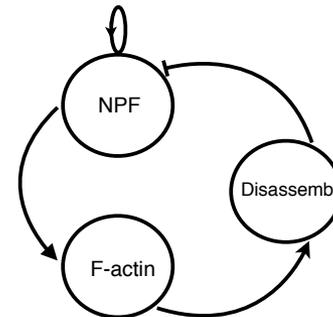
Stochastic-growth simulations



Whitelam et al, 2009

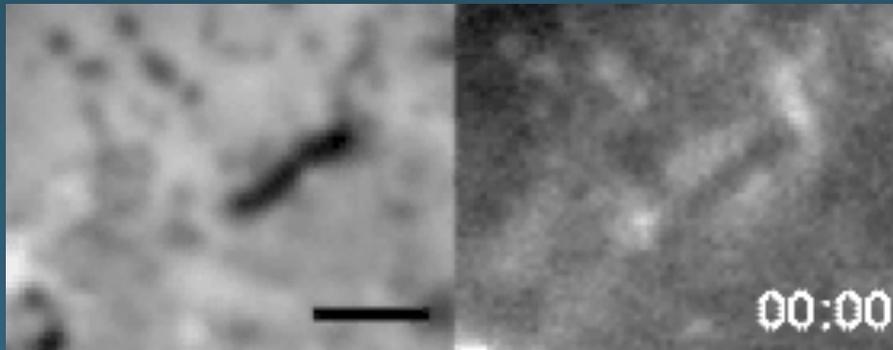


Weiner et al 2007;
Dobrovinski and Kruse 2008



Possible model for actin dynamics
in endocytosis

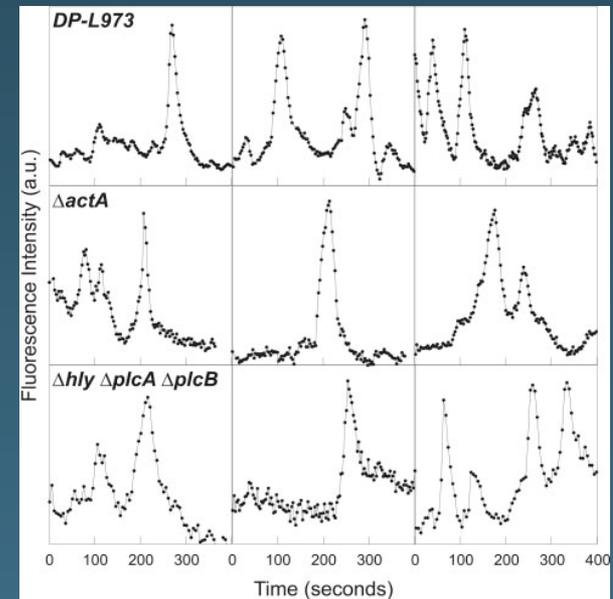
Listeria Phagosomes Can Exhibit Spontaneous Oscillations



Phase

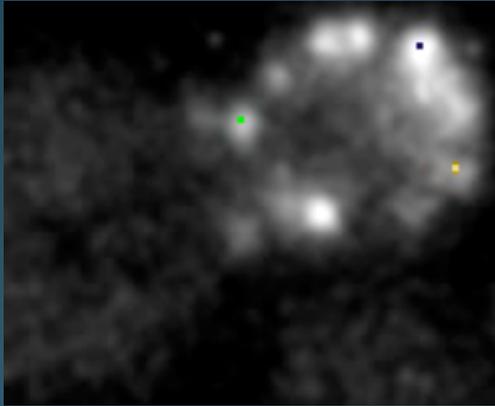
GFP-actin

(Theriot lab 2004)



Could these be due to a similar combination of positive and negative feedback?

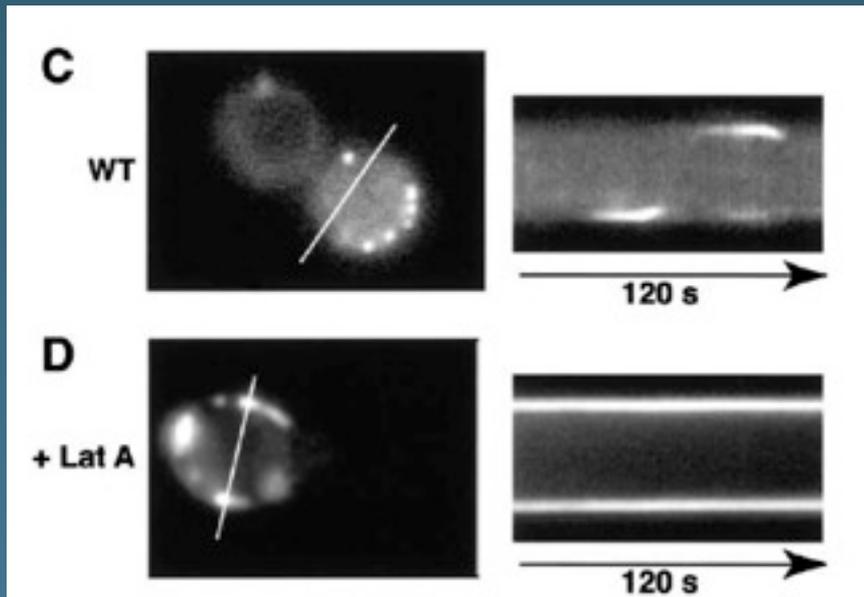
Does Such a Mechanism Describe Dynamic Actin Patches in Yeast?



(Cooper lab 2006)

Fluorescence of a pre-NPF (Sla1)

(Kaksonen et al 2003)



Sla1 patch disassembly
requires F-actin

But patch assembly does
not require F-actin

Modeling a Transient Actin Patch

NPF

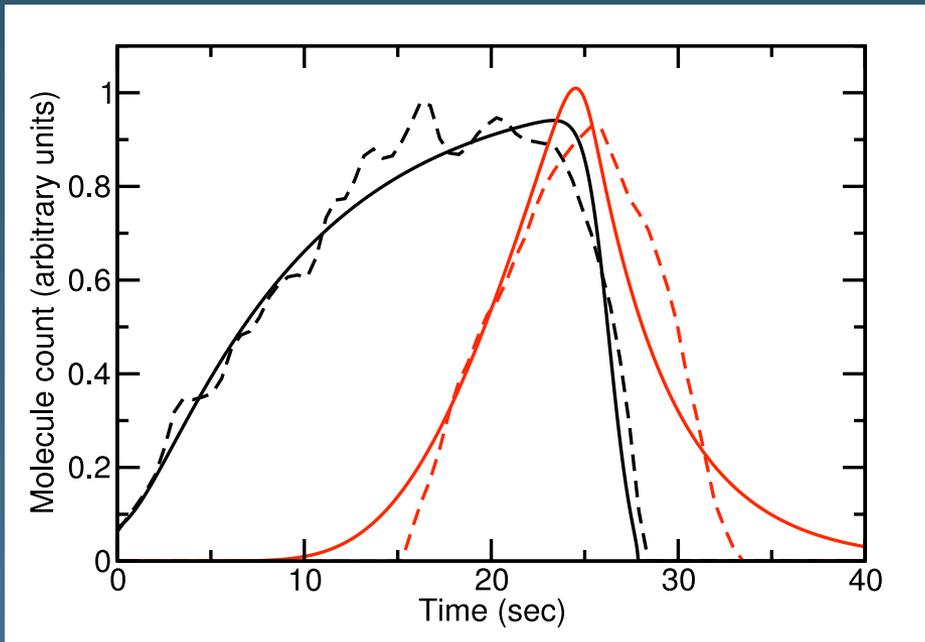
F-actin

Disassembly Proteins

$$\frac{d[N]}{dt} = k_N^+(N_0 - [N]) - k_N^+ N_0 \exp[\epsilon_s(1/\sqrt{N} - 1/\sqrt{N_c})] - k_N^- [D]^i [N]$$

$$\frac{d[F]}{dt} = k_F^+ [N]^j - k_F^- [F]$$

$$\frac{d[D]}{dt} = k_D^+ [F] - k_D^- [D]$$

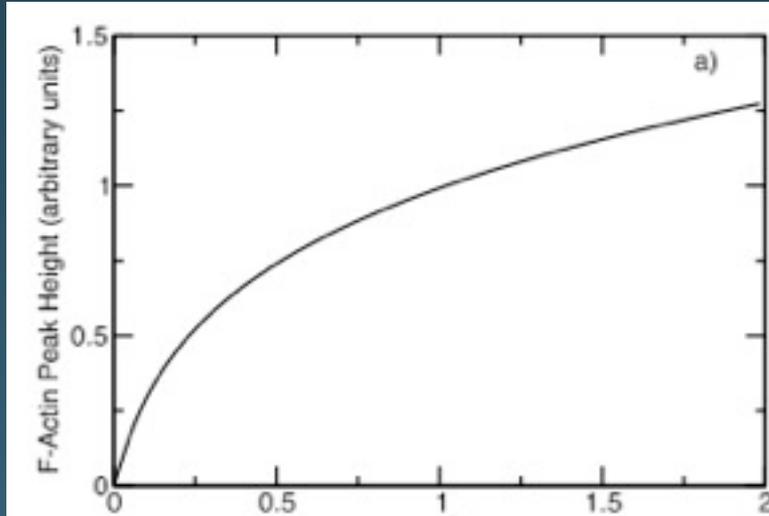


Fit to experimental time courses from Drubin lab

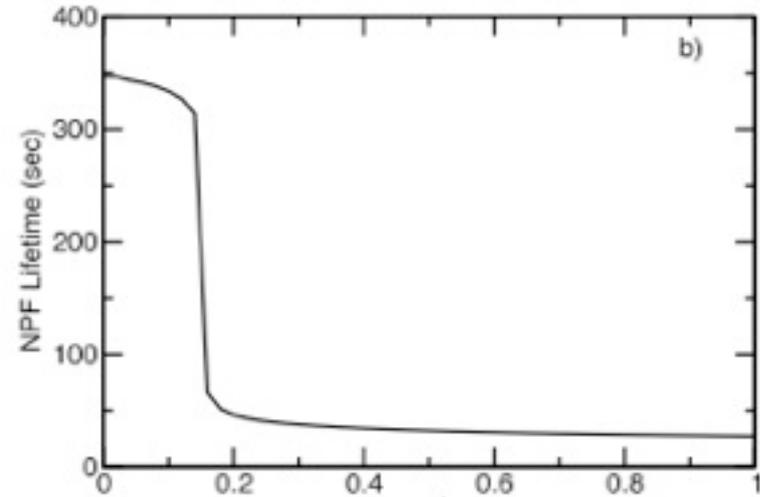
Black: NPF

Red: actin

Effect of Branching Rate on Patch Properties



Branching rate



Branching rate

Actin peak height increases slowly with branching rate - relates to NPF mutation experiments?

NPF lifetime drops abruptly with branching rate

Conclusions

Known actin biochemistry leads to spontaneous formation of waves and patches

Characteristic transitions in dynamic behavior are seen with varying actin concentration

Negative feedback of F-actin on membrane proteins might be a common mechanism leading to oscillating or transient behavior

Currently working on including myosin in network model

Supported by joint DMS/NIGMS initiative in mathematical biology