

Title: Monte Carlo Simulations of the Interaction Between a Self-Avoiding Polymer and a Membrane

Date: Apr 23, 2009 04:00 PM

URL: <http://pirsa.org/09040028>

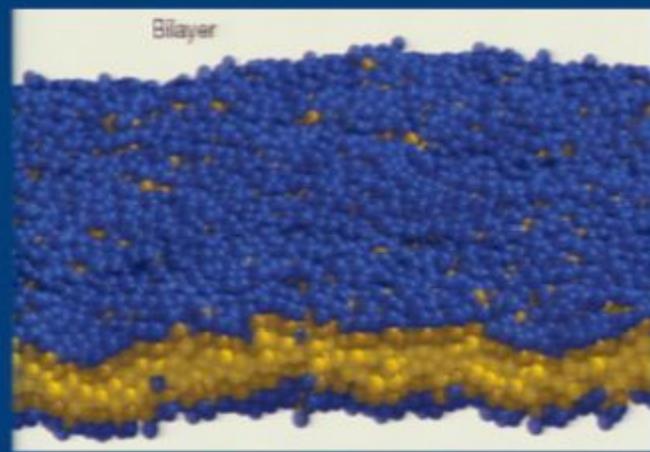
Abstract:

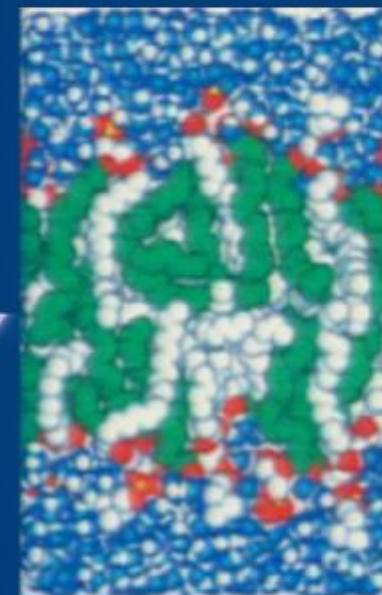
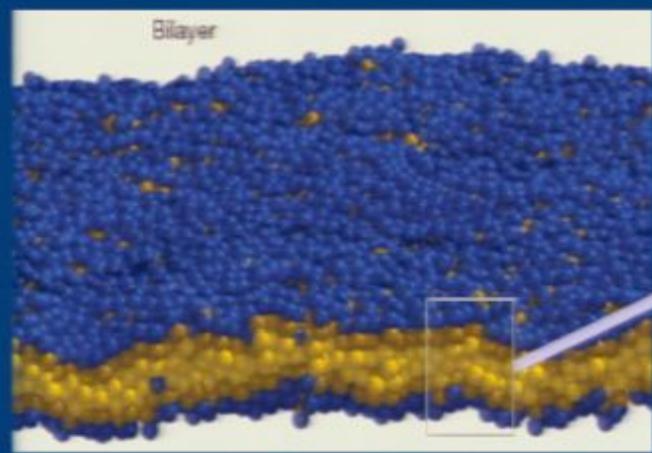
# **Structures of membranes interacting with macromolecules**

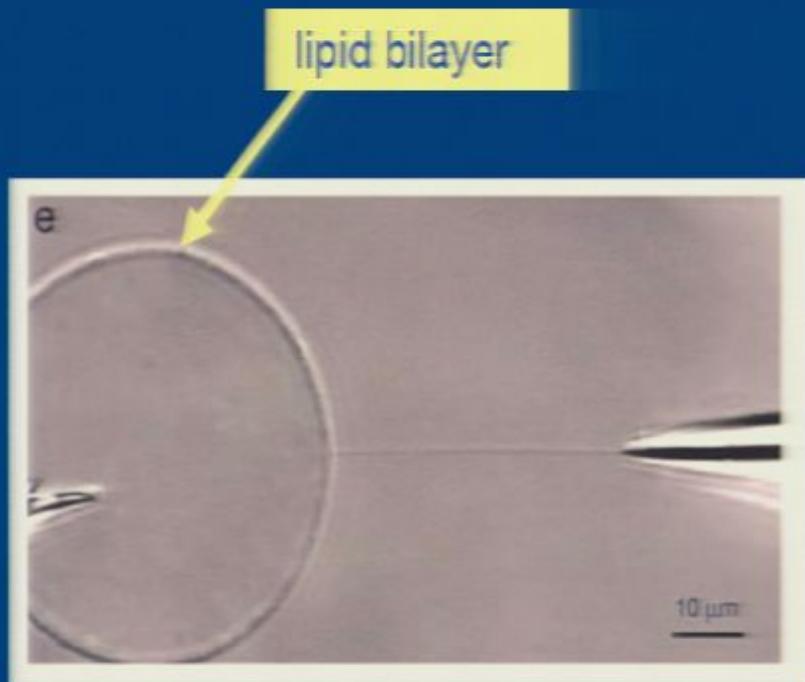
**Jeff Z. Y. Chen**

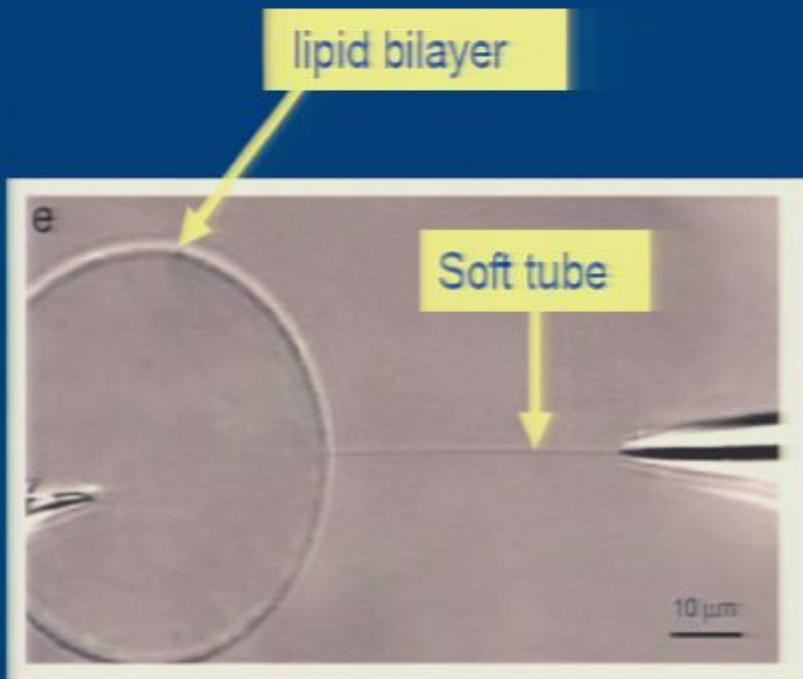
**Department of Physics & Astronomy  
University of Waterloo**

- Introduction to tubular membrane
- DNA confined in a soft tube
- Nanoparticles on a soft sheet



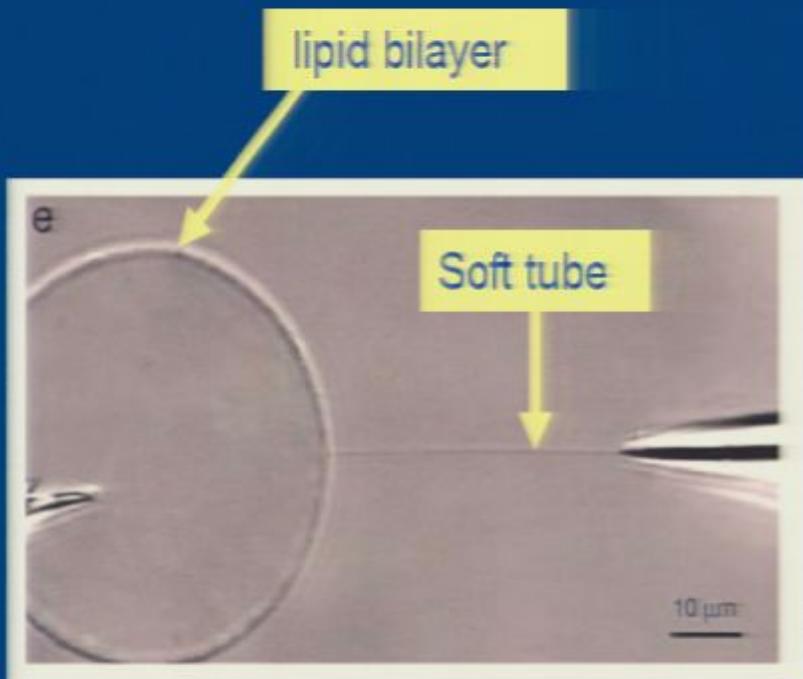






## Experiments: membrane tubes

- Borghi, Rossier and Brochard-Wyart, *Europhys Lett* **64**, 837 (2003)
- Tokarz, etc, *PNAS* **102**, 9127 (2005)
- Borghi, Kremer, Askovic and Brochard-Wyart, *Europhys Lett* **75**, 666 (2006)



## Experiments: membrane tubes

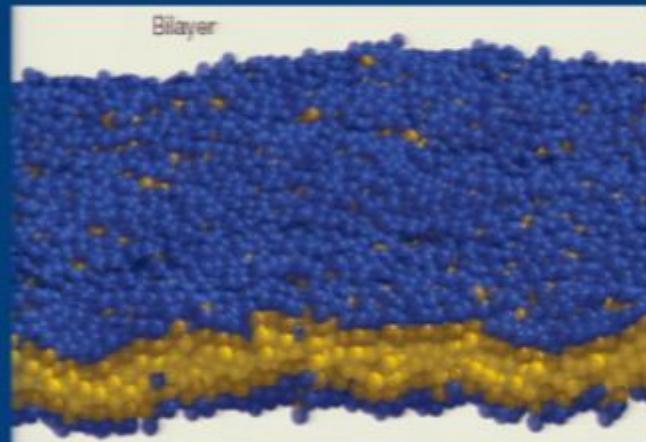
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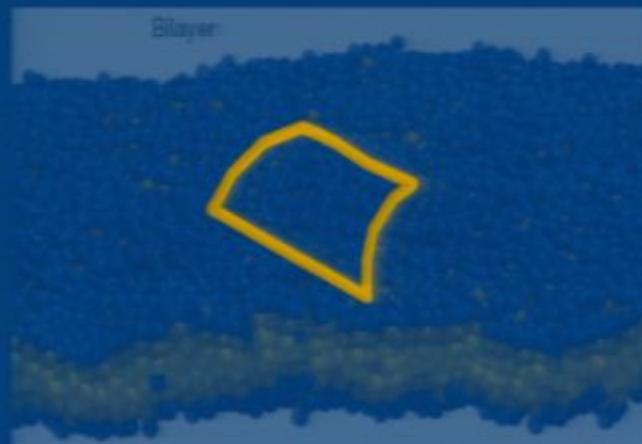
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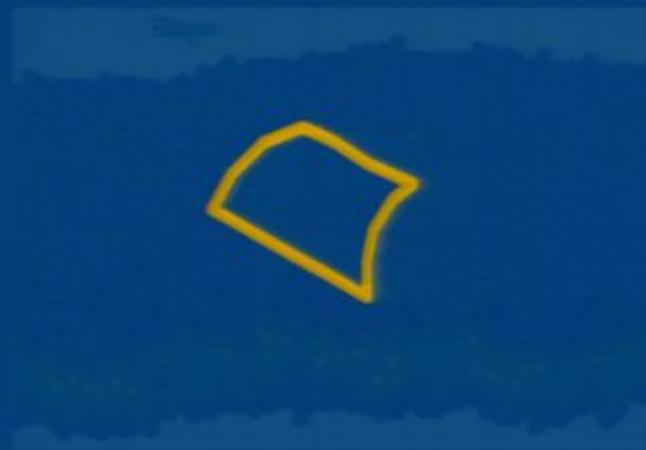
Energy model: **Helfrich model** for a fluid membrane



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■ Geometrical:

area:  $\Delta A$

## Energy model: Helfrich model for a fluid membrane



### ■ Geometrical:

area:  $\Delta A$

principal curvatures:  $1/r_1$  and  $1/r_2$

### ■ Physical parameters:

surface tension:  $\sigma$

bending energy:  $\kappa$

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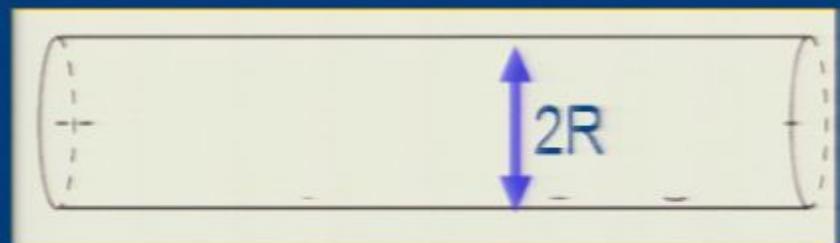
$$\text{Helfrich energy} = \Delta A \left[ \sigma + (\kappa/2) (1/r_1 + 1/r_2)^2 \right]$$

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- $\beta\kappa = 20$
- $\sigma a^2/\kappa: 10^{-3}$  to  $10^3$

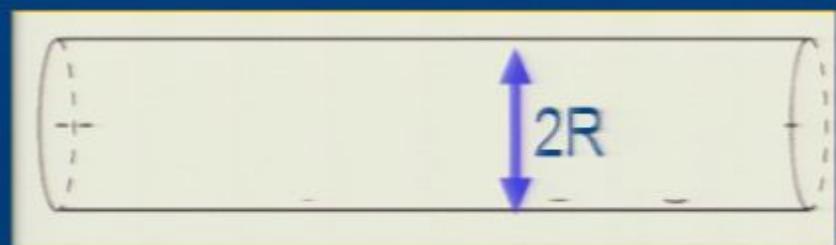
### Energy model: Helfrich model for a perfect cylinder

$$\text{Helfrich energy} = \Delta A [ \sigma + (\kappa/2) (1/r_1 + 1/r_2)^2 ]$$



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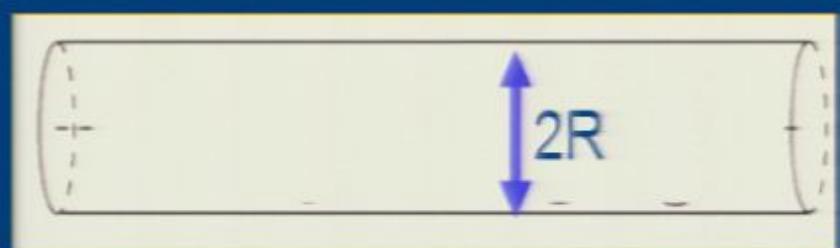
$$1/r_1 = 1/R$$

$$1/r_2 = 0$$

$$E = 2\pi LR [\sigma + \kappa/(2R^2)]$$

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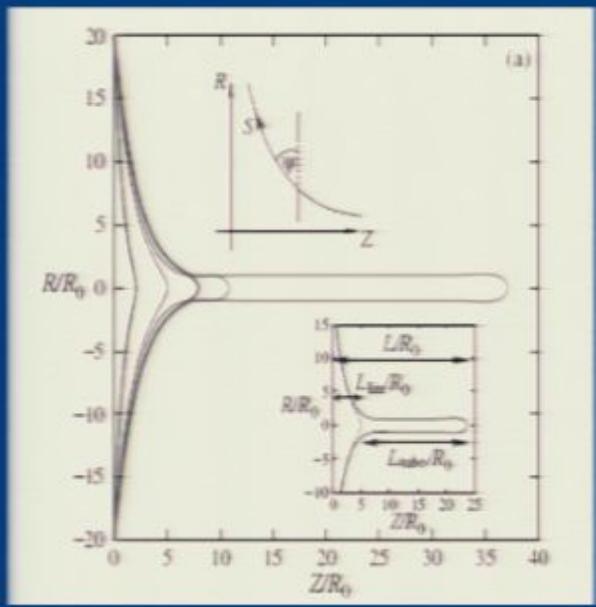
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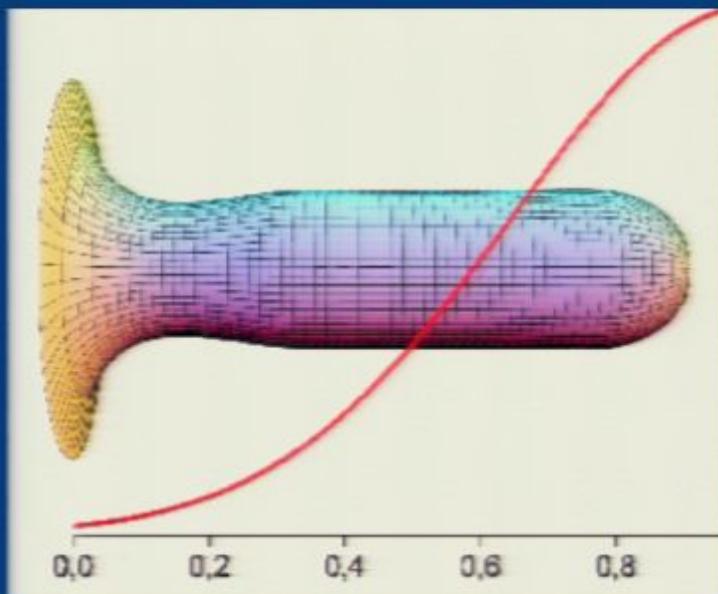
$$E = 2\pi LR [\sigma + \kappa/(2R^2)]$$

Minimization:  $dE/dR = 0$

$$\text{Equilibrium: } R_0^2 = \kappa/2\sigma$$



Derenyi, Julicher and Prost,  
PRL 88, 238101 (2002)



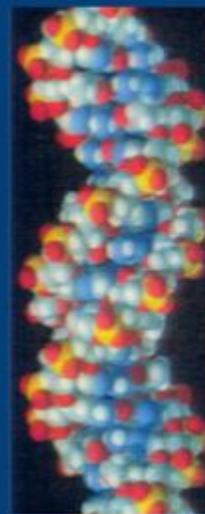
Campelo and Hernandez-Machado, PRL, 100,  
158103(2008)

# **TOPIC 1: Swollen-to-globular transition of a self-avoiding polymer confined in a soft tube**

## DNA and polymer models



## DNA and polymer models



Self-avoiding  
polymer



## DNA and polymer models

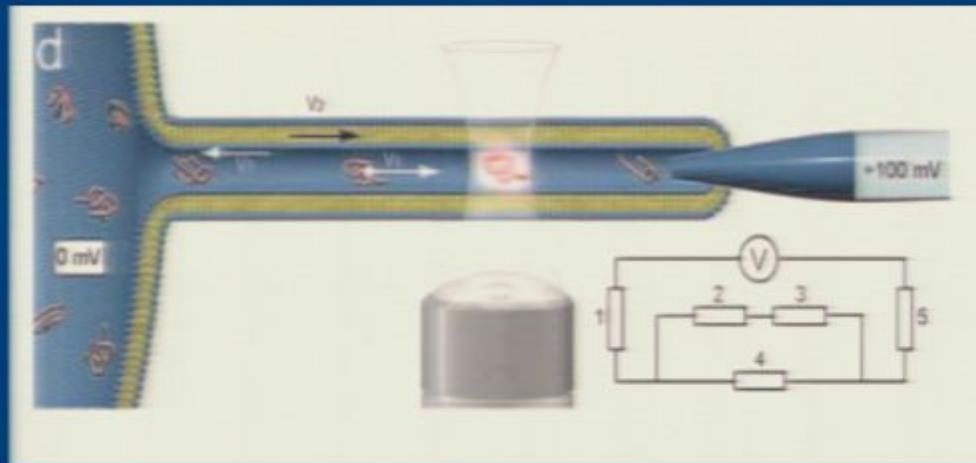
Self-avoiding  
polymer



“DNA physics”

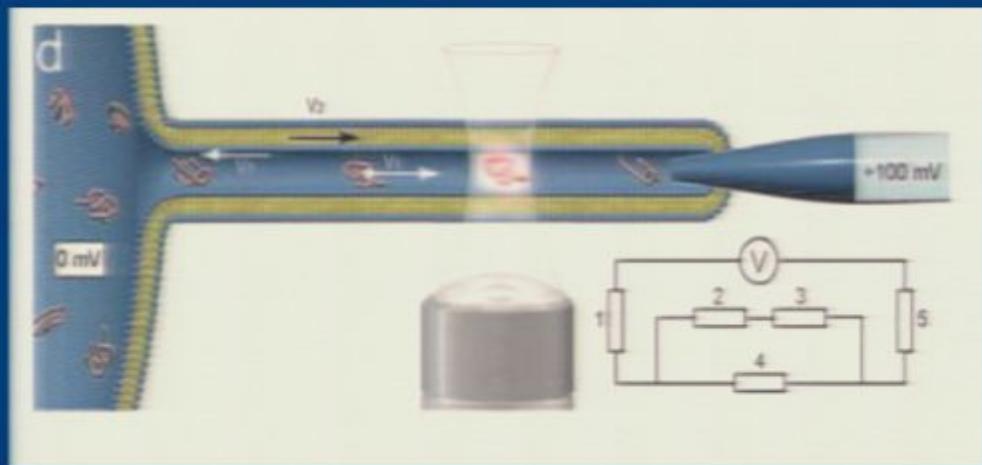
Polymer physics

## DNA in a soft tube



Michał Tokarz, Björn  
Akerman, Jessica Olofsson,  
Jean-François Joanny, Paul  
Dommersnes, and Owe  
Orwar, PNAS 102, 9127  
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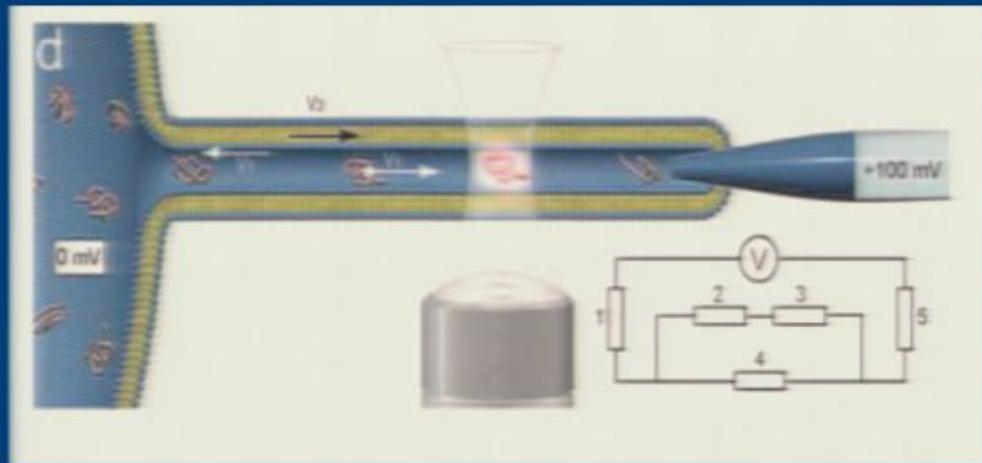
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- case 1: fluorescence light intensity = constant
- case 2: fluorescence light intensity = DNA length

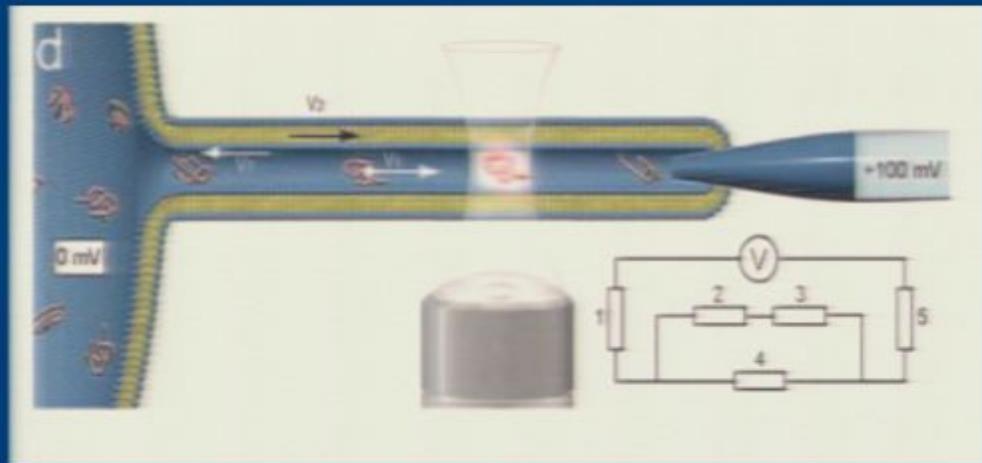
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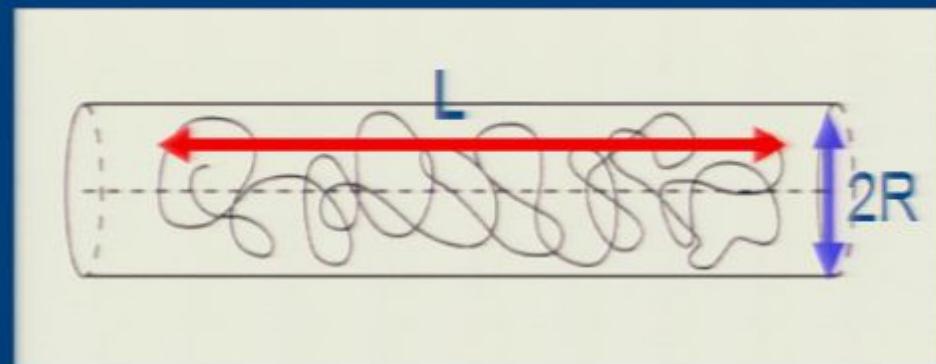


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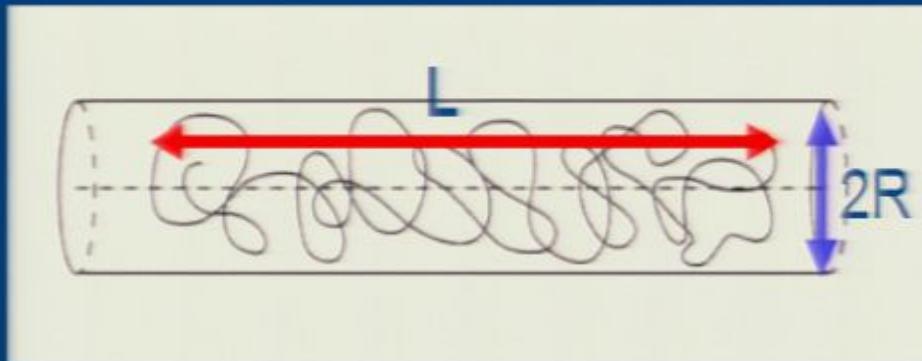
A self-avoiding polymer confined in a soft tube

## Polymer confined in a hard tube



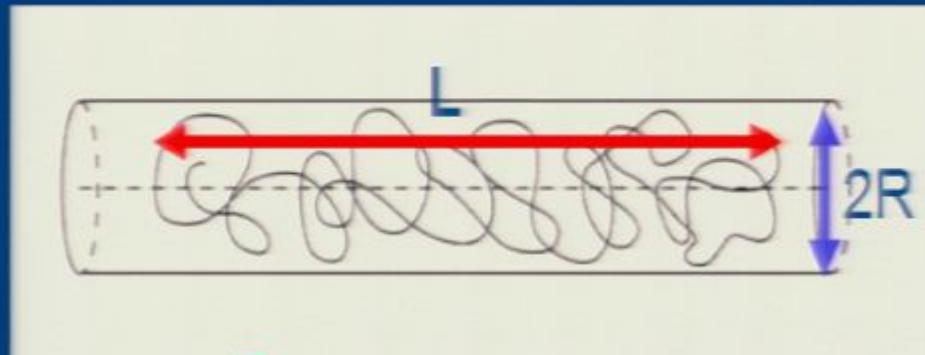
## Polymer confined in a hard tube

Total polymer length  $N$



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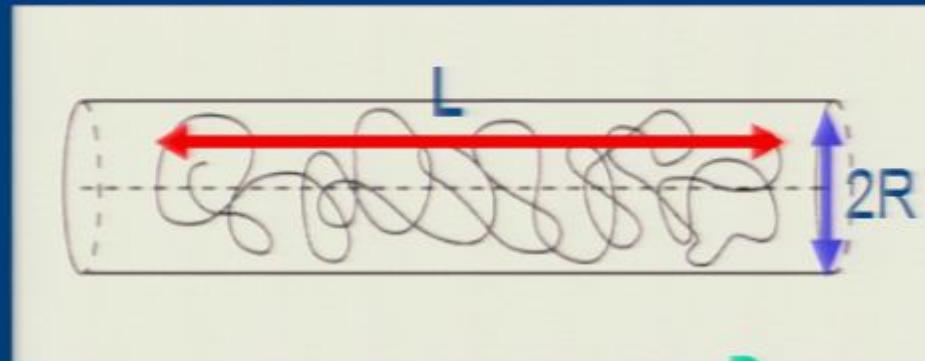
$$L = N R^{-2/3}$$

$D$



## Polymer confined in a hard tube

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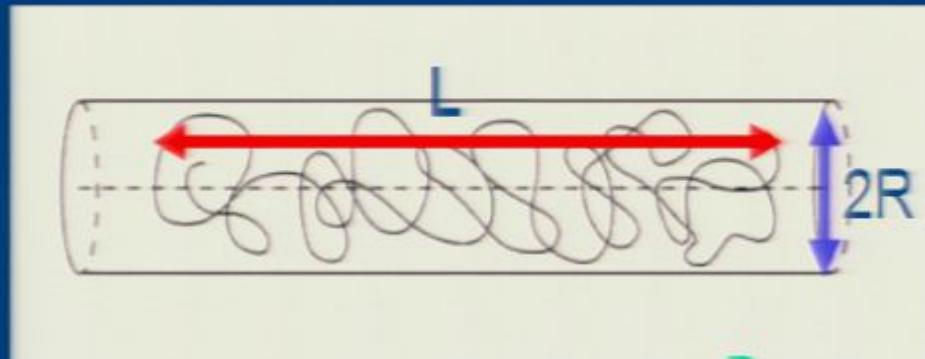
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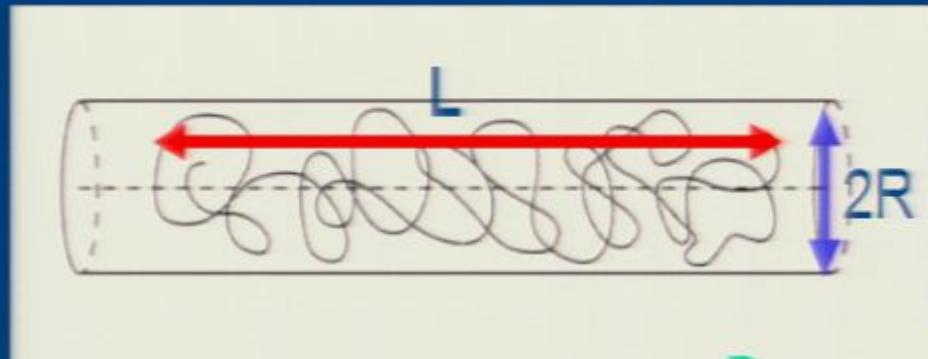
- Signal strength  $\sim D^* R^{**2} \sim [N/(L^* R^{**2})] = D^* R^{**(2/3)} = N\text{-independent, weak}$

$$D$$
  




## Polymer confined in a hard tube

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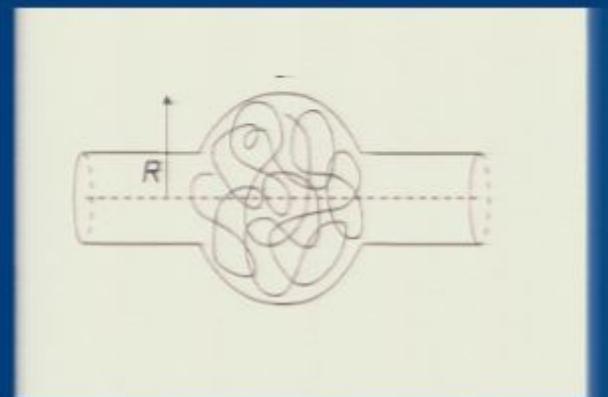


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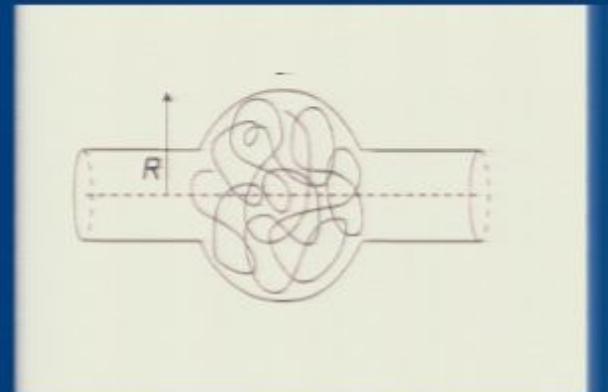
- Signal strength  $\sim D^* R^{**2} \sim [N/(L^* R^{**2})] = D^* R^{**(2/3)} = N\text{-independent, weak}$
- However, signal strength  $\sim N$  experimentally in some cases.



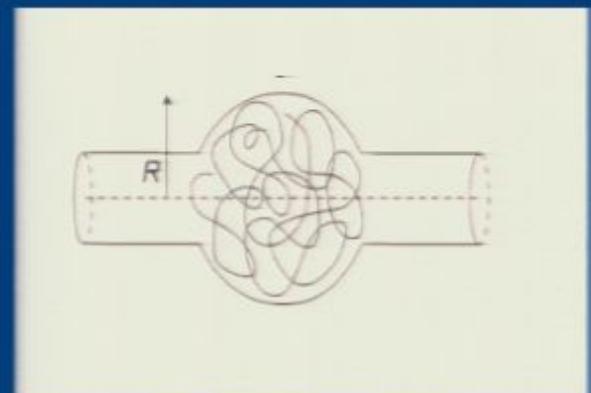
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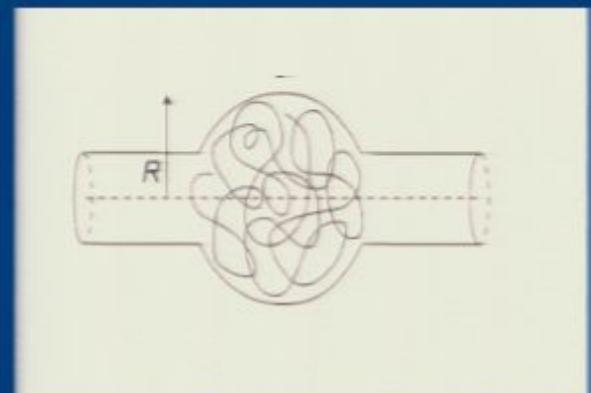


Under what circumstance would signal strength  $\sim N$ ?



- DNA in a globular state  
--- detecting the entire DNA hence signal strength is proportional to  $N$ .

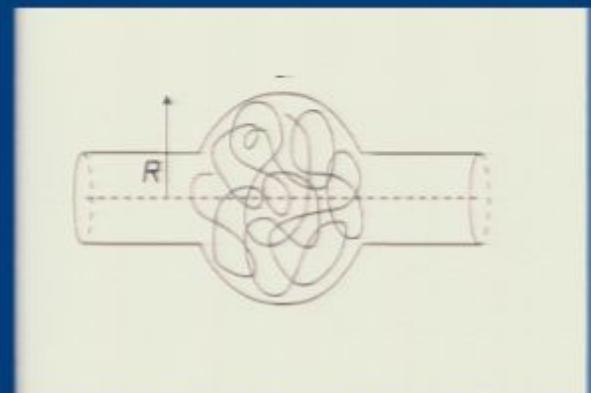
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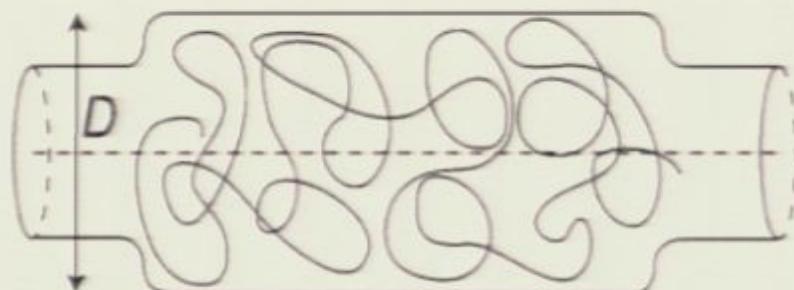


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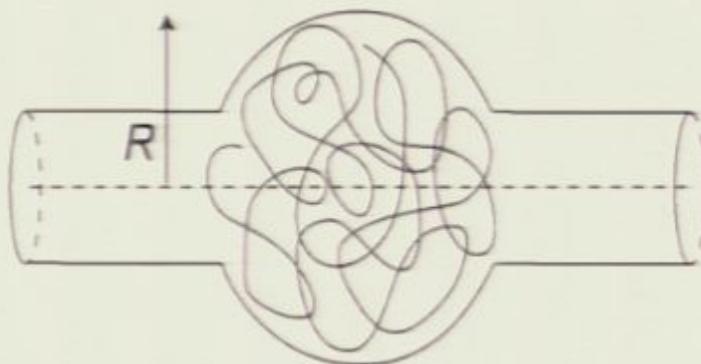
- The diameter of the soft tube is dilated in this regime.

- Theory and simulation:  
simultaneous consideration of polymer and soft tube.

## Swollen to globular transition

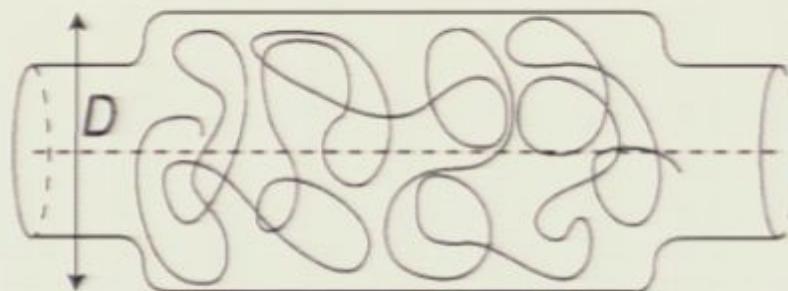


■ Swollen  
(elongated)

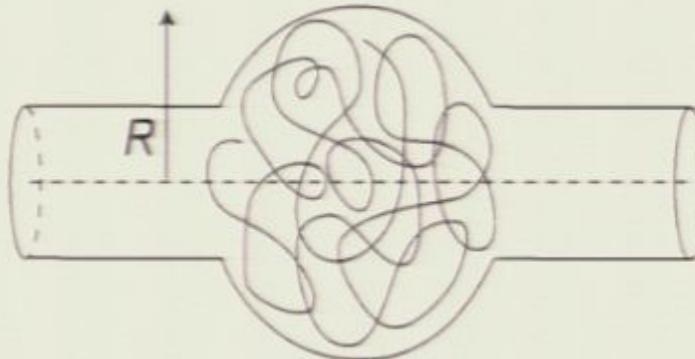


■ Globular

## Swollen to globular transition

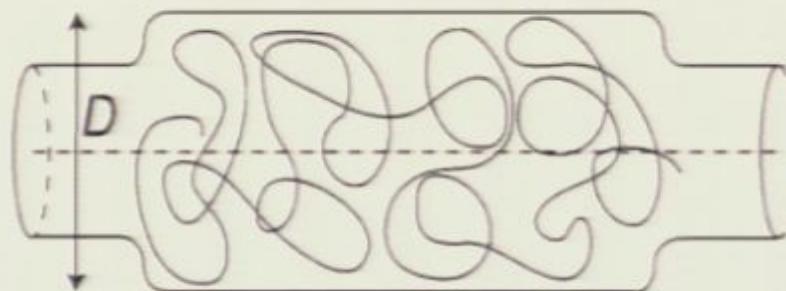


■ Swollen  
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"Snake eating a  
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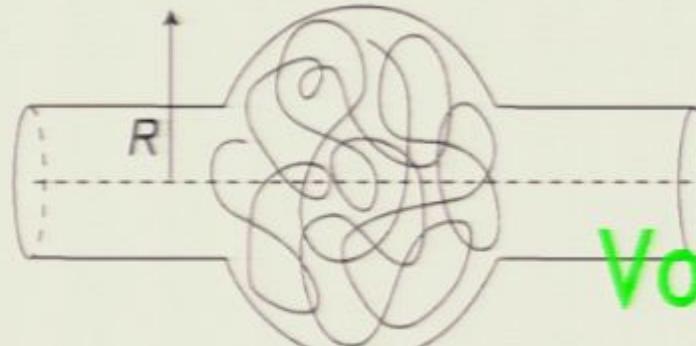


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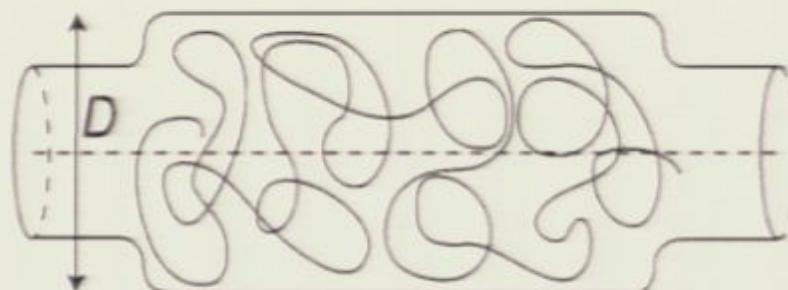


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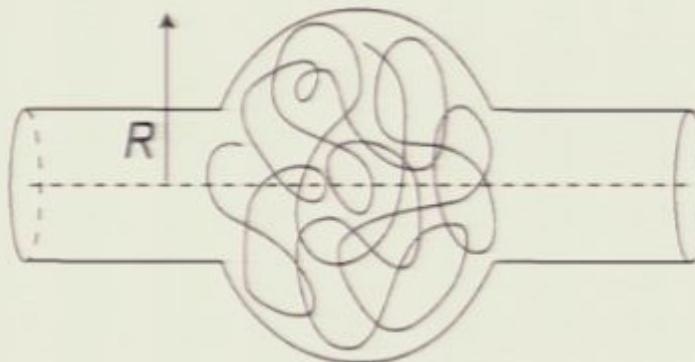
Volume



## Swollen to globular transition



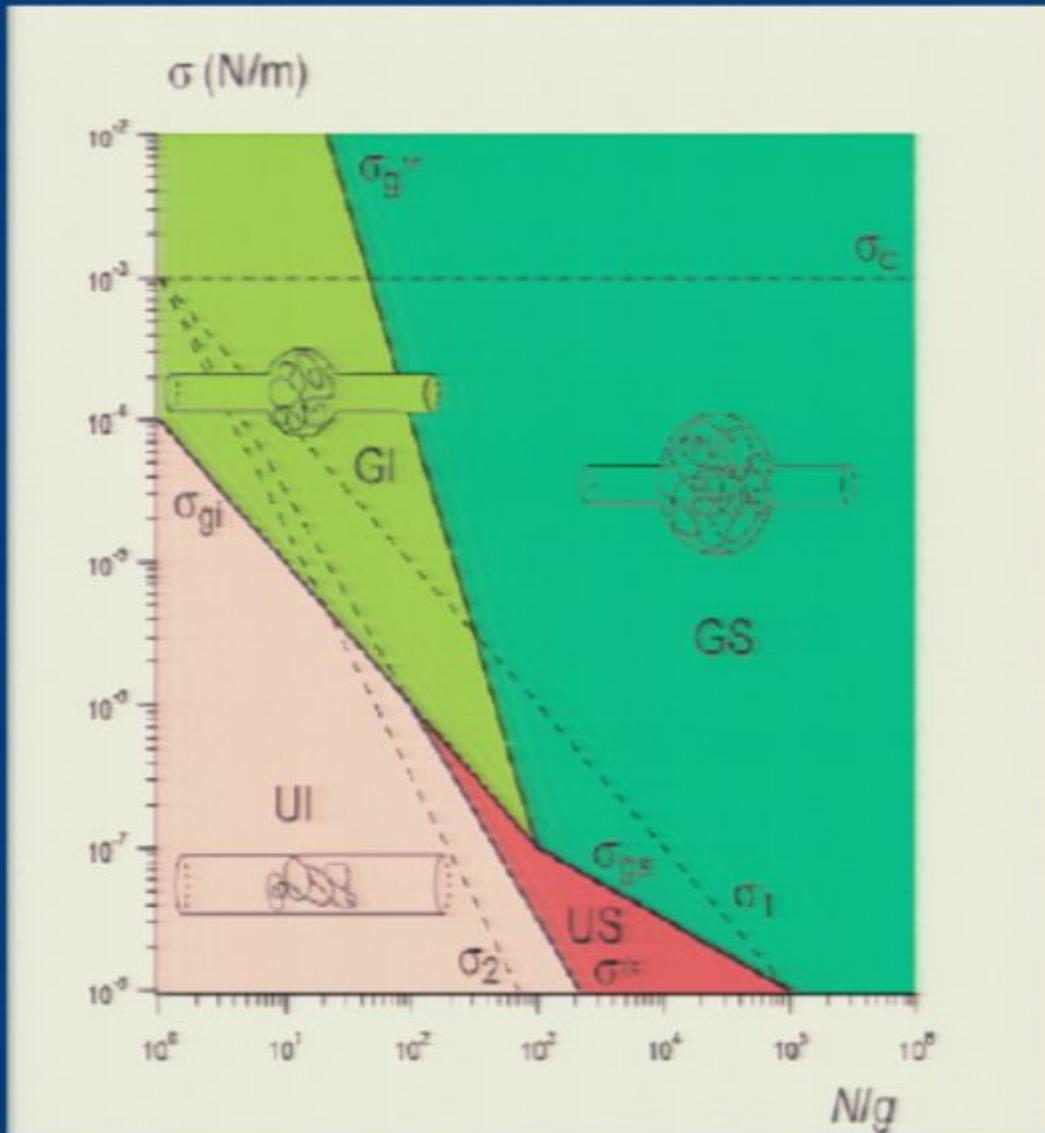
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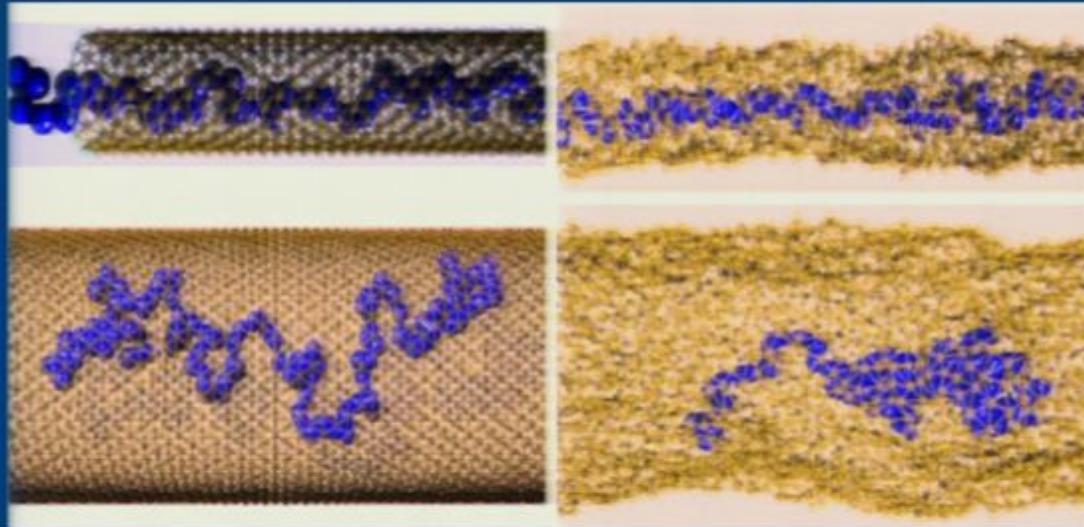
■ Globular

## Swollen to globular transition

by Brochard-Wyart, Tanaka, Borghi and de Gennes, Langmuir (2005)

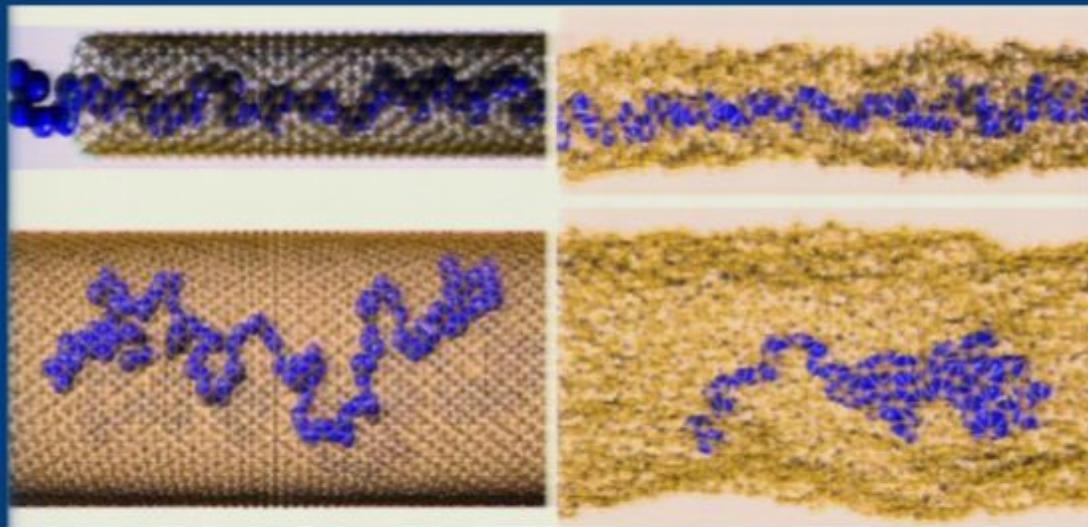


Computer simulation by Avramova and Milchev



Avramova and  
Milchev,  
*J. Chem. Phys.*  
**124**, 024909 (2006)

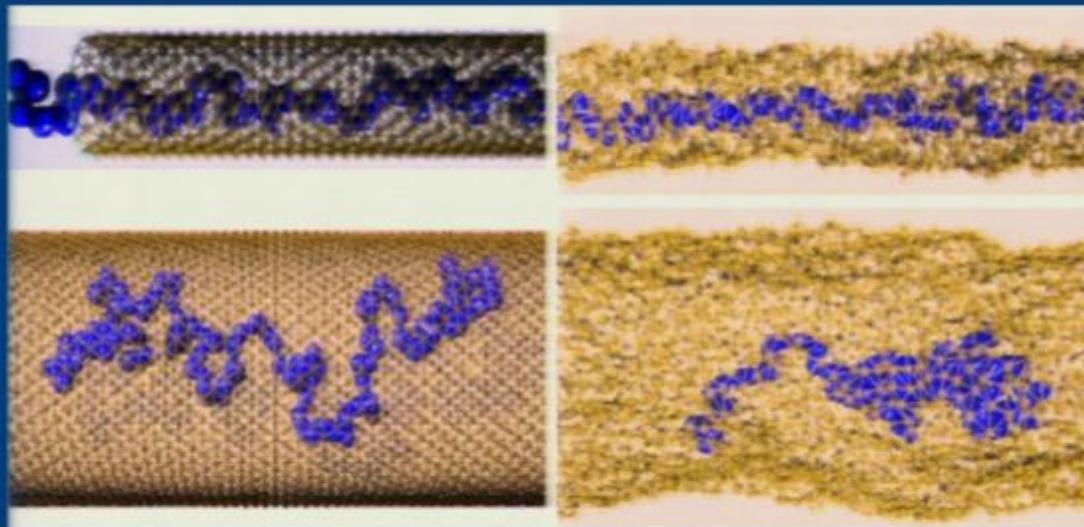
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- Tube must be modeled differently from the theory

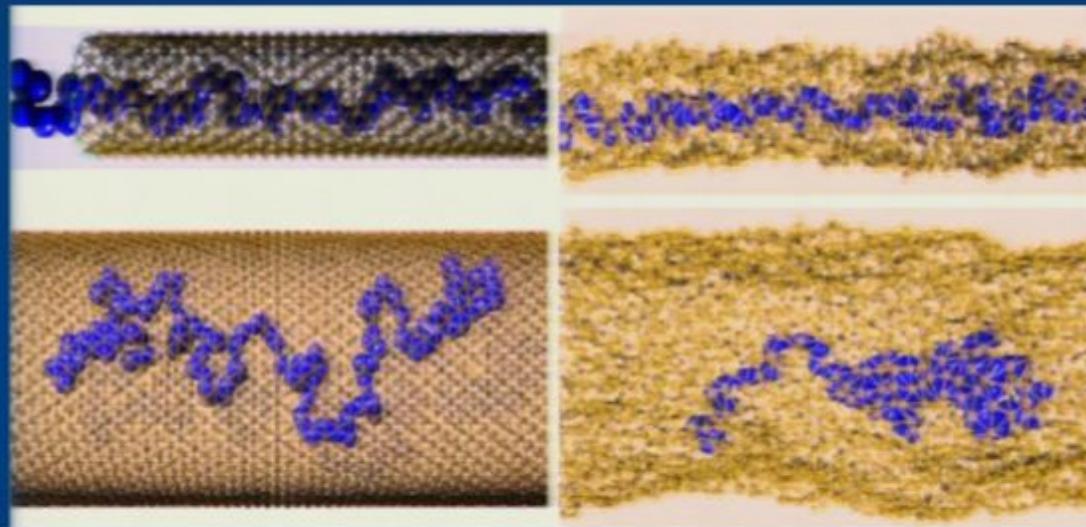
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- Tube must be modeled differently from the theory
- No observations of the structural transition

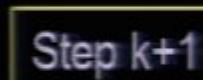
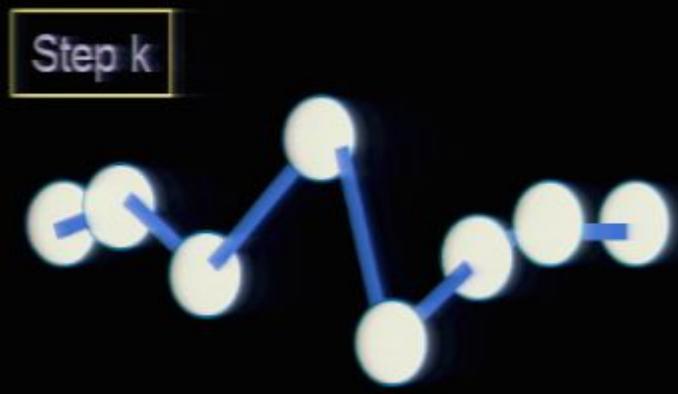
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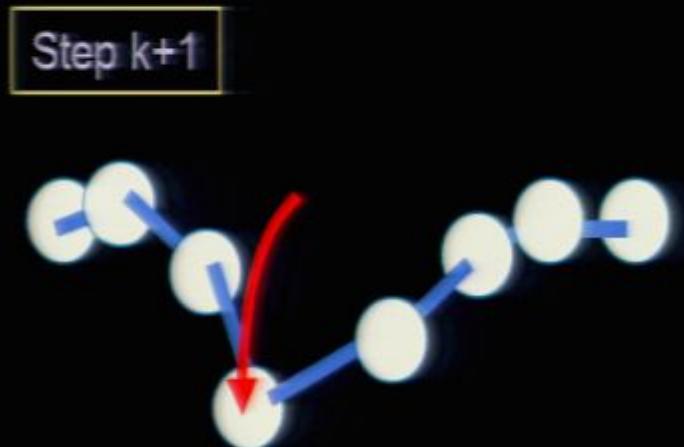
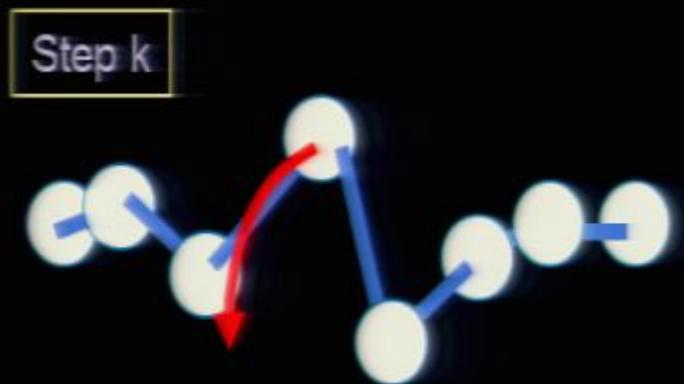
Avramova and  
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- Tube must be modeled differently from the theory
- No observations of the structural transition
- Expensive in modeling the tube (a M\*M problem)

## Monte Carlo Simulation of a self-avoiding chain

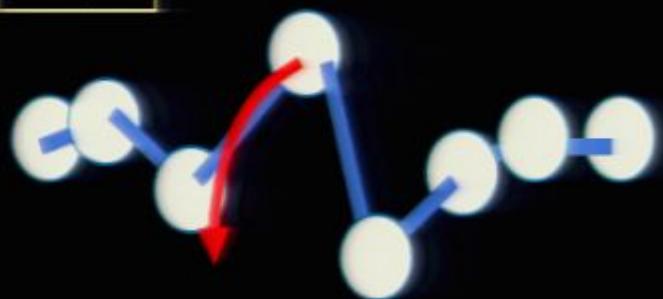


## Monte Carlo Simulation of a self-avoiding chain



## Monte Carlo Simulation of a self-avoiding chain

Step k

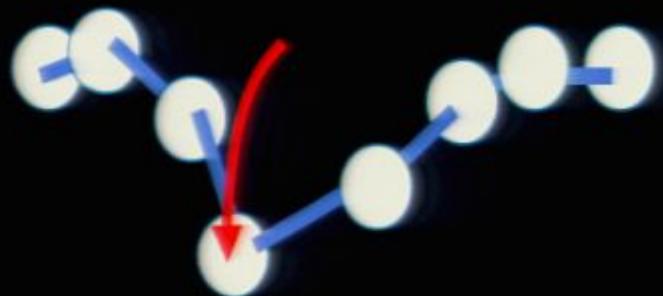


Basic parameters:

Bond length:  $a$

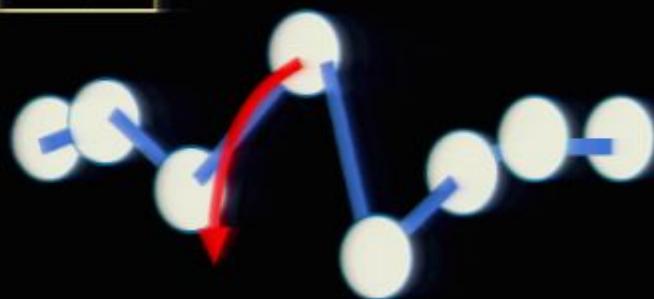
Total no. of

Step k+1



## Monte Carlo Simulation of a self-avoiding chain

Step k

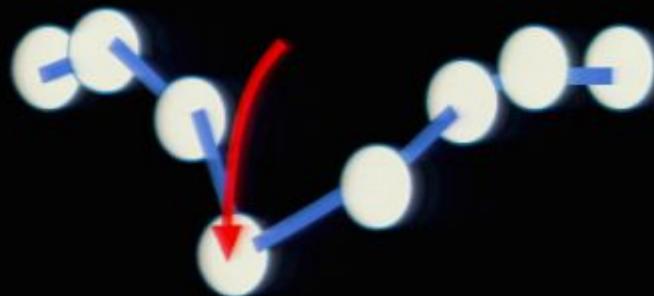


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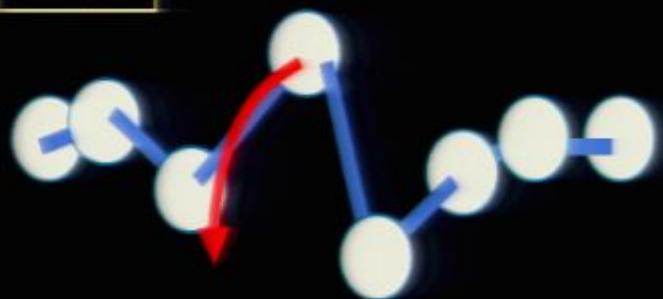
Total no. of monomers:  $N$

Step  $k+1$



## Monte Carlo Simulation of a self-avoiding chain

Step k

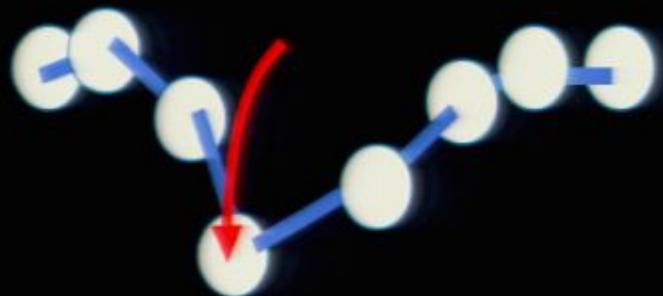


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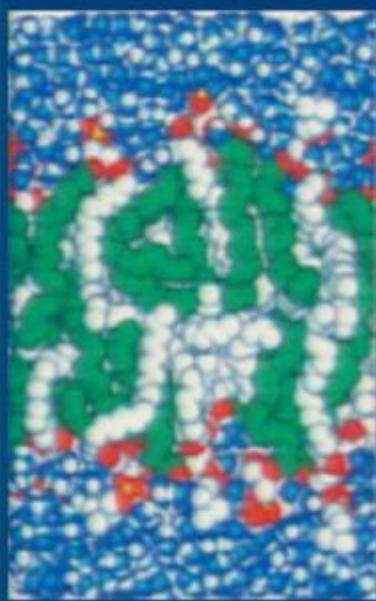


Excluded-volume diameter:  $D (=a)$

## Lipid bilayer models

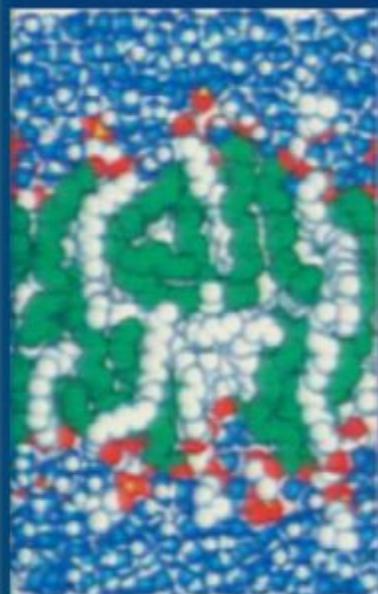
## Lipid bilayer models

### All-atom model

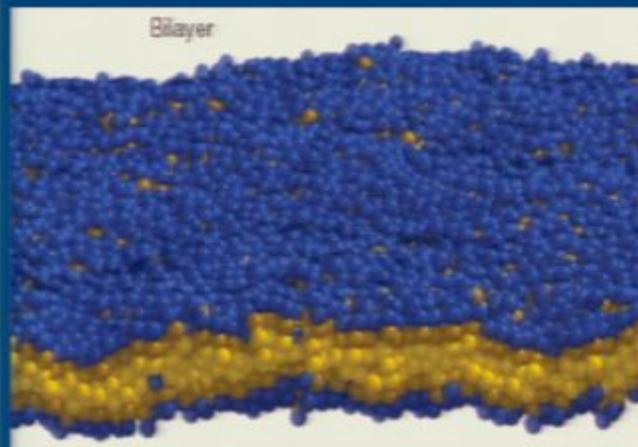


## Lipid bilayer models

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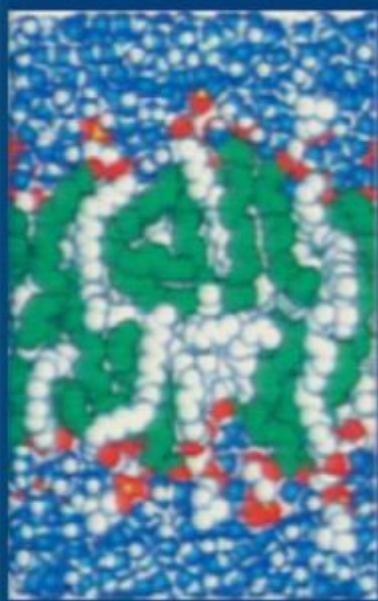


### Coarse grained

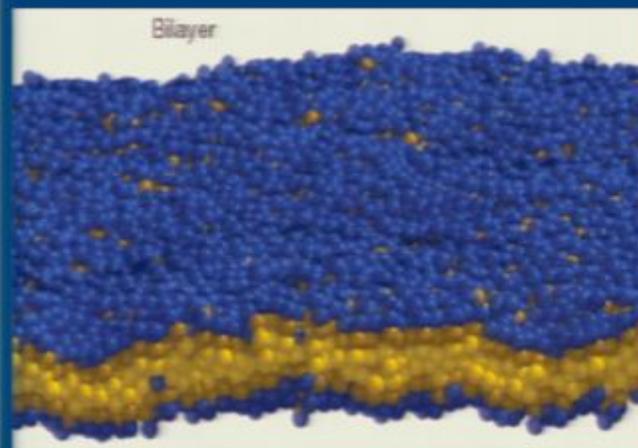


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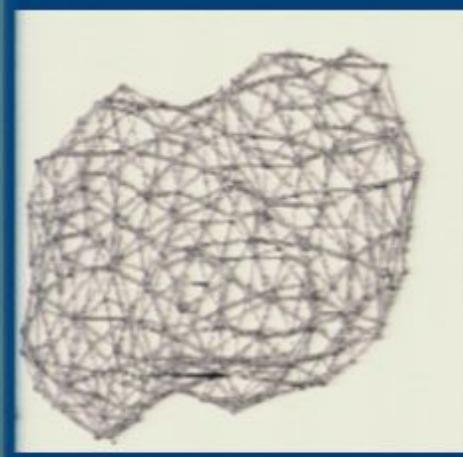
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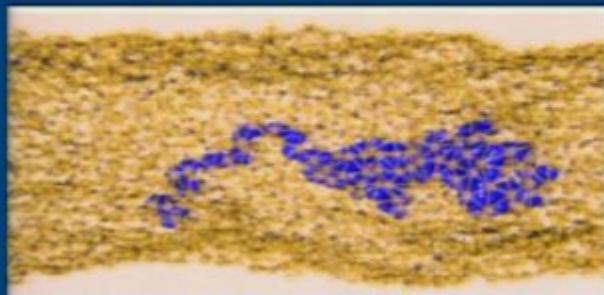
### Coarse grained



### Elastic sheet

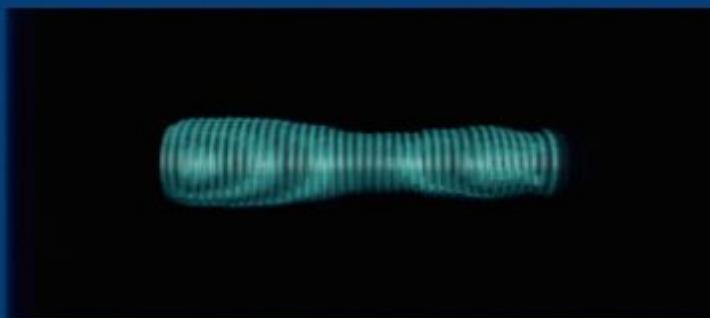


## Representation of a tube membrane



### Elastic sheet techniques

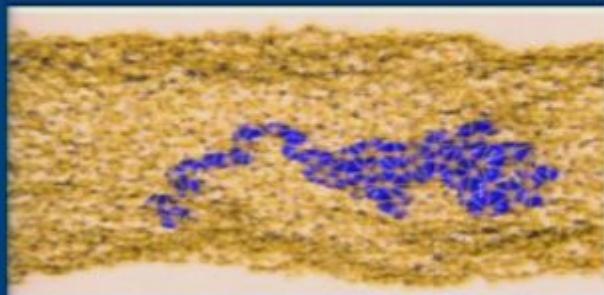
- $M^2$  nodes needed
- Complex to maintain a tube



### Our technique

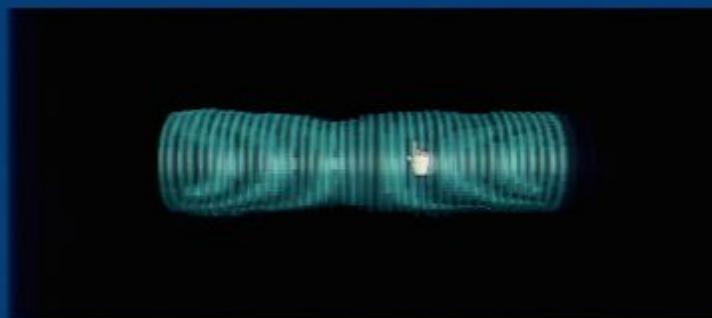
- Only  $M$  nodes needed
- Use Helfrich model directly

## Representation of a tube membrane



### Elastic sheet techniques

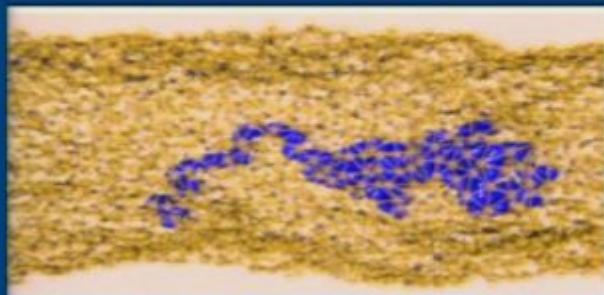
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### Our technique

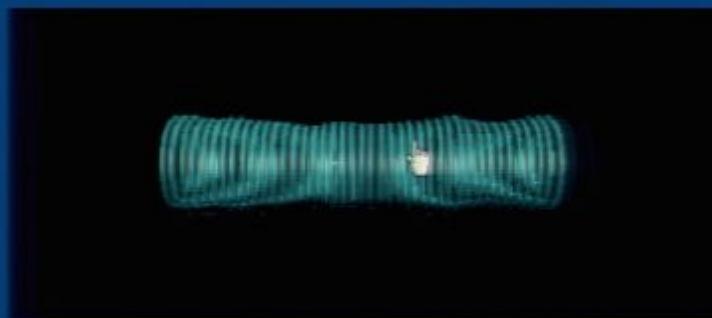
- Only  $M$  nodes needed
- Use Helfrich model directly

## Representation of a tube membrane



### Elastic sheet techniques

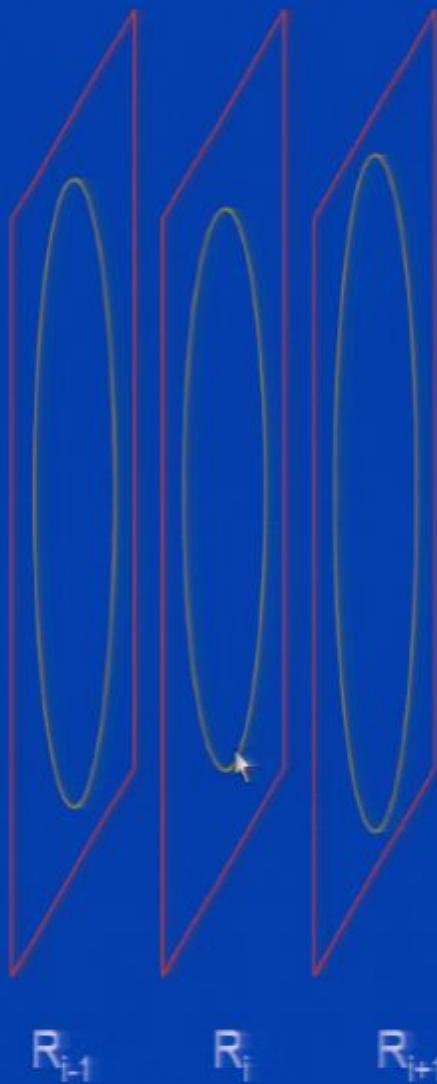
- $M^2$  nodes needed
- Complex to maintain a tube



### Our technique

- Only  $M$  nodes needed
- Use Helfrich model directly

## Helfrich energy for a soft tube



- Geometrical:

area:  $\Delta A(R_{i-1}, R_i, R_{i+1})$

inverse curvatures:  $r_1(R_{i-1}, R_i, R_{i+1})$   
 $r_2(R_{i-1}, R_i, R_{i+1})$

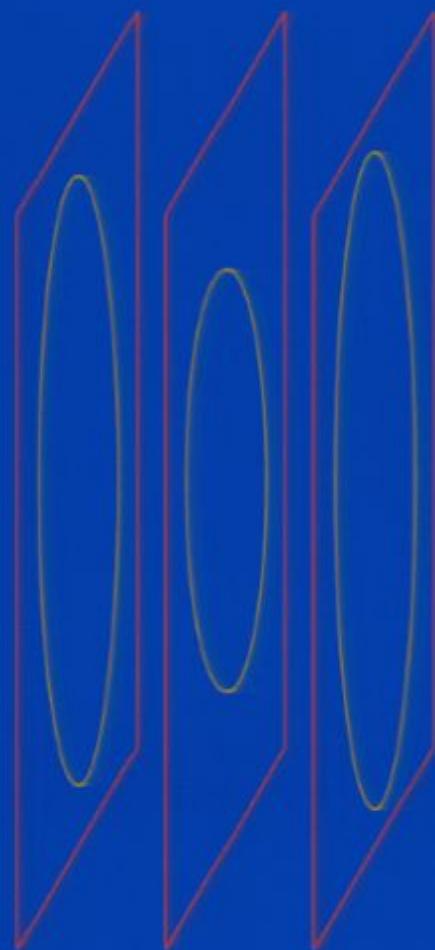
- $E_i$

$$= \Delta A [\sigma + (\kappa/2) (1/r_1 + 1/r_2)^2 ]$$

= function of  $R_{i-1}, R_i, R_{i+1}$  with two physical parameters  $\kappa$  and  $\sigma$

- $E = \sum_{i=1}^M E_i$

## Monte Carlo simulation for a tube membrane confining a polymer



- Select a random  $i$
- Move  $R_i^2$
- Calculate energy difference between the new and old systems
- Evaluate the acceptability of the new configuration using

$$P = \exp [ - \beta \Delta E ]$$

(Metropolis Monte Carlo)

- Move the polymer inside
- Go back to the first step

Always check to see if the polymer is confined inside

## Parameters in the model

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- $N, a$  (polymer)
- $\kappa, \sigma$  (membrane)

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## Reduced Parameters in the model

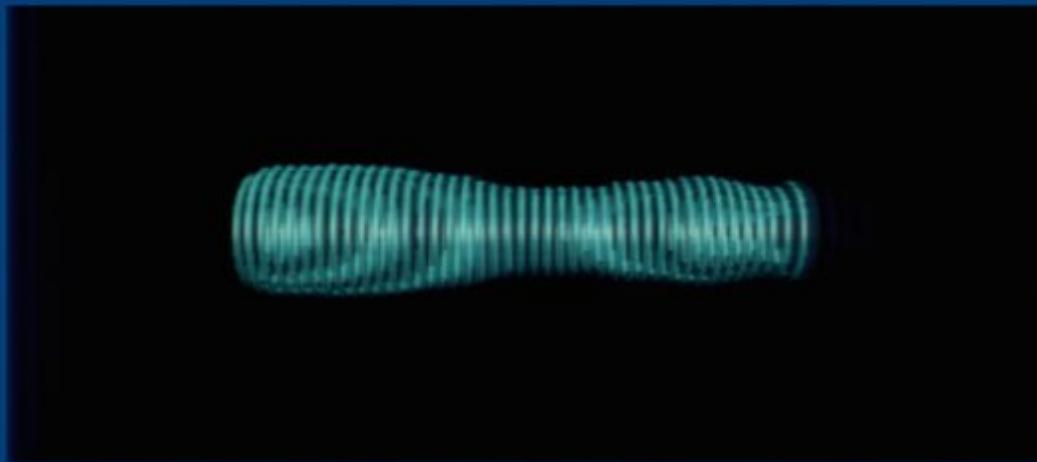
## Parameters in the model

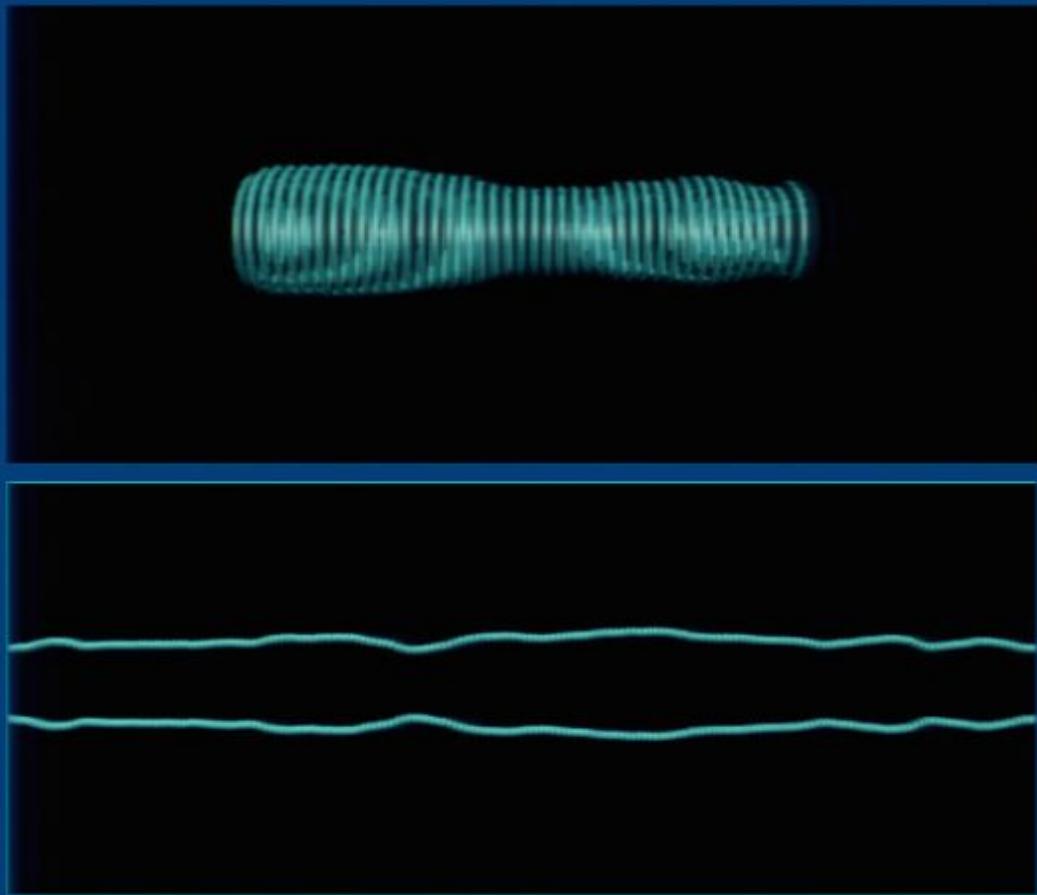
- $N, a$  (polymer)
- $\kappa, \sigma$  (membrane)
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## Reduced Parameters in the model

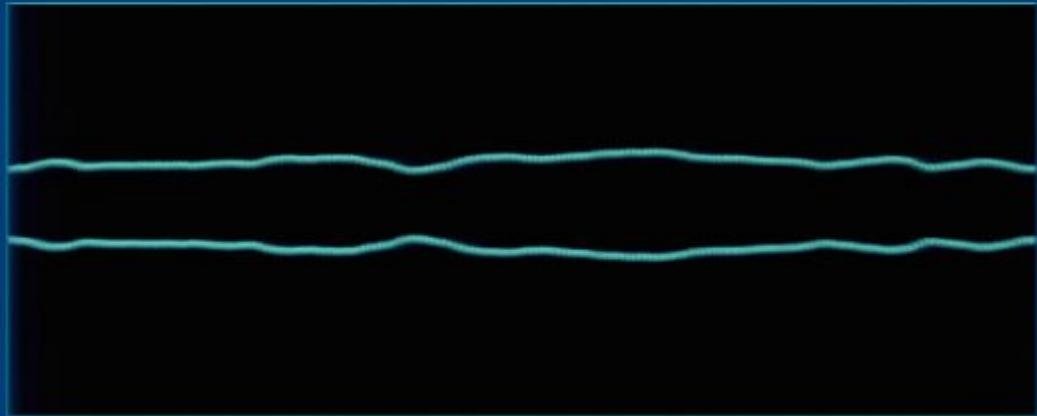
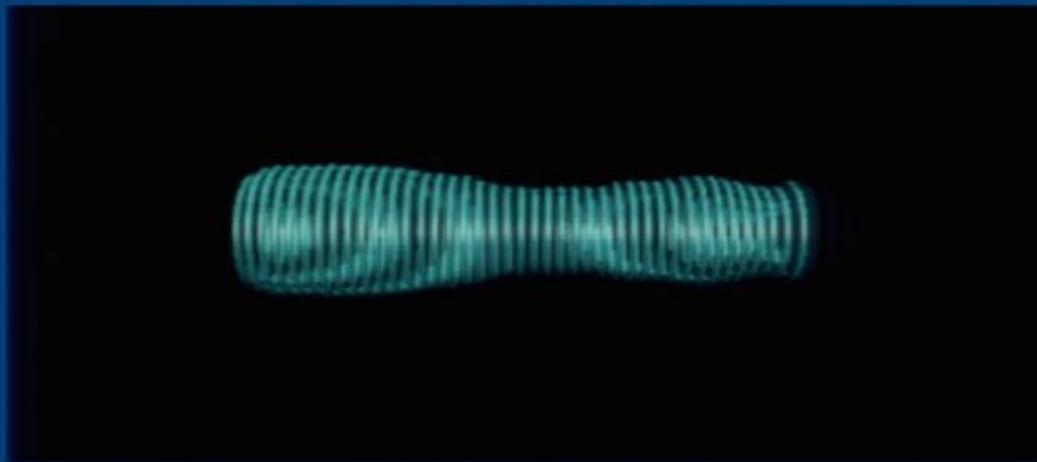
- $N$  (polymer)
- $\beta\sigma a^2$
- $\beta\kappa$

(we FIX  $\beta\kappa$  in the remainder of this talk)

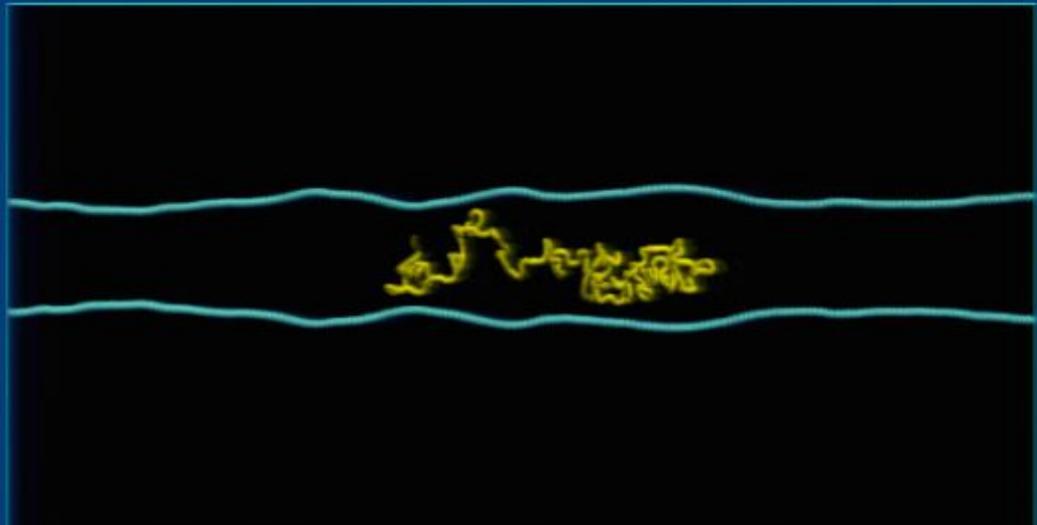




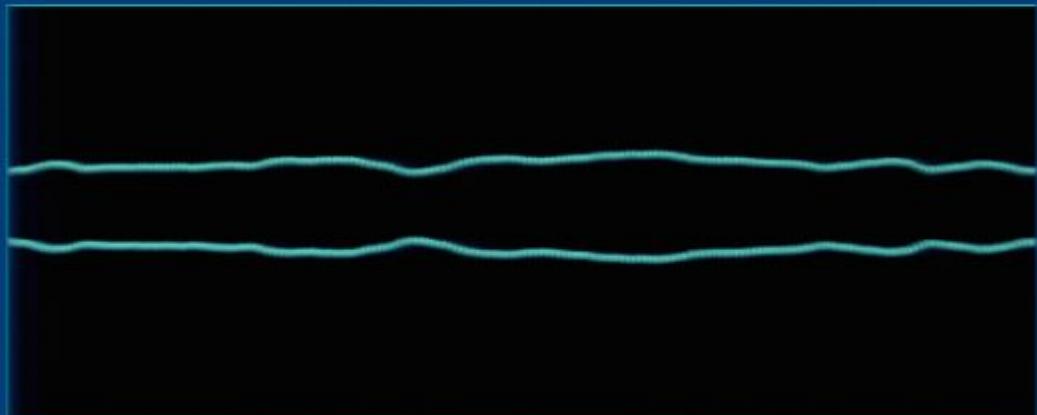
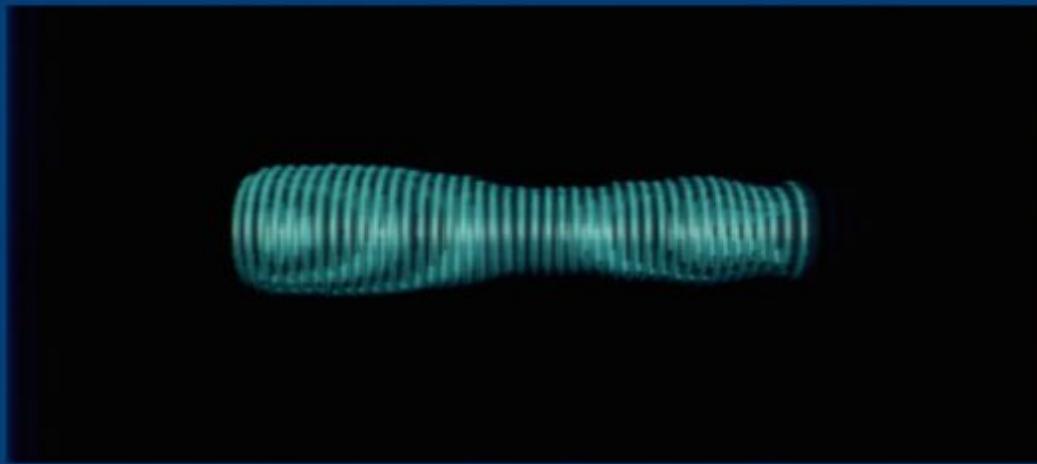
Equilibrium:  
 $R_0^2 = \kappa/2\sigma$



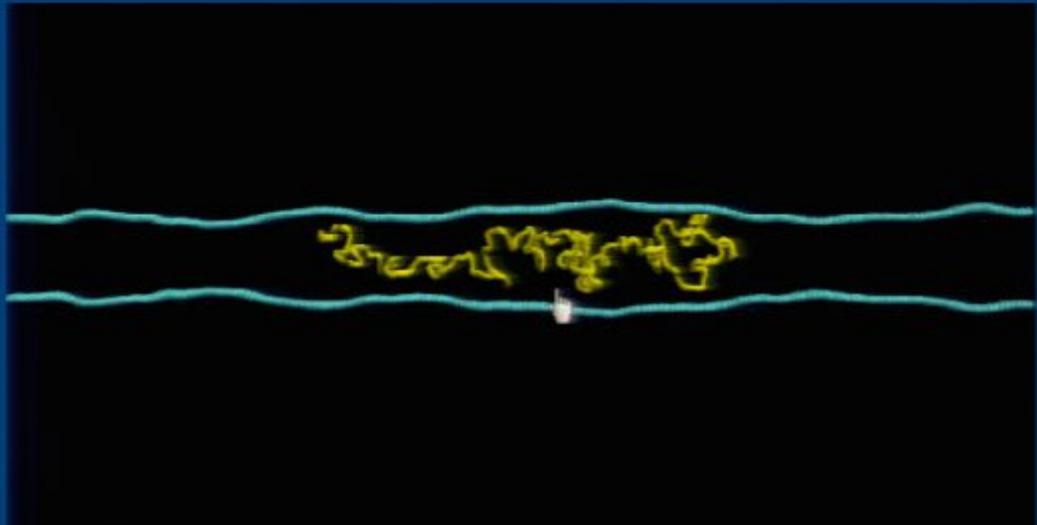
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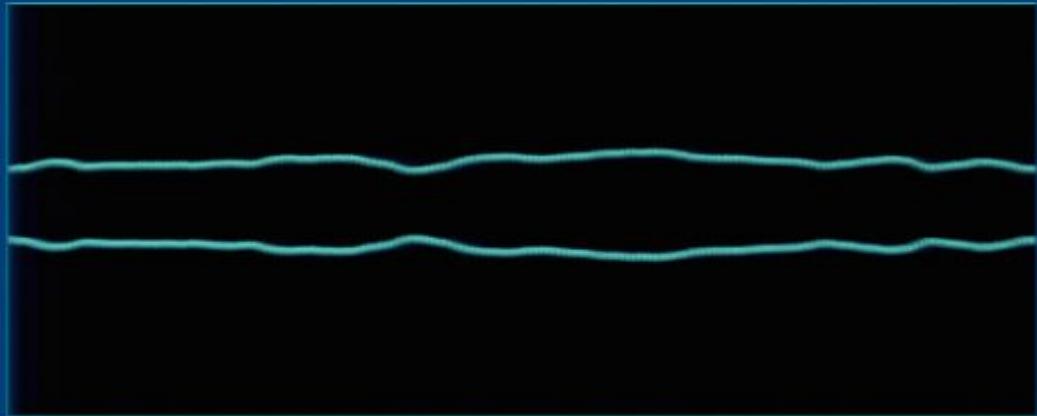
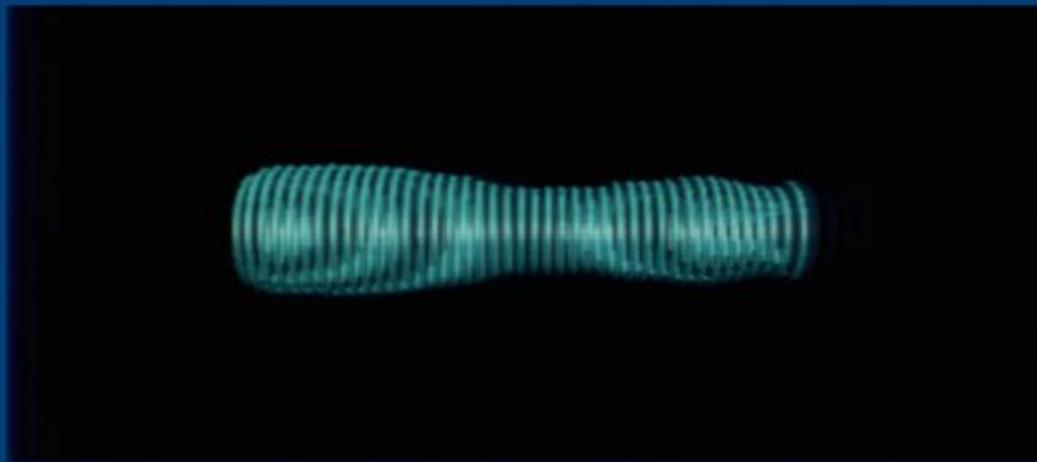
Increase  $\sigma$



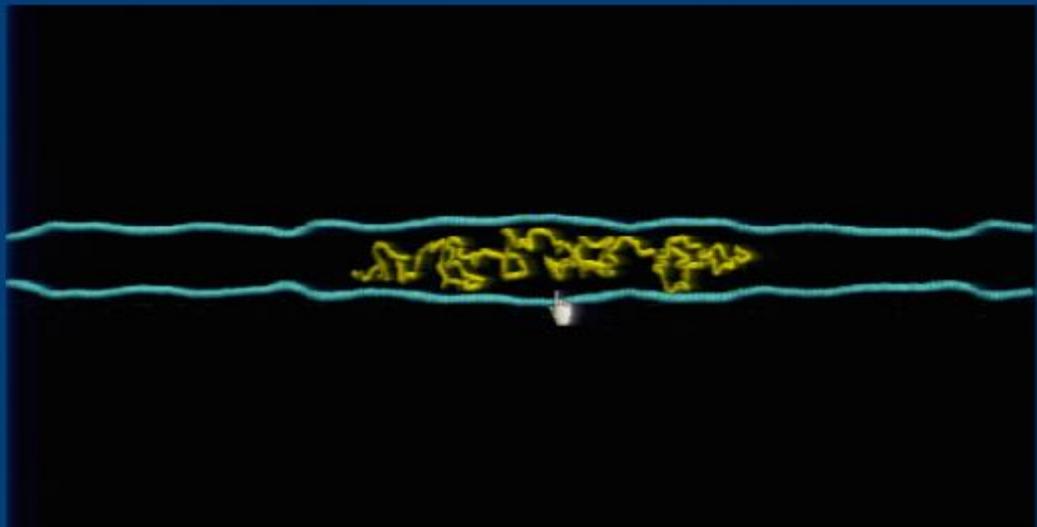
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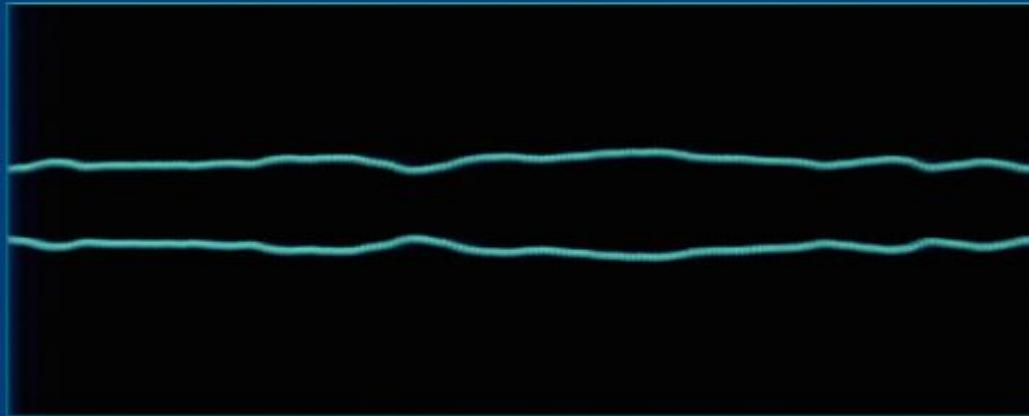
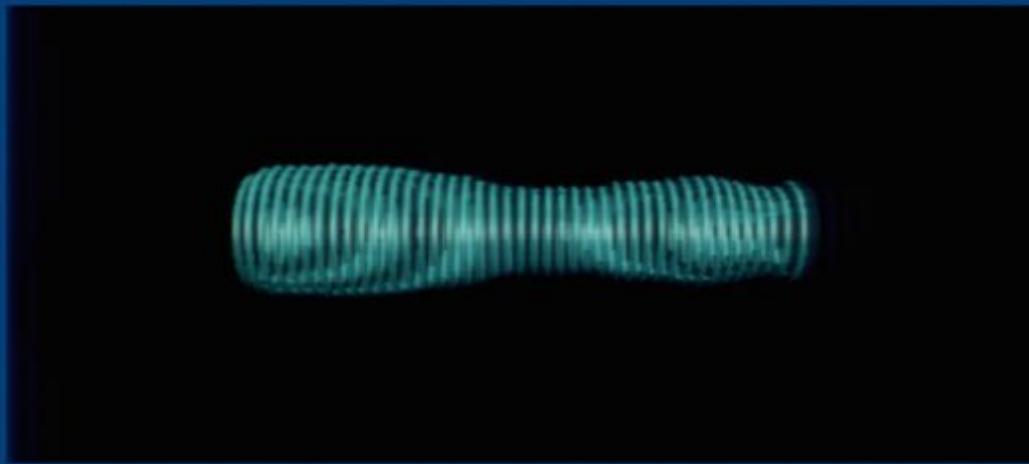
Increase  $\sigma$



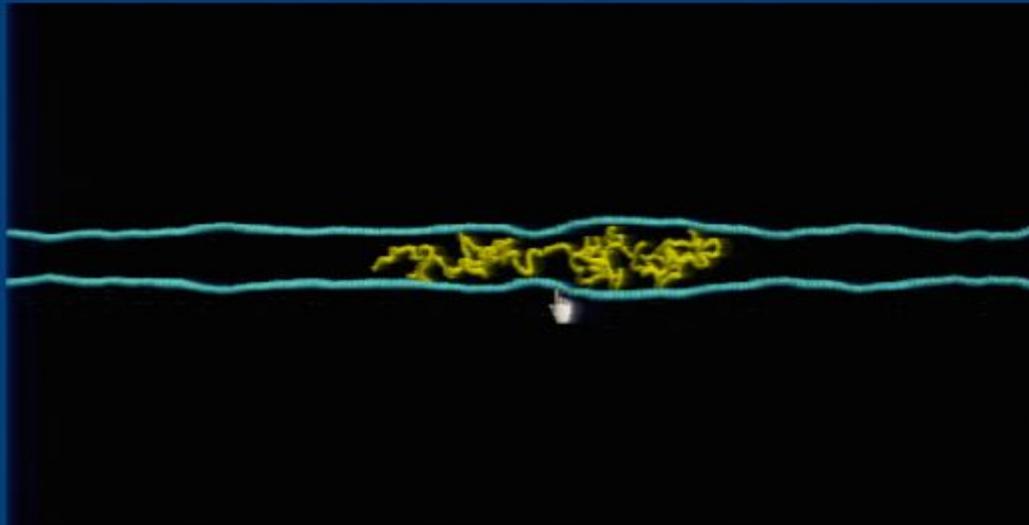
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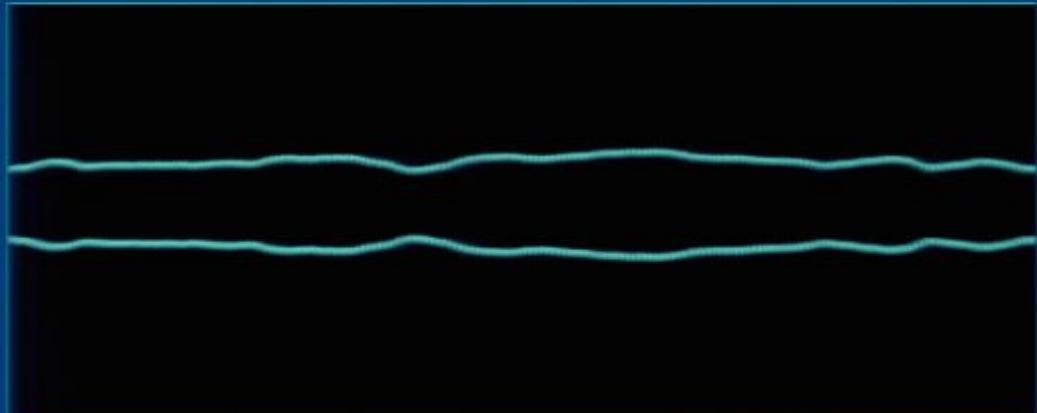
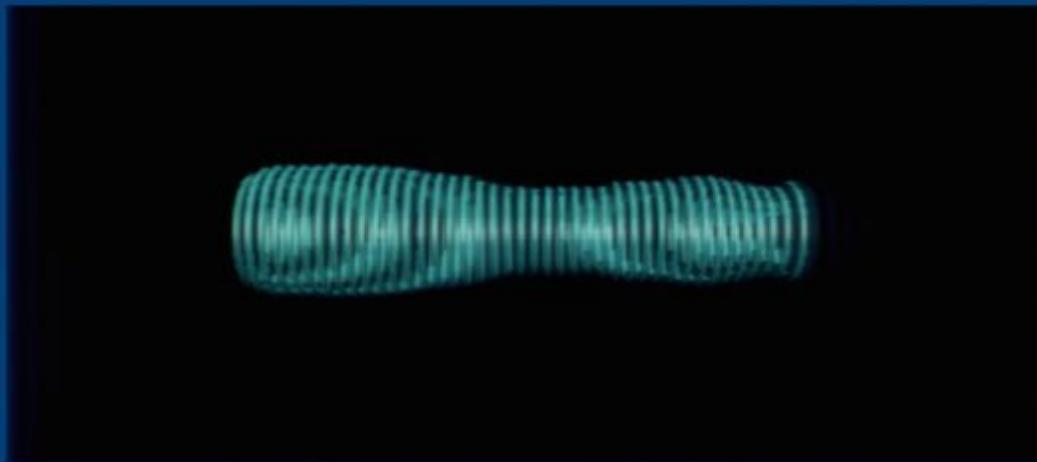
Increase  $\sigma$



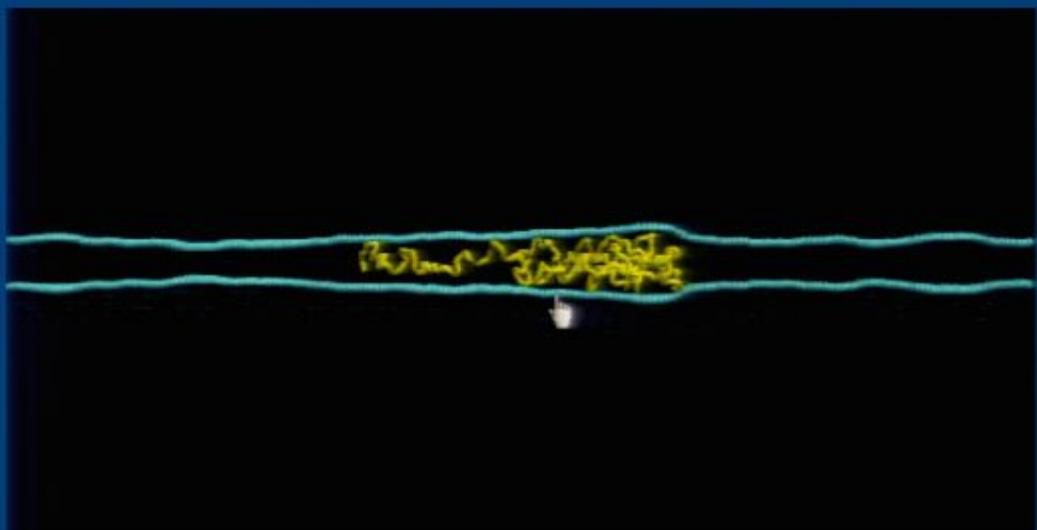
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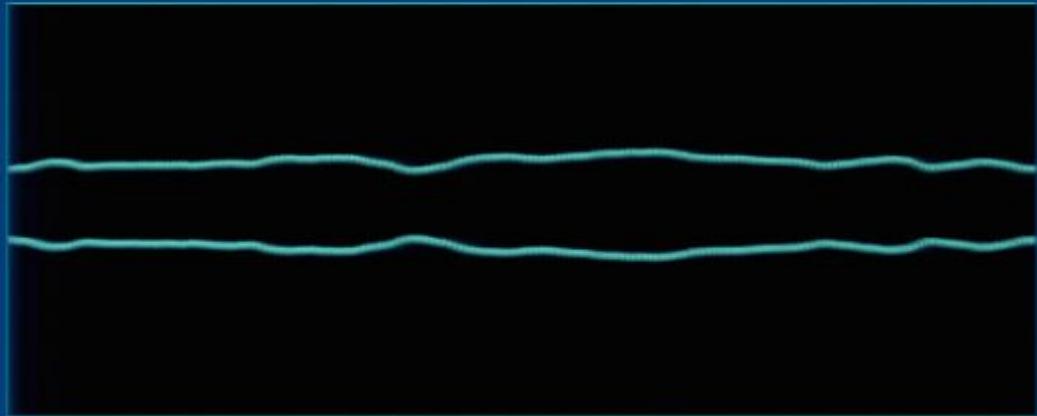
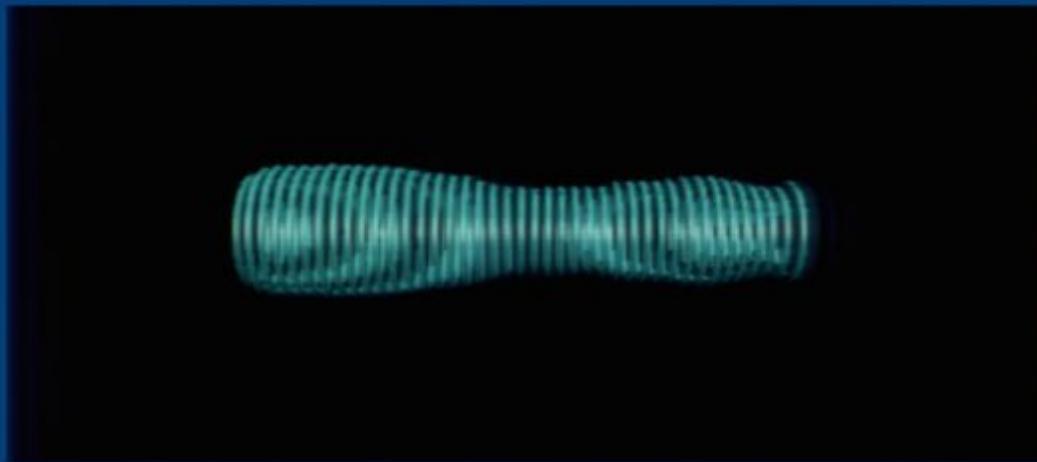
Increase  $\sigma$



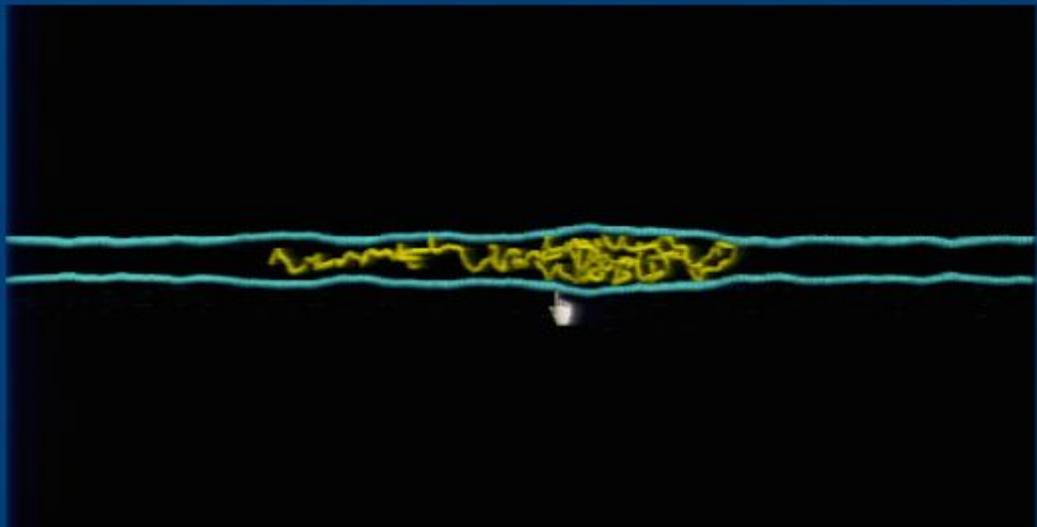
Equilibrium:  
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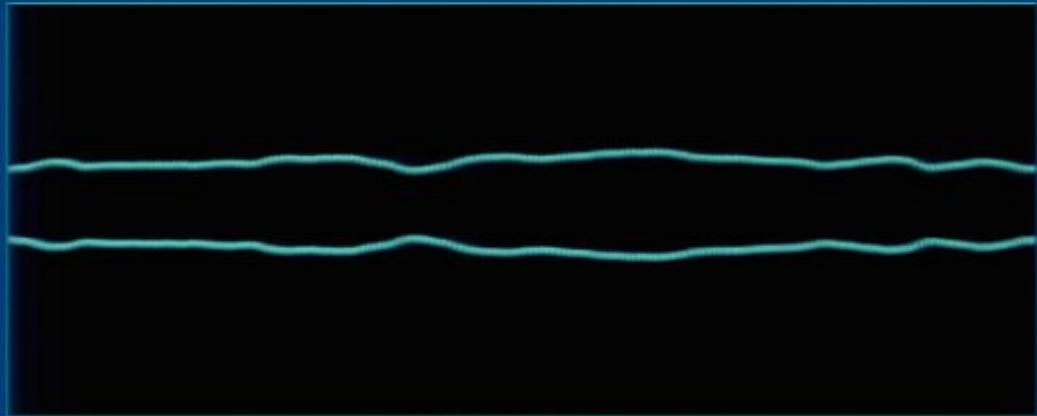
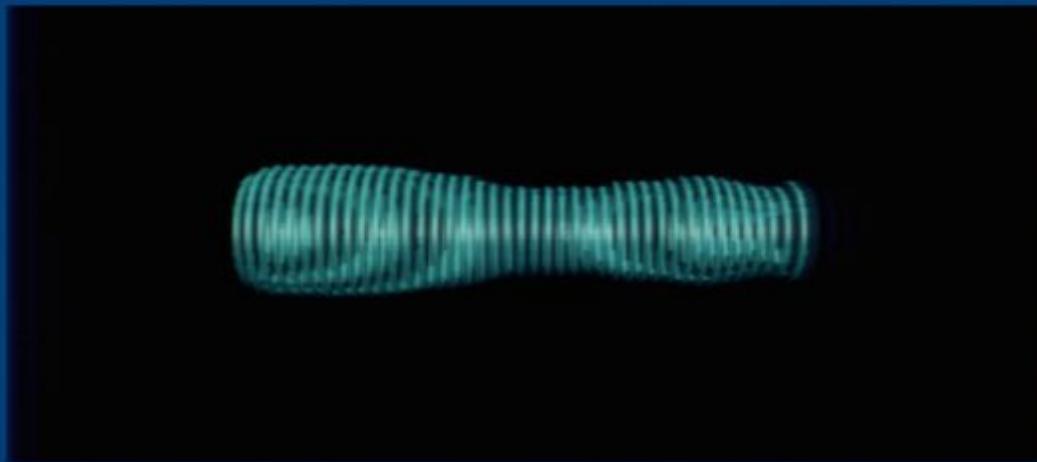
Increase  $\sigma$



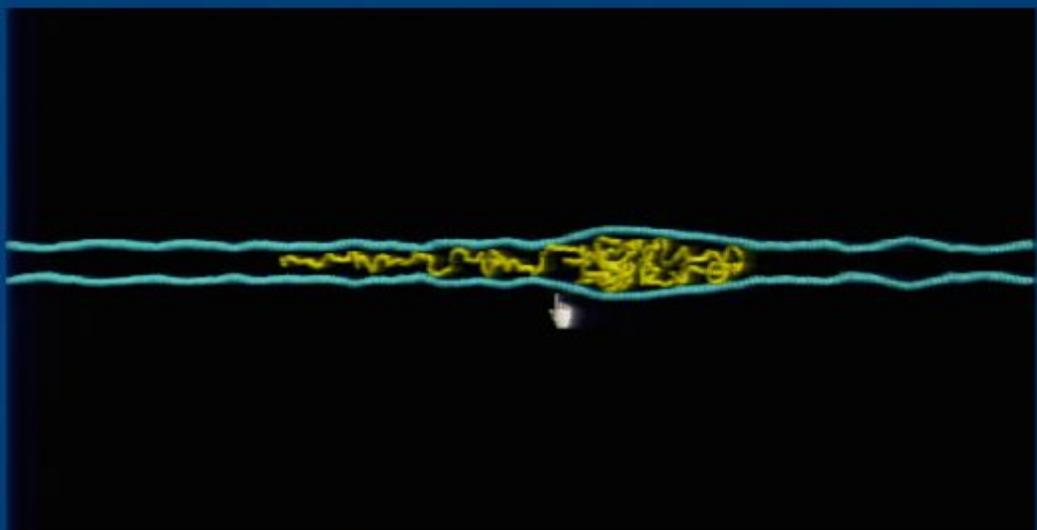
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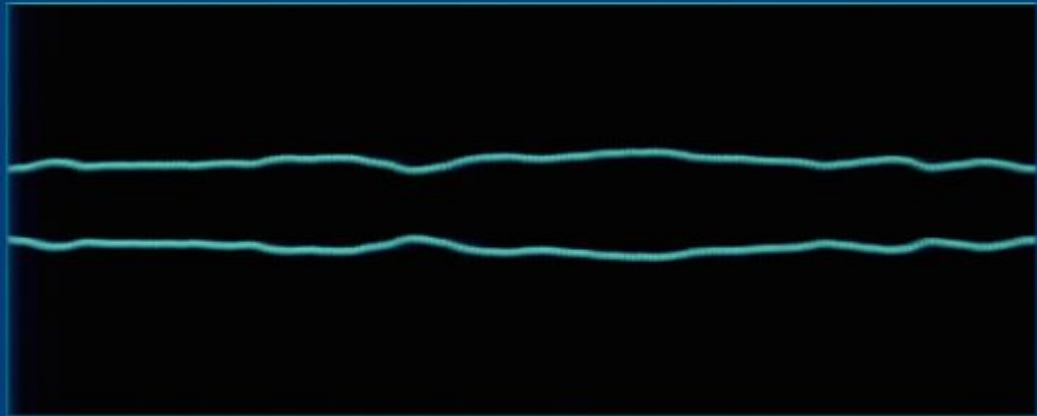
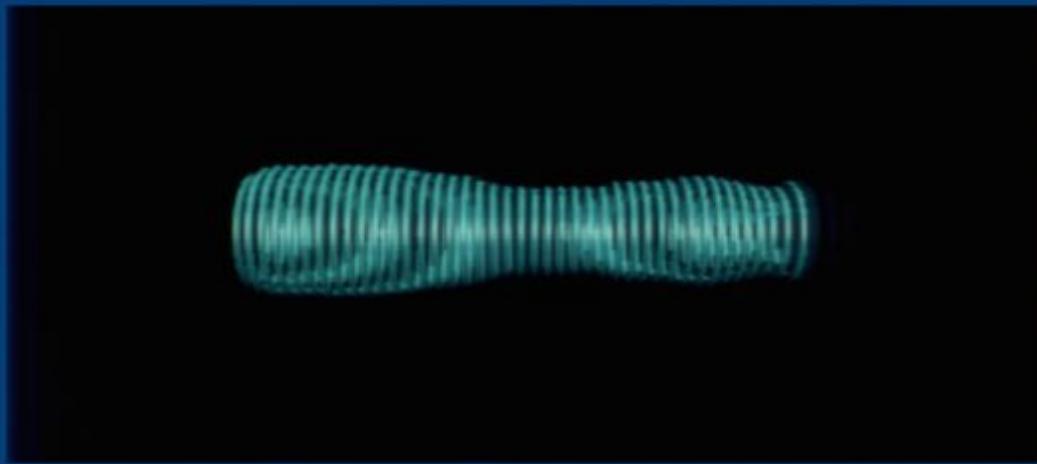
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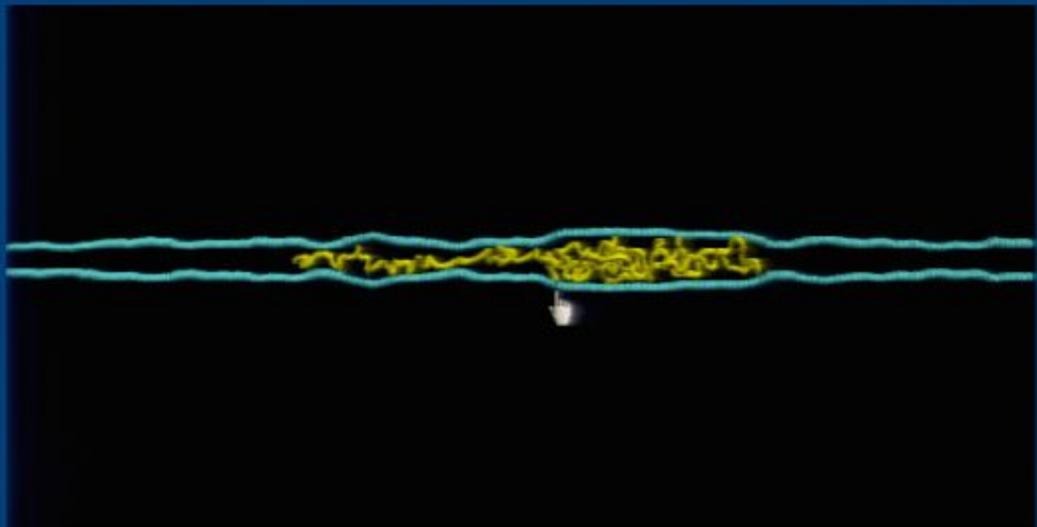
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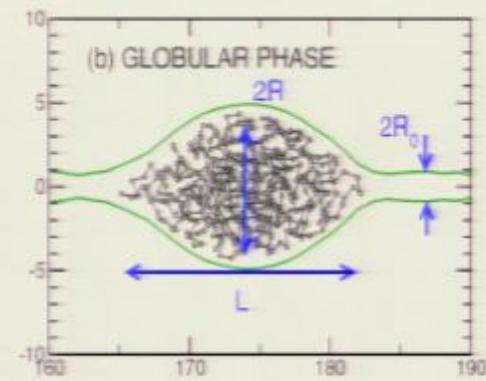
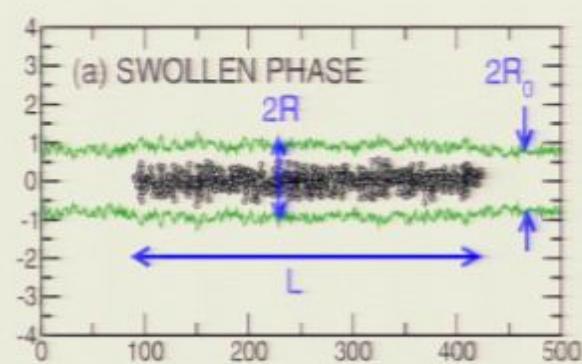
Increase  $\sigma$

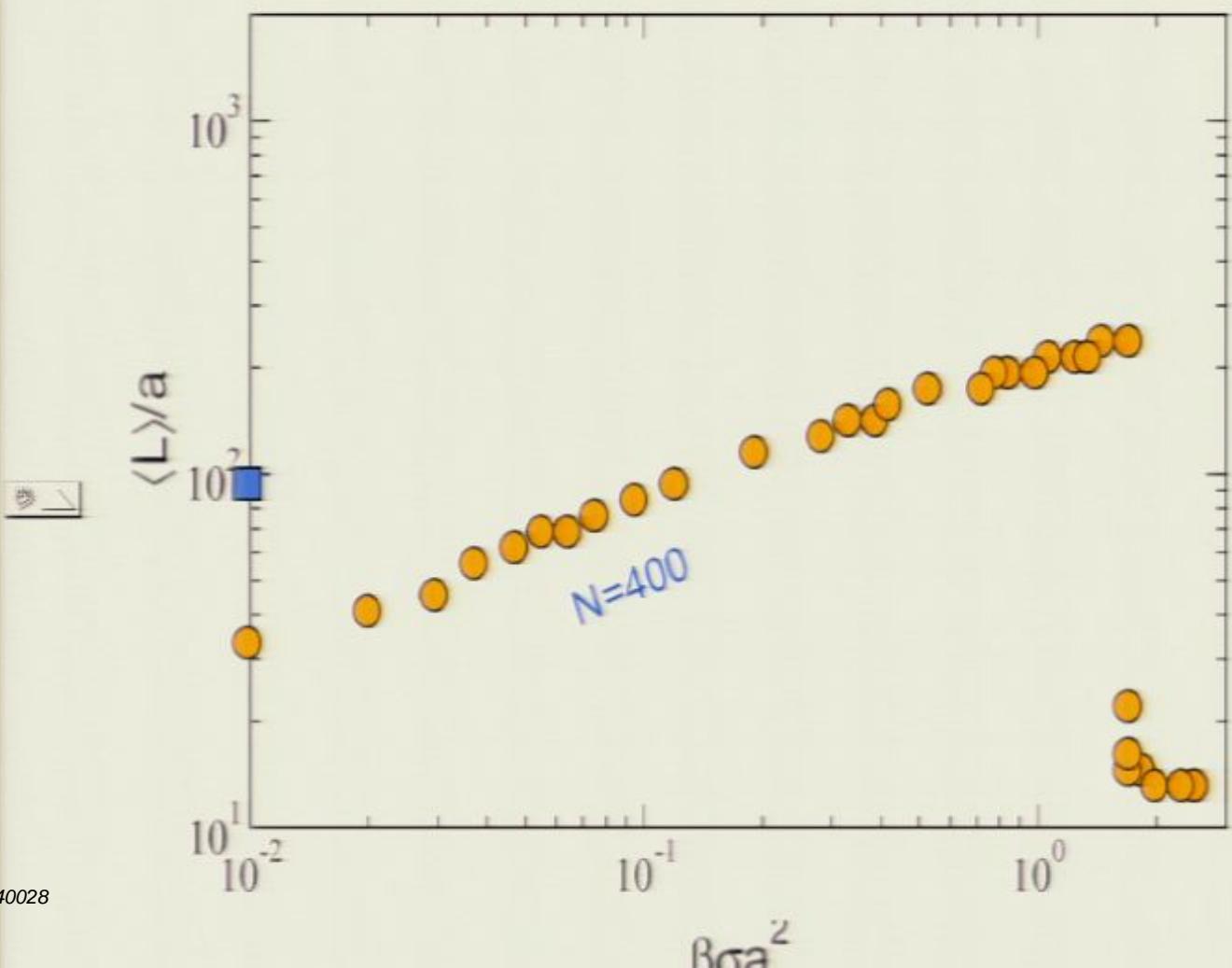
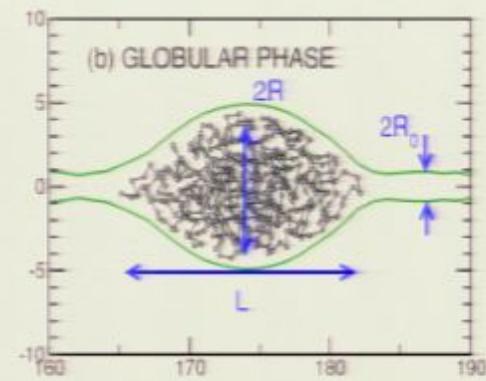
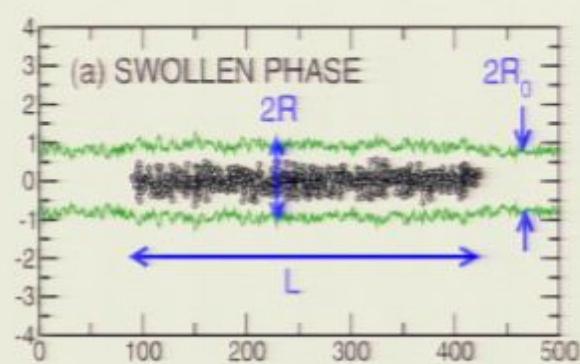


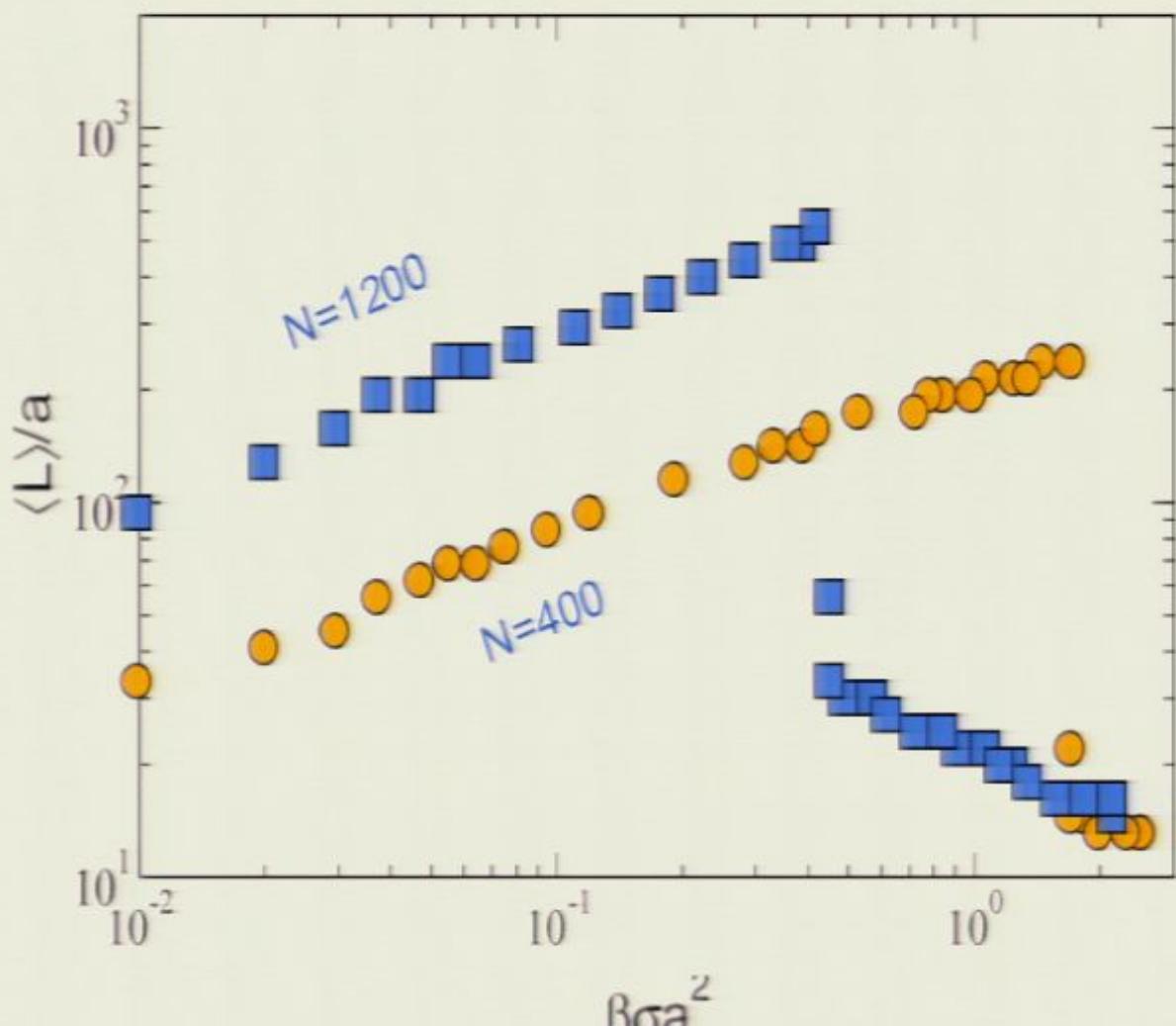
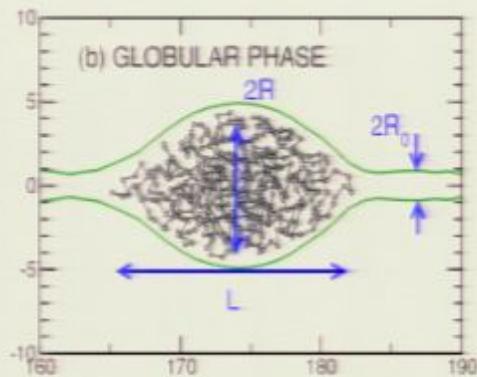
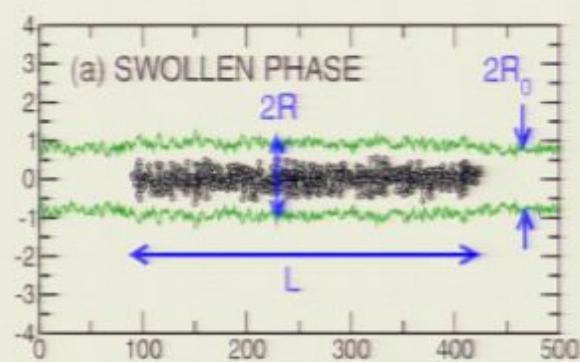
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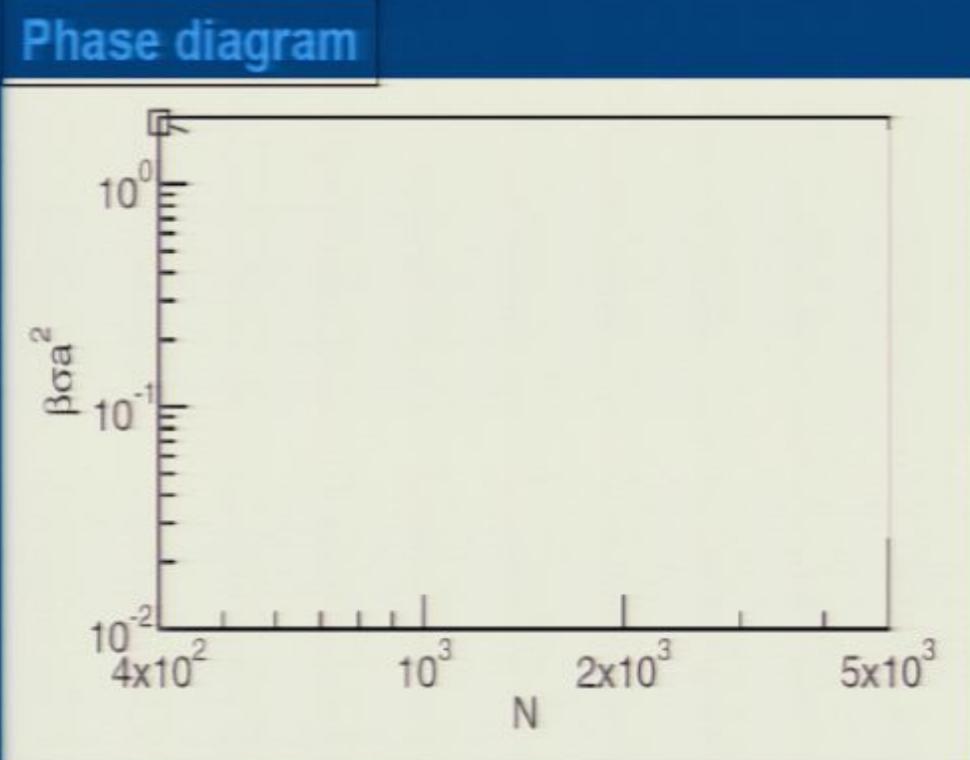
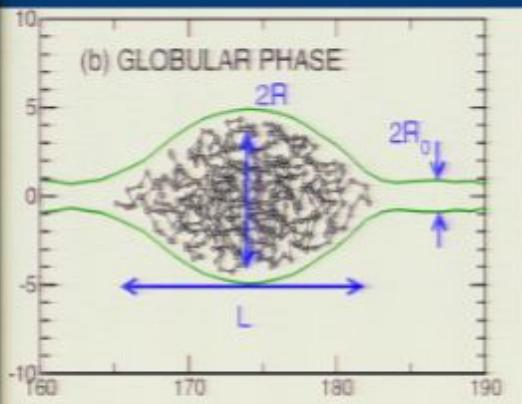
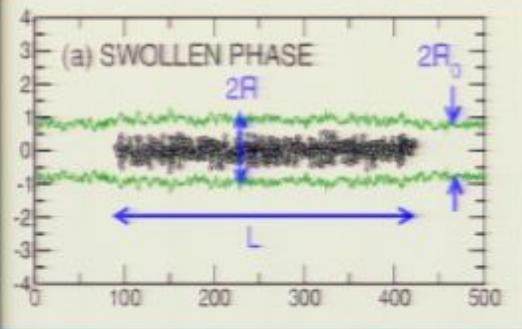


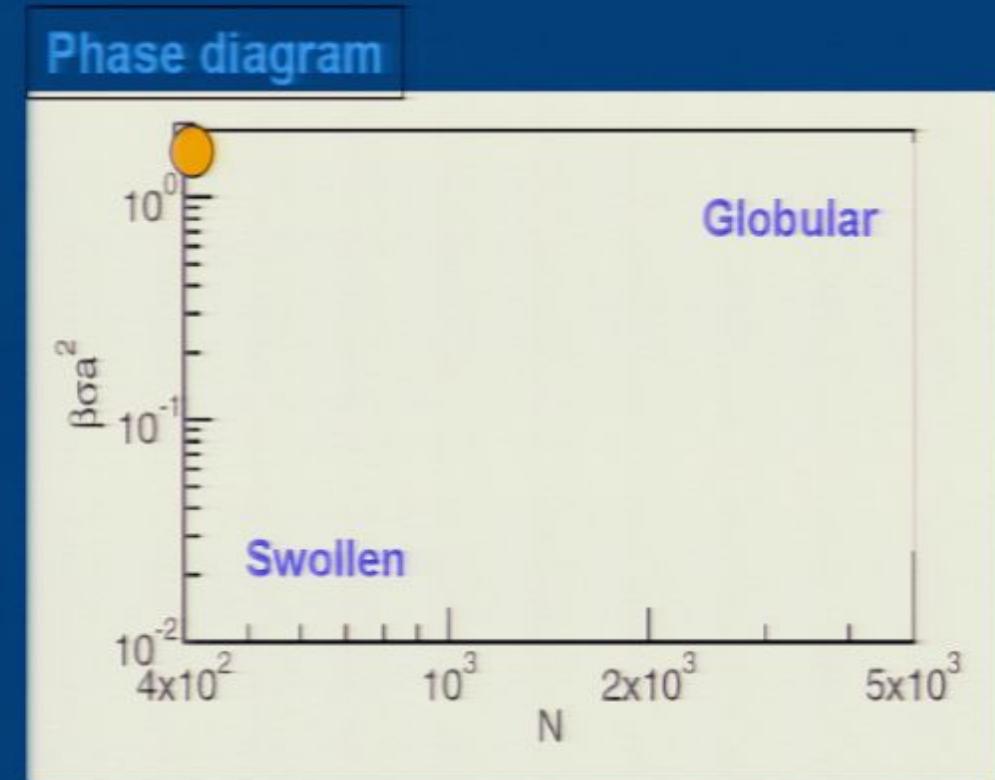
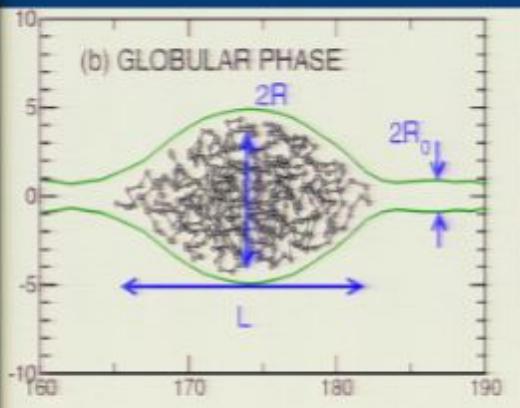
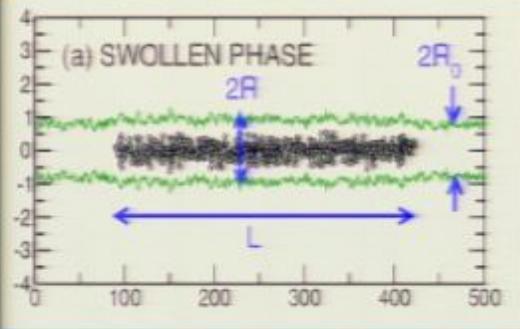
Increase  $\sigma$

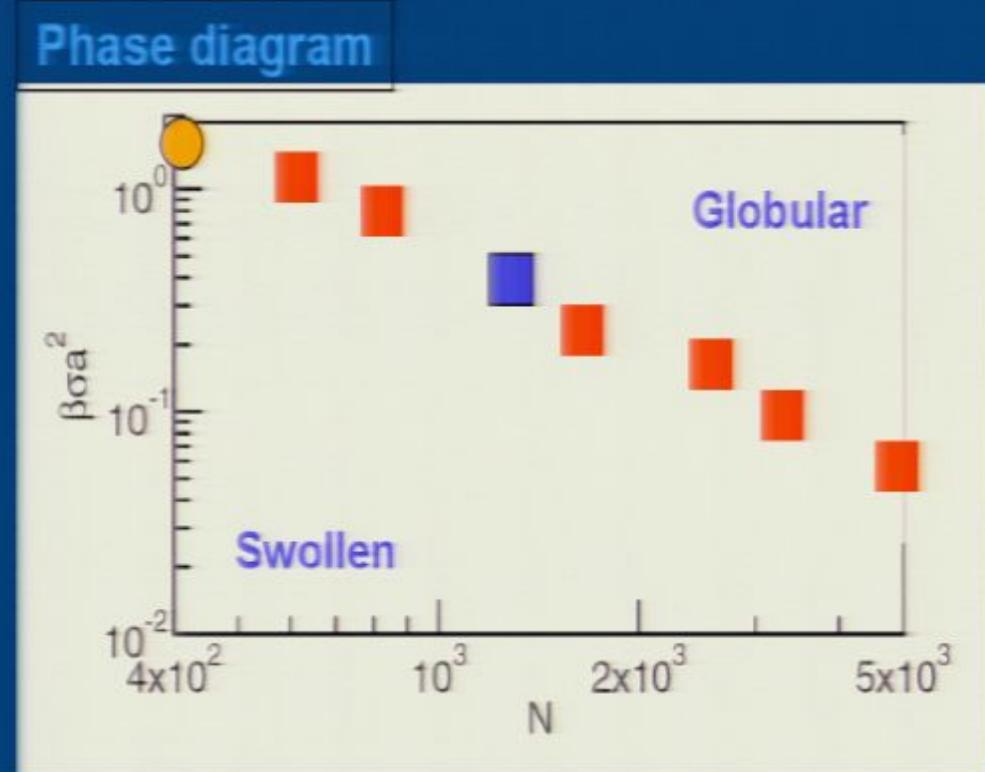
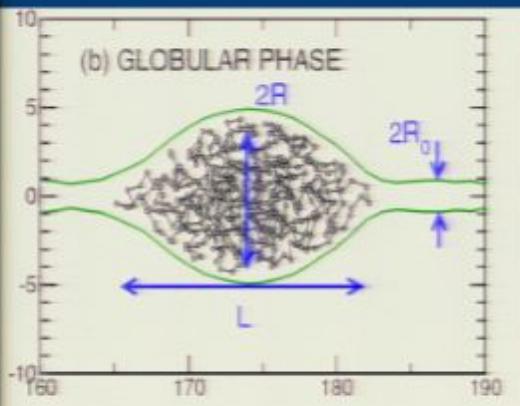
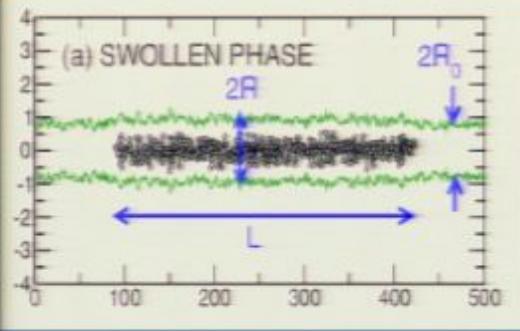




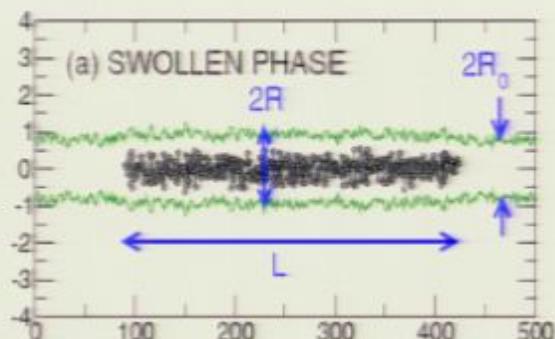






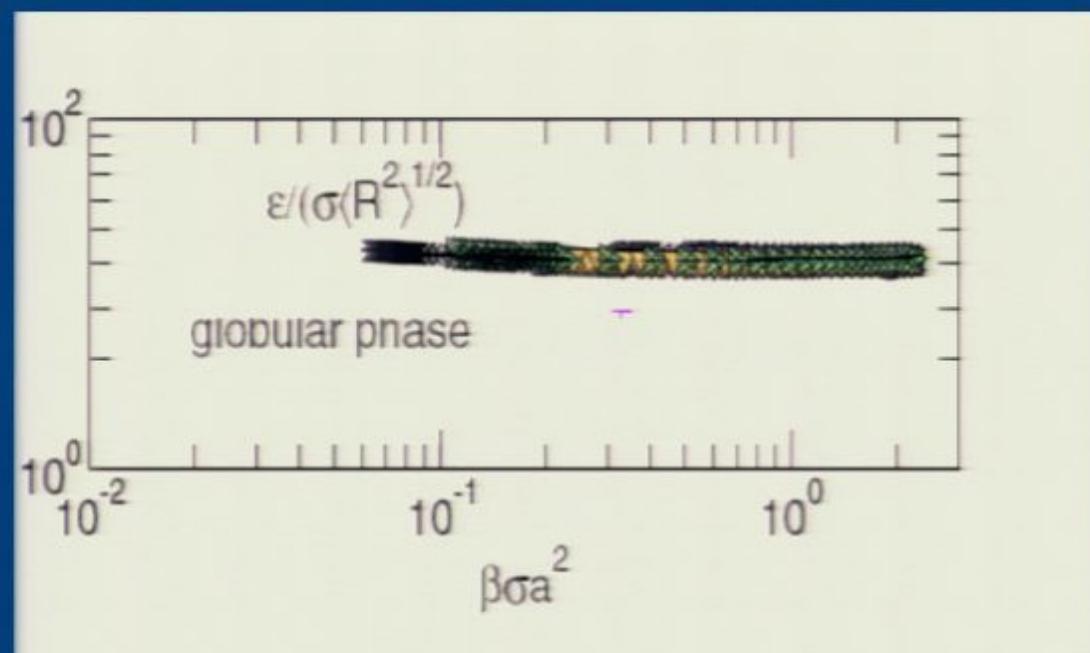
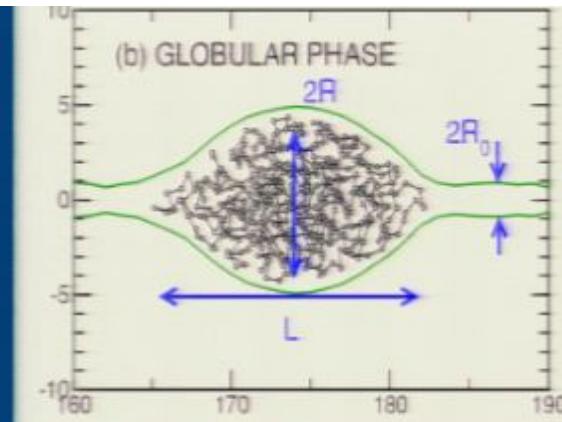


## Properties of the swollen phase



$$\text{Equilibrium: } R_0^2 = \kappa / 2\sigma$$

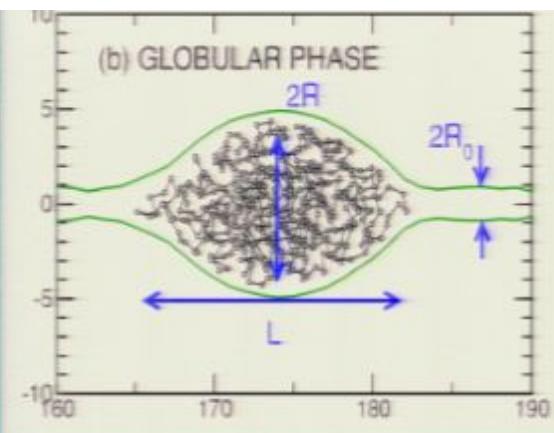
## Properties of the globular phase



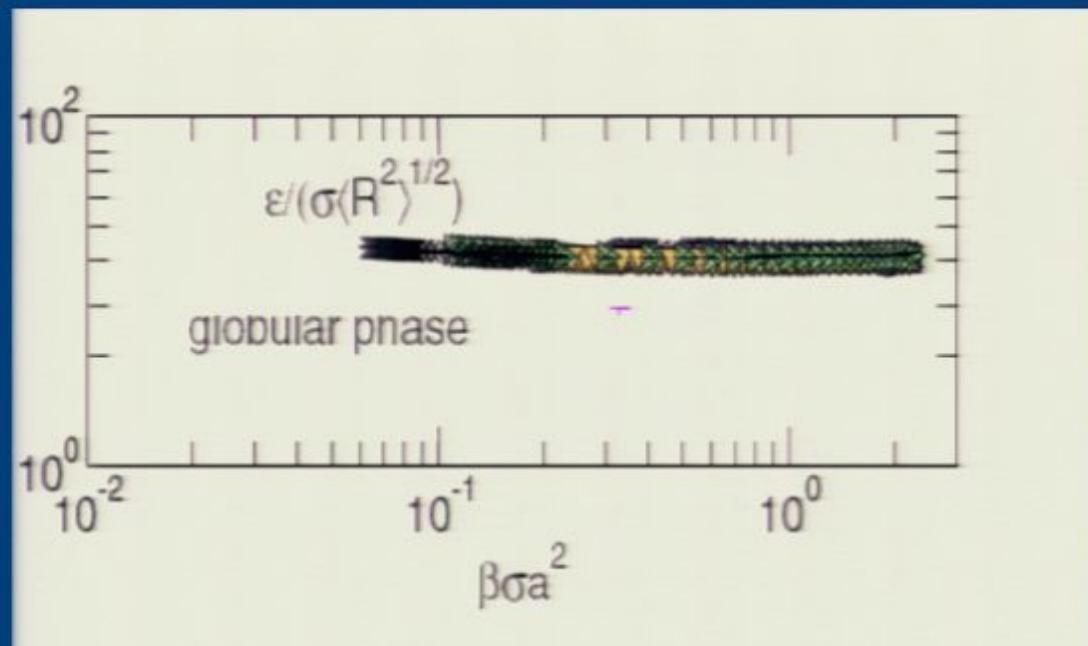
## Properties of the globular phase

$$\text{Helfrich energy} = \Delta A [\sigma + (\kappa/2) (1/r_1 + 1/r_2)^2] \\ \sim \Delta A \sigma + \dots$$

$$\text{Total energy} = 2\pi \langle R^2 \rangle^{1/2} L \sigma$$

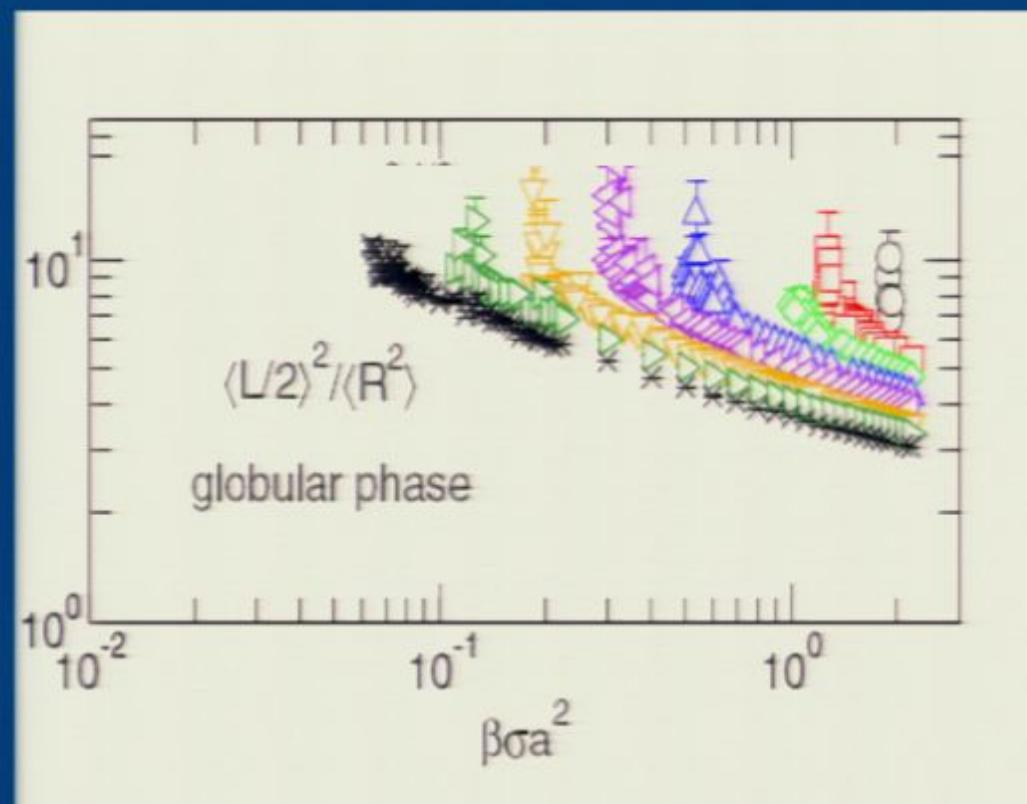
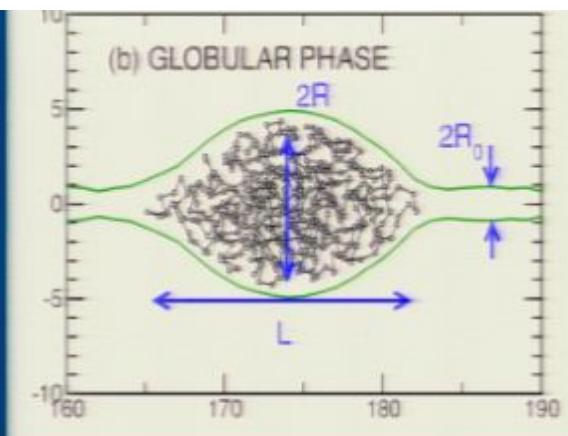


$$\text{Total energy}/L = \sigma + \epsilon/\sigma^{1/2}$$



## Properties of the globular phase

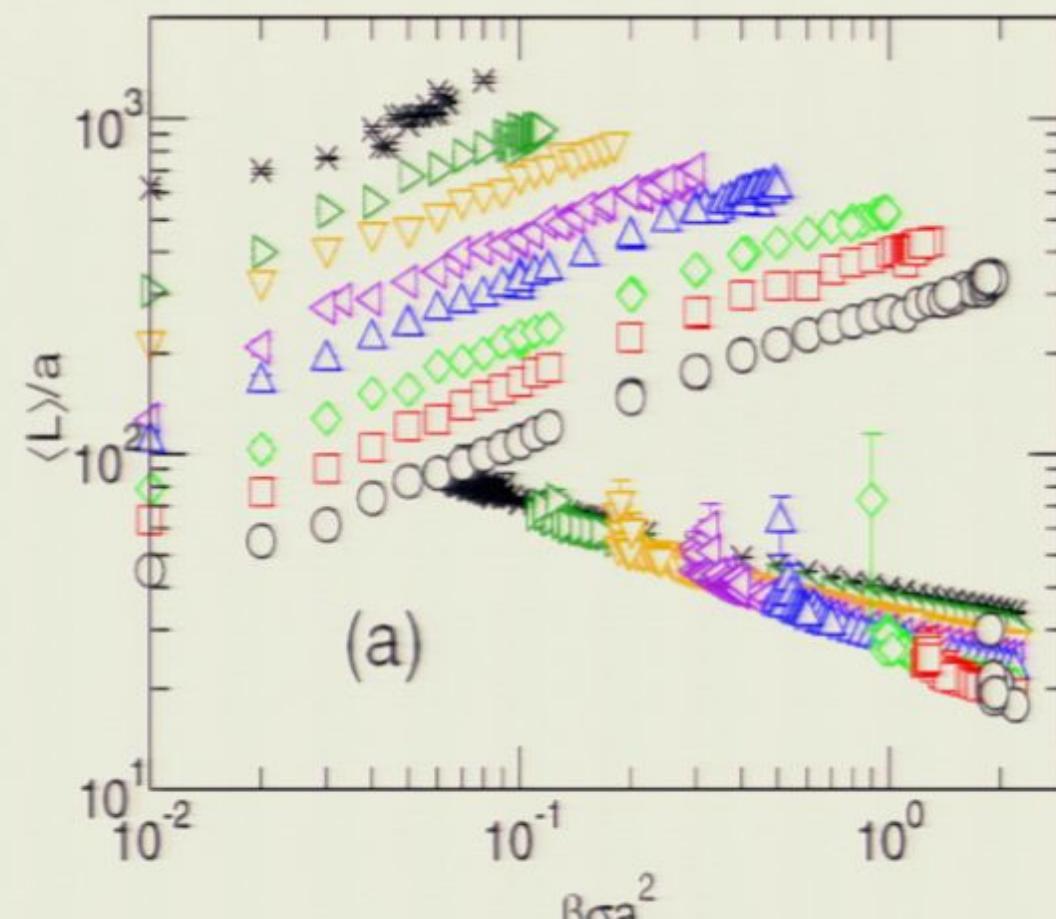
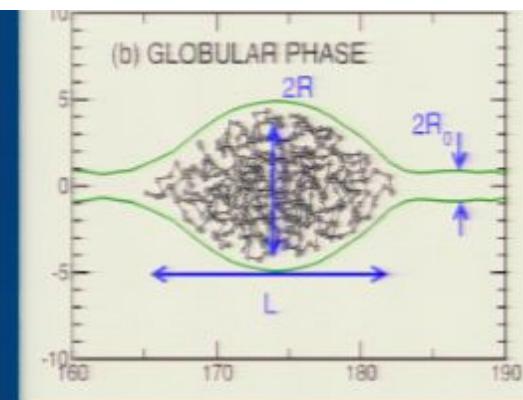
Scaling theory:  $L \sim R$



## Properties of the globular phase

$$\langle L \rangle = \text{Constant } N^\alpha \sigma^{-\beta}$$

■ Scaling theory:  $\alpha=2/5$ ,  $\beta=1/5$



# TOPIC 2:

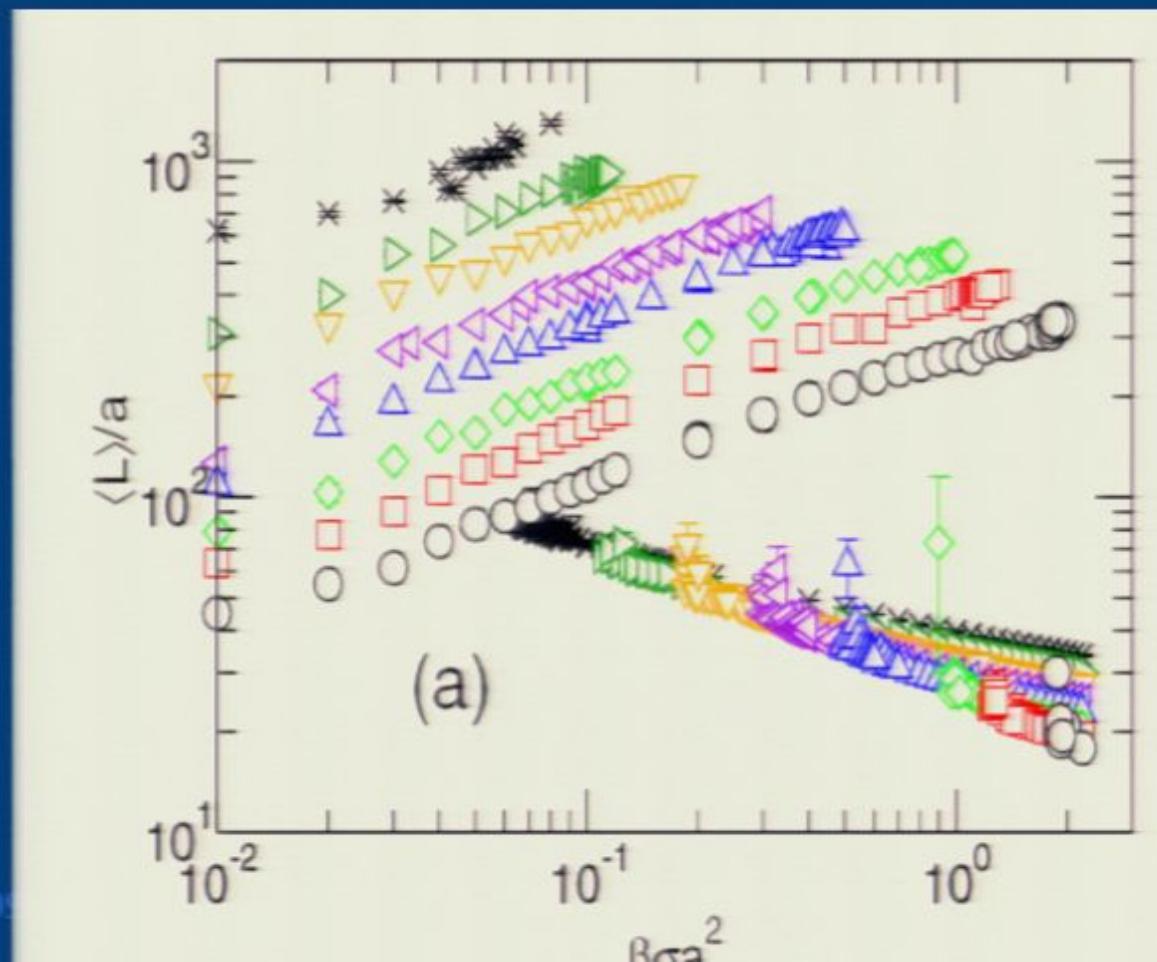
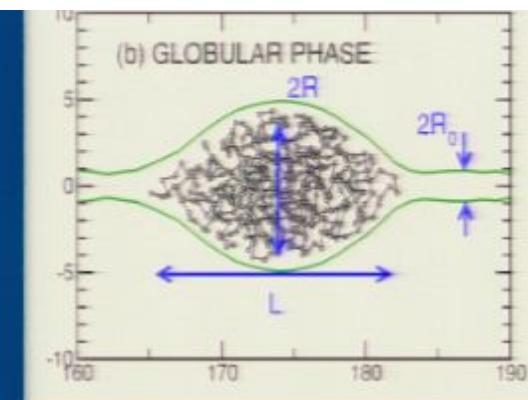
## Nanospheres attracted to a flat membrane

## Properties of the globular phase

$$\langle L \rangle = \text{Constant } N^\alpha \sigma^{-\beta}$$

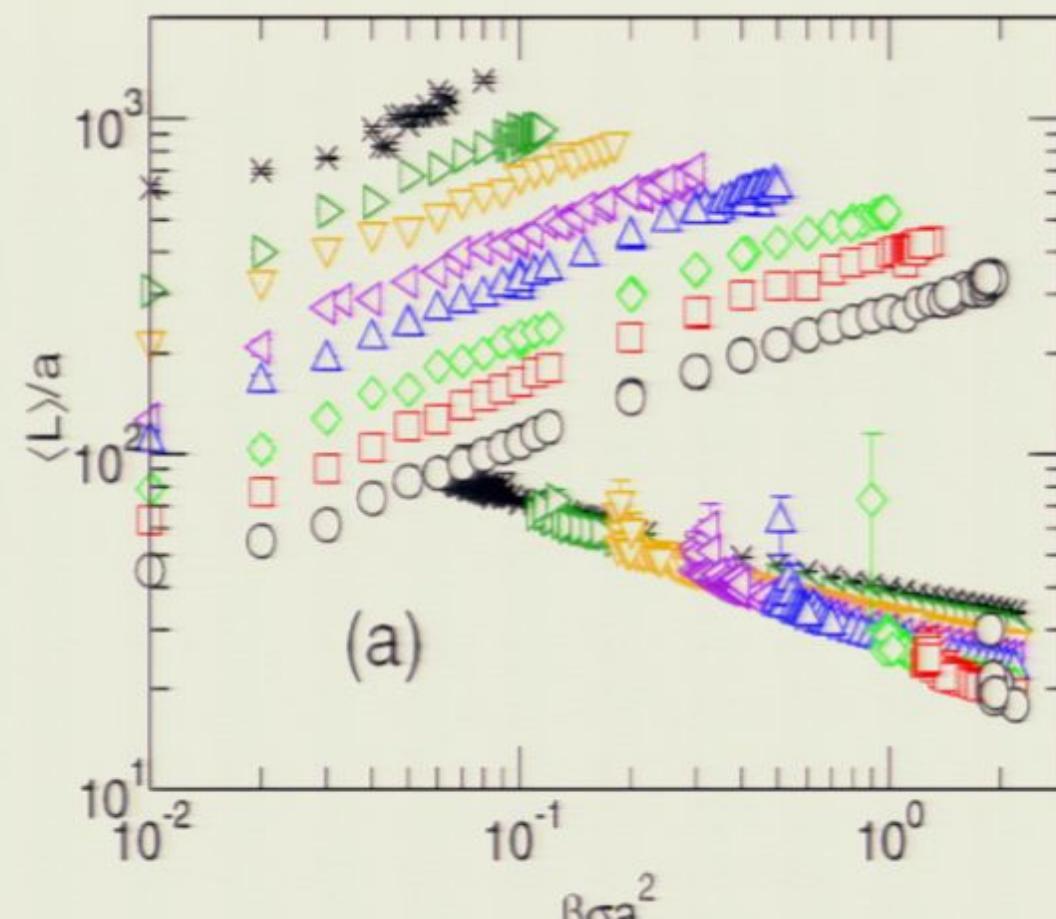
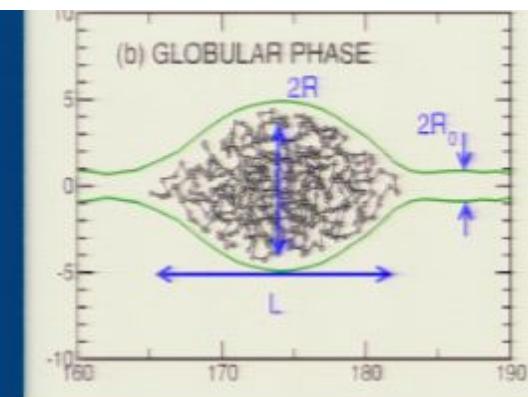
■ Scaling theory:  $\alpha=2/5$ ,  $\beta=1/5$

■ Simulation: much stronger dependence

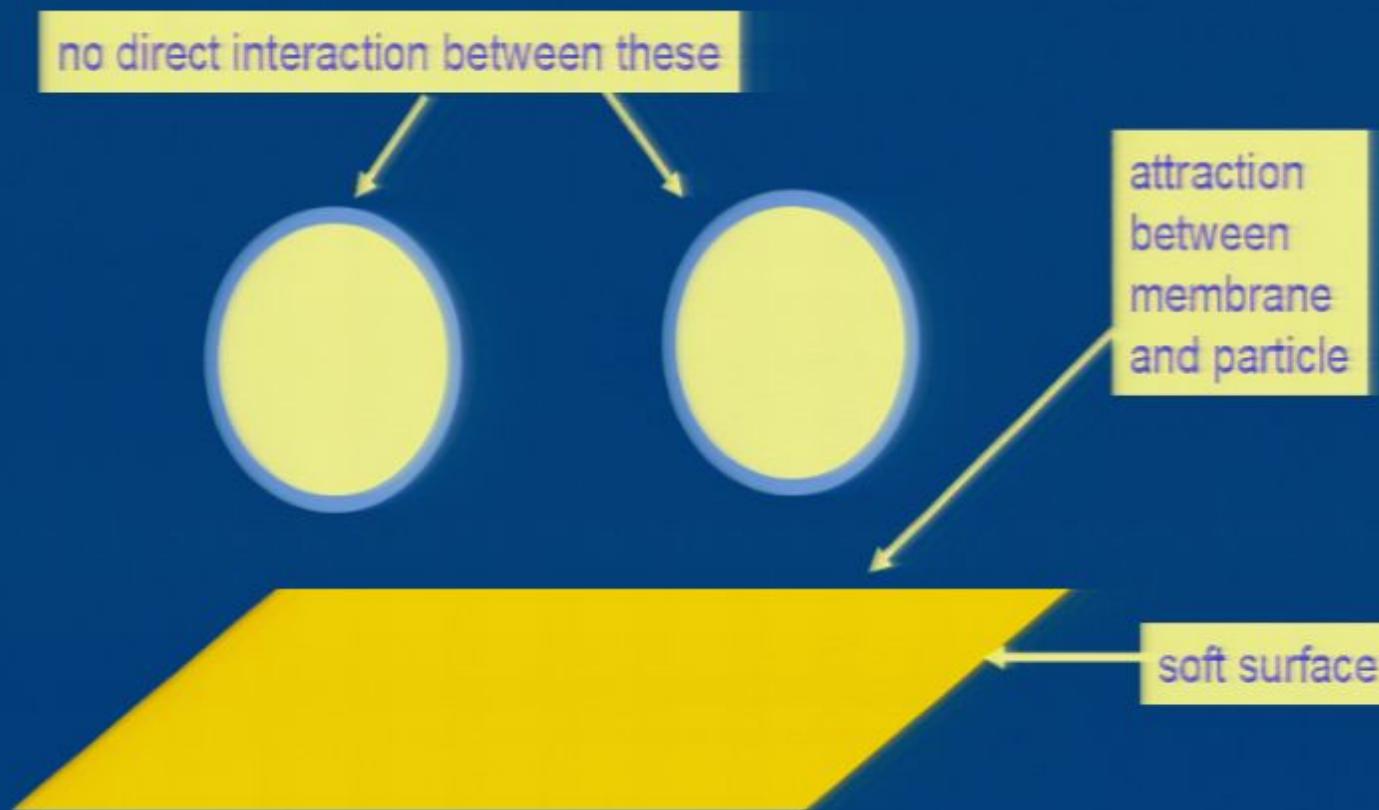


## Properties of the globular phase

$$\langle L \rangle = \text{Constant } N^\alpha \sigma^{-\beta}$$



### Adsorption of two colloid particles on a soft membrane



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- Under what conditions would the colloid particles be adsorbed onto the surface.

### Adsorption of two colloid particles on a soft membrane

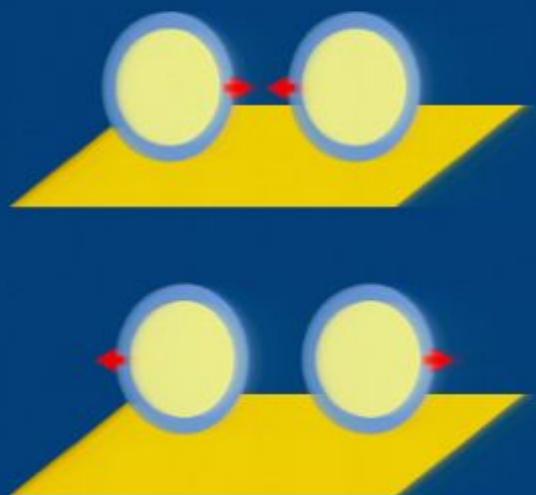


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### Adsorption of two colloid particles on a soft membrane



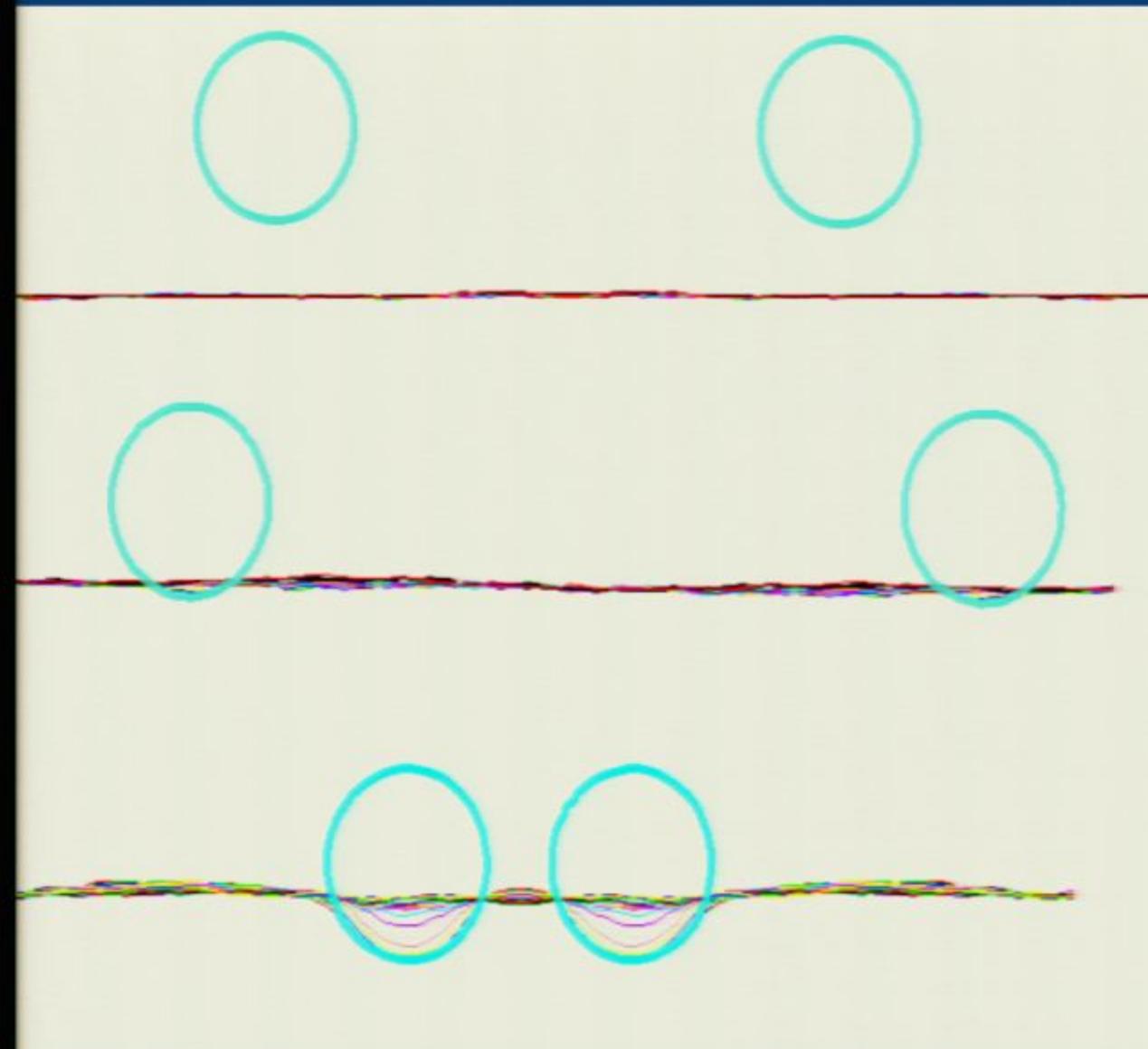
- Under what conditions would the colloid particles be adsorbed onto the surface.



- After the adsorption, is there a membrane induced capillary force between the particles? Attraction? Repulsion?

## Why do we study this model?

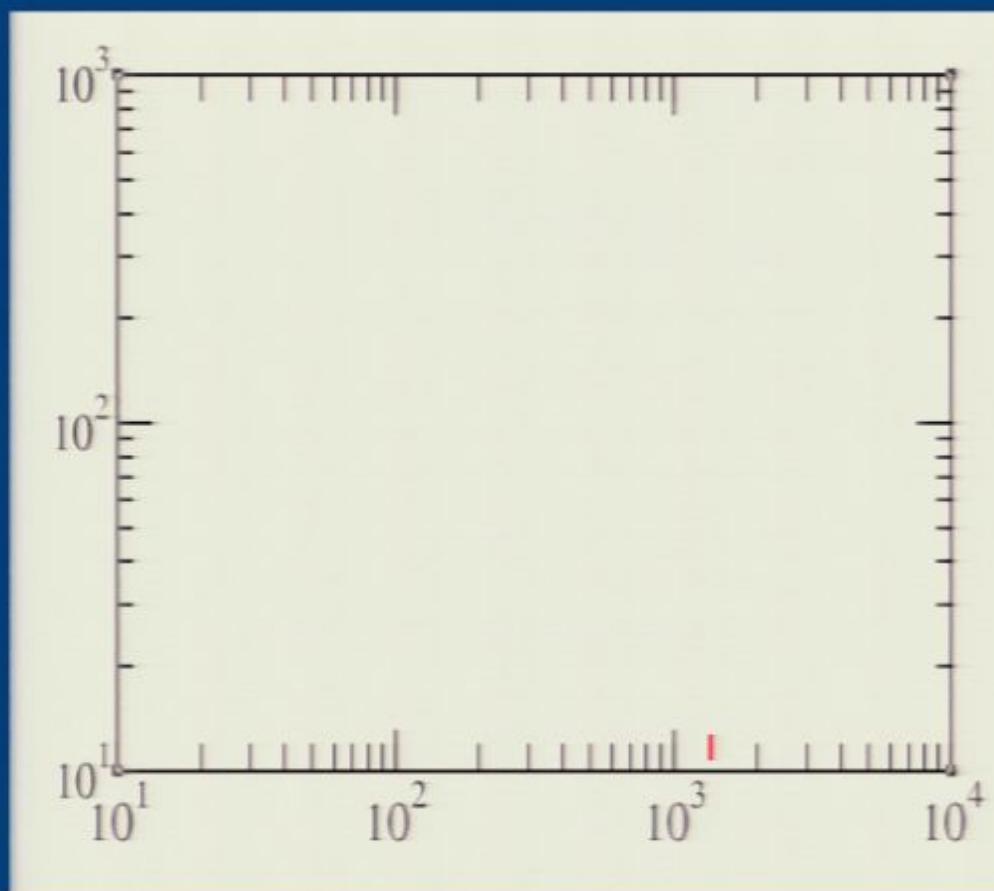
## Structures founded in the simulation



■ Free state

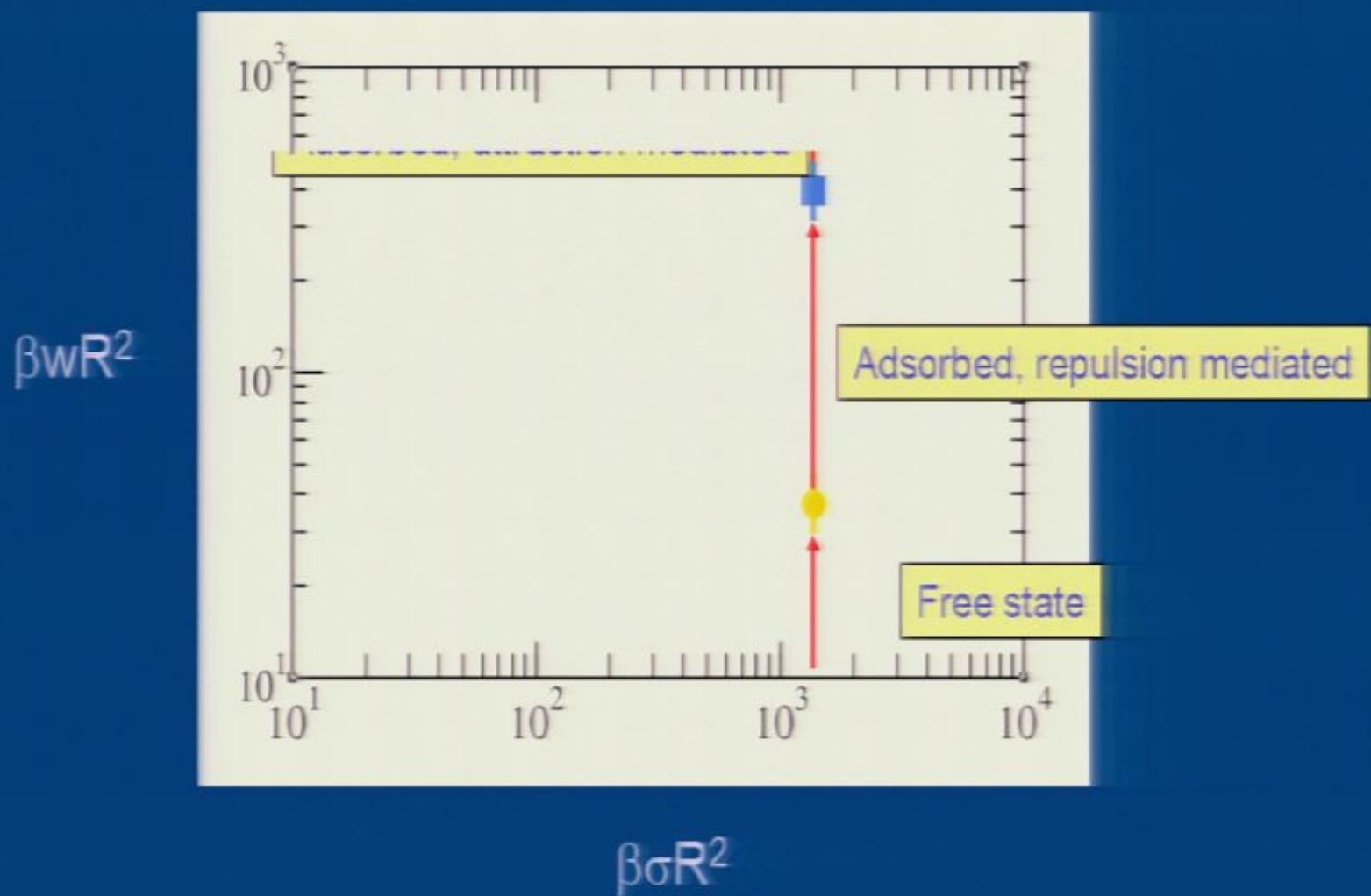
## State diagram

$\beta w R^2$

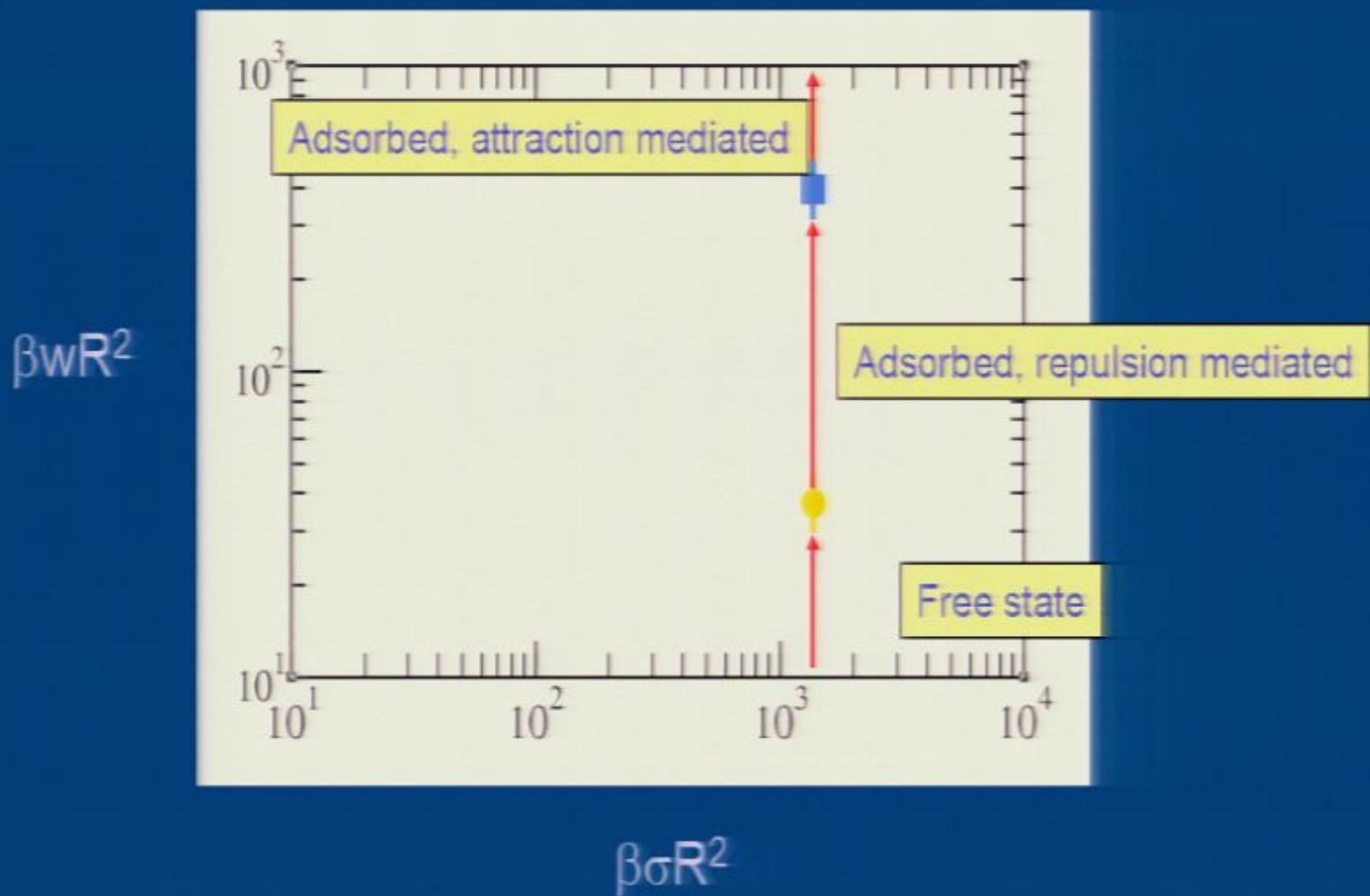


$\beta \sigma R^2$

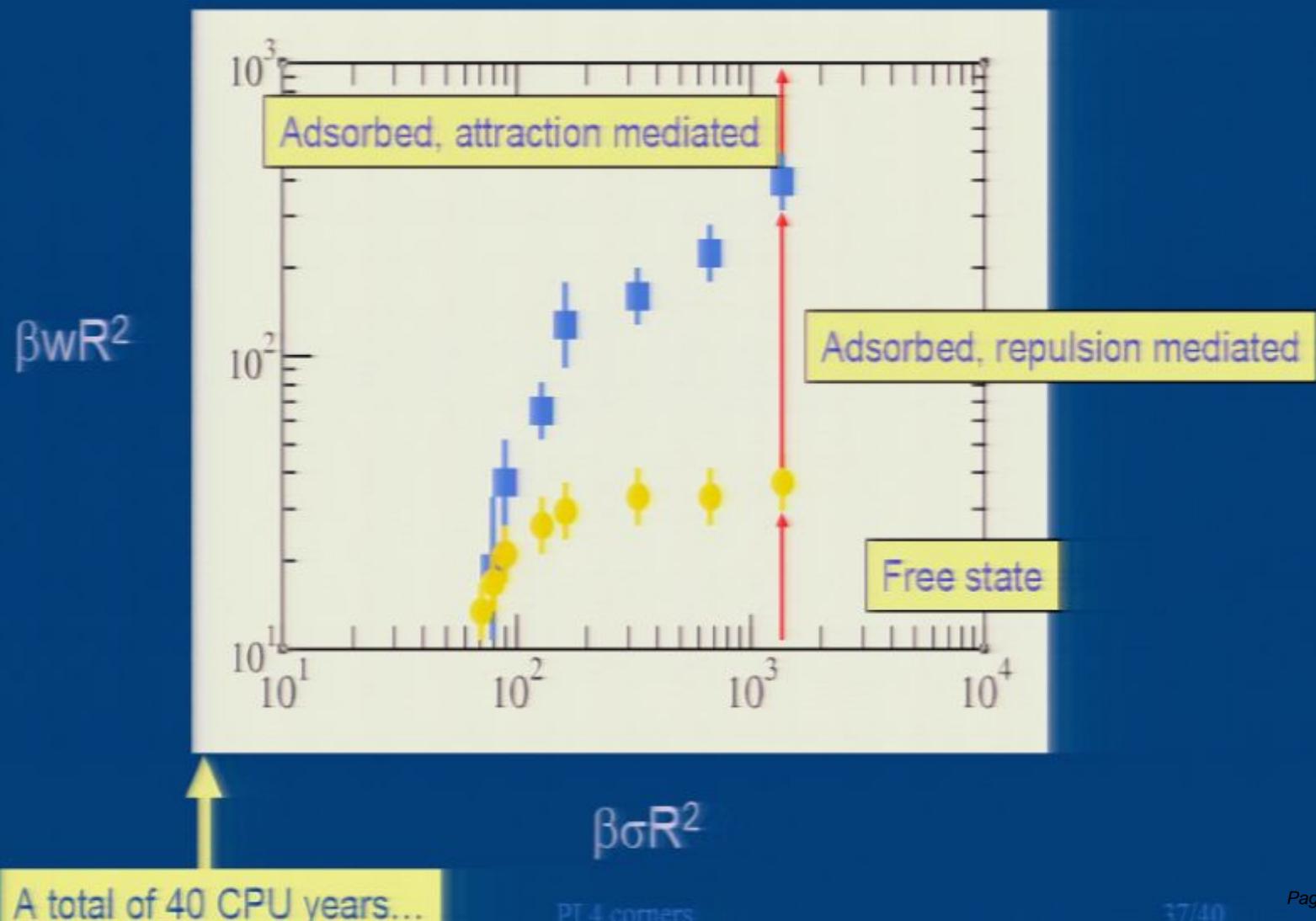
## State diagram



## State diagram



## State diagram



Transition from free to adsorption states

$$\text{Helfrich energy} = \Delta A [\sigma + (\kappa/2) (1/r_1 + 1/r_2)^2]$$

$$\beta F = -\beta w A$$

■ Adsorption energy

$$+ \beta \sigma R^2 (A^2 / 4\pi R^4)$$

■ Free energy change due to surface stretching

$$+ (\beta \kappa/2) (2/R)^2 A$$

■ surface bend

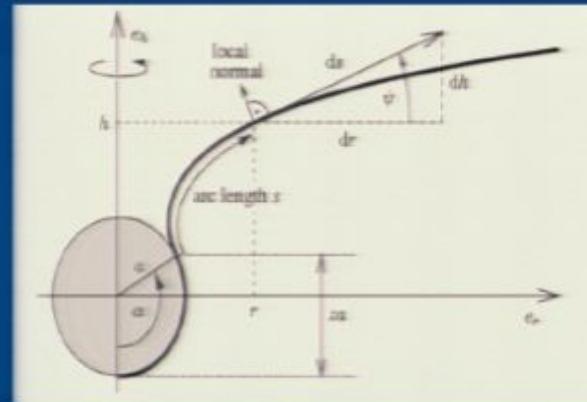
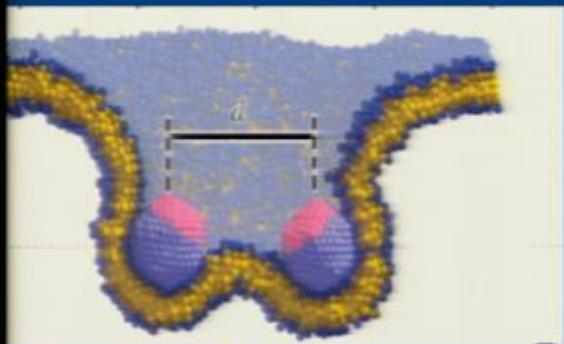
+ ...

■ F of unadsorbed part; fluctuations

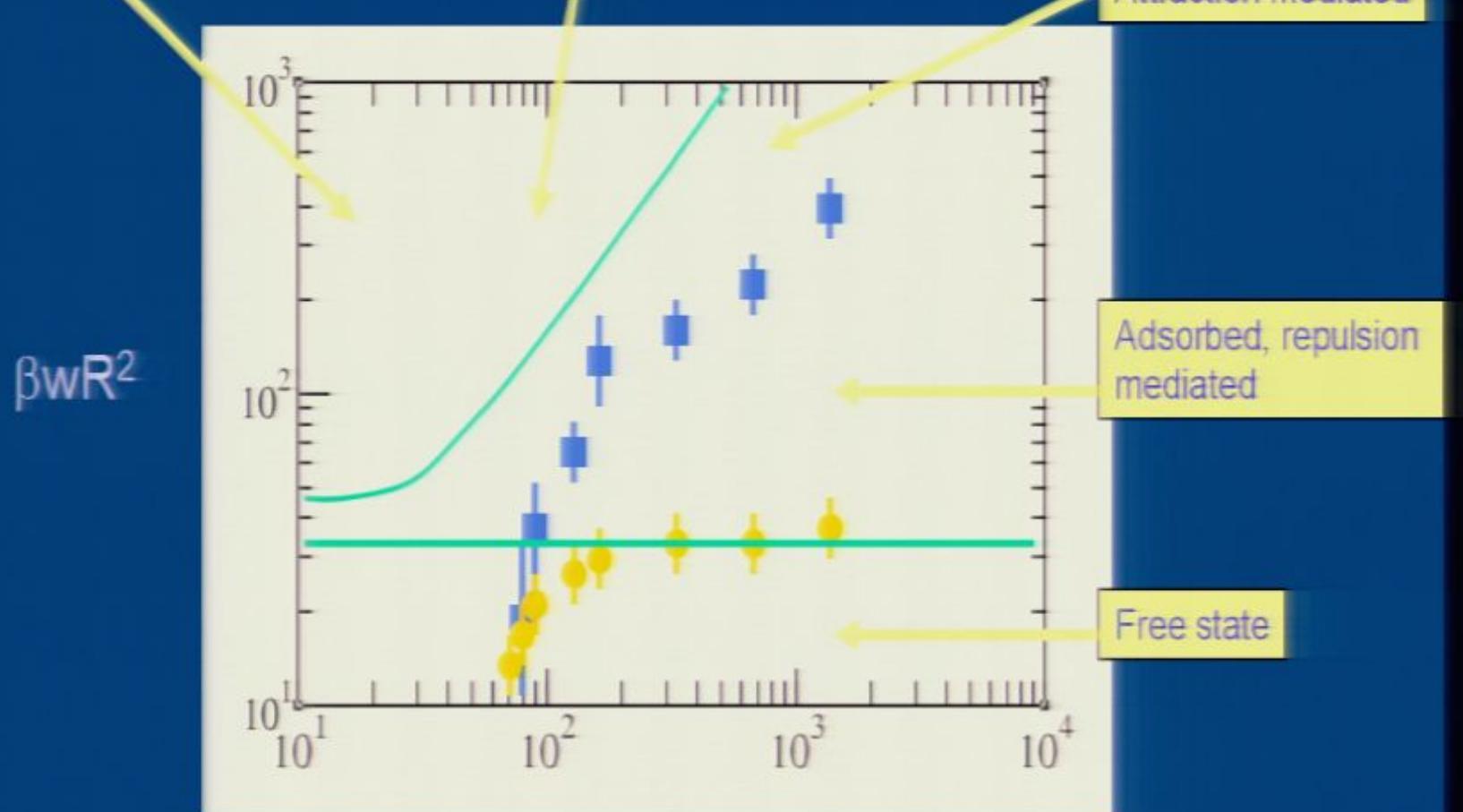


$\beta w R^2 = 2\beta \kappa$  free-to-adsorption transition

## SUMMARY



Adsorbed,  
Attraction mediated



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- We have developed a useful computational procedure that can effectively handle a soft membrane simulation.

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- We have developed a useful computational procedure that can effectively handle a soft membrane simulation.
- The procedure enable us to identify a **first-order** swollen to globular transition for the first time, in polymer-tube system.
- Understanding a simple membrane/macromolecule system can be theoretically and computationally challenging.