

# Multi-scale modelling of flow-induced crystallisation in polymers.

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School of Mathematical Sciences, University of Nottingham.



The University of  
**Nottingham**

# Acknowledgements

## Collaborators

- Daniel Read, Oliver Harlen and Chinmay Das (University of Leeds)
- Muhammad Anwar (Institute of Space Technology, Islamabad)
- Claire McIlroy (University of Lincoln)
- Peter Olmsted (Georgetown University)
- Kenny Jolley (Loughborough University)
- Matthew Hamer
- Dow Chemicals
- SCG Chemicals
- Autodesk

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- EPSRC (EP/G048827/1) and (EP/P005403/1)
- School of Mathematical Sciences, University of Nottingham (PhD Studentship)

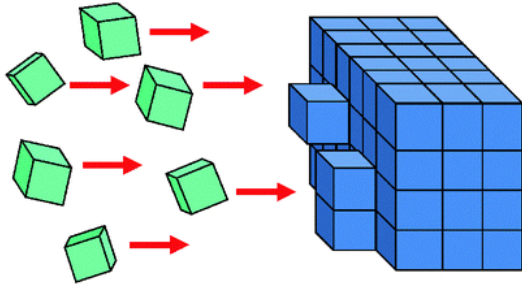


The University of  
**Nottingham**



# Crystallisation in polymers

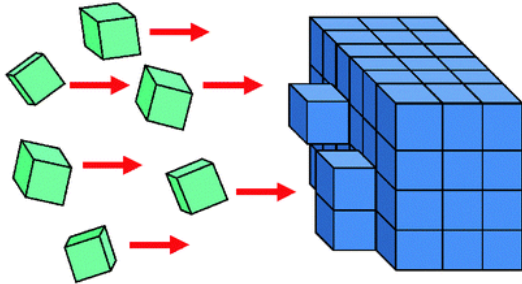
## SMALL MOLECULES



a

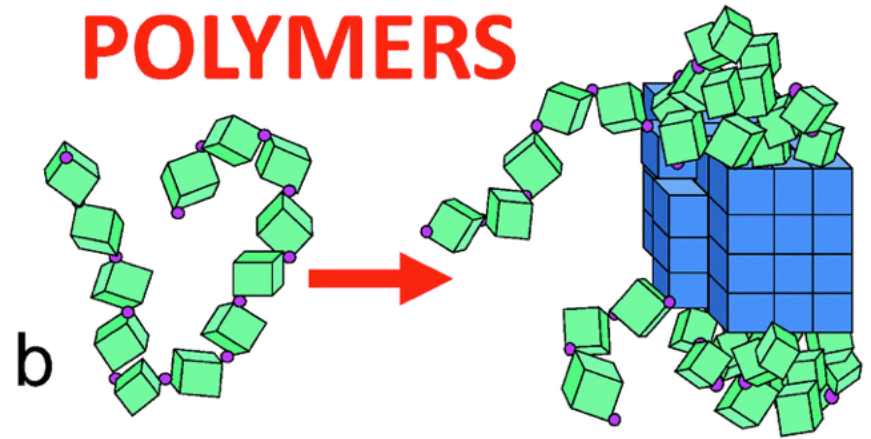
# Crystallisation in polymers

**SMALL MOLECULES**



a

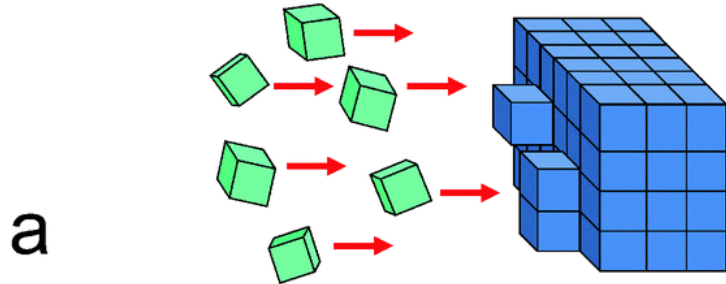
**POLYMERS**



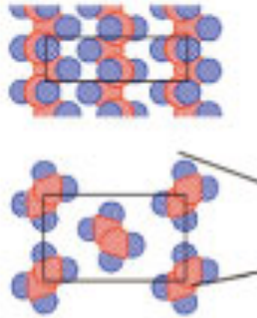
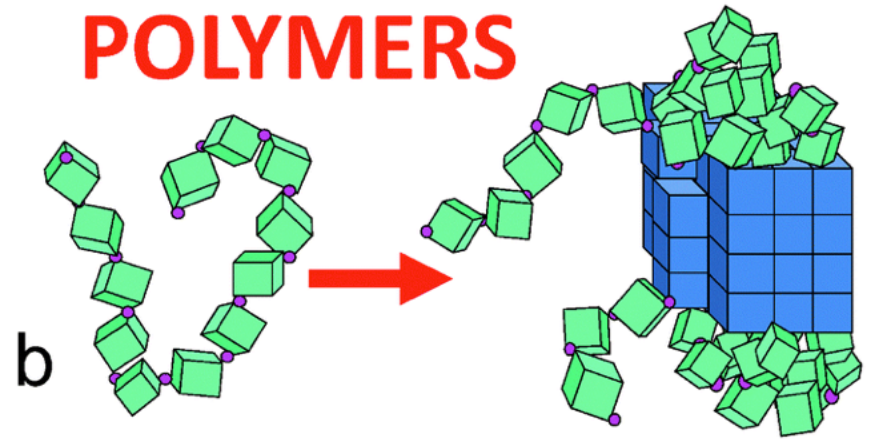
b

# Crystallisation in polymers

**SMALL MOLECULES**



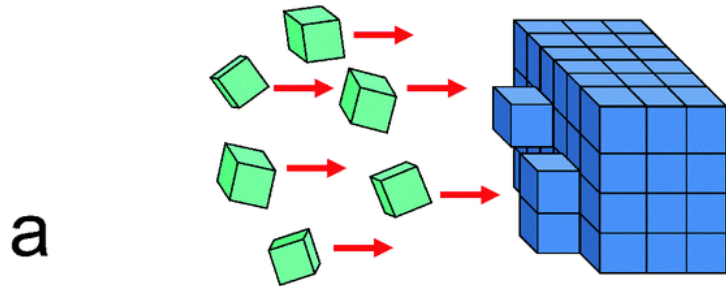
**POLYMERS**



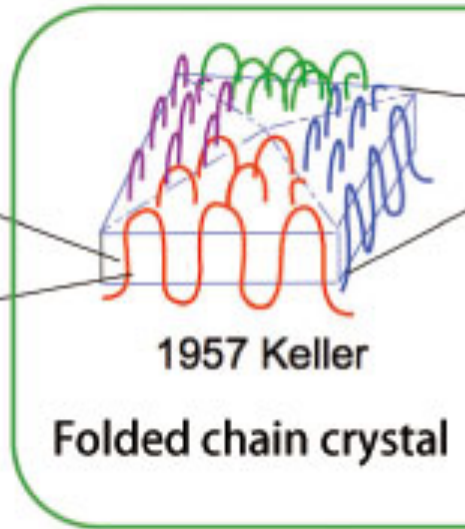
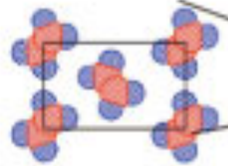
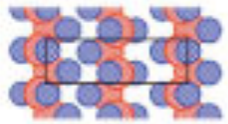
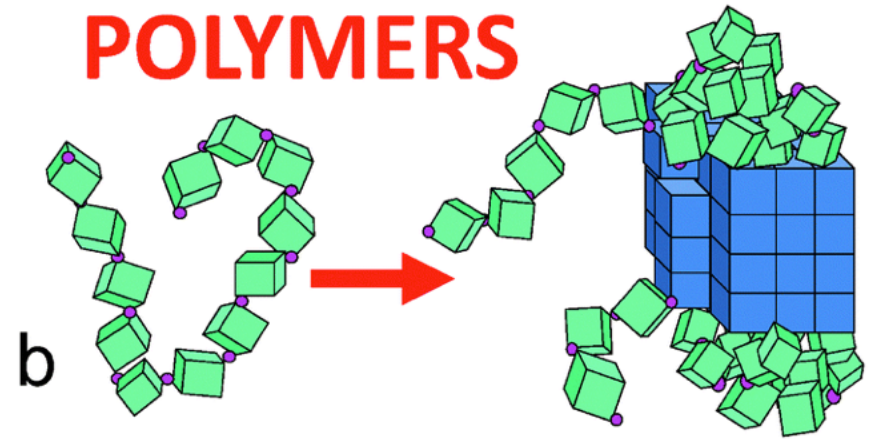
Crystalline structure

# Crystallisation in polymers

## SMALL MOLECULES



## POLYMERS



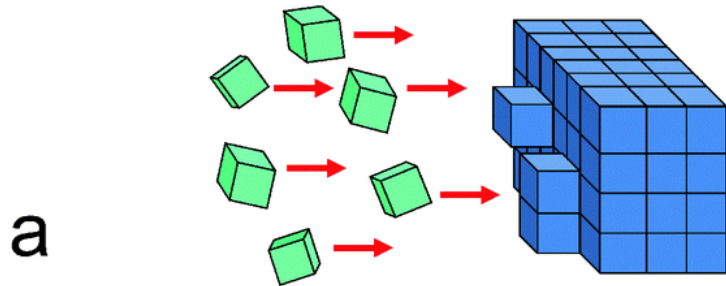
1957 Keller

Crystalline structure

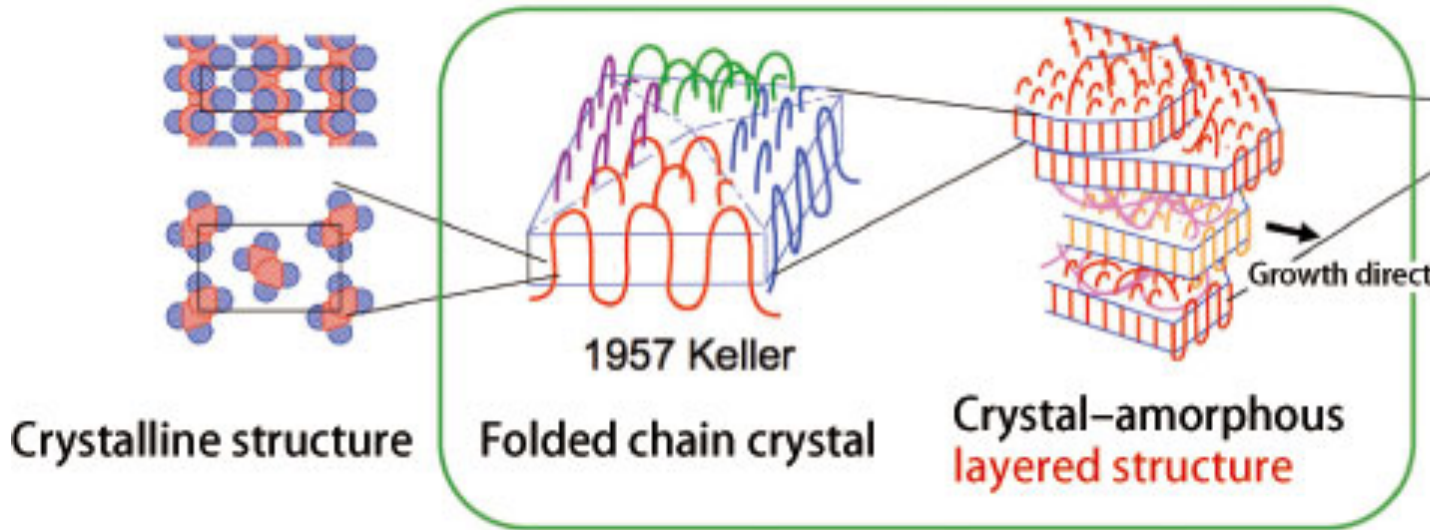
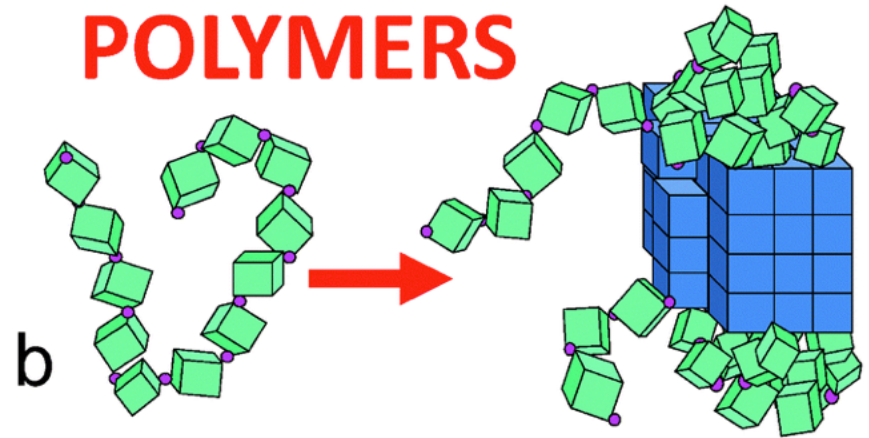
Folded chain crystal

# Crystallisation in polymers

## SMALL MOLECULES

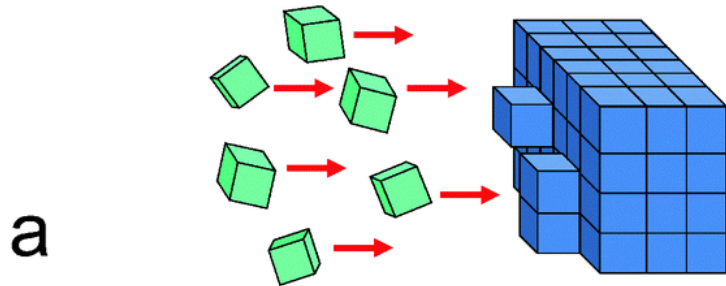


## POLYMERS

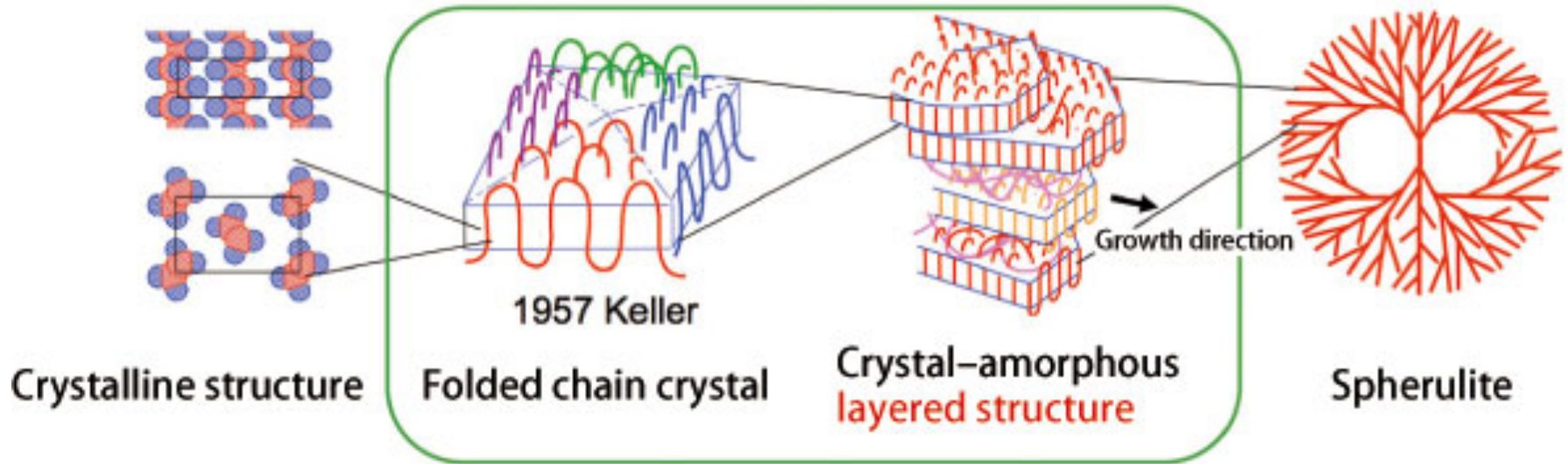
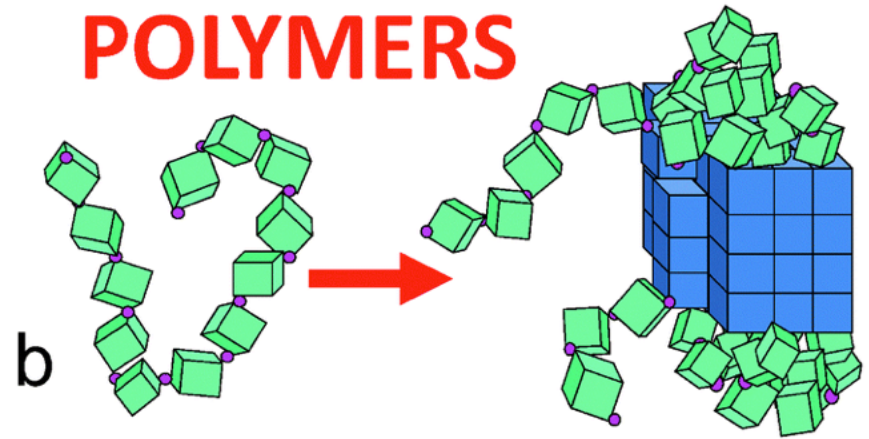


# Crystallisation in polymers

## SMALL MOLECULES



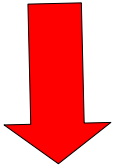
## POLYMERS





# Polymer crystallisation during flow

Deep slit ;  $T=130^{\circ}\text{C}$ ;  $\dot{\gamma}_{w,app.} \approx 100 \text{ s}^{-1}$

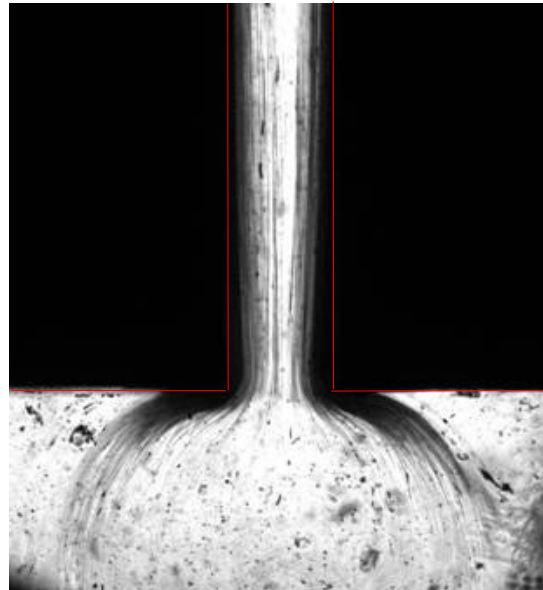
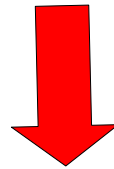
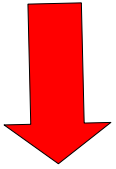


180kg/mol HDPE melt

Scelsi et al., Rheol Acta(2008).

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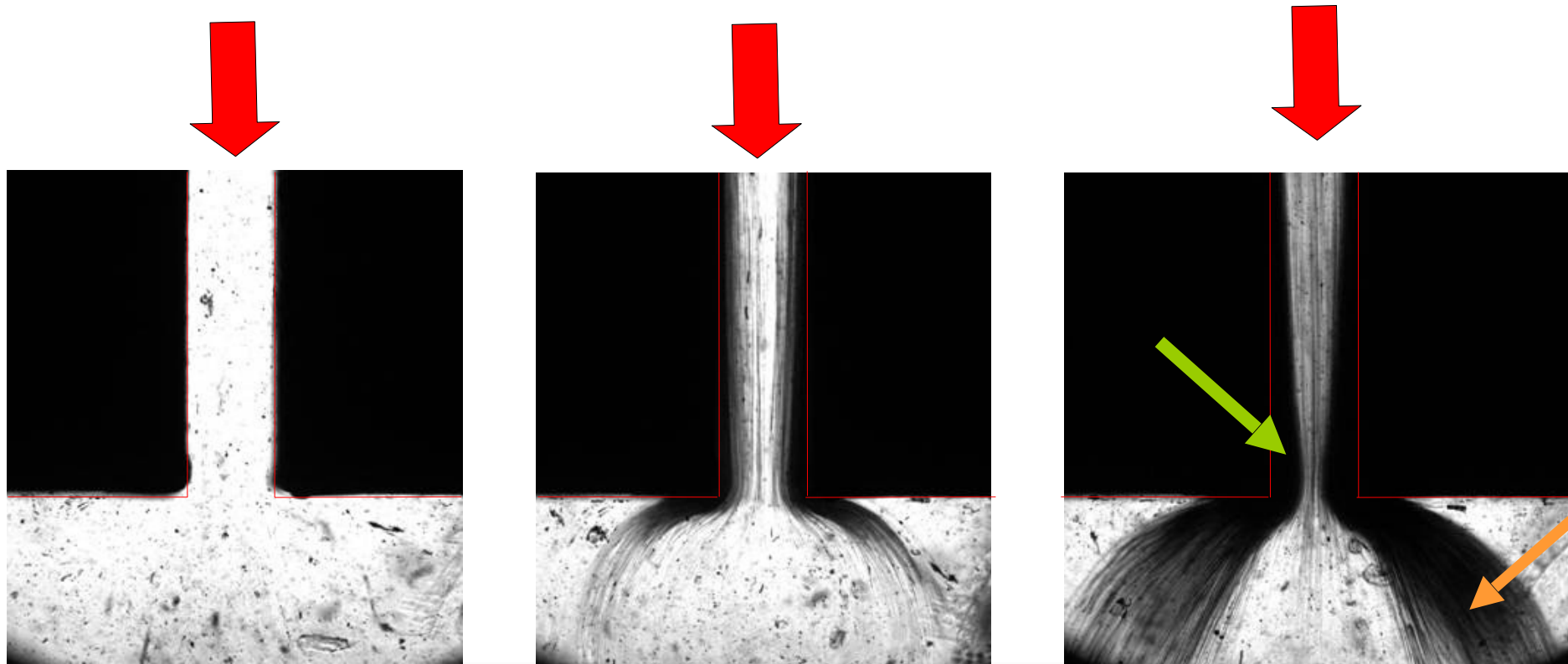


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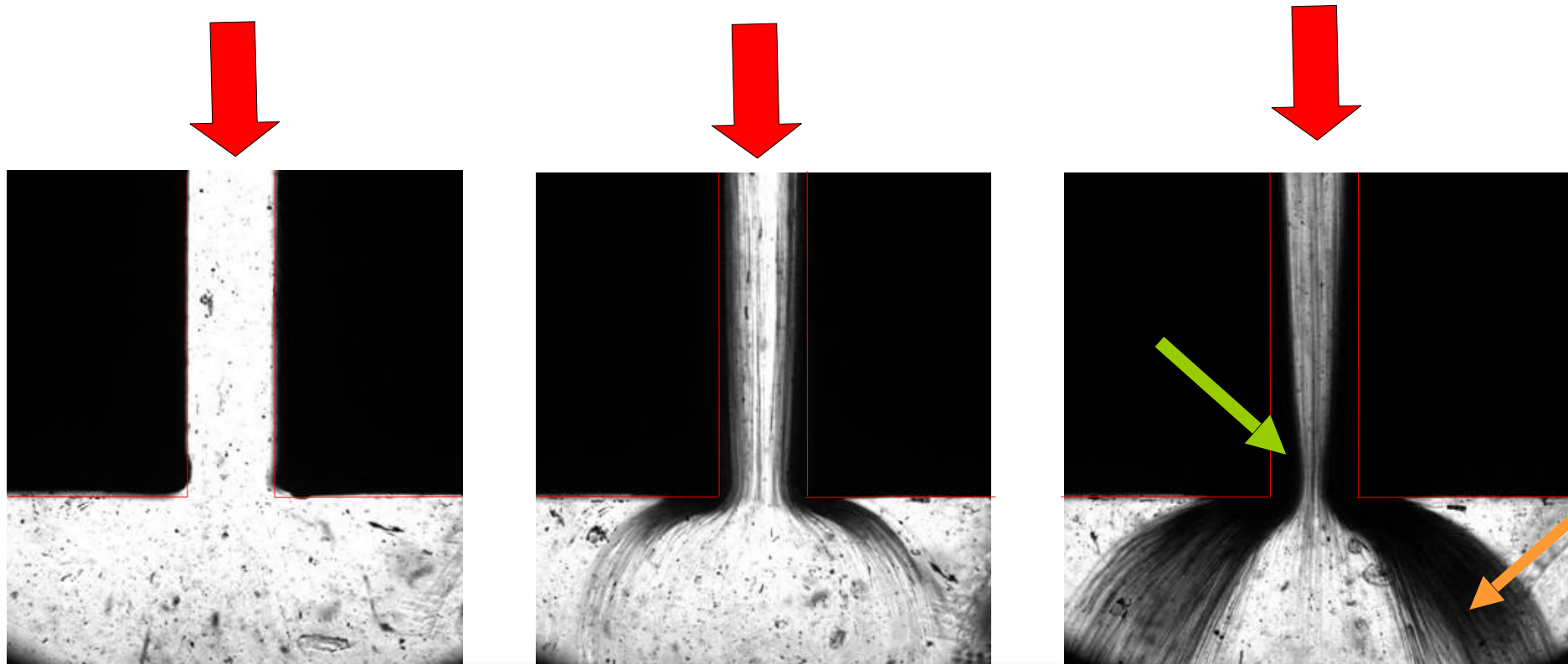
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Crystallisation is enhanced in regions of strong flow

# Polymer crystallisation during flow

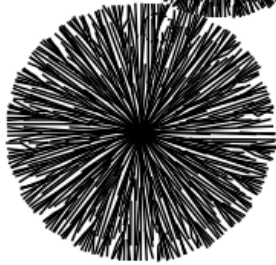
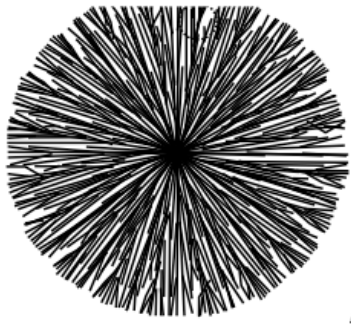
Deep slit ;  $T=130^{\circ}\text{C}$ ;  $\dot{\gamma}_{w,app.} \approx 100 \text{ s}^{-1}$



Crystallisation is enhanced in regions of strong flow

Key processing control variables are  
temperature, flow-rate and molecular weight

# Effect of flow

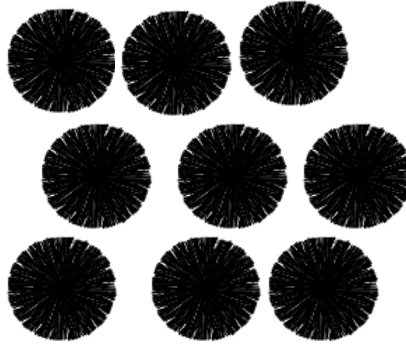
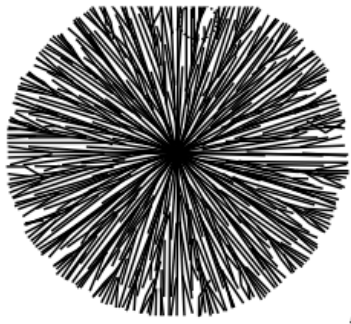


Quiescent  
(no flow)



Increasing flow rate

# Effect of flow



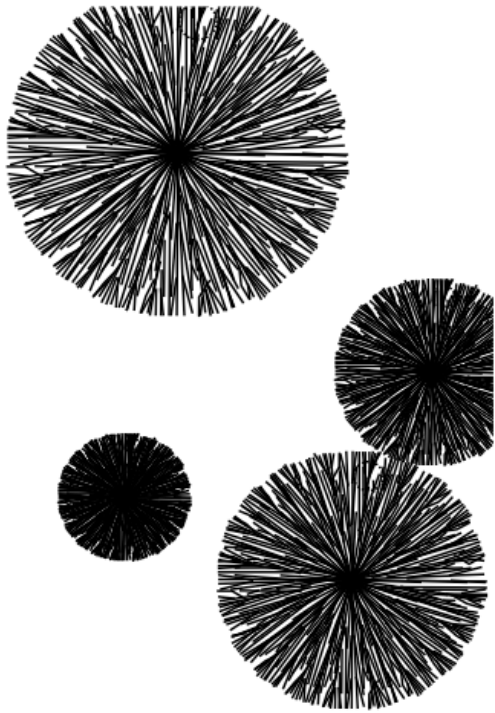
Quiescent  
(no flow)

Enhanced  
nucleation

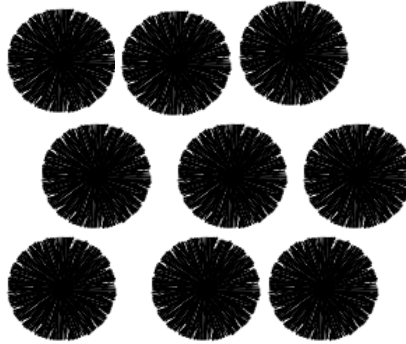
Increasing flow rate



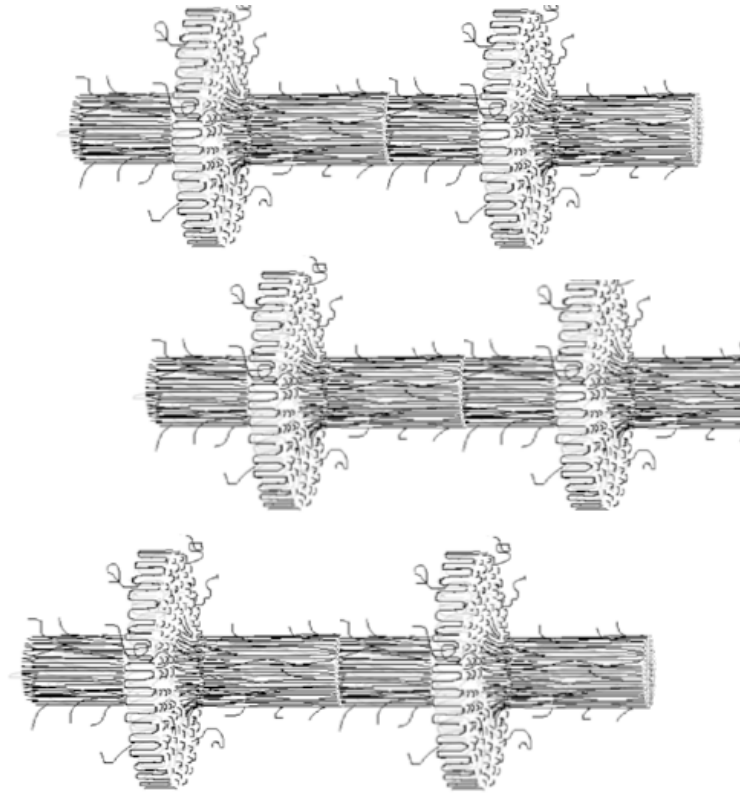
# Effect of flow



Quiescent  
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Enhanced  
nucleation

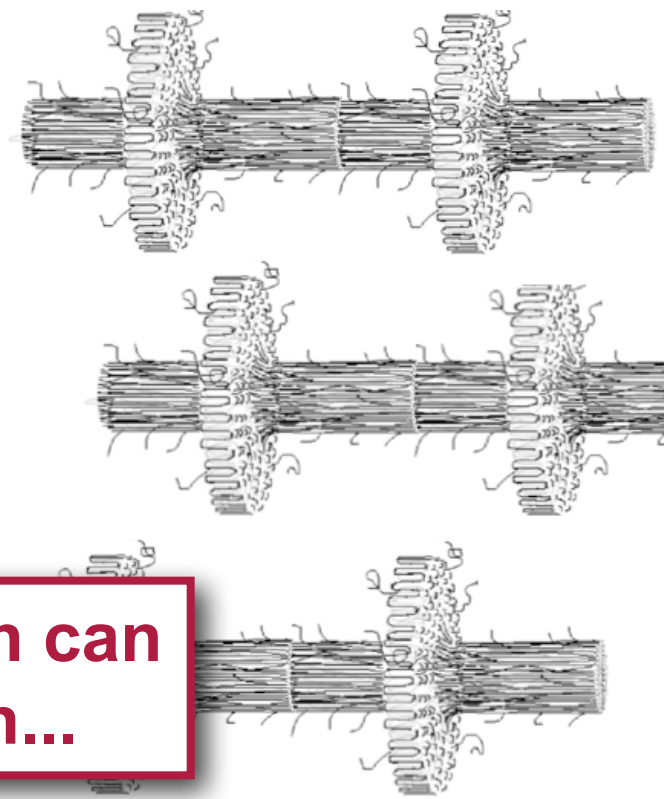
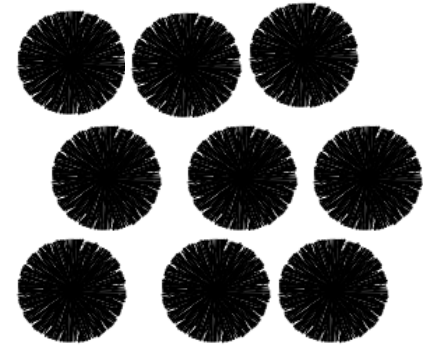
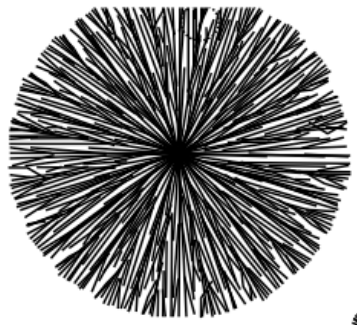


Shish  
kebabs

Increasing flow rate



# Effect of flow



**Controlling crystallisation can be the difference between...**

Quiescent  
(no flow)

Enhanced  
nucleation

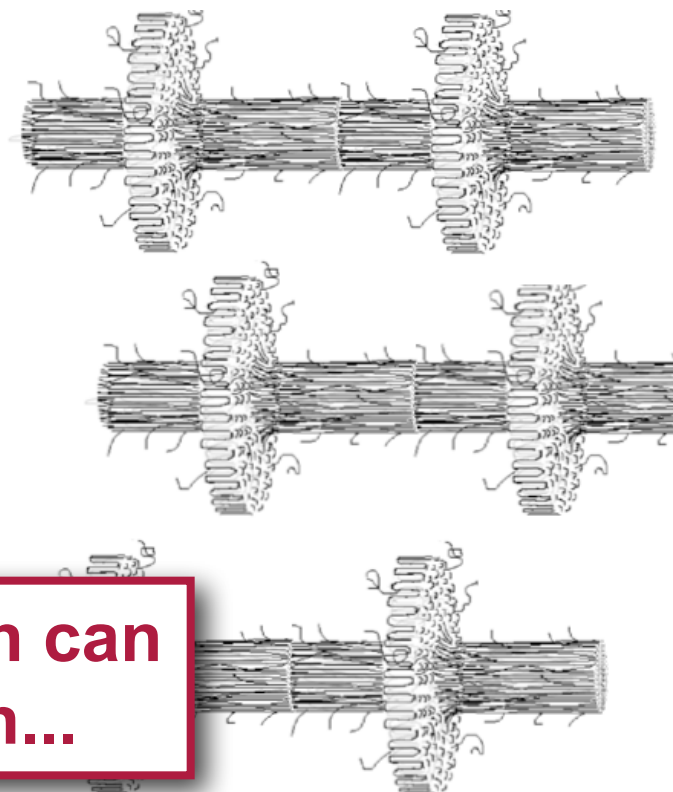
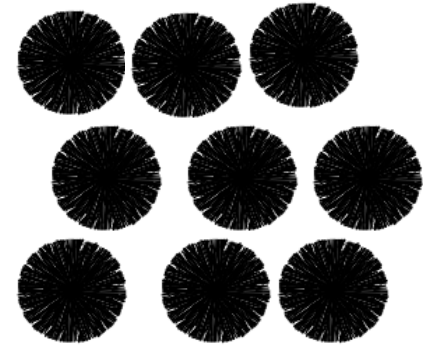
Shish  
kebabs



Increasing flow rate



# Effect of flow



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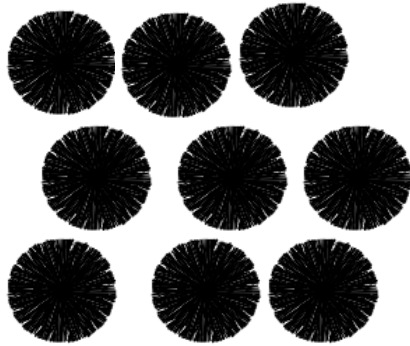
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Increasing flow rate

# Effect of flow



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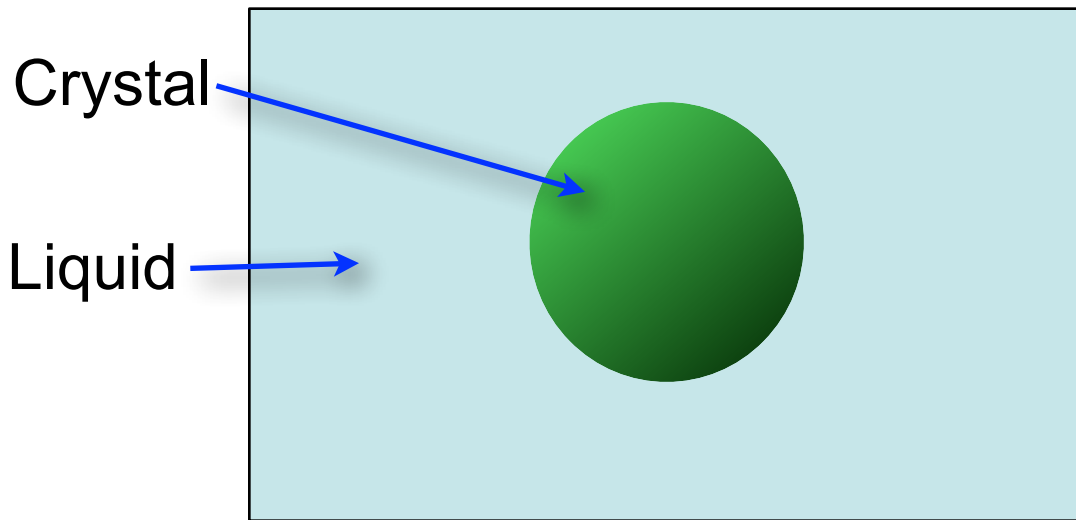
Enhanced  
nucleation

Shish  
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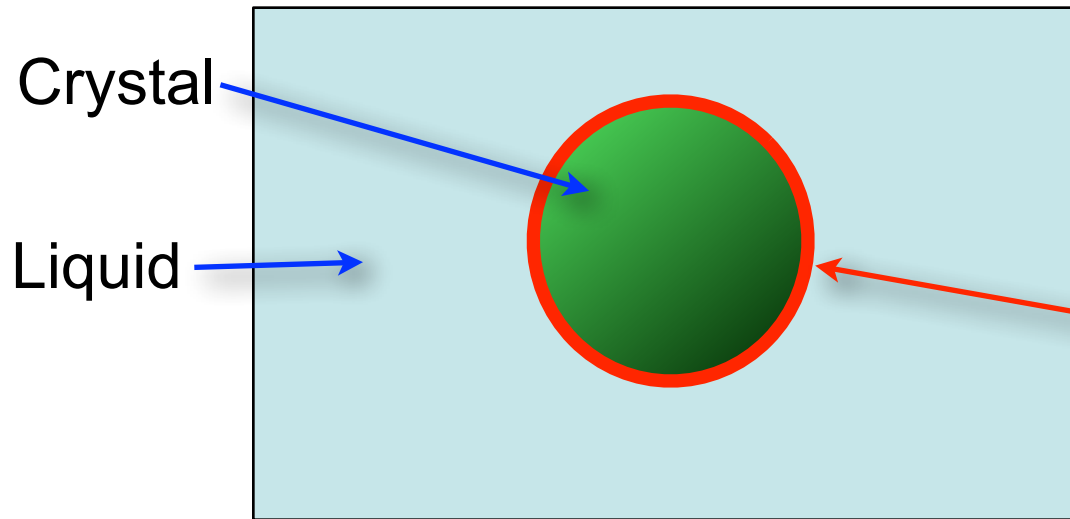
Increasing flow rate

# Nucleation



Bulk crystallisation  
lowers the free  
energy

# Nucleation



Bulk crystallisation lowers the free energy  
... but the interface has a cost.

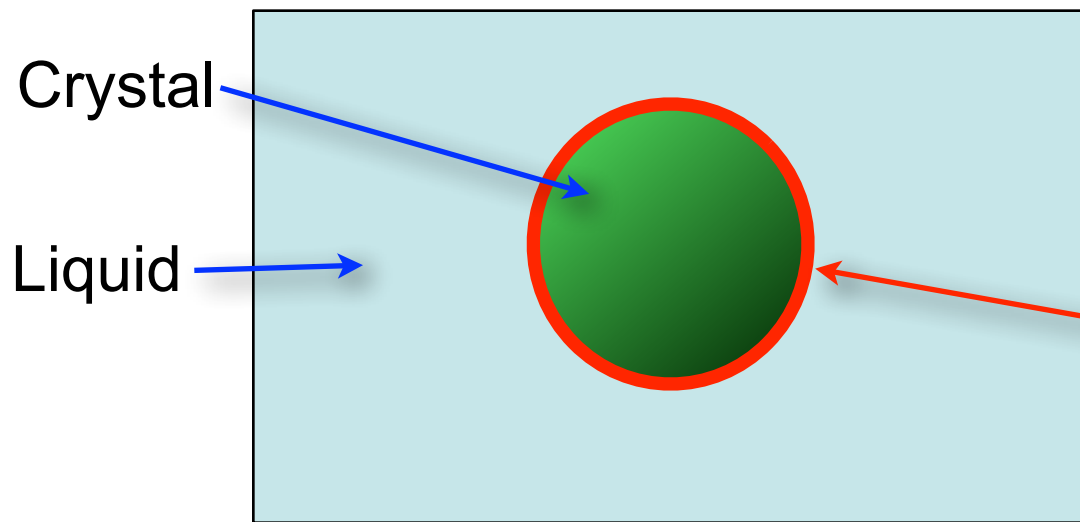
$$F = \underbrace{AR^2}_{\text{Surface cost}} - \underbrace{BR^3}_{\text{Bulk gain}}$$

Surface cost

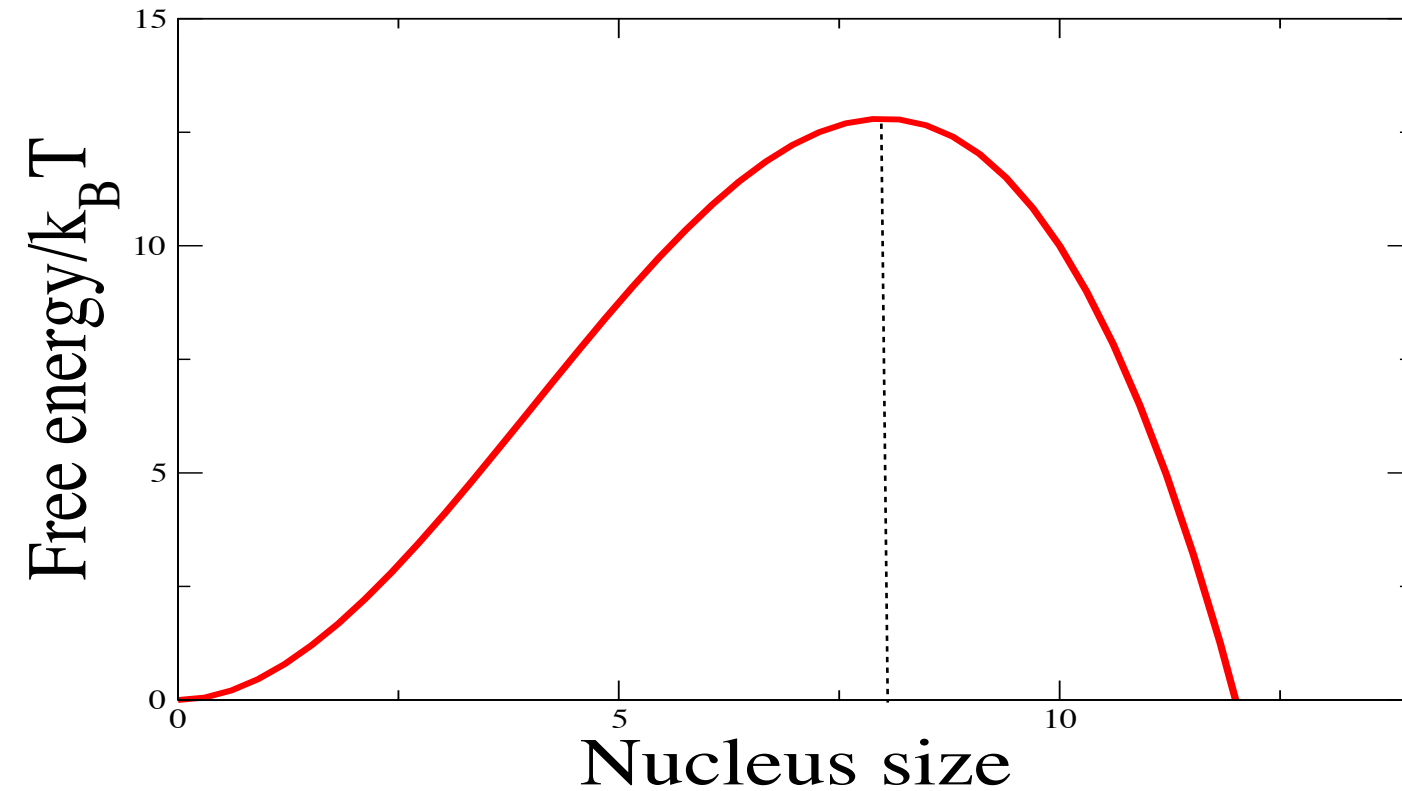
F= Free energy

R= Nucleus  
radius

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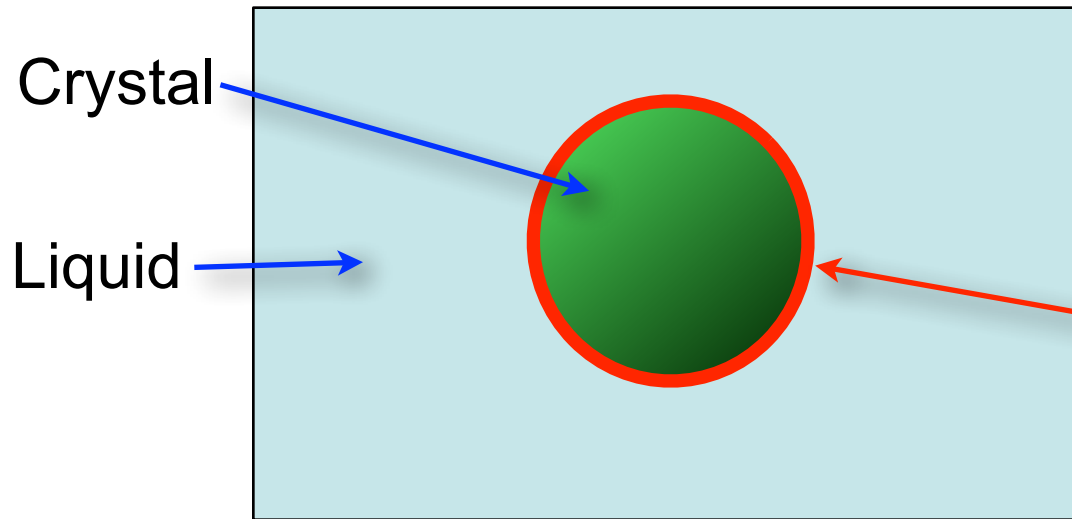
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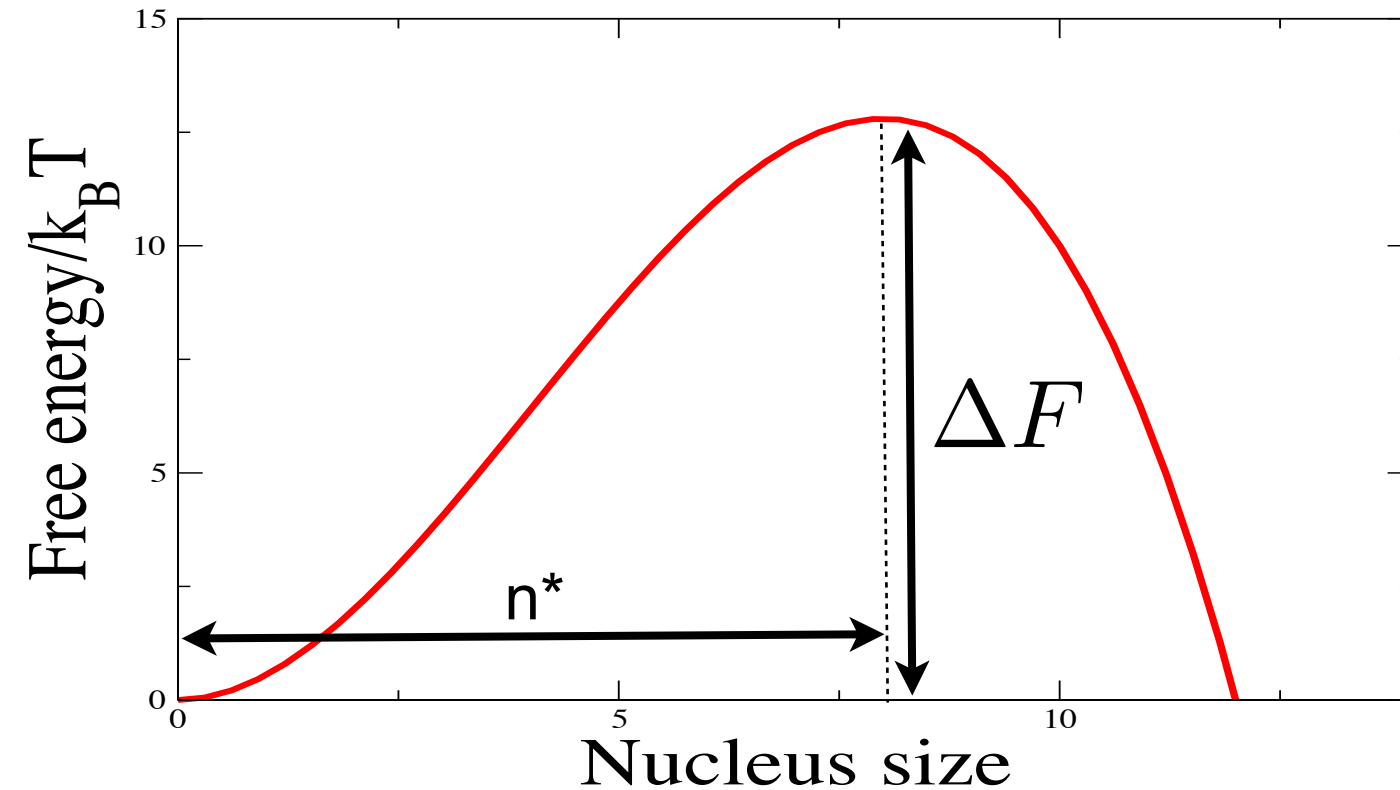
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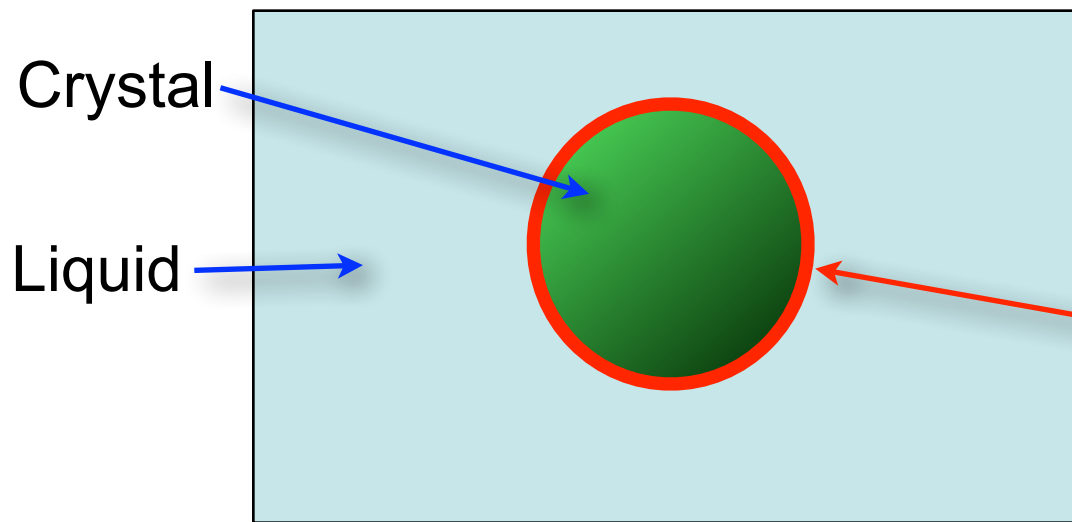
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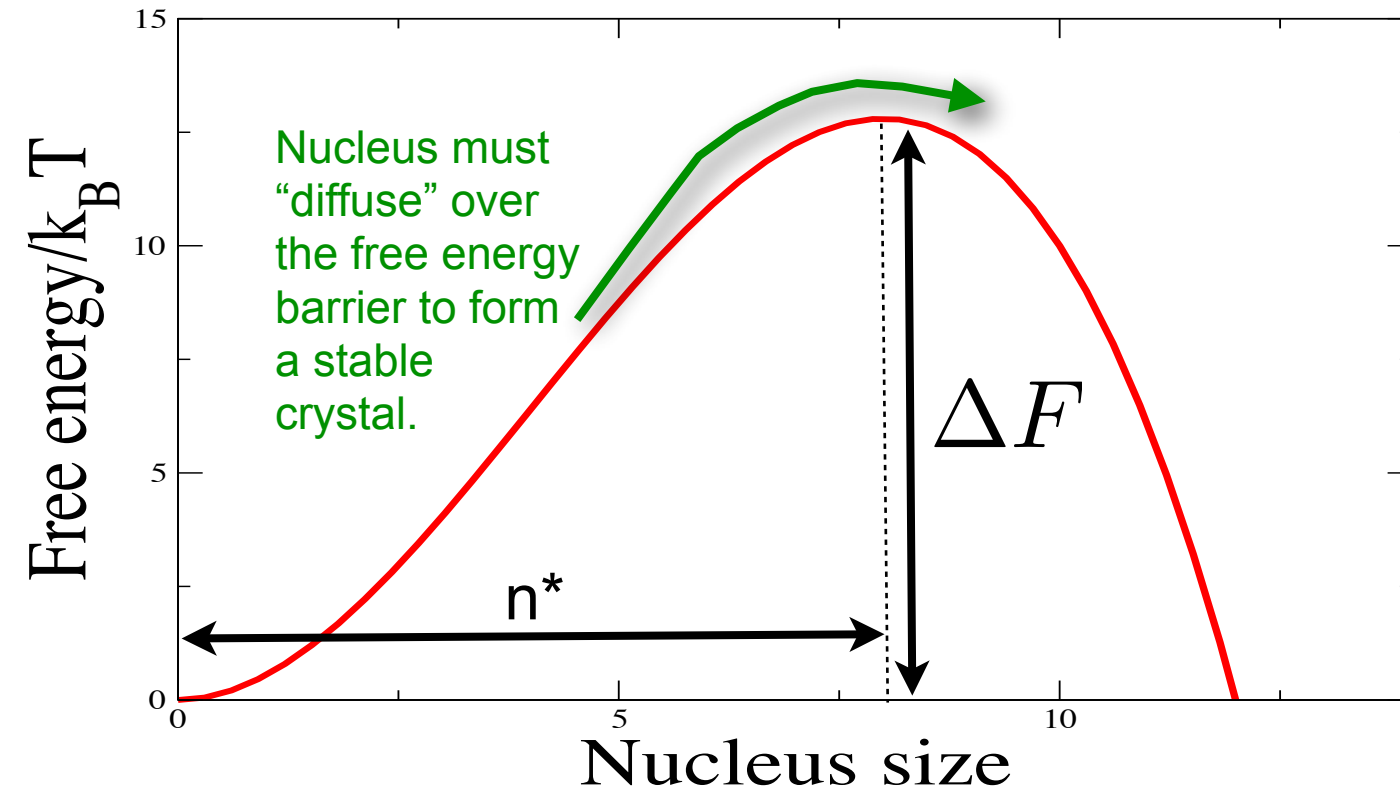
F= Free energy

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# Nucleation



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Bulk gain

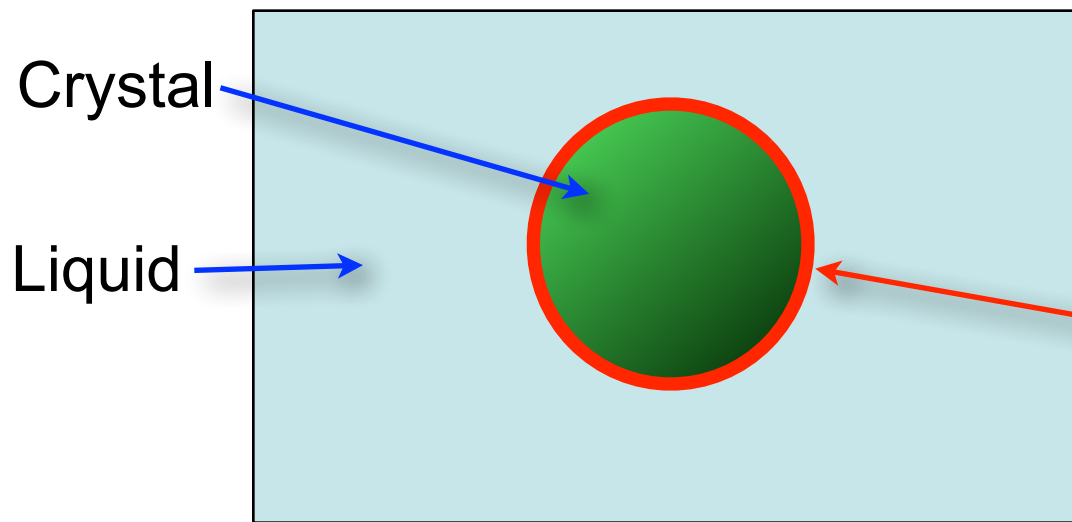
$$F = \underbrace{AR^2}_{\text{Surface cost}} - \underbrace{BR^3}_{\text{Bulk gain}}$$

Surface cost

F= Free energy

R= Nucleus radius

# Nucleation



Bulk crystallisation lowers the free energy  
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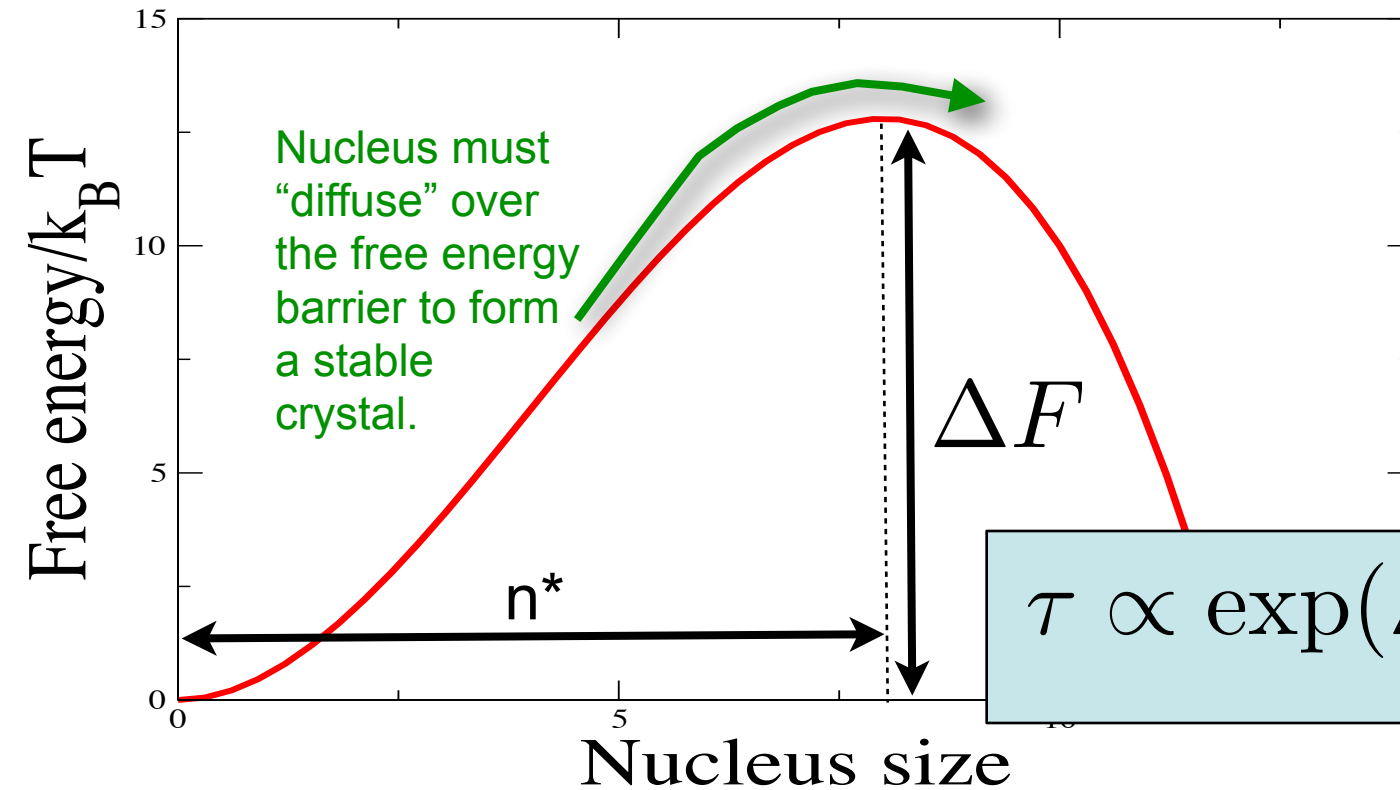
Bulk gain

$$F = \underbrace{AR^2}_{\text{Surface cost}} - \underbrace{BR^3}_{\text{Bulk gain}}$$

Surface cost

F= Free energy

R= Nucleus radius



$$\tau \propto \exp(\Delta F / k_B T)$$



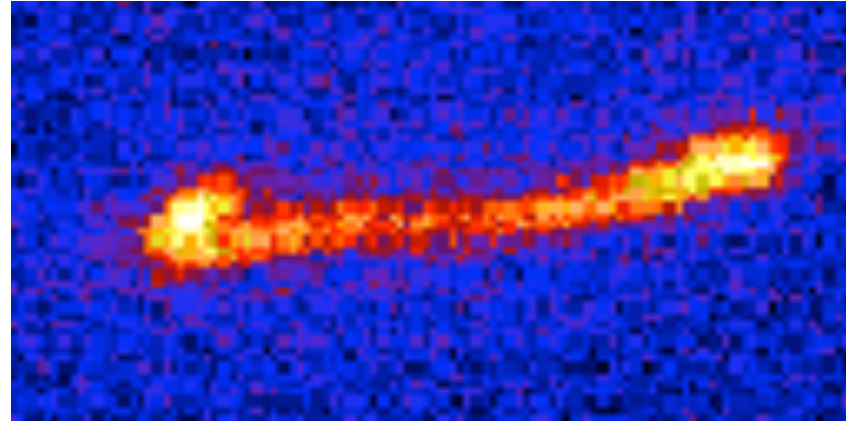
How rare is nucleation?



How rare is nucleation?

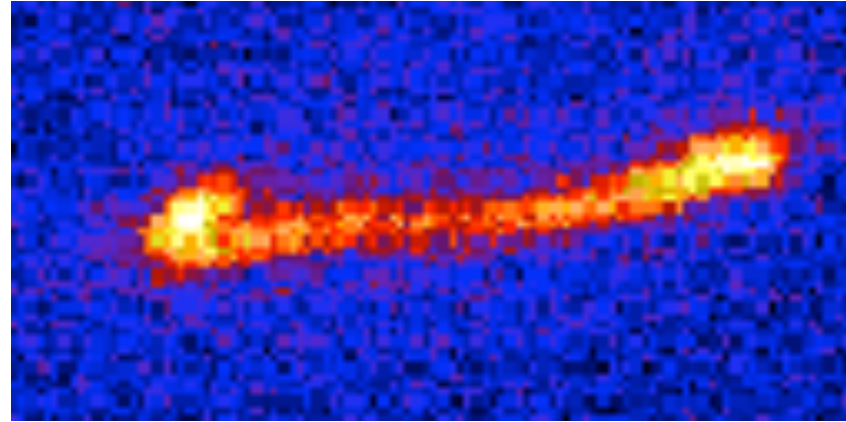


# Polymer nucleation during flow



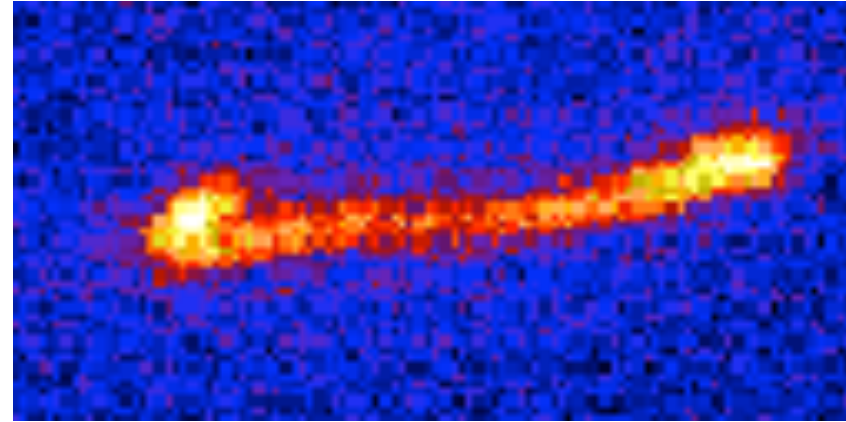
Experiments by Teixeira et al *Macromolecules*  
(2005) vol. 38 (2) pp. 581-592

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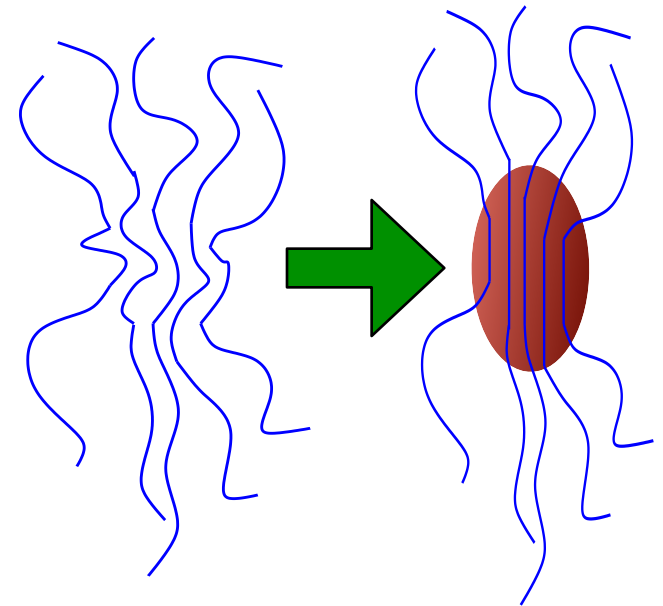


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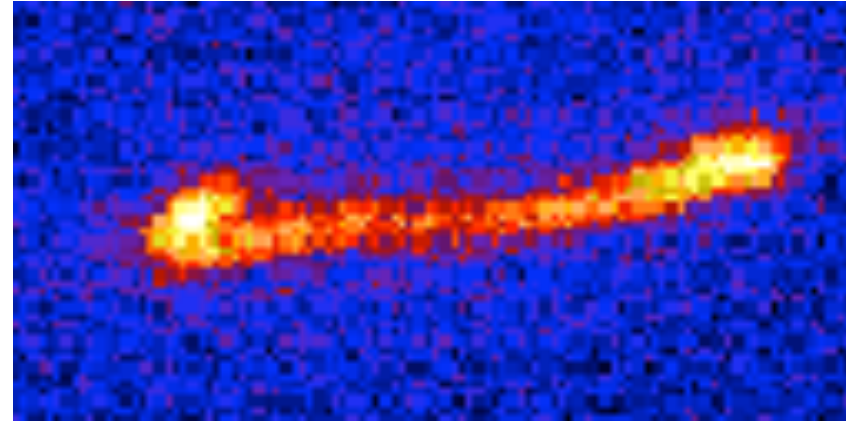
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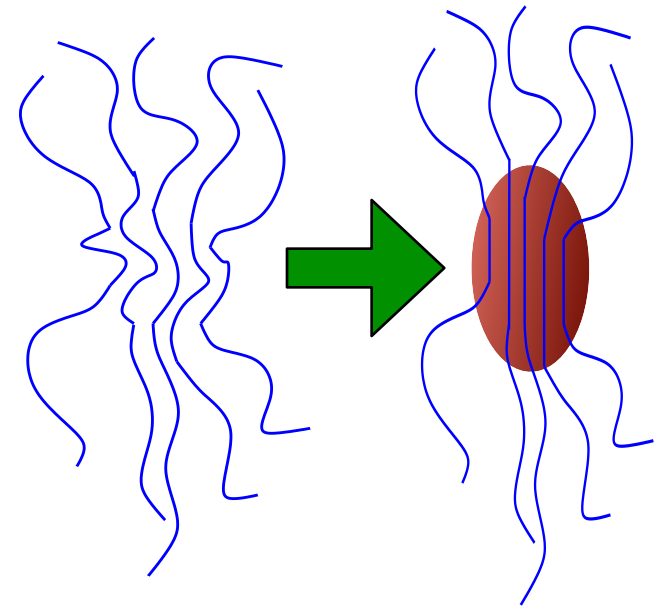
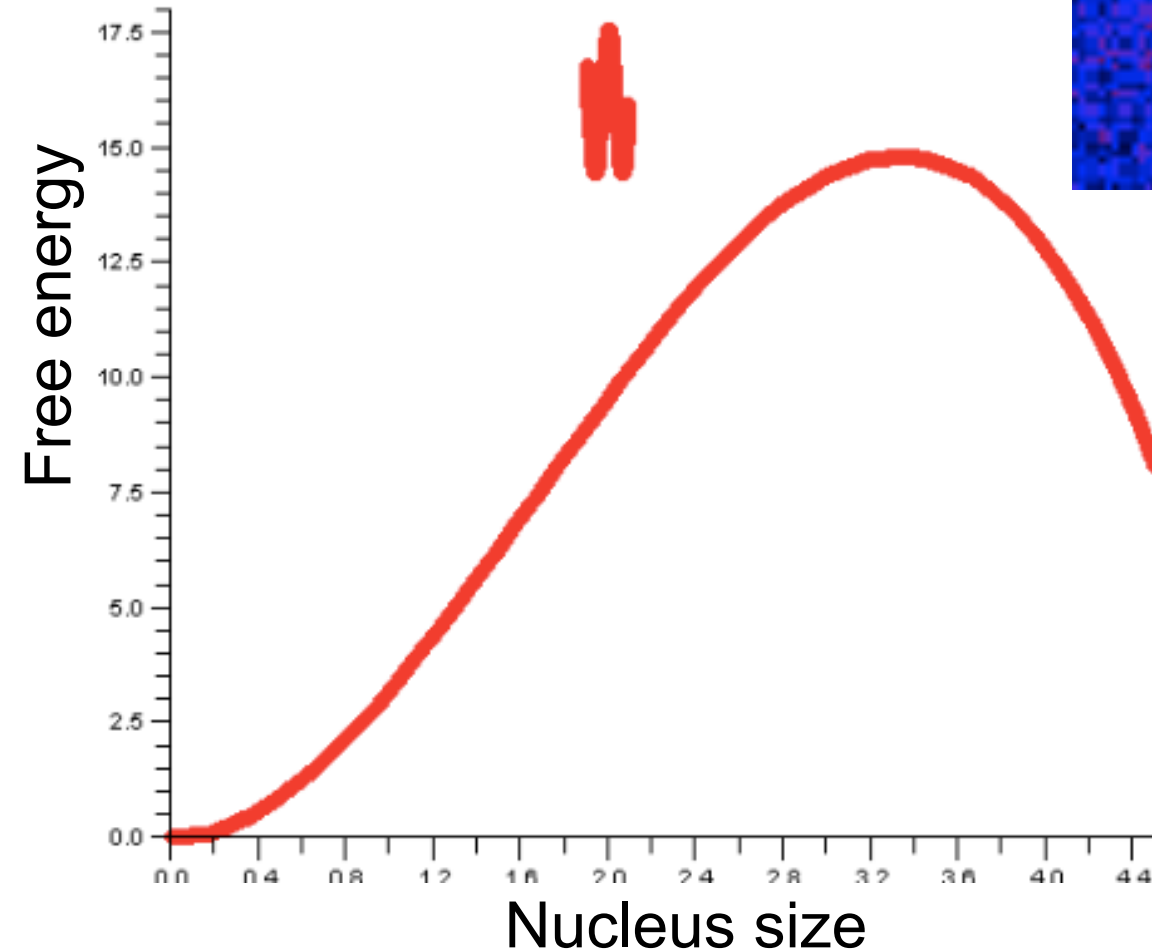


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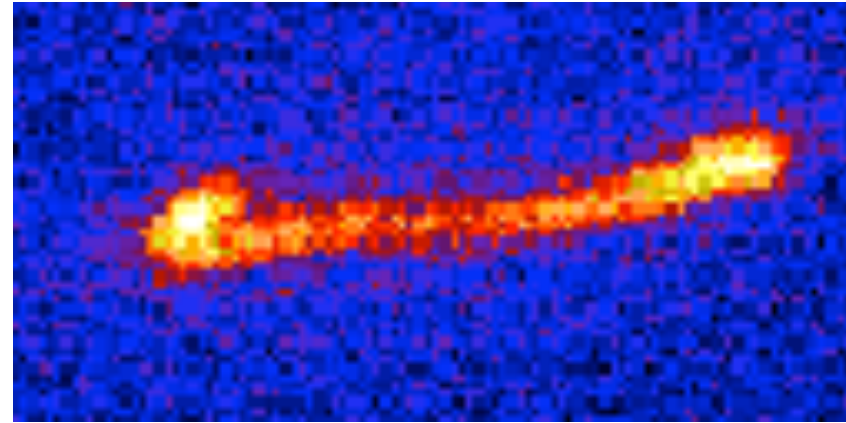


Experiments by Teixeira et al *Macromolecules* (2005) vol. 38 (2) pp. 581-592

$t=0.$

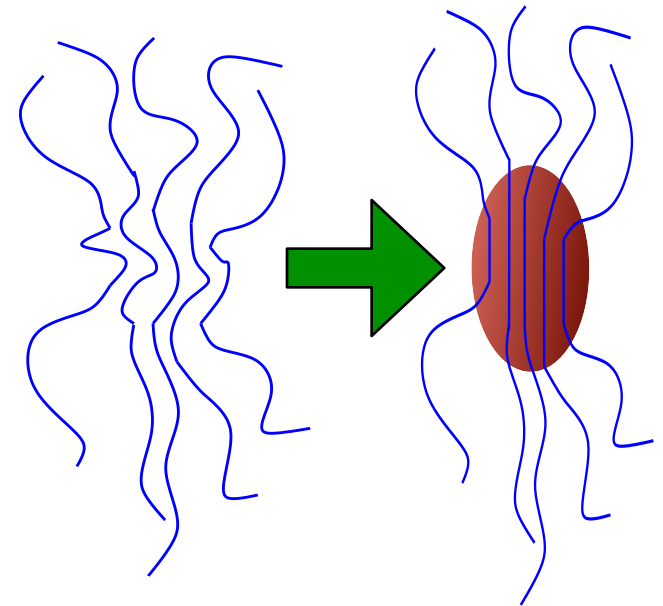
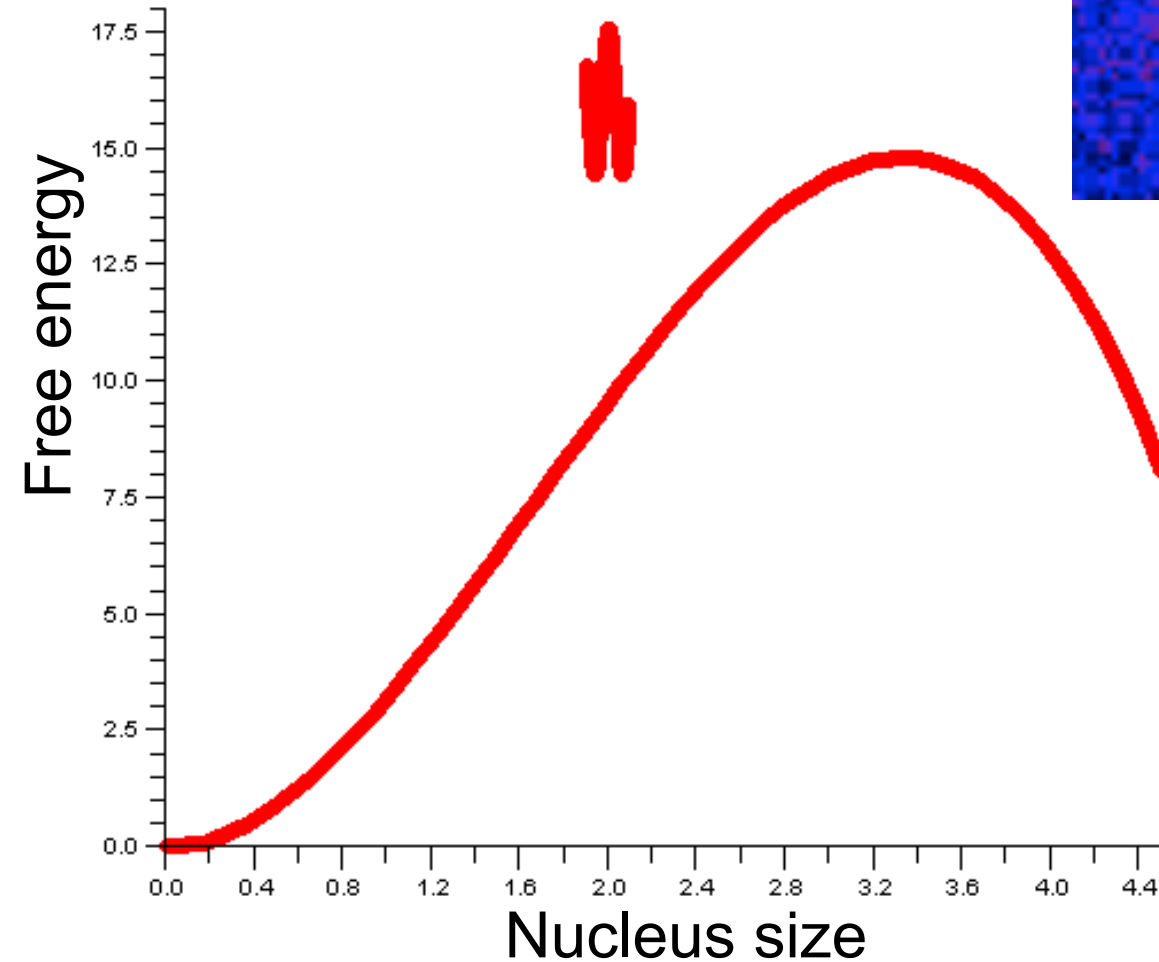


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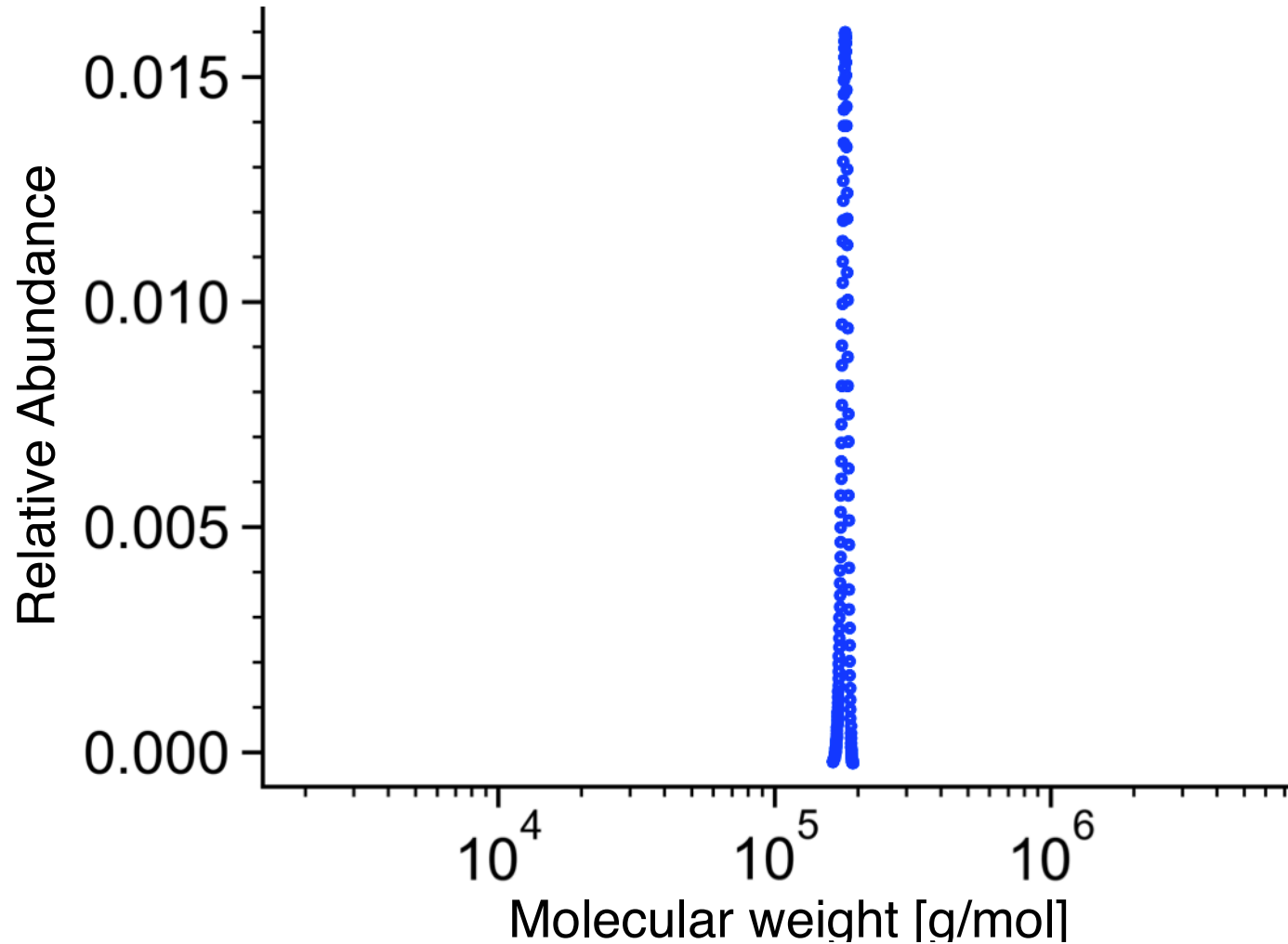
Experiments by Teixeira et al *Macromolecules* (2005) vol. 38 (2) pp. 581-592

$t = 0.$



# Polydispersity

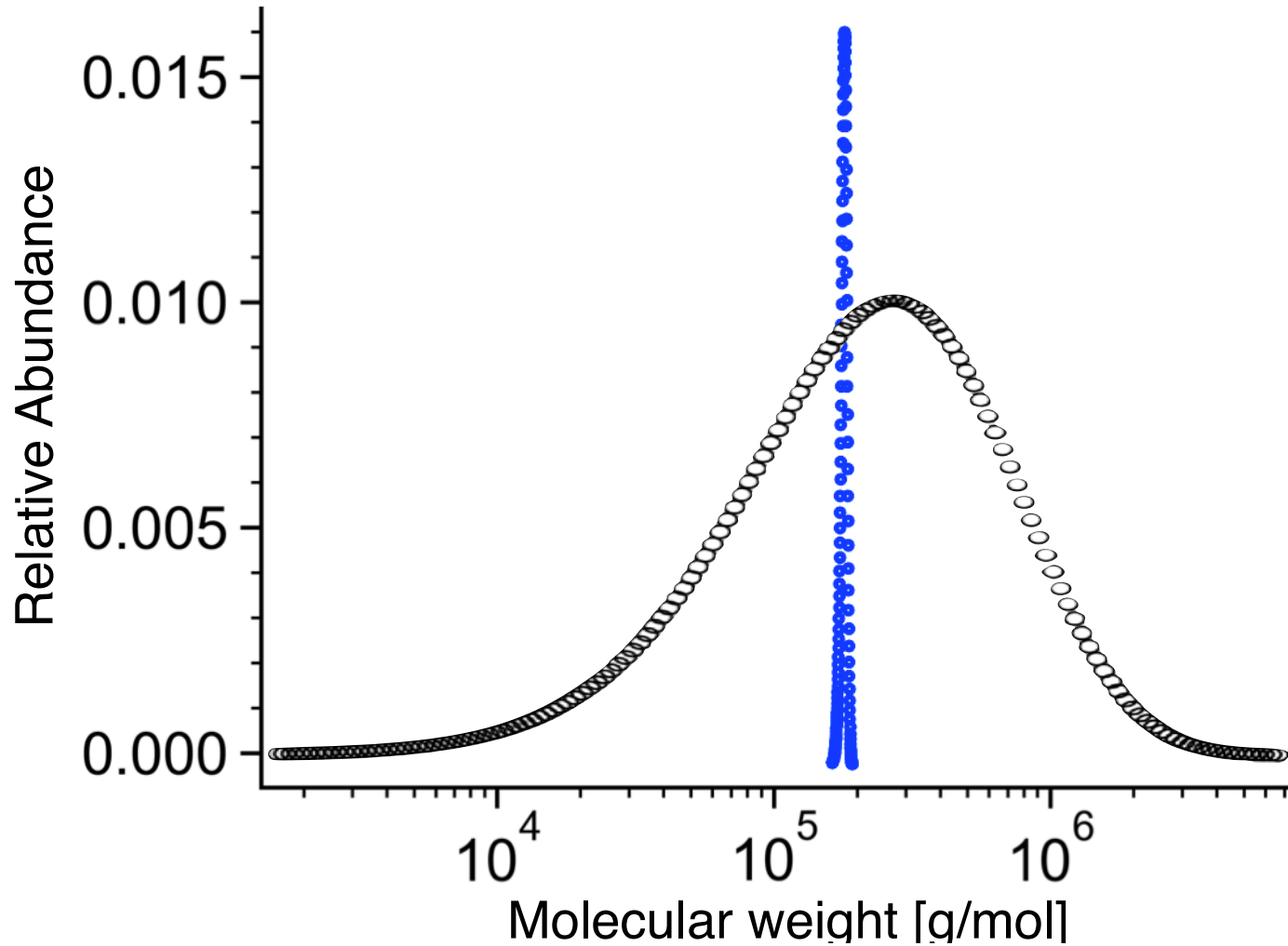
Model Polymers  
Real Polymers





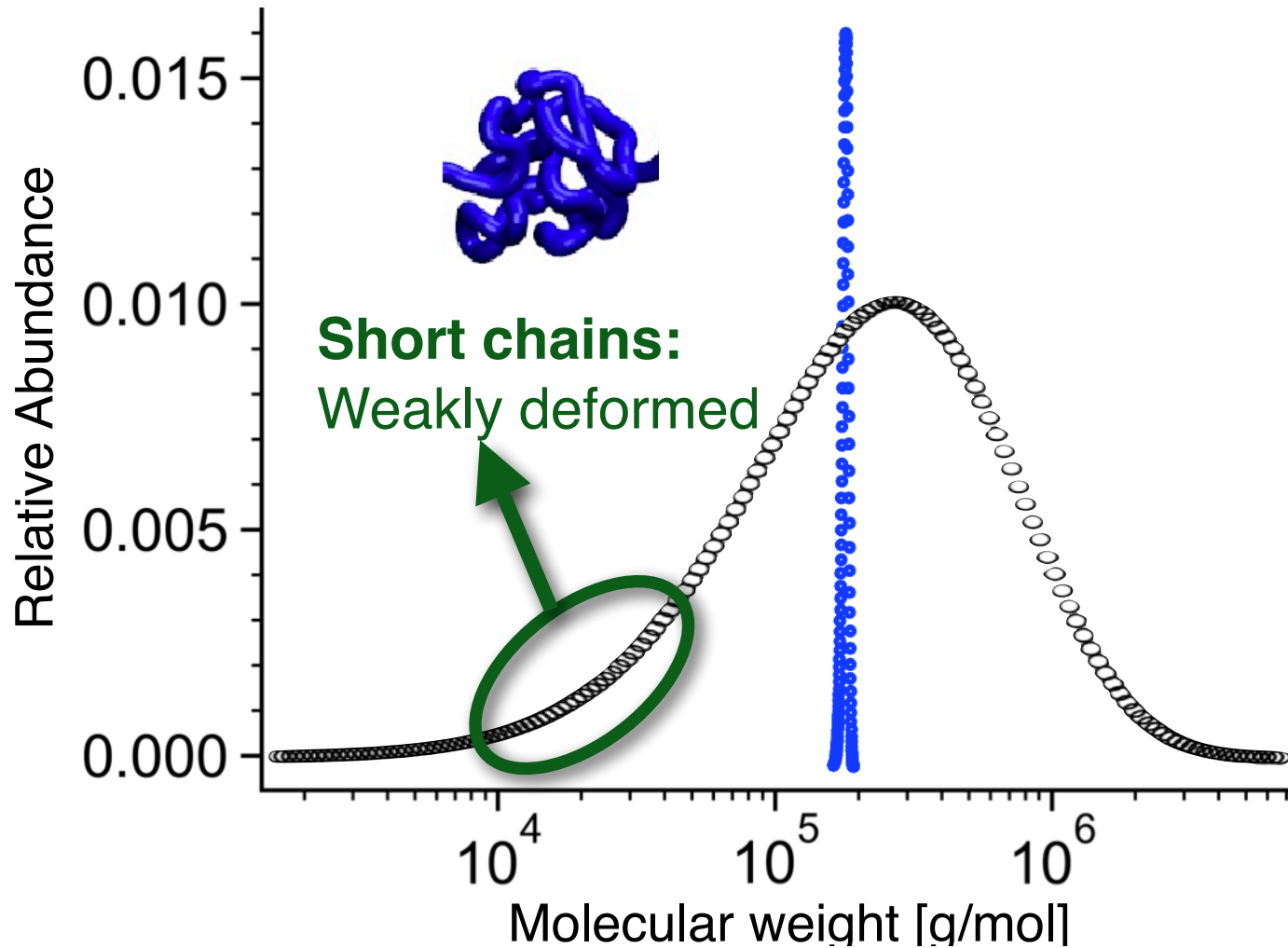
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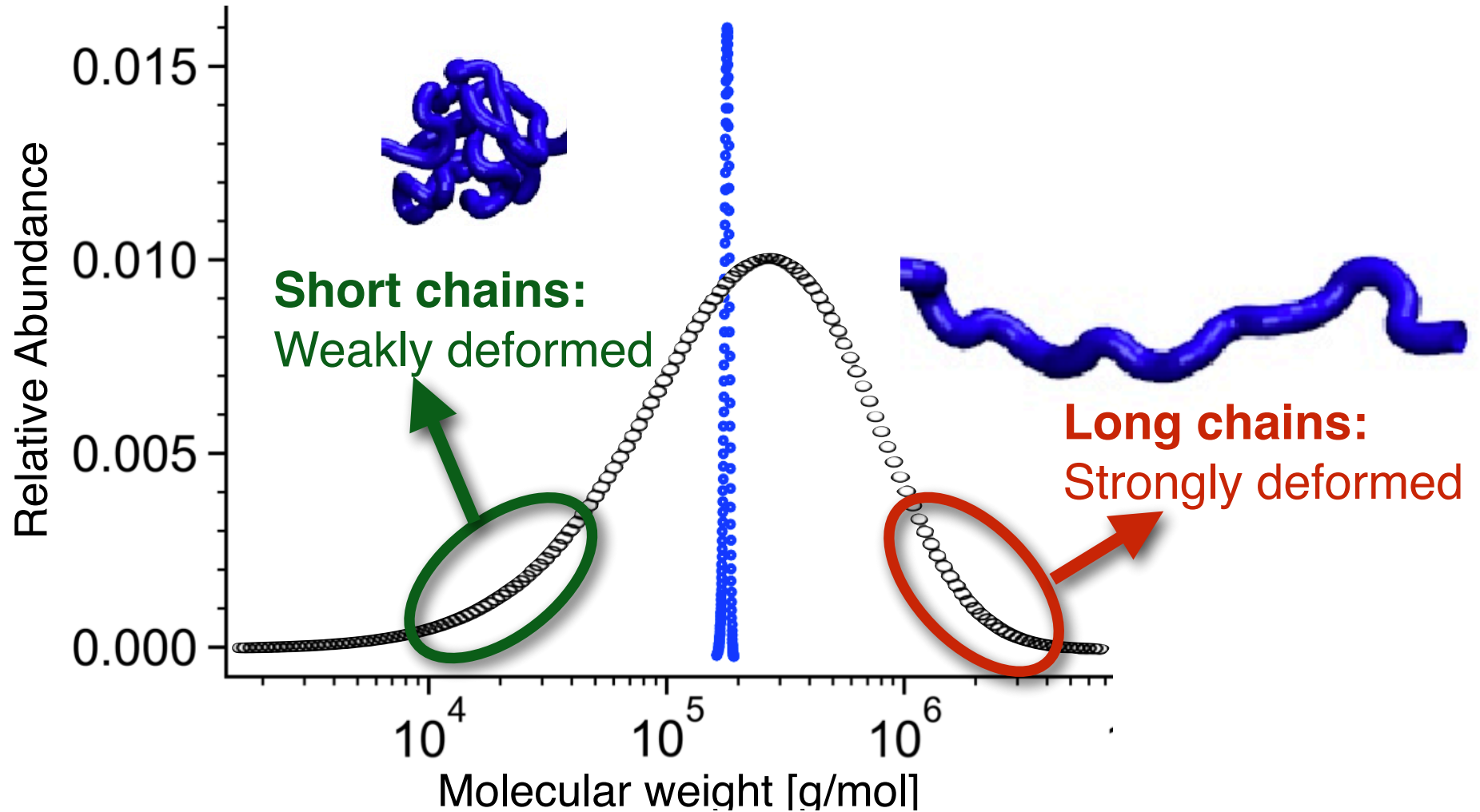
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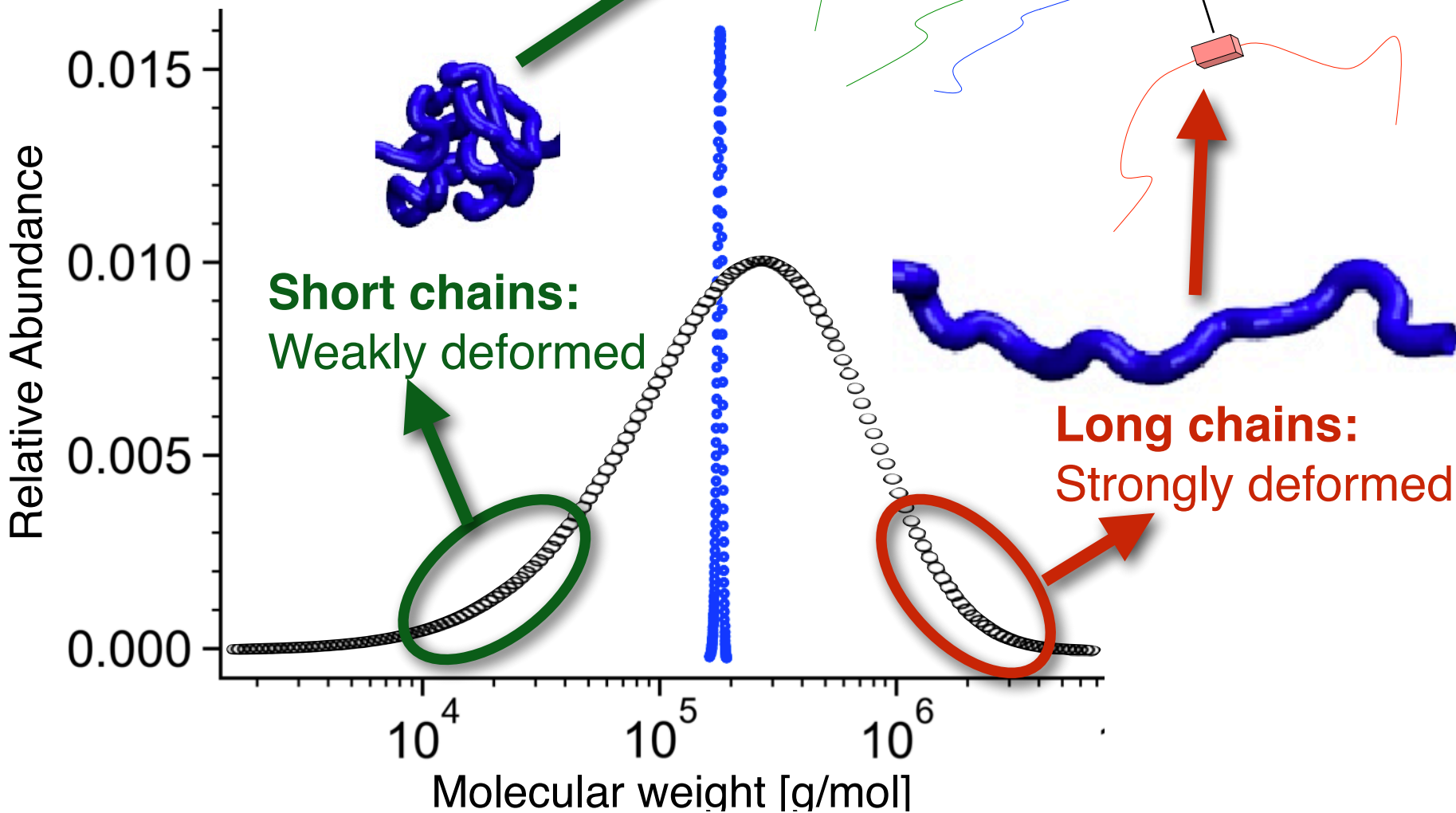
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# Polydispersity

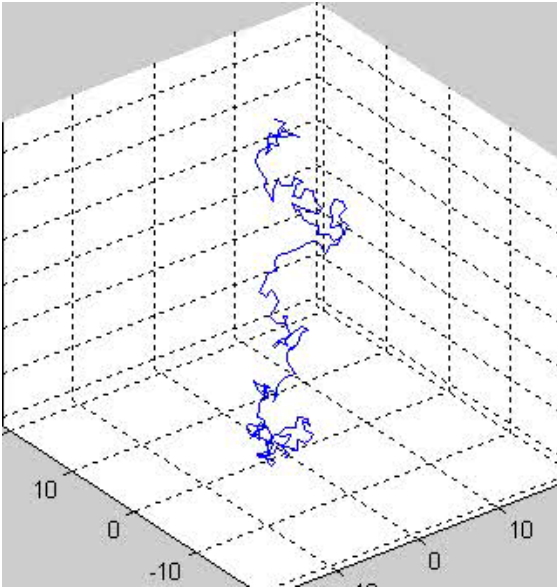
Model Polymers  
Real Polymers



# Two computational difficulties

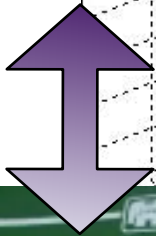
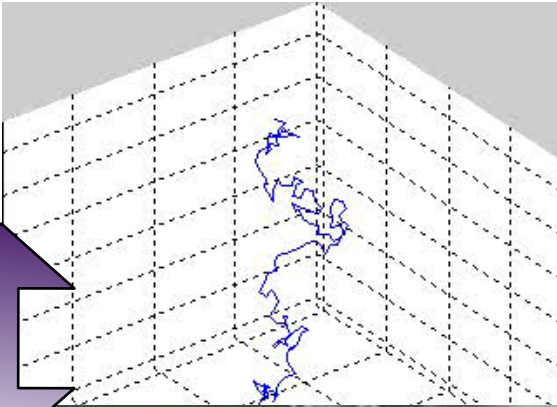
# Two computational difficulties

## Slow diffusion



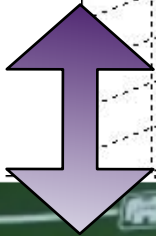
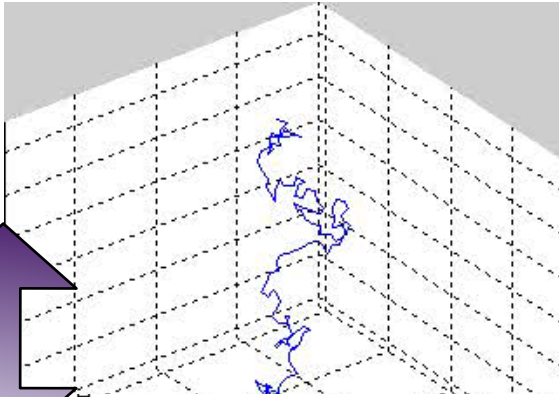
# Two computational difficulties

Slow diffusion



# Two computational difficulties

Slow diffusion



Nucleation

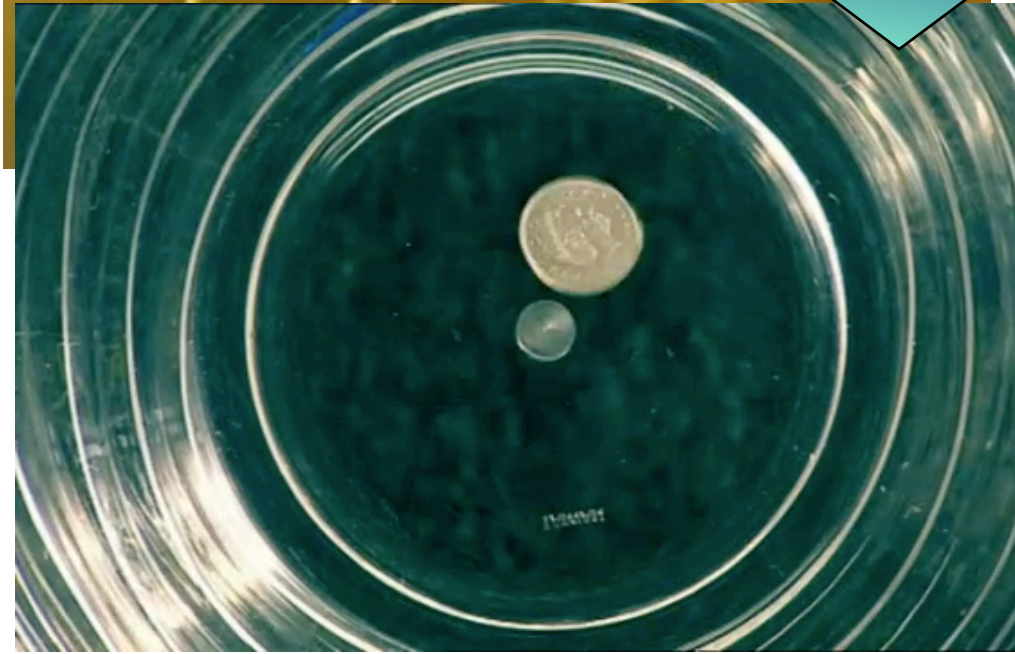
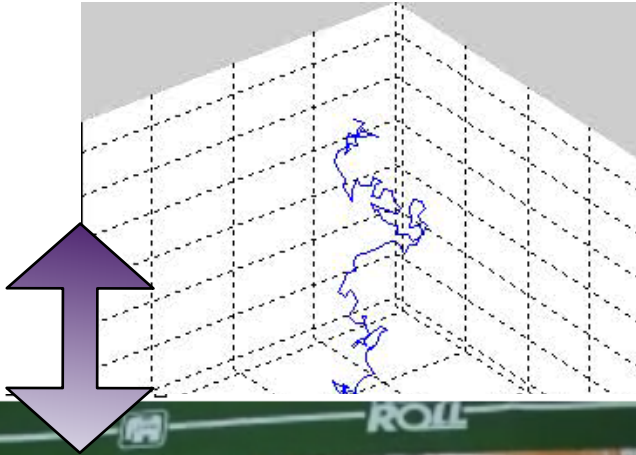




# Two computational difficulties

Slow diffusion

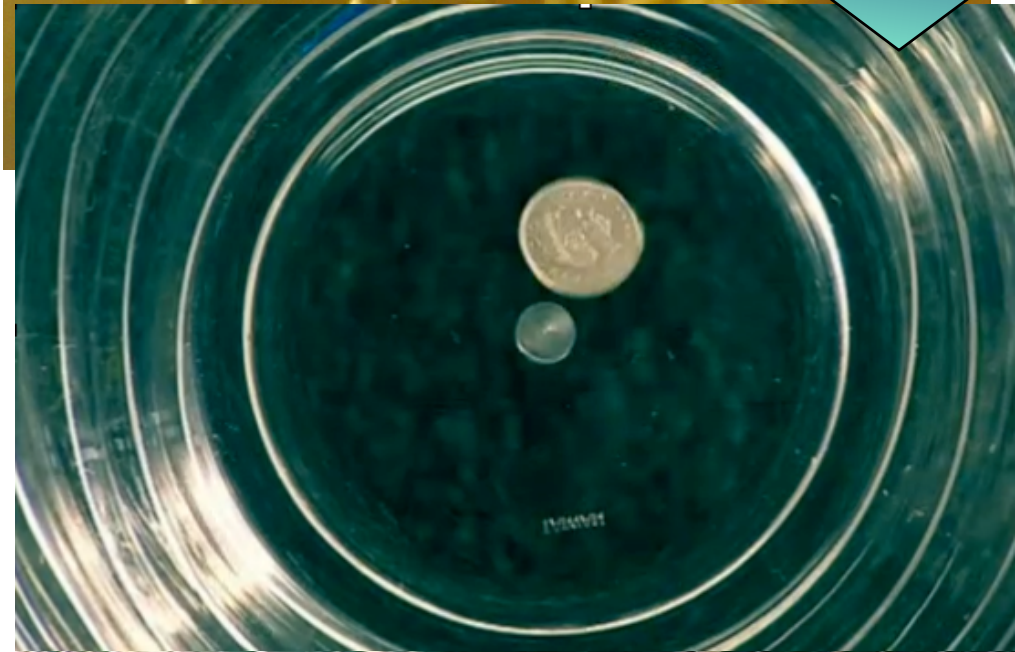
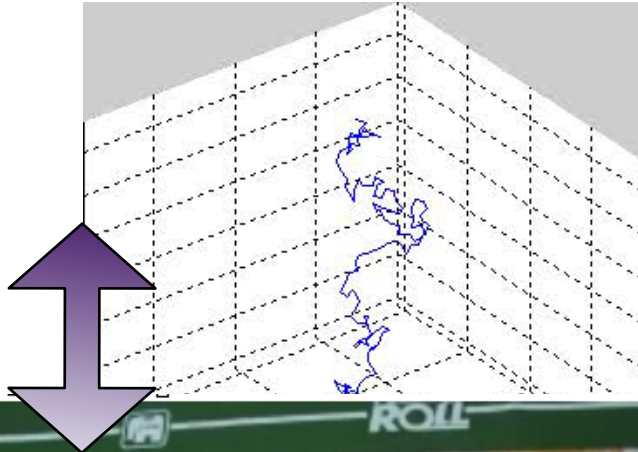
Nucleation



# Two computational difficulties

Slow diffusion

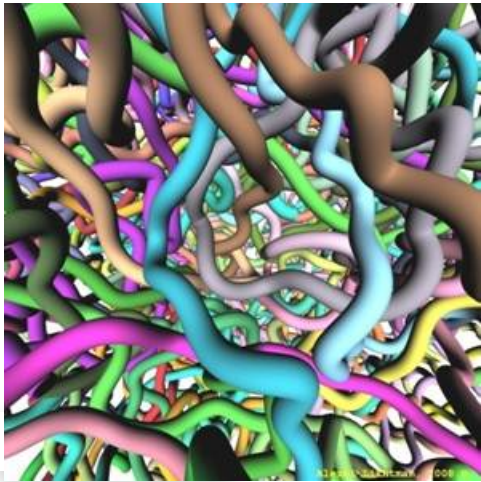
Nucleation



# Multiscale modelling in flow-induced crystallisation

Increasing computational speed

Molecular  
dynamics  
simulations



Continuous description  
of the position and  
momentum of of all  
atoms.

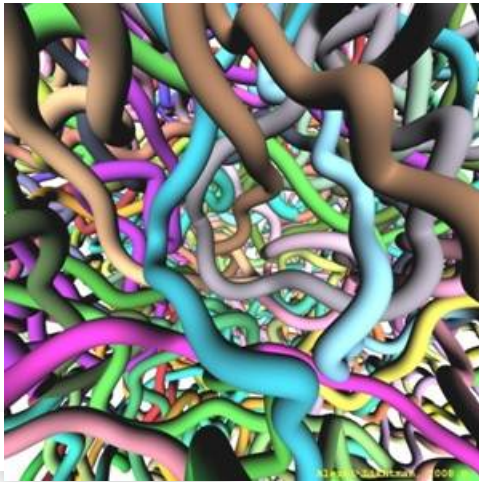
Increasing detail

# Multiscale modelling in flow-induced crystallisation

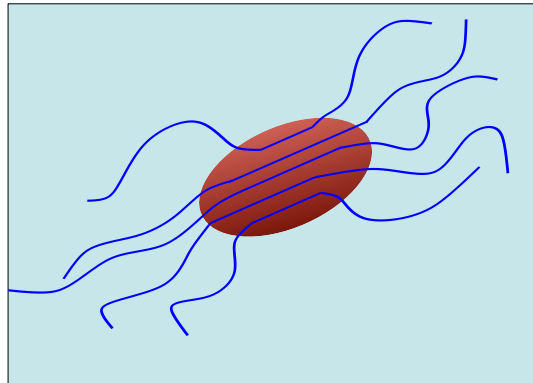
Increasing computational speed

Molecular dynamics simulations

The GO model (kinetic Monte Carlo)



Continuous description of the position and momentum of all atoms.



Limited resolution and discretised nucleus

Increasing detail

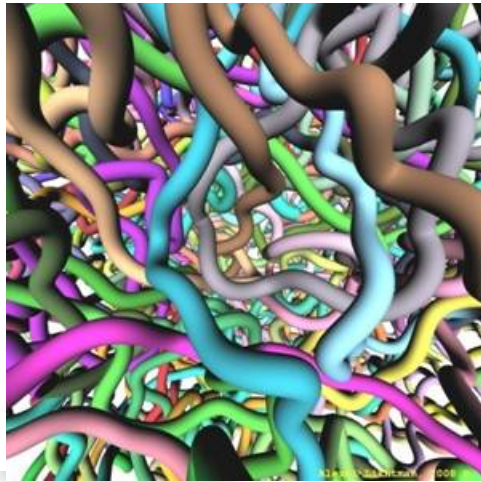
# Multiscale modelling in flow-induced crystallisation

Increasing computational speed

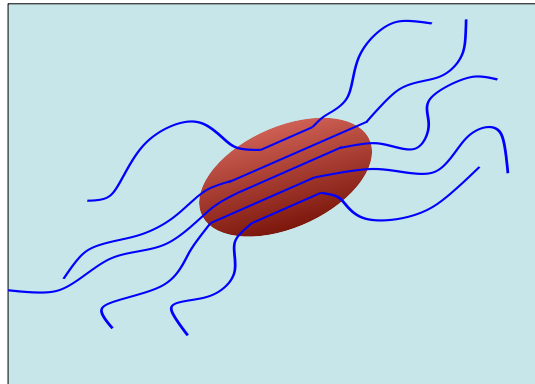
Molecular  
dynamics  
simulations

The GO model  
(kinetic Monte  
Carlo)

Macroscale  
continuum  
models



Continuous description  
of the position and  
momentum of all  
atoms.



Limited resolution  
and discretised  
nucleus

$$\begin{aligned}\dot{\phi}_3 &= 8\pi\dot{N}(T), \\ \dot{\phi}_2 &= G(T)\phi_3, \\ \dot{\phi}_1 &= G(T)\phi_2, \\ \dot{\phi}_0 &= G(T)\phi_1, \\ \dot{N} &= \dot{N}_0 \exp[\eta(\lambda^2 - 1)]\end{aligned}$$

Continuum  
description through  
deterministic  
differential equations

Increasing detail

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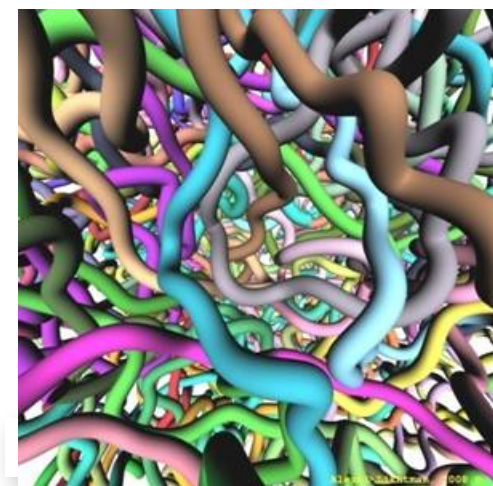
Increasing computational speed

Molecular dynamics simulations

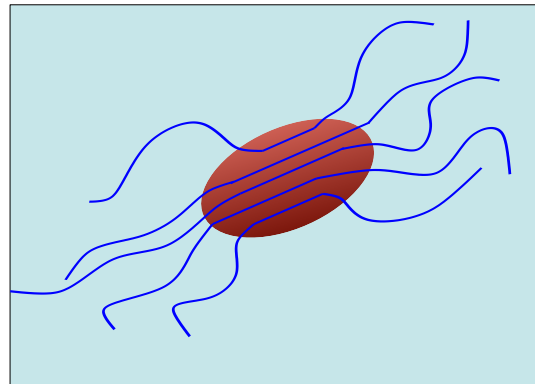
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Macroscale continuum models

Finite element calculations



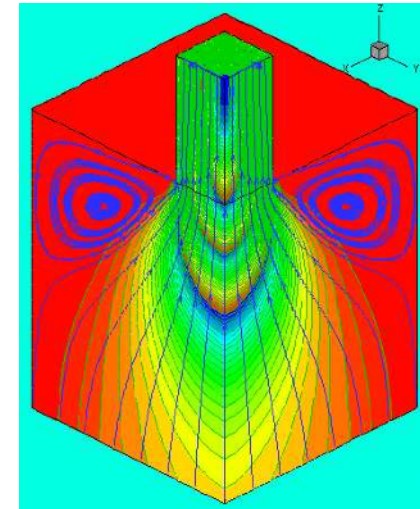
Continuous description of the position and momentum of all atoms.



Limited resolution and discretised nucleus

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Continuum description through deterministic differential equations



Computational fluid dynamics of polymer crystallisation during processing

Increasing detail

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Increasing computational speed 

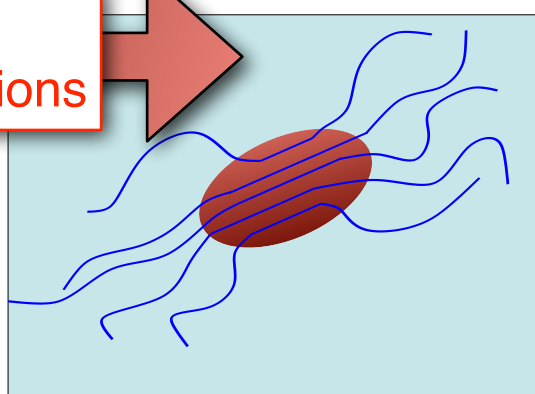
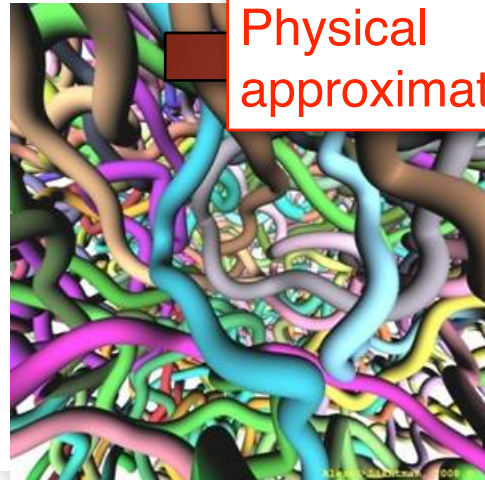
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Physical approximations 

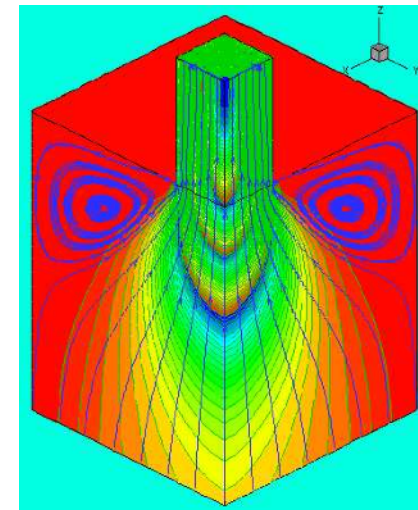


Continuous description of the position and momentum of all atoms.

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Computational fluid dynamics of polymer crystallisation during processing

Increasing detail 

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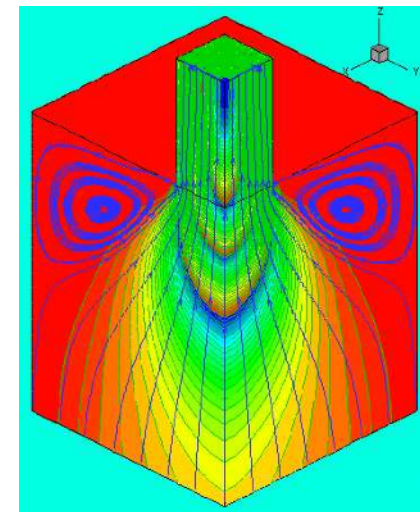
Mathematical approximations 

$$\begin{aligned} \dot{\phi}_3 &= 8\pi\dot{N}(T), \\ \dot{\phi}_2 &= G(T)\phi_3, \\ \dot{\phi}_1 &= G(T)\phi_2, \\ \dot{\phi}_0 &= G(T)\phi_1, \\ &\dots \end{aligned}$$

Limited resolution and discretised nucleus

Continuum description through deterministic differential equations

Continuous description of the position and momentum of all atoms.



Computational fluid dynamics of polymer crystallisation during processing

Increasing detail 



# Multiscale modelling in flow-induced crystallisation

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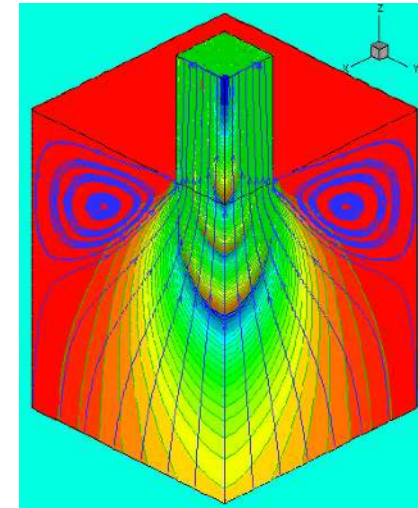
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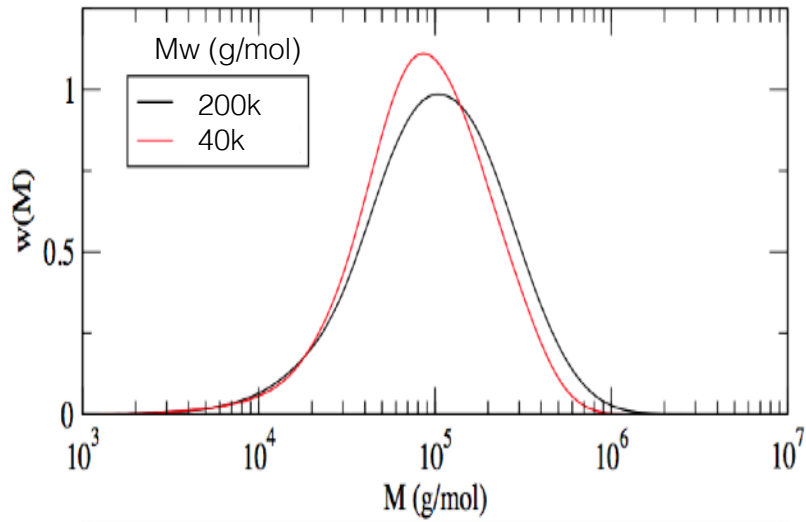
Computational fluid dynamics of polymer crystallisation during

Continuous description of the position and momentum of all atoms.

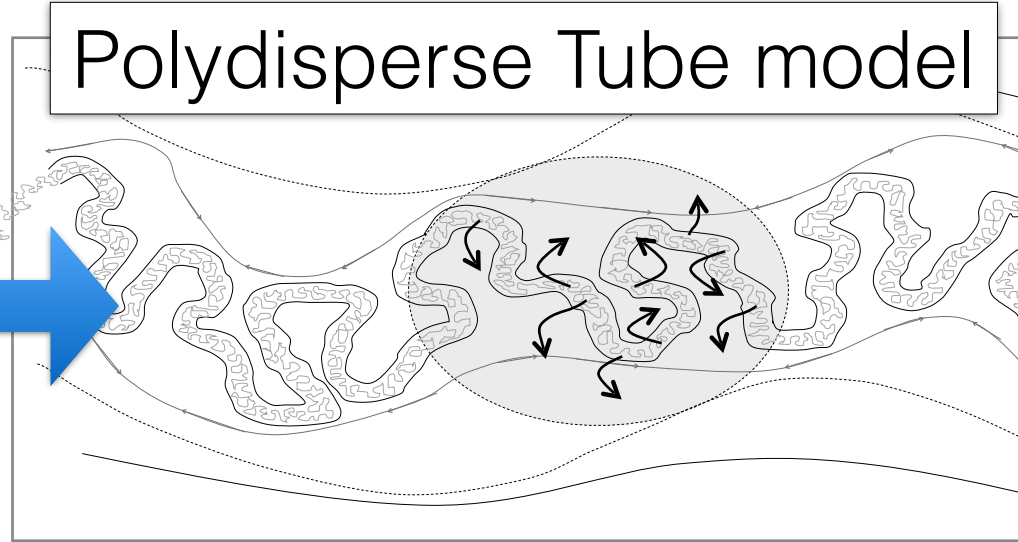
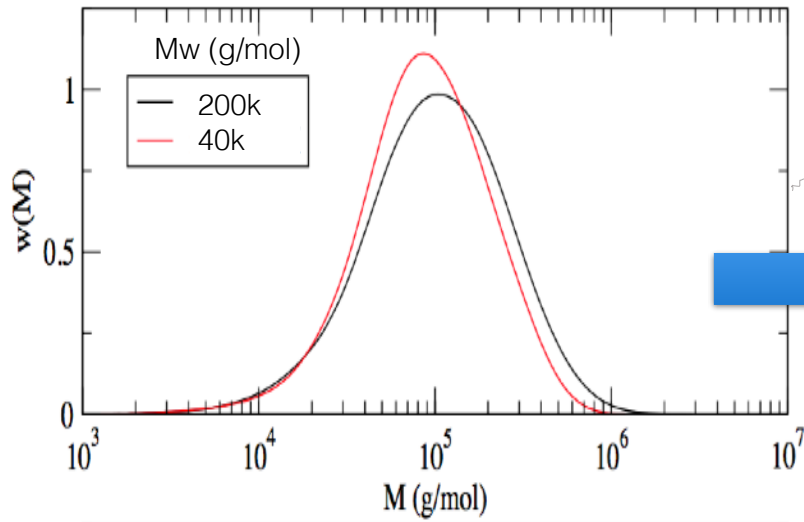
Techniques developed can assist more detailed simulations 

Increasing detail 

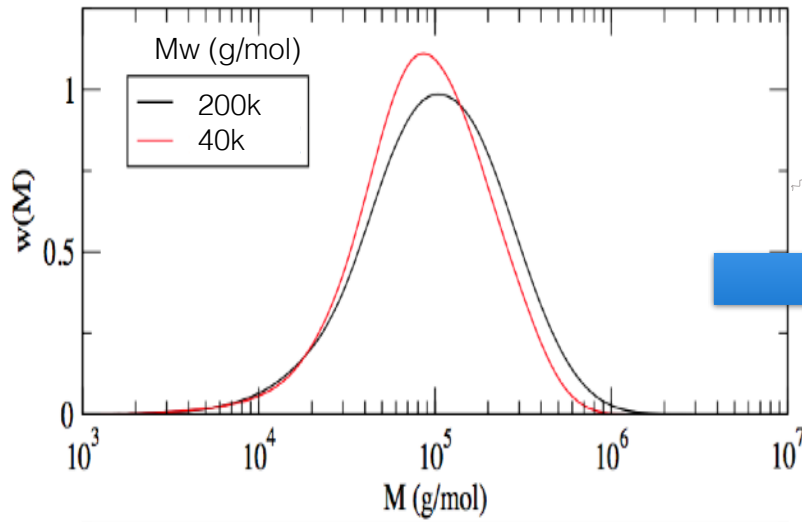
# Rheological modelling (flow only)



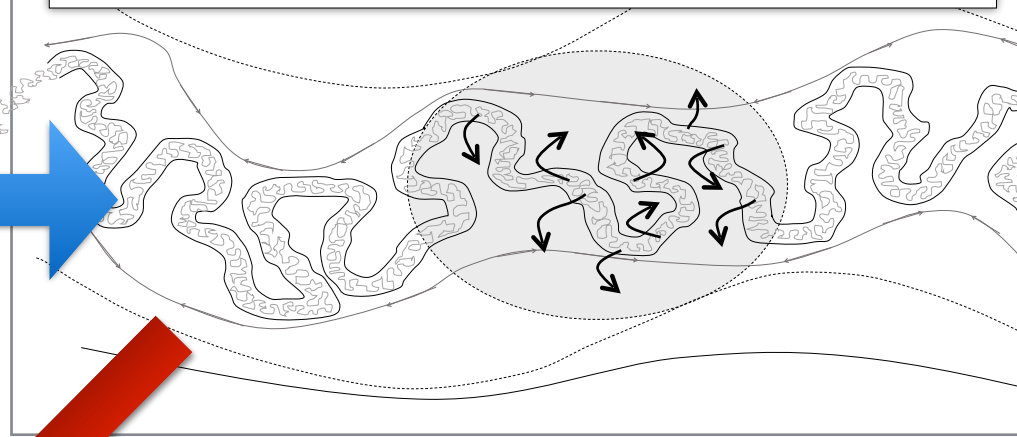
# Rheological modelling (flow only)



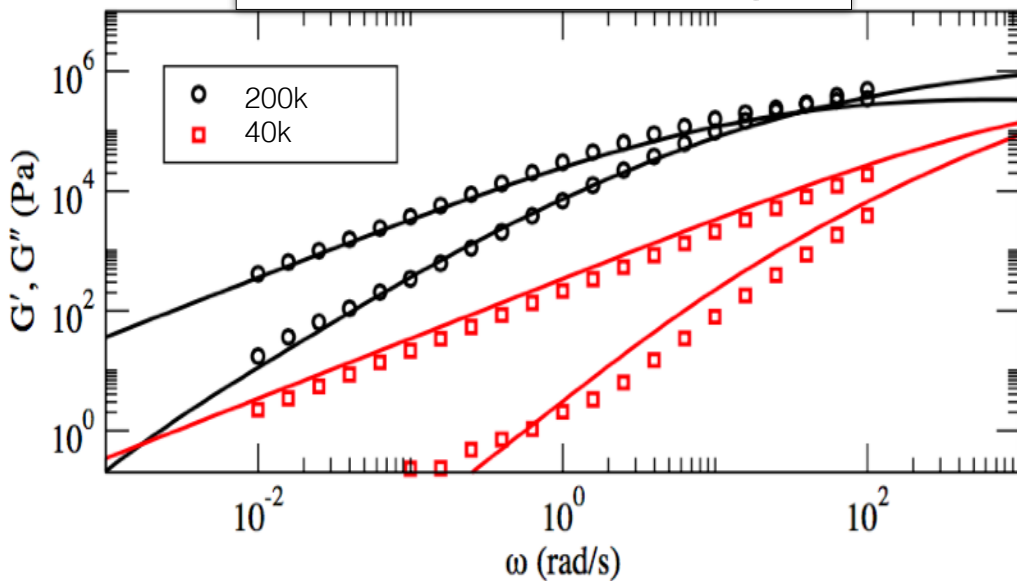
# Rheological modelling (flow only)



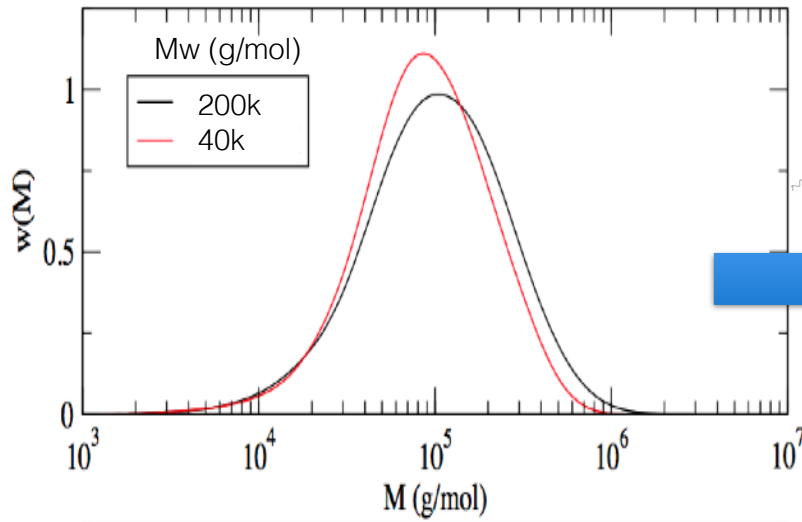
Polydisperse Tube model



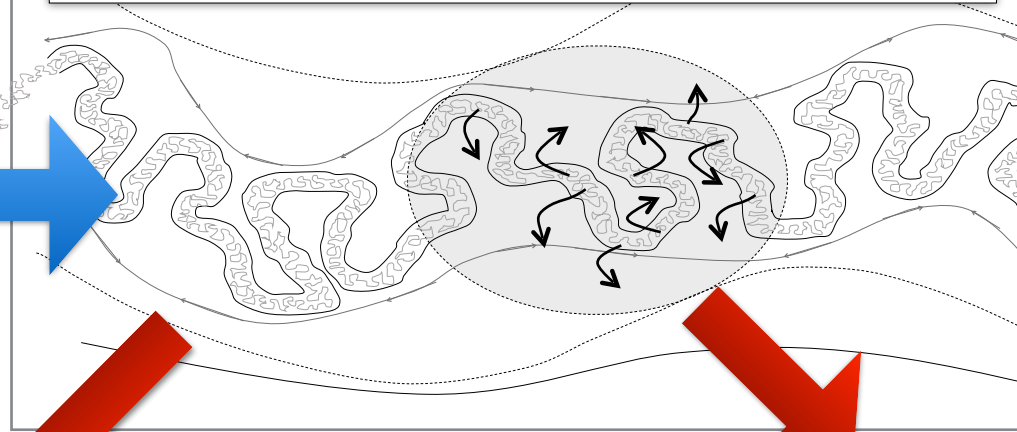
Linear rheology



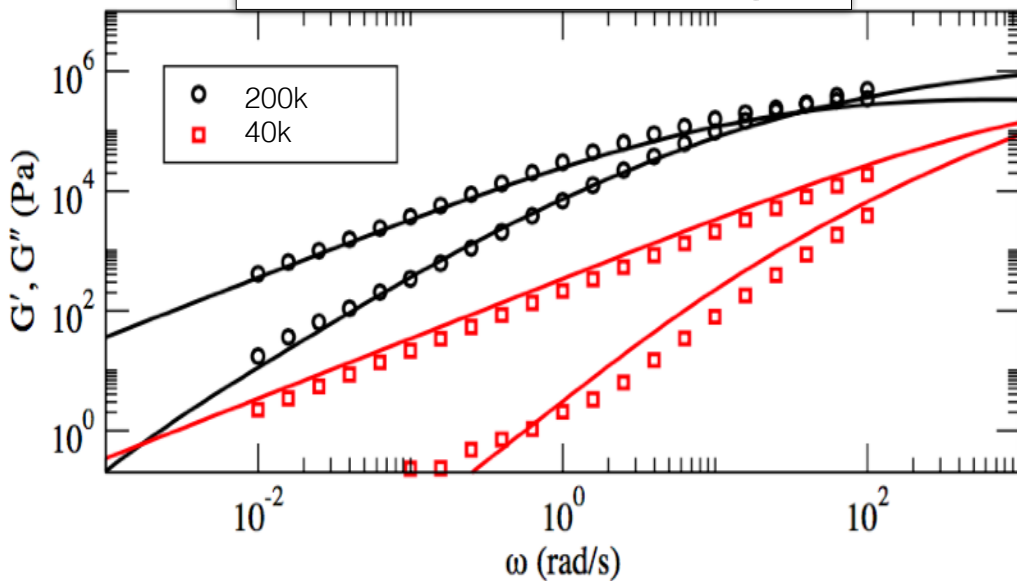
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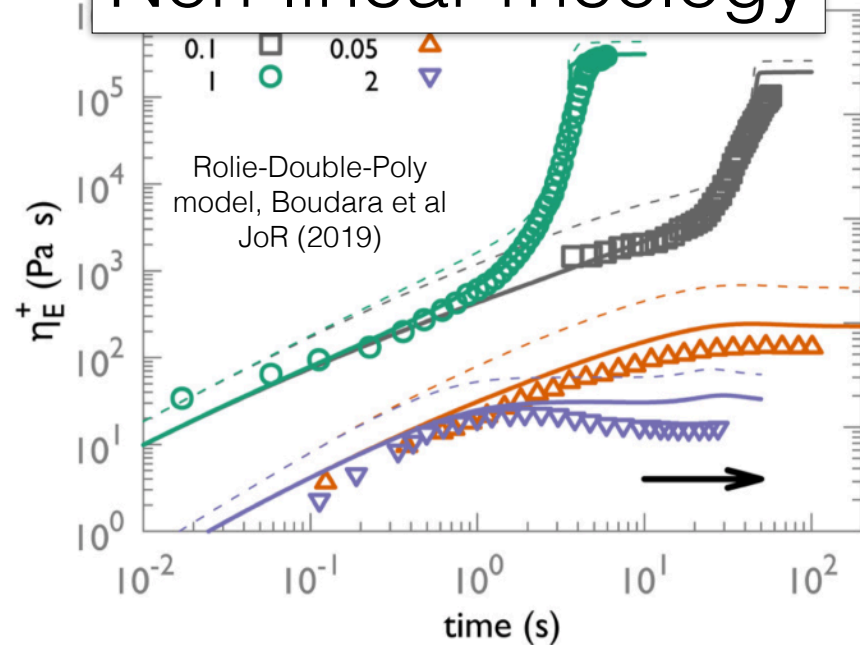
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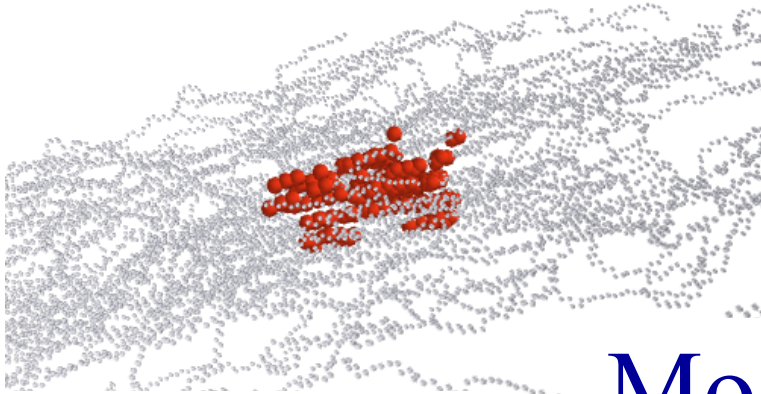


Non-linear rheology



# Modelling nucleation

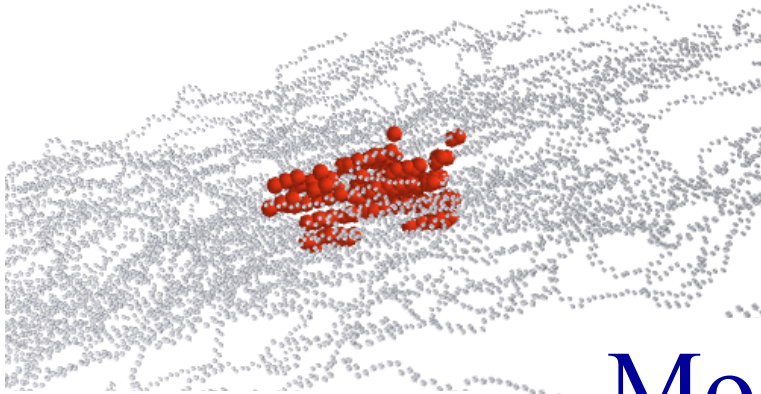
## (a) Molecular dynamics



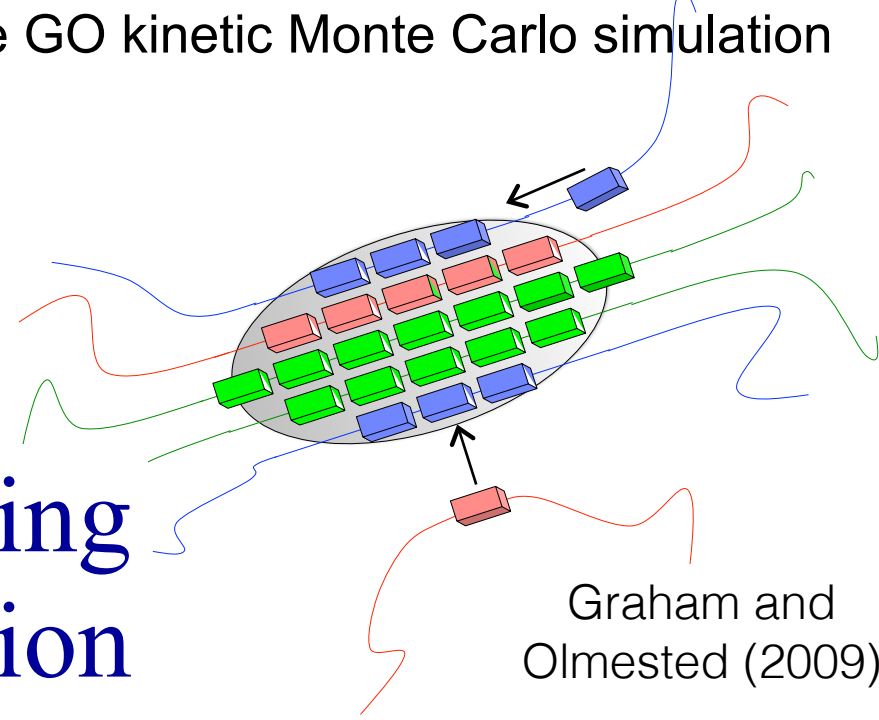
Rutledge et al (2013) & (2016)  
Schilling et al (2013) & (2014)  
Anwar and Graham (2019)

# Modelling nucleation

(a) Molecular dynamics



(b) The GO kinetic Monte Carlo simulation



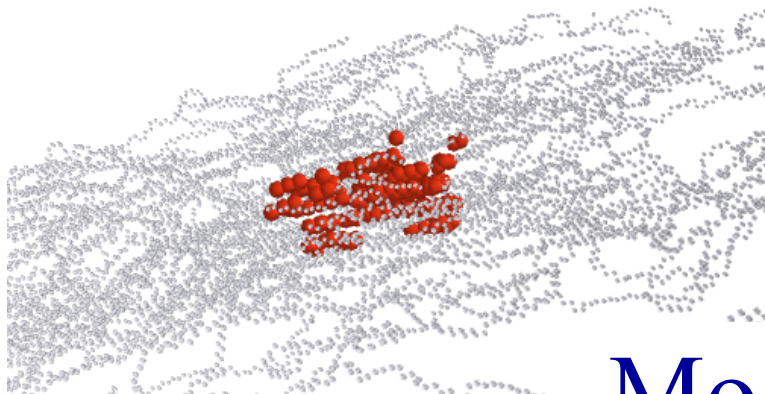
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# Modelling nucleation

Graham and  
Olmsted (2009)

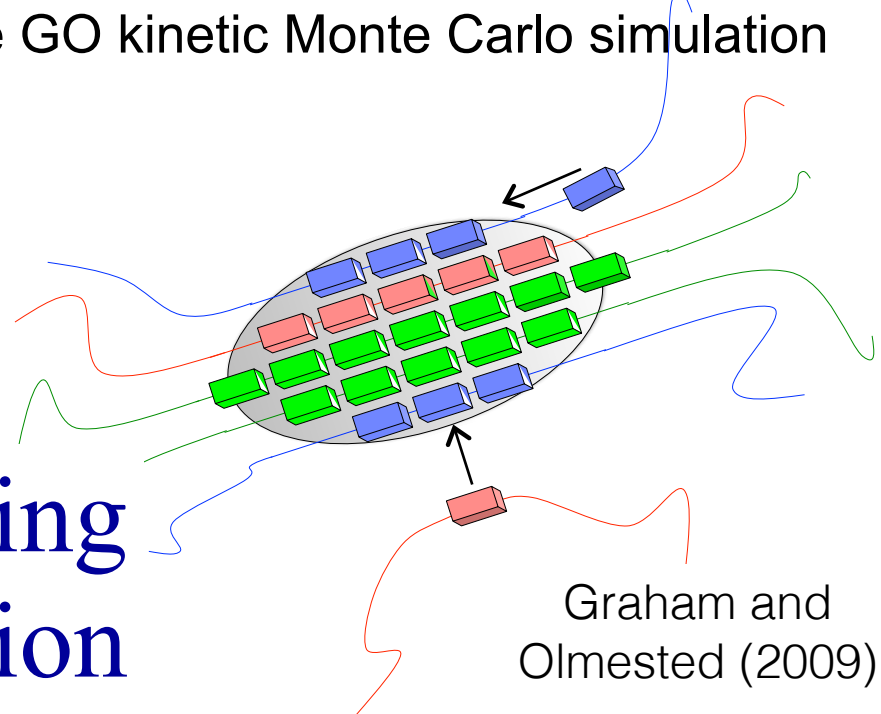


(a) Molecular dynamics



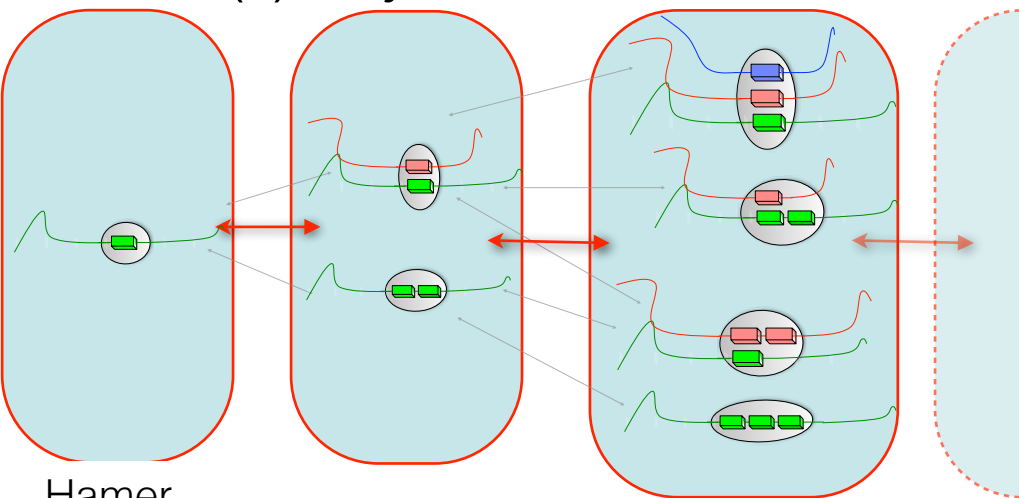
Rutledge et al (2013) & (2016)  
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# Modelling nucleation

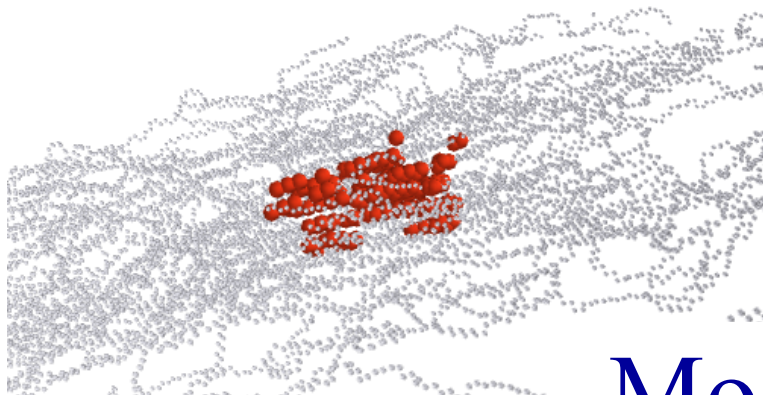
(c) Projection to 1D



Hamer,  
Wattis and  
Graham  
(2012)

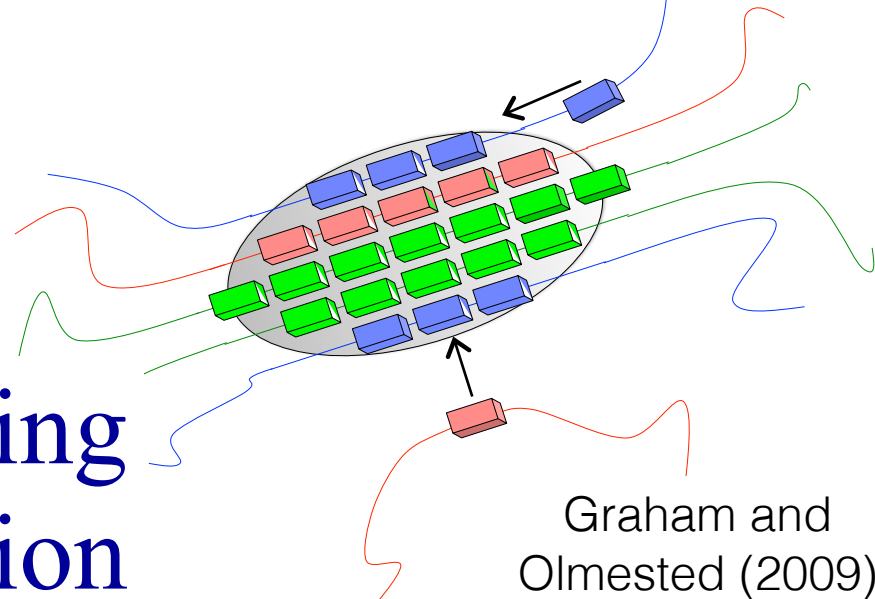
2  
3  
4  
Total nucleus size

(a) Molecular dynamics



Rutledge et al (2013) & (2016)  
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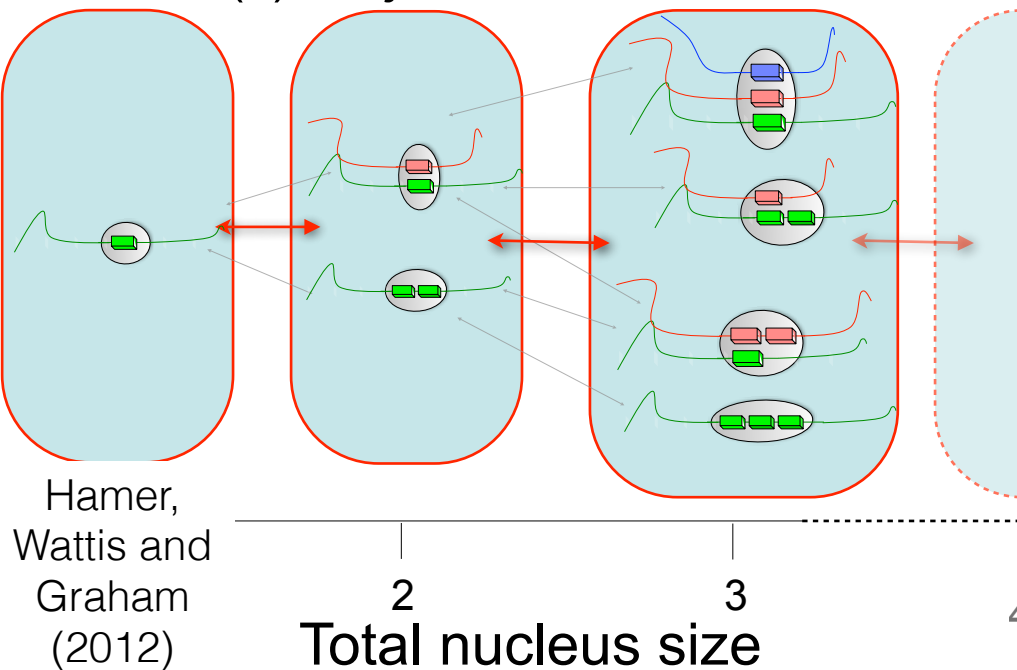
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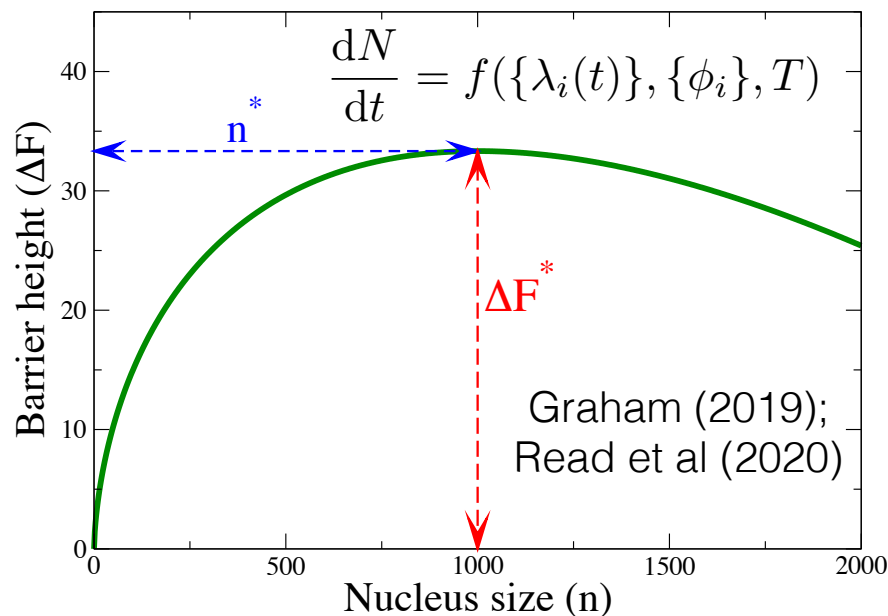
Graham and Olmsted (2009)

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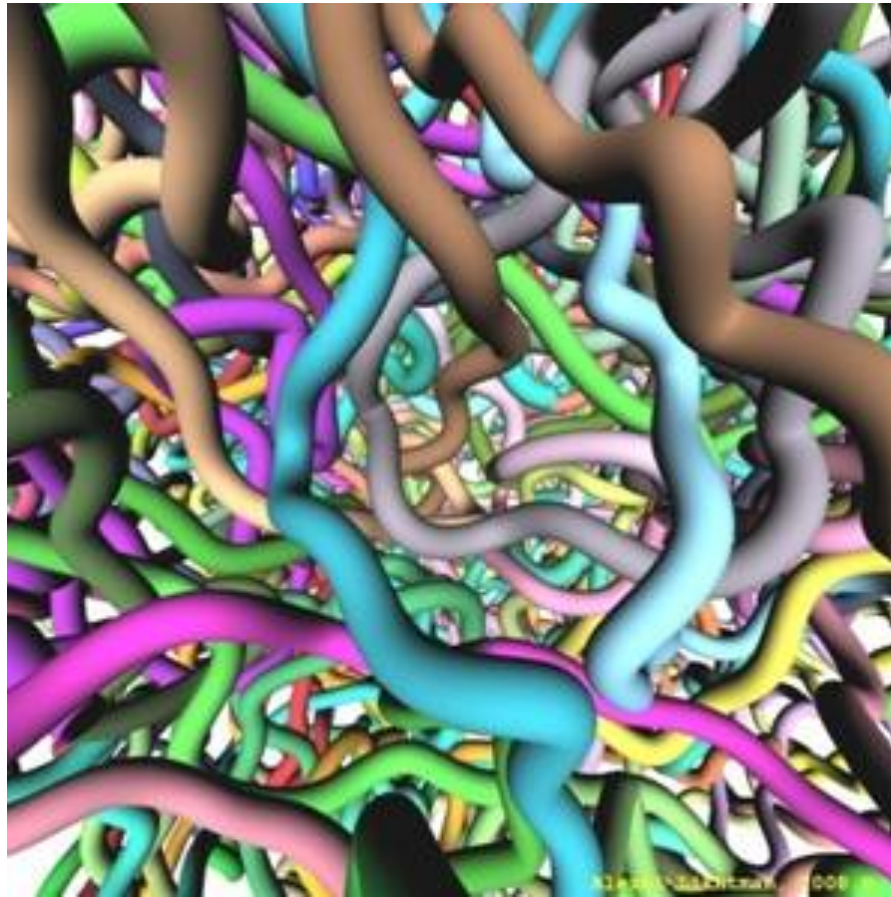
(c) Projection to 1D



(d) Continuum modelling

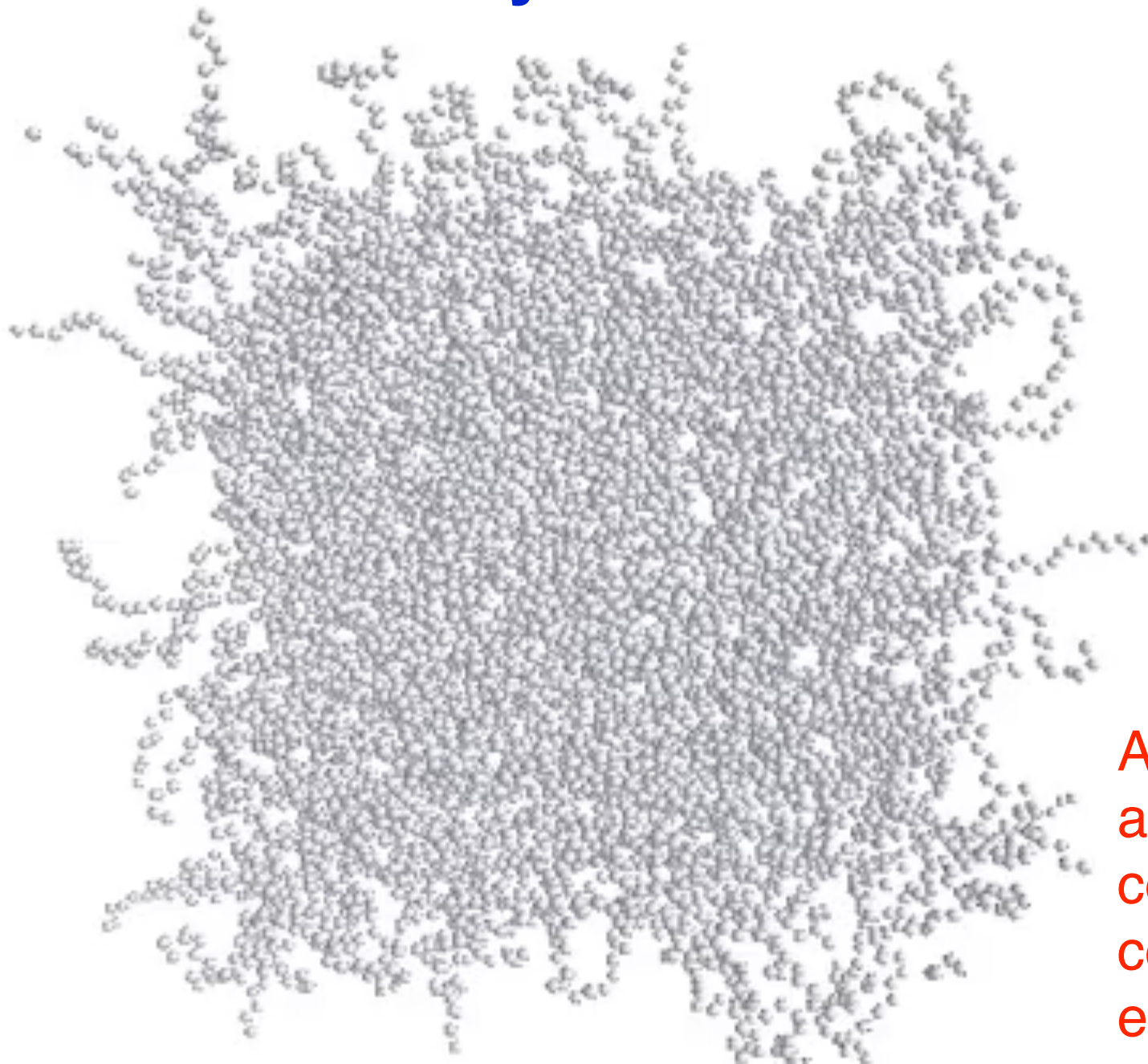


# Molecular Dynamics - Monodisperse polymers



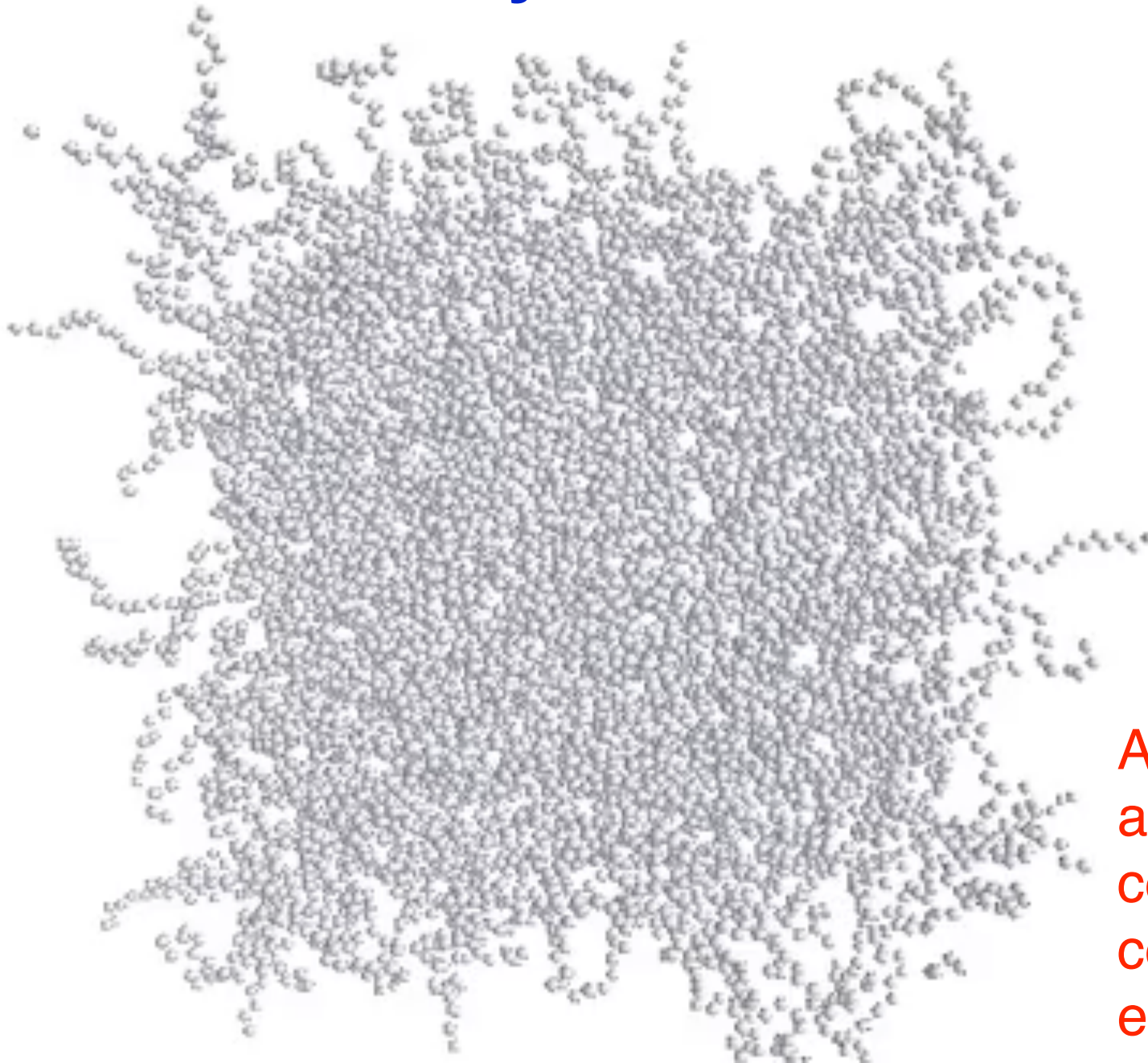
Continuous description of the position and momentum of all atoms.

# Molecular dynamics: flow & nucleation



All simulations  
at high under-  
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# Molecular dynamics: flow & nucleation

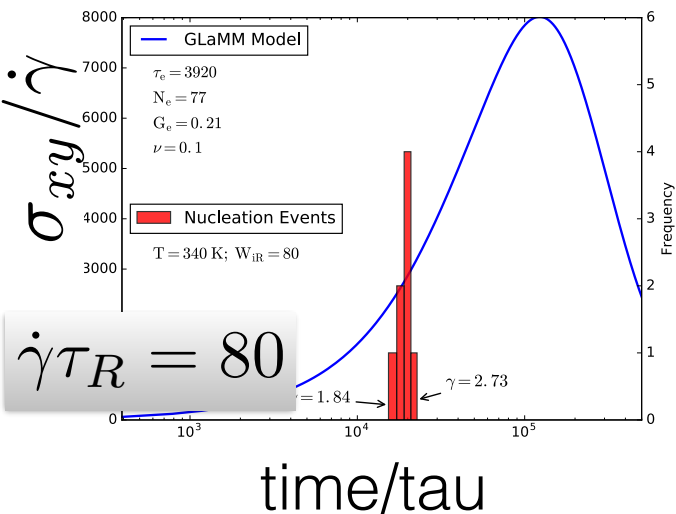
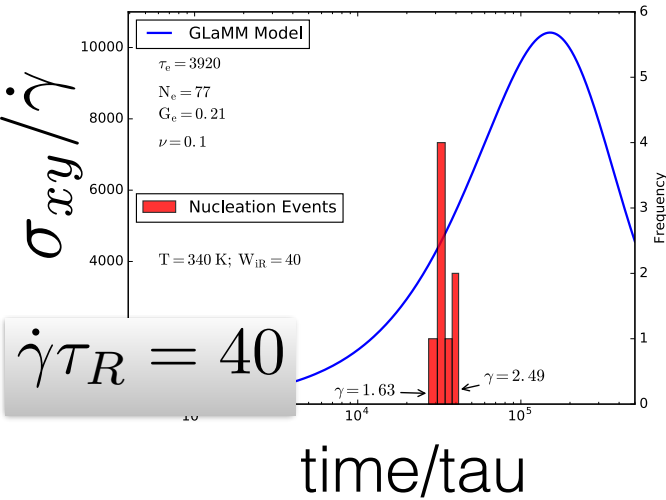


All simulations  
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# Molecular dynamics:

## The timing of nucleation events during the stress transient

$$T = 340K$$

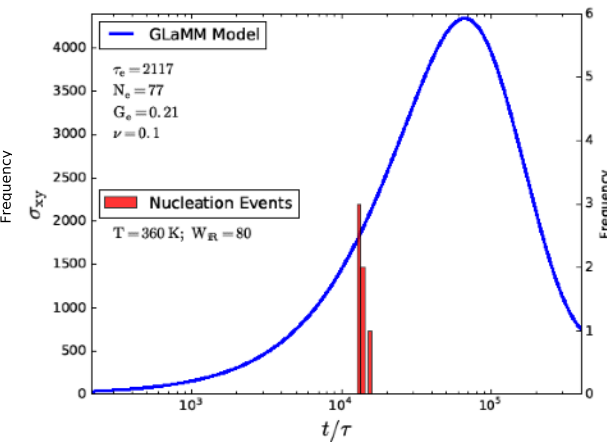
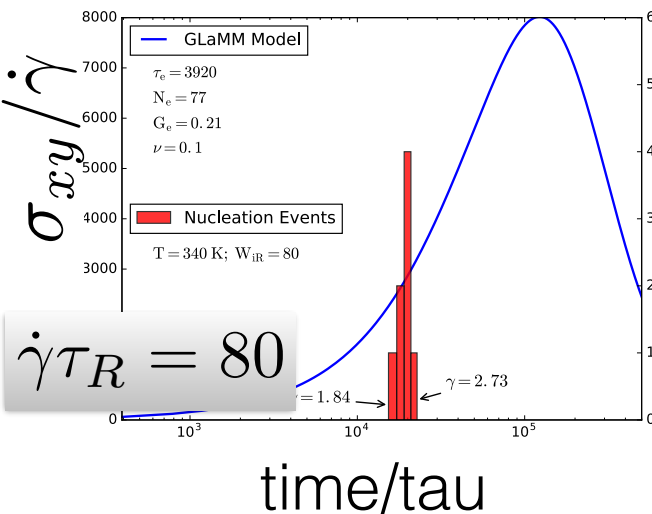
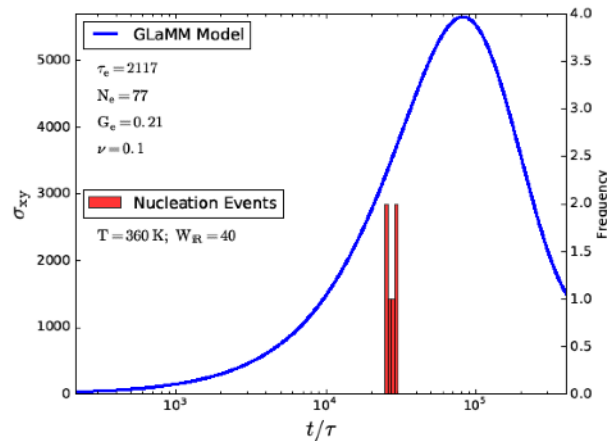
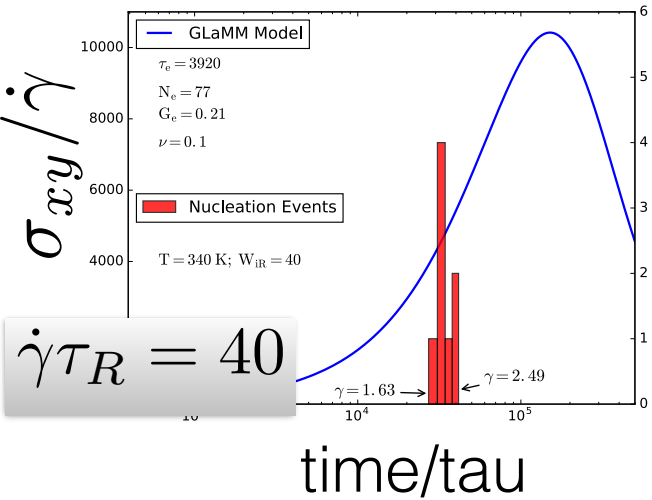


# Molecular dynamics:

## The timing of nucleation events during the stress transient

$T = 340K$

$T = 360K$



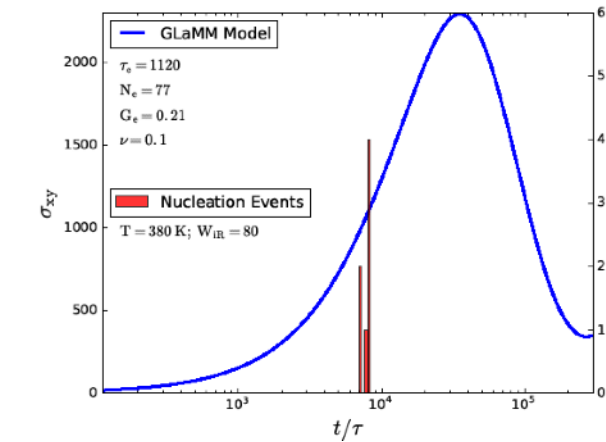
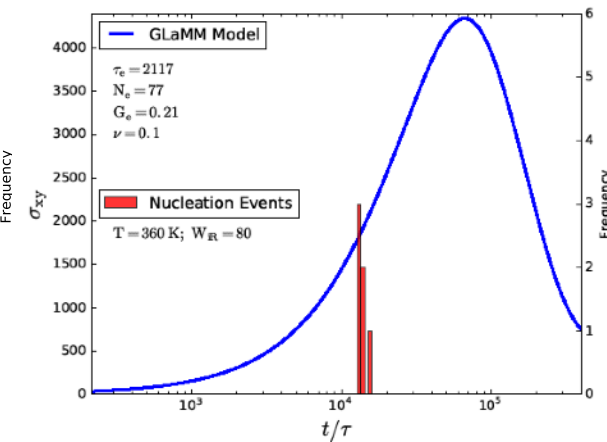
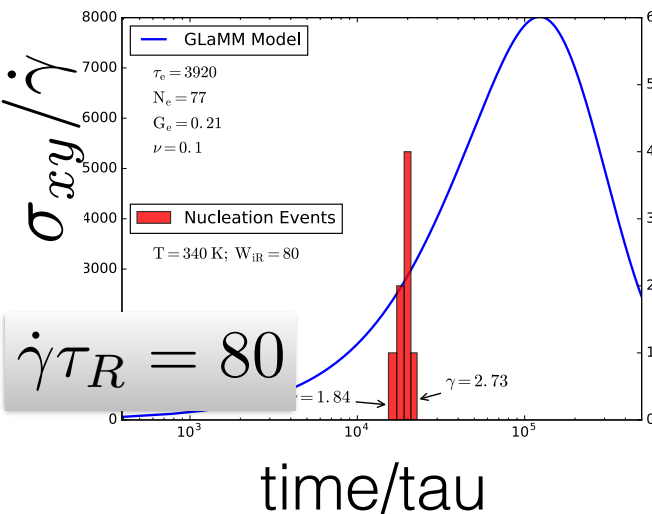
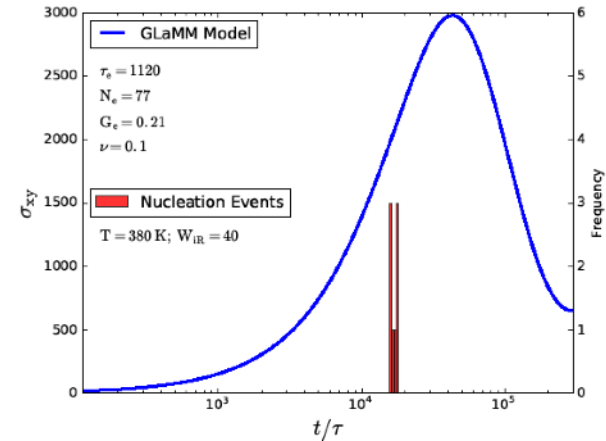
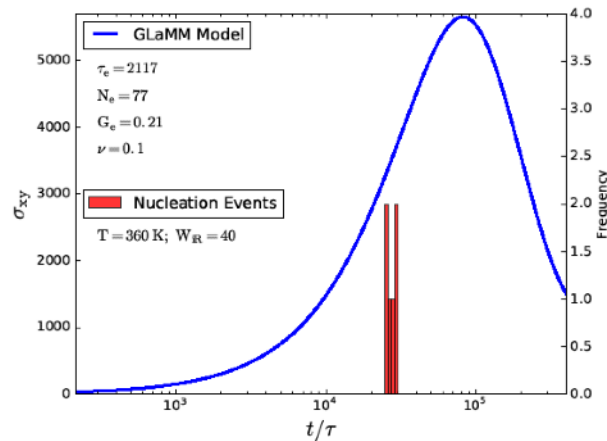
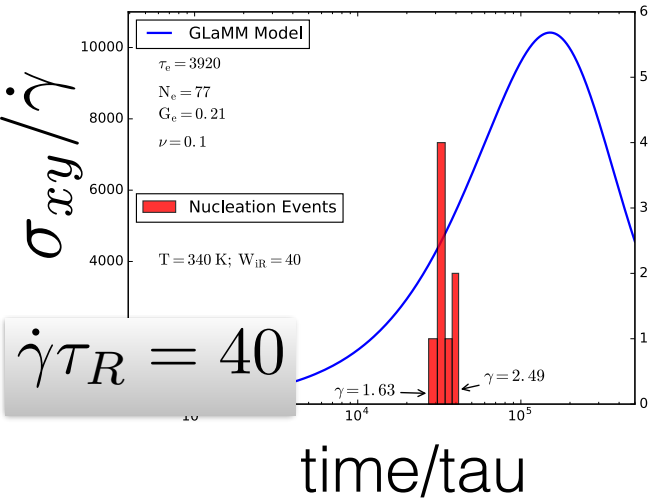
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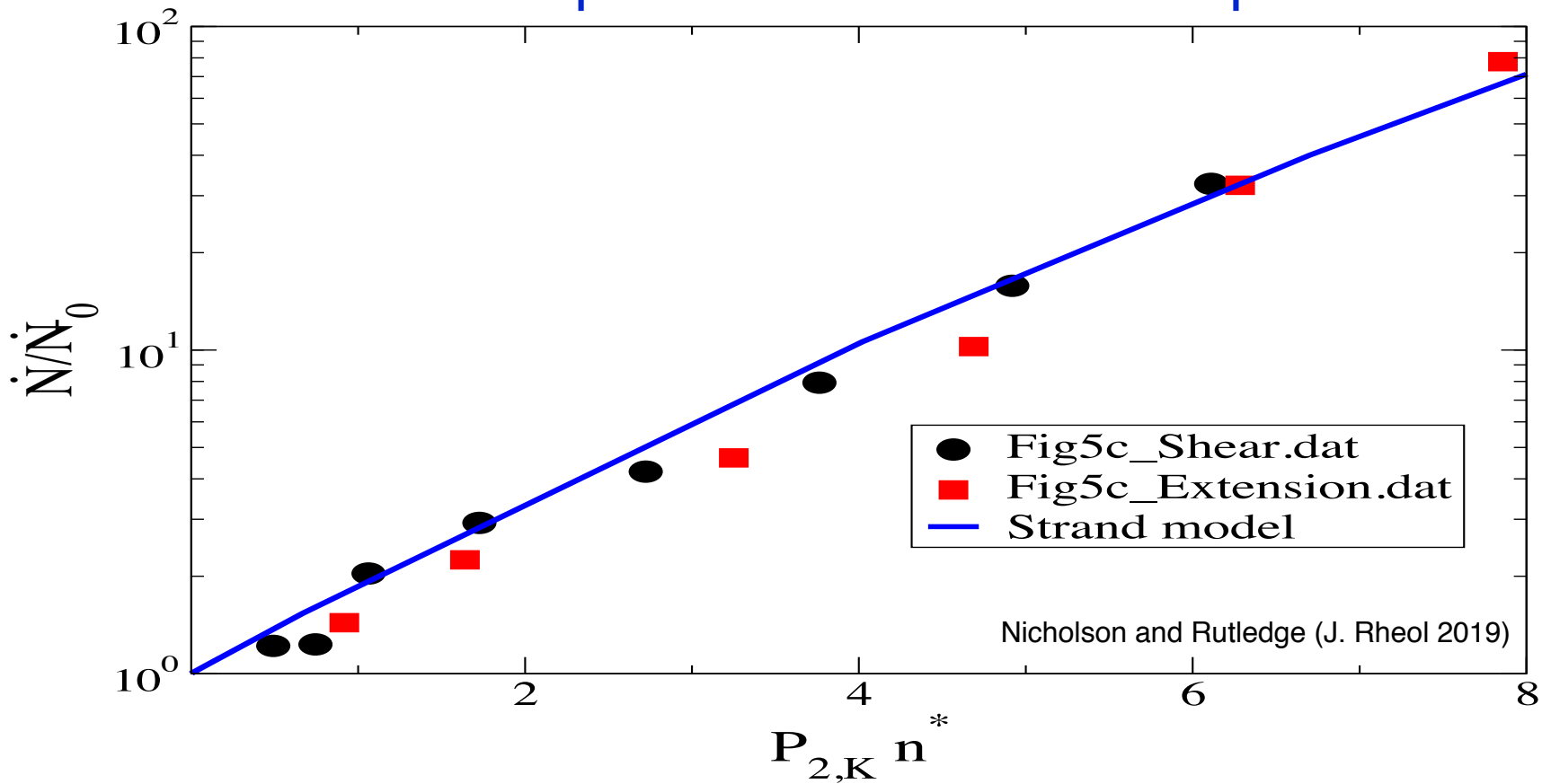
$T = 380K$





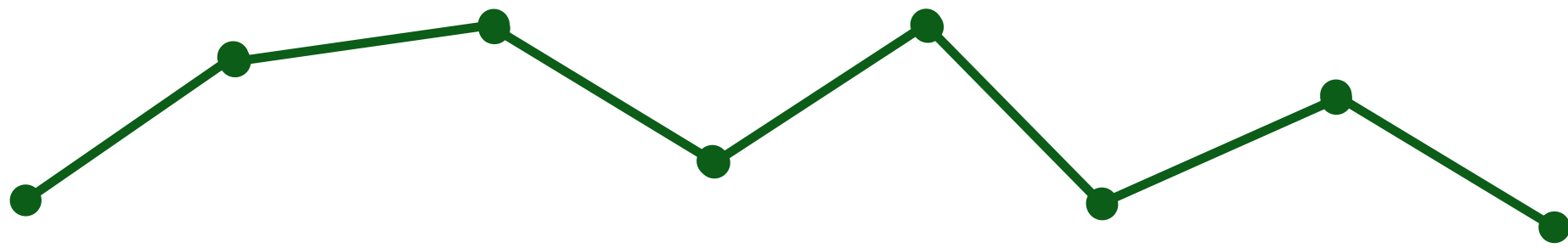
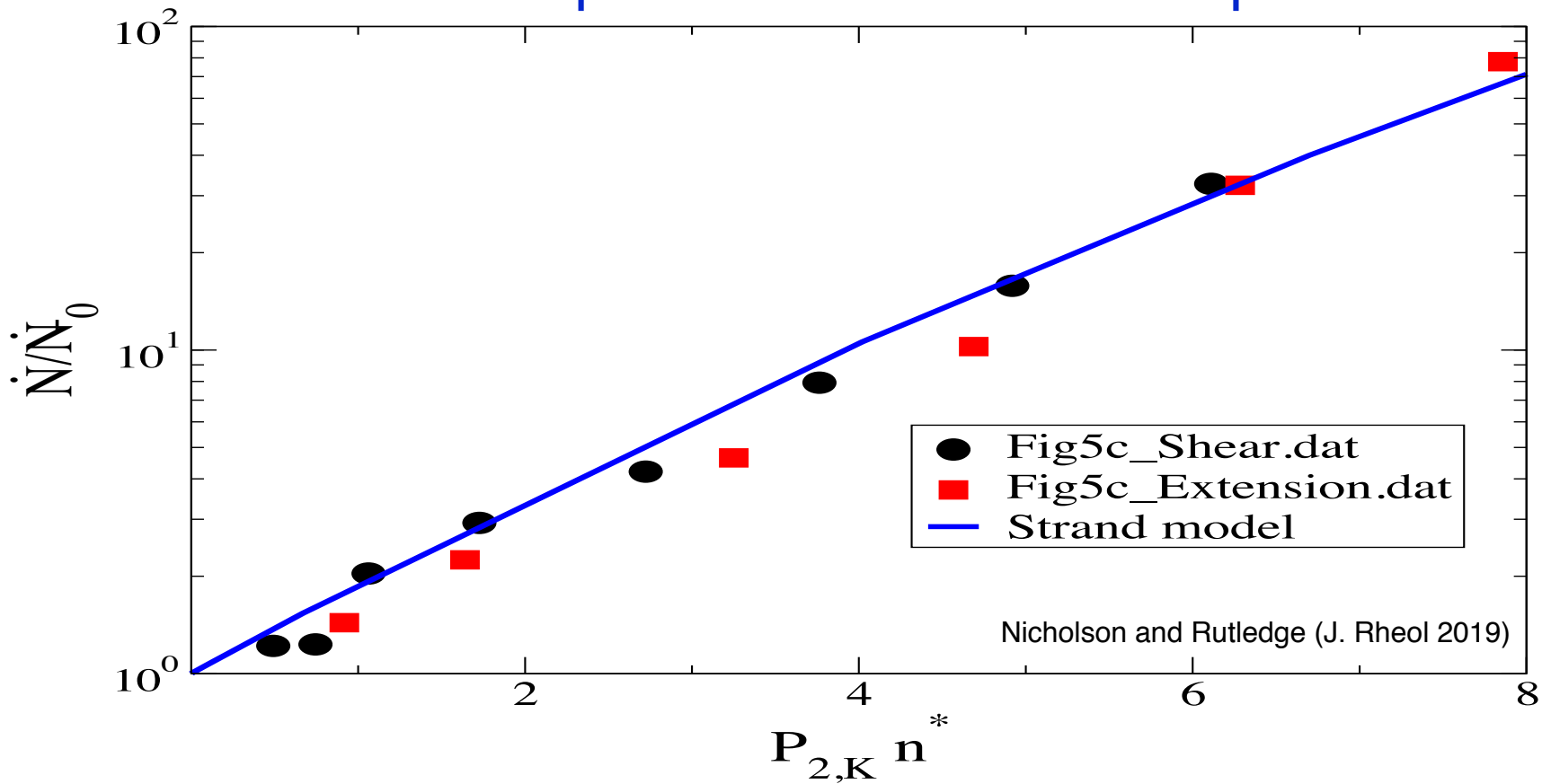
# A key result from Nicholson and Rutledge

Nucleation rate is exponential in the Kuhn step nematic order



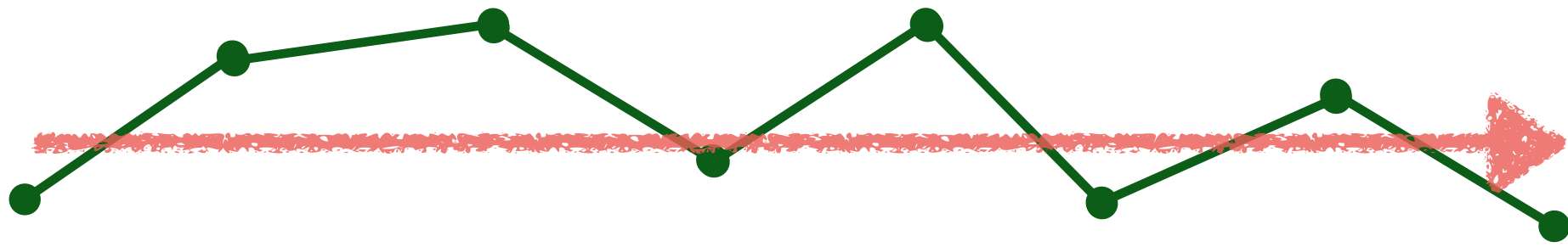
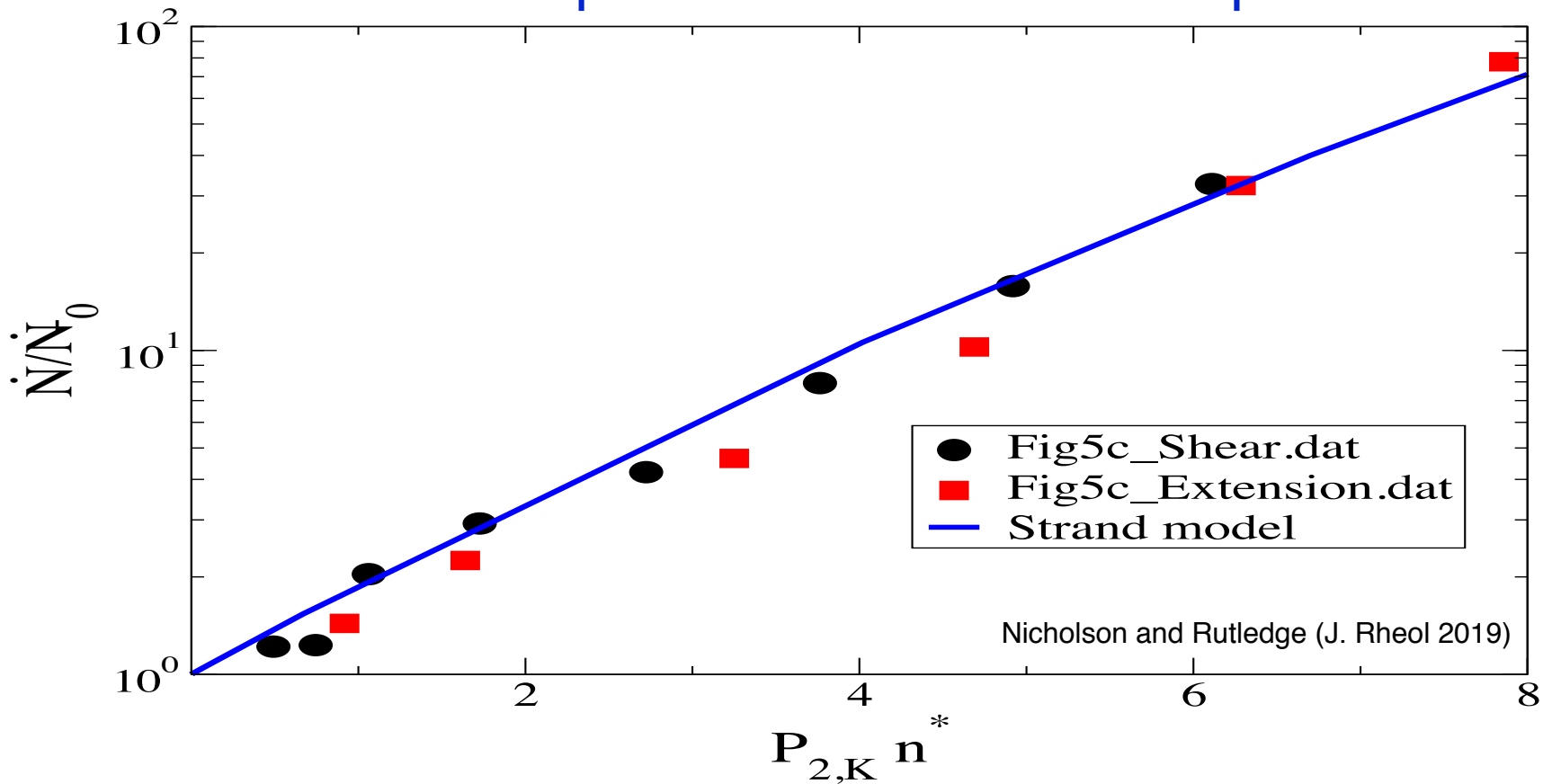
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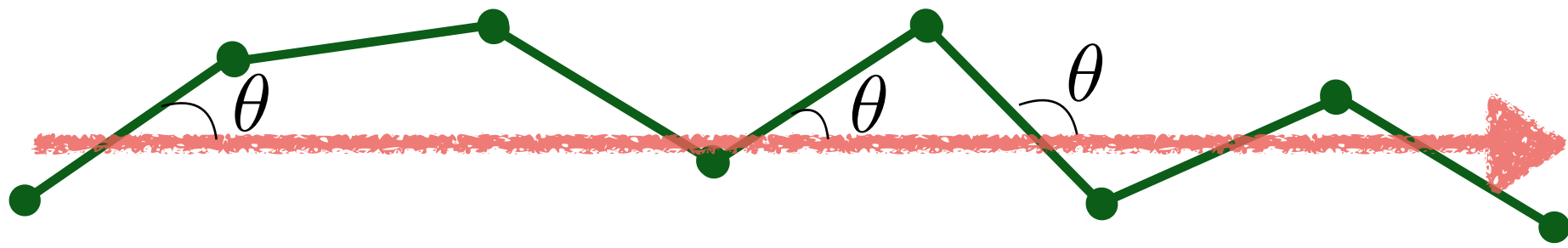
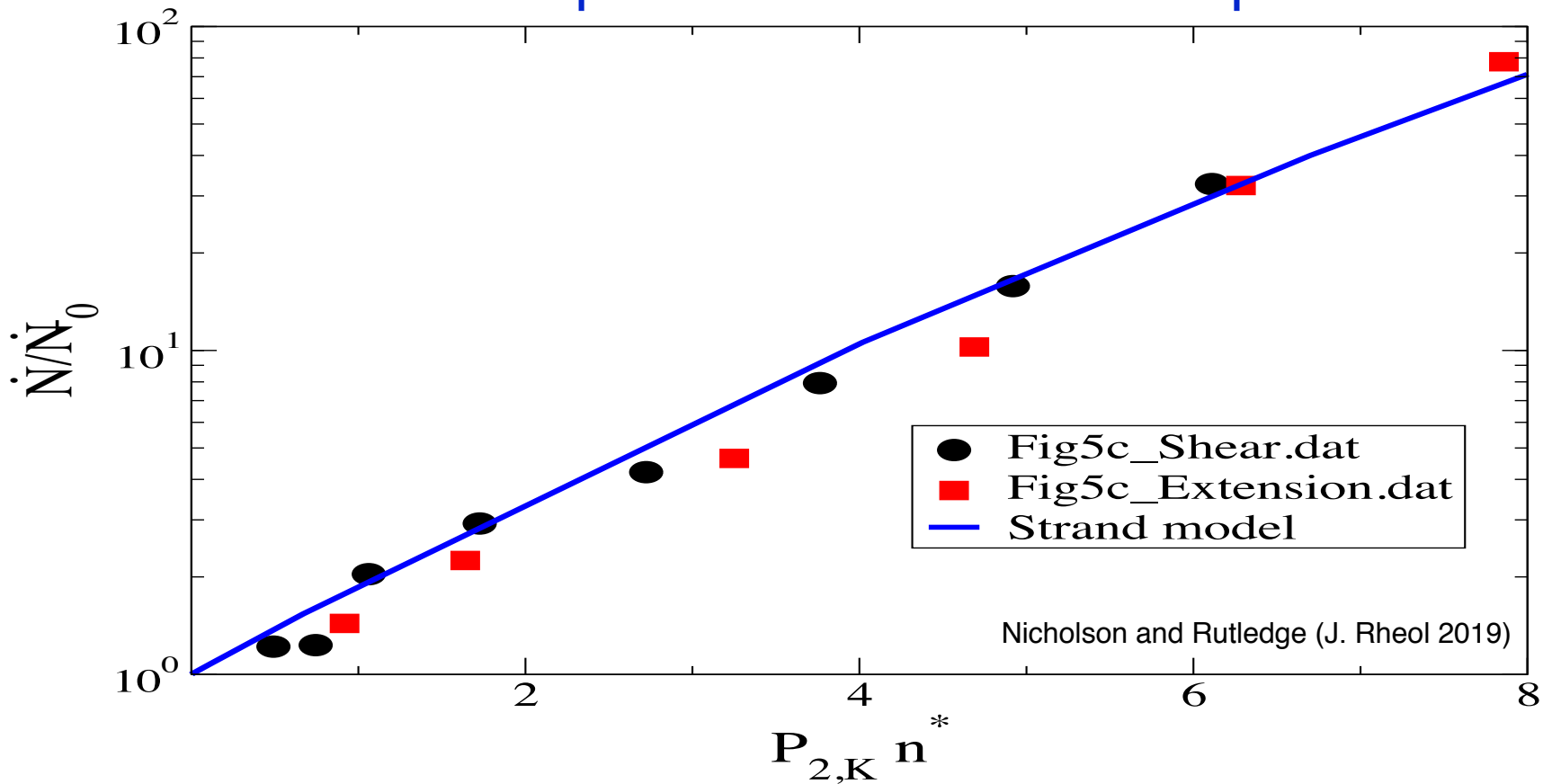
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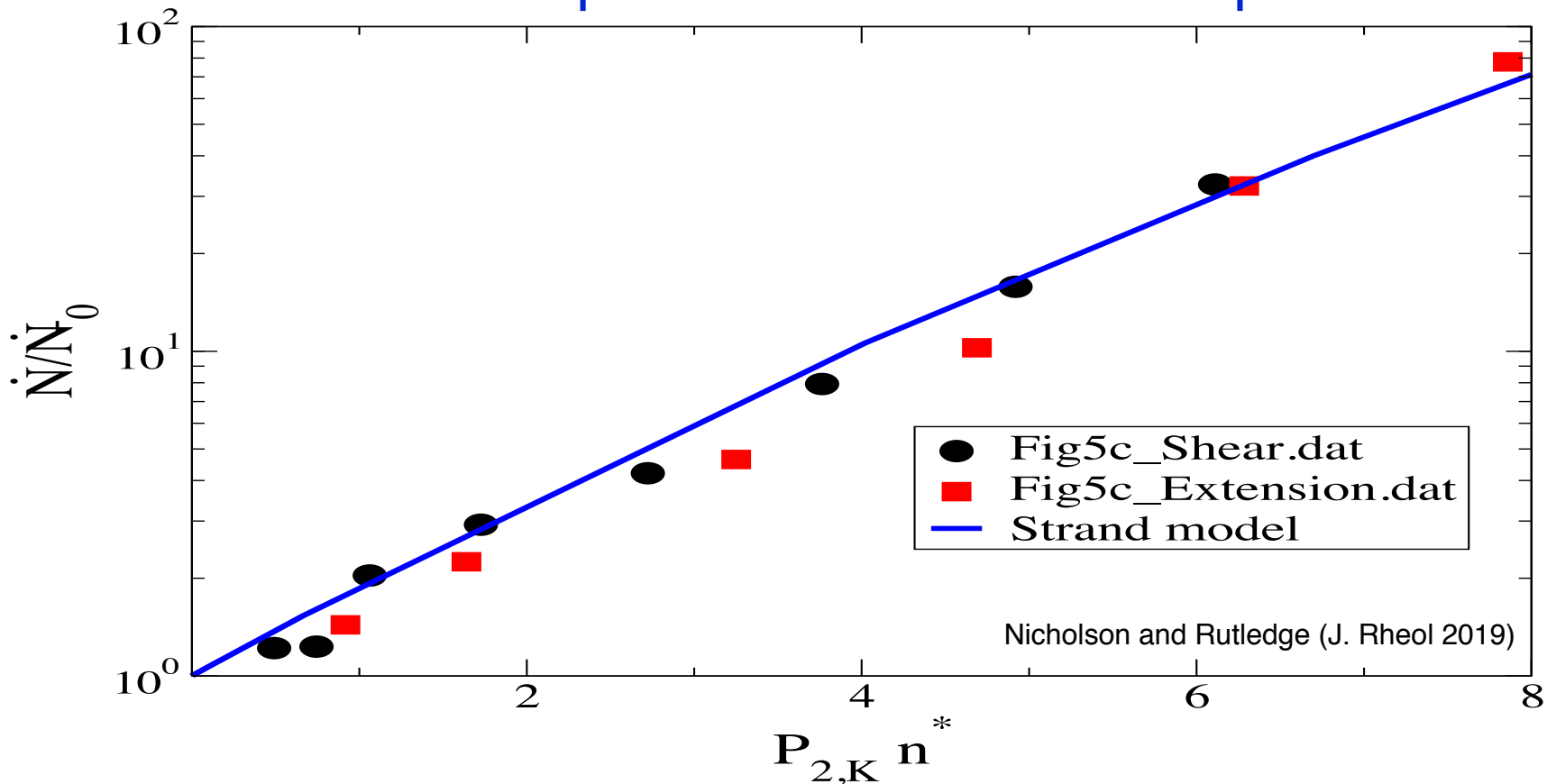
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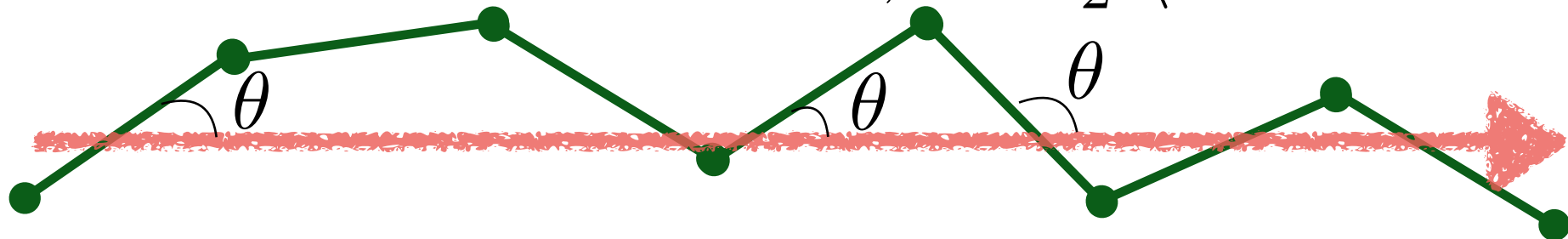
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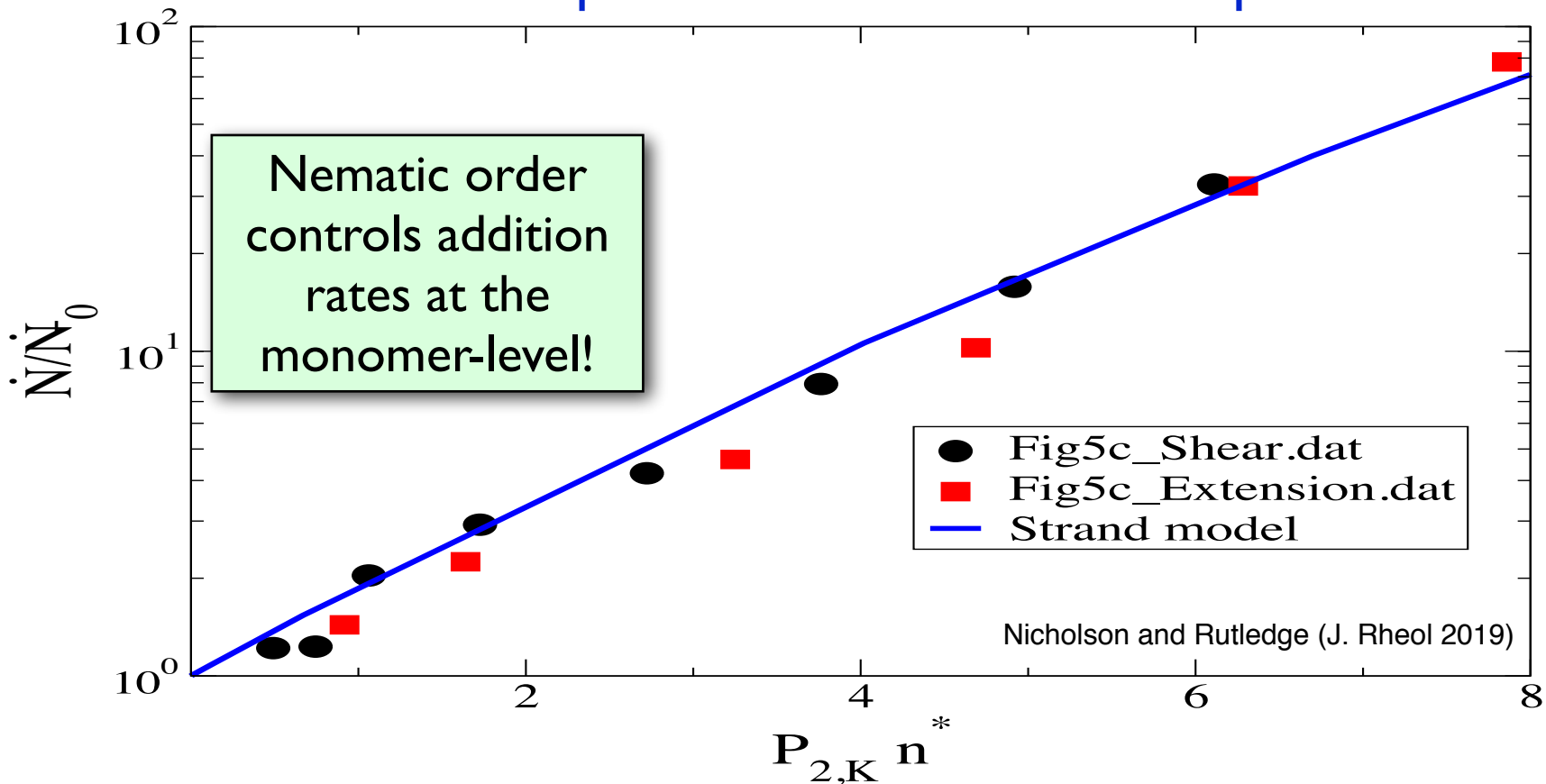
Nicholson and Rutledge (J. Rheol 2019)

$$P_{2,K} = \frac{1}{2} \langle 3 \cos^2 \theta - 1 \rangle$$

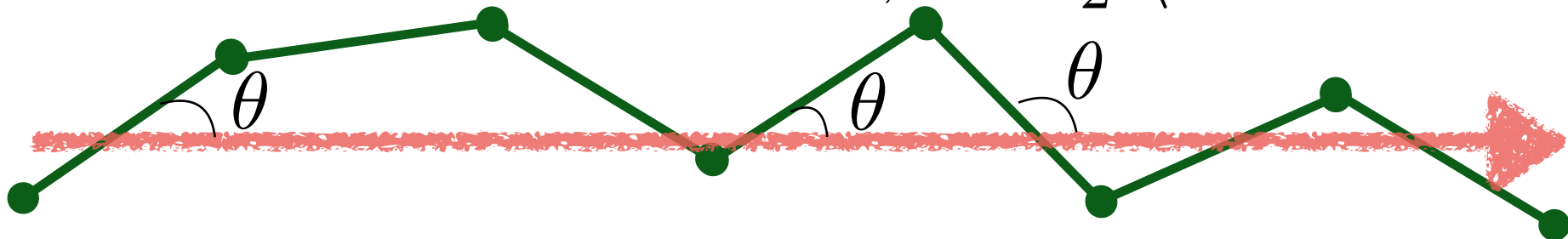


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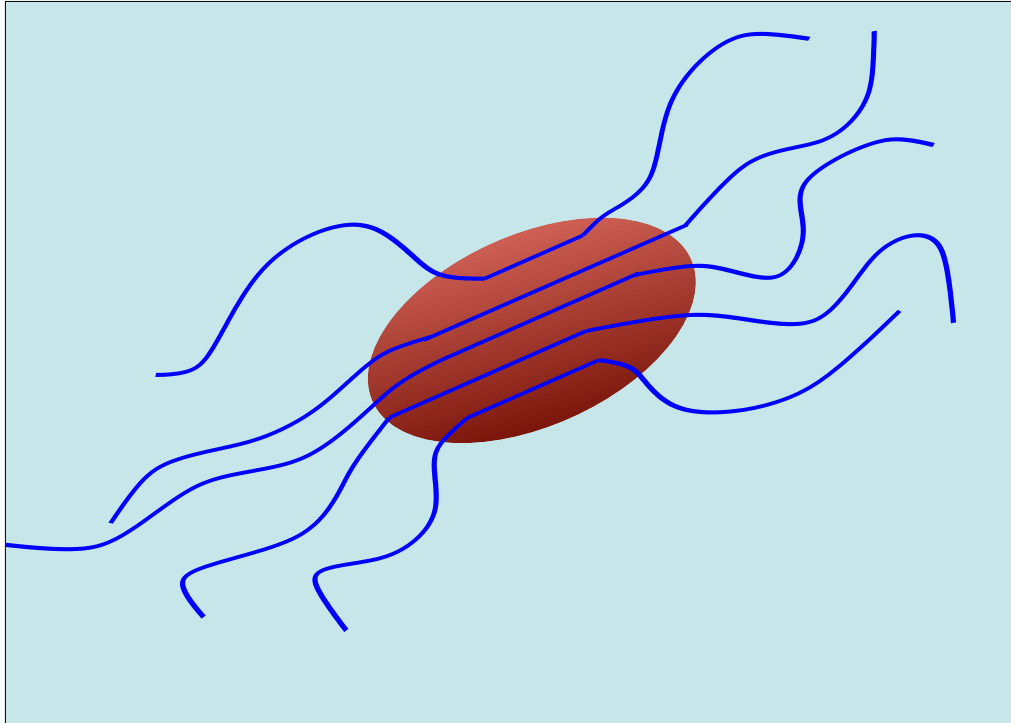
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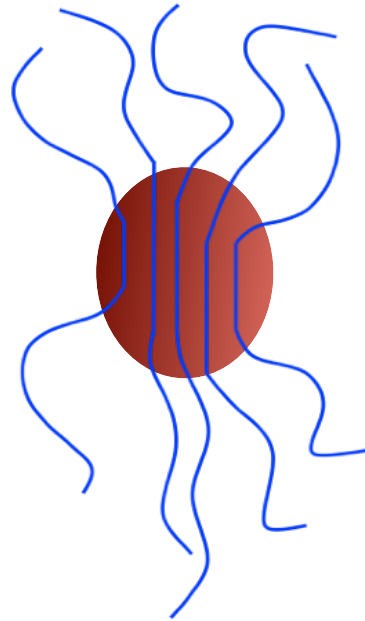


# GO model (kinetic Monte-Carlo)



Limited resolution and  
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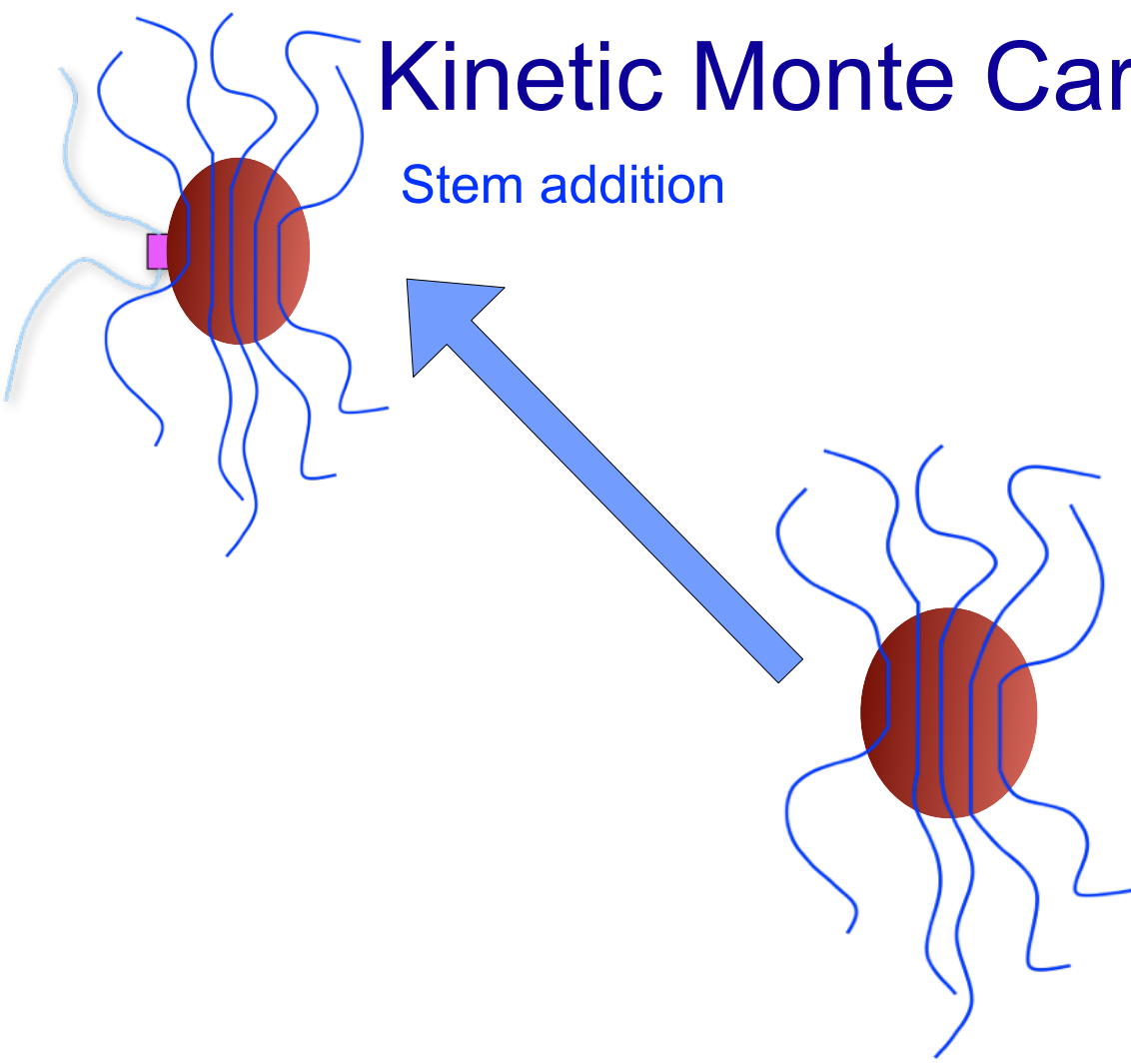
# Kinetic Monte Carlo model



Graham and Olmsted,  
Phys. Rev. Lett. **115**  
115707 (2009)



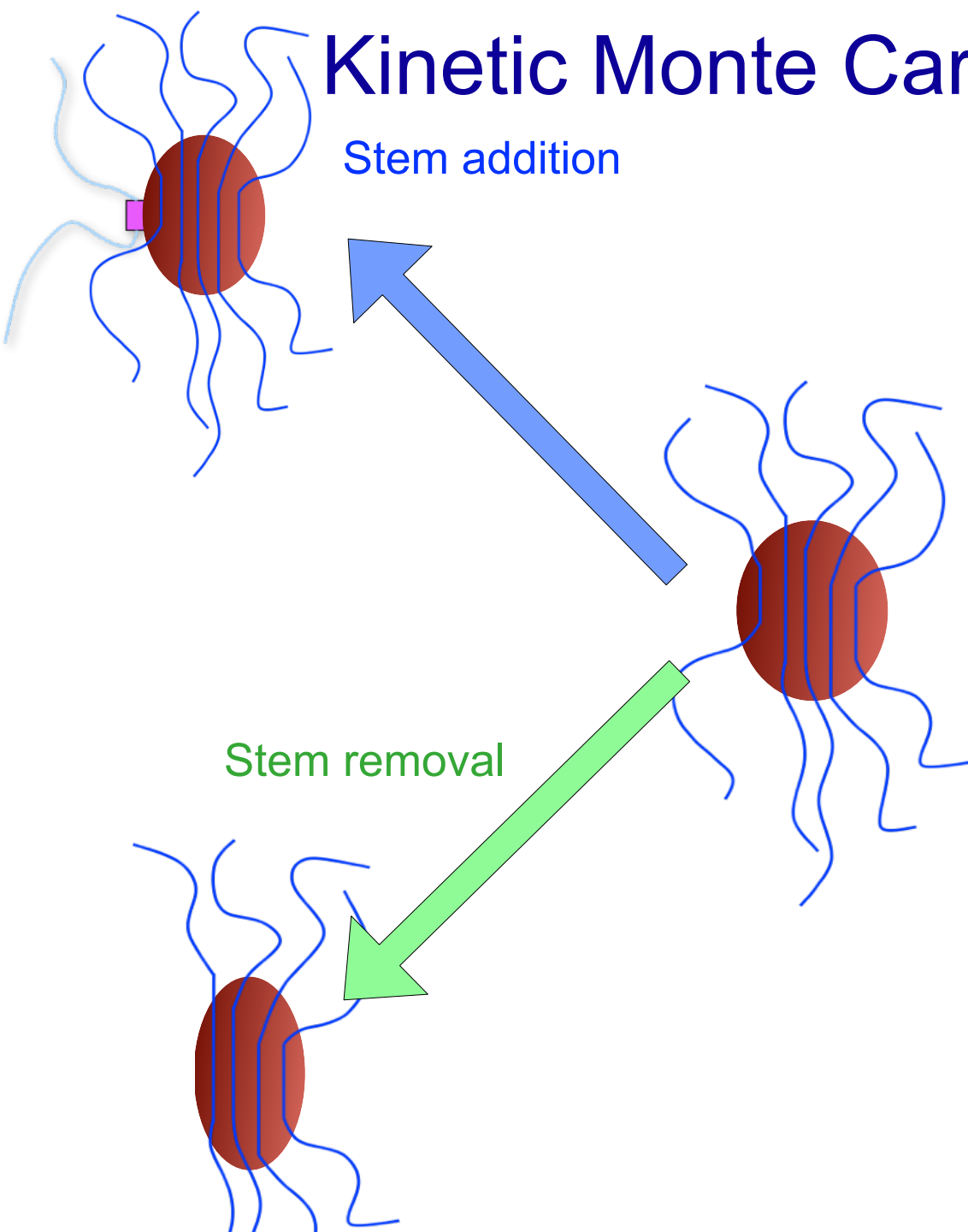
# Kinetic Monte Carlo model



Stem addition

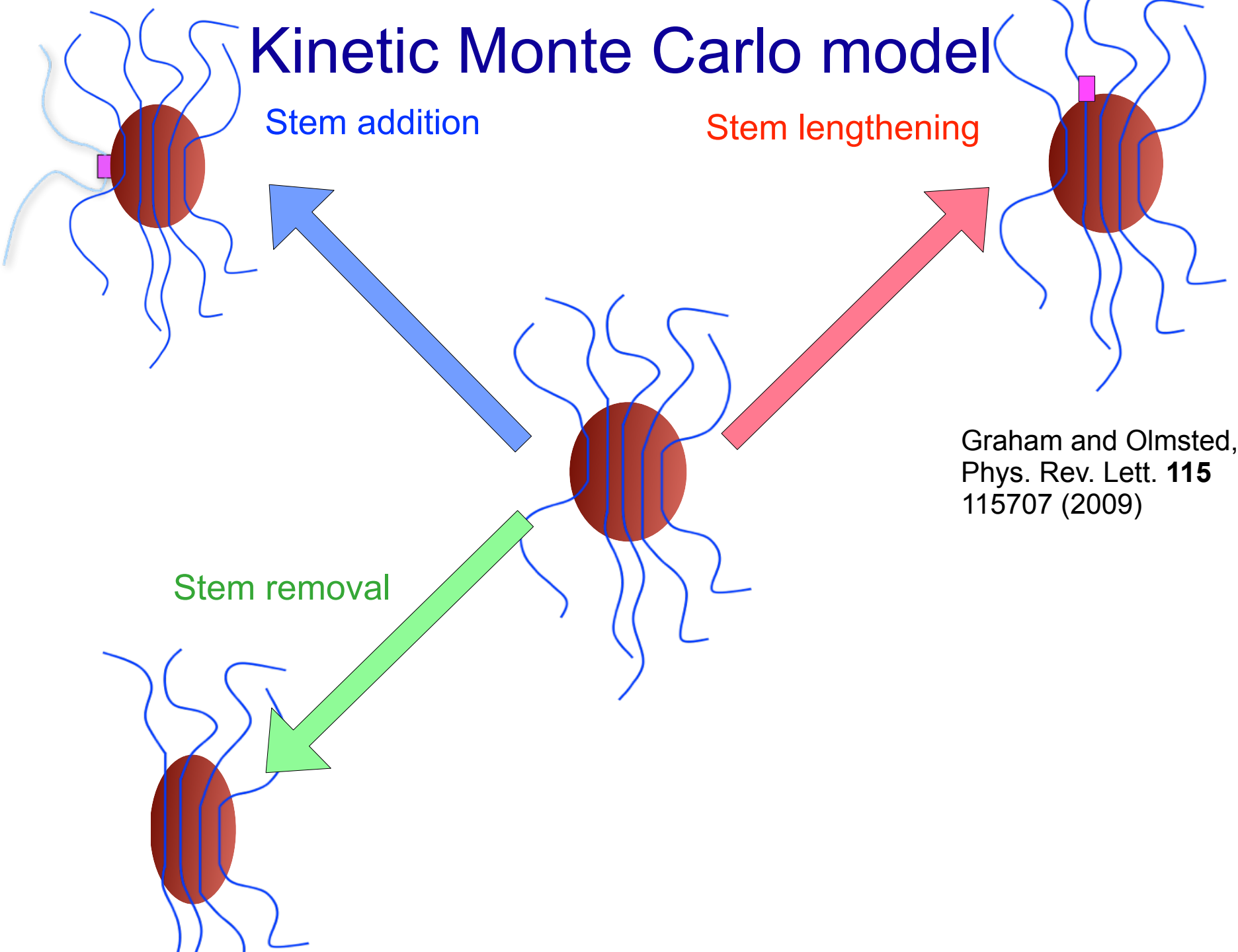
Graham and Olmsted,  
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115707 (2009)

# Kinetic Monte Carlo model



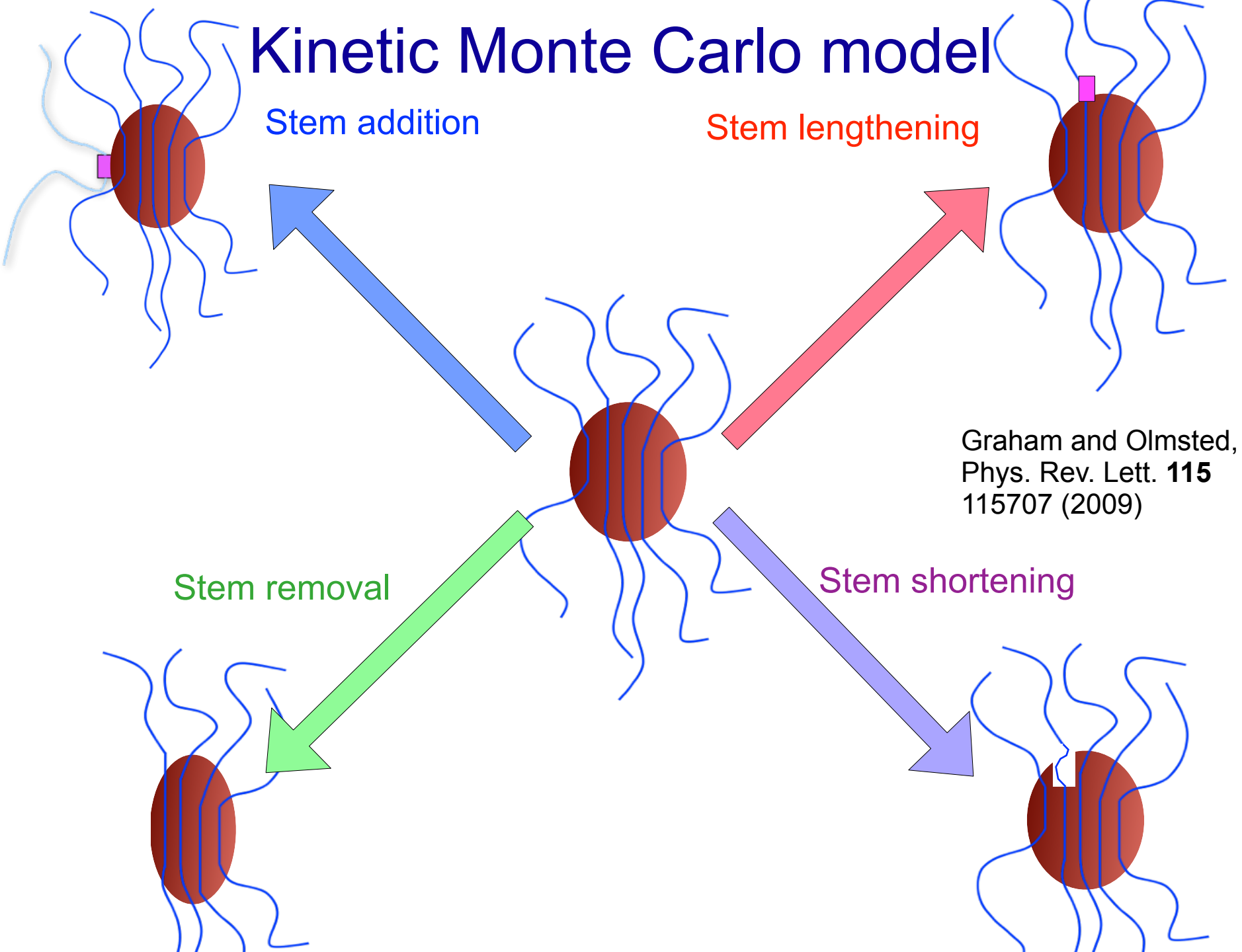
Graham and Olmsted,  
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115707 (2009)

# Kinetic Monte Carlo model



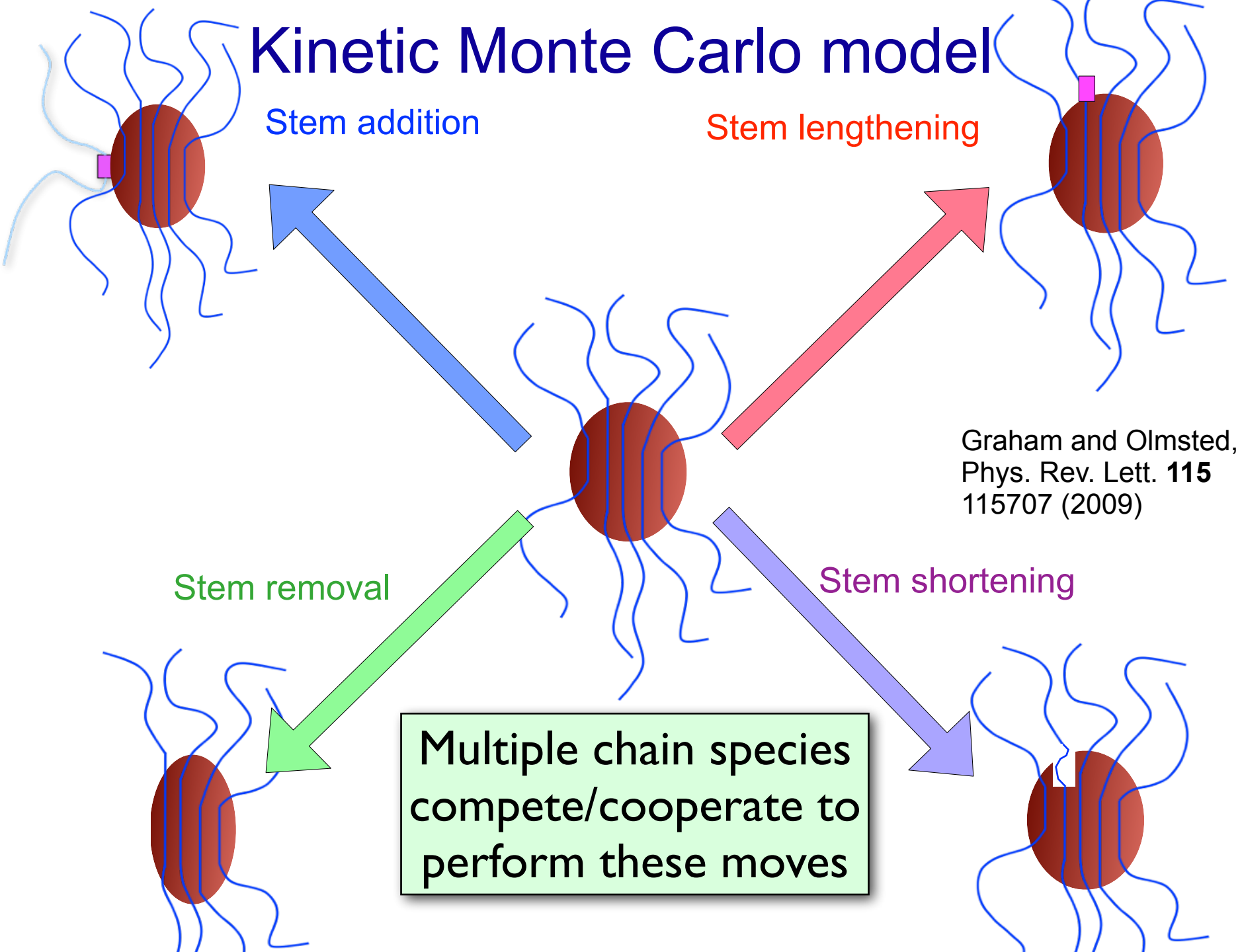
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Graham and Olmsted,  
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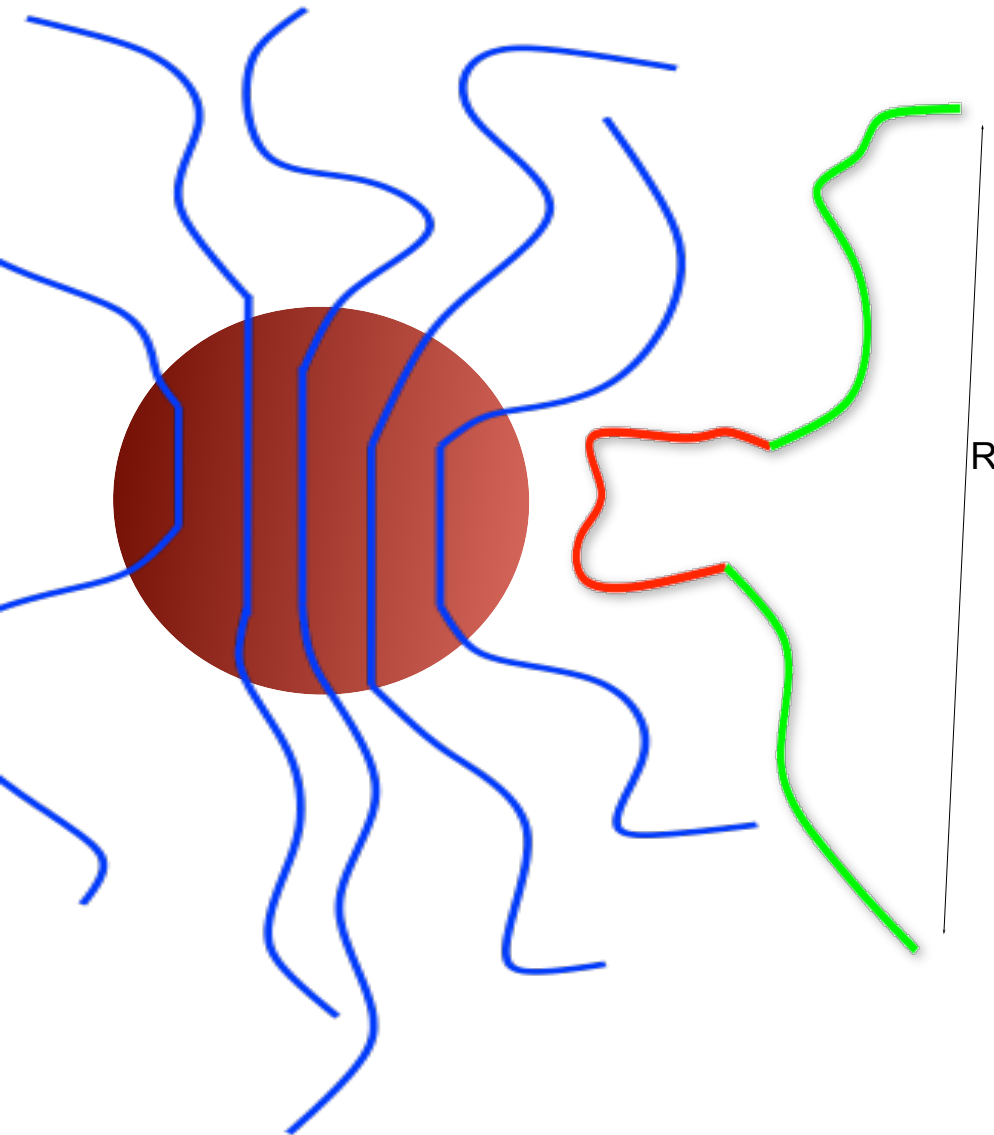
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115707 (2009)

# Influence of flow - chain deformation

As computed by a  
molecular flow theory.



Flow imposes an  
average molecular  
strain,  $\langle RR \rangle$ .

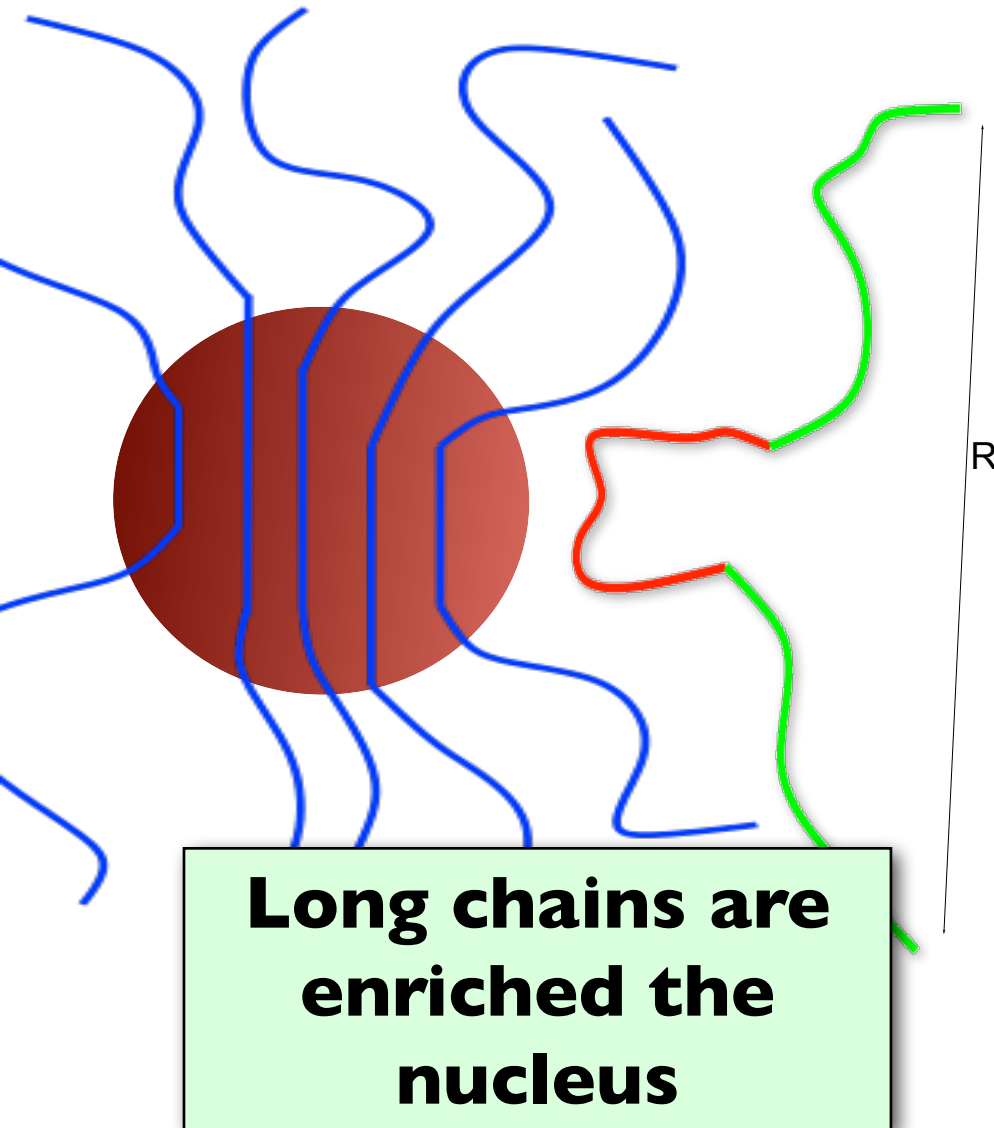


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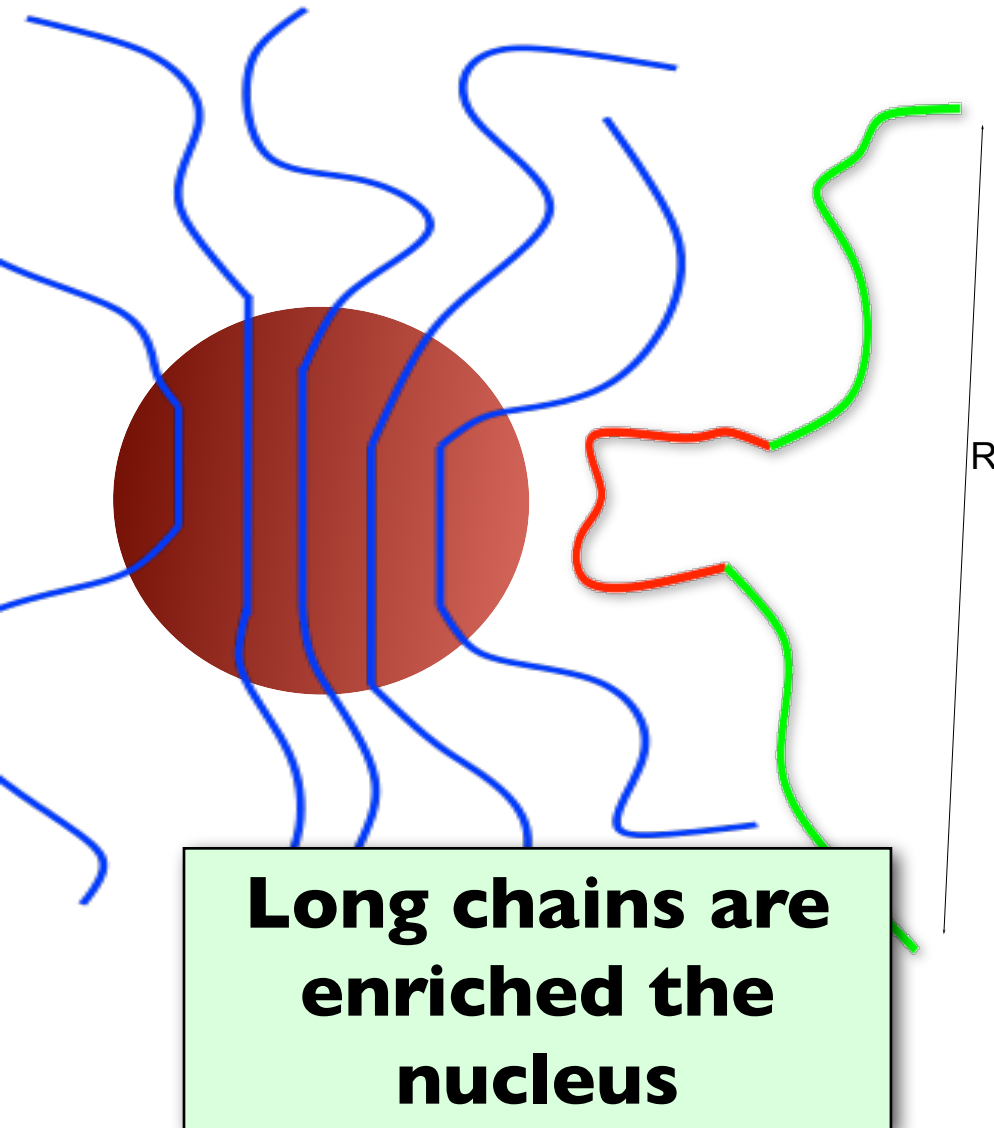


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Monomer

attachment rate:  $k_i^+ \propto \exp(\Delta f_i)$

**Long chains are  
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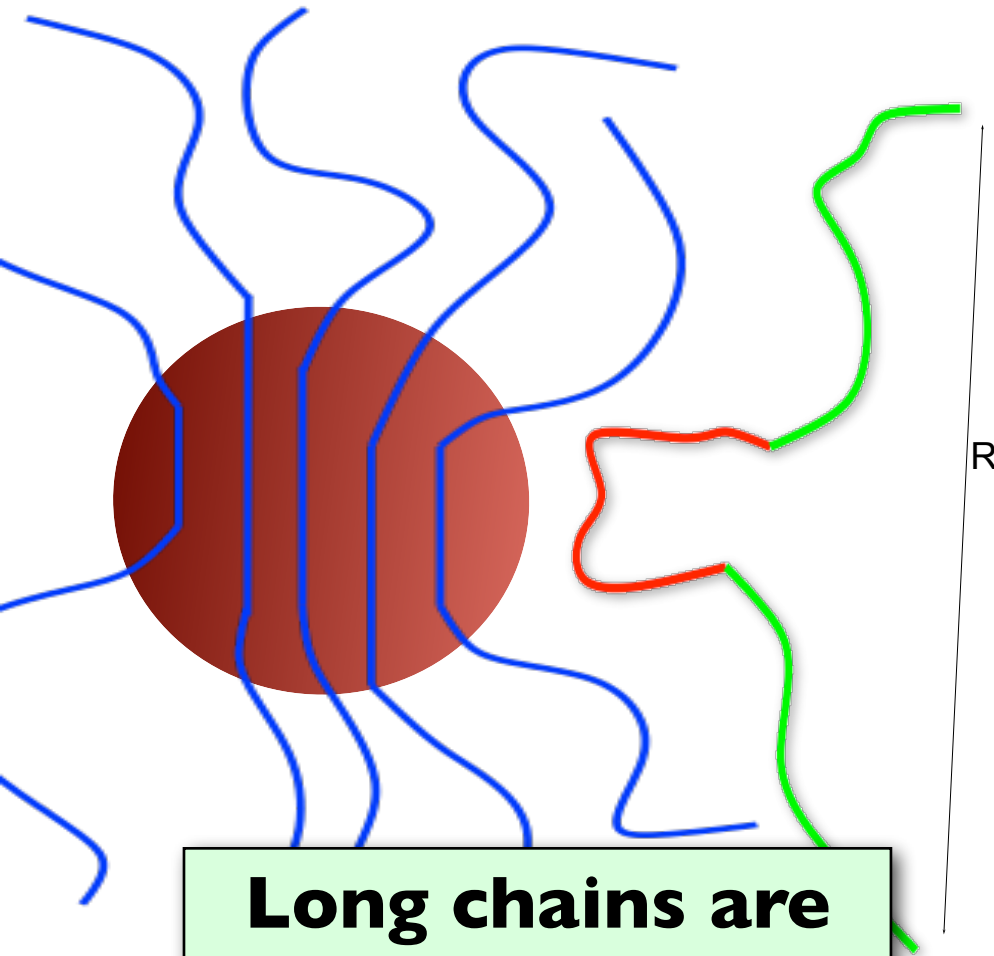


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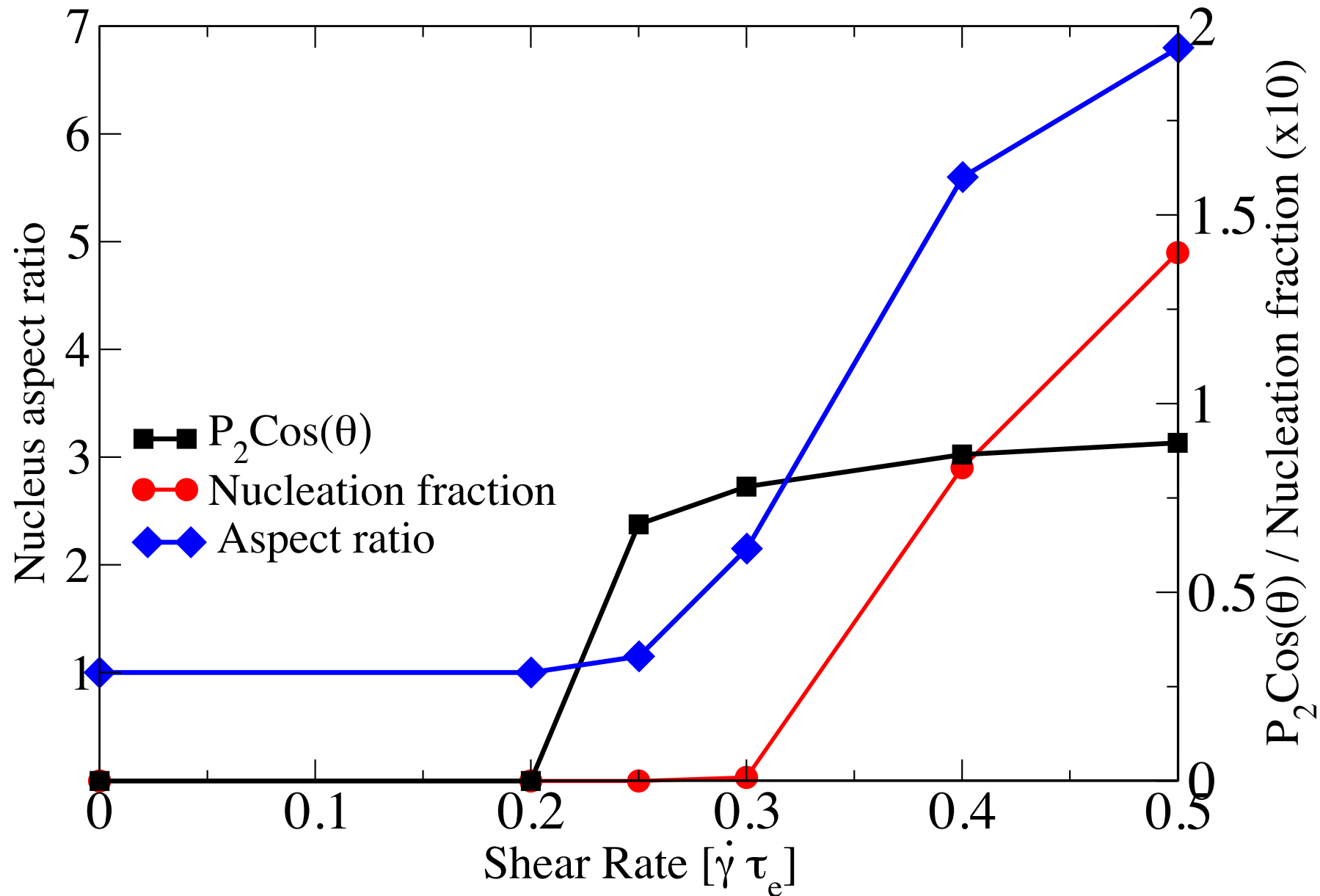
attachment rate:  $k_i^+ \propto \exp(\Delta f_i)$

Free energy  
change on  
deformation:

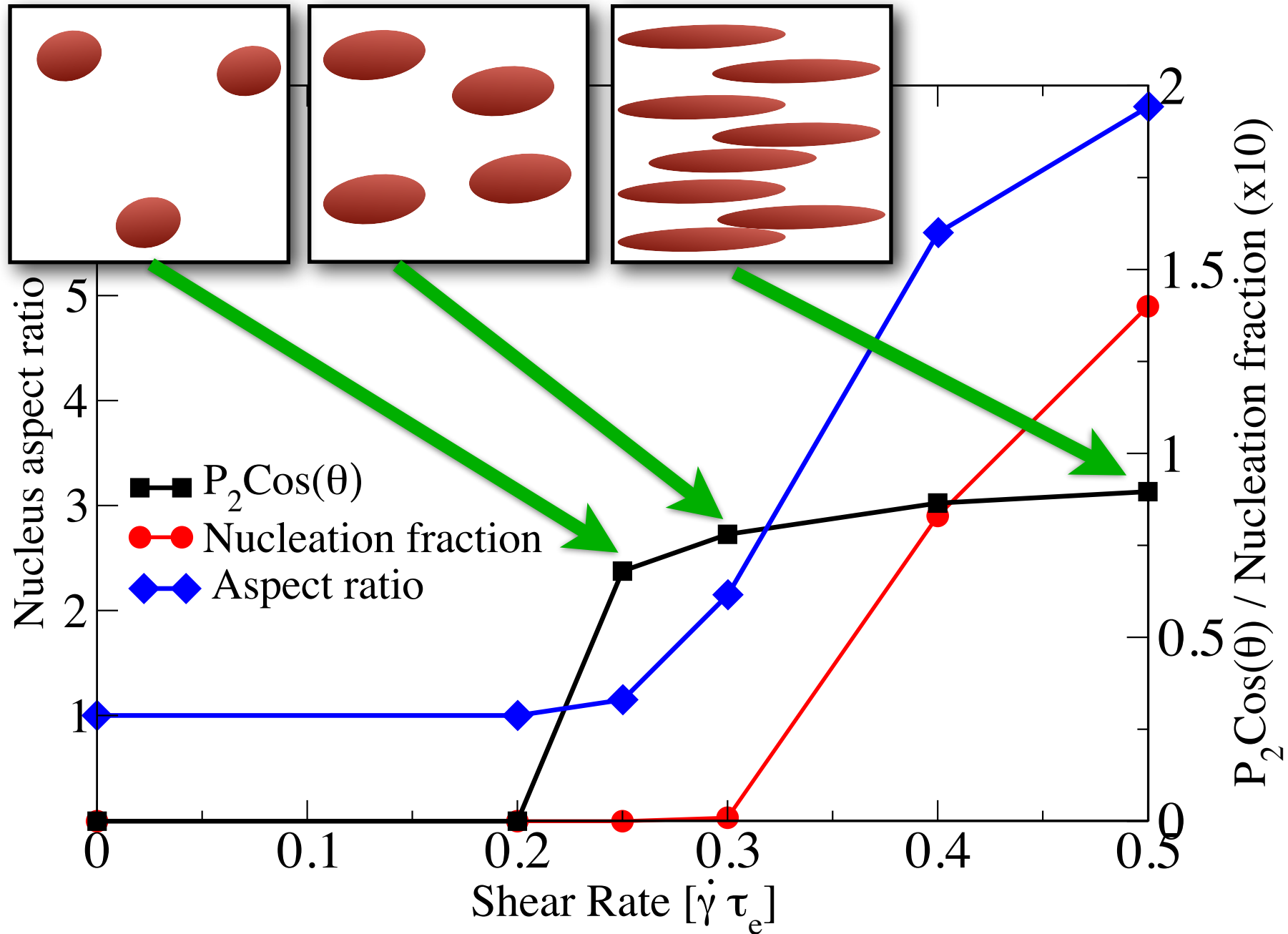
$\Delta f_i = \Gamma P_{2,K,i}$

Nematic order

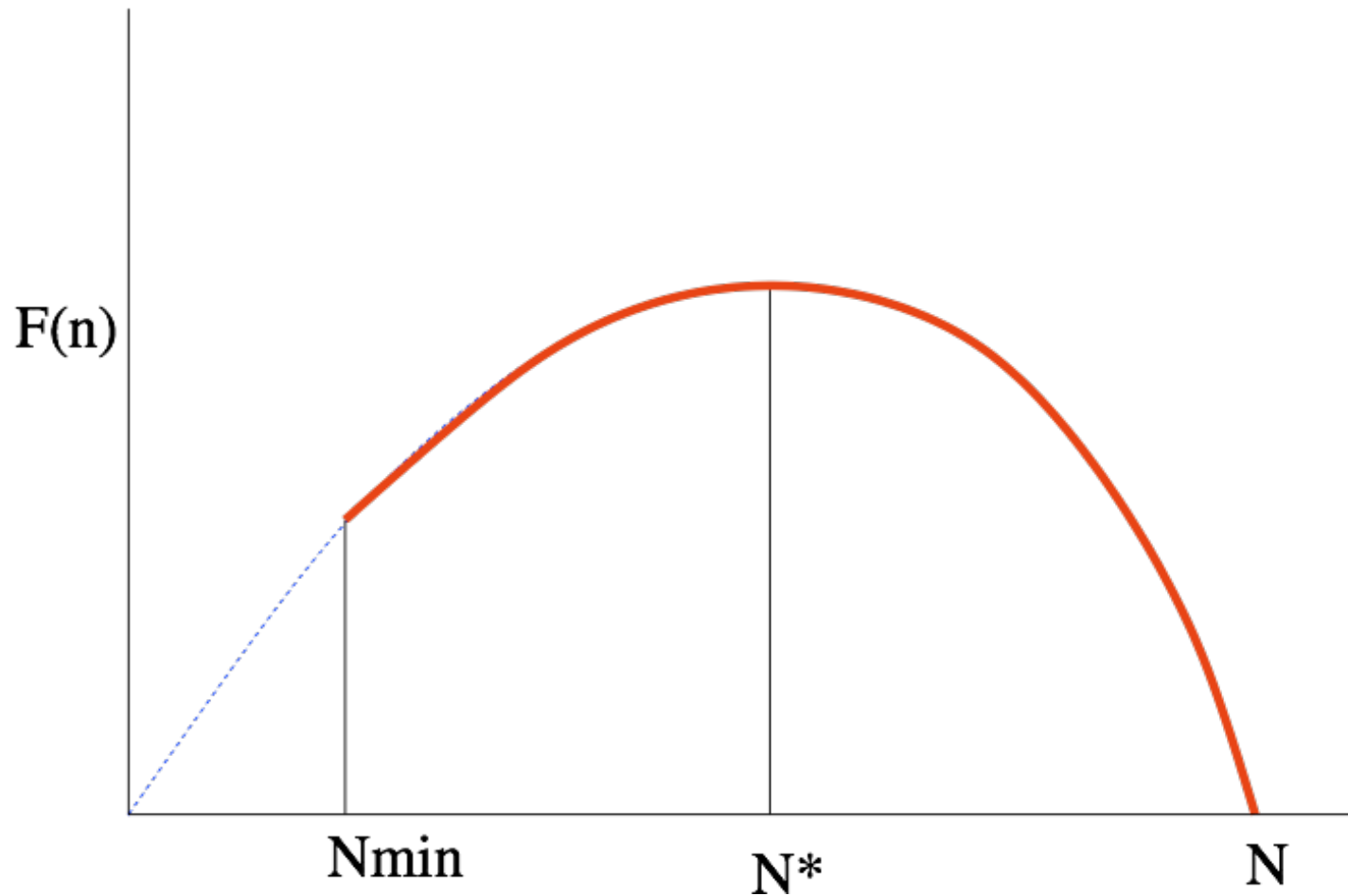
# Effect of flow on nucleation



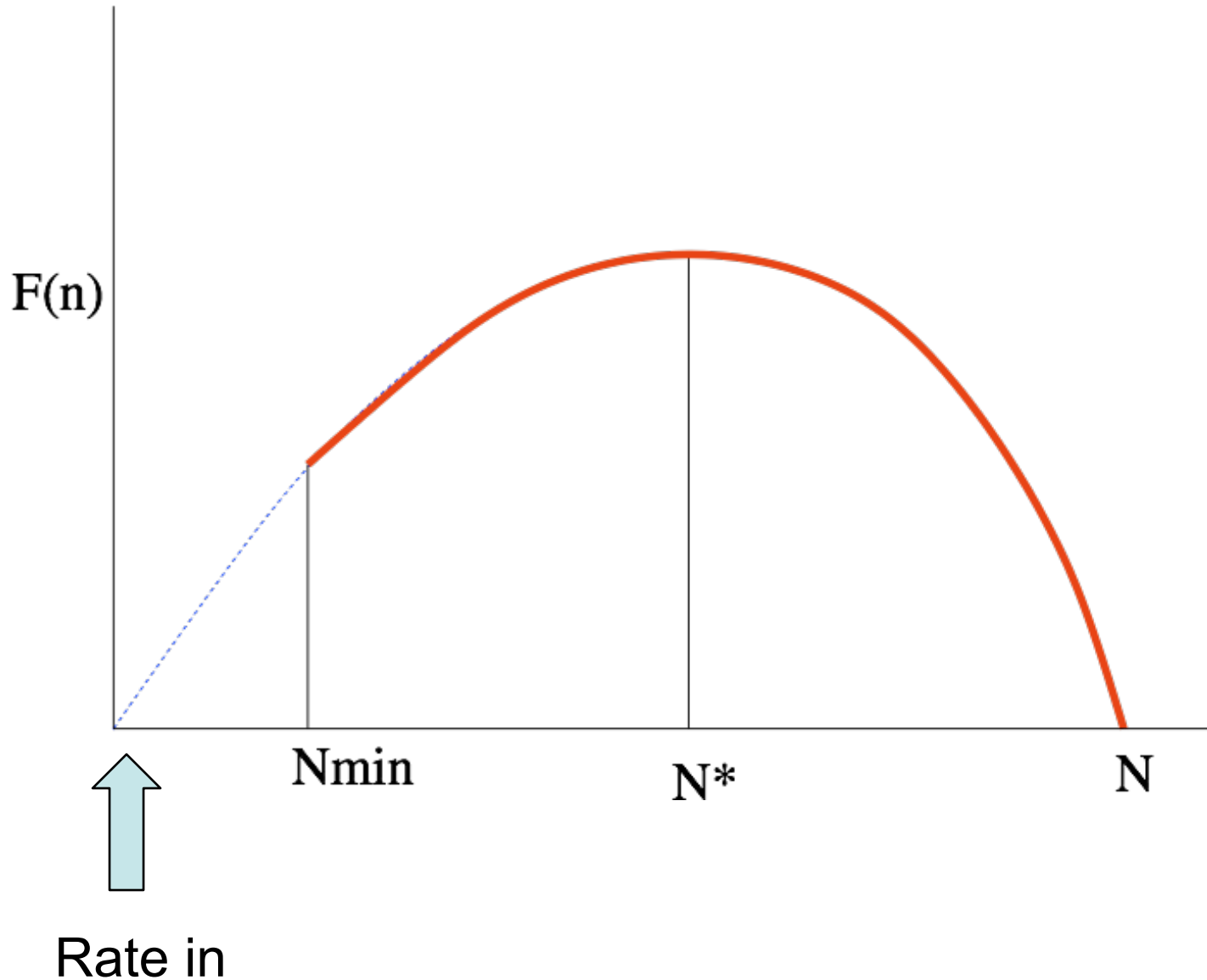
# Effect of flow on nucleation



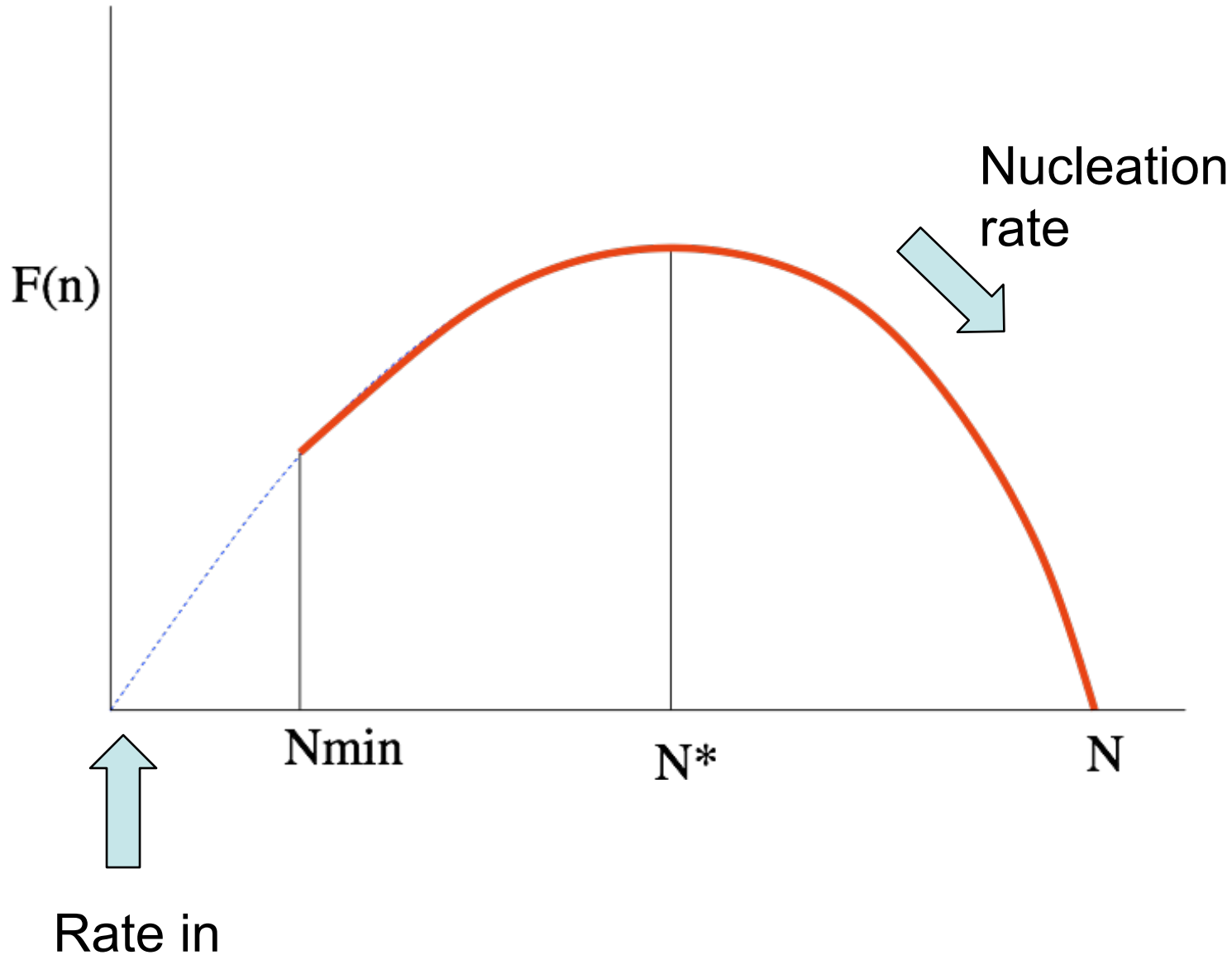
# a) Faster nucleation simulation



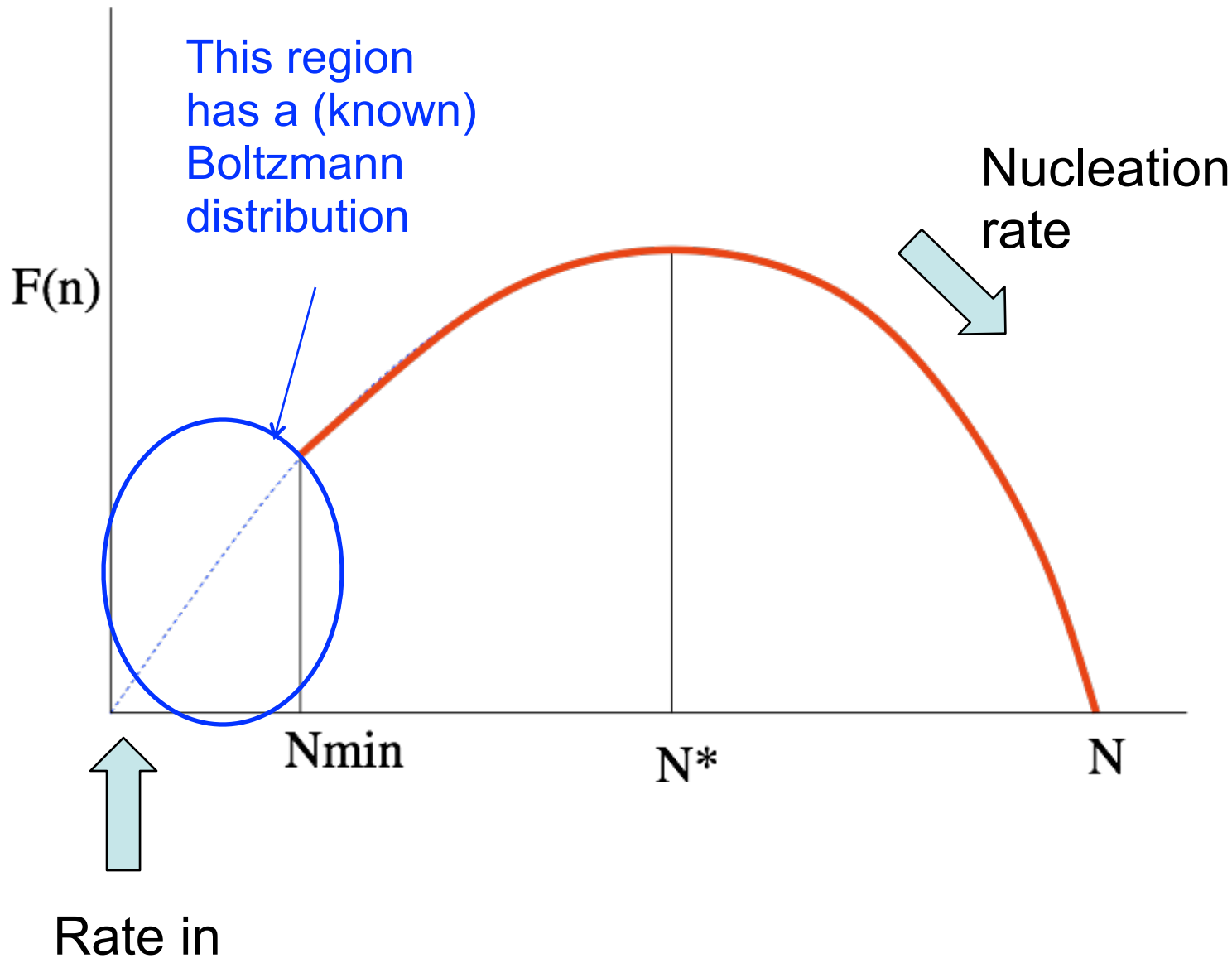
# a) Faster nucleation simulation



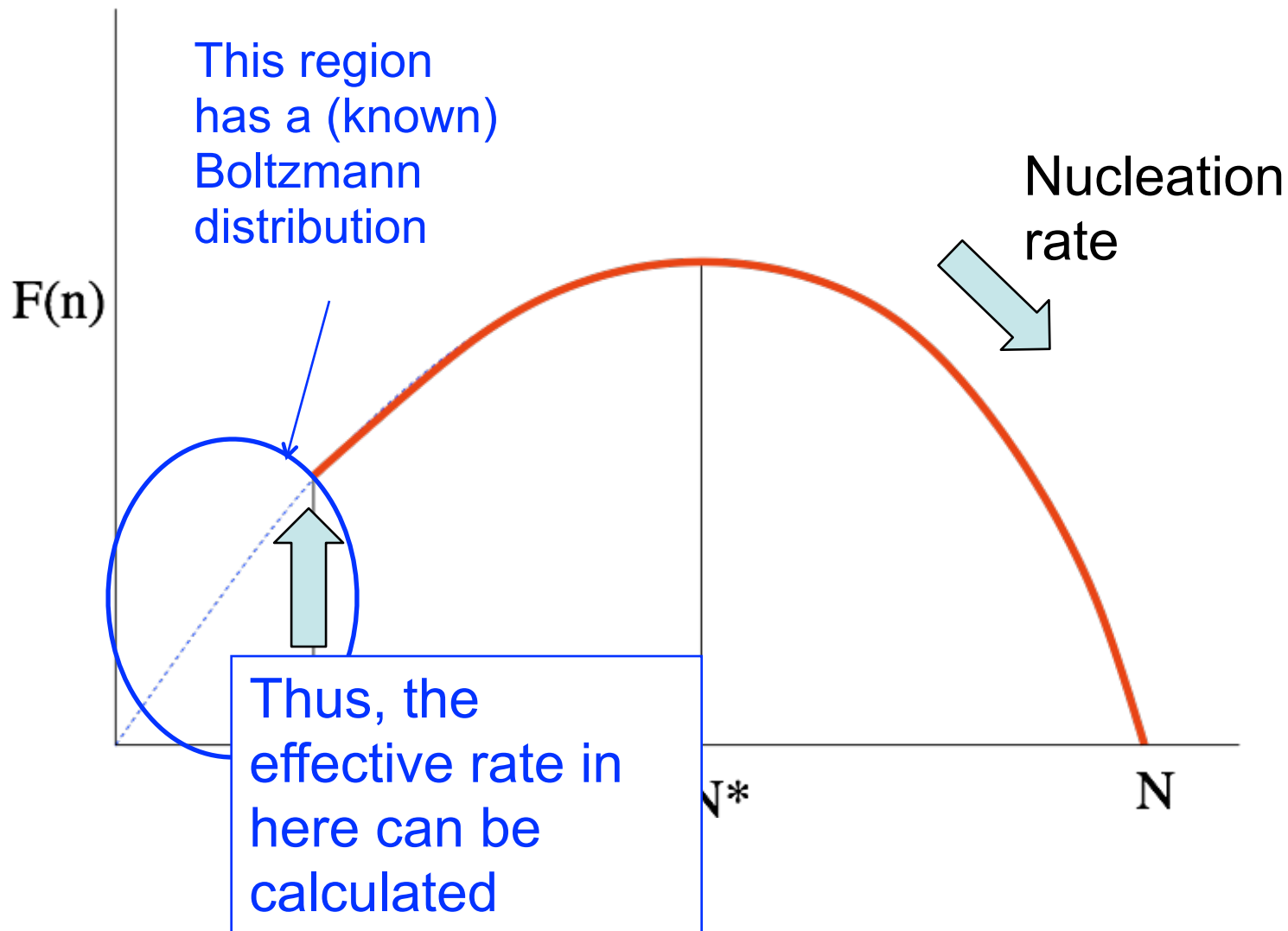
# a) Faster nucleation simulation



# a) Faster nucleation simulation

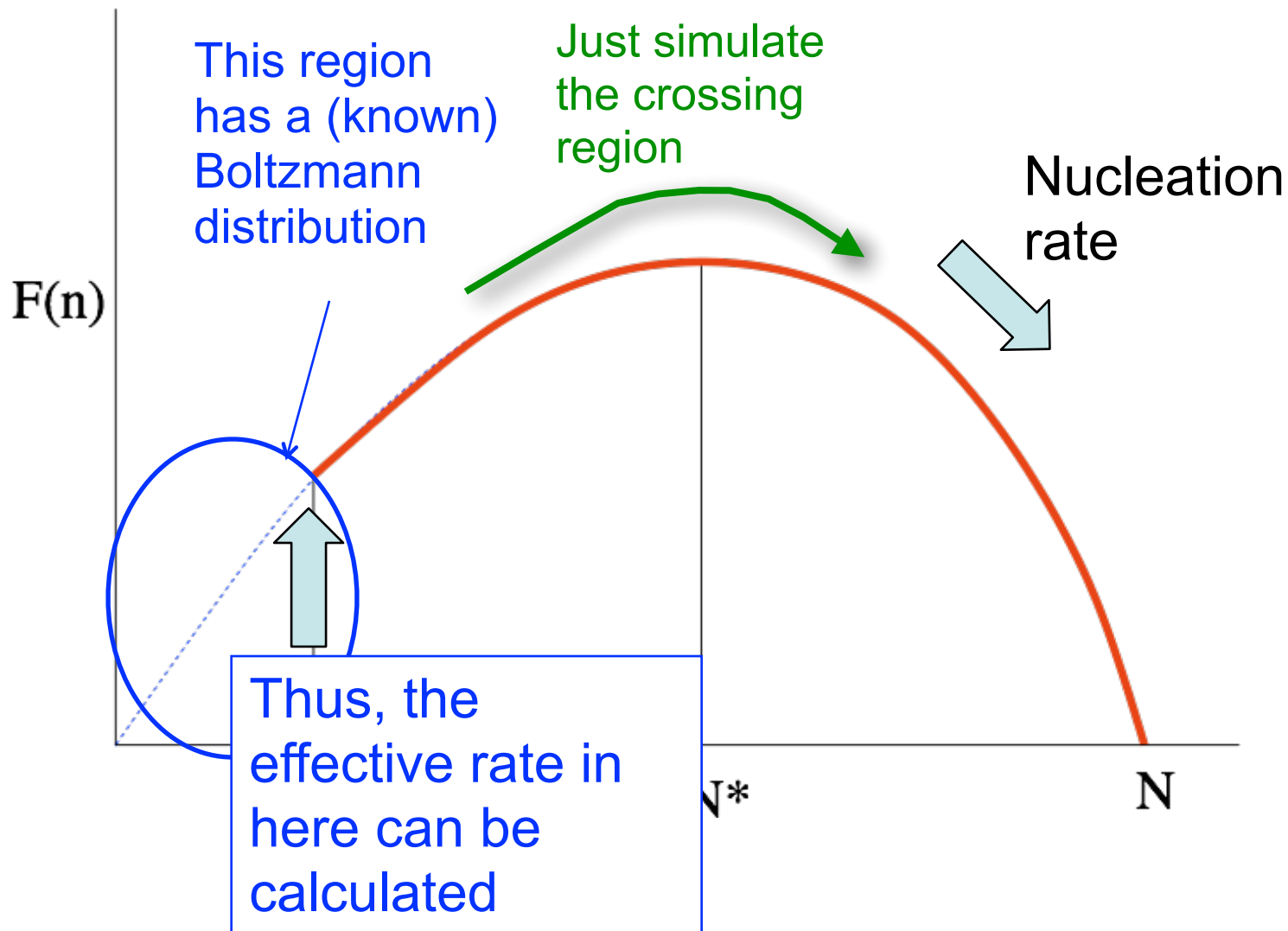


# a) Faster nucleation simulation

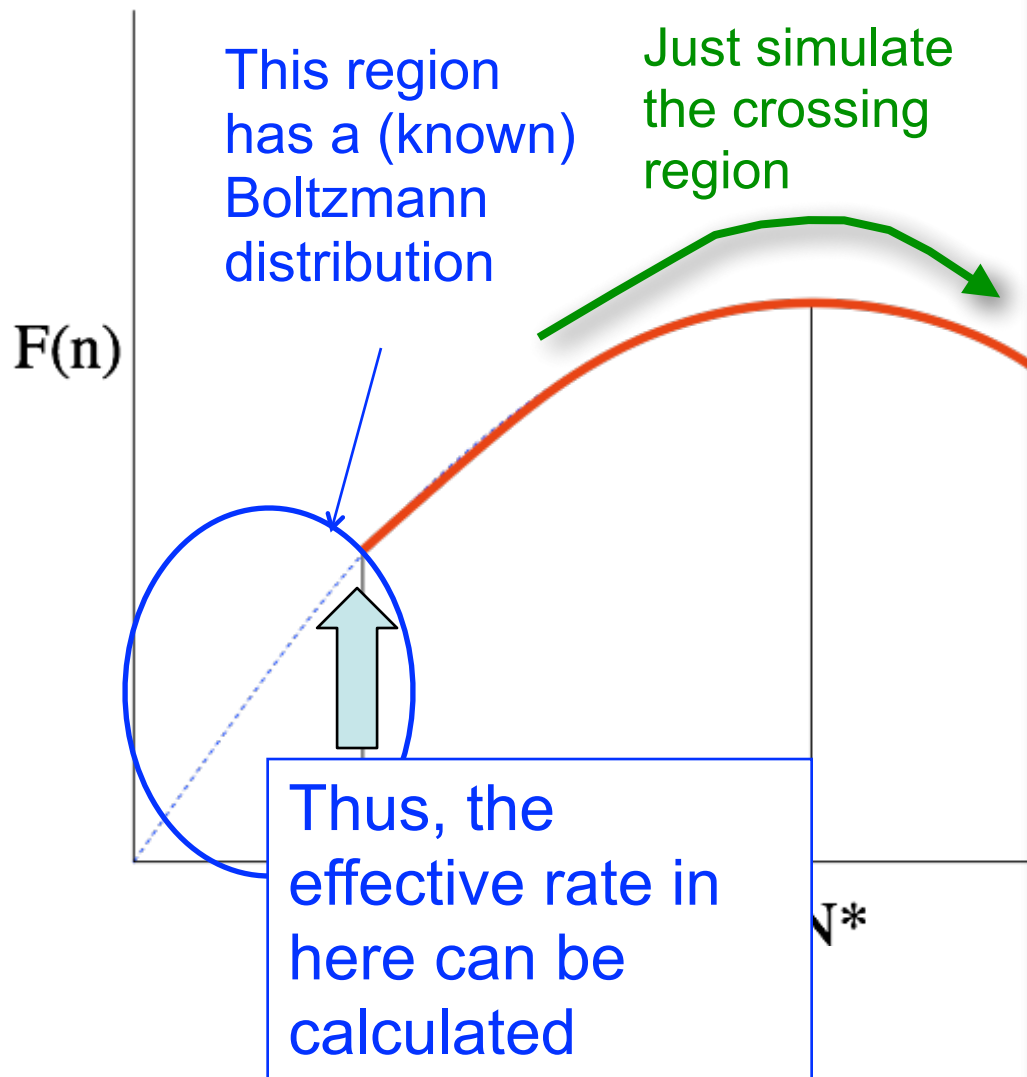




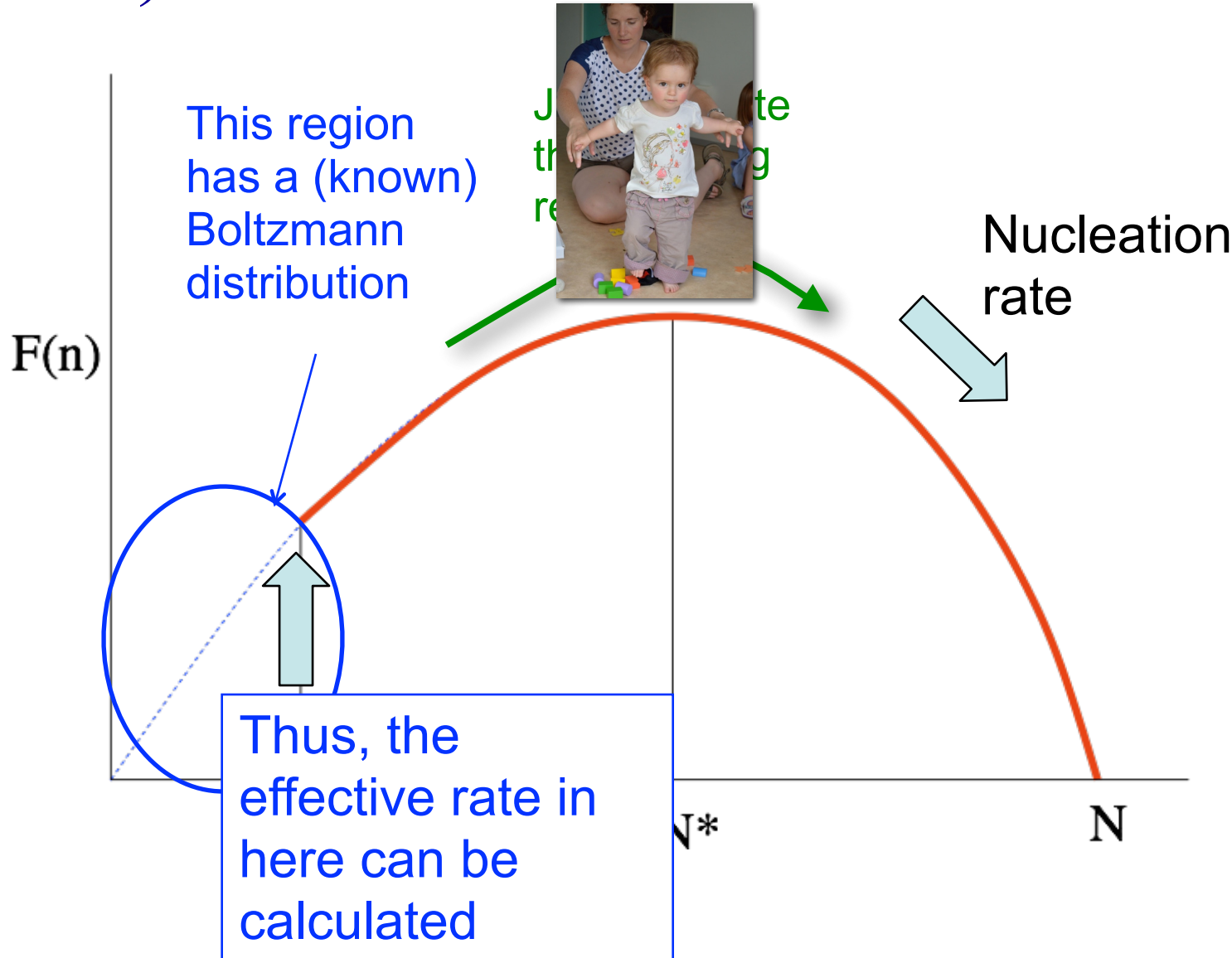
# a) Faster nucleation simulation



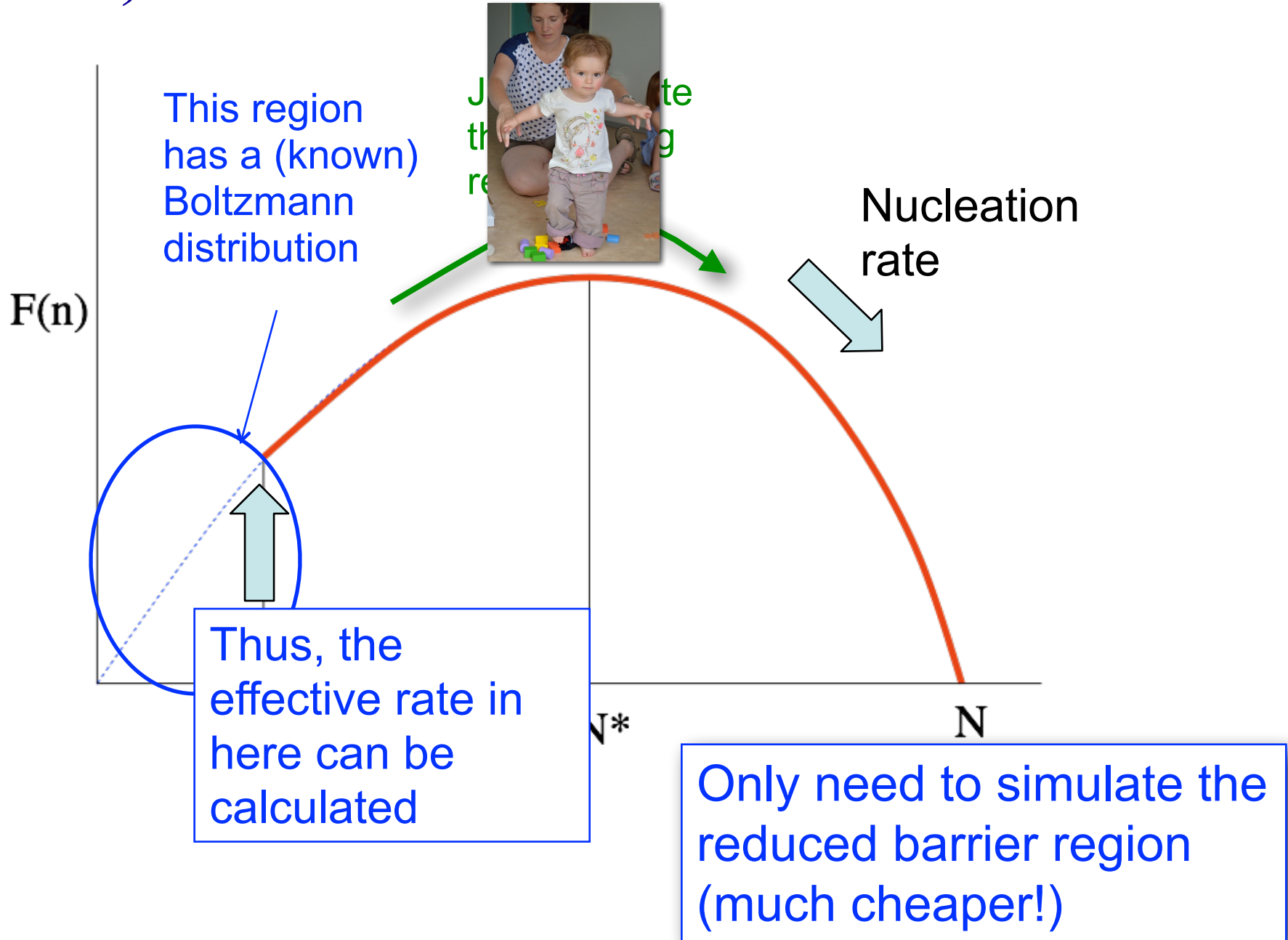
# a) Faster nucleation simulation



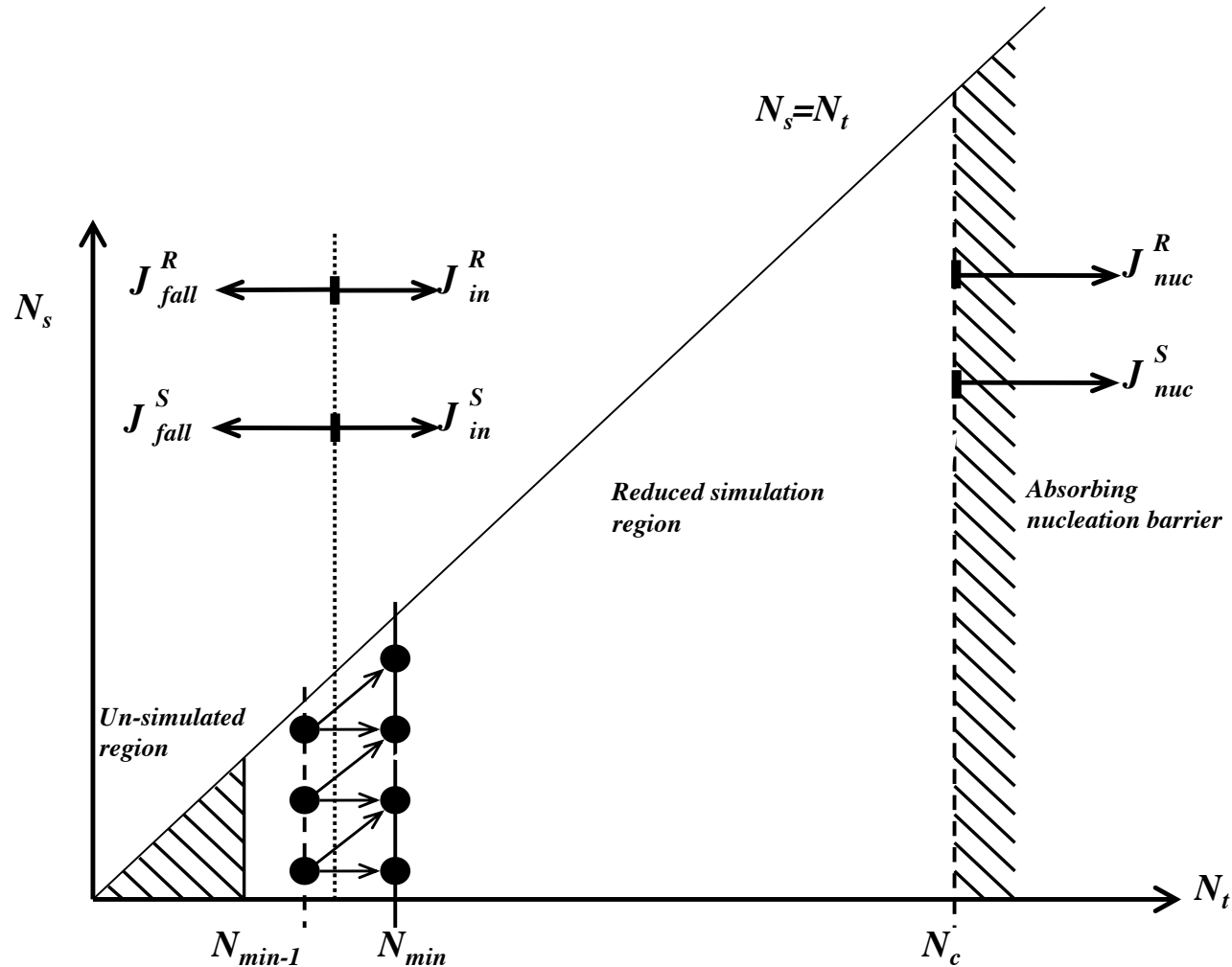
# a) Faster nucleation simulation



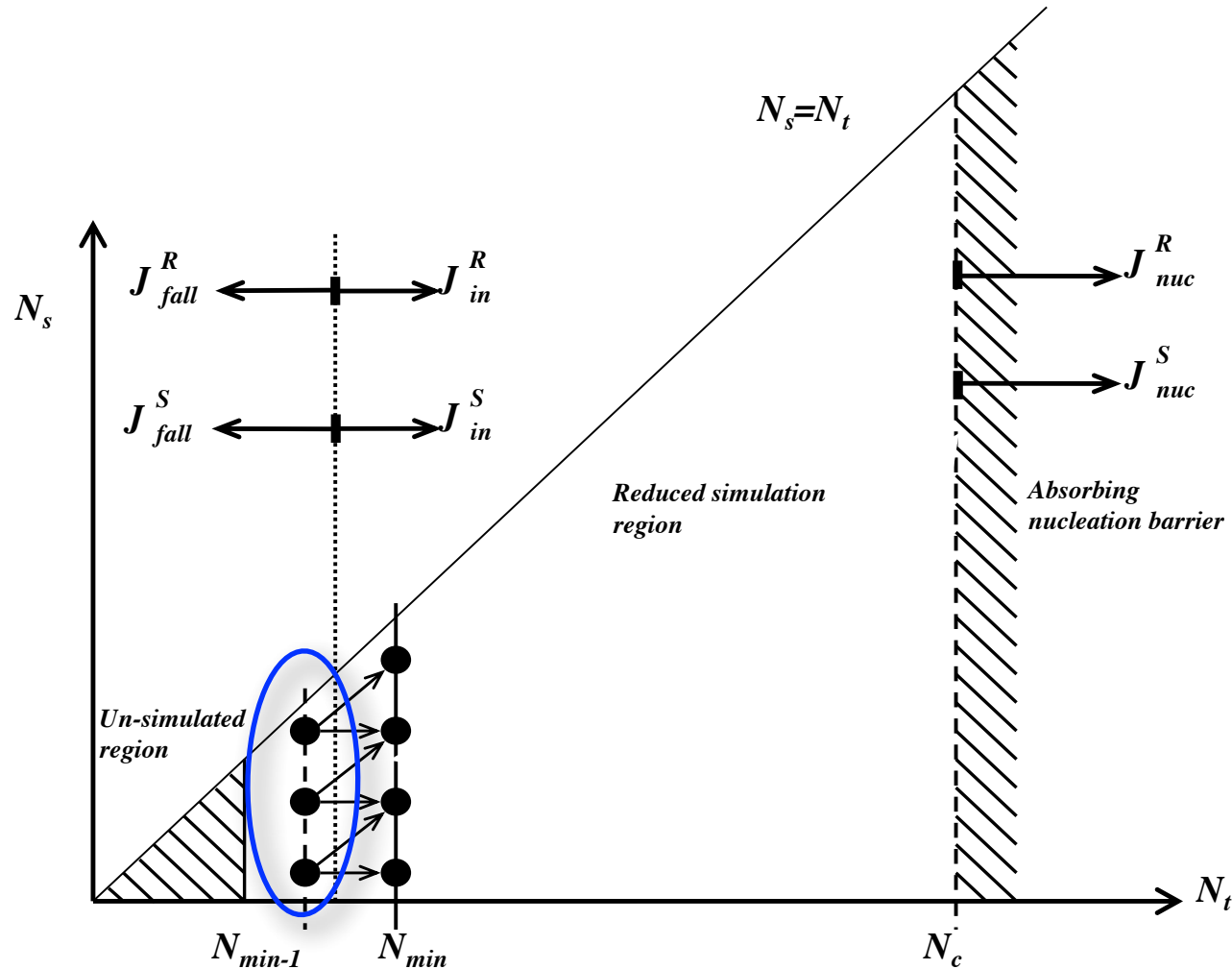
# a) Faster nucleation simulation



# Generalisation to higher dimensions



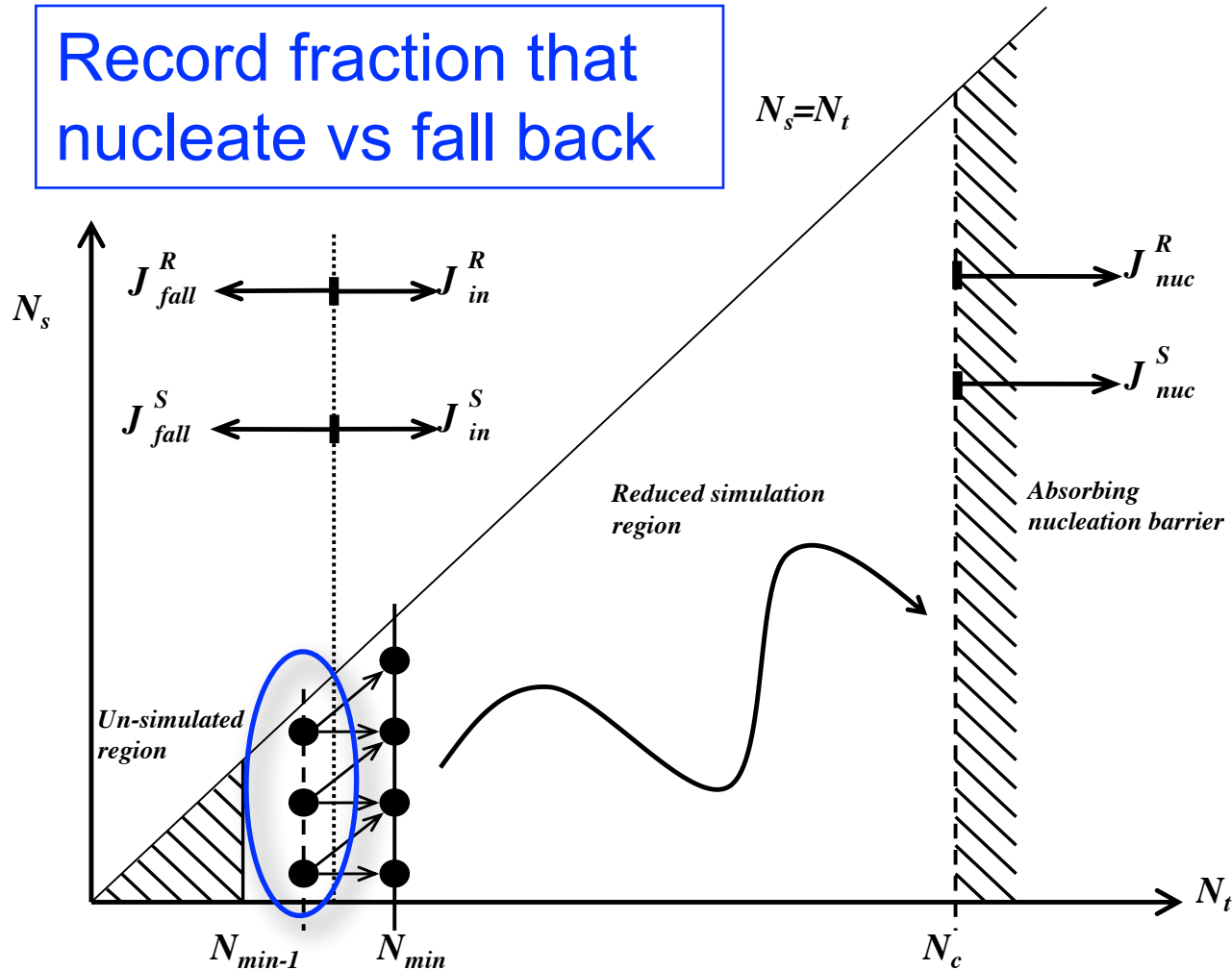
# Generalisation to higher dimensions



Inject nuclei with  
a Boltzmann  
distribution

# Generalisation to higher dimensions

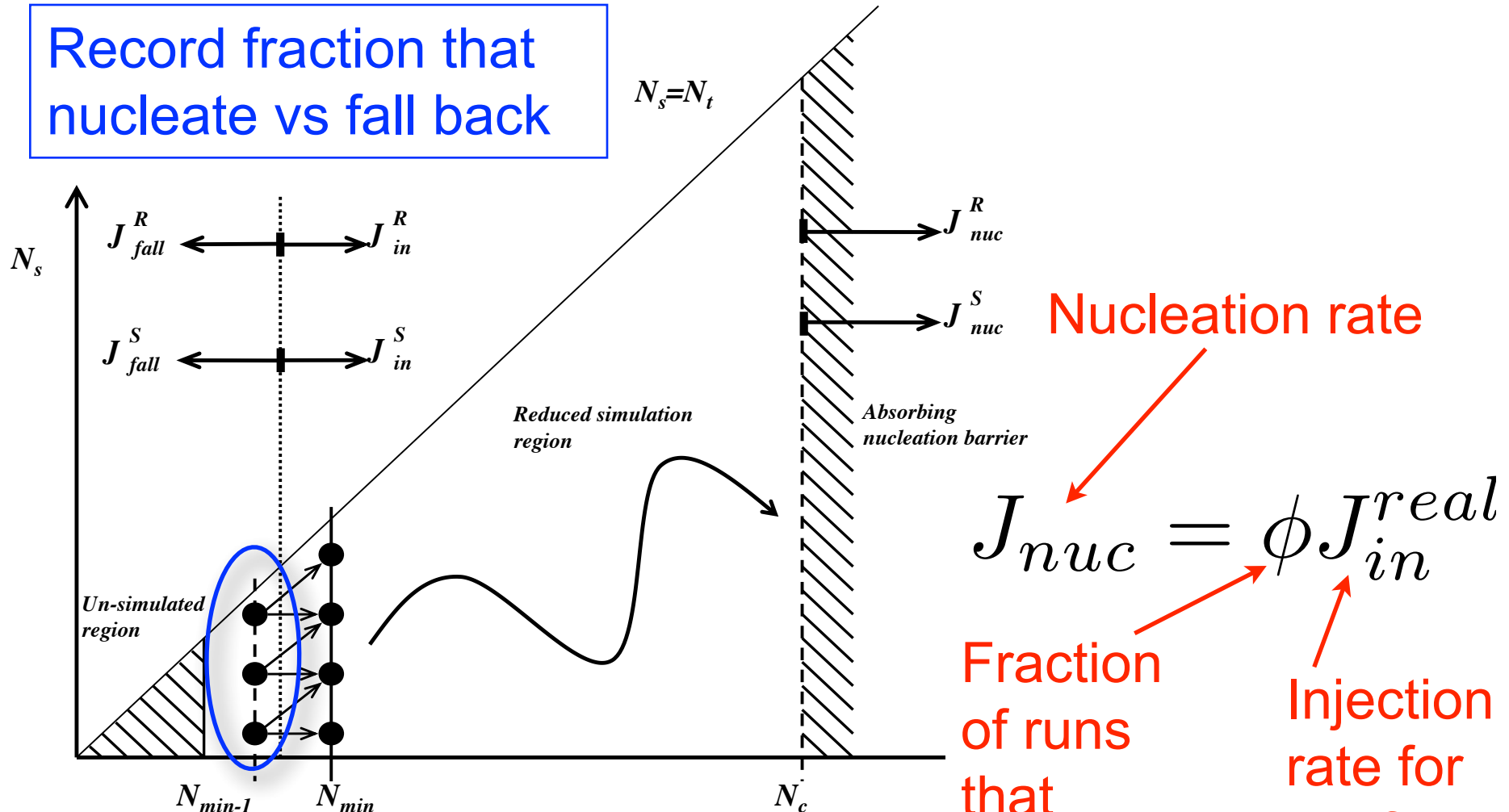
Record fraction that nucleate vs fall back



Inject nuclei with a Boltzmann distribution

# Generalisation to higher dimensions

Record fraction that nucleate vs fall back



Nucleation rate

$$J_{nuc} = \phi J_{in}^{real}$$

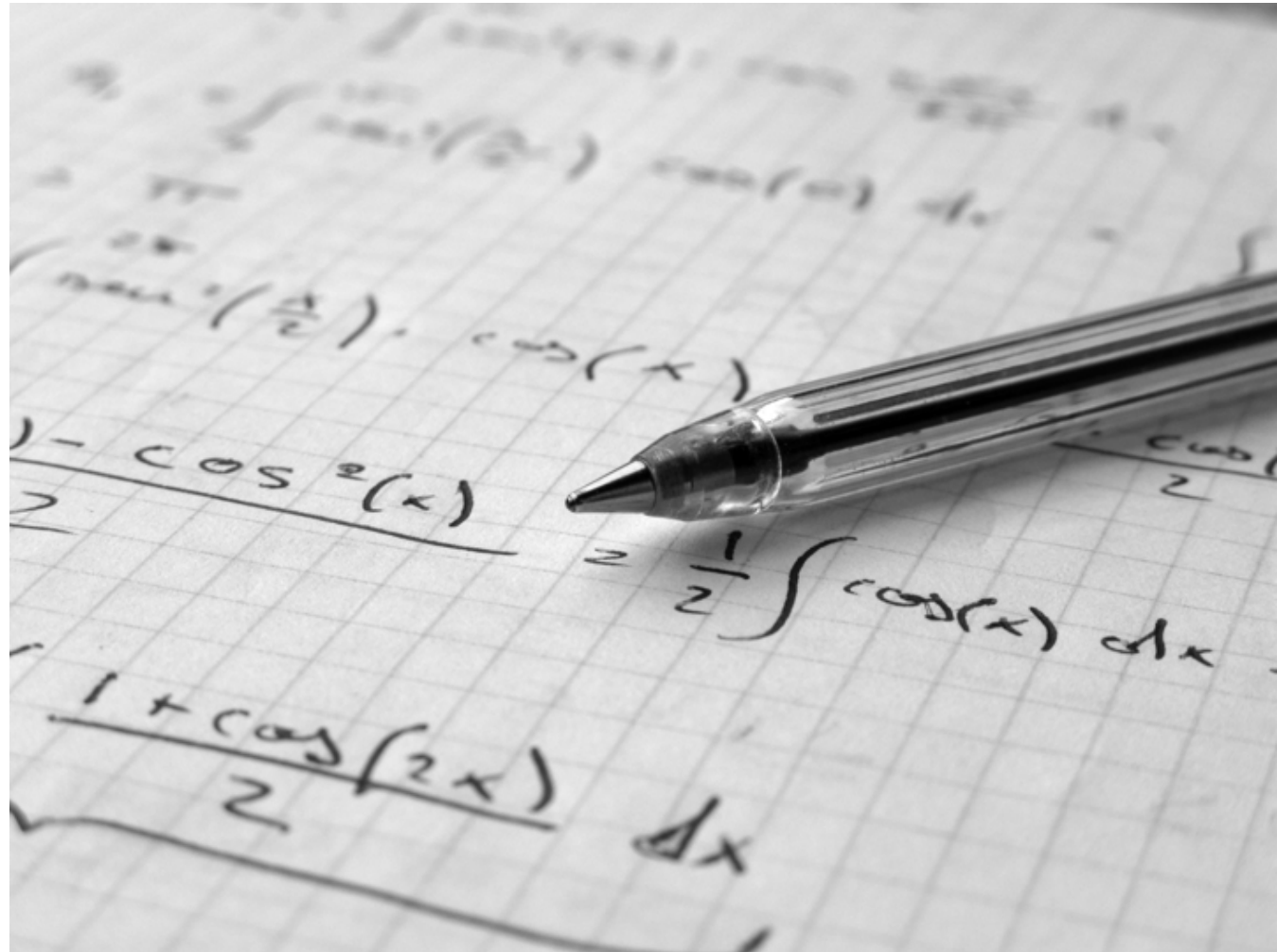
Fraction of runs that nucleate

Injection rate for the full system

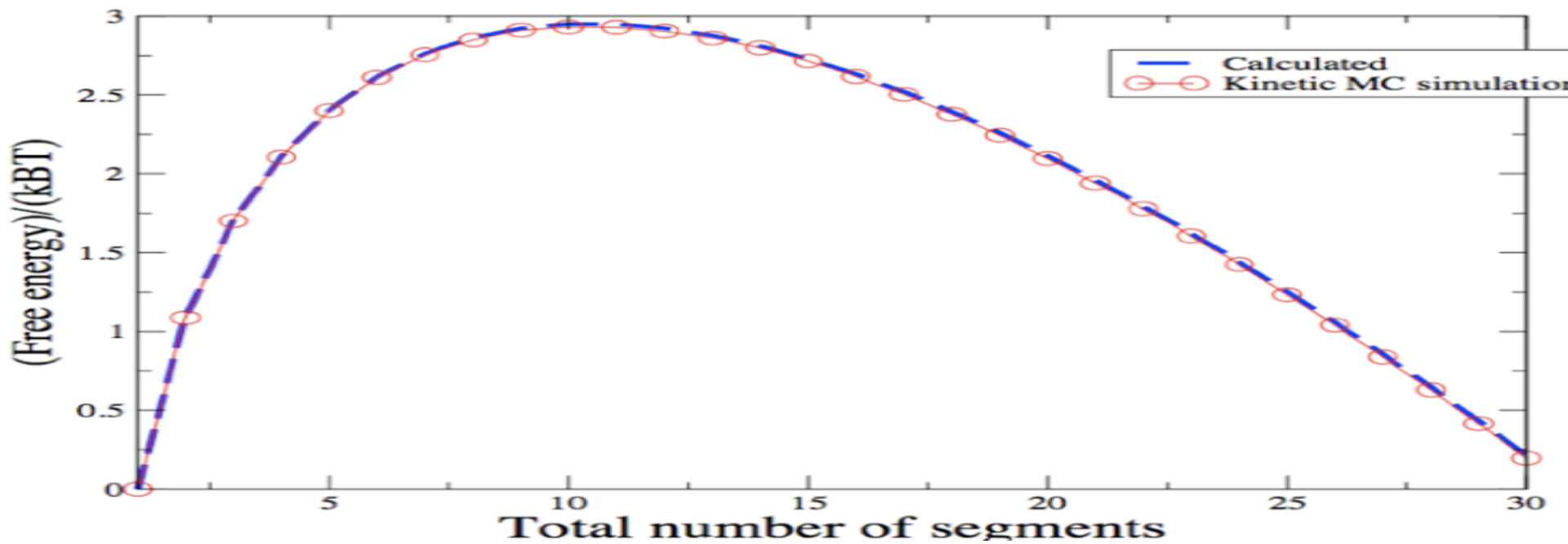
Inject nuclei with a Boltzmann distribution



# Continuum model (nearly analytic)

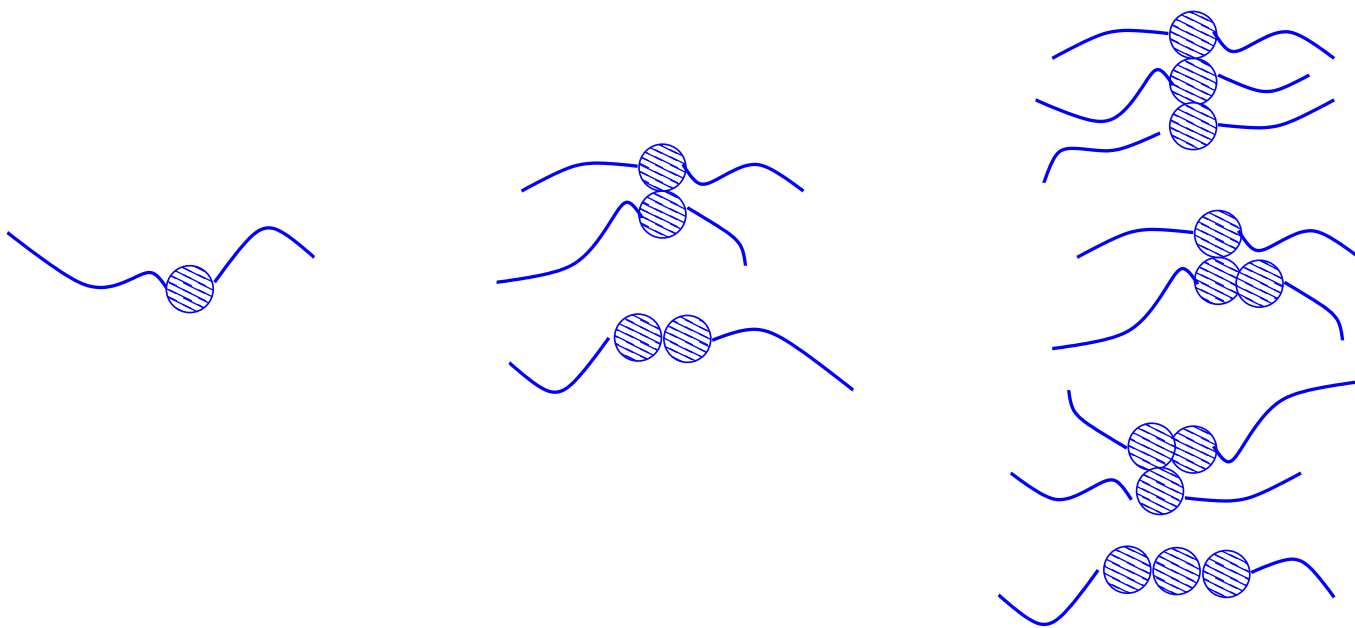
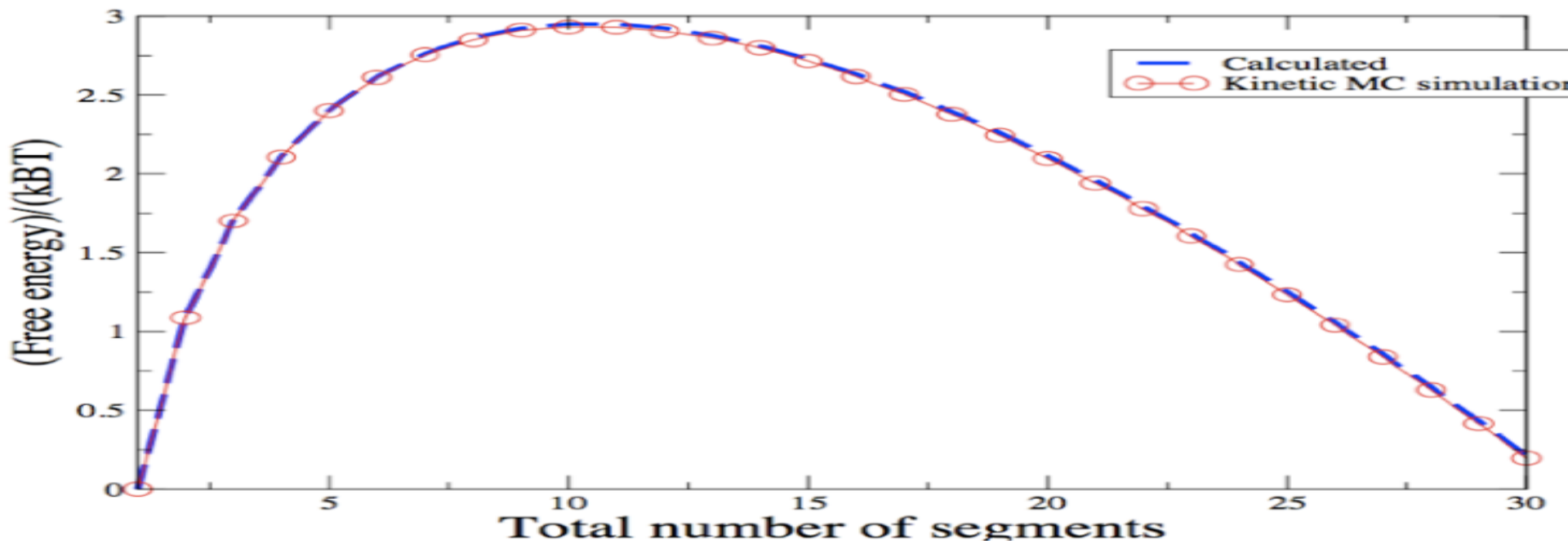


# Projecting the GO model to a 1D system



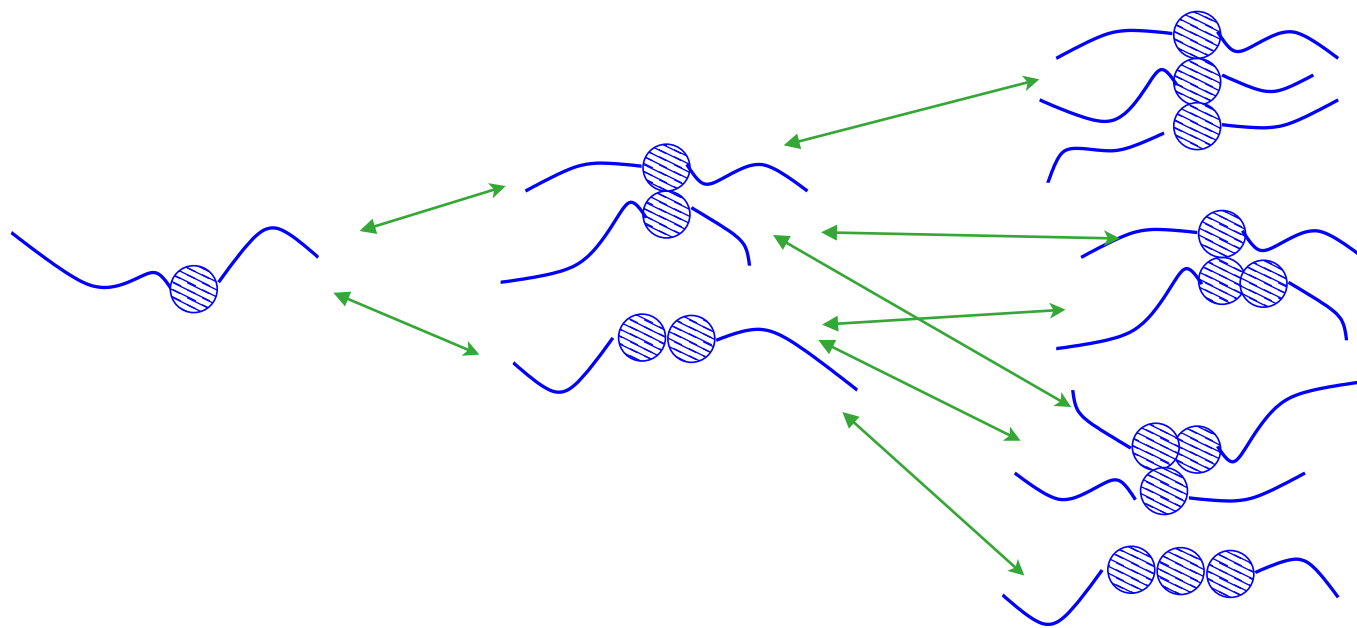
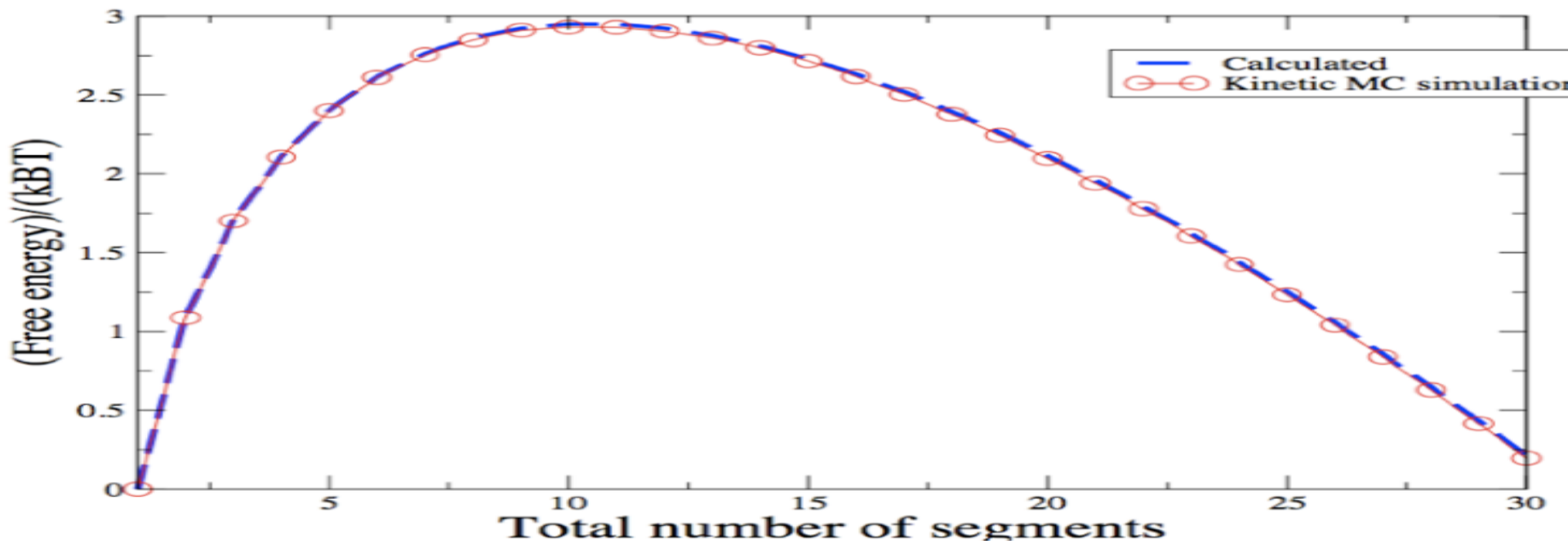
Hamer, Wattis and  
Graham, *Soft Matter*  
**8**, 11396 (2012)

# Projecting the GO model to a 1D system



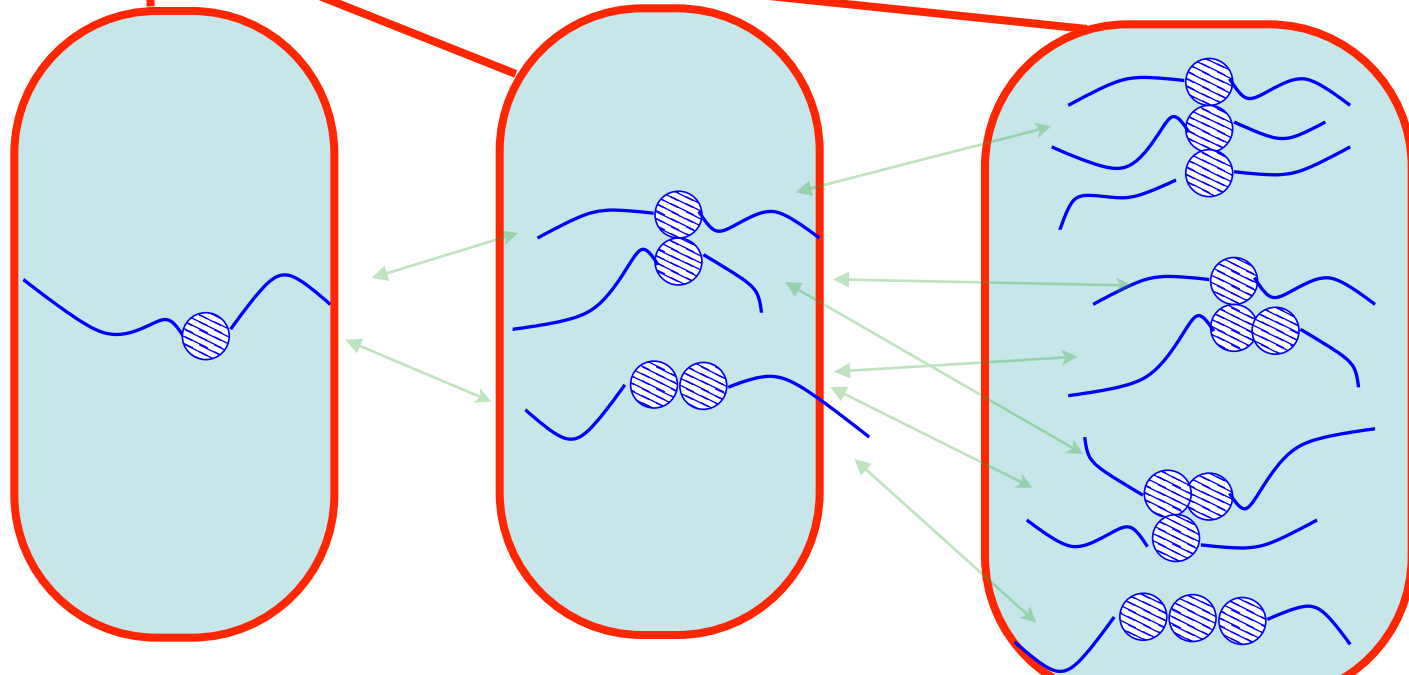
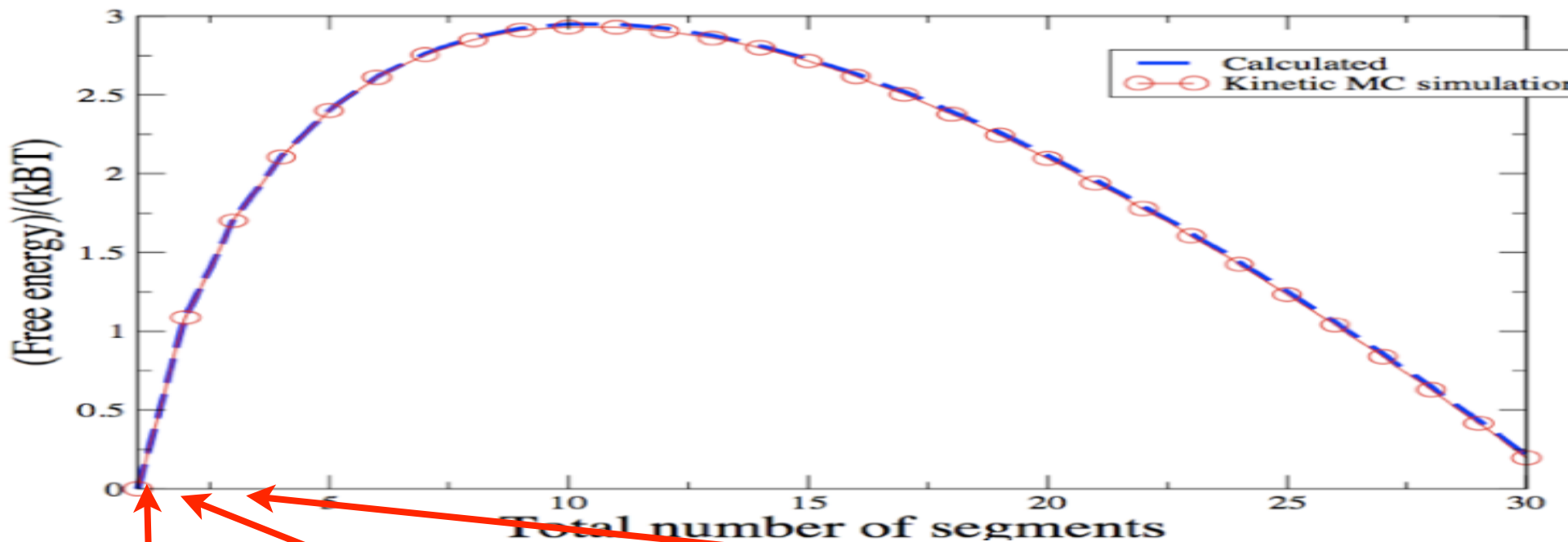
Hamer, Wattis and  
Graham, *Soft Matter*  
**8**, 11396 (2012)

# Projecting the GO model to a 1D system



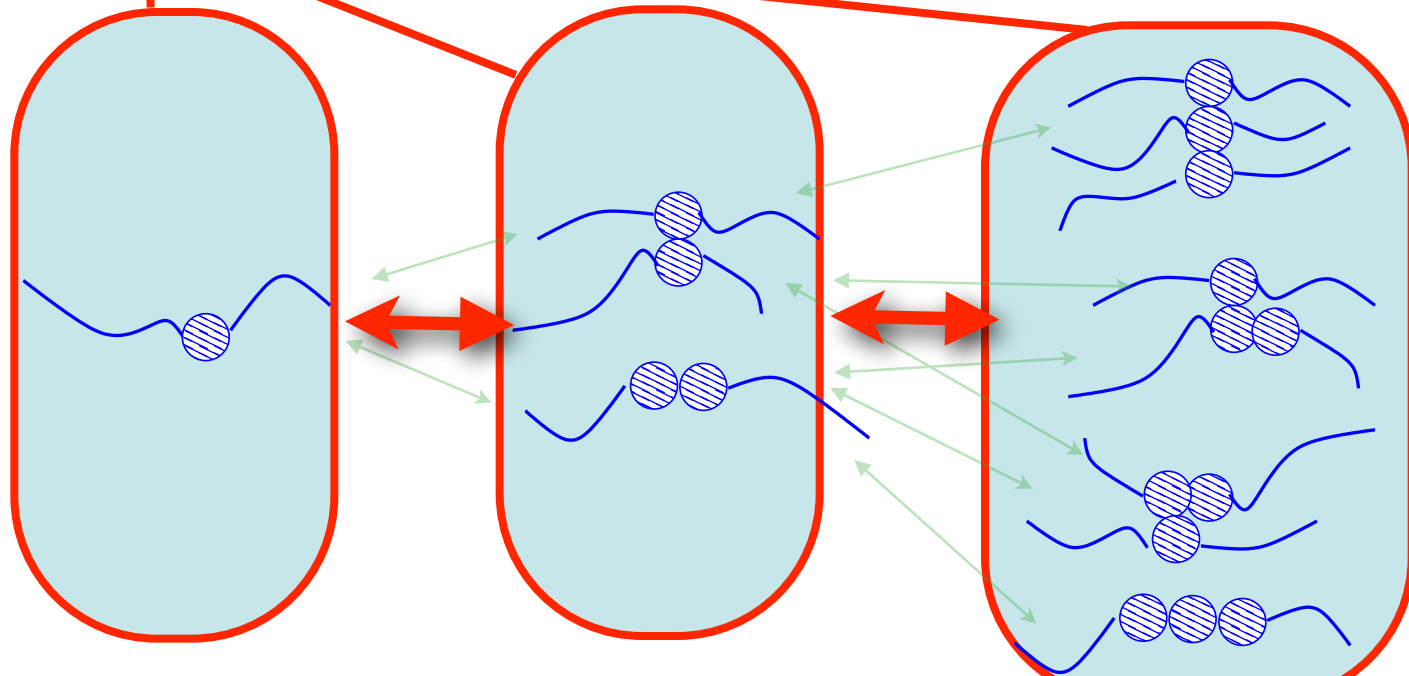
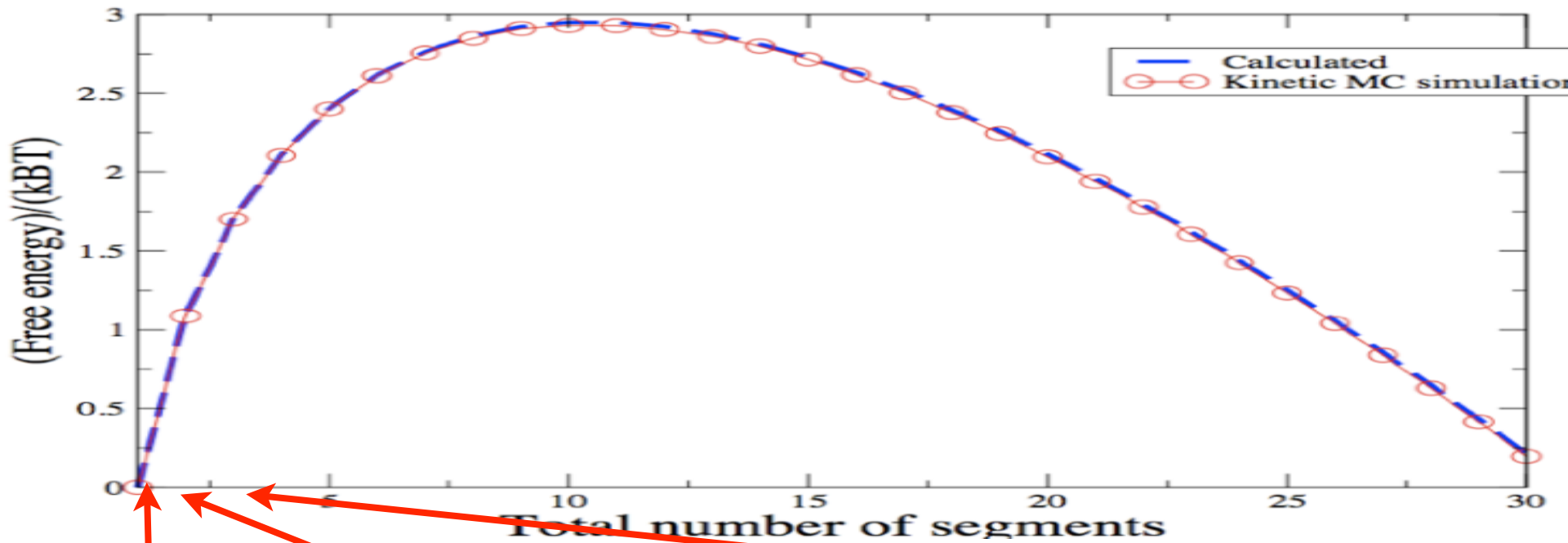
Hamer, Wattis and  
Graham, *Soft Matter*  
**8**, 11396 (2012)

# Projecting the GO model to a 1D system



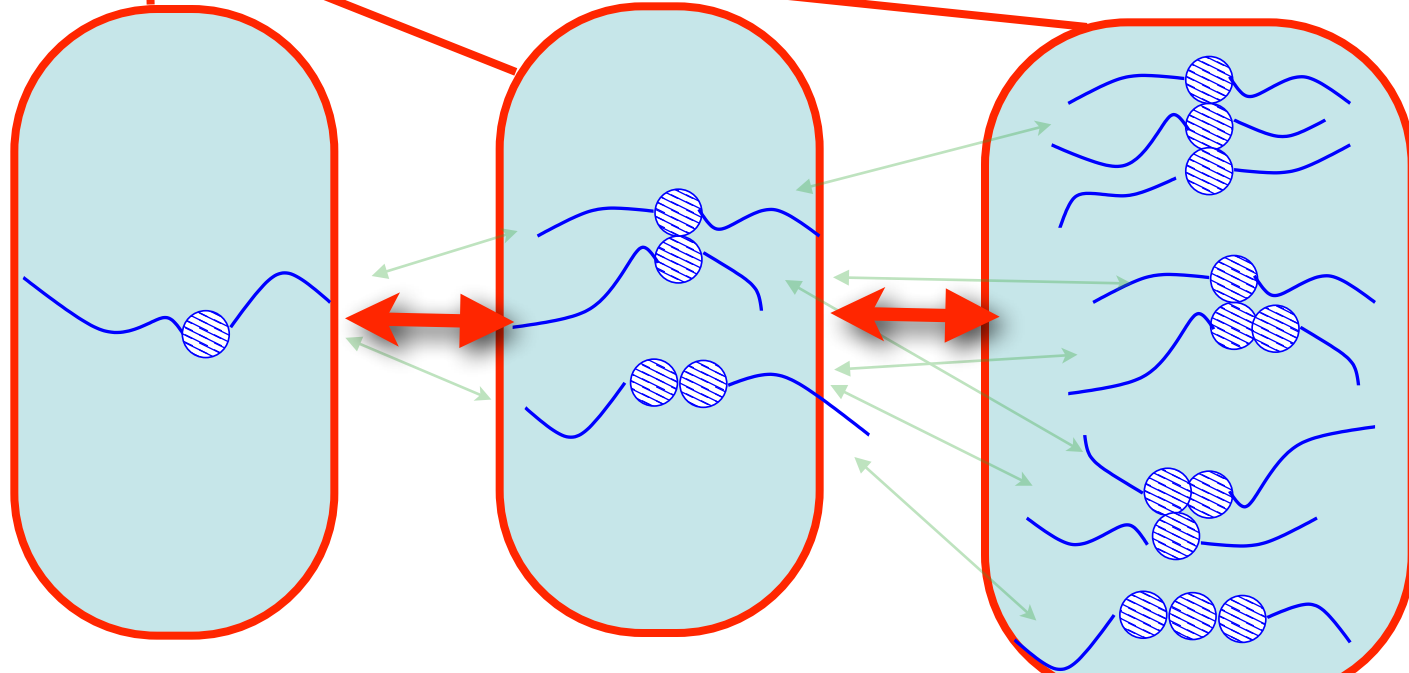
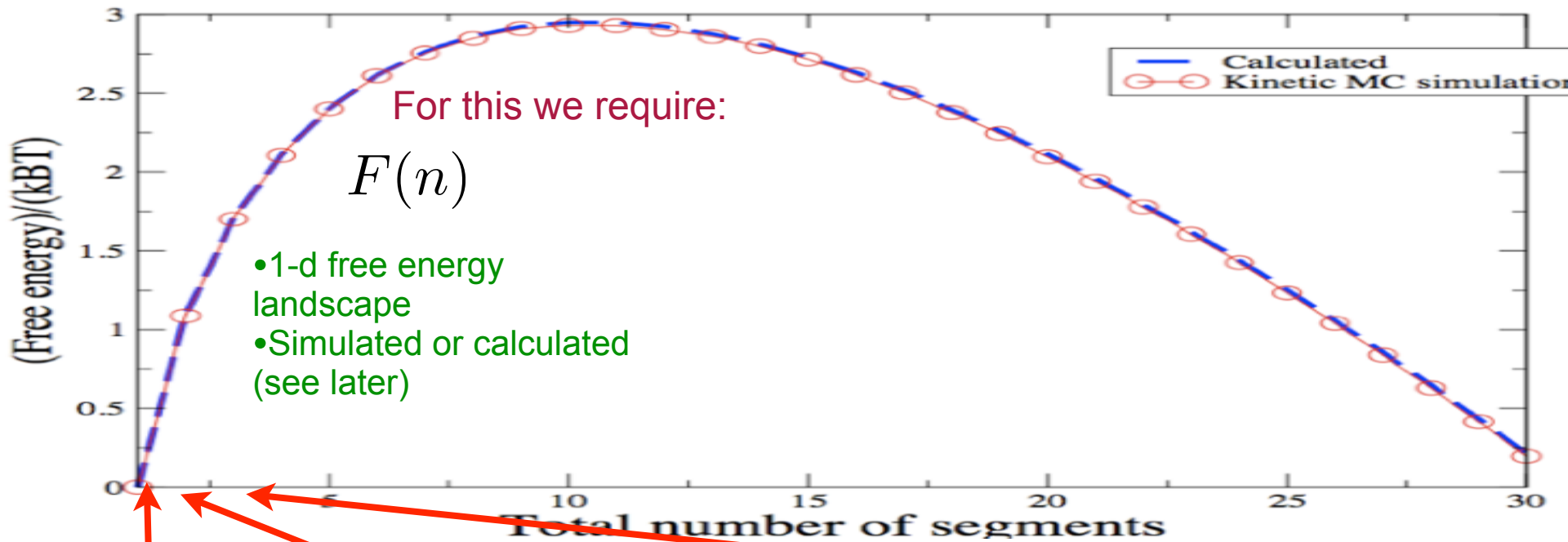
Hamer, Wattis and  
Graham, *Soft Matter*  
**8**, 11396 (2012)

# Projecting the GO model to a 1D system



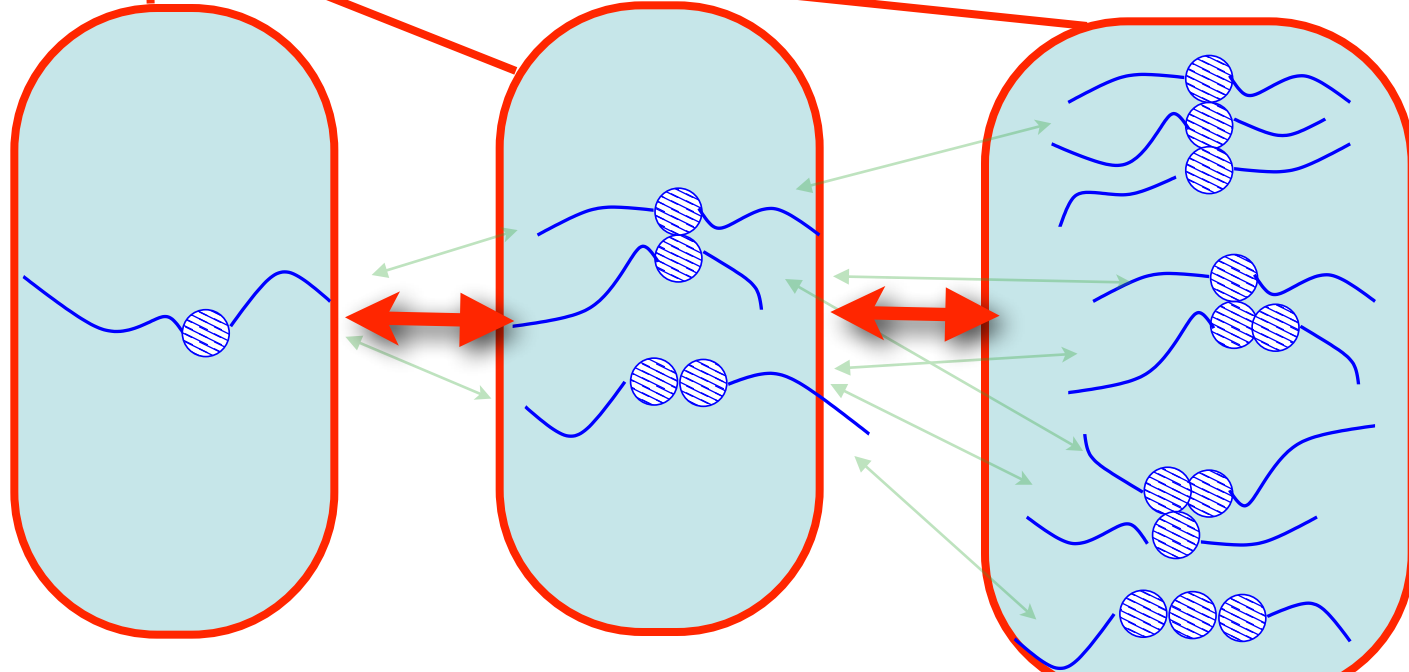
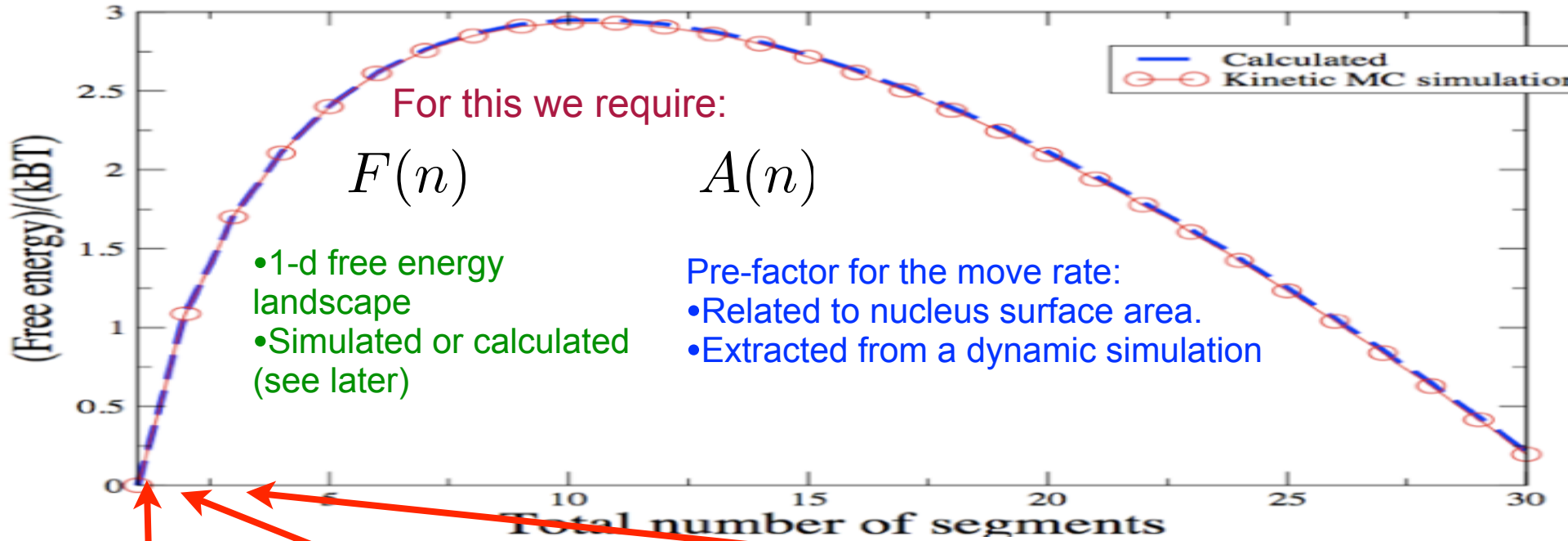
Hamer, Wattis and  
Graham, *Soft Matter*  
**8**, 11396 (2012)

# Projecting the GO model to a 1D system



Hamer, Wattis and Graham, *Soft Matter* **8**, 11396 (2012)

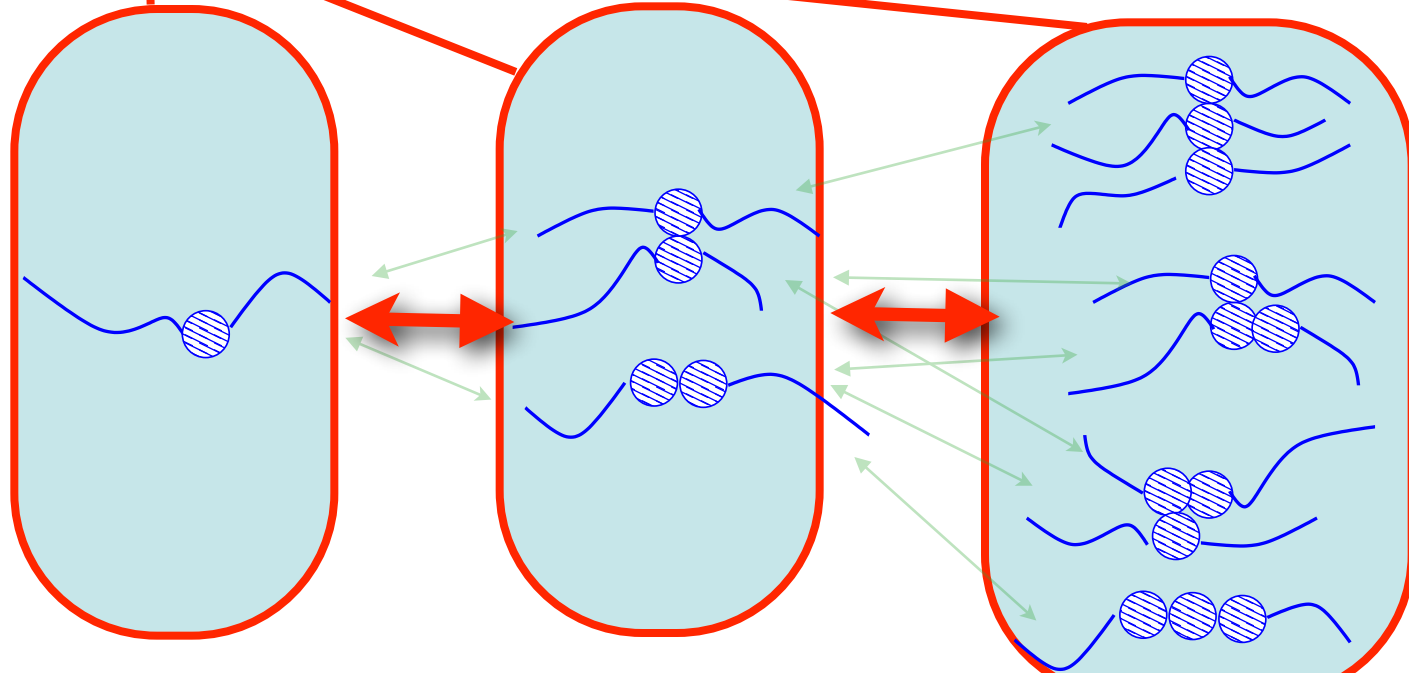
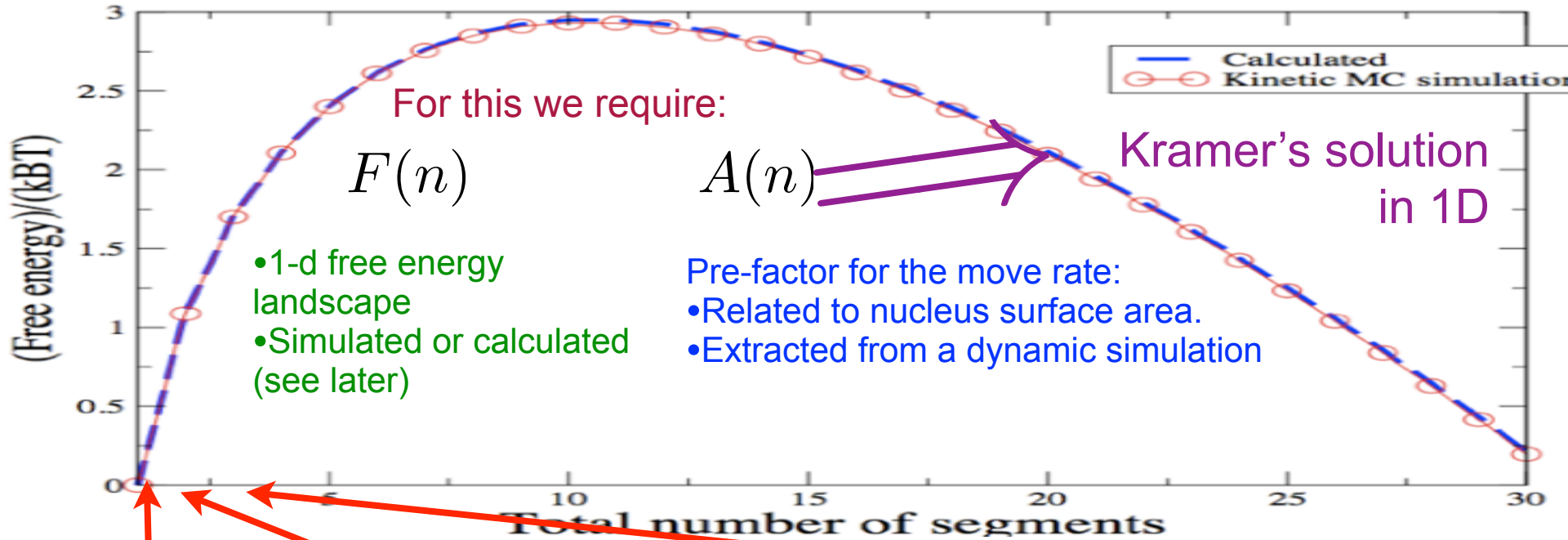
# Projecting the GO model to a 1D system



Hamer, Wattis and Graham, *Soft Matter* **8**, 11396 (2012)

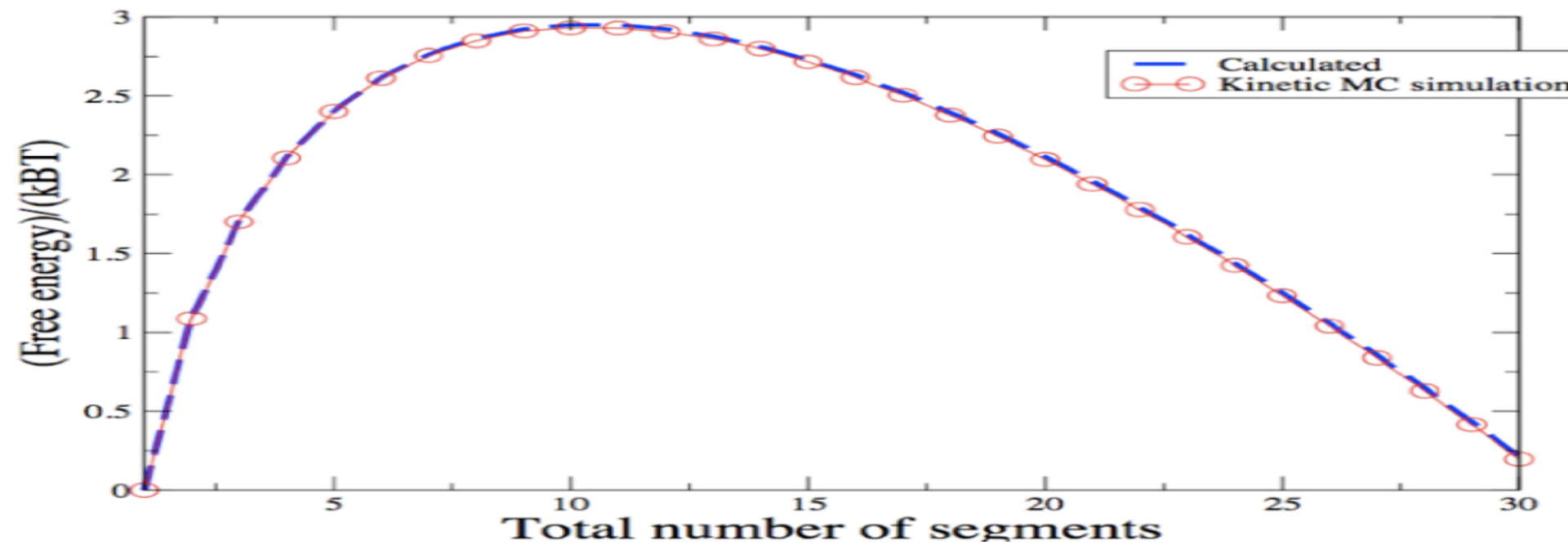


# Projecting the GO model to a 1D system

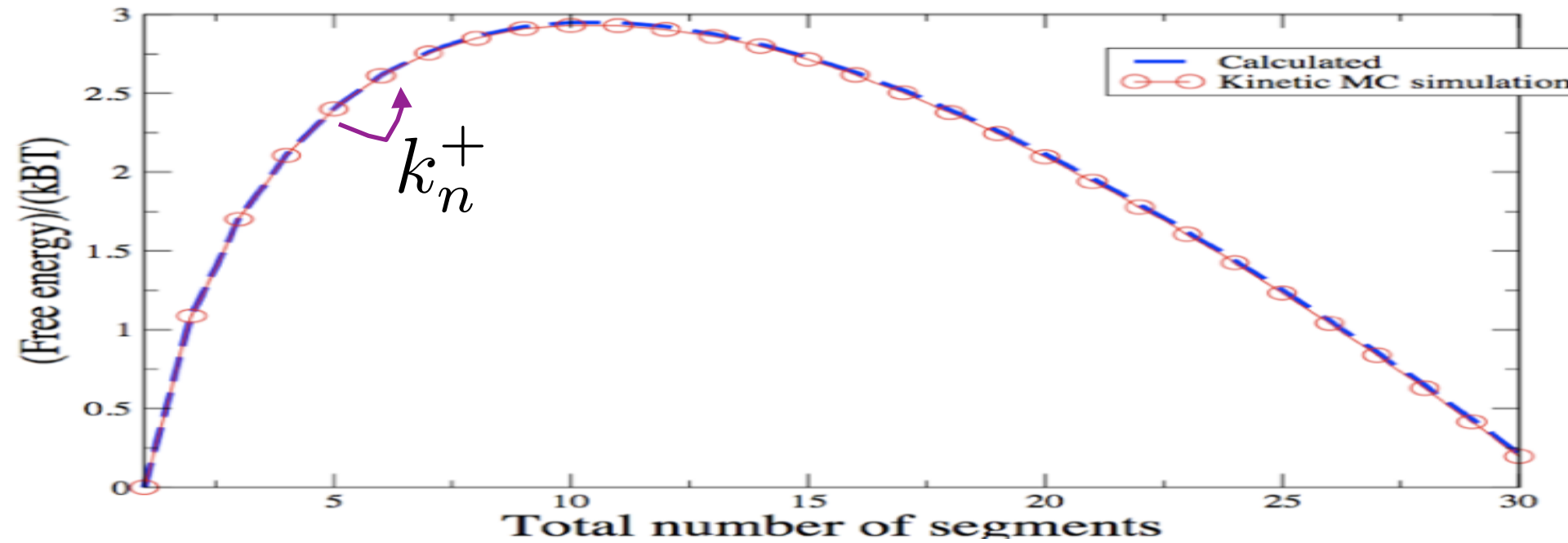


Hamer, Wattis and Graham, *Soft Matter* **8**, 11396 (2012)

# Projecting the GO model to a 1D system

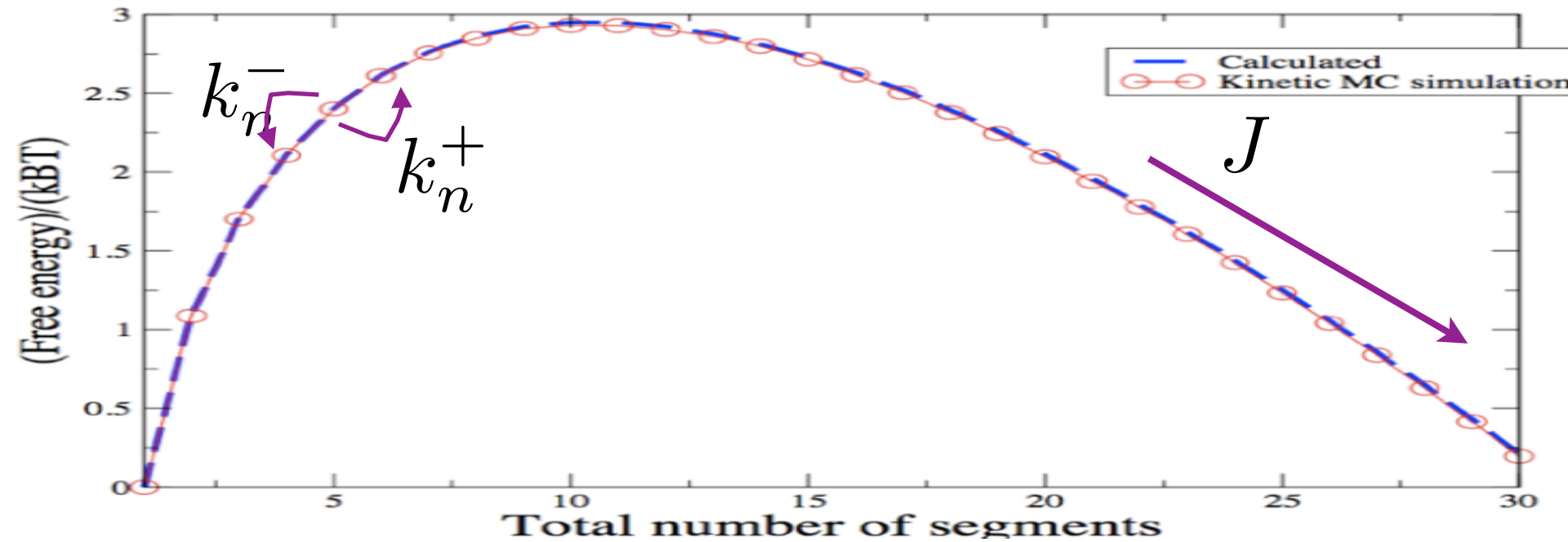


# Projecting the GO model to a 1D system



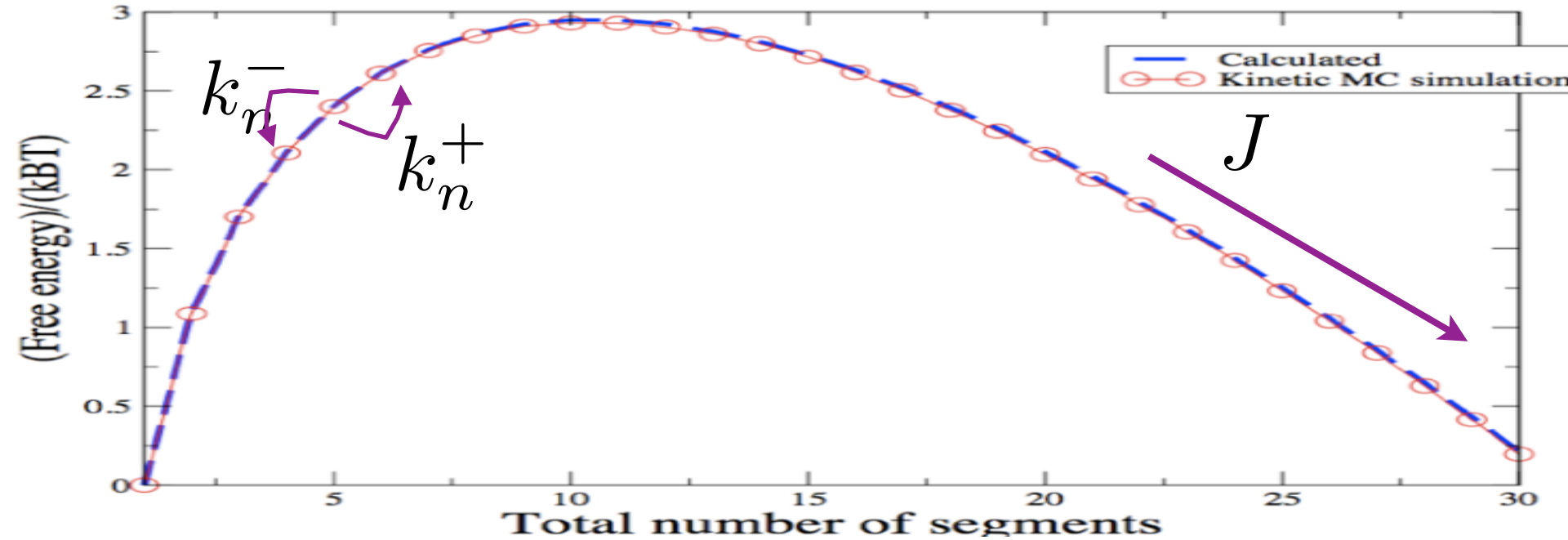
$$k_n^+ = A(n) \exp(-\Delta F(n))$$

# Projecting the GO model to a 1D system



$$k_n^+ = A(n) \exp(-\Delta F(n)) \quad k_n^- = A(n-1)$$

# Projecting the GO model to a 1D system

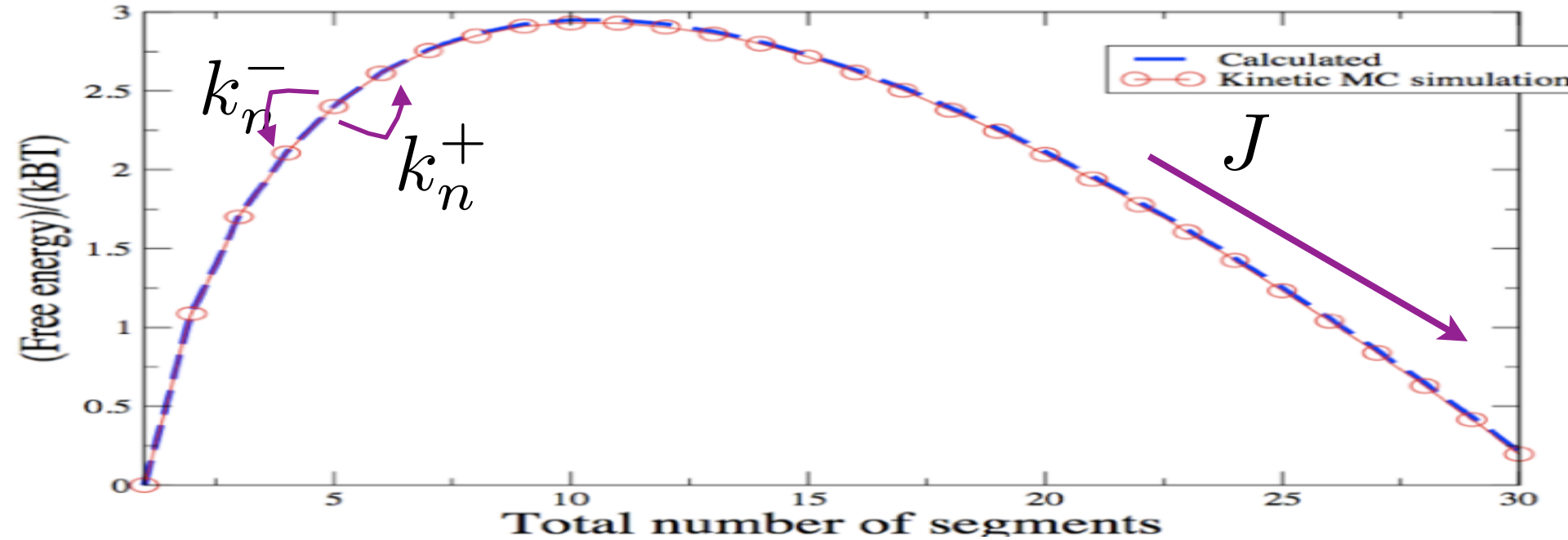


$$k_n^+ = A(n) \exp(-\Delta F(n)) \quad k_n^- = A(n-1)$$

In steady state

$$J = k_n^+ P_n - k_{n+1}^- P_{n+1}$$

# Projecting the GO model to a 1D system

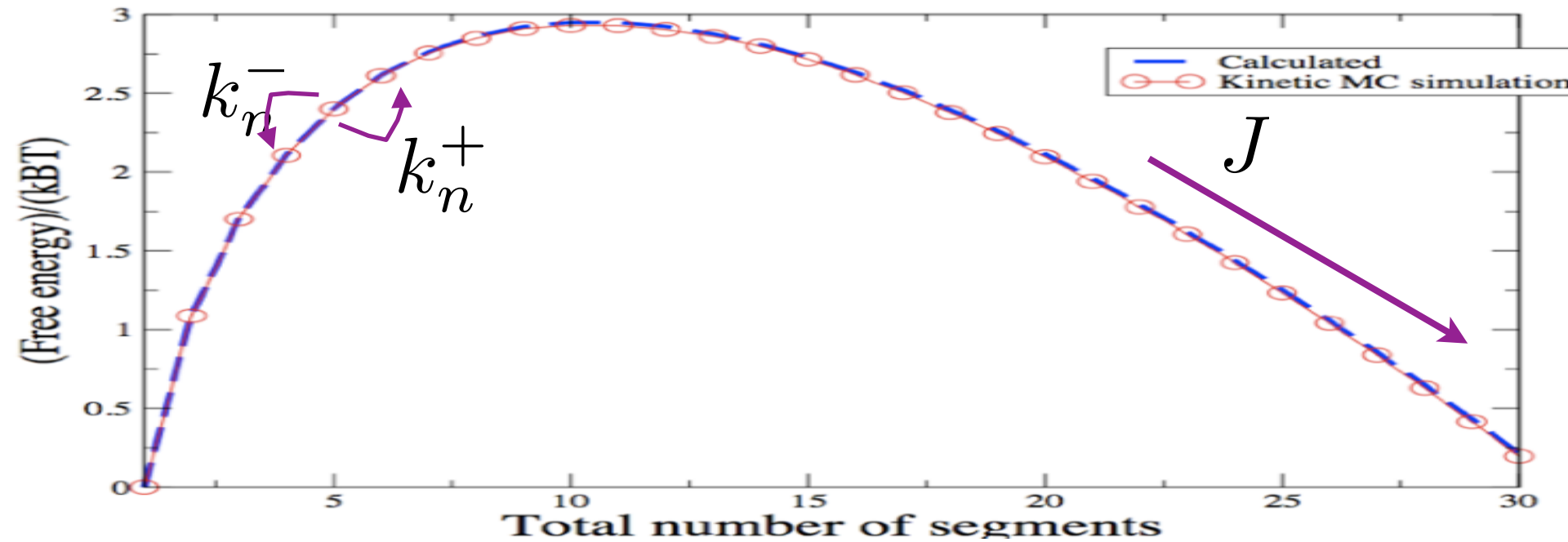


$$k_n^+ = A(n) \exp(-\Delta F(n)) \quad k_n^- = A(n-1)$$

In steady state  $J = k_n^+ P_n - k_{n+1}^- P_{n+1}$

$$A(n) = \frac{J}{P_n \exp(-\Delta F(n)) - P_{n+1}}$$

# Projecting the GO model to a 1D system



$$k_n^+ = A(n) \exp(-\Delta F(n)) \quad k_n^- = A(n-1)$$

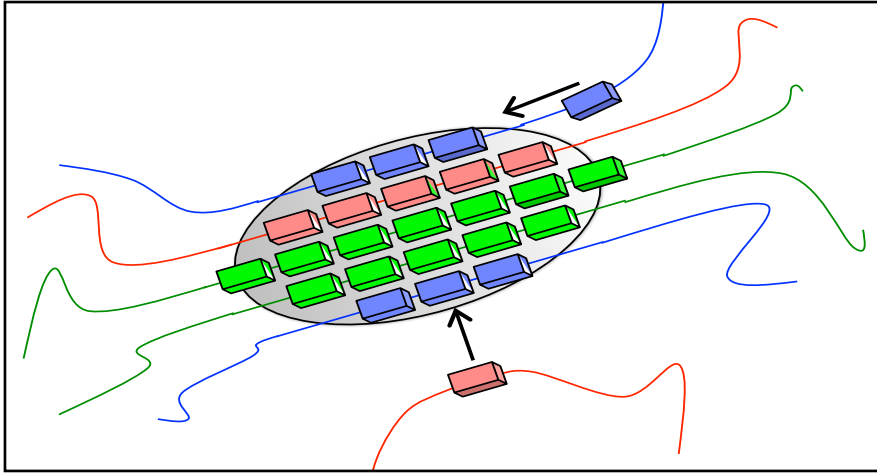
In steady state

$$J = k_n^+ P_n - k_{n+1}^- P_{n+1}$$

$$A(n) = \frac{J}{P_n \exp(-\Delta F(n)) - P_{n+1}}$$

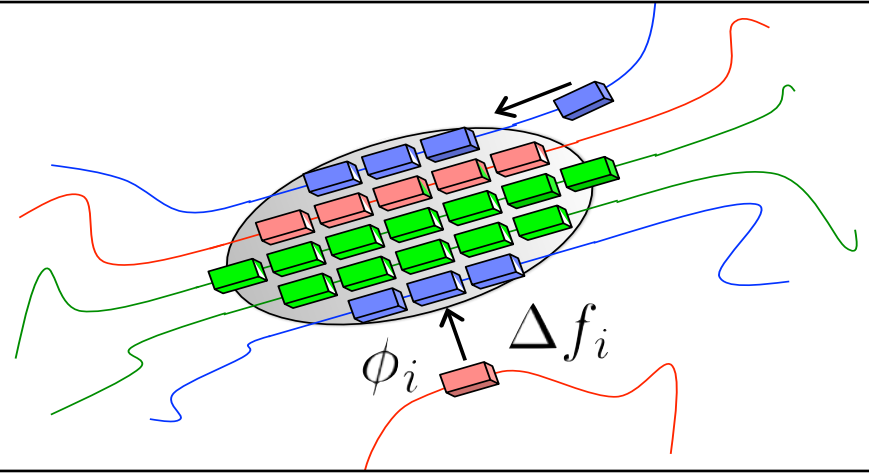
Can be extracted from our simulations

# Model for the barrier under flow (polySTRAND)



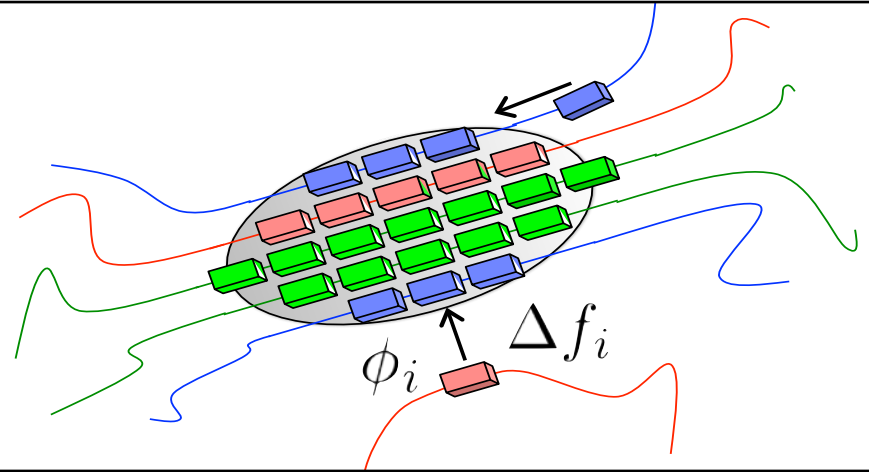


# Model for the barrier under flow (polySTRAND)



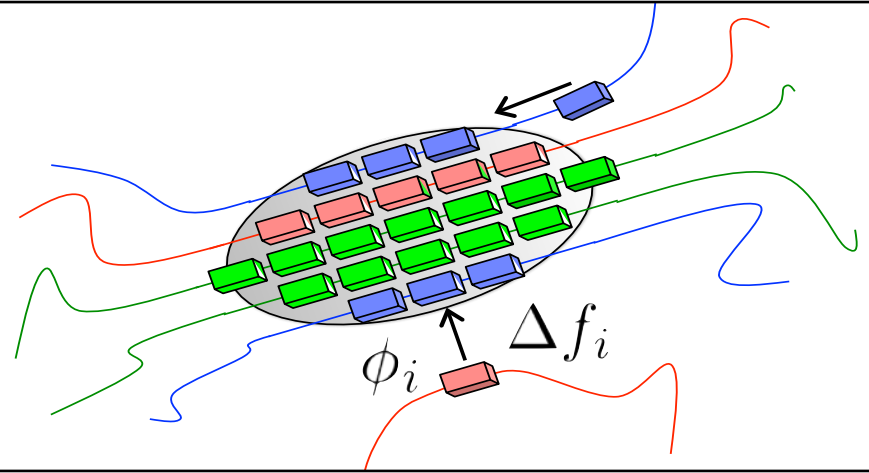
$Df_i$ =nematic order  
 $\phi_i$ =melt fraction of  $i$

# Model for the barrier under flow (polySTRAND)



$Df_i$ =nematic order  
 $\phi_i$ =melt fraction of  $i$   
 $N_T$ =total monomers  
 $N_s$ =total stems  
 $S$ =surface area

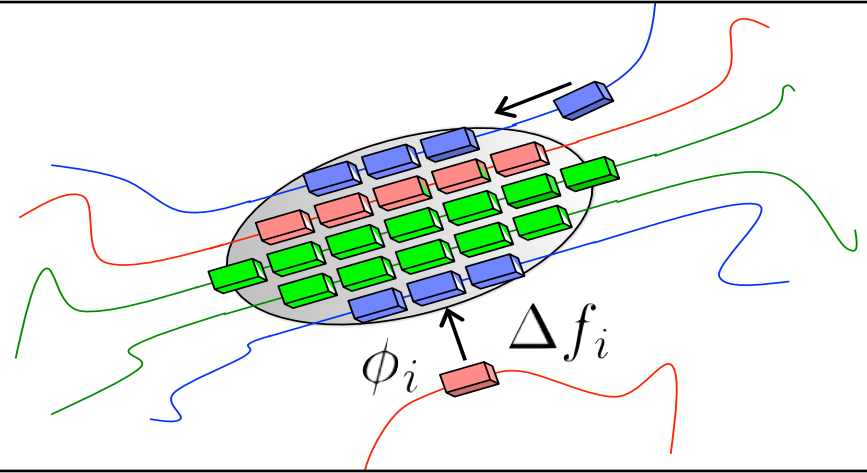
# Model for the barrier under flow (polySTRAND)



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$e_B$ =Bulk free energy gain  
 $\mu_s$ =surface area cost

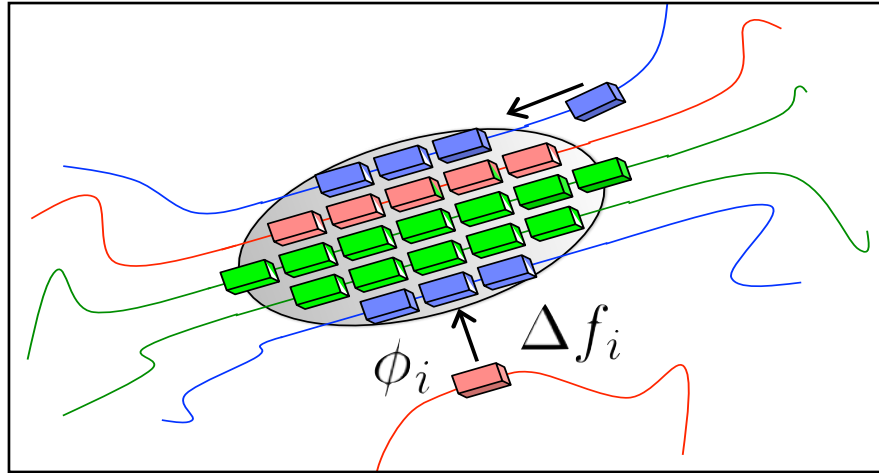
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 $v_i$ =fraction of  $i$  monomers  
 $w_i$ =fraction of  $i$  stems  
 $q=N_T/N_S$

# Model for the barrier under flow (polySTRAND)



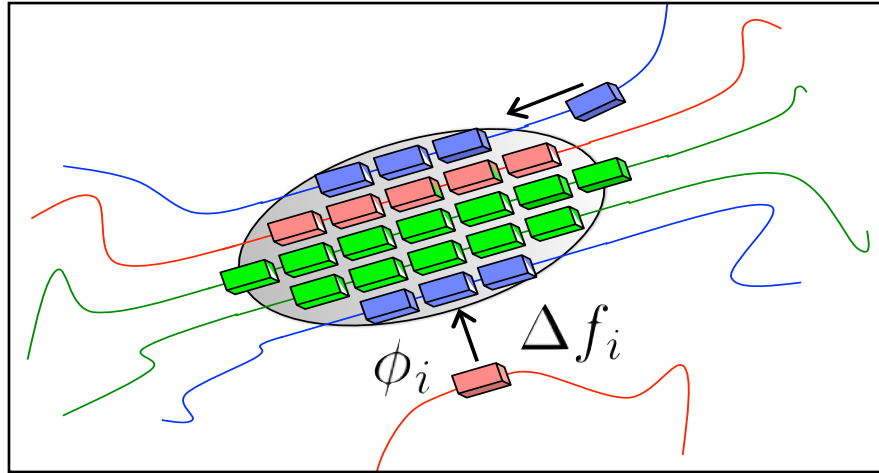
$Df_i$ =nematic order  
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 $N_T$ =total monomers  
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$\epsilon_B$ =Bulk free energy gain  
 $\mu_s$ =surface area cost  
 $v_i$ =fraction of  $i$  monomers  
 $w_i$ =fraction of  $i$  stems  
 $q=N_T/N_S$

Nucleus free energy

$$\begin{aligned}
 \frac{F(N_T, N_S, \{w_i\}, \{v_i\})}{k_B T} = & N_T \sum_i [q w_i (2 \log w_i - \log \phi_i) - v_i \log v_i + (v_i - q w_i) \log(v_i - q w_i) - v_i \Delta f_i] \\
 & + N_S \log q - \epsilon_B N_T + \mu_S S(N_T, N_S).
 \end{aligned}$$

# Model for the barrier under flow (polySTRAND)



$Df_i$ =nematic order  
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Nucleus free energy

Stem entropy change:  
melt  $\rightarrow$  nucleus

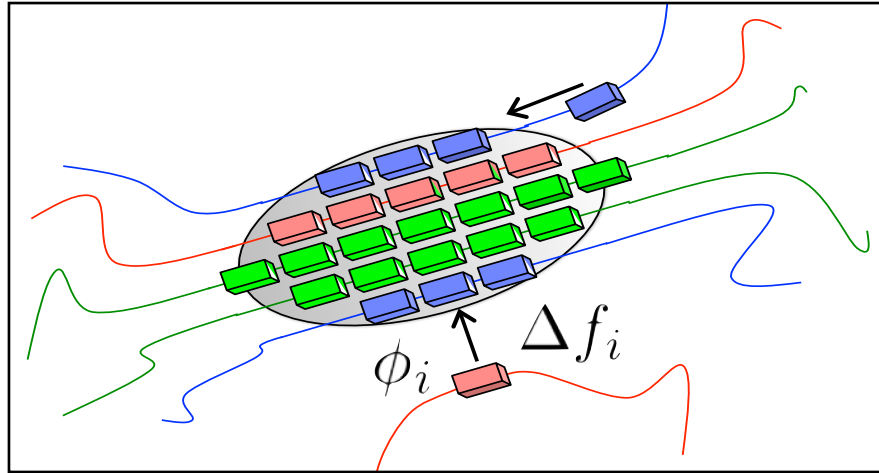
Stem/monomer  
arrangement entropy

$$\frac{F(N_T, N_S, \{w_i\}, \{v_i\})}{k_B T} = N_T \sum_i \left[ \underbrace{q w_i (2 \log w_i - \log \phi_i)}_{\text{Stem entropy change: melt} \rightarrow \text{nucleus}} - \underbrace{v_i \log v_i + (v_i - q w_i) \log (v_i - q w_i)}_{\text{Stem/monomer arrangement entropy}} - \underbrace{v_i \Delta f_i}_{\text{Flow-induced ordering}} \right] + N_S \log q - \underbrace{\epsilon_B N_T}_{\text{Bulk gain}} + \underbrace{\mu_S S(N_T, N_S)}_{\text{Surface cost}}.$$

Sum over species

Flow-induced ordering

# Model for the barrier under flow (polySTRAND)



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 $v_i$ =fraction of  $i$  monomers  
 $w_i$ =fraction of  $i$  stems  
 $q=N_T/N_S$

Minimise with constraint  $\sum_i w_i = \sum_i v_i = 1$

Nucleus free energy

Stem entropy change:  
melt  $\rightarrow$  nucleus

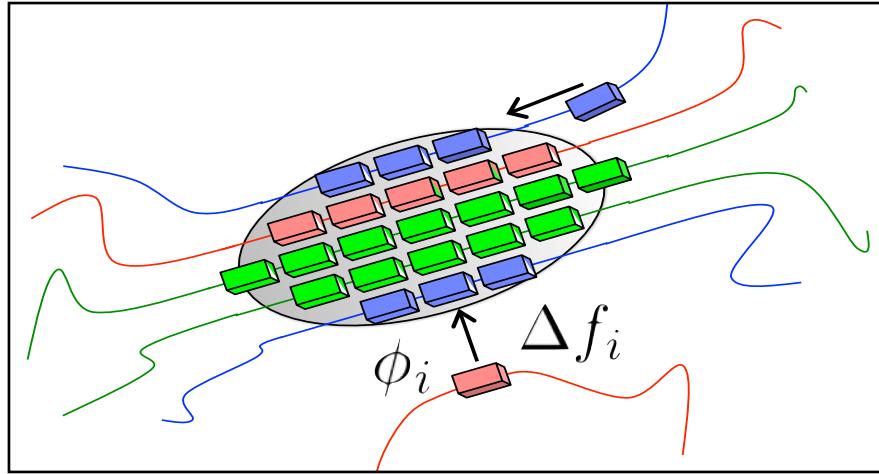
Stem/monomer  
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$$\frac{F(N_T, N_S, \{w_i\}, \{v_i\})}{k_B T} = N_T \sum_i \underbrace{[q w_i (2 \log w_i - \log \phi_i)]}_{\text{Stem entropy change}} - \underbrace{v_i \log v_i + (v_i - q w_i) \log(v_i - q w_i)}_{\text{Stem/monomer arrangement entropy}} - \underbrace{v_i \Delta f_i}_{\text{Flow-induced ordering}} + N_S \log q - \underbrace{\epsilon_B N_T}_{\text{Bulk gain}} + \underbrace{\mu_S S(N_T, N_S)}_{\text{Surface cost}}.$$

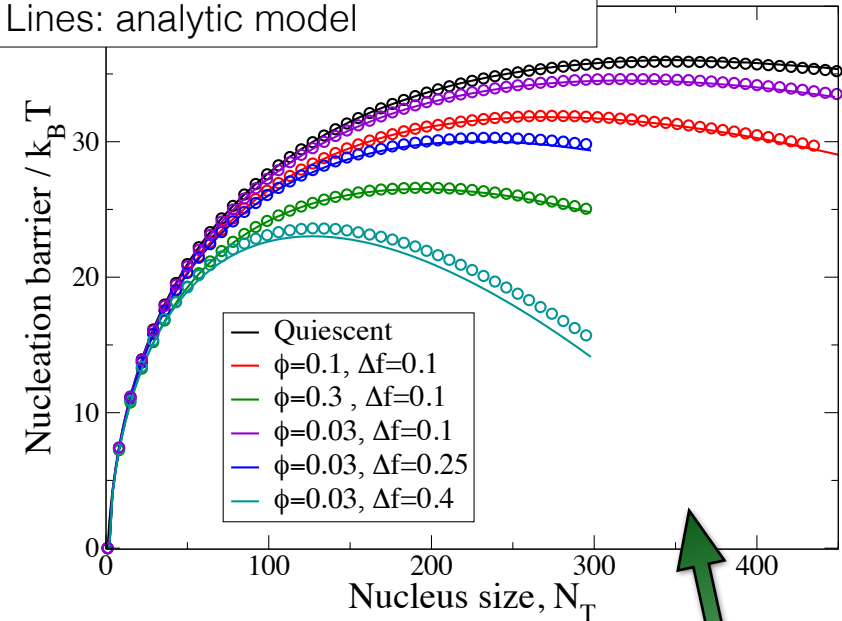
Sum over species

Flow-induced ordering

# Model for the barrier under flow (polySTRAND)



Symbols: GO model simulations  
Lines: analytic model



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Nucleus free energy

Stem entropy change:  
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Stem/monomer  
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$$\frac{F(N_T, N_S, \{w_i\}, \{v_i\})}{k_B T} = N_T \sum_i [q w_i (2 \log w_i - \log \phi_i) - v_i \log v_i + (v_i - q w_i) \log (v_i - q w_i) - v_i \Delta f_i] + N_S \log q - \epsilon_B N_T + \mu_S S(N_T, N_S).$$

Sum over  
species

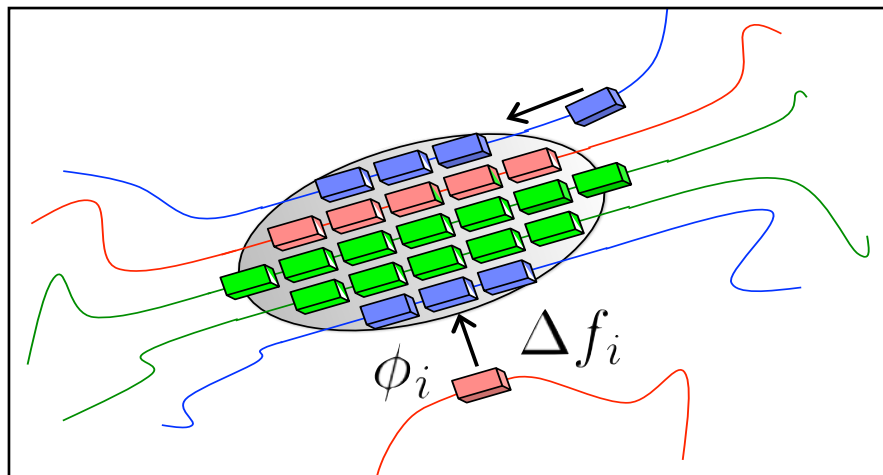
Bulk gain

Surface cost

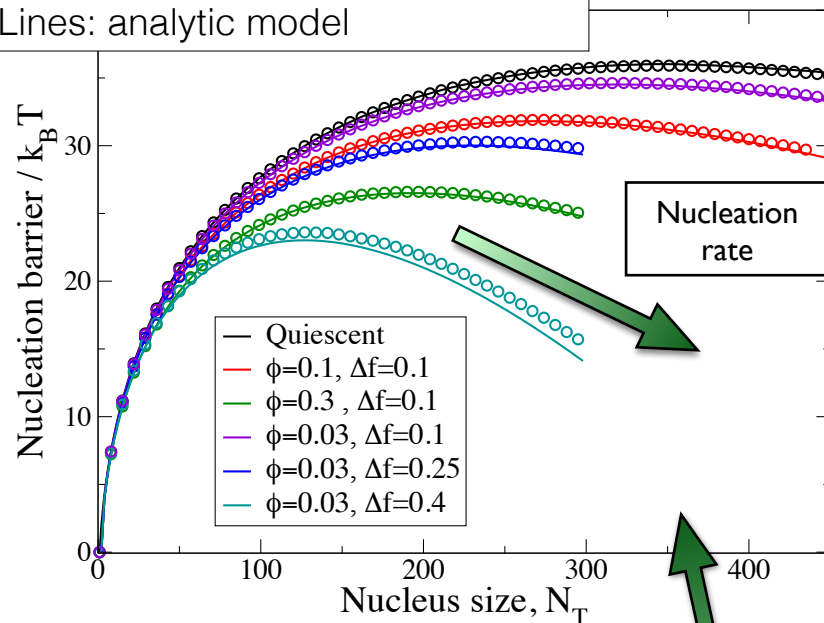
Flow-  
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# Model for the barrier under flow (polySTRAND)



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Stem/monomer  
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Sum over  
species

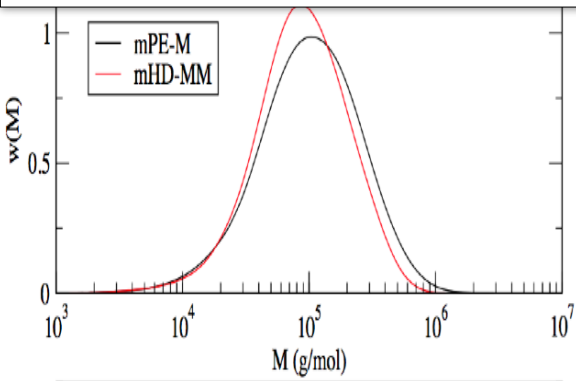
Bulk gain

Surface cost

Flow-  
induced  
ordering

# Model summary

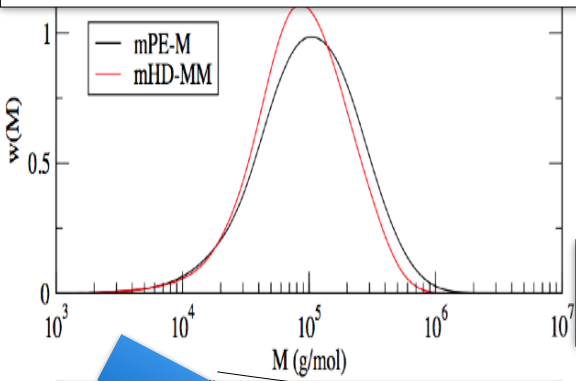
Molecular weight distribution



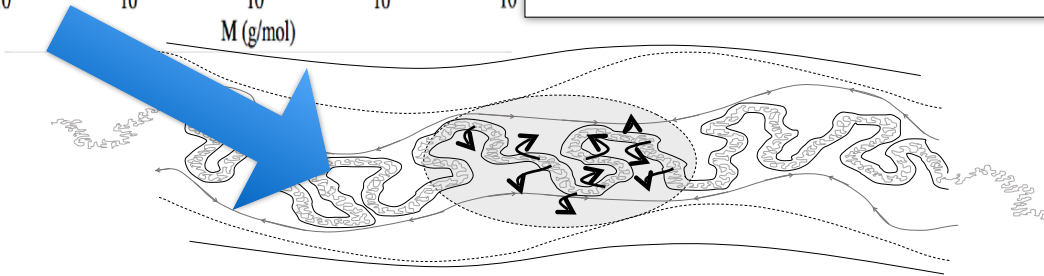
Read, McIlroy, Das,  
Harlen and Graham,  
Phys. Rev. Lett. **124**  
147802 (2020)

# Model summary

Molecular weight distribution



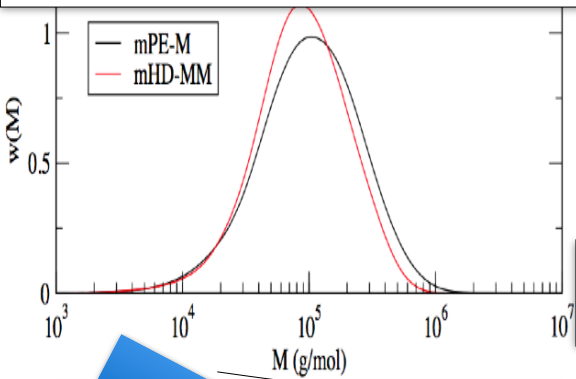
Flow model



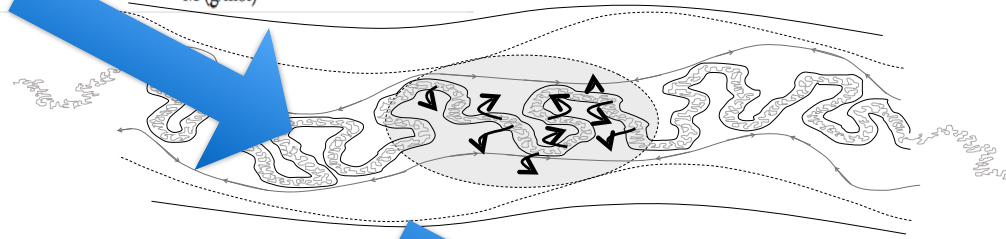
Read, McIlroy, Das,  
Harlen and Graham,  
Phys. Rev. Lett. **124**  
147802 (2020)

# Model summary

Molecular weight distribution



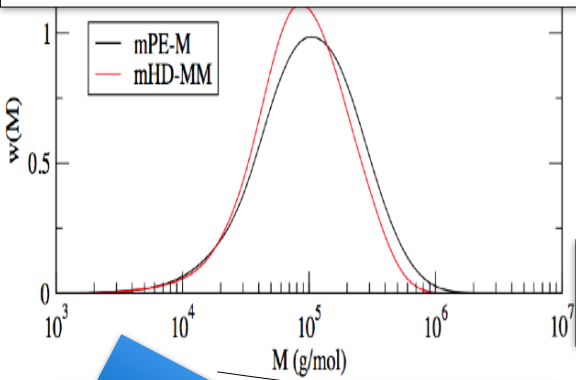
Flow model



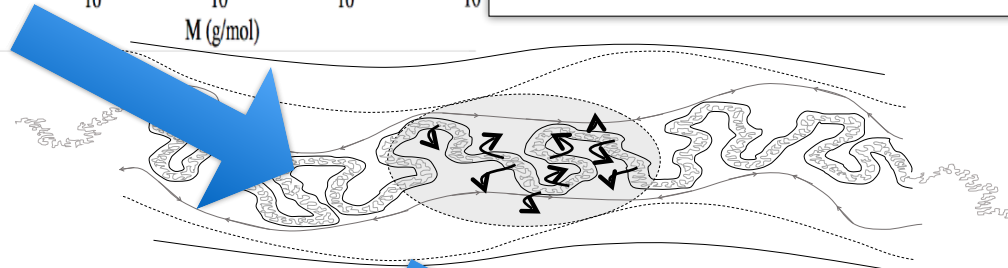
Chain configurations

# Model summary

Molecular weight distribution

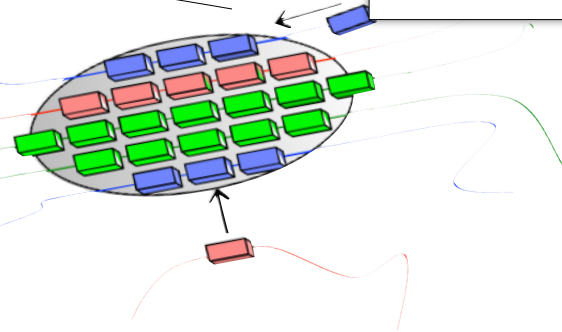


Flow model



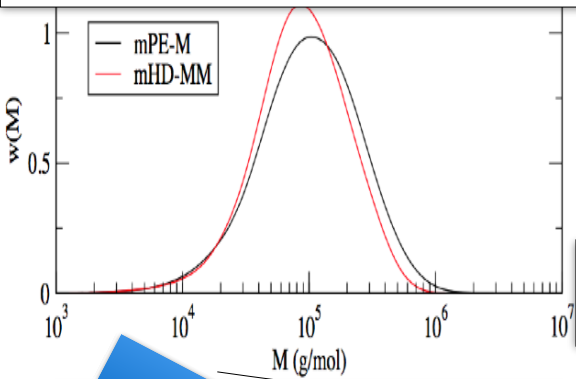
STRAND model

Chain configurations

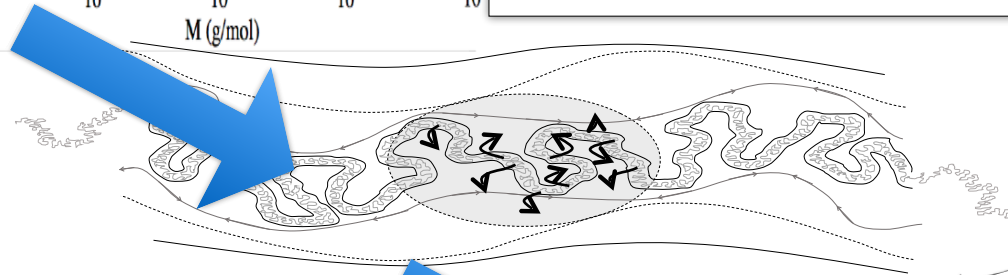


# Model summary

Molecular weight distribution

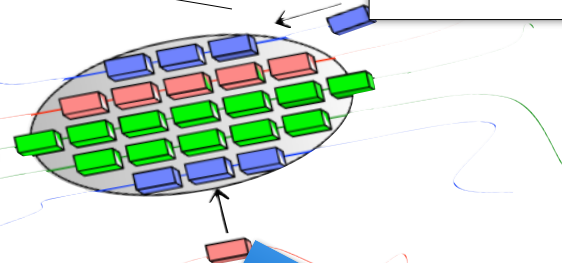


Flow model



STRAND model

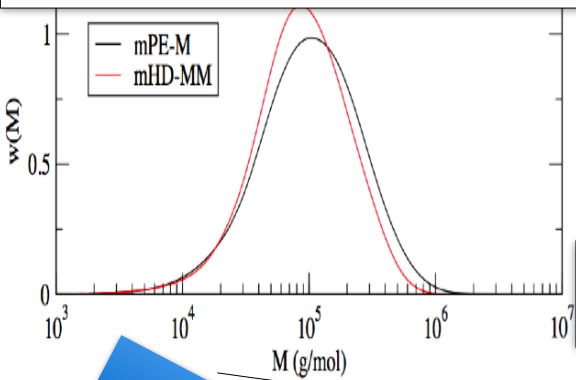
Chain configurations



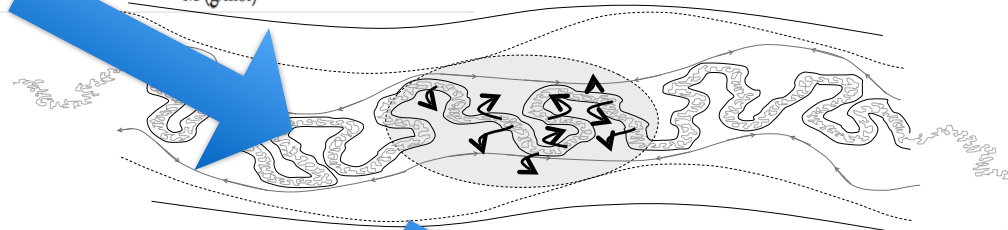
Nucleation barrier

# Model summary

Molecular weight distribution

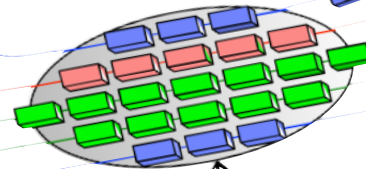


Flow model



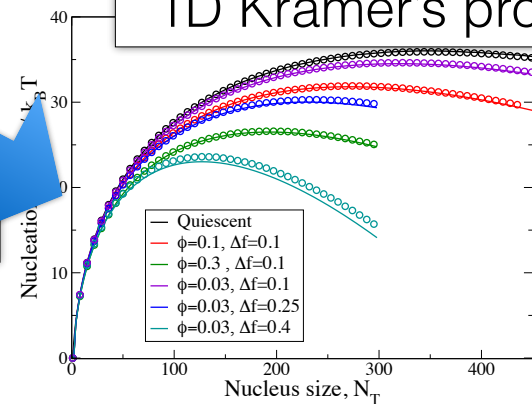
STRAND model

Chain configurations



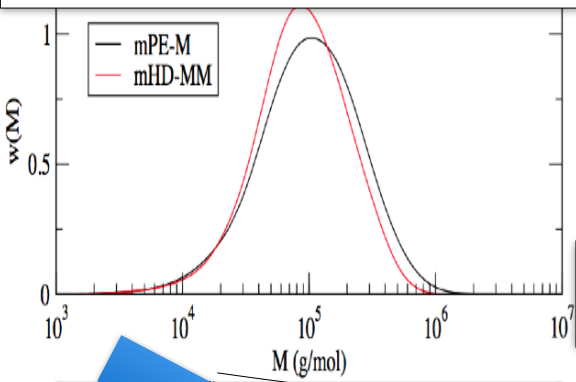
Nucleation barrier

1D Kramer's problem

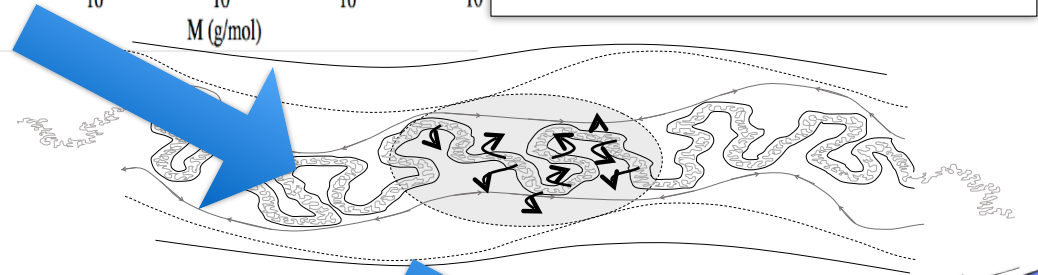


# Model summary

Molecular weight distribution

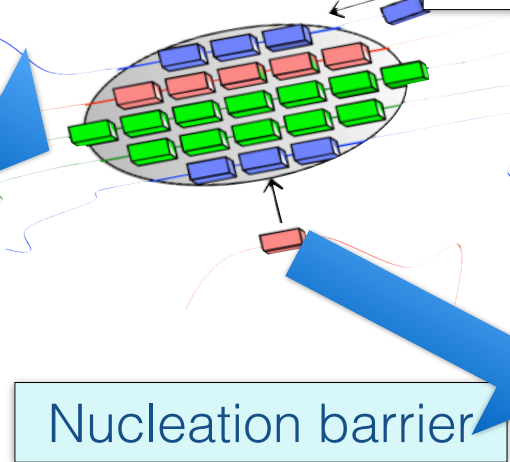


Flow model

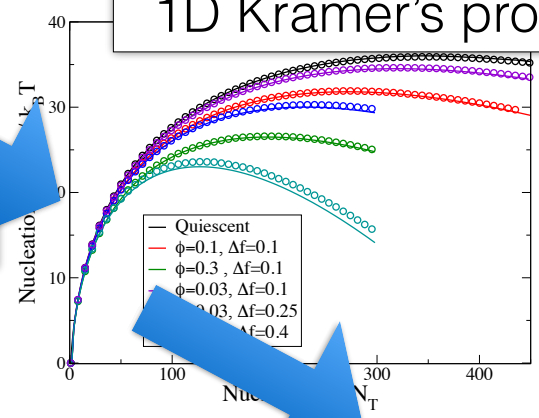


STRAND model

Chain configurations



1D Kramer's problem



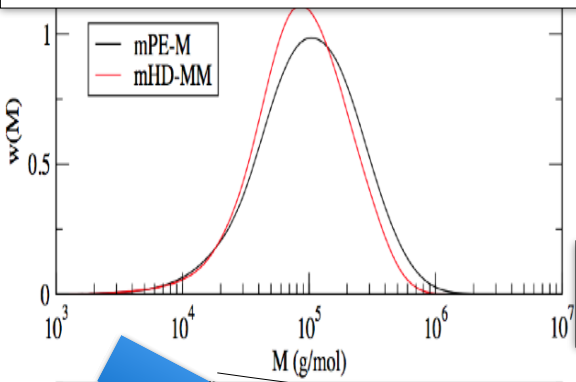
Nucleation Rate

Read, McIlroy, Das, Harlen and Graham, Phys. Rev. Lett. **124** 147802 (2020)

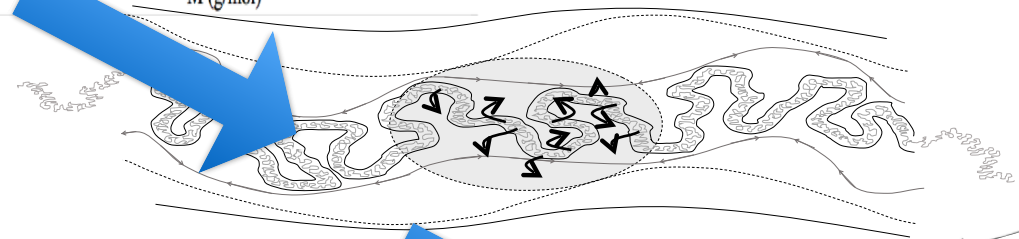


# Model summary

Molecular weight distribution

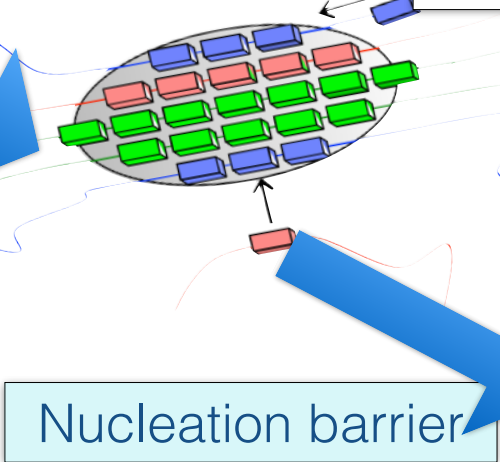


Flow model

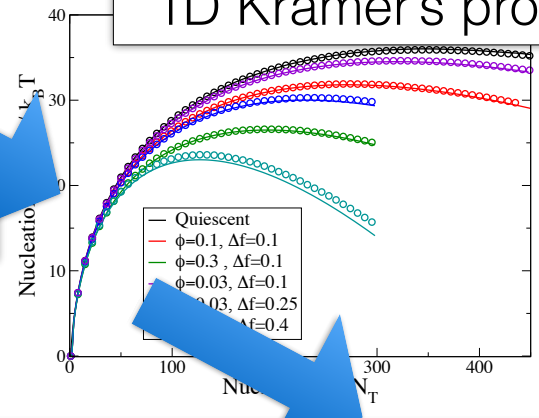


STRAND model

Chain configurations



1D Kramer's problem



Nucleation barrier

Also predicts long chain enhancement

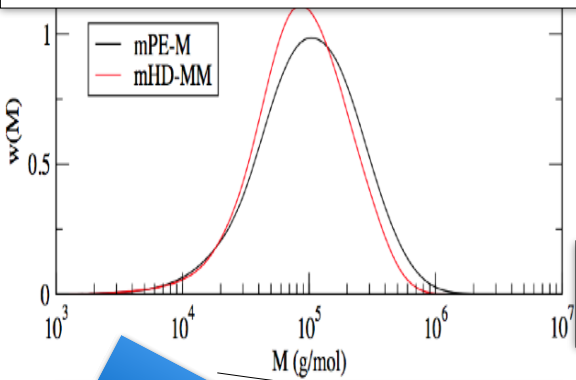
Nucleation Rate

Read, McIlroy, Das, Harlen and Graham, Phys. Rev. Lett. **124** 147802 (2020)

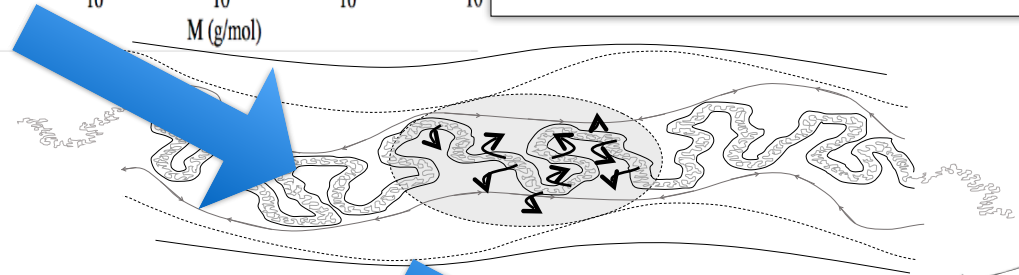
# Model summary

All steps are numerically simple/cheap - no stochastic simulation required

Molecular weight distribution

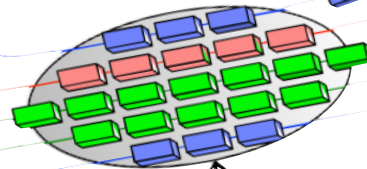


Flow model



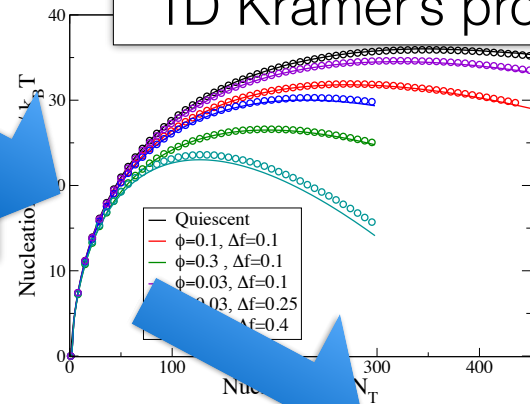
STRAND model

Chain configurations



Nucleation barrier

1D Kramer's problem



Also predicts long chain enhancement

Nucleation Rate

# Comparison with experiments

# Direct observation of nucleation during steady shear



Continuous Shear  
Flow

Measurements on an isotactic polypropylene resin by  
Coccorullo *et al.* *Macromolecules* (2008) and Pantani *et al.*  
*Macromolecules* (2010)

# Direct observation of nucleation during steady shear

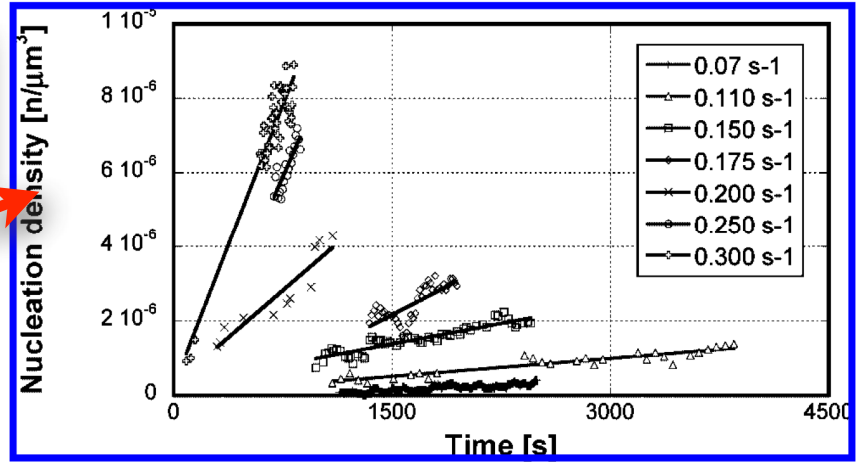


Continuous Shear  
Flow

Measurements on an isotactic polypropylene resin by  
Coccorullo *et al.* *Macromolecules* (2008) and Pantani *et al.*  
*Macromolecules* (2010)

# Direct observation of nucleation during steady shear

Count nuclei as they appear



Continuous Shear Flow

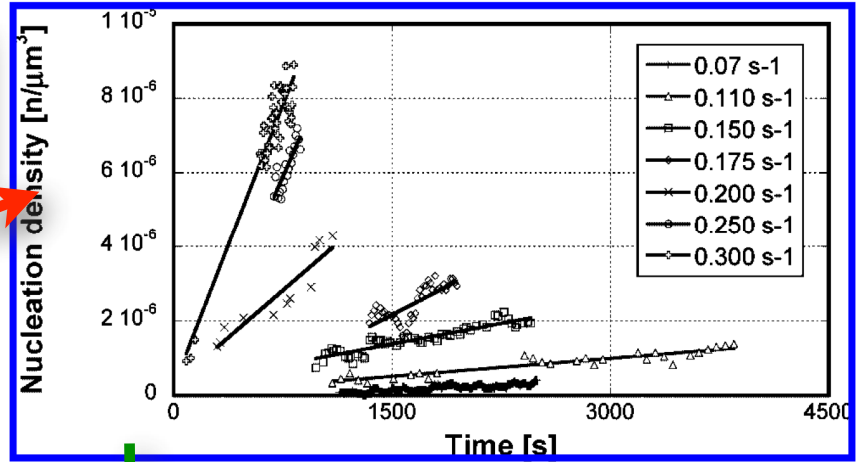
Measurements on an isotactic polypropylene resin by Coccorullo *et al.* *Macromolecules* (2008) and Pantani *et al.* *Macromolecules* (2010)

# Direct observation of nucleation during steady shear

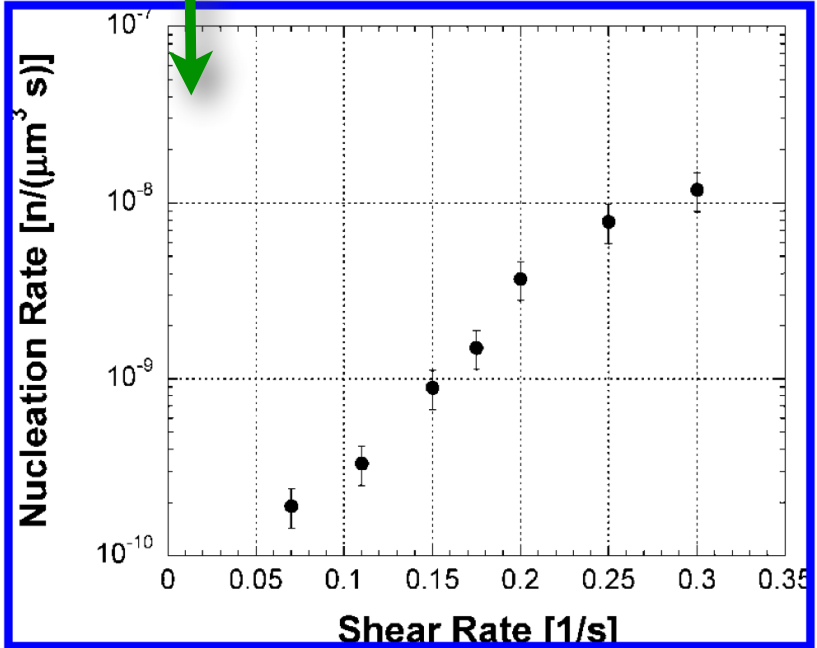
Count nuclei as they appear



Continuous Shear Flow

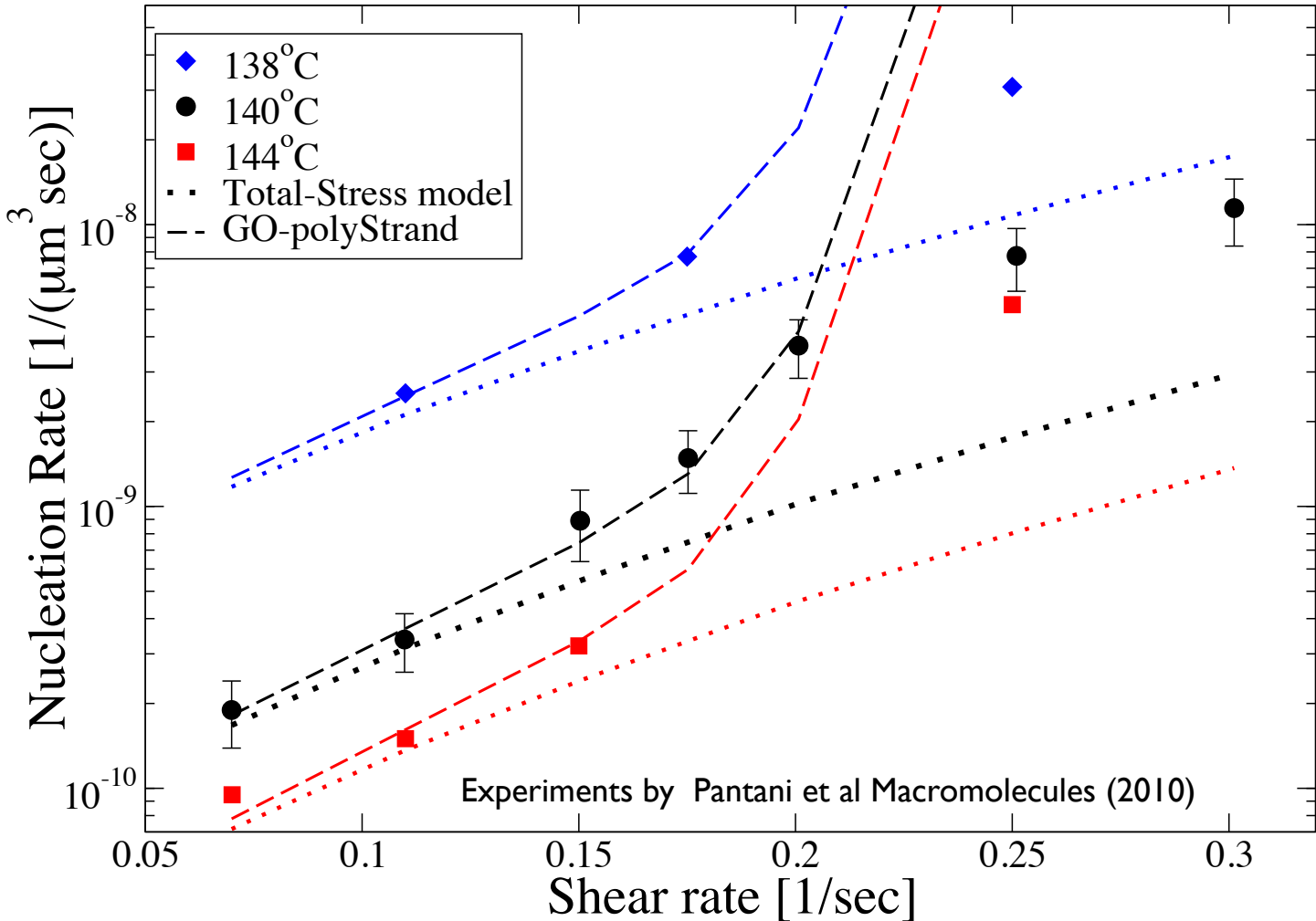


Steady nucleation rate against shear rate



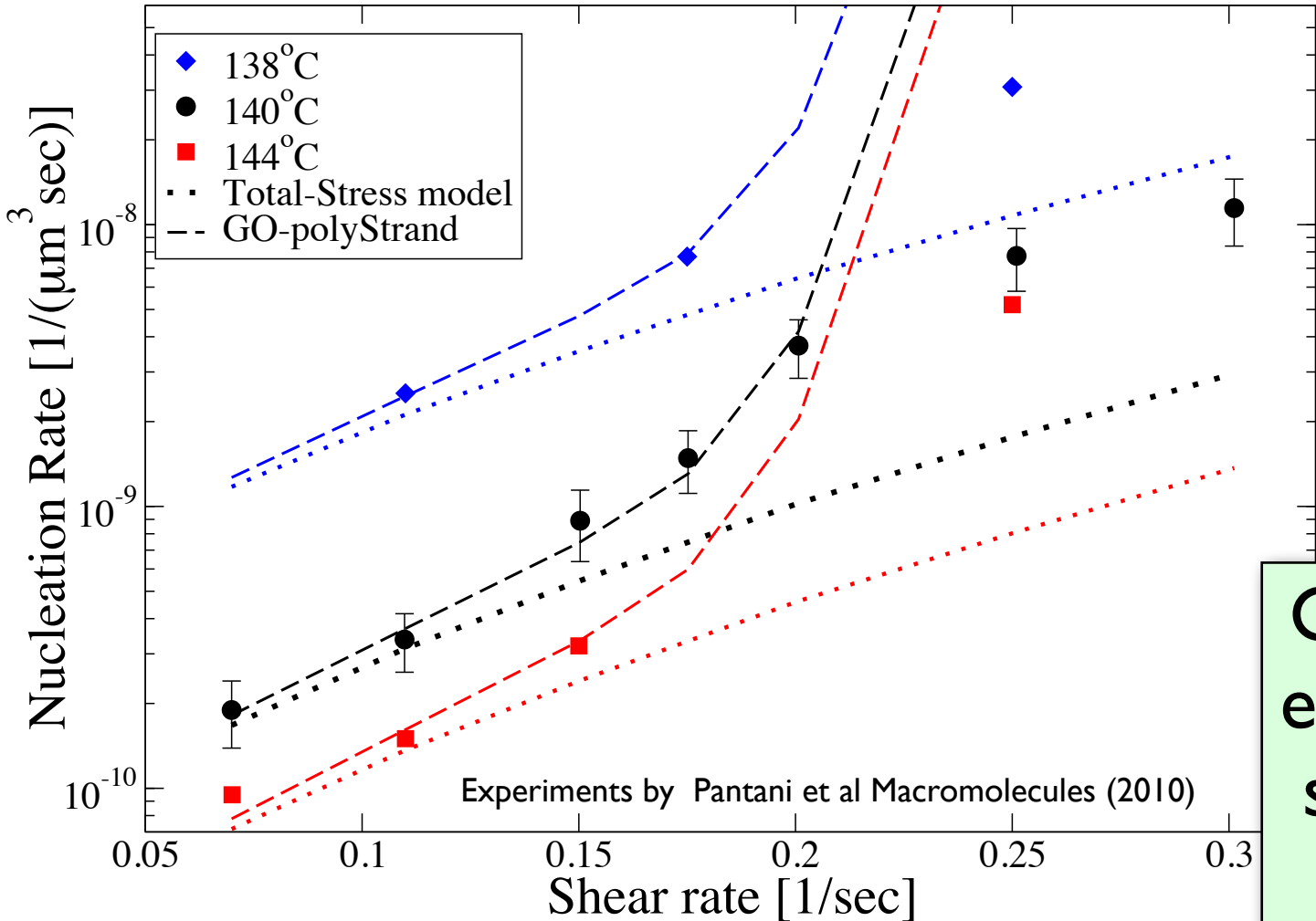
Measurements on an isotactic polypropylene resin by Coccorullo *et al.* Macromolecules (2008) and Pantani *et al.* Macromolecules (2010)

# Direct observation of nucleation during steady shear



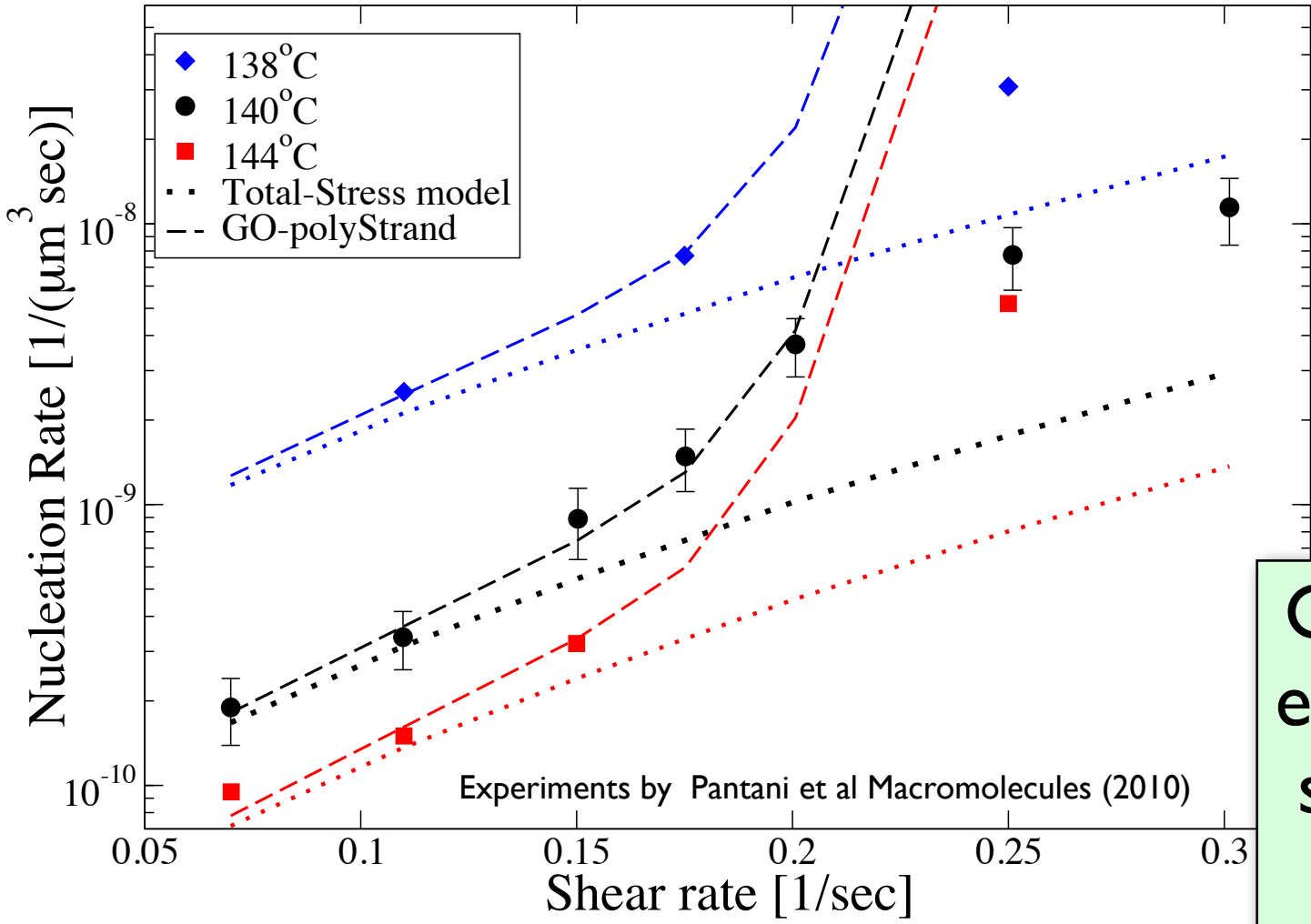


# Direct observation of nucleation during steady shear



Curvature is experimental signature of long chain enhancement

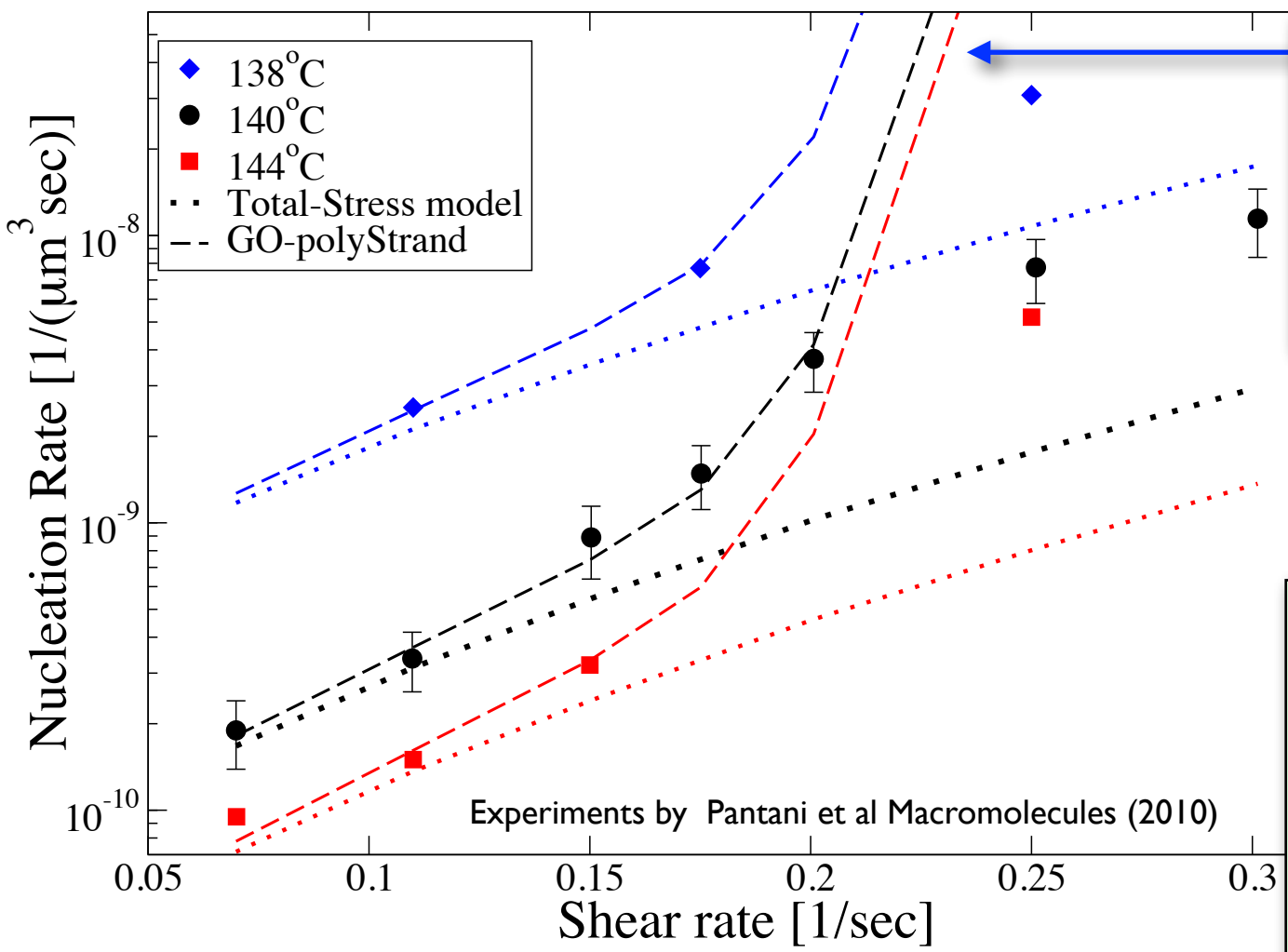
# Direct observation of nucleation during steady shear



Curvature is experimental signature of long chain enhancement

Effect of temperature is successfully predicted

# Direct observation of nucleation during steady shear



Model overpredicts the enhancement at higher rates

Curvature is experimental signature of long chain enhancement

Effect of temperature is successfully predicted

Read, McIlroy, Das, Harlen and Graham, Phys. Rev. Lett. **124** 147802 (2020)

# Smooth-polyStrand model

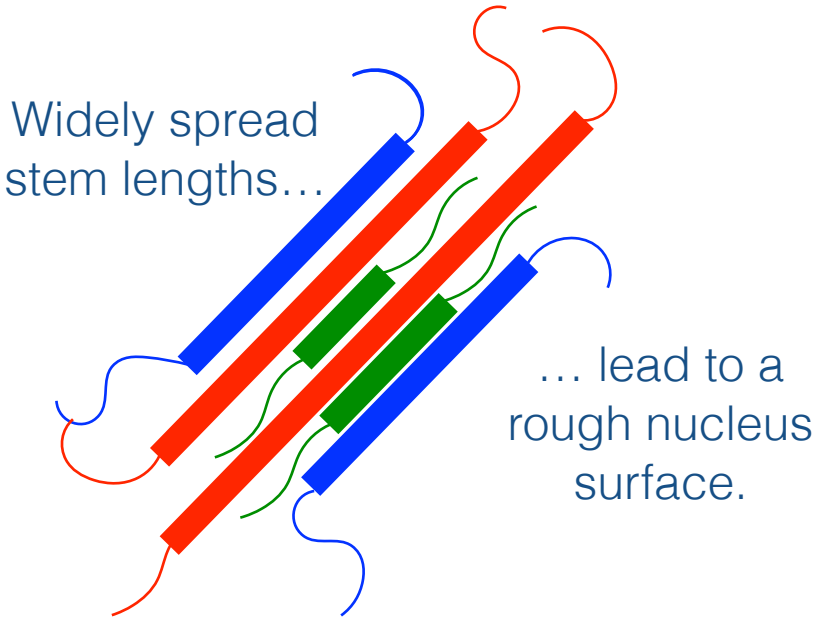
Nucleus roughness penalty

Long chain depletion

# Smooth-polyStrand model

**Nucleus roughness penalty**

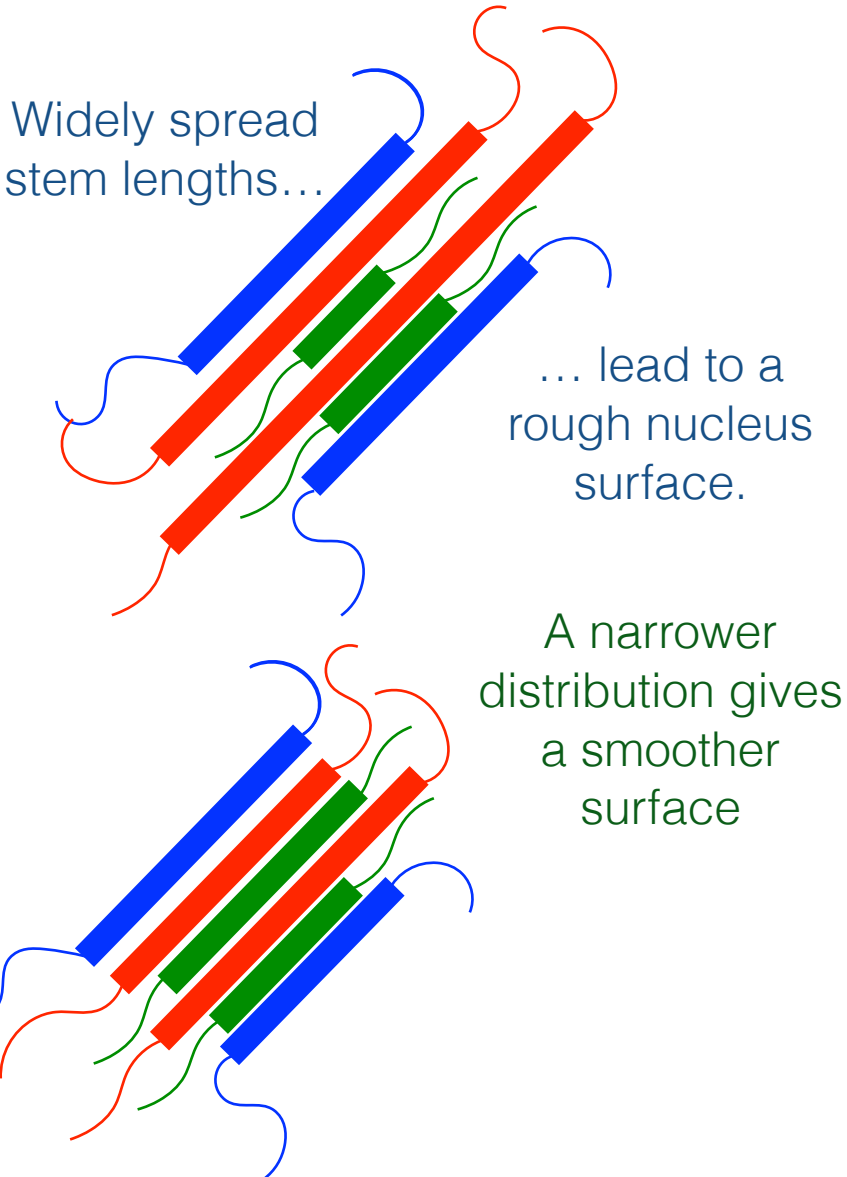
**Long chain depletion**



# Smooth-polyStrand model

**Nucleus roughness penalty**

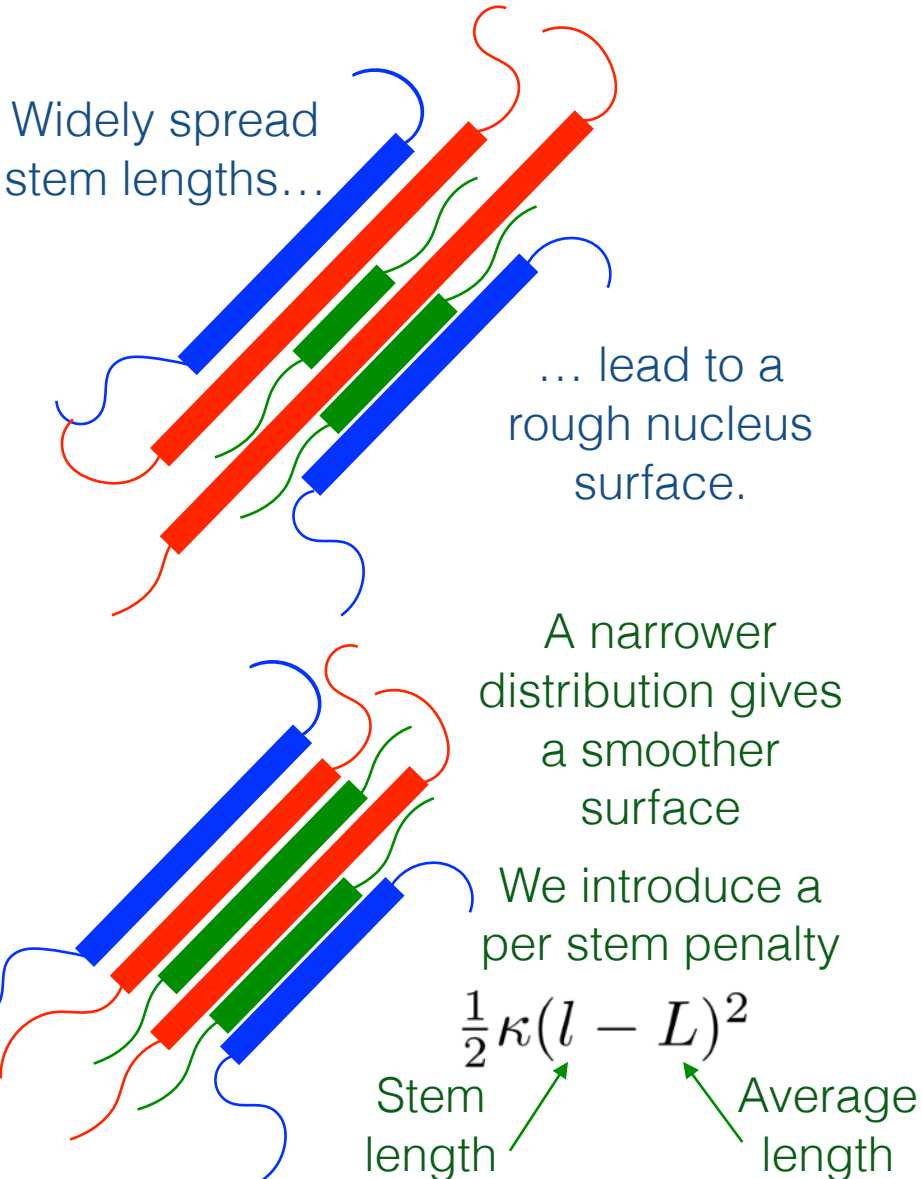
**Long chain depletion**



# Smooth-polyStrand model

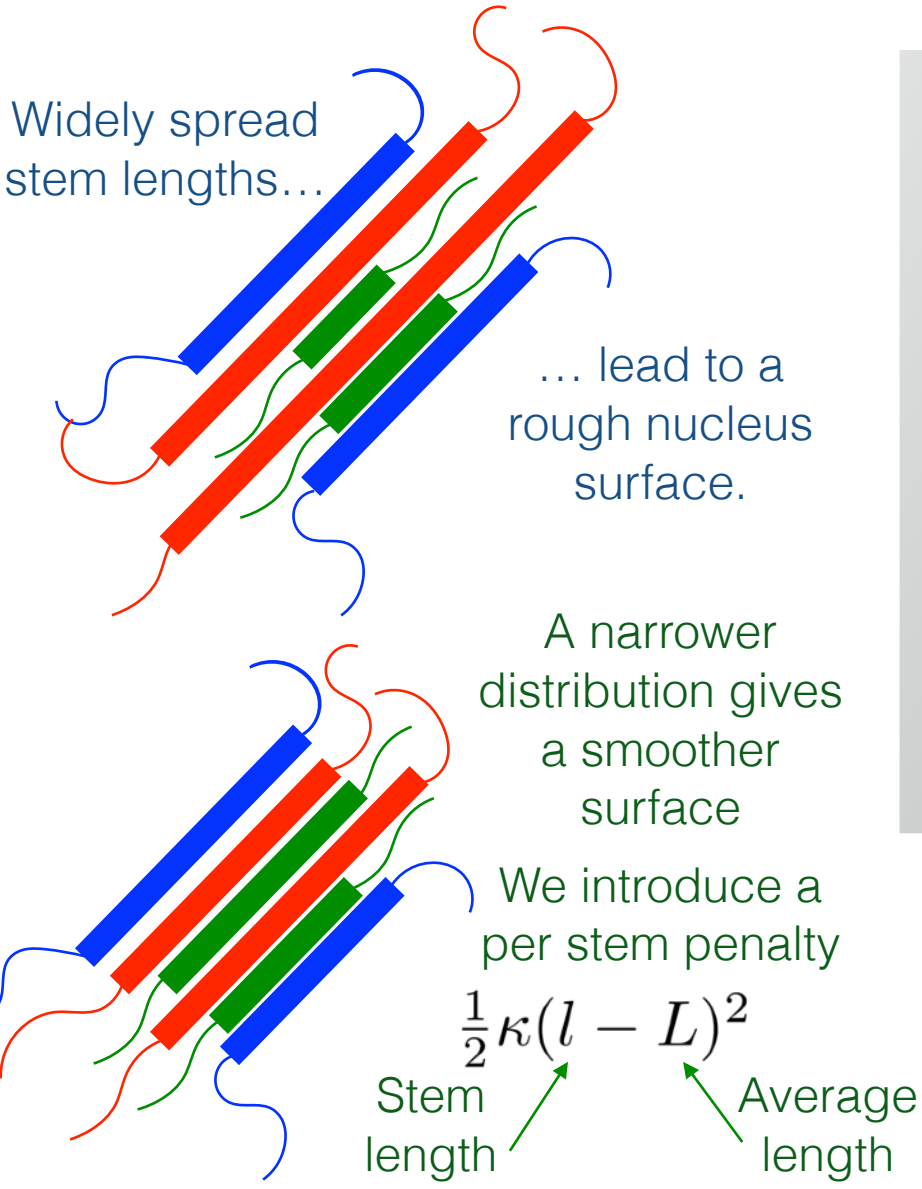
Nucleus roughness penalty

Long chain depletion

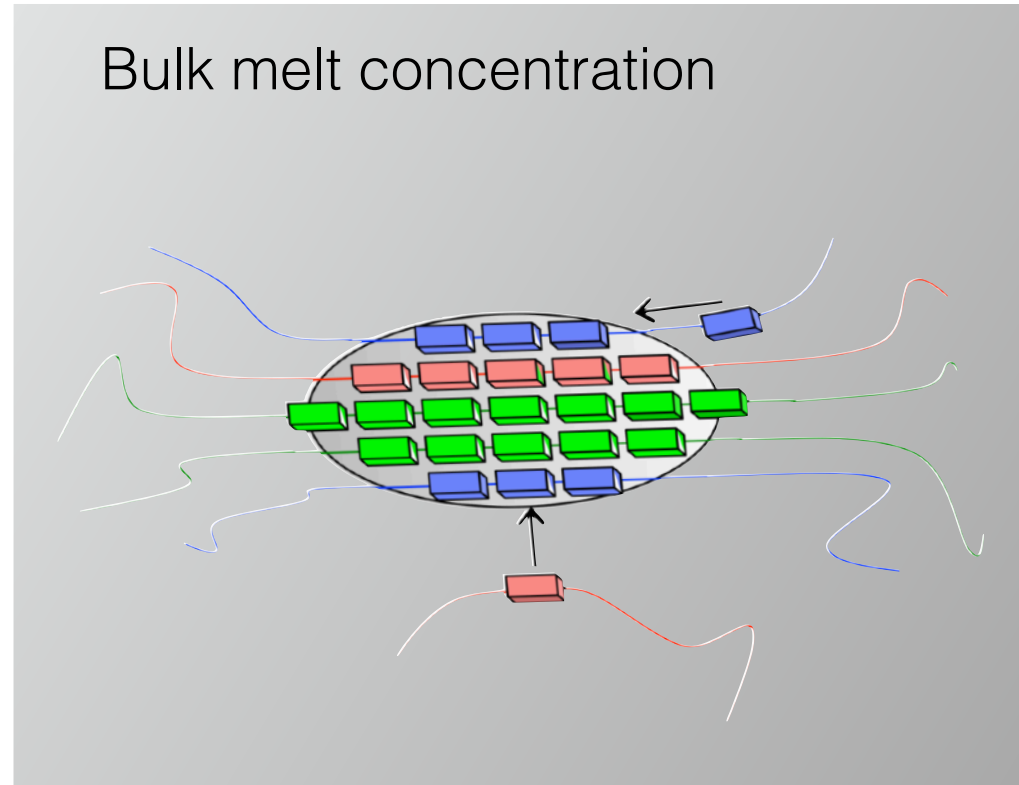


# Smooth-polyStrand model

## Nucleus roughness penalty



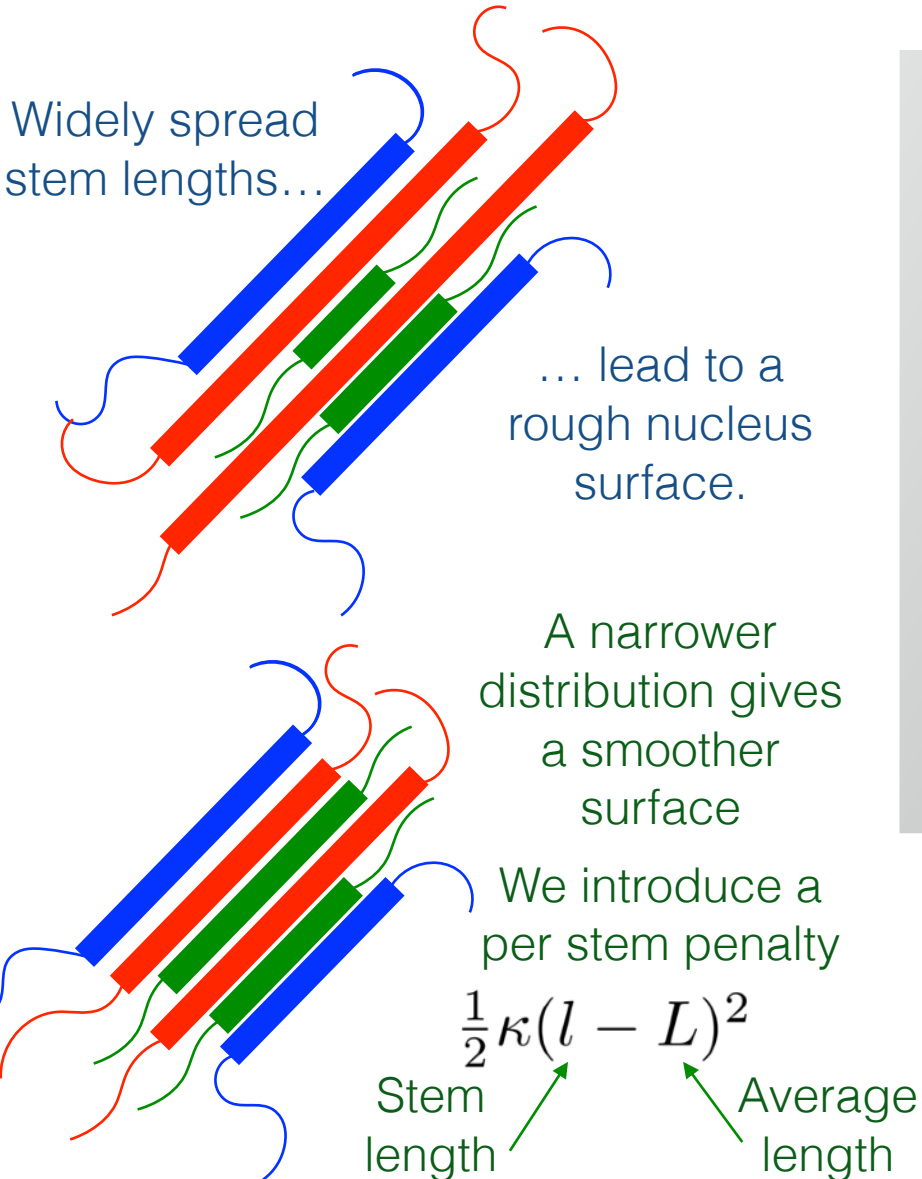
## Long chain depletion



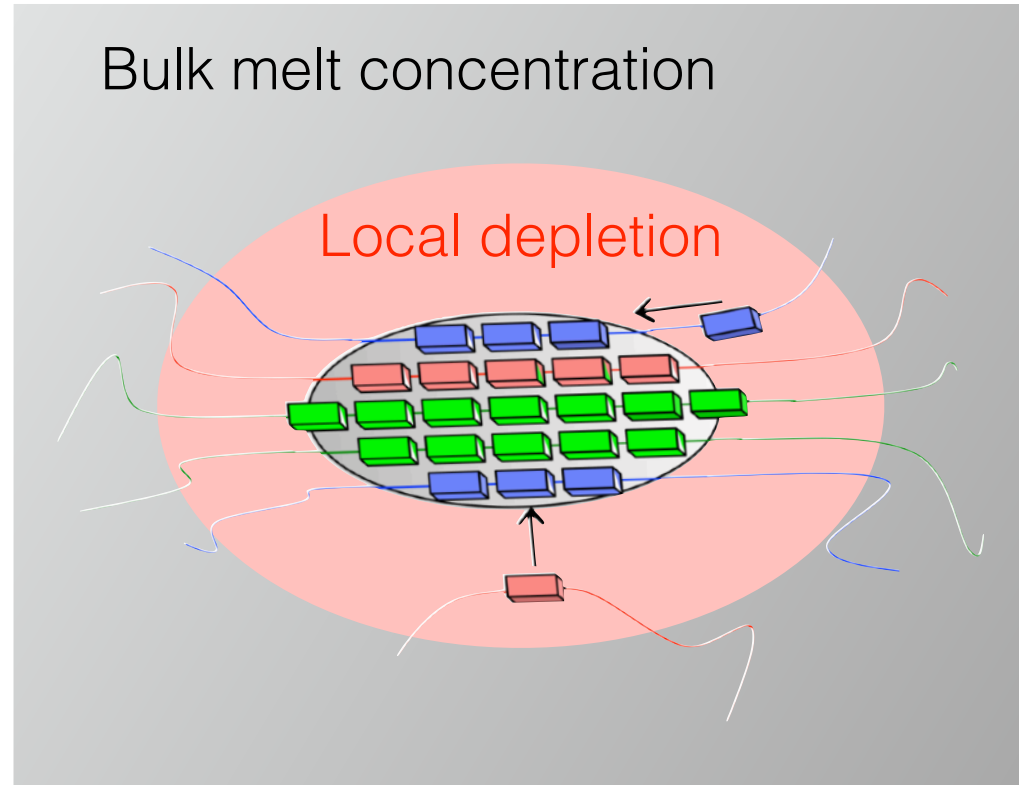


# Smooth-polyStrand model

## Nucleus roughness penalty

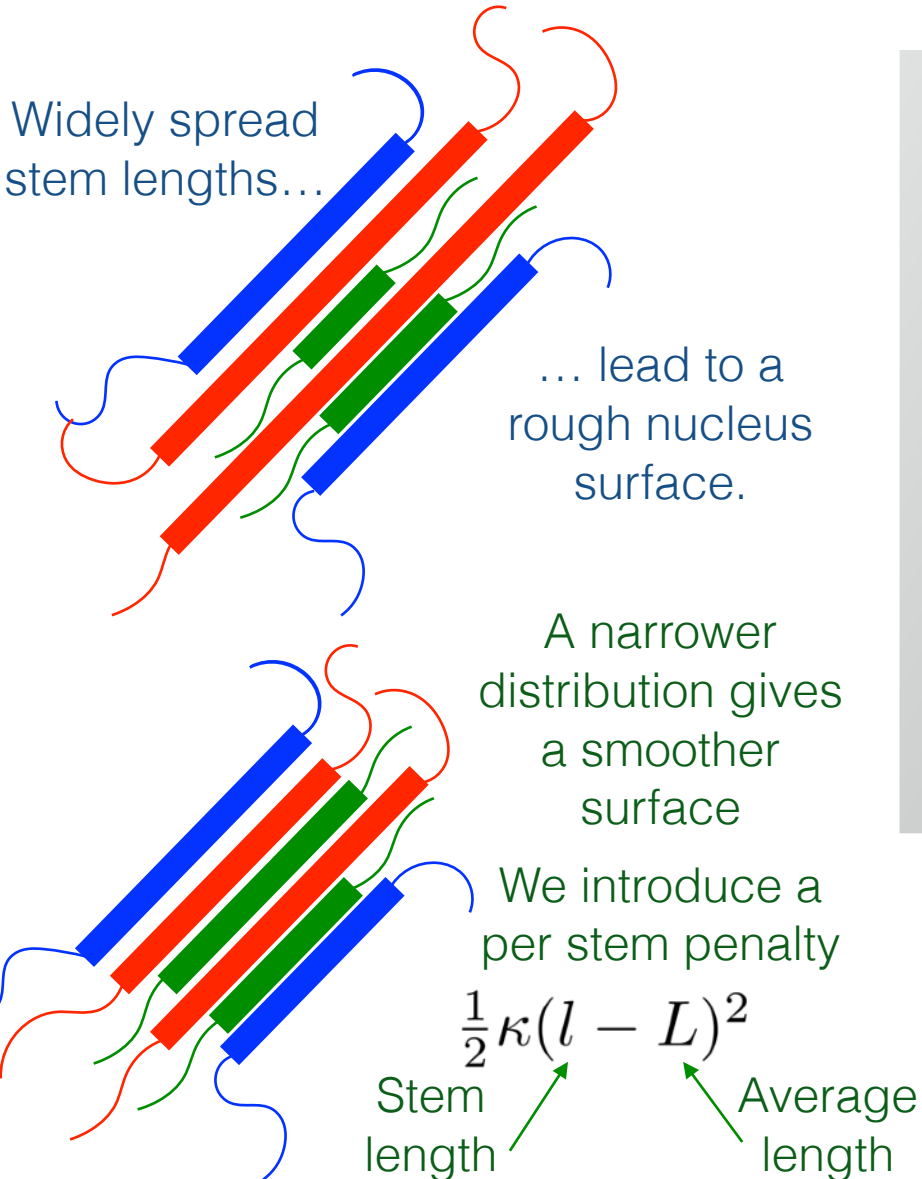


## Long chain depletion

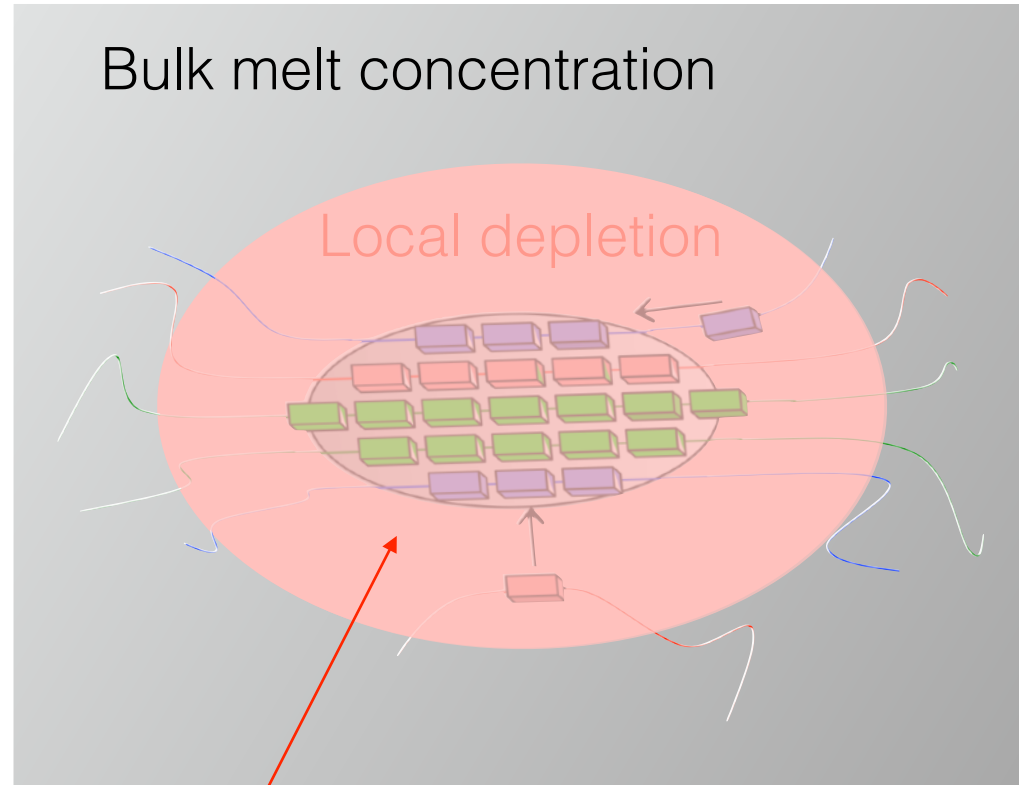


# Smooth-polyStrand model

## Nucleus roughness penalty



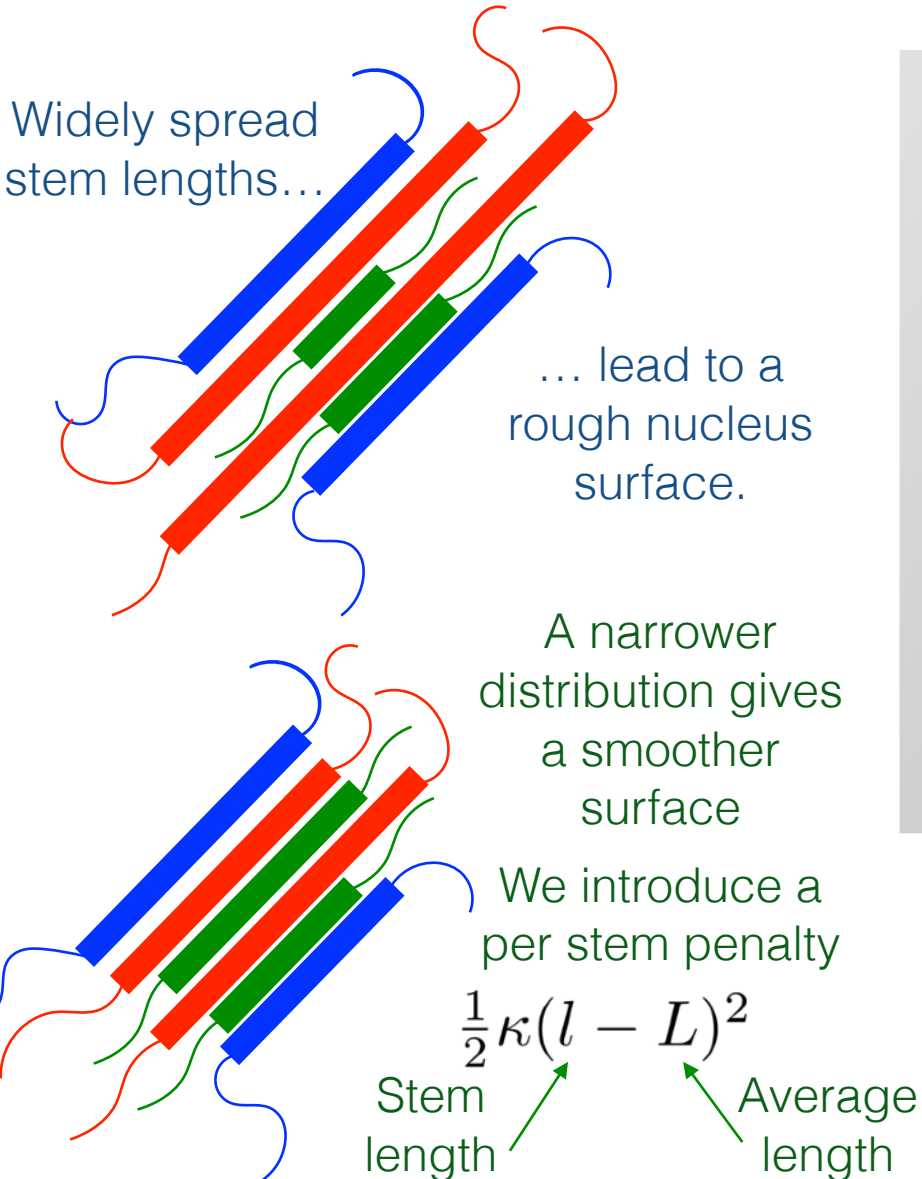
## Long chain depletion



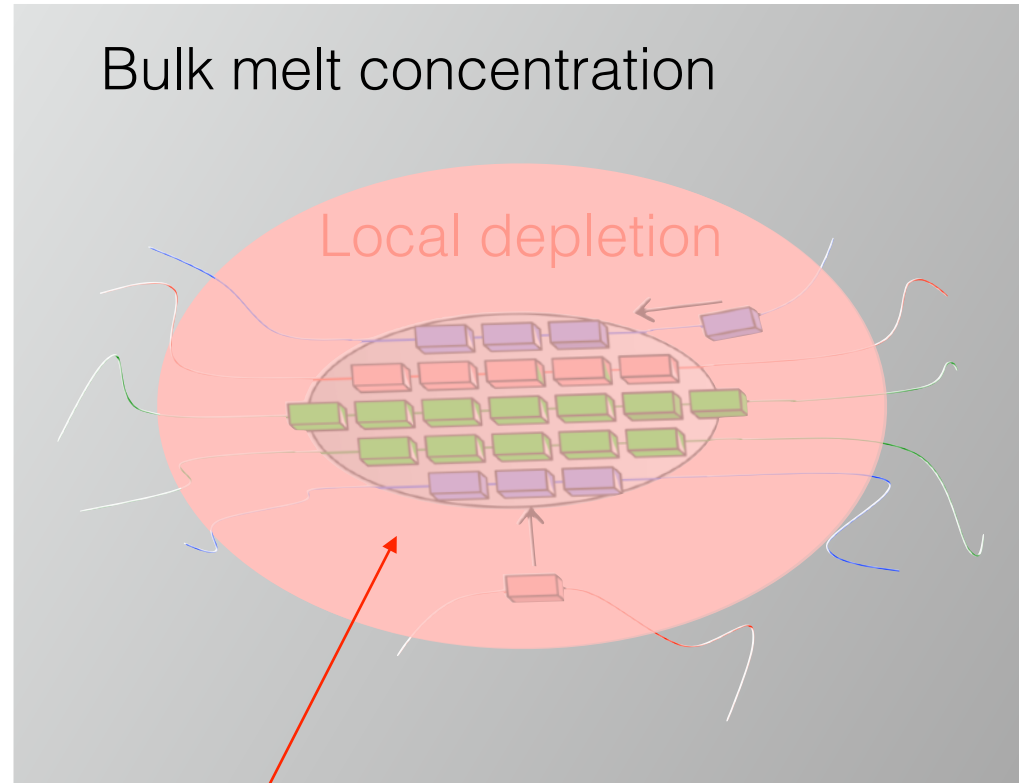
Stems are conserved over this region [nucleus + depletion region]

# Smooth-polyStrand model

## Nucleus roughness penalty



## Long chain depletion

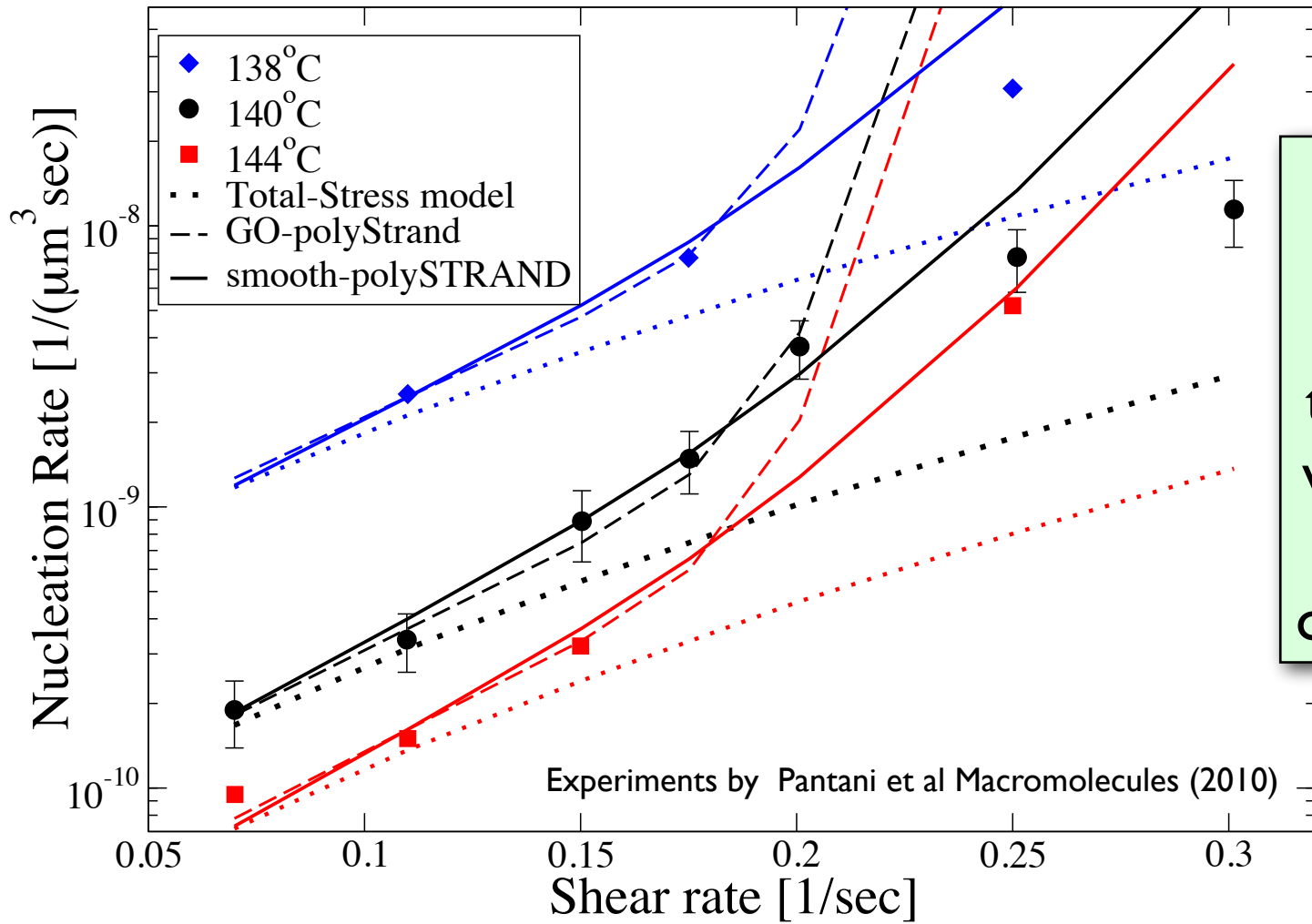


Stems are conserved over this region [nucleus + depletion region]

$$Q_s = Q_{s0} N_s$$

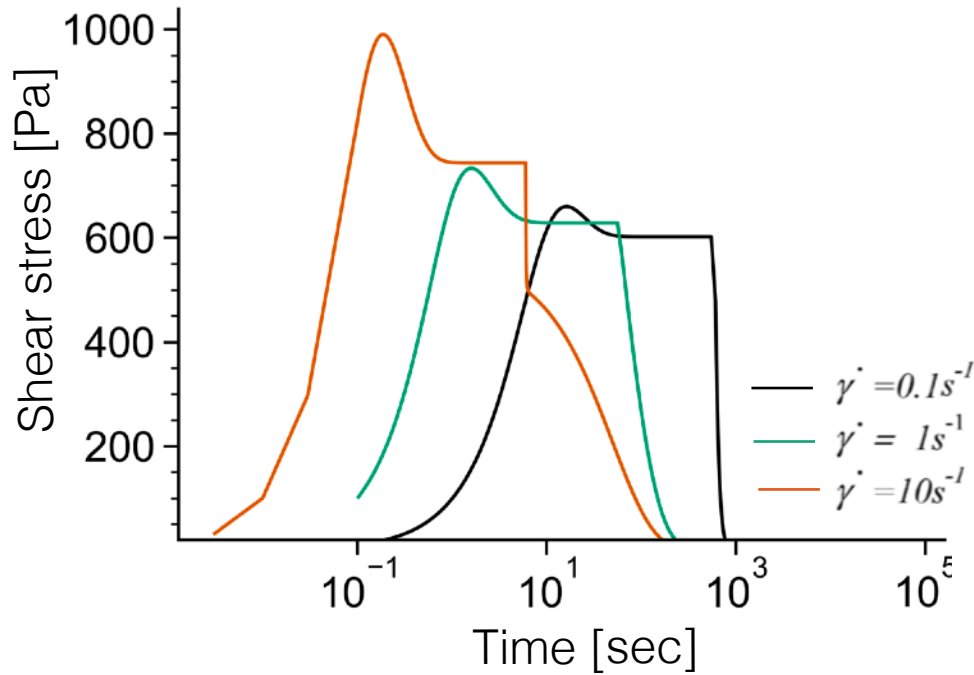
Number of available stems,  $Q_s$ , grows with nucleus size,  $N_s$ .

# Direct observation of nucleation during steady shear



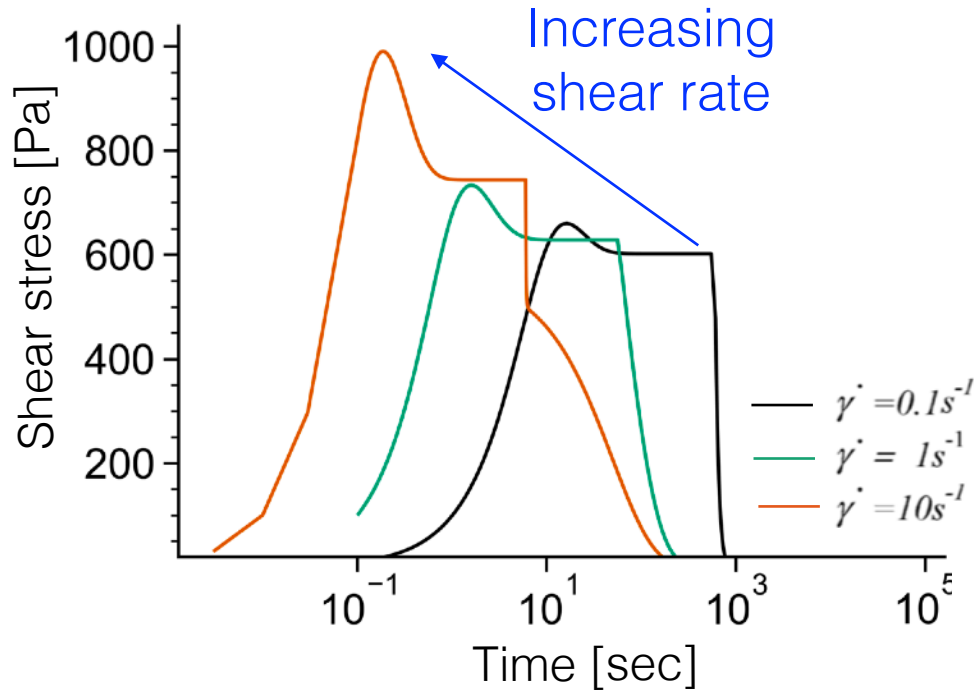
The smooth-polyStrand model retains the curvature, while relieving some of the overprediction

# Shear pulse experiments



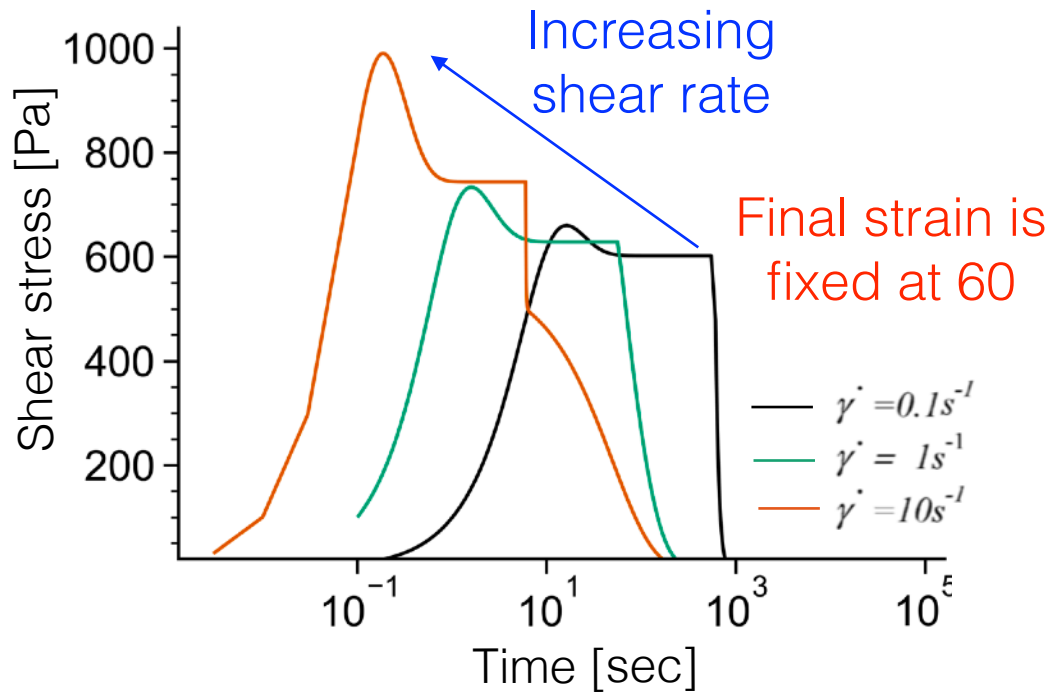
Experiments on isotactic poly(1-butene) [ $M_w = 116-398$  kg/mol] by Acierno et al Rheol Acta (2003)

# Shear pulse experiments



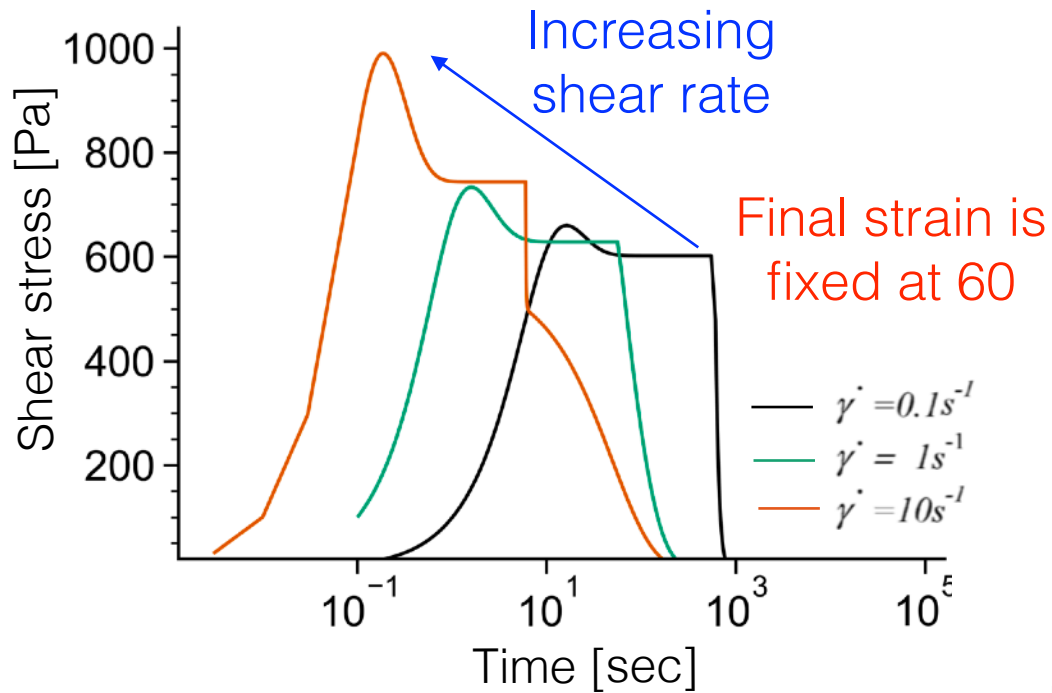
Experiments on isotactic poly(1-butene) [Mw=116-398 kg/mol] by Acierno et al Rheol Acta (2003)

# Shear pulse experiments

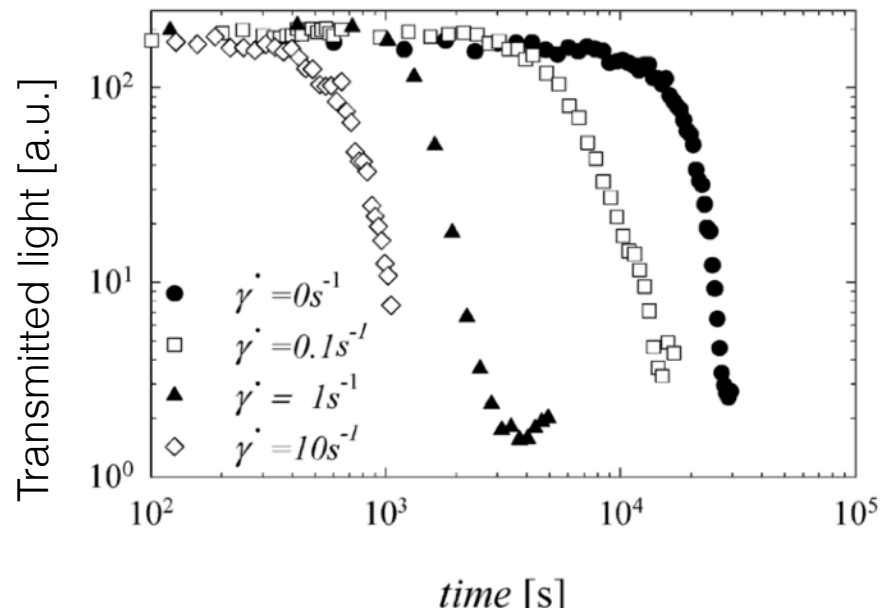


Experiments on isotactic poly(1-butene) [ $M_w = 116\text{-}398$  kg/mol] by Acierno et al Rheol Acta (2003)

# Shear pulse experiments

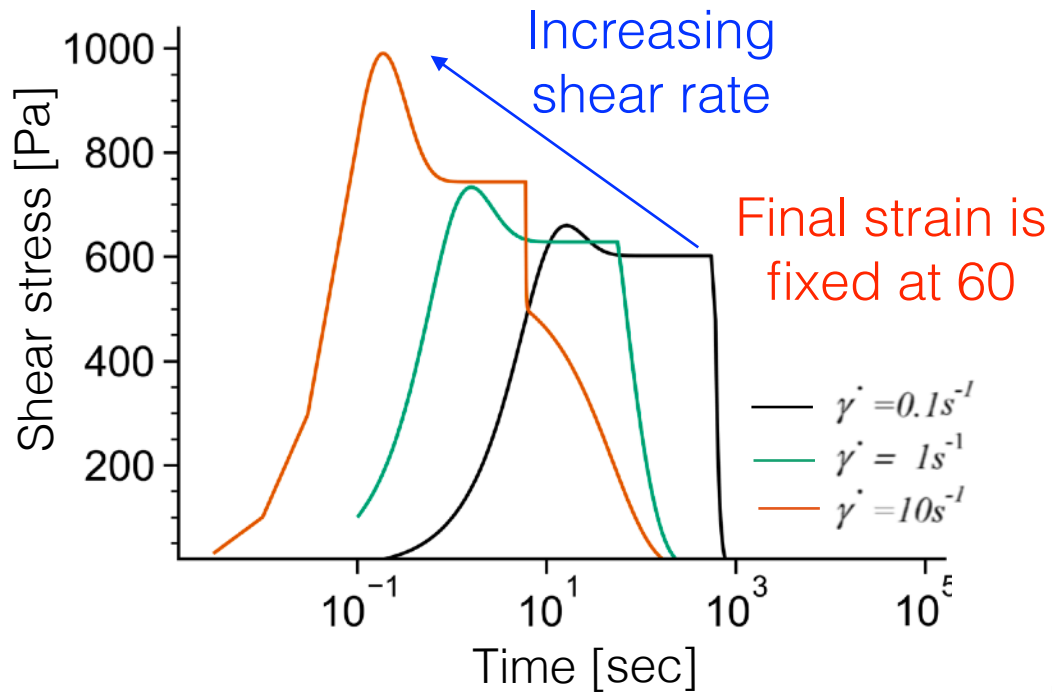


Experiments on isotactic poly(1-butene) [ $M_w = 116-398$  kg/mol] by Acierno et al Rheol Acta (2003)

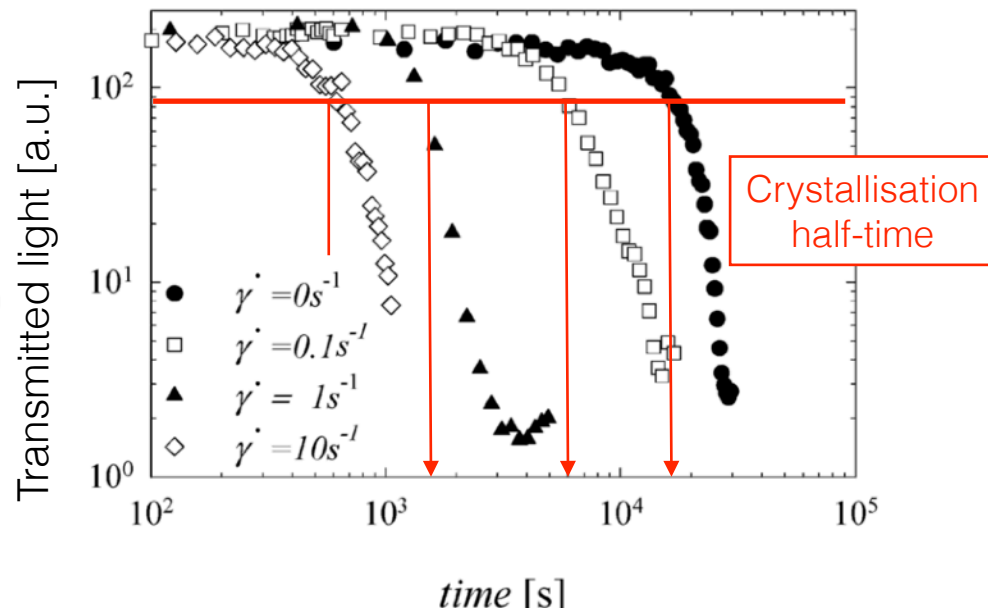




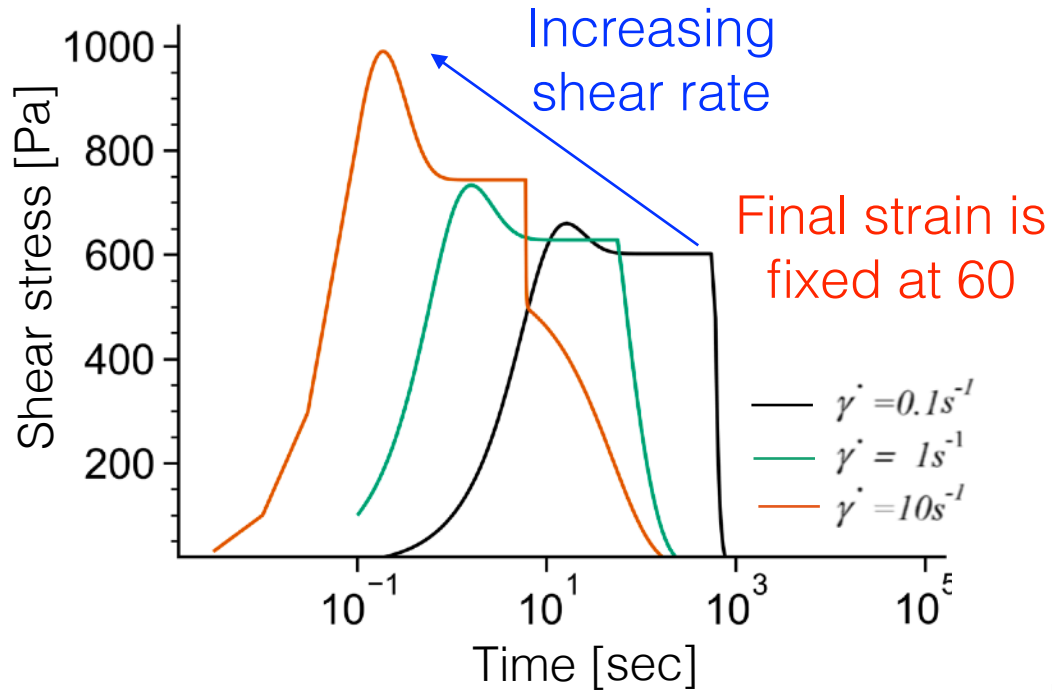
# Shear pulse experiments



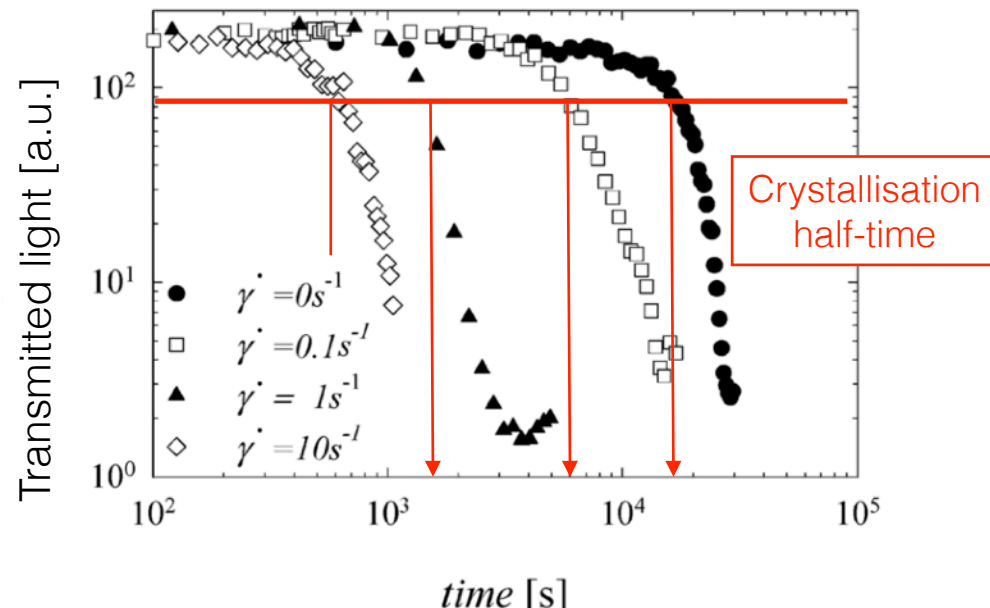
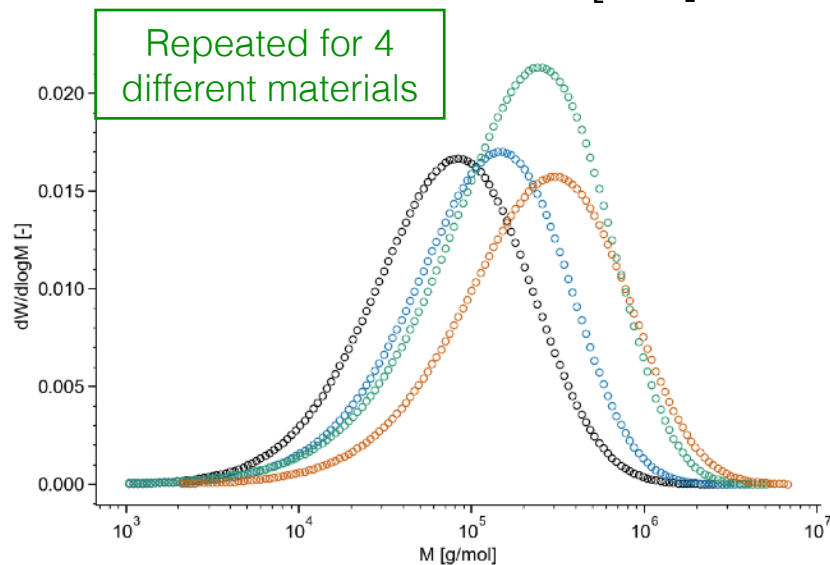
Experiments on isotactic poly(1-butene) [ $M_w = 116-398$  kg/mol] by Acierno et al Rheol Acta (2003)



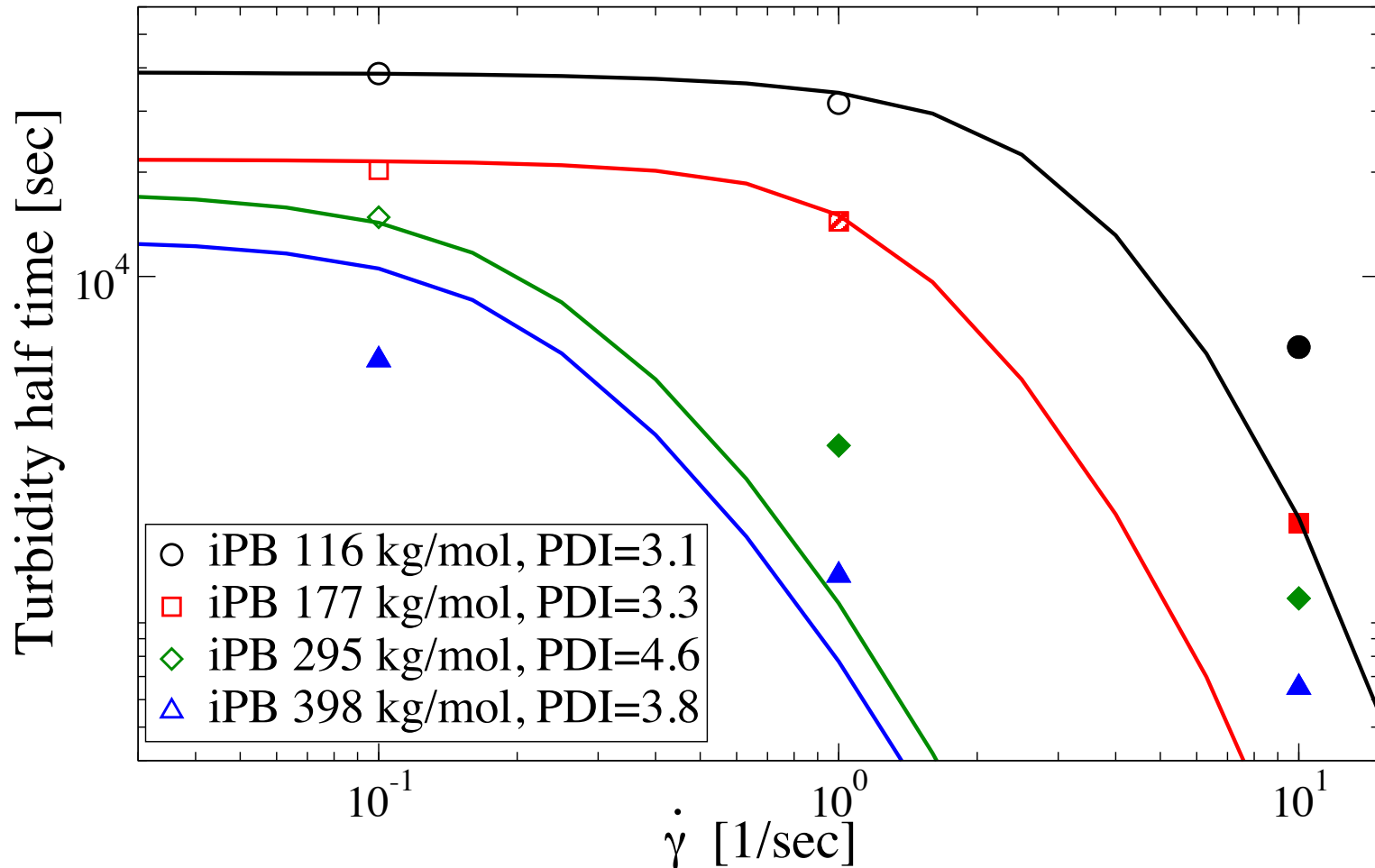
# Shear pulse experiments



Experiments on isotactic poly(1-butene) [ $M_w = 116-398$  kg/mol] by Acierno et al Rheol Acta (2003)



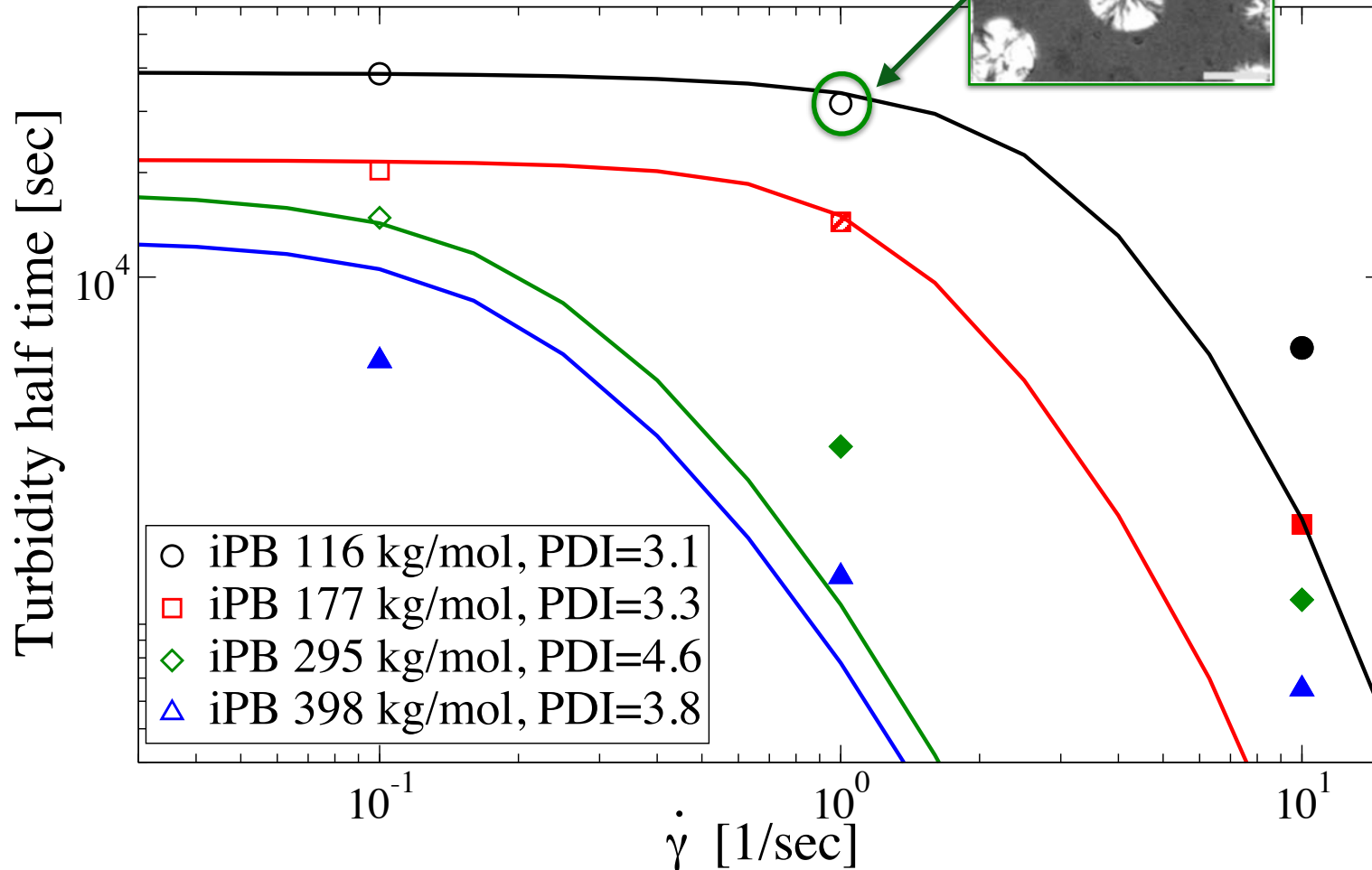
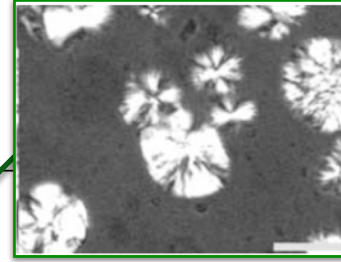
# Shear pulse experiments



- The same FIC parameters are used for all 4 curves
- Model captures variation due to molecular weight distribution

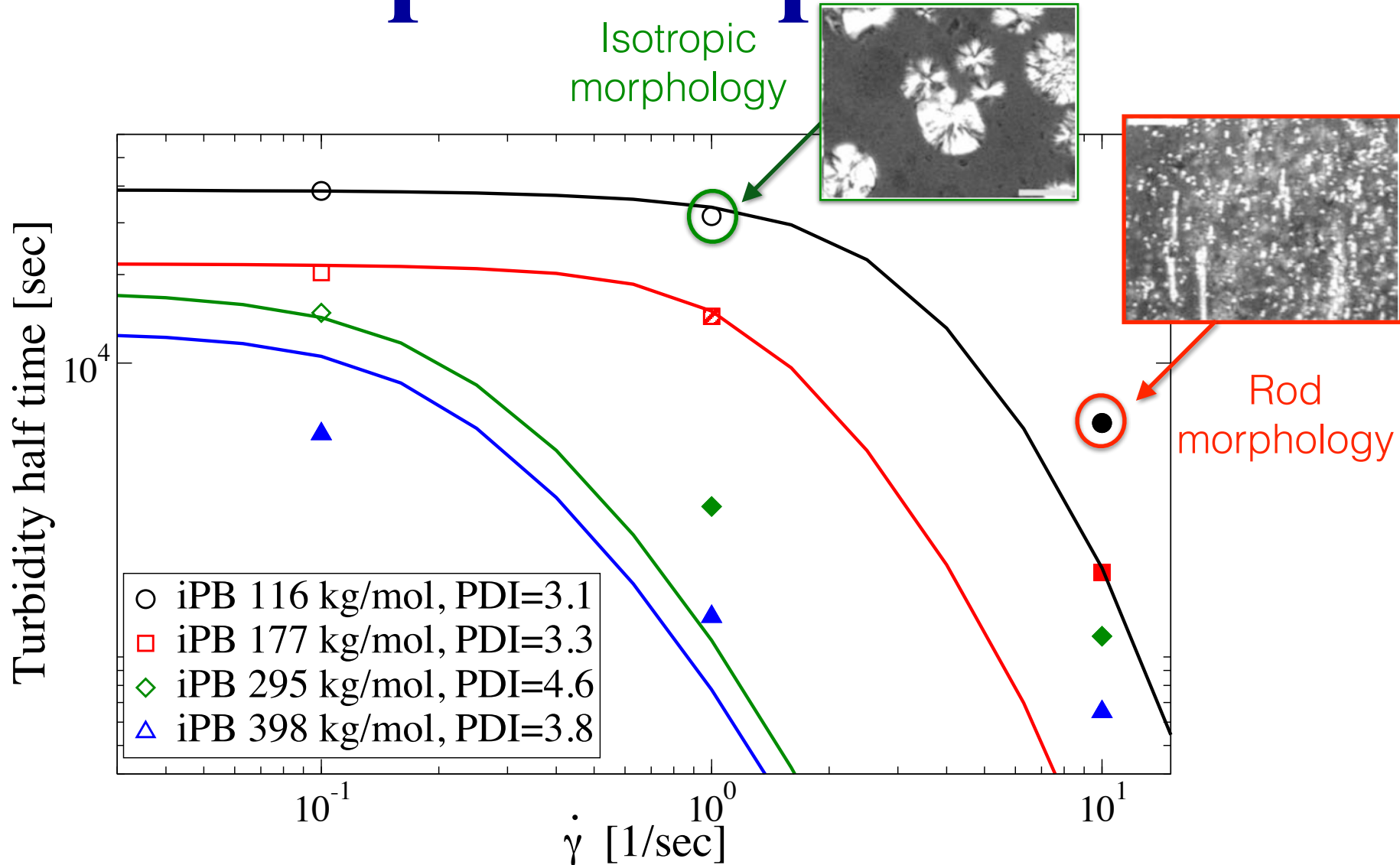
# Shear pulse experiments

Isotropic morphology



- The same FIC parameters are used for all 4 curves
- Model captures variation due to molecular weight distribution

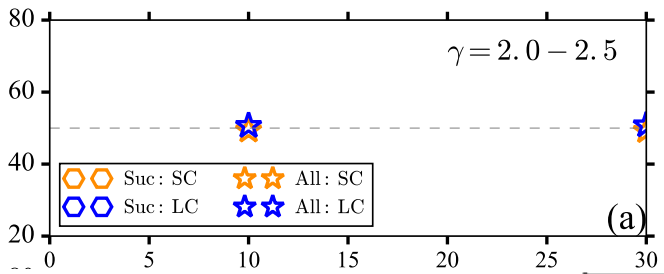
# Shear pulse experiments



- The same FIC parameters are used for all 4 curves
- Model captures variation due to molecular weight distribution

**Is long chain enhancement  
seen in Molecular Dynamics  
simulations?**

Nucleus Composition

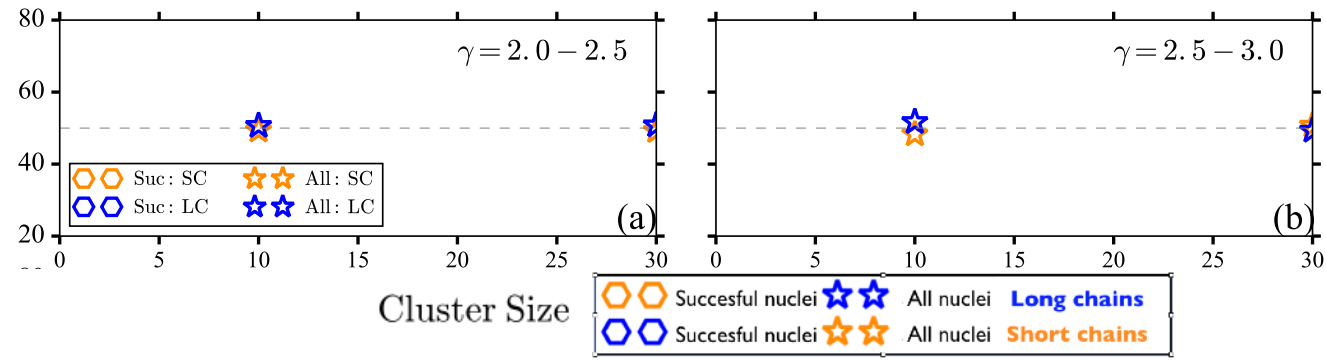


Cluster Size



MD simulations of bidisperse PE - 50:50 C1000/C125 chains  $\dot{\gamma}\tau_{R, long} = 10$

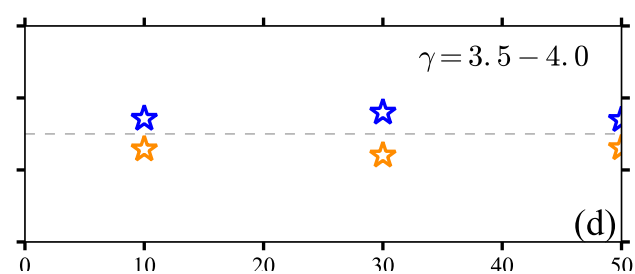
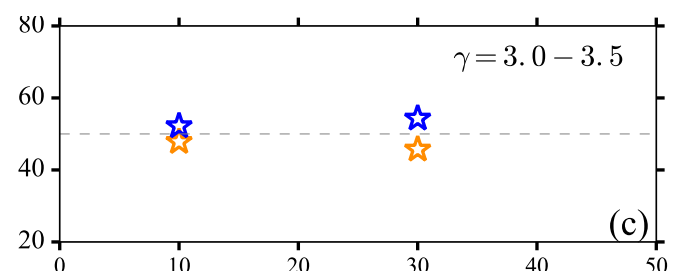
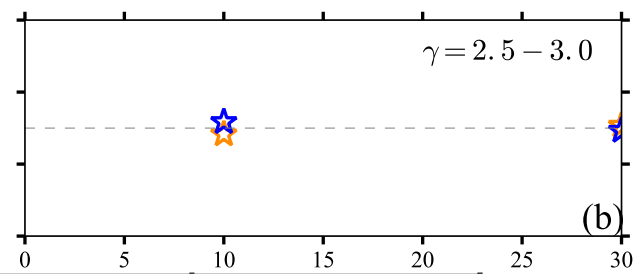
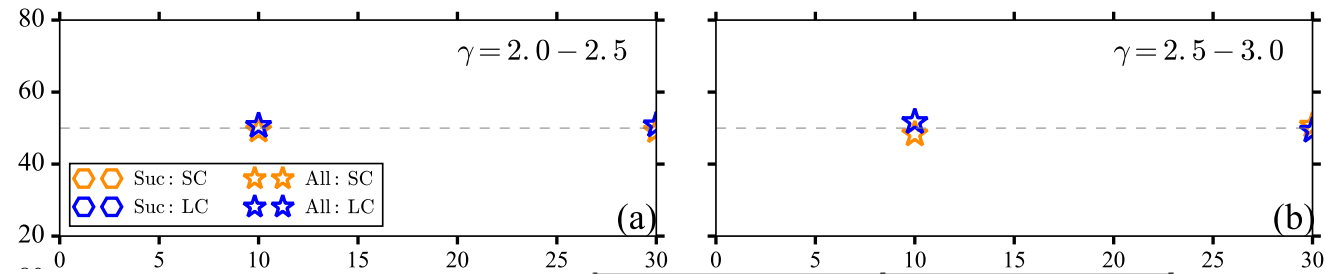
Nucleus Composition



MD simulations of bidisperse PE - 50:50 CI000/CI25 chains  $\dot{\gamma}\tau_{R, \text{long}} = 10$

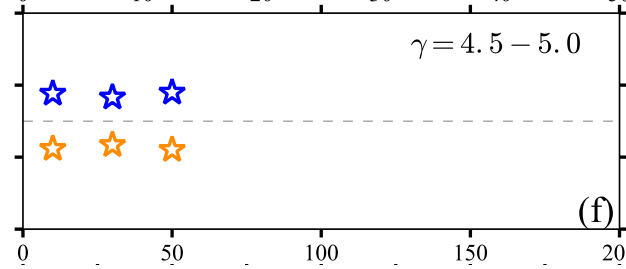
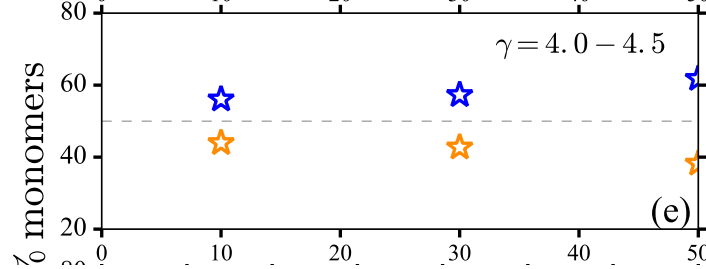
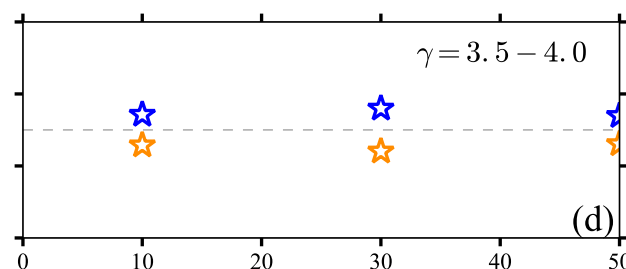
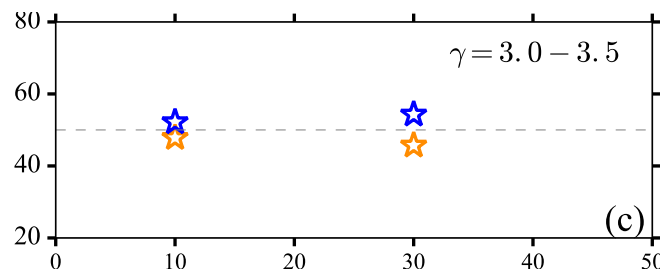
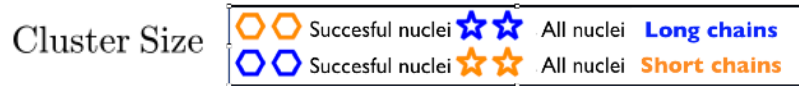
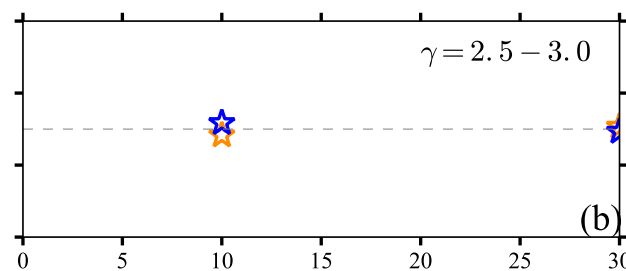
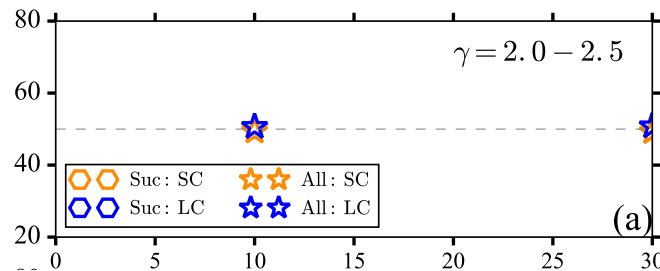


Nucleus Composition

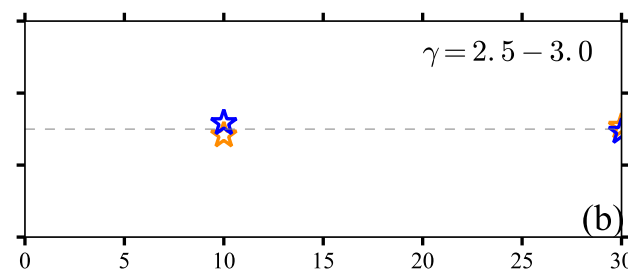
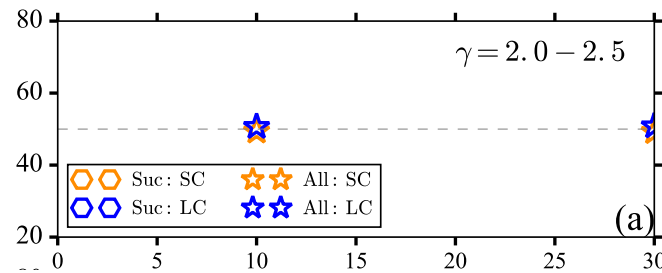


MD simulations of bidisperse PE - 50:50 CI000/CI25 chains  $\dot{\gamma}\tau_{R, \text{long}} = 10$

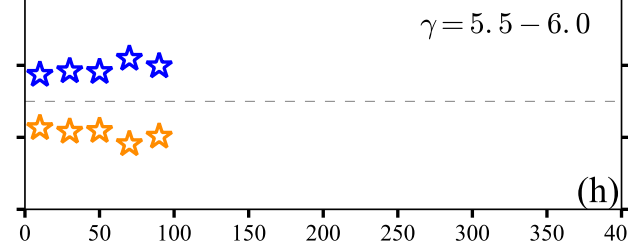
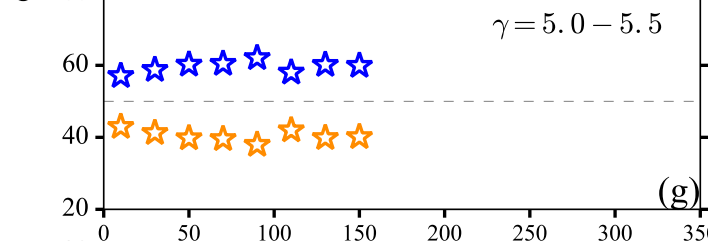
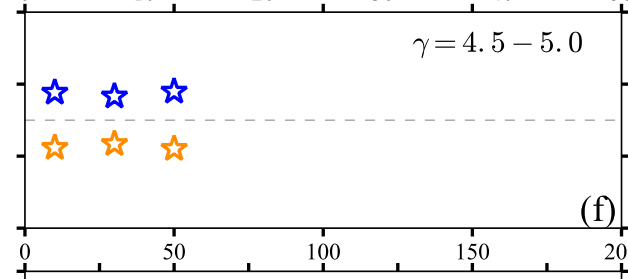
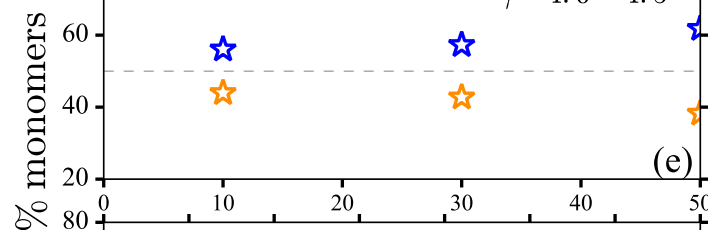
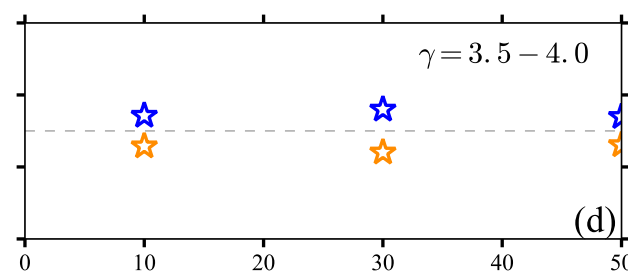
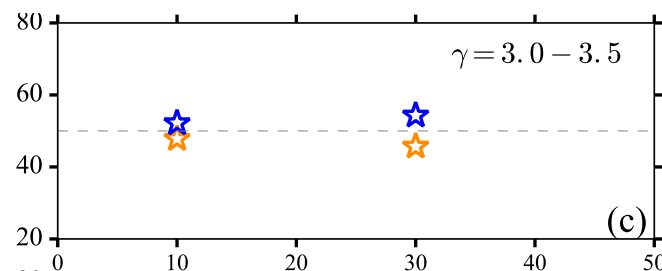
Nucleus Composition



MD  
simulations  
of bidisperse  
PE - 50:50  
CI000/CI25  
chains  
 $\dot{\gamma} \tau_{R, \text{long}} = 10$



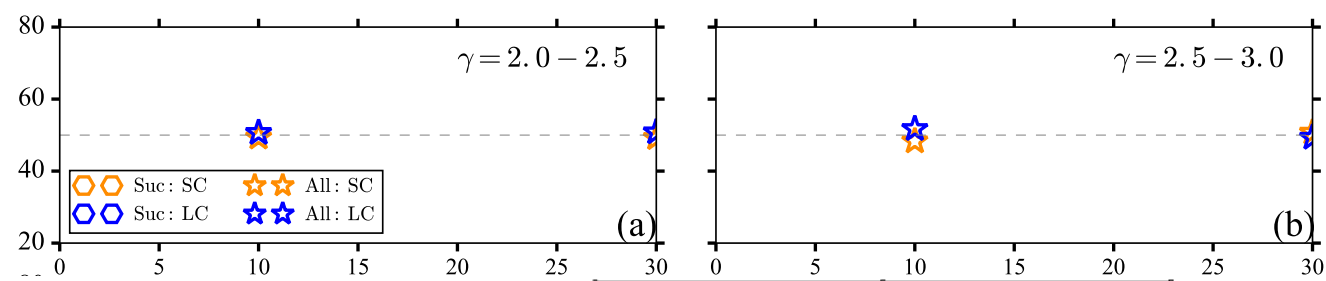
Cluster Size



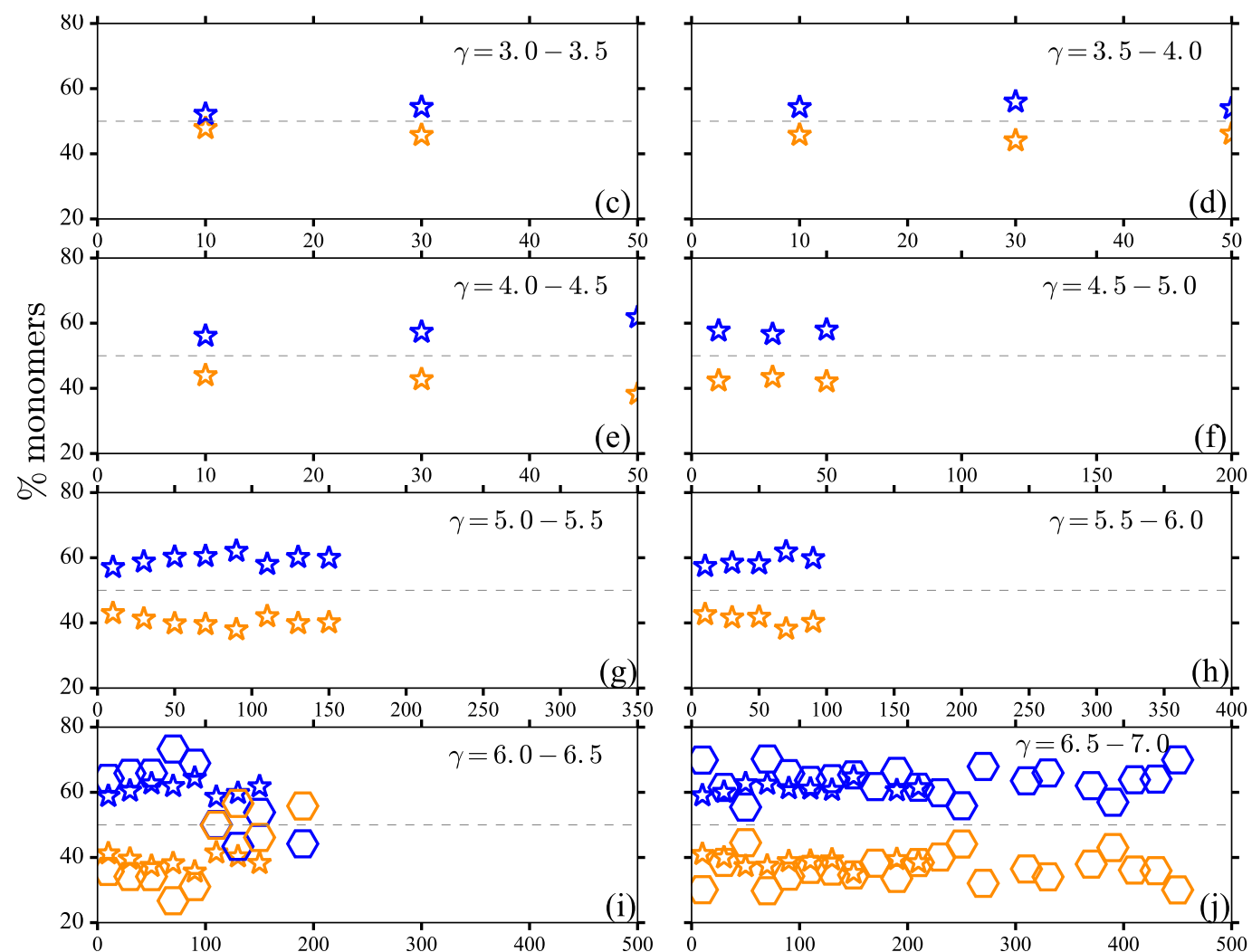
% monomers

MD  
 simulations  
 of bidisperse  
 PE - 50:50  
 CI000/CI25  
 chains  
 $\dot{\gamma}\tau_{R, \text{long}} = 10$

Nucleus Composition



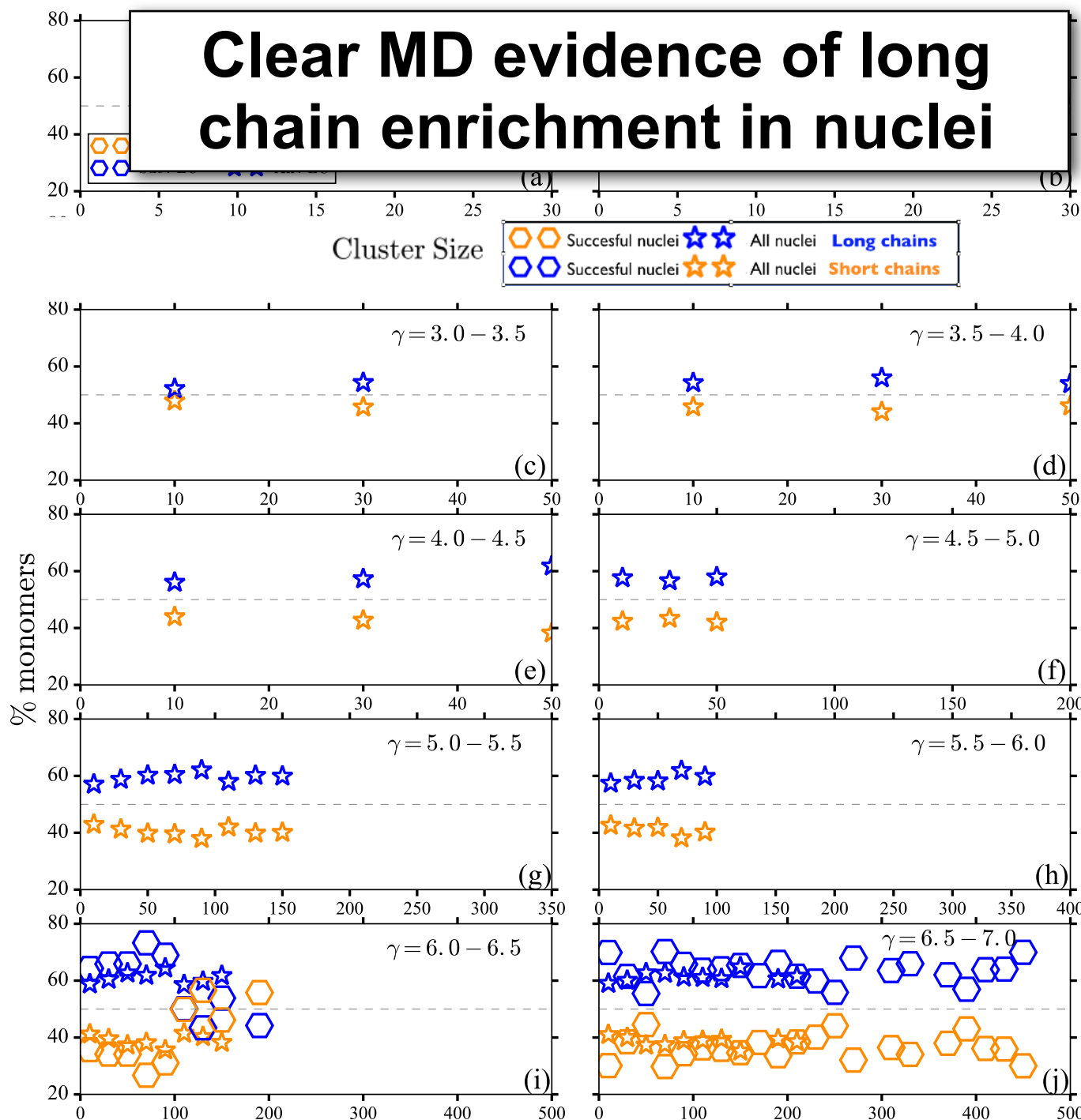
MD simulations of bidisperse PE - 50:50 C1000/C125 chains  $\dot{\gamma} \tau_{R, \text{long}} = 10$



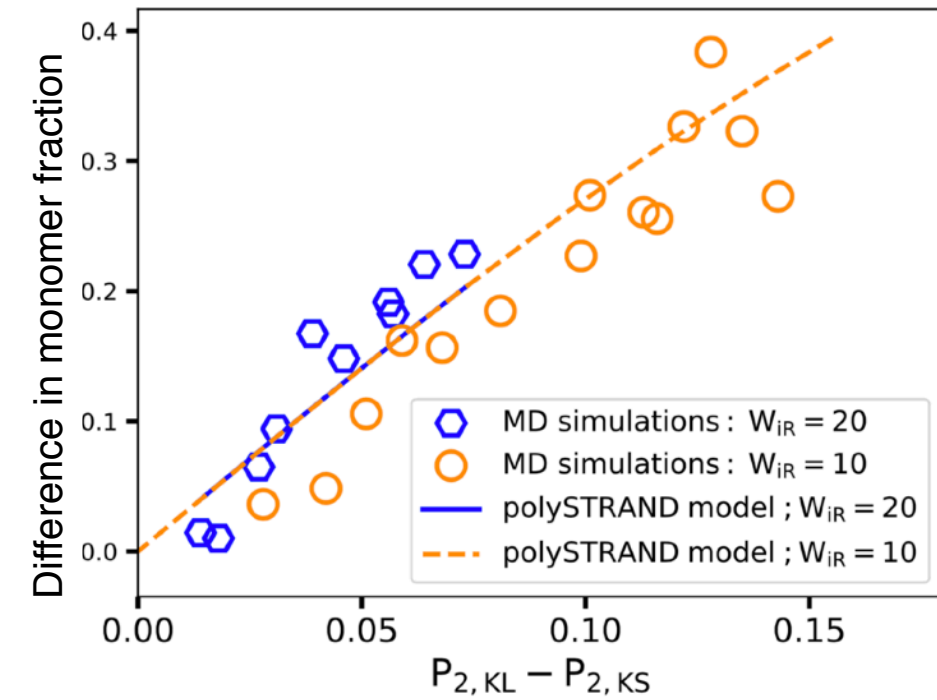
# Clear MD evidence of long chain enrichment in nuclei

MD simulations of bidisperse PE - 50:50 C1000/C125 chains

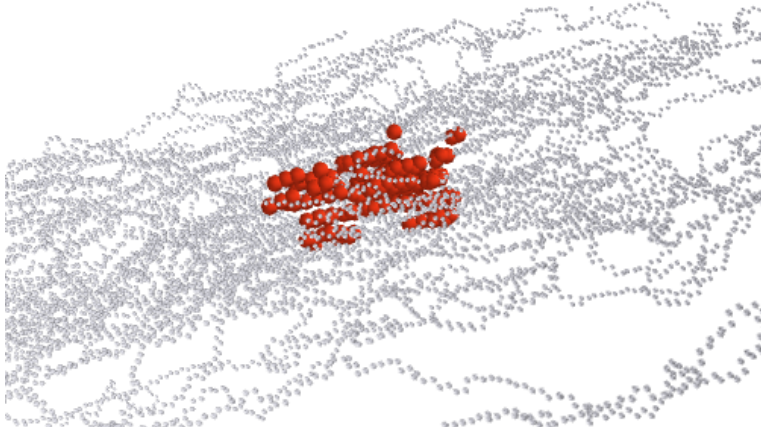
$$\dot{\gamma} \tau_{R, \text{long}} = 10$$



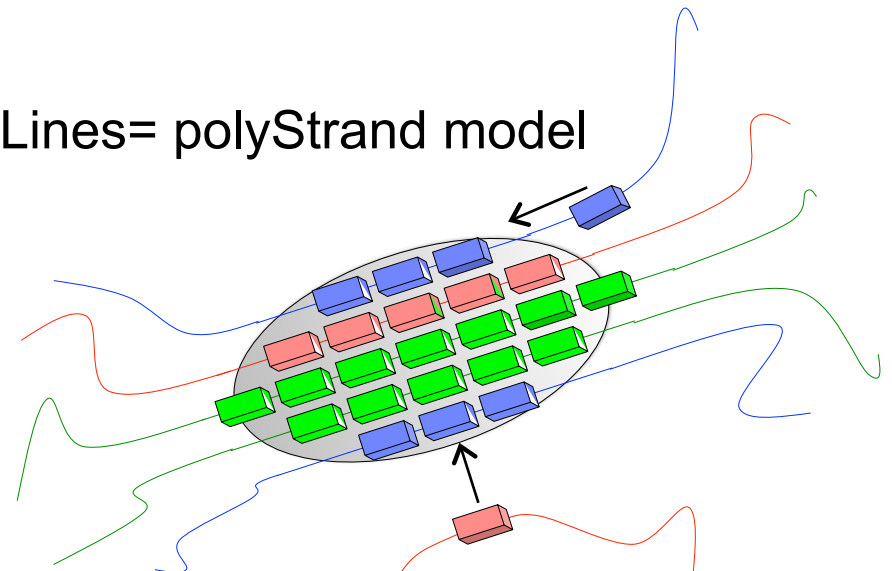
# Quantifying the long chain enhancement



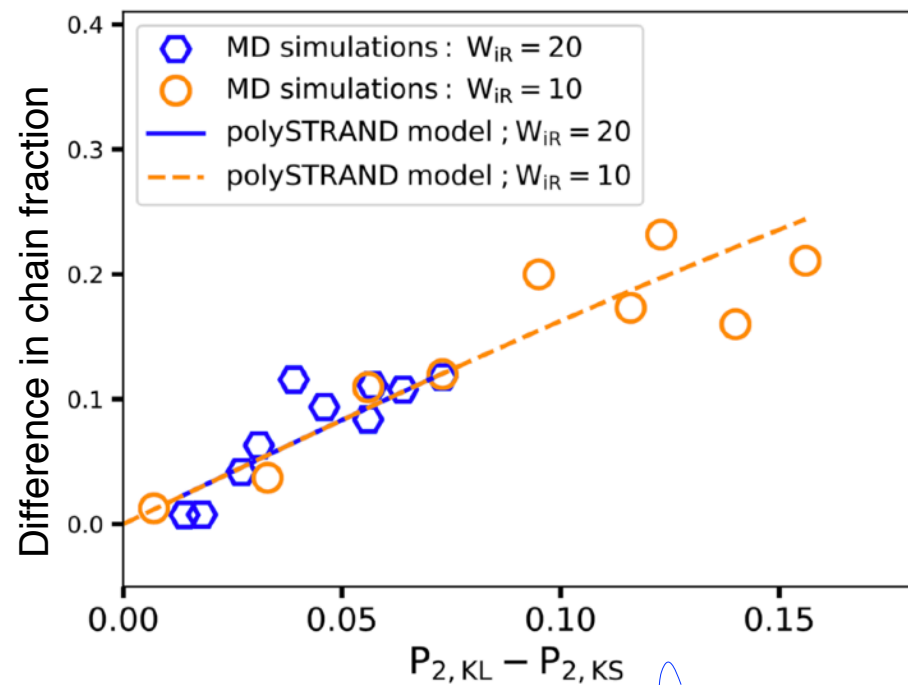
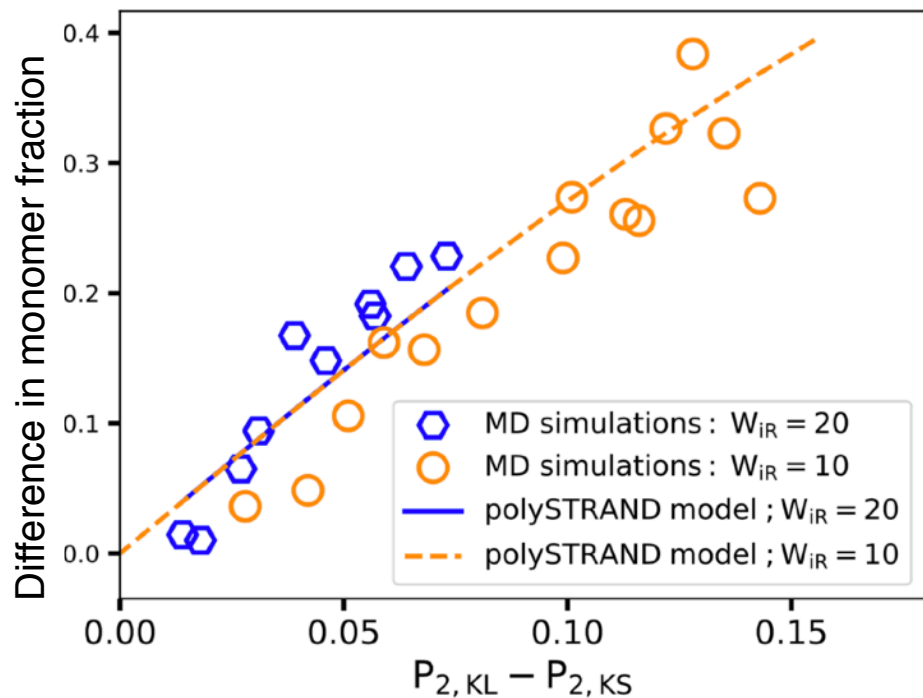
Symbols= Molecular dynamics



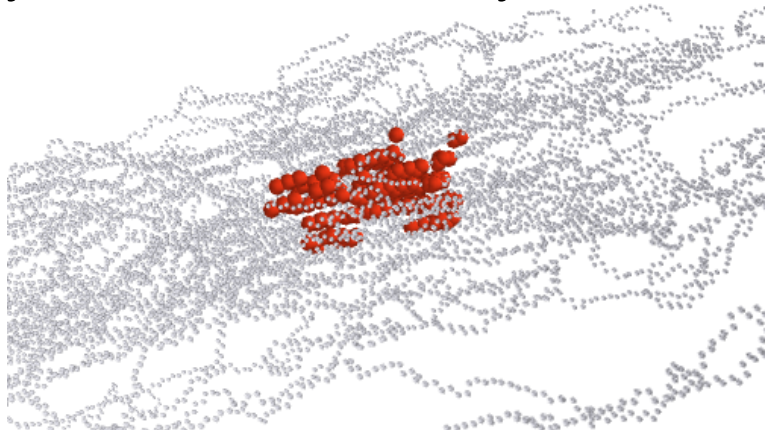
Lines= polyStrand model



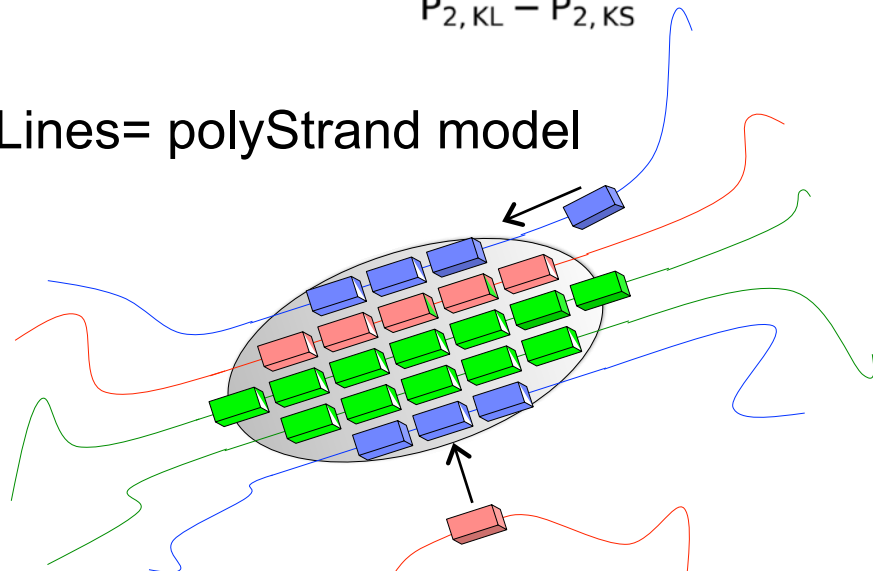
# Quantifying the long chain enhancement



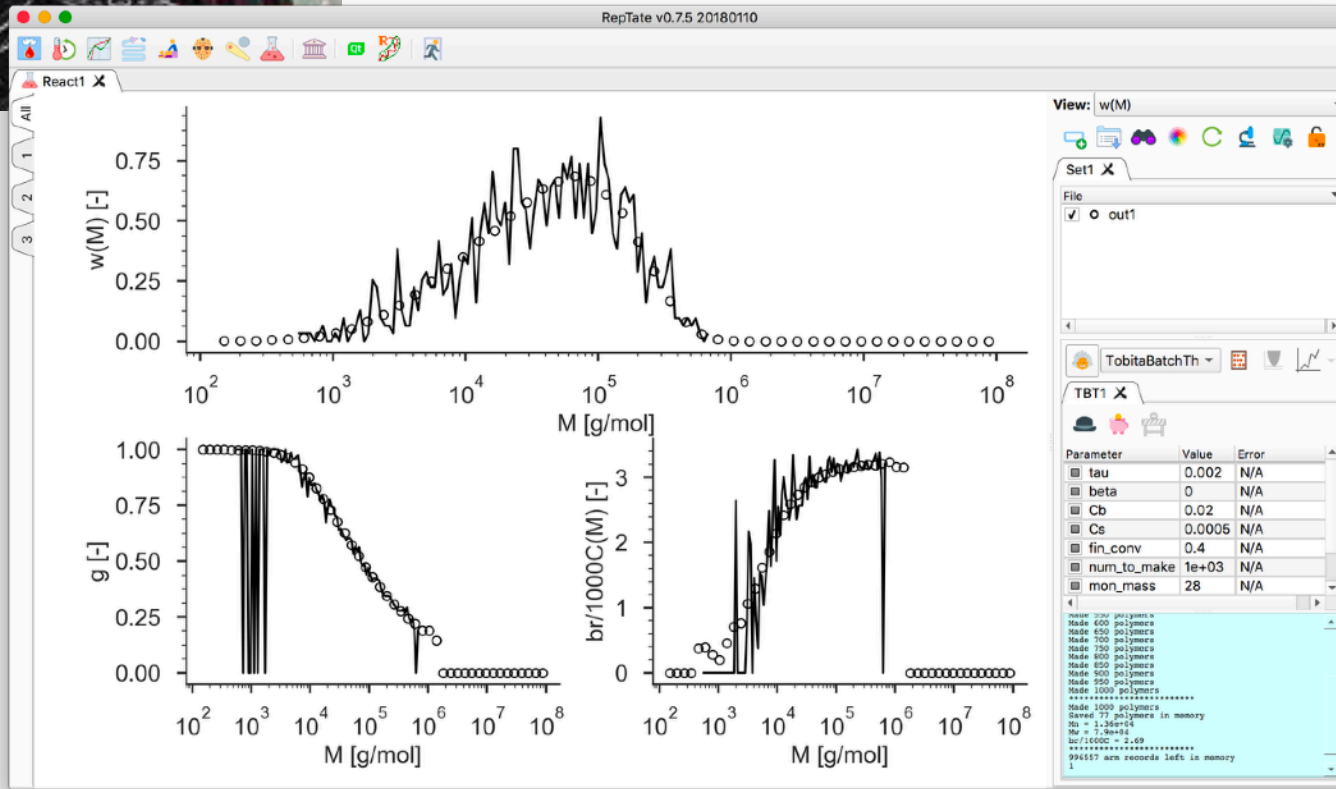
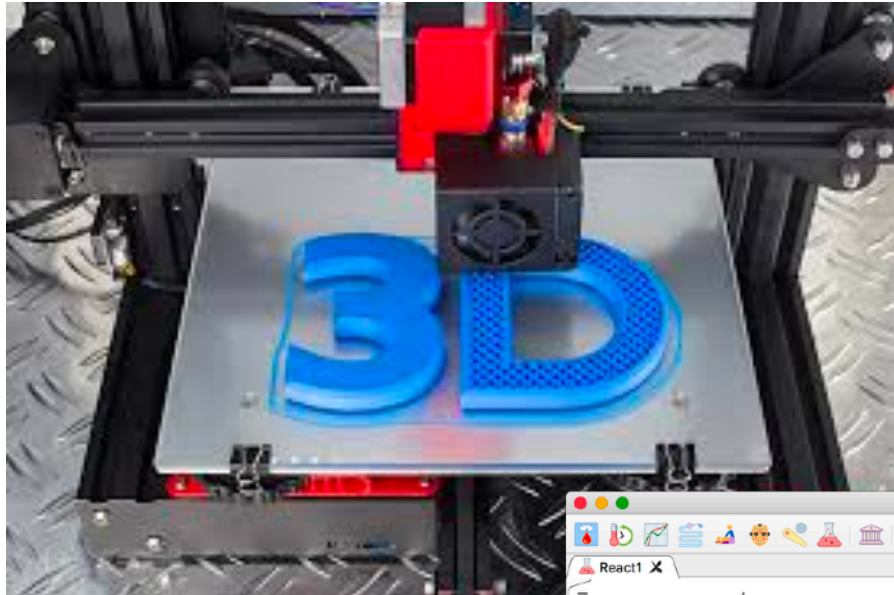
Symbols= Molecular dynamics



Lines= polyStrand model



# Applications





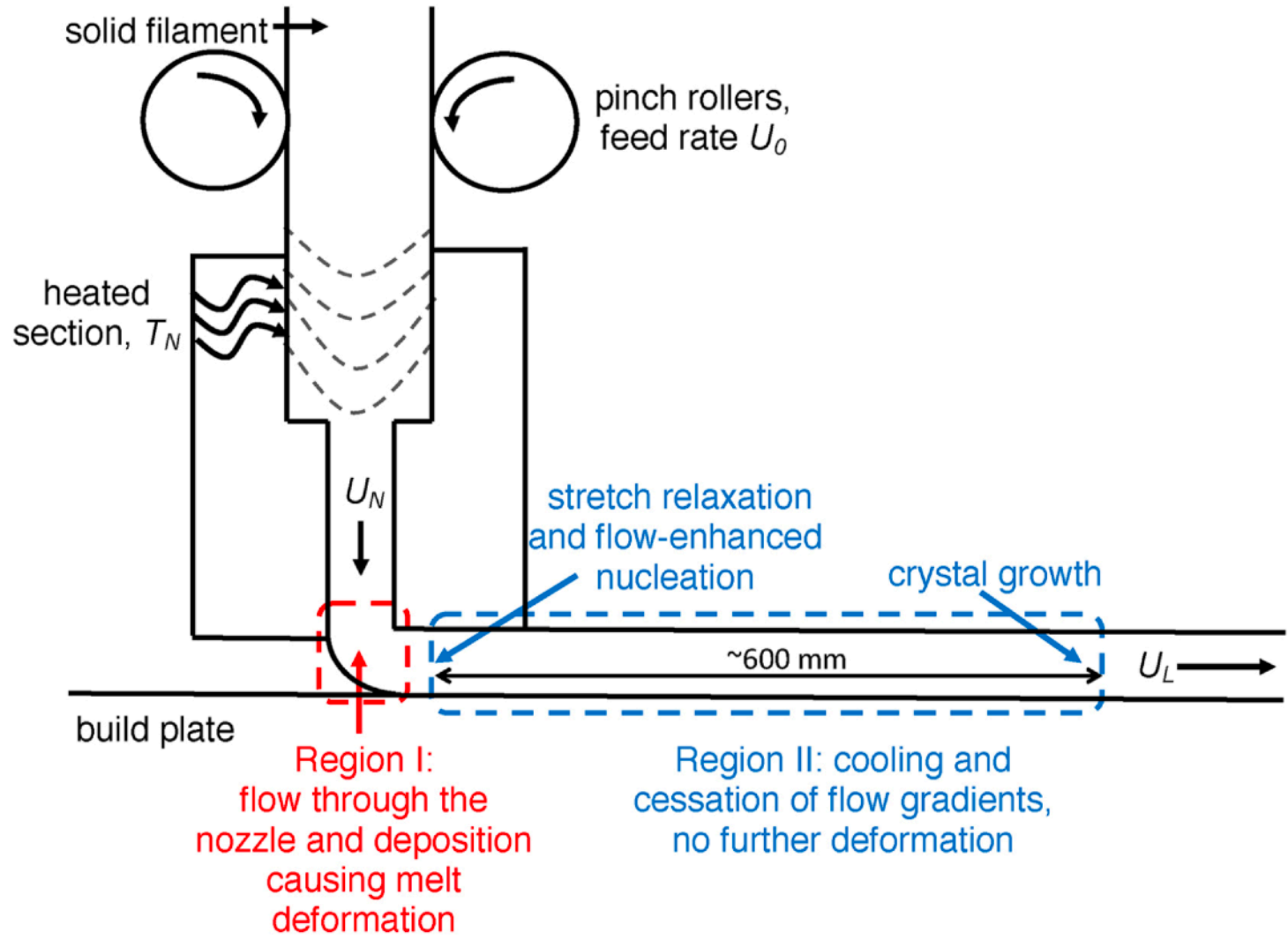
# Modelling crystallisation in 3D-printing



# Modelling crystallisation in 3D-printing

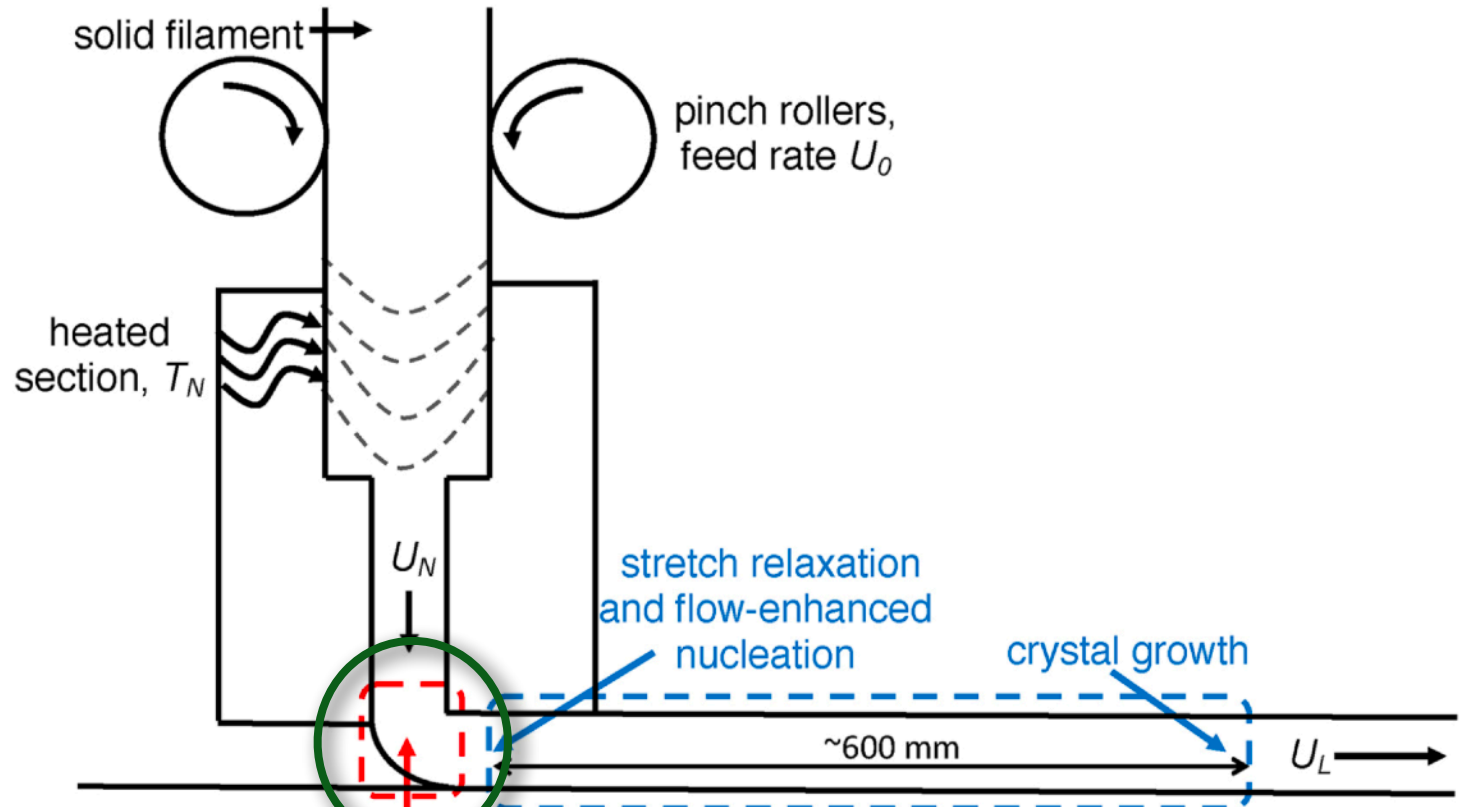


# Modelling crystallisation in 3D-printing



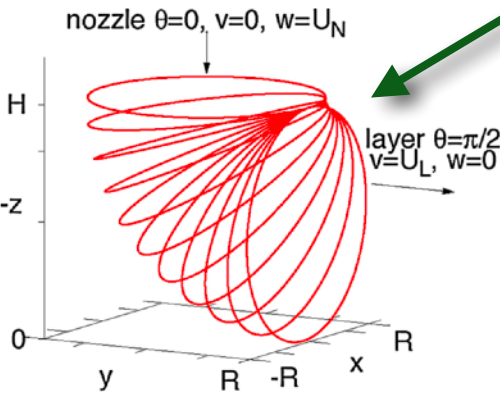
McIlroy and Graham, Additive Manufacturing 24, 323 (2018).

# Modelling crystallisation in 3D-printing

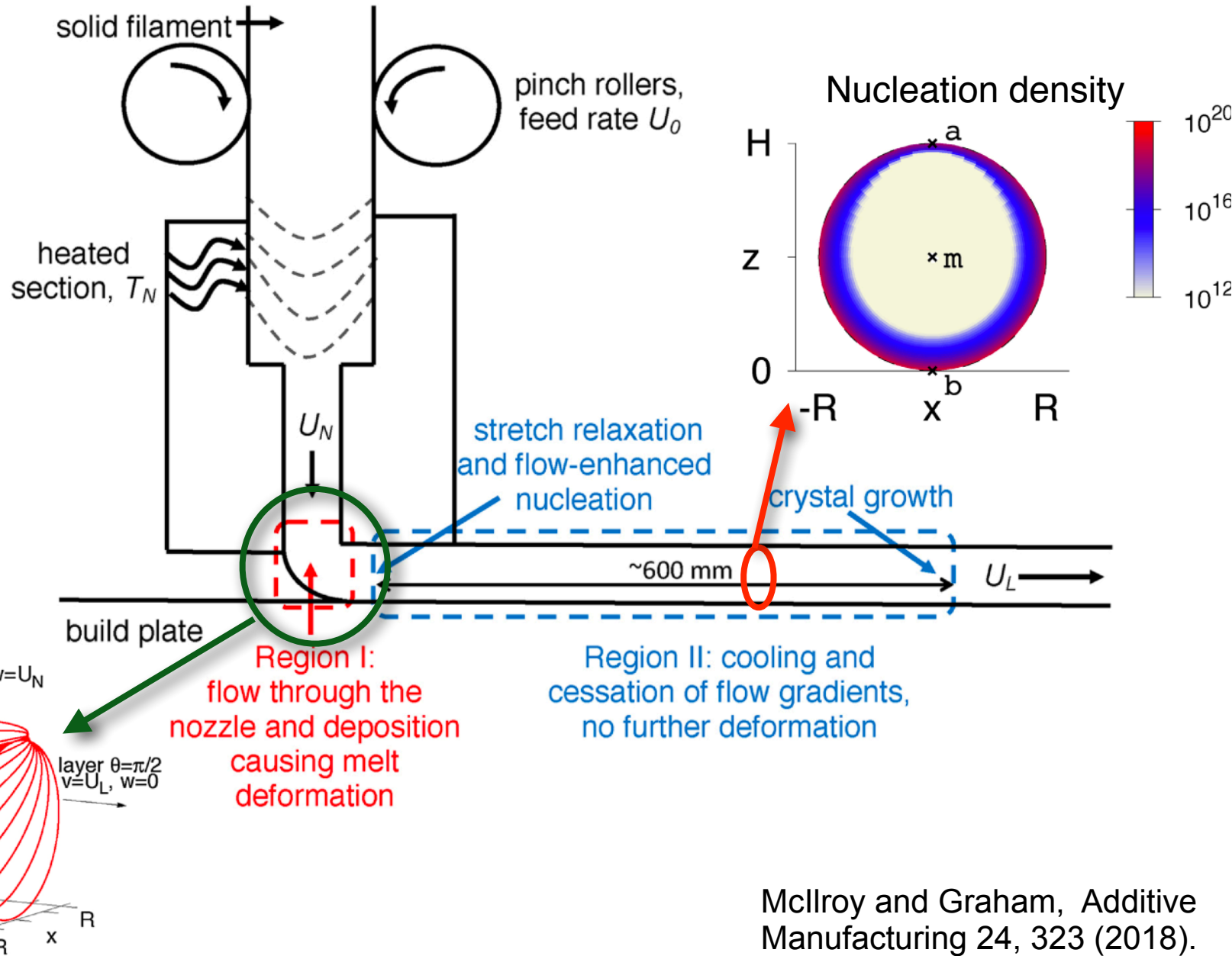


**Region I:**  
 flow through the nozzle and deposition causing melt deformation

**Region II:** cooling and cessation of flow gradients, no further deformation

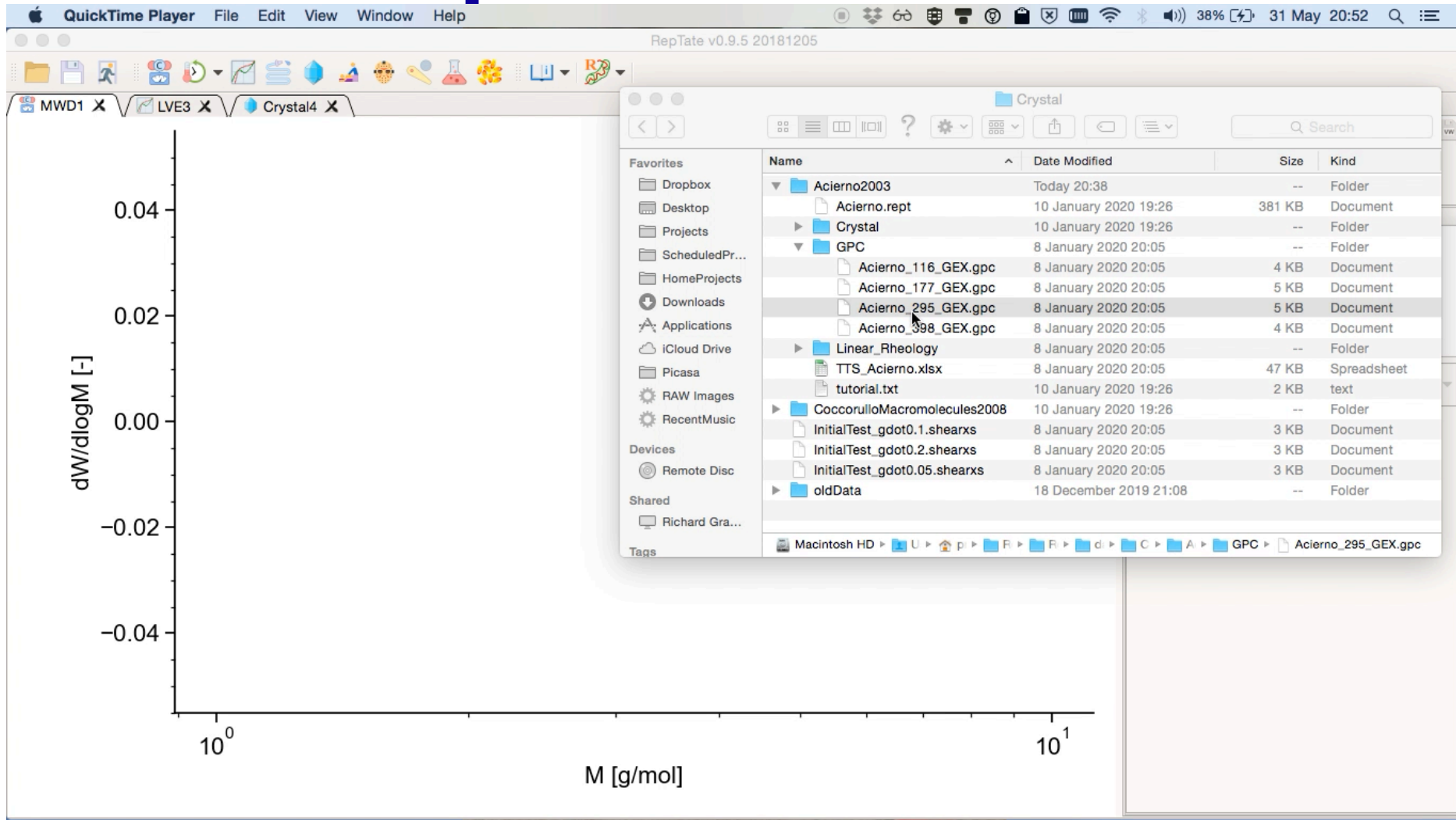


# Modelling crystallisation in 3D-printing



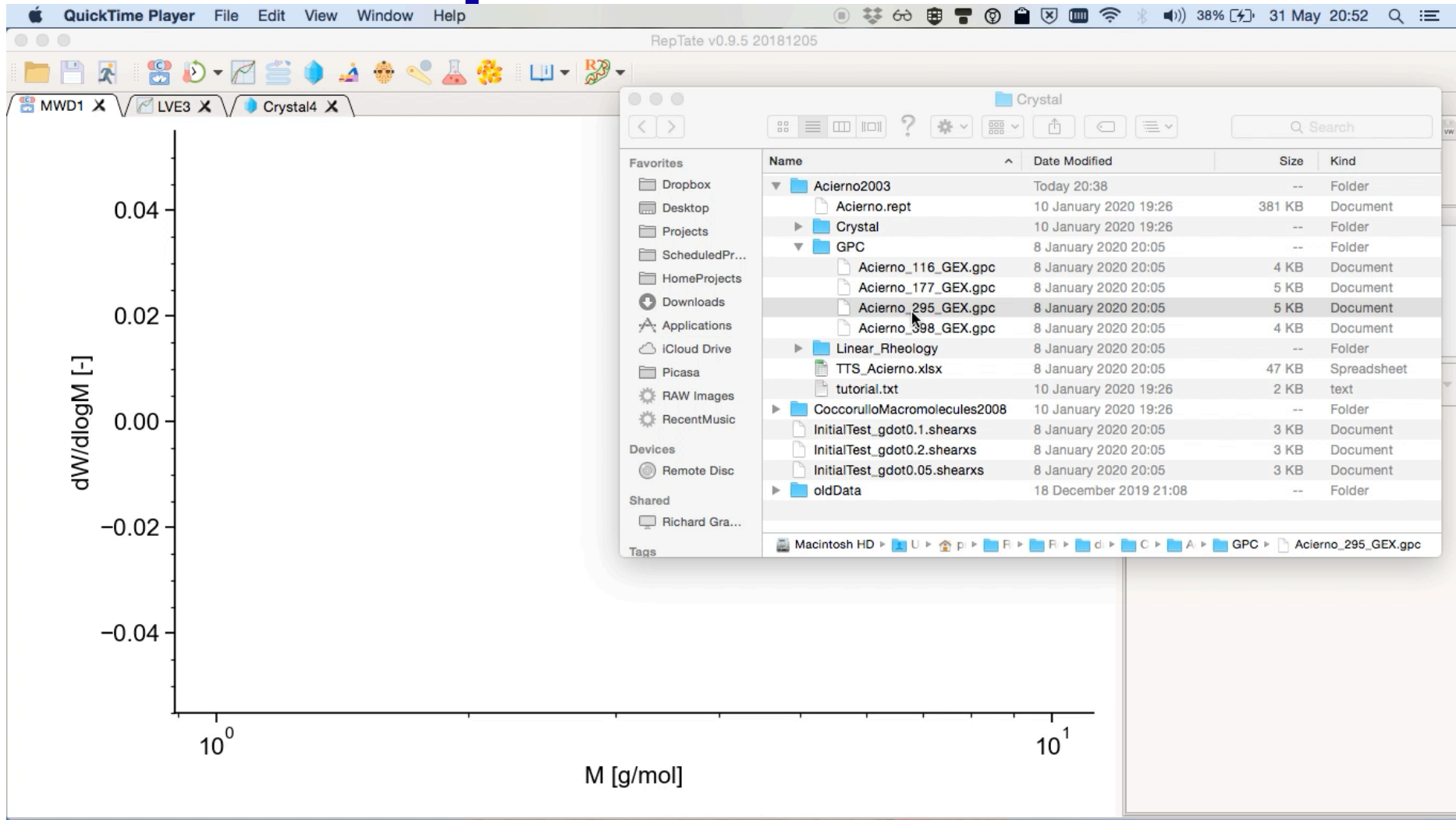
McIlroy and Graham, Additive Manufacturing 24, 323 (2018).

# RepTate software



Freely available from: <http://reptate.readthedocs.io/>

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# Summary

- **Flow-induced crystallisation:** In polymers flow can very strongly enhance the rate of crystallisation.
- **Modelling is challenging:** due to very wide spread of length and timescales
- **Multi-scale modelling:** MD and kinetic Monte Carlo simulations used to systematically derive a continuum model of nucleation.
- **Continuum model:** Nearly analytic expression for the nucleation rate; quantitatively consistent with MD and the GO model
- **Agreement with experiments:** Agrees with experiments. Correctly captures variation with flow rate, temperature and molecular weight.
- **Enhancement of long chains:** Model predicts long chains are over-presented in nuclei of crystallisation; confirmed in MD simulations; experimental signature identified.



