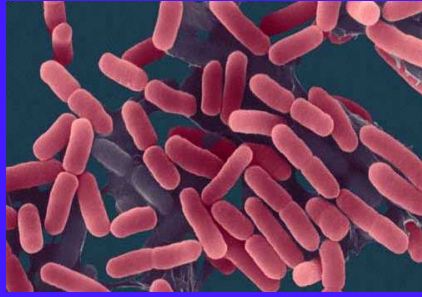


Accurate Division in *E. coli*



E. Coli [www.dontkneel.com]

Andrew Rutenberg
Simon deVet

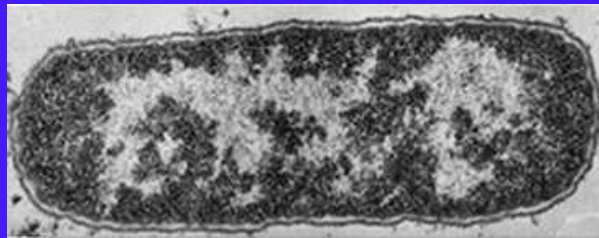
Department of Physics
Dalhousie University

Martin Howard

Department of Mathematics
Imperial College

Length and Time-scales

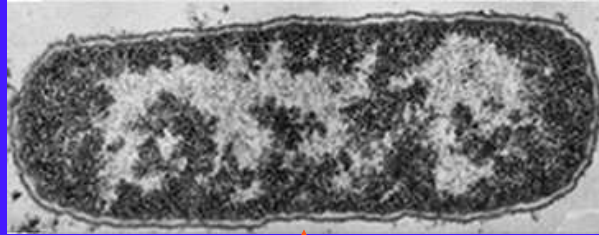
- **small rod** (2-4 μm in length, 1-1.5 μm in diameter)
- **divides accurately every hour**



E. coli [Albani et al]

Length and Time-scales

- small rod (2-4 μm in length, 1-1.5 μm in diameter)
- divides accurately every hour



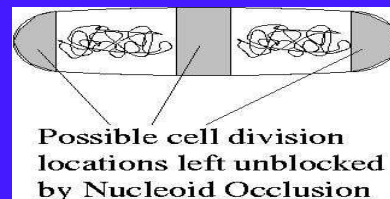
E. coli [Alberts *et al.*]

↑
WHY HERE?

Qualitative Model

“Nucleoid Occlusion”

- midcell division
- minicell mutants
- no guillotining

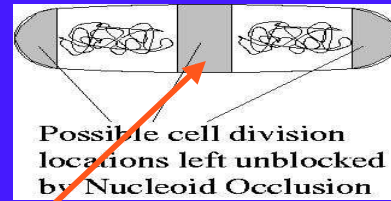


[Res. Microbiol. **141**, p39 (1990).]

Qualitative Model

“Nucleoid Occlusion”

- midcell division
- minicell mutants
- no guillotining



[Res. Microbiol. 141, p39 (1990).]

How is midcell selected?

Experimental interactions:

MinC

- inhibits division
- recruited to membrane by MinD

MinE

- recruited by MinD
- releases MinD

MinD

- binds to membrane
- recruits **MinE**
- released by **MinE**

[labs of Larry Rothfield, Joe Lutkenhaus, Piet de Boer, and others]

Phenomenology:

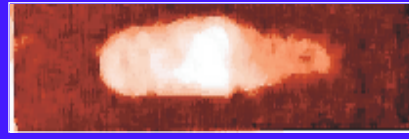
MinC/MinD

- suppress division
- mostly at cell poles
- ~1600 copies per cell



MinE

- suppresses MinD
- mostly at midcell
- ~1600 copies per cell



[Hale, Meinhardt, de Boer,
EMBO 20, p1563 (2001).]

Phenomenology:

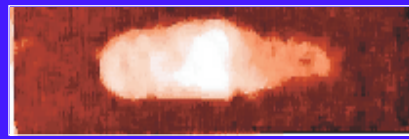
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MinE

- suppresses MinD
- mostly at midcell
- ~1600 copies per cell



[Hale, Meinhardt, de Boer,
EMBO 20, p1563 (2001).]

Min Model:

Membrane



Bulk

Bulk MinD $\frac{\partial D}{\partial t}$

Membrane MinD $\frac{\partial d}{\partial t}$

Bulk MinE $\frac{\partial E}{\partial t}$

Membrane MinE $\frac{\partial e}{\partial t}$

Min Model:

DIFFUSION • all proteins diffuse along the bacterium

• no diffusion when membrane associated ★

Bulk MinD $\frac{\partial D}{\partial t} = D_D \frac{\partial^2 D}{\partial x^2}$

Membrane MinD $\frac{\partial d}{\partial t} =$

Bulk MinE $\frac{\partial E}{\partial t} = D_E \frac{\partial^2 E}{\partial x^2}$

Membrane MinE $\frac{\partial e}{\partial t} =$

Min Model:

REACTION

- MinD goes onto membrane spontaneously, comes off due to **MinE**
- **MinE** goes onto membrane due to MinD, comes off spontaneously

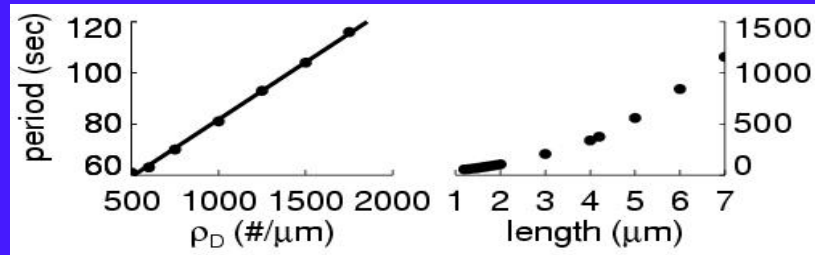
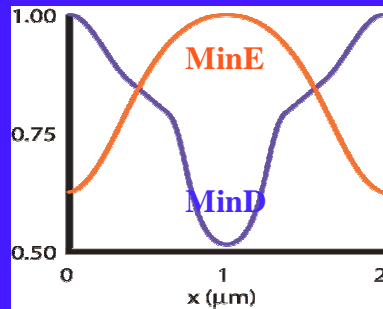
Bulk MinD $\frac{\partial \mathbf{D}}{\partial t} = D_D \frac{\partial^2 \mathbf{D}}{\partial x^2} - \frac{\sigma_1 \mathbf{D}}{1 + \sigma'_1 \mathbf{e}} + \sigma_2 \mathbf{e} \mathbf{d}$

Membrane MinD $\frac{\partial \mathbf{d}}{\partial t} = \dots + \dots - \dots$

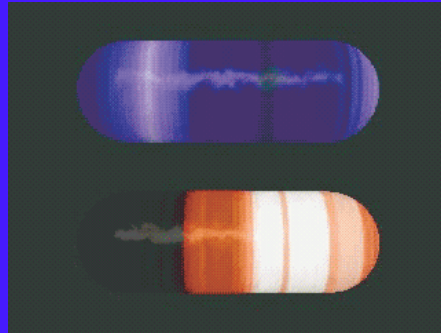
Bulk MinE $\frac{\partial \mathbf{E}}{\partial t} = D_E \frac{\partial^2 \mathbf{E}}{\partial x^2} - \sigma_3 \mathbf{E} \mathbf{D} + \frac{\sigma_4 \mathbf{e}}{1 + \sigma'_4 \mathbf{D}}$

Membrane MinE $\frac{\partial \mathbf{e}}{\partial t} = \dots + \dots - \dots$

Results:

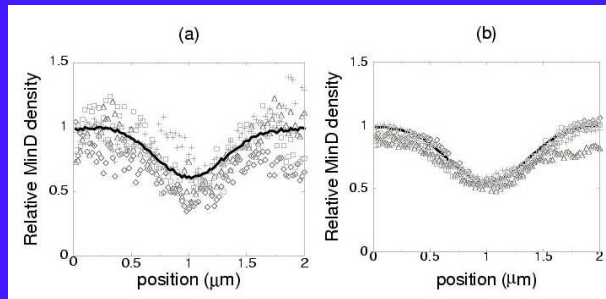


Shot-noise:

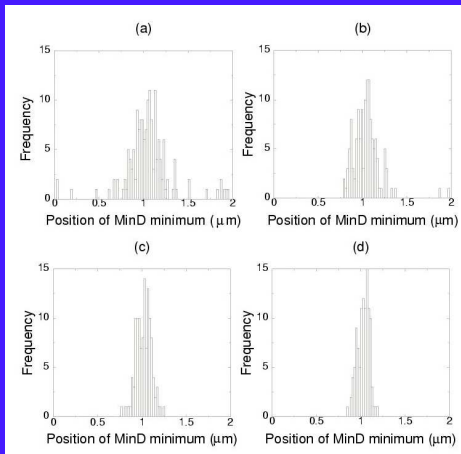


Number of MinD ~200,

number of MinD= 1500



Shot-noise:

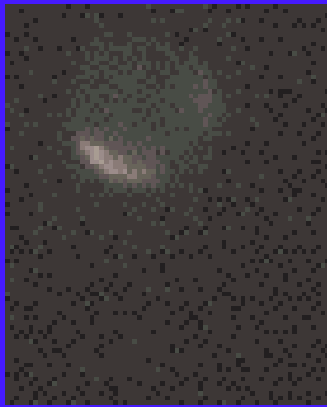


more MinD leads to more precise midpoint positioning in every oscillation.

[Howard and Rutenberg, PRL **90**, p128102 (2003).]

Number of MinD ~200, 400, 800, 1500
for (a), (b), (c), (d)

Division of round cells:



[Corbin, Yu, and Margolin]

GFP-MinD in round *E. coli*

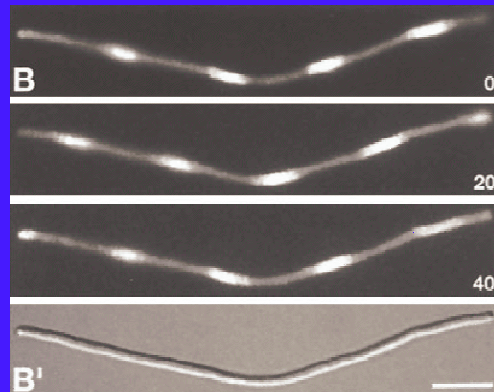
- oscillation axis wanders
- deformations lock axis

“Post-genomics”

- Quantitative model
- Dynamic compartmentalization:
spatial regulation without organelles
- division accuracy imposes expression levels
- *in vivo* biochemistry

Tuning the model

- frequency tunable via time and parameter rescaling
- wavelength tunable via density and parameter rescaling



gfp-MinD
[Raskin and de Boer,
PNAS 96, p4971 (1999).]

5micron scale

Min Models

<u>paper</u>	<u># parameters</u>
Meinhardt, deBoer, PNAS, 98 (2001).	17
Wingreen, Huang, Meir, [01/2003, KITP biomolecular networks]	9
Kruse, Biophys J, 82 (2002).	6
Howard, Rutenberg, deVet, PRL 87, (2001).	8

Linear Instability

- spontaneous oscillations
- exponential onset
- preliminary screening
- e.g.

$$\frac{\partial \mathbf{D}}{\partial t} = D_D \frac{\partial^2 \mathbf{D}}{\partial x^2} - \frac{\sigma_1 \mathbf{D}}{1 + \sigma_1' \mathbf{e}} + \sigma_2 \mathbf{e} \mathbf{d}$$

Filaments:

MinD

MinE

Decorated
or
Assembled?

[Shih, Le, Rothfield PNAS **100**, p7865 (2003).]