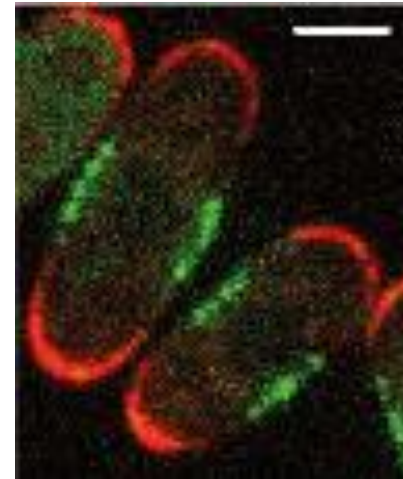
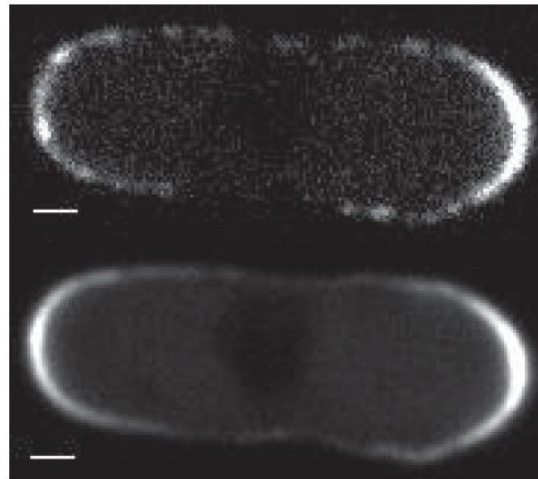
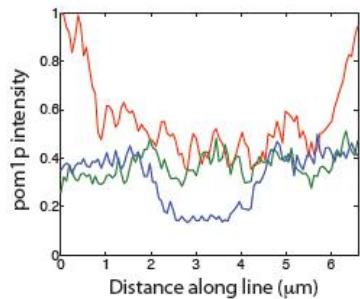
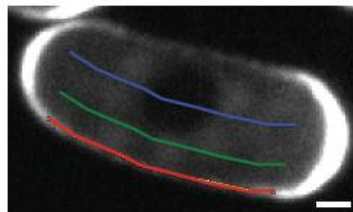


Dynamics of intracellular gradients and cell size control in fission yeast



Martin Howard

**Computational & Systems Biology
John Innes Centre, UK**

Acknowledgements

Theory:

Timothy Saunders (John Innes Centre/EMBL)

Andrew Angel (John Innes Centre), Filipe Tostevin (Imperial/AMOLF)

Experiments:

Kally Pan, Ignacio Flor Parra, Fred Chang (Columbia Univ)

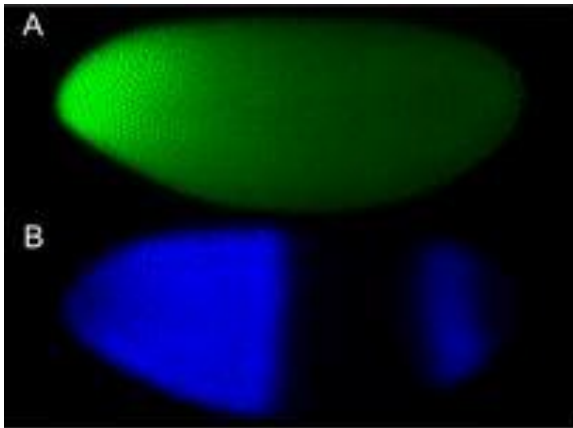
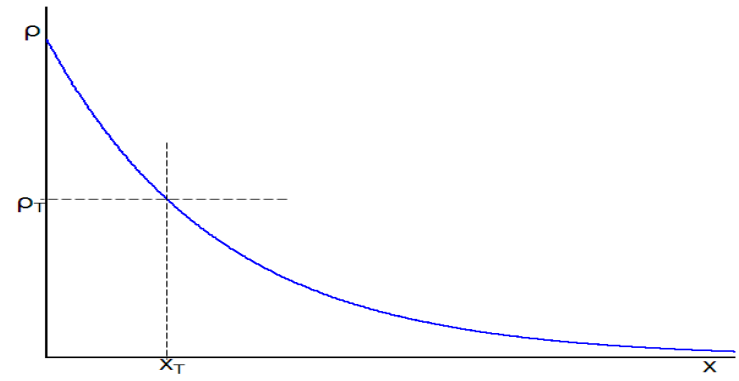
Yinghua Guan and Jagesh Shah (Harvard)

£££:



Concentration gradients in biology

- How to measure **position** in biological systems?
- Use **gradients** of protein concentration ...



Bcd/hb in *Drosophila*

- Use local concentration to drive a switch
- Familiar in **developmental biology**

- Importance increasingly understood in **subcellular** context

A simple model of a concentration gradient

- Source: production at $x=0$ at rate J
- Diffusion: Diffusion constant D
- Disassociation/Degradation: at rate μ

$$\frac{\partial \rho}{\partial t} = D \nabla^2 \rho - \mu \rho + J \delta(x)$$

- At steady-state:

$$\rho(x) = \frac{J \lambda}{D} \exp\left(-\frac{x}{\lambda}\right) \quad \text{where} \quad \lambda = \sqrt{D / \mu}$$

- **Very simple model**, but is it biologically relevant in single cells?

Can gradients exist inside single cells?

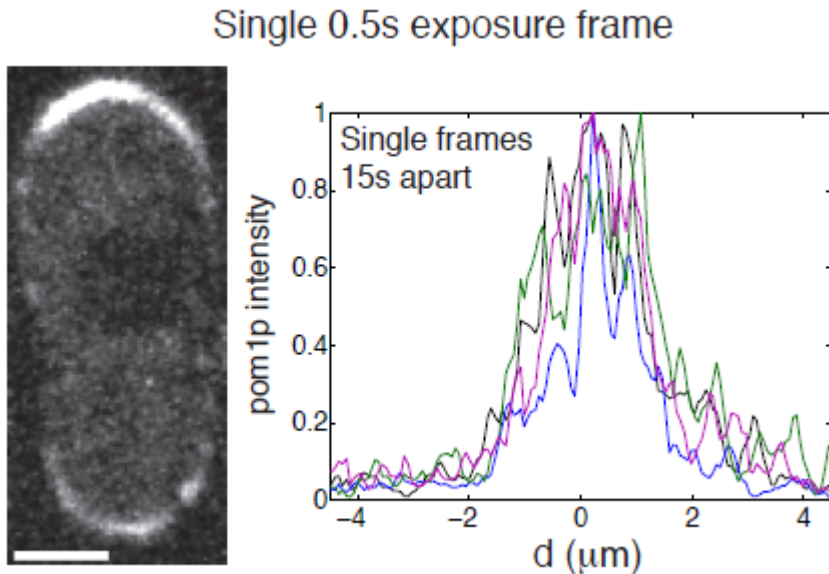
- Need $\lambda = \sqrt{D/\mu}$ to be a few μm
- Suppose gradient is on membrane
- With $D \sim 0.1 \mu\text{m}^2\text{s}^{-1}$
- For $\mu \sim 0.1 \text{s}^{-1} \Rightarrow \lambda \sim 1 \mu\text{m}$
- Short dwell time $\sim 10 \text{s}$
- But this isn't protein lifetime!
- Protein can be recycled from cytoplasm back to membrane
- Subcellular gradients are likely to be plausible/ubiquitous

**Brown & Kholodenko
FEBS Lett. (1999)**

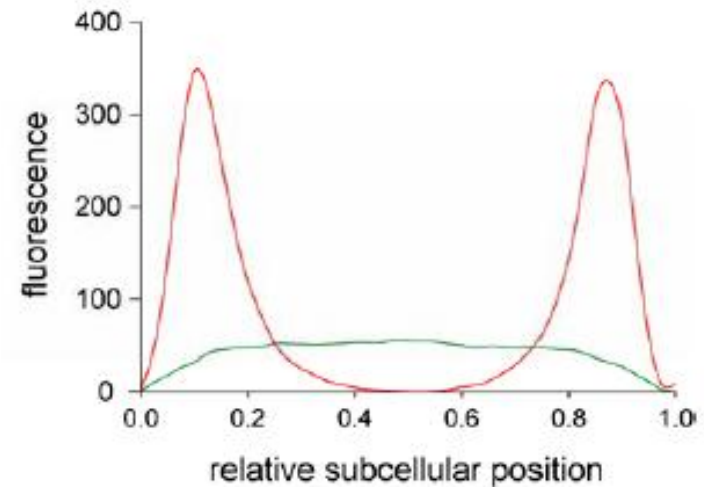
Examples: intracellular gradients

- Cell division:

Pom1p in fission yeast



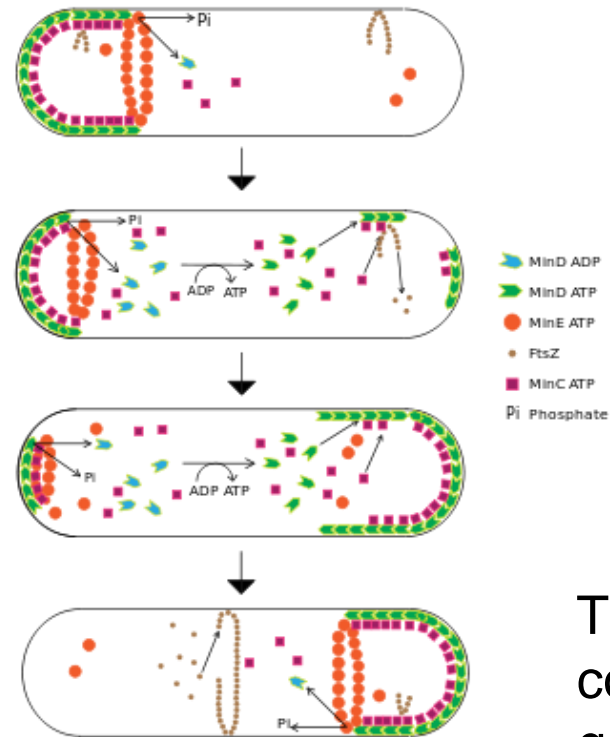
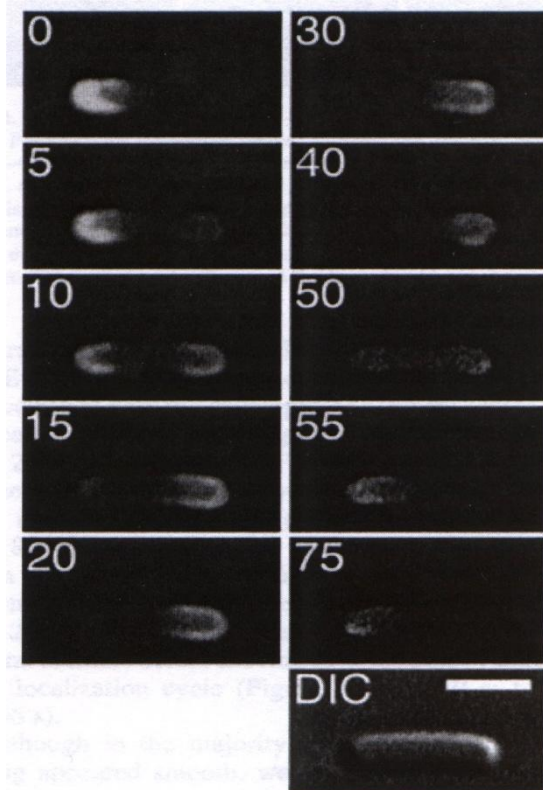
MipZ in *Caulobacter*



Thanbichler & Shapiro,
Cell (2006)

Examples: intracellular gradients

- MinCDE system in *E. coli*:



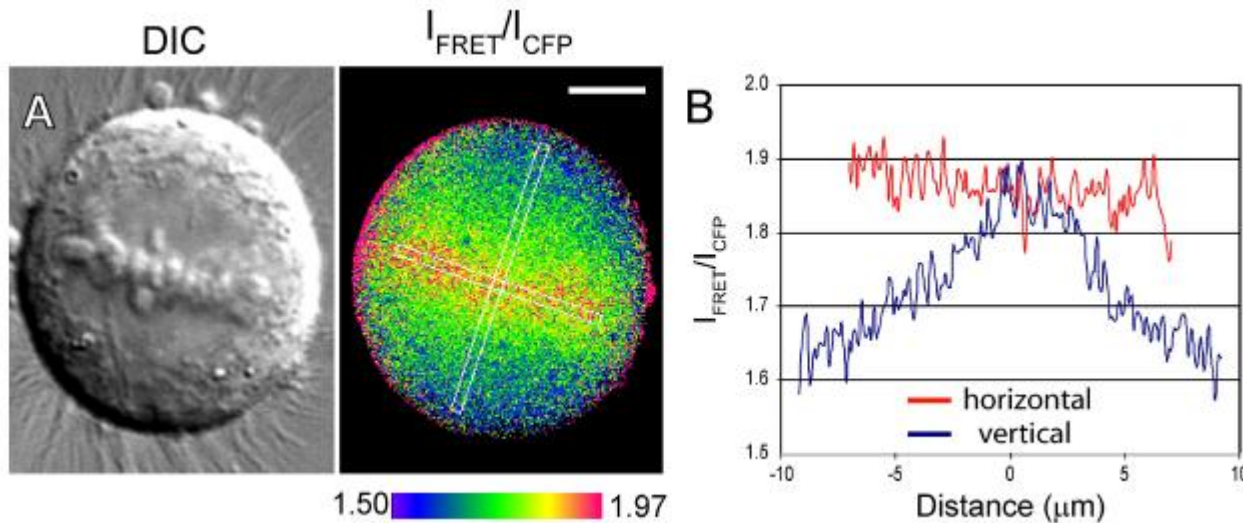
Time-averaged
concentration
gradient!

de Boer et al. PNAS (1999)

Howard et al. PRL (2001) also K. Kruse

Examples: intracellular gradients

- RanGTP system:

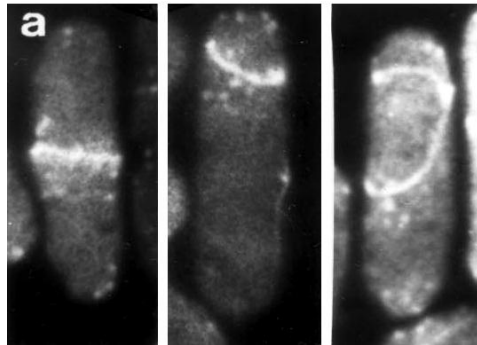


- RanGTP promotes microtubule nucleation/stabilization, aiding in spindle assembly

O'Connell et al JCB (2009)

Cell division positioning in *S. pombe*

- Nucleus pushed to cell centre via microtubules
- Nucleus then signals to cell cortex via **Mid1p**
- **Mid1p** localizes to broad band overlying the nucleus, shuttles in and out of nucleus and ring
- **Mid1p** then controls location of cytokinetic machinery



F-actin rings in wt and *mid1* mutant

- *mid1* mutant forms rings at random sites not linked to nucleus

Pom1p is an inhibitory gradient

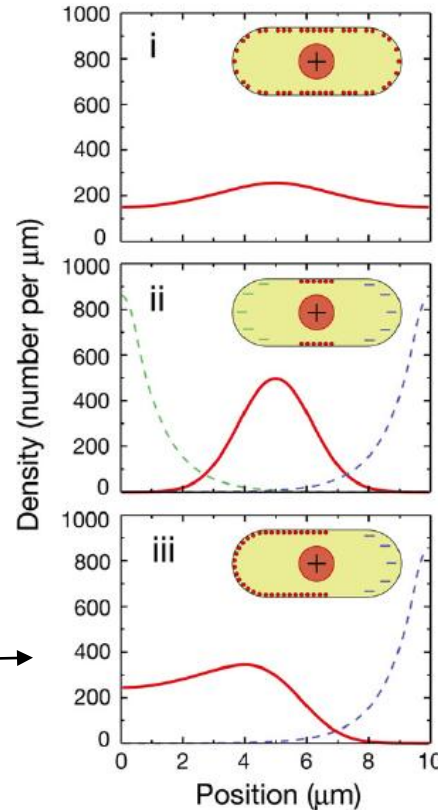
- Rapid cytoplasmic diffusion \Rightarrow nearly homogeneous Mid1p distribution

- Assume existence of polar inhibitor gradients

- If one polar inhibitor is removed, expect Mid1p to spread to that end of cell

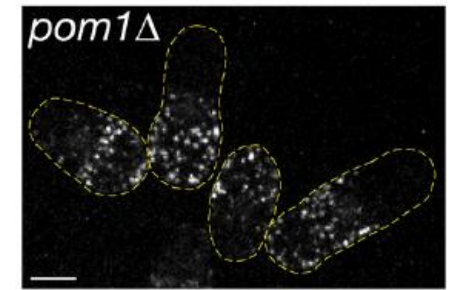
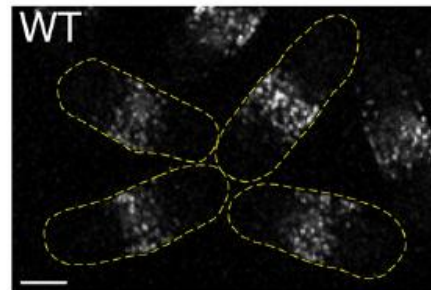
- *pom1* mutant exhibits cell division defect

- Predict that one of polar “morphogens” is Pom1p

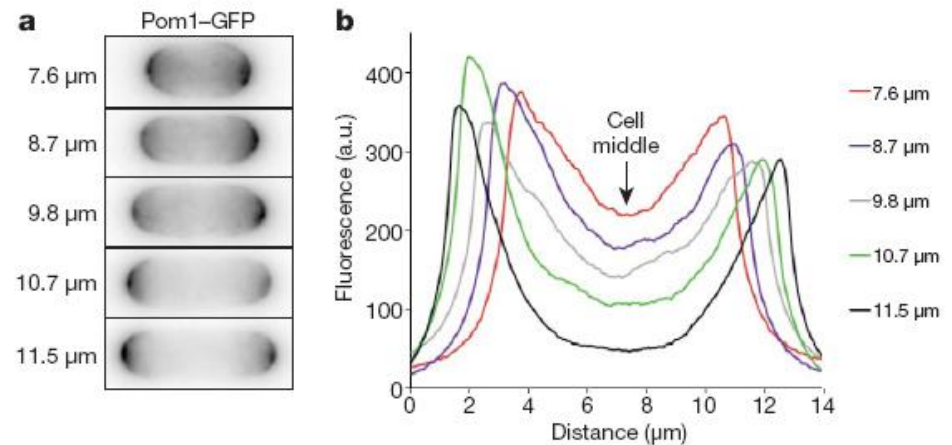
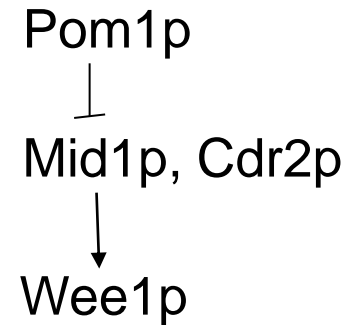
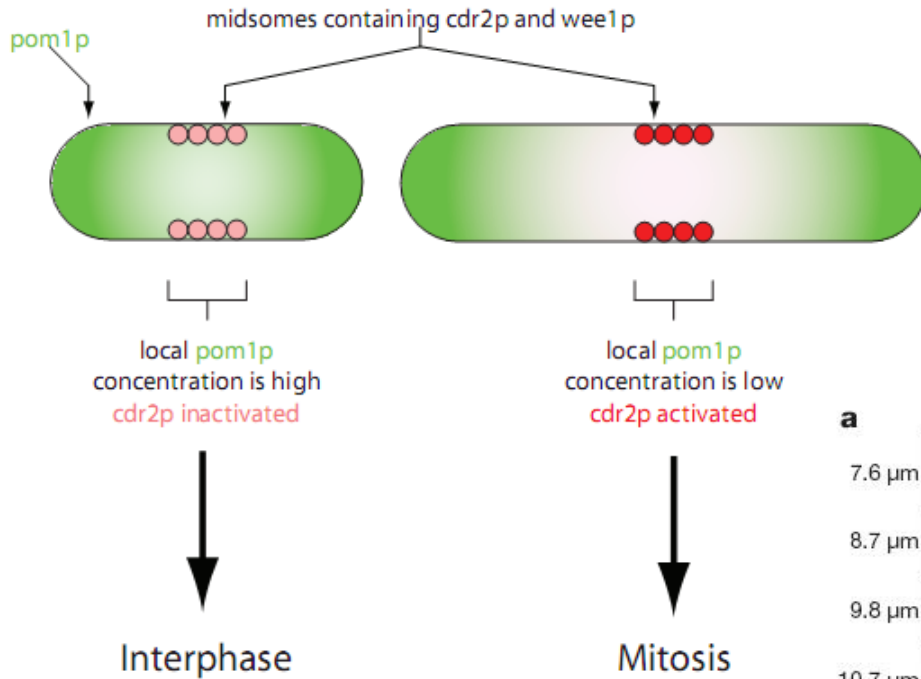


Padte et al. *Curr. Biol.* (2006)

Also Paoletti lab (2006)



Pom1p gradient as a cell-size sensor?



Moseley et al, Nature (2009)

Martin & Berthelot-Grosjean, Nature (2009)

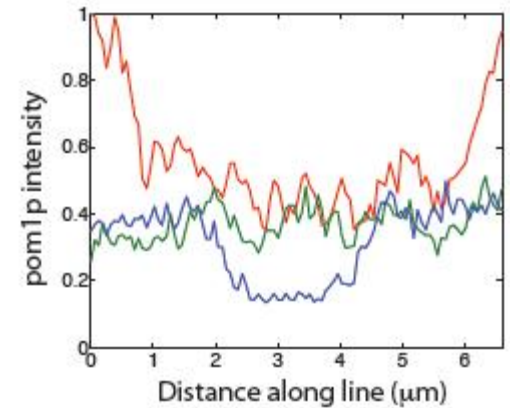
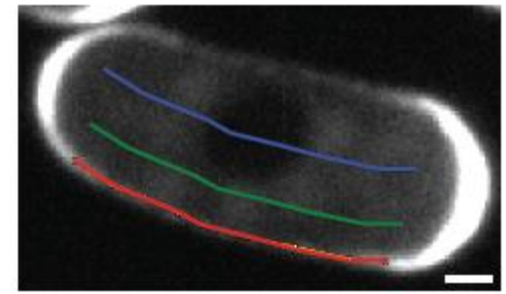
Two important questions!

- What is the mechanism of gradient formation?
 - Simple source, diffusion, disassociation leading to exponential decay with distance?
- How are fluctuations reduced?
 - Gradient imparts positional/size information, so noise must be filtered out
- To answer these questions we need reliable measurements of the gradient profile...

Saunders et al Dev. Cell (2012)
Howard, Trends Cell Biol. (2012)

Extracting the pom1p profile

- Pom1p forms a cortical gradient
- Also cytoplasmic population, but excluded from nucleus



Blue contour is final mask

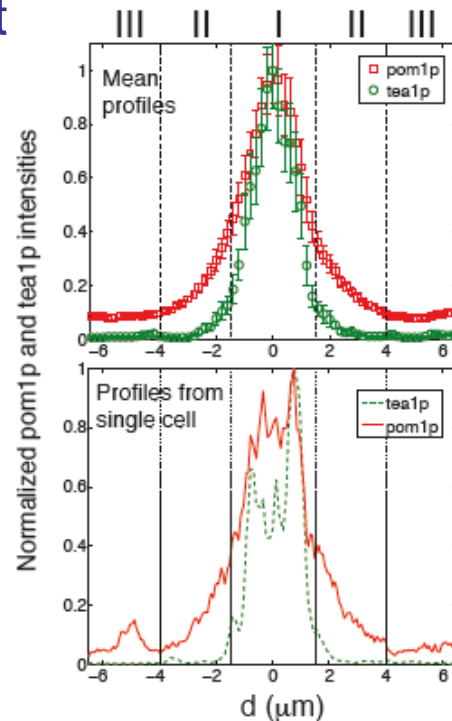
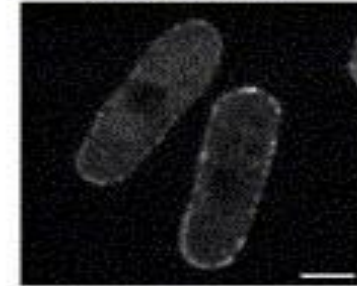
- Computer generates initial mask, using phase image
- Then contour/pixels adjusted to maximise overlap with fluorescent signal
- Finally, human intervention needed to finalise mask

Making the gradient: tea1p dependent injection

- Pom1p homogeneously distributed in *tea1* mutant

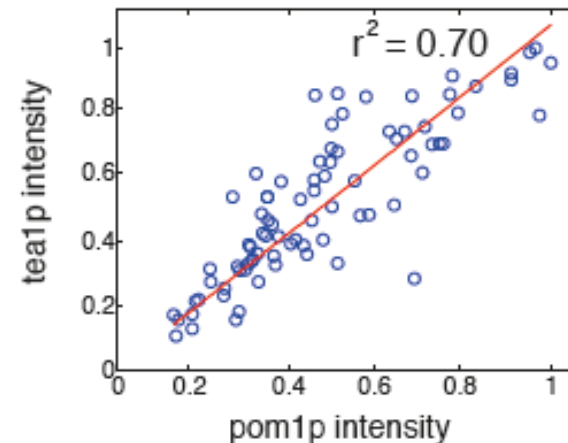
tea1 mutant

pom1-GFP

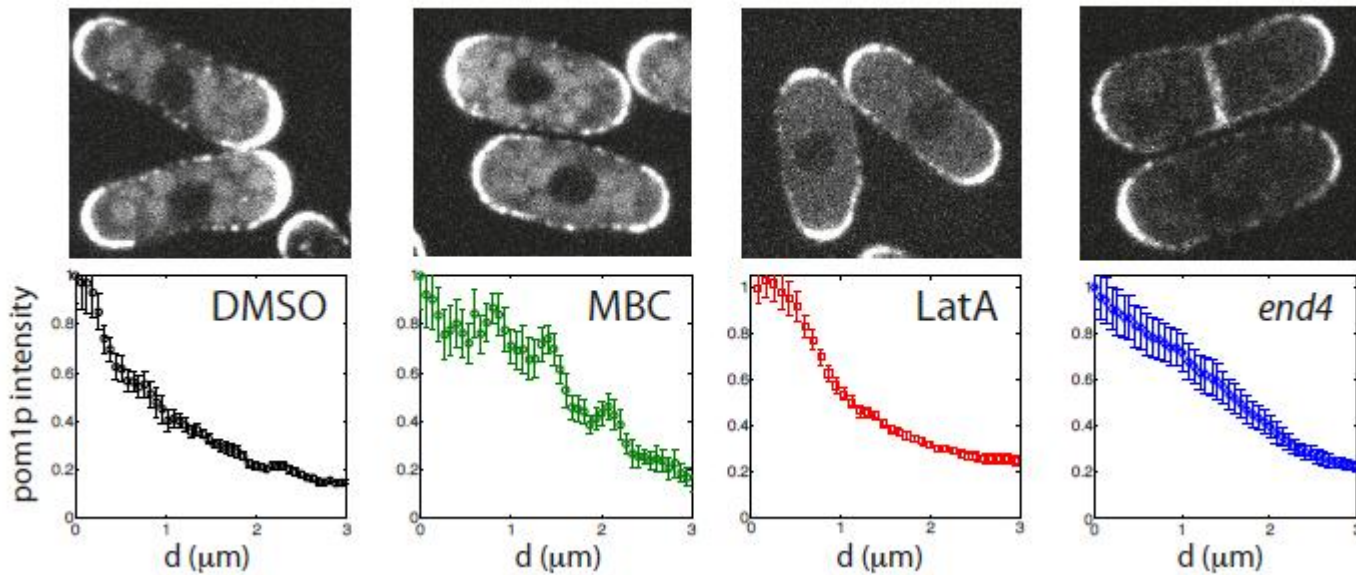


- Pom1p injected at poles and spreads along membrane

- Maximum Pom1p levels proportional to Tea1p consistent with insertion role at tips

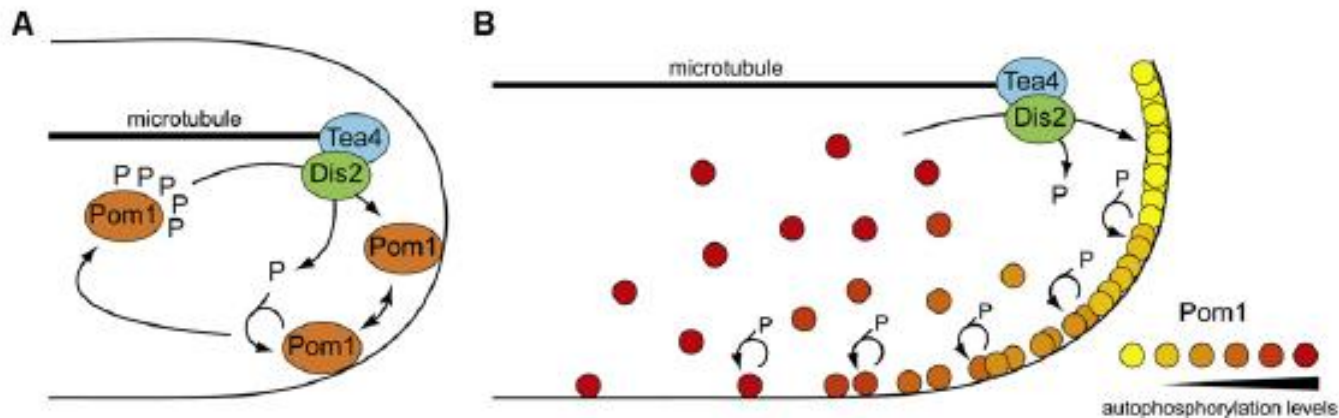


Gradient does not depend on cytoskeleton/endocytosis



- Gradient unaffected by cytoskeletal/endocytic perturbation

Gradient formation involves autophosphorylation

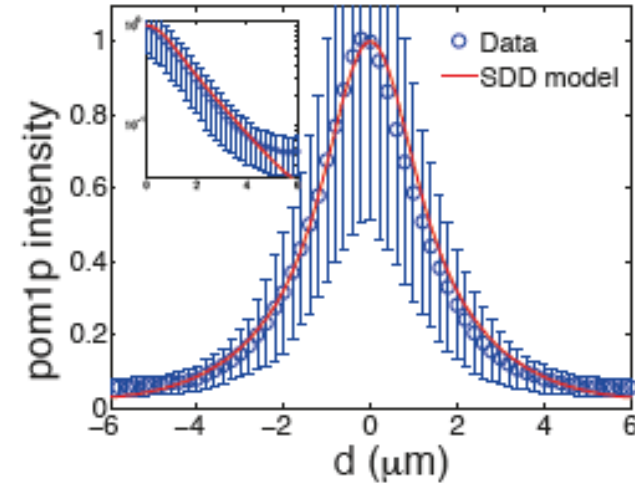


Hachet et al Cell (2011)

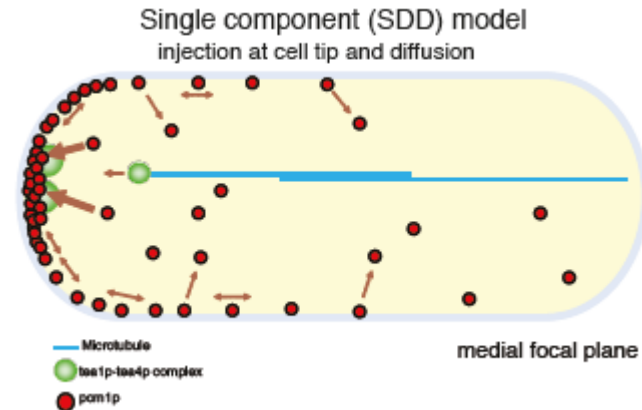
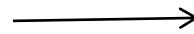
- Pom1p dephosphorylated by Dis2p at microtubule tips, allowing incorporation into membrane
- Subsequently autophosphorylates, losing affinity for membrane

Gradient profile

- Away from tips, profile mostly consistent with **exponential** decay

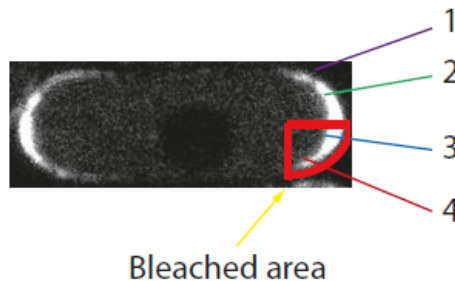
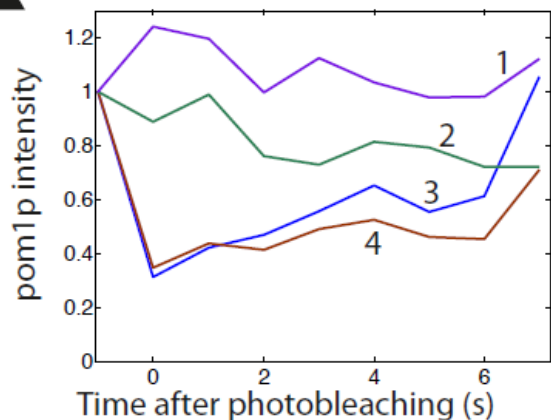
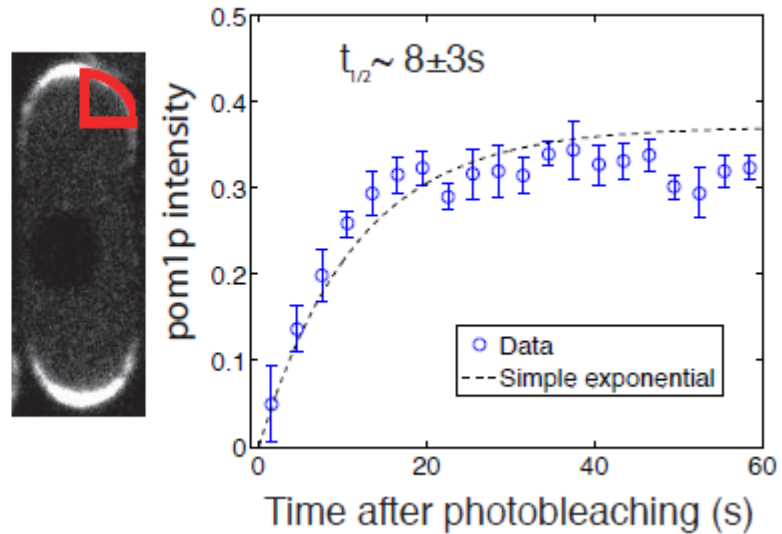
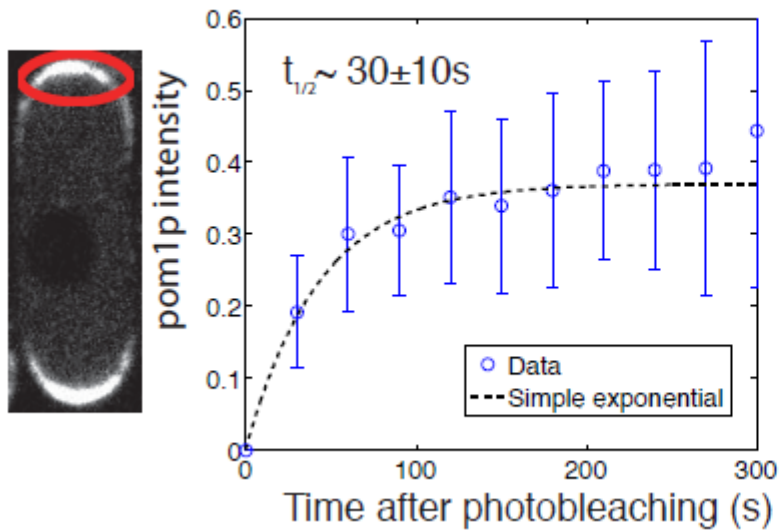


- Consistent with SDD model



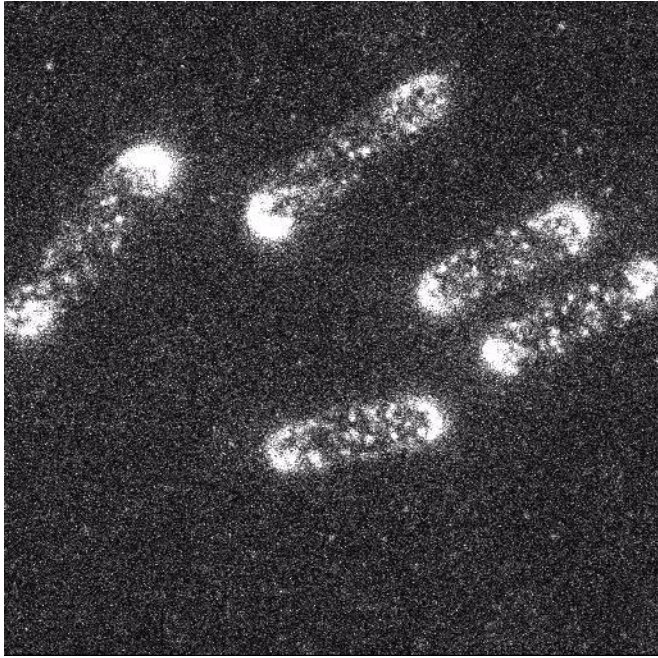
- So it's all very simple ...

FRAP experiments



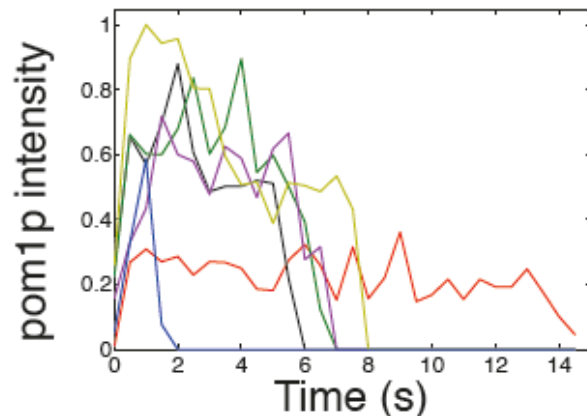
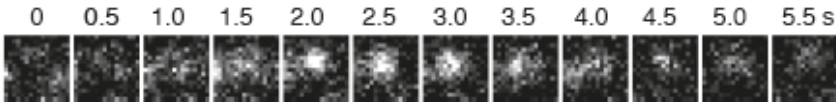
- Consistent with recovery through diffusion from unbleached half-tip
- lifetime $\sim 30s$
- diffusion $\sim 0.1 \mu m^2 s^{-1}$

Imaging reveals a different timescale



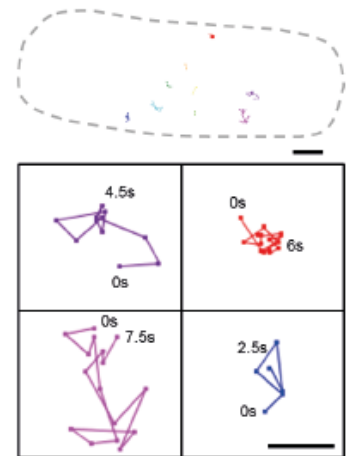
Pom1p forms clusters!

- Pom1p appears to concentrate into blobs that move around cortex ...
- Movie from slice through upper cortex of cell



Cluster tracking:

- Lifetime $\sim 4 \pm 2$ s
- Diffusive tracks
- $D \sim 0.01 \mu\text{m}^2 \text{s}^{-1}$
- Problem!

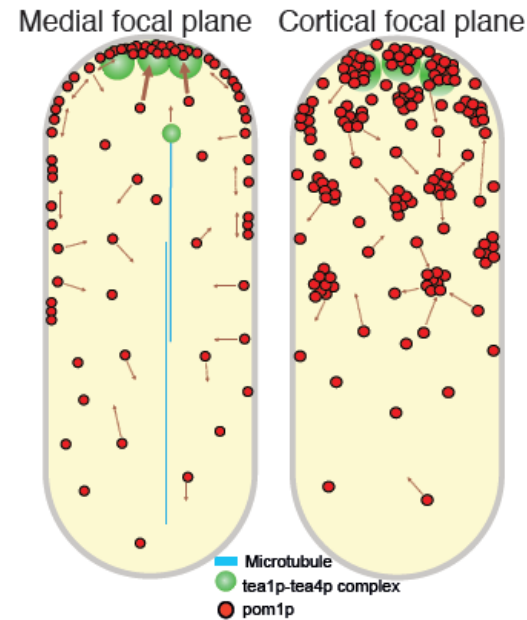


SDD model fails!

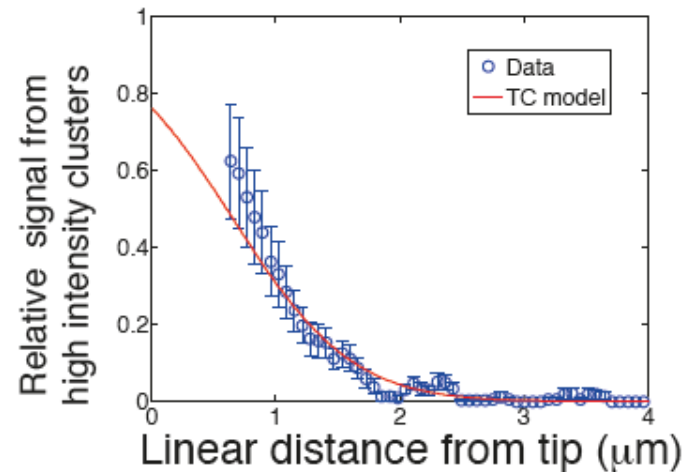
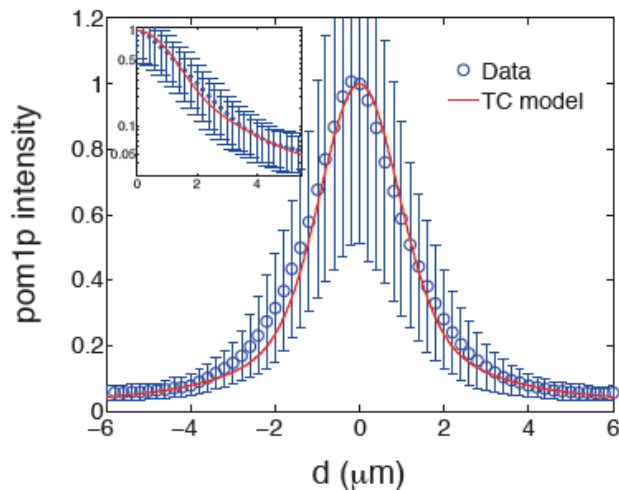
- **SDD** model only has a single lifetime (membrane dwell time)
- Cannot explain two timescales, order of magnitude apart
- Cannot different diffusion constants, factor of 5 different
- Hypothesis: more than one “species” of **Pom1p** on membrane

Two state (TS) model for pom1p dynamics

- Slow-diffusing, clustered form (S) and fast-diffusing form (F)



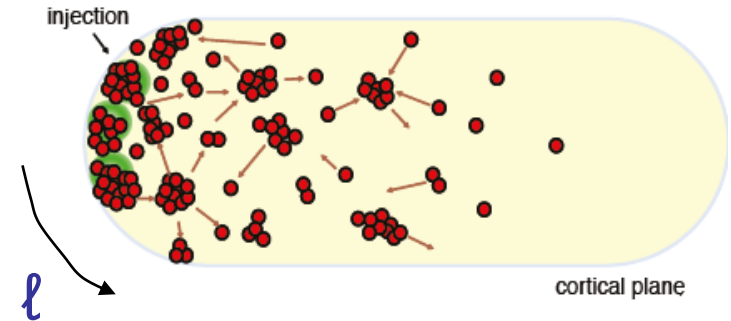
- Fits well to mean intensity, FRAP data, aggregate lifetime, fraction of high intensity clusters



TS model technicalities

- Assume symmetry about long axis of cell
- Use arc-length coordinate ℓ :

$$\nabla^2 = \partial_\ell^2 + 1/(R \tan(\ell/R)) \partial_\ell$$



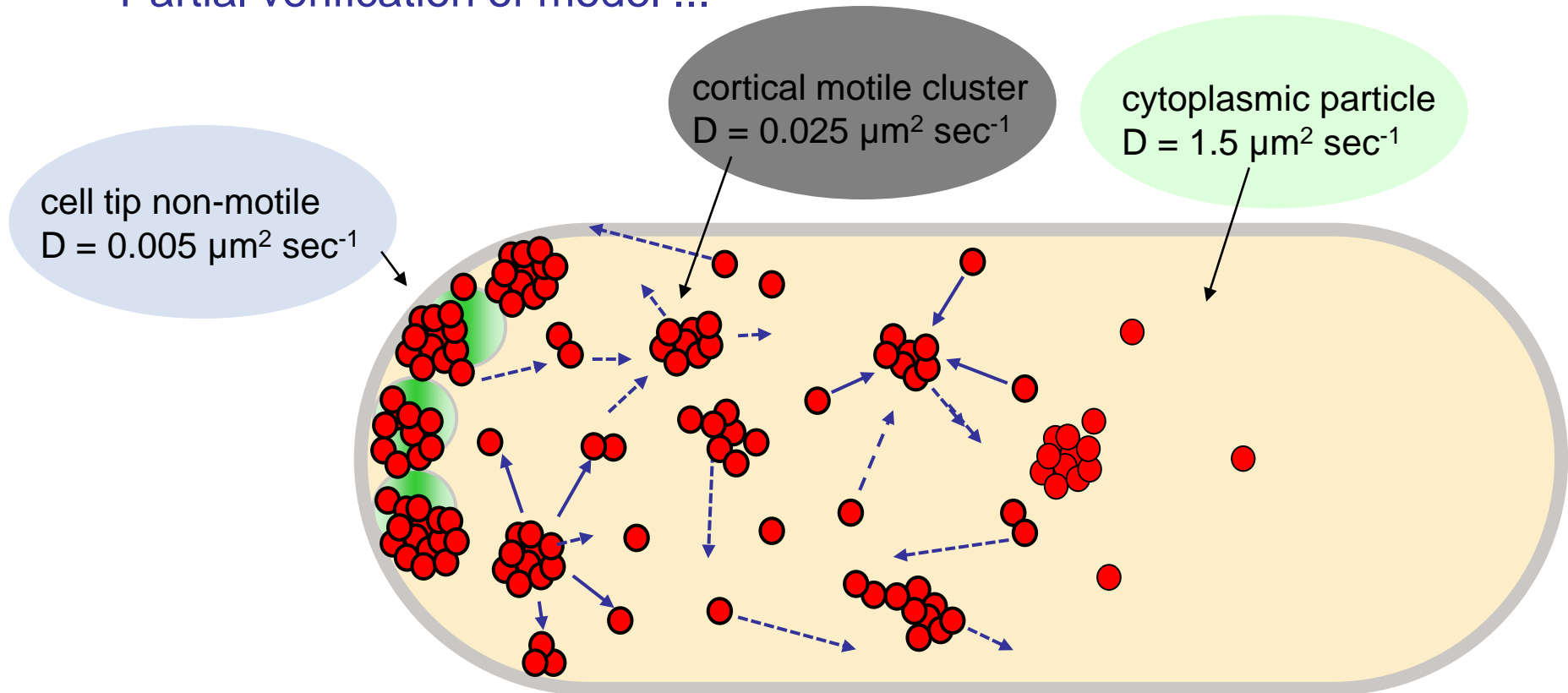
- Simplified equations, for slow and fast-diffusing components:

$$\partial_t \rho_s = D_s \nabla^2 \rho_s - \alpha \rho_s + \beta \rho_f \rho_s + (1 - \epsilon) J f(d/\sigma),$$

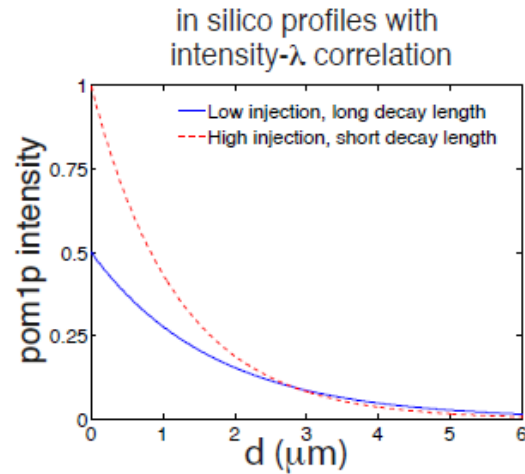
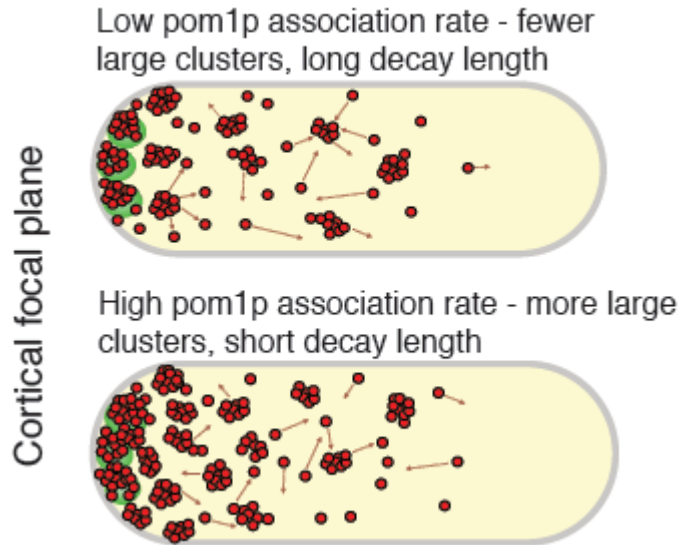
$$\partial_t \rho_f = D_f \nabla^2 \rho_f + \alpha \rho_s - \beta \rho_f \rho_s - \mu \rho_f + \epsilon J f(d/\sigma).$$

Measuring Pom1p membrane mobility

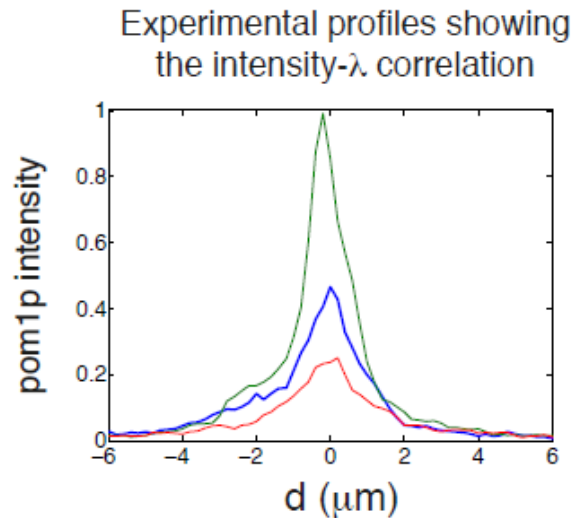
- Model predicts (at least) two membrane diffusion constants
- Use **FCS** to measure **Pom1p** membrane diffusion
- Find wide range of **Pom1p** diffusion constants
- Partial verification of model ...



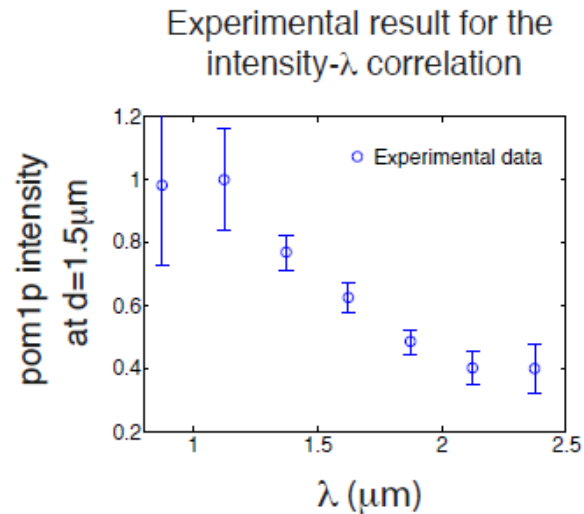
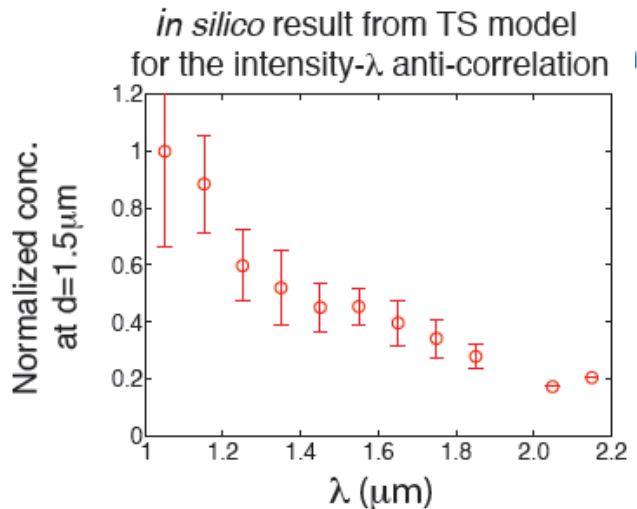
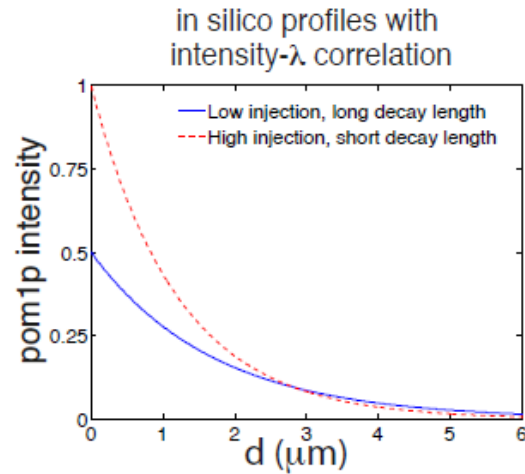
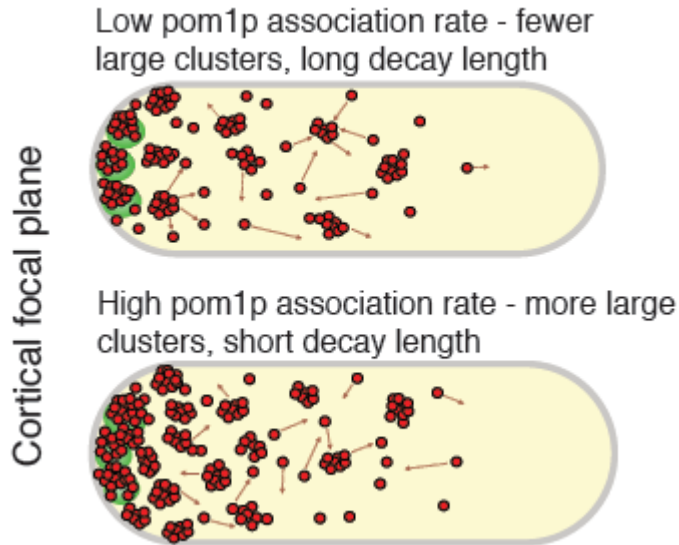
Model prediction: anticorrelation between amplitude and decay length



- Anticorrelation seems to be present in experiments



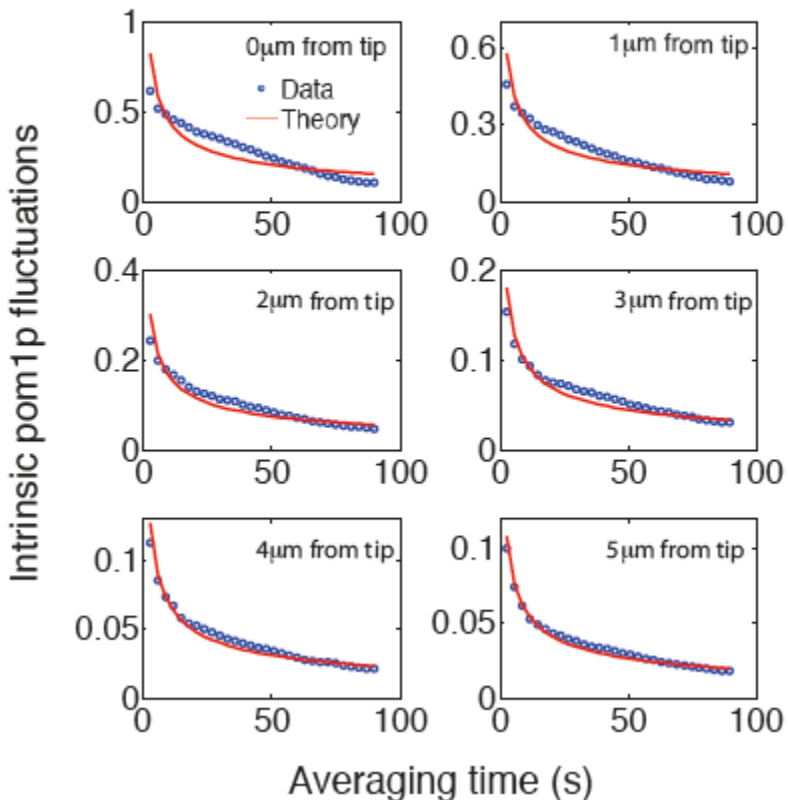
Model prediction: anticorrelation between amplitude and decay length



Reducing intrinsic fluctuations

- Time averaging can reduce intrinsic fluctuations: **Tostevin, ten Wolde, Howard PLoS Comput Biol (2007)**

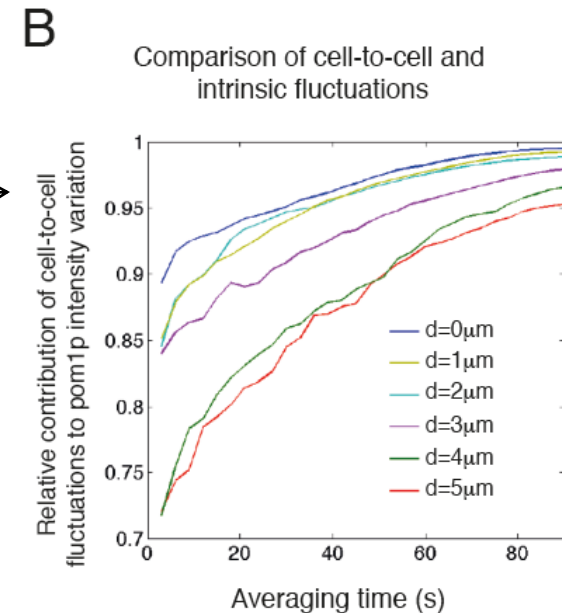
Time-averaging of intrinsic fluctuations at different positions in the cell



- Error reduced by $1/\sqrt{t}$

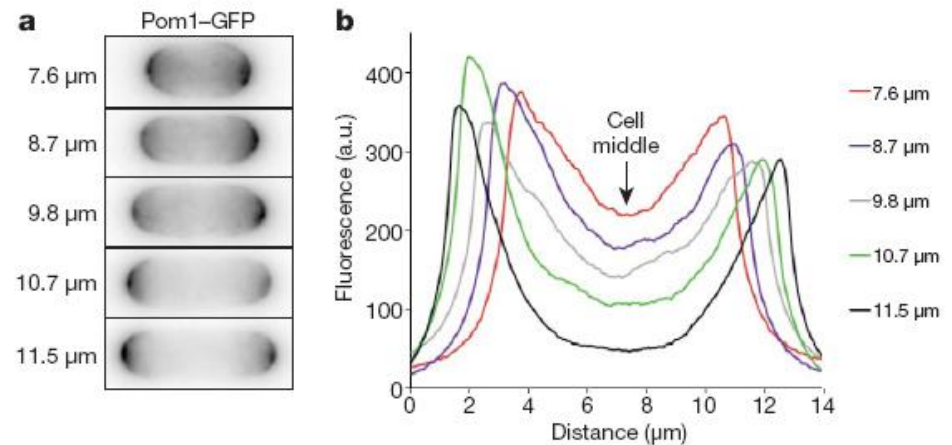
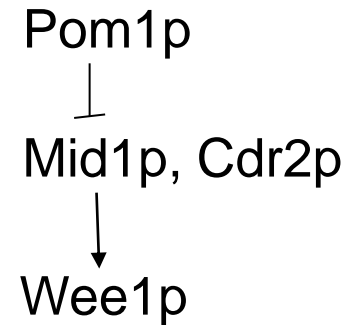
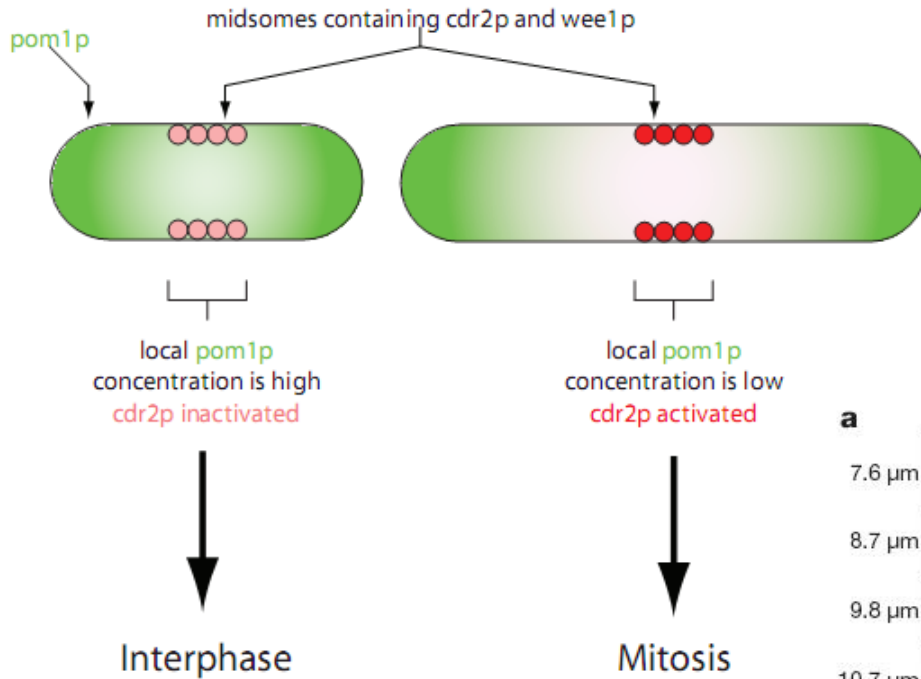
• Cell-to-cell variations dominate after about 10-20s of averaging

• First quantification of intrinsic noise



More general discussion of noisy gradients
Extrinsic & intrinsic noise: Saunders & Howard PRE (2009)
Pre-steady-state gradients: Saunders & Howard Phys. Biol. (2009)

Pom1p gradient as a cell-size sensor?

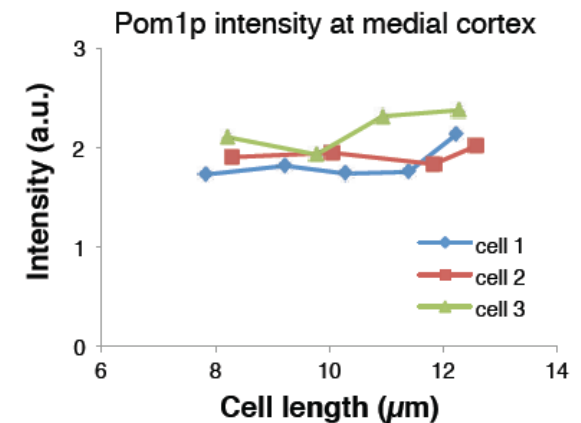
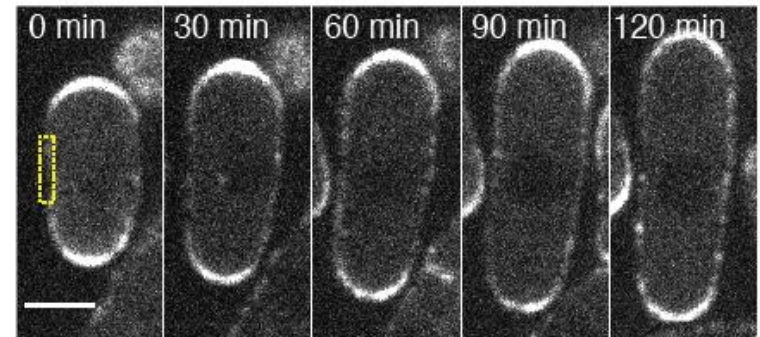
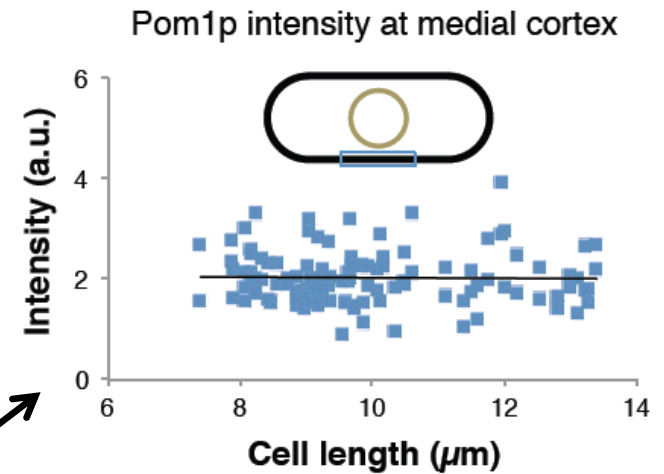


Moseley et al, Nature (2009)

Martin & Berthelot-Grosjean, Nature (2009)

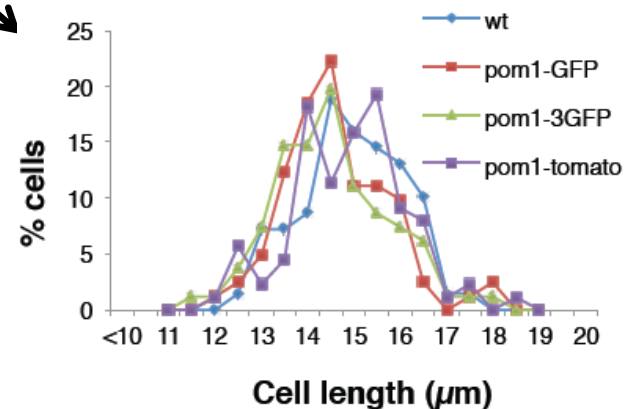
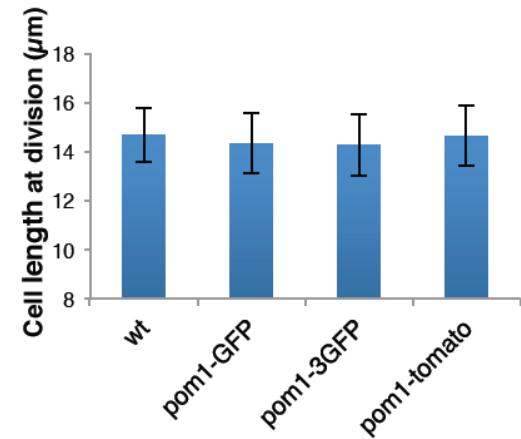
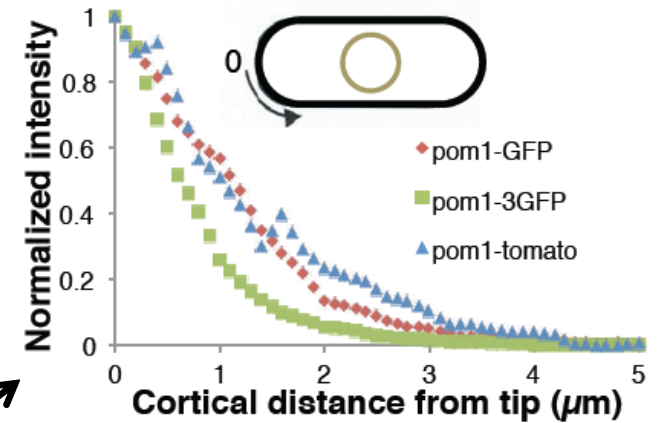
Pom1p gradient as a cell-size sensor?

- Our data does **not** support this hypothesis
- **Pom1p** levels at medial cortex almost constant for different cell lengths in population of cells
- Same results holds when single cells are monitored



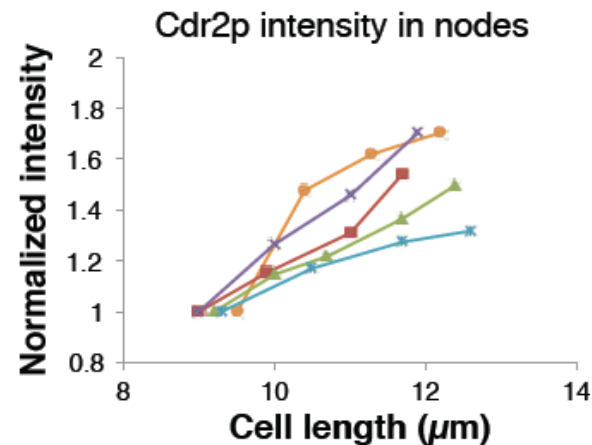
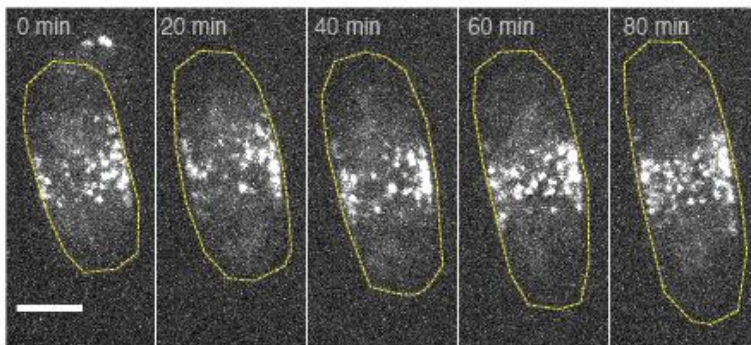
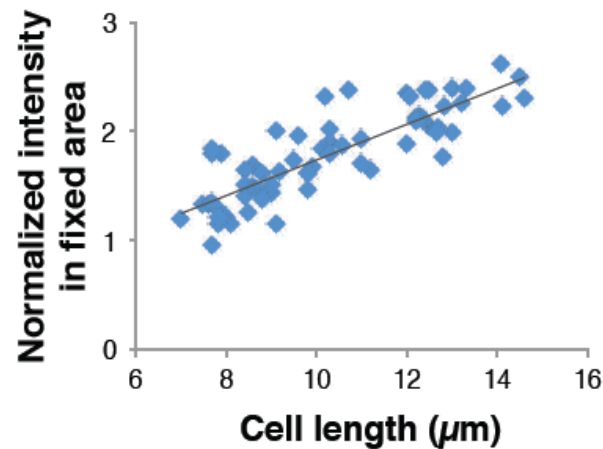
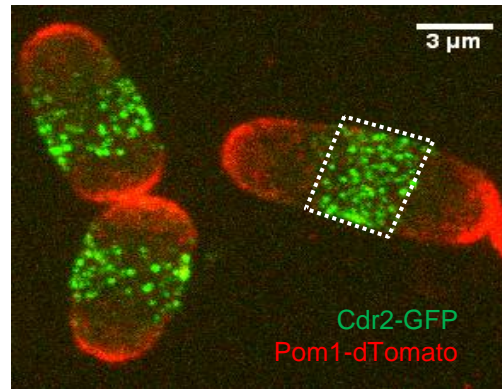
Pom1p gradient as a cell-size sensor?

- Our data does **not** support this hypothesis
- Manipulating gradient length scale via different **fluorescent tags** does not perturb size control
- Role of **Pom1p** more consistent as **tip inhibitor**



What is sensing cell size?!

- Could it be related to medial accumulation of Cdr2p?
- Medial density of Cdr2p, rises with cell length in cell population and in individual cells



Conclusions

- “Dissection” of Pom1p gradient
 - more complex mechanism of gradient formation!
 - could clustering be a general feature?
- Mechanism of cell size control
 - surprisingly does not centrally involve Pom1p
 - instead involves Cdr2p accumulation in medial zone of fixed area