

Phase separation of DNA-bound proteins

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Phase separation of polymer-bound particles, David et al., arXiv:1811.09234

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Par**ABS** partition system

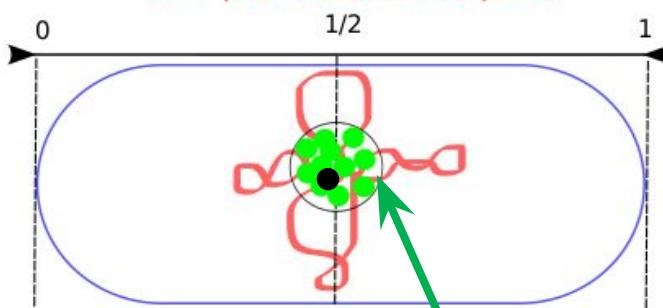
ParABS partition system and bacterial DNA segregation

Replication

Segregation

Division

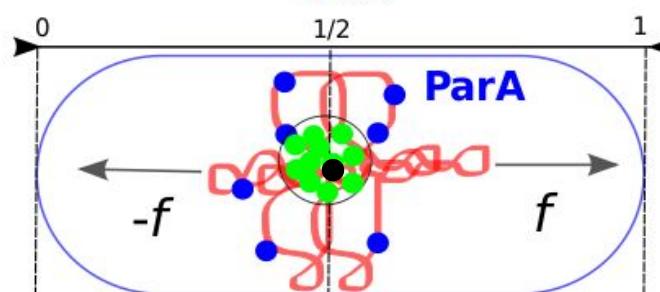
Step 1. Formation of the partition complex



- ParA
- ParB
- parS

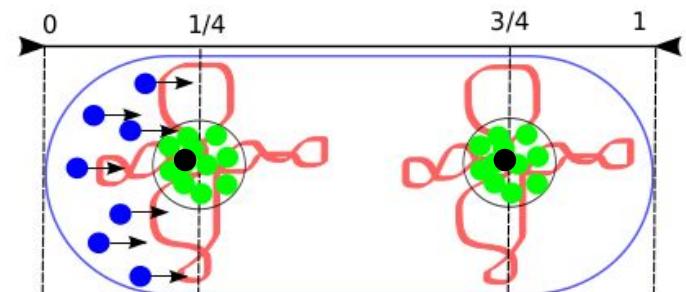
ParBS
complexes

Step 2. Separation of the copies of DNA



opposite forces

Step 3. Positioning



oscillations of ParA

Source: Jean-Charles Walter

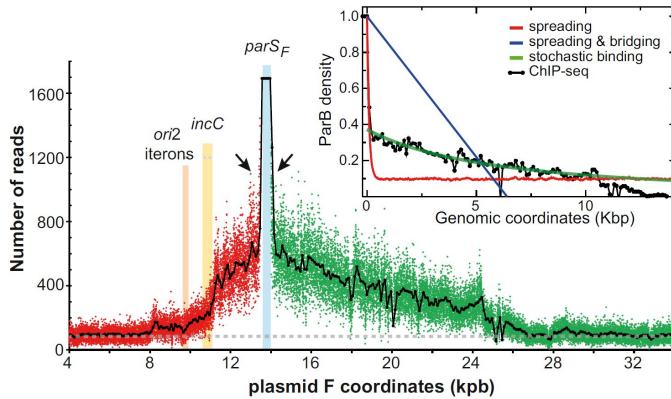
Overview

Overview of ParABS modeling (non-exhaustive)

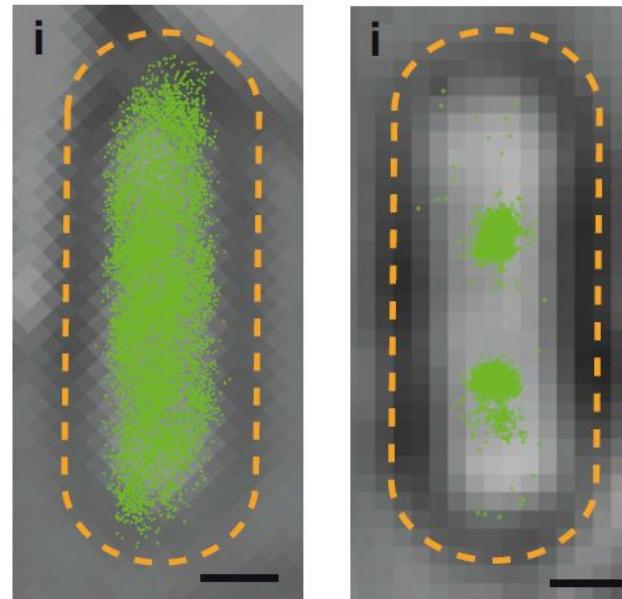
Experimental data /
Phenomena

ParBS complexes formation

Sanchez et al., Cell Syst 2015,
Debaugny et al., Mol Syst Biol 2018

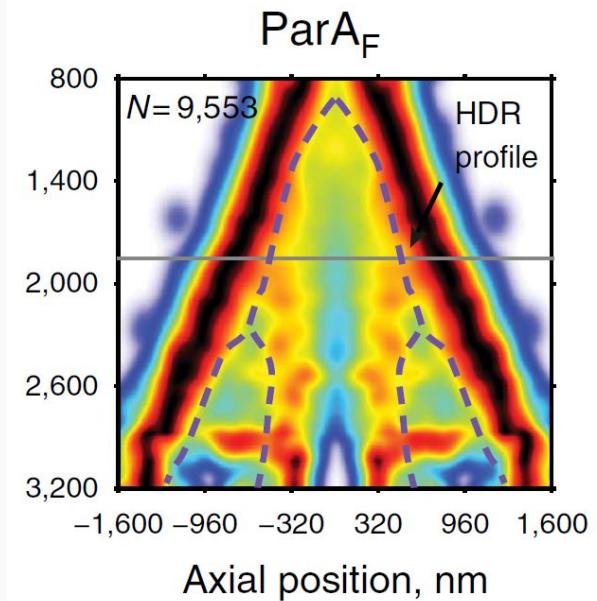


Sanchez et al., Cell Syst 2015



Positioning mechanism

Le Gall et al., Nat Com 2016



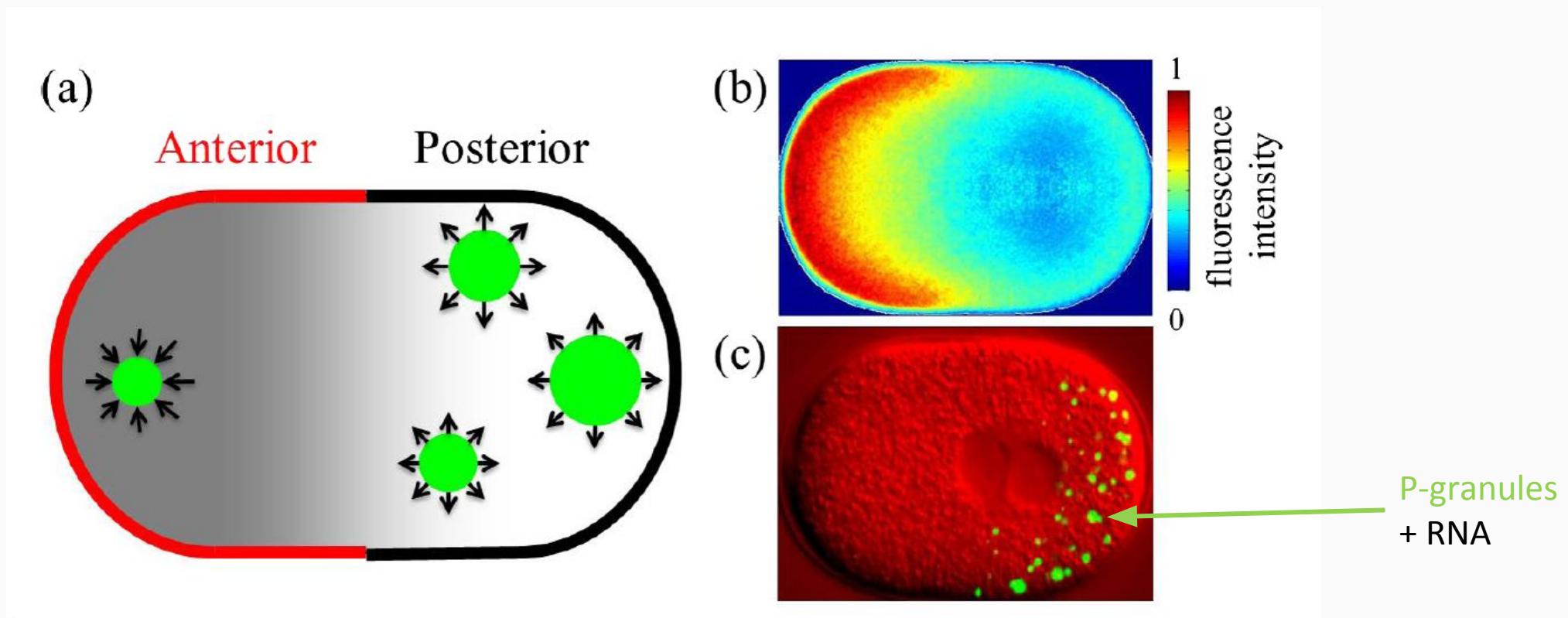
- Spreading and/or Bridging
(Broedersz et al., PNAS 2014)
- Stochastic Binding
(Sanchez et al. Cell Syst 2015)
 - Looping and Clustering
(Walter et al., NJP 2018)

- Spreading and/or Bridging
(Broedersz et al., PNAS 2014)
- Long Range Lattice Gas
(David et al., re-submitted PRL)

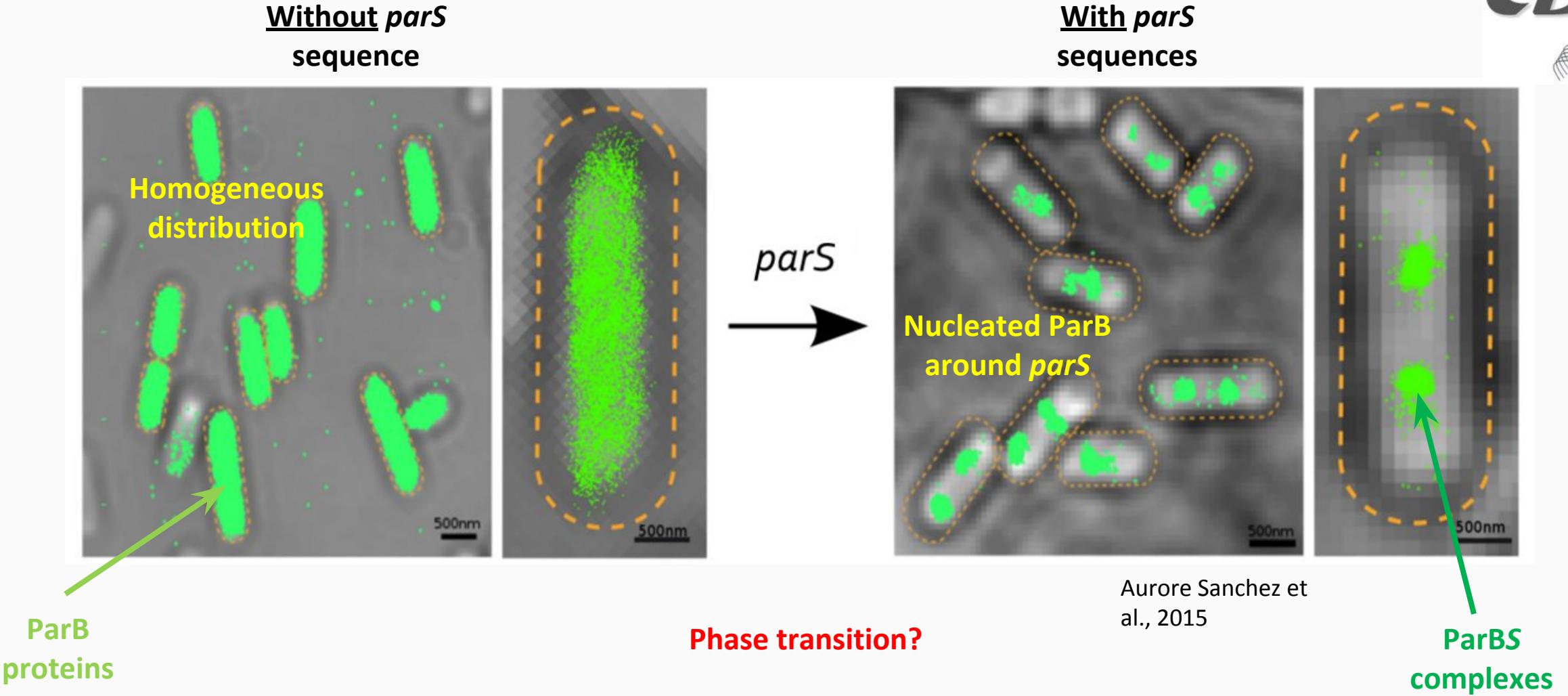
- Proteophoresis force model
(Walter et al., PRL 2017)

Phase separation in biological systems

Phase Separation → Non-membrane compartments → Confine and organize chemical reactions in intracellular space



ParBS complexes formation: phase separation?



1. Are phase transitions even possible?
2. Where are we in the phase diagram?

DNA + interacting DNA-bound particles = Spreading and Bridging

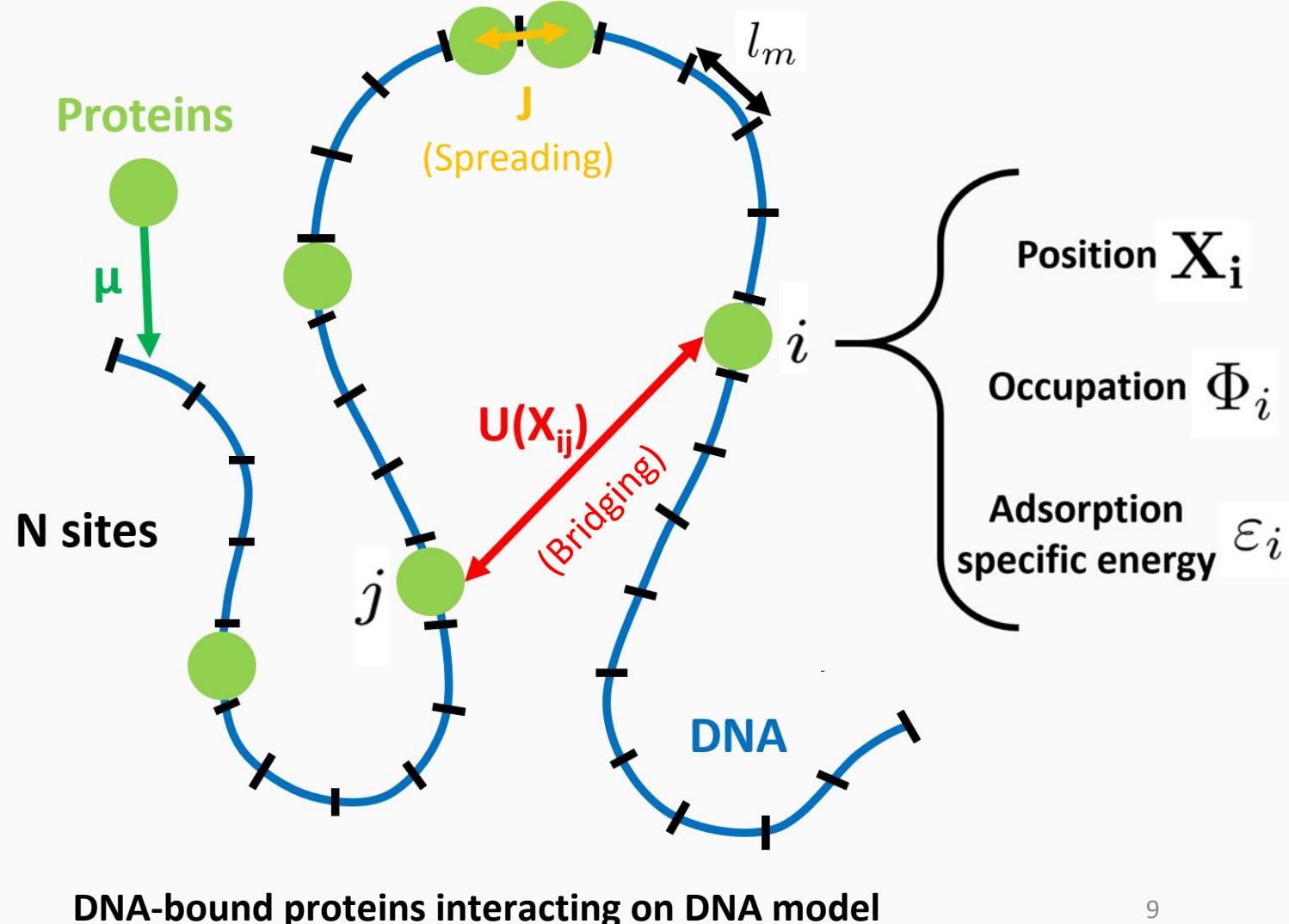
3D DNA-polymer spatial distribution



1D protein occupation distribution

Broedersz et al., PNAS 2014

Spreading & Bridging Model



The S'n'B is an Hamiltonian approach

**Short Range Lattice Gas
(Spreading)**

$$H_{\text{SRLG}}[\Phi_i] = -J \sum_{i=1}^{N-1} \Phi_{i+1}\Phi_i - \sum_{i=1}^N (\mu - \varepsilon_i) \Phi_i$$

$$H[\Phi_i, \mathbf{X}_i] = H_P[\mathbf{X}_i] + H_{\text{SRLG}}[\Phi_i] + H_B[\Phi_i, \mathbf{X}_i]$$

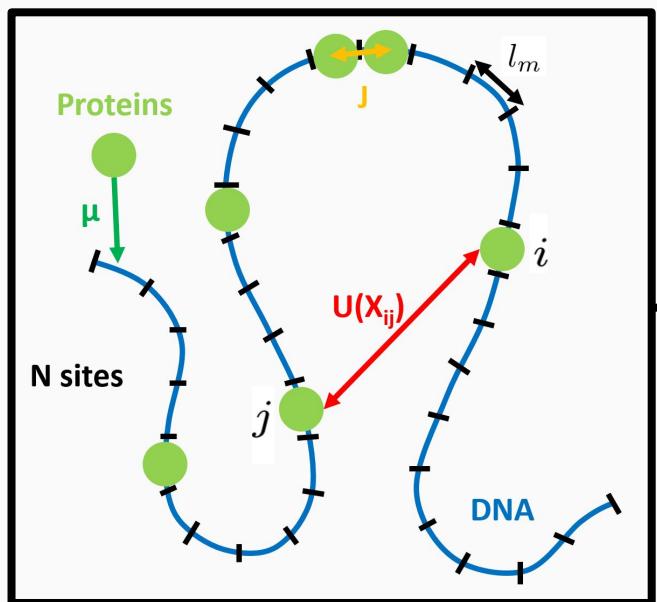
3D Polymer

**Coupling between
polymer and lattice gas
(Bridging)**

$$H_B[\Phi_i, \mathbf{X}_i] = \frac{1}{2} \sum_{i,j}^N \Phi_i U(X_{ij}) \Phi_j$$

From the S'n'B to the Long Range Lattice Gas model

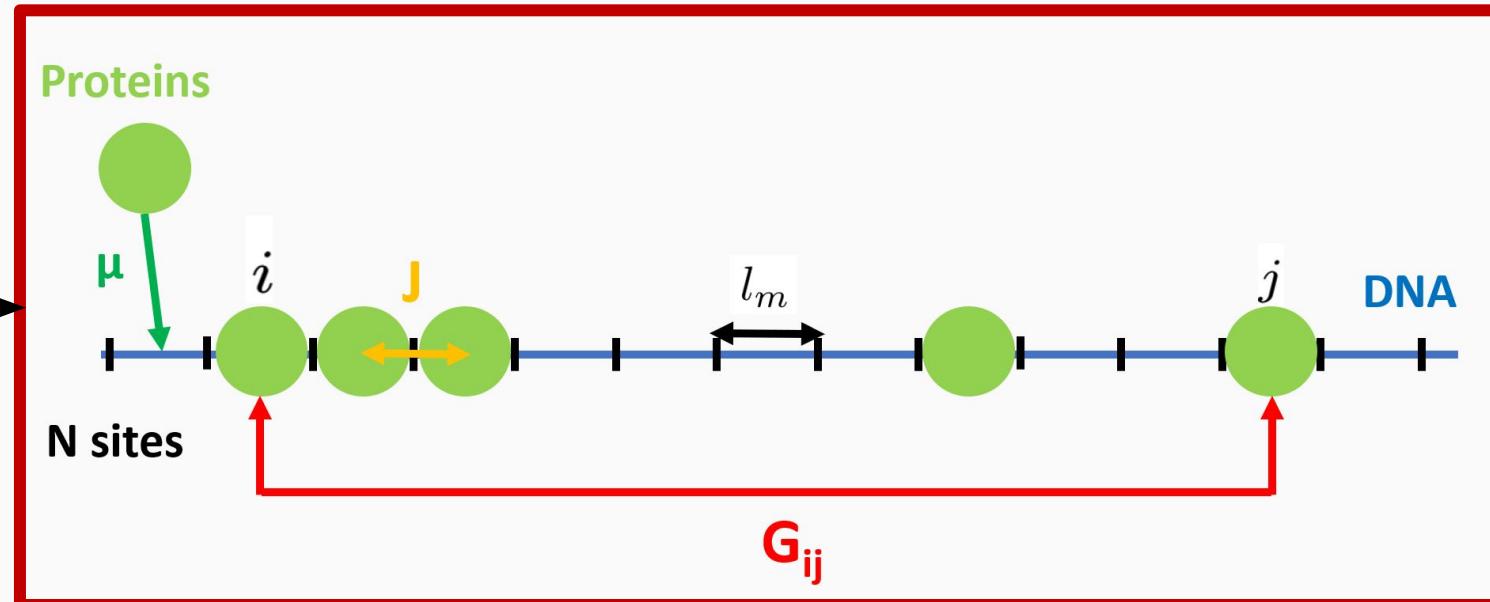
Spreading & Bridging



Cluster expansion,
two-body
interactions only
(low density)

3D → 1D

Long Range Lattice Gas



$$\mathcal{F}_{\text{LRLG}}[\Phi_i] = H_{\text{SRLG}}[\Phi_i] - \frac{1}{2} \sum_{i,j}^N \Phi_i G_{ij} \Phi_j$$

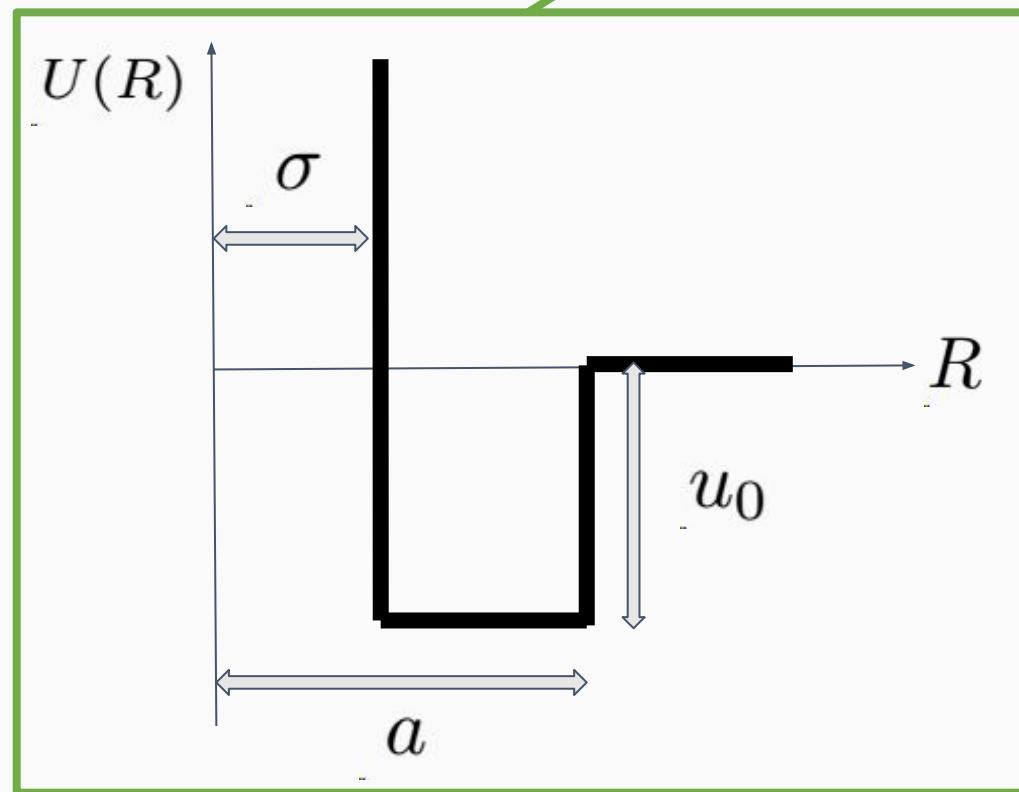
Definition of the kernel G_{ij}

2nd order virial
expansion
coefficient

$$G_{ij} = 4\pi\beta^{-1} \int_0^{+\infty} dR R^2 \left[e^{-\beta U(R)} - 1 \right] P_{ij}(R)$$

Spatial potential

Short range
hard core +
square well



Polymer
distribution
function \Leftrightarrow Chain
connectivity !

Criterion for phase transitions/separation

Asymptotic behavior

$$G_{ij} \underset{|i-j| \rightarrow +\infty}{\longrightarrow} K_{SW} |i - j|^{-\alpha}$$

$$\alpha = (3 + g)\nu$$

Enhancement exponent
of the polymer partition
function

$$\mathcal{Z}_P \sim \bar{\mu}^N N^{\nu g}$$

Polymer size
exponent

$$R_{ij} \sim |i - j|^\nu$$

PDF asymptotic
behavior

$$P_{ij}(R) \sim \frac{R^g}{R_{ij}^{3+g}}$$

Phase transition only if $1 < \alpha < 2$

Freeman J. Dyson, 1969



Gaussian Polymer:
 $\alpha_{GP} = 1.5$

Self-Avoiding Polymer:
 $\alpha_{SAP} \approx 1.92$

1. Are phase transitions even possible?

2. Where are we in the phase diagram?

Phase separation



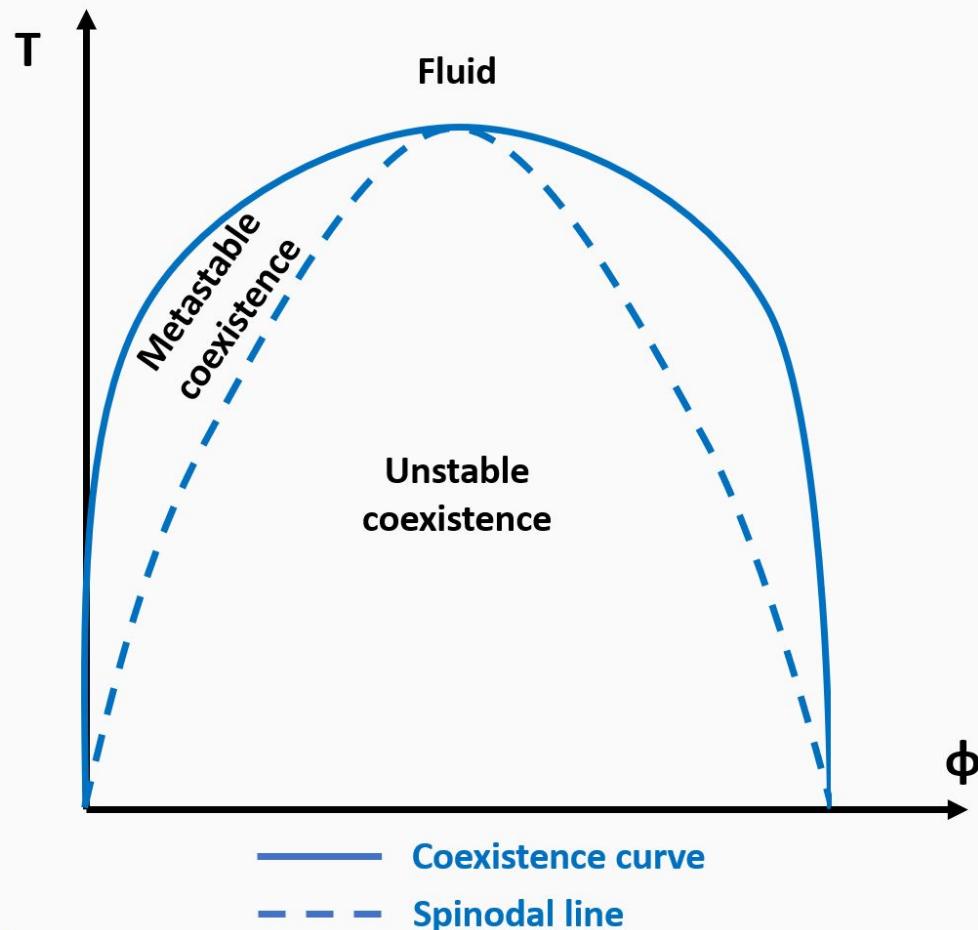
Coexistence of high ϕ_h and low occupation ϕ_l states

- Coexistence curve:

$$\begin{cases} P(T, \Phi_l) = P(T, \Phi_h) \\ \mu(T, \Phi_l) = \mu(T, \Phi_h) \end{cases}$$

- Spinodal line:

$$\left(\frac{\partial P}{\partial \Phi} \right)_T = 0$$



Variational method

$$\mathcal{F}_{\text{LRLG}}[\Phi_i] = H_0 + \Delta H \longrightarrow \Omega_{\text{LRLG}} \leq \Omega_V = \Omega_0 + \langle \Delta H \rangle_0$$

Hamiltonian of reference = 1D SRLG

$$\left\{ \begin{array}{l} H_0 = -J \sum_{i=1}^{N-1} \Phi_{i+1} \Phi_i - \mu_0 \sum_{i=1}^N \Phi_i \\ \Delta H = -\frac{1}{2} \sum_{i,j}^N \Phi_i G_{ij} \Phi_j - (\mu - \mu_0) \sum_{i=1}^N \Phi_i \end{array} \right.$$

Minimization

$$\left(\frac{\partial \Omega_V}{\partial \mu_0} \right)_{\mu_0=\mu_0^*}$$

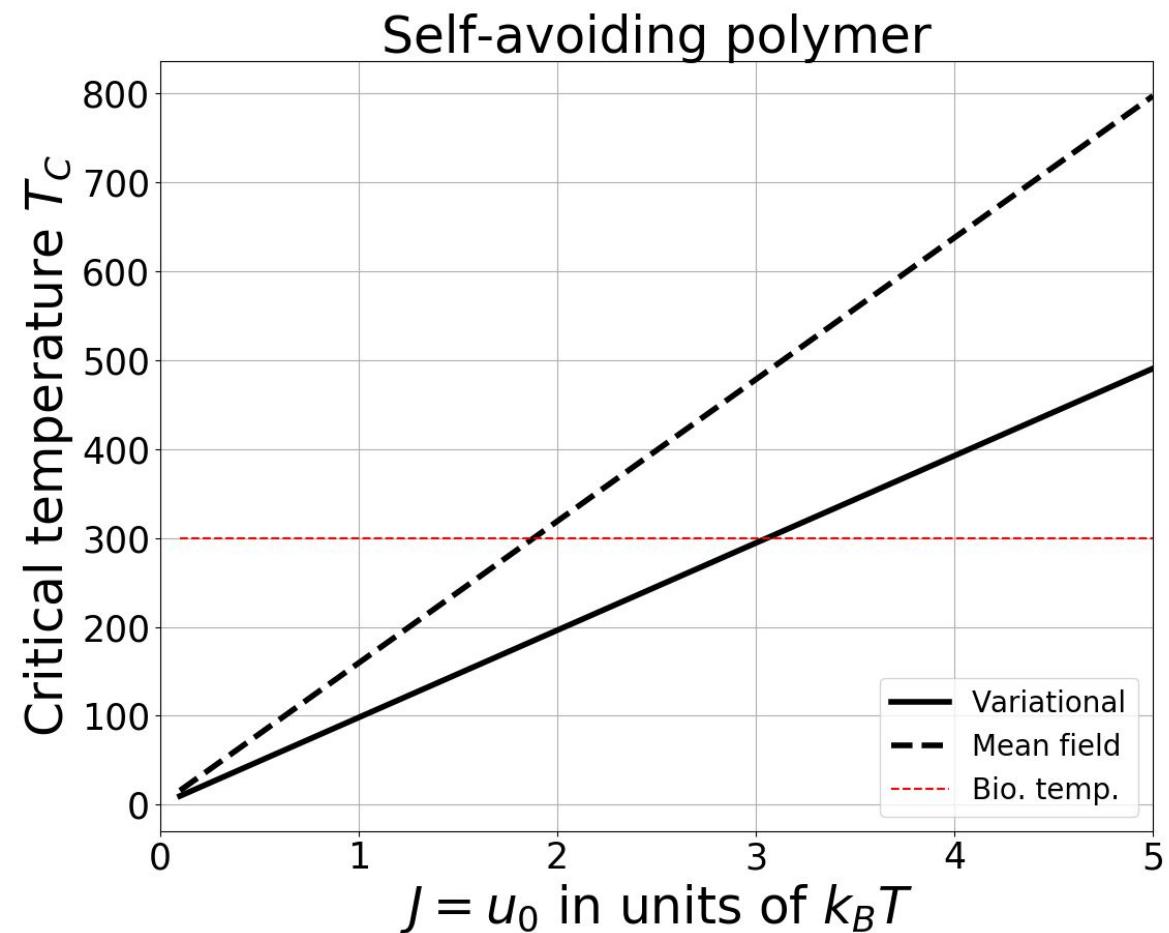
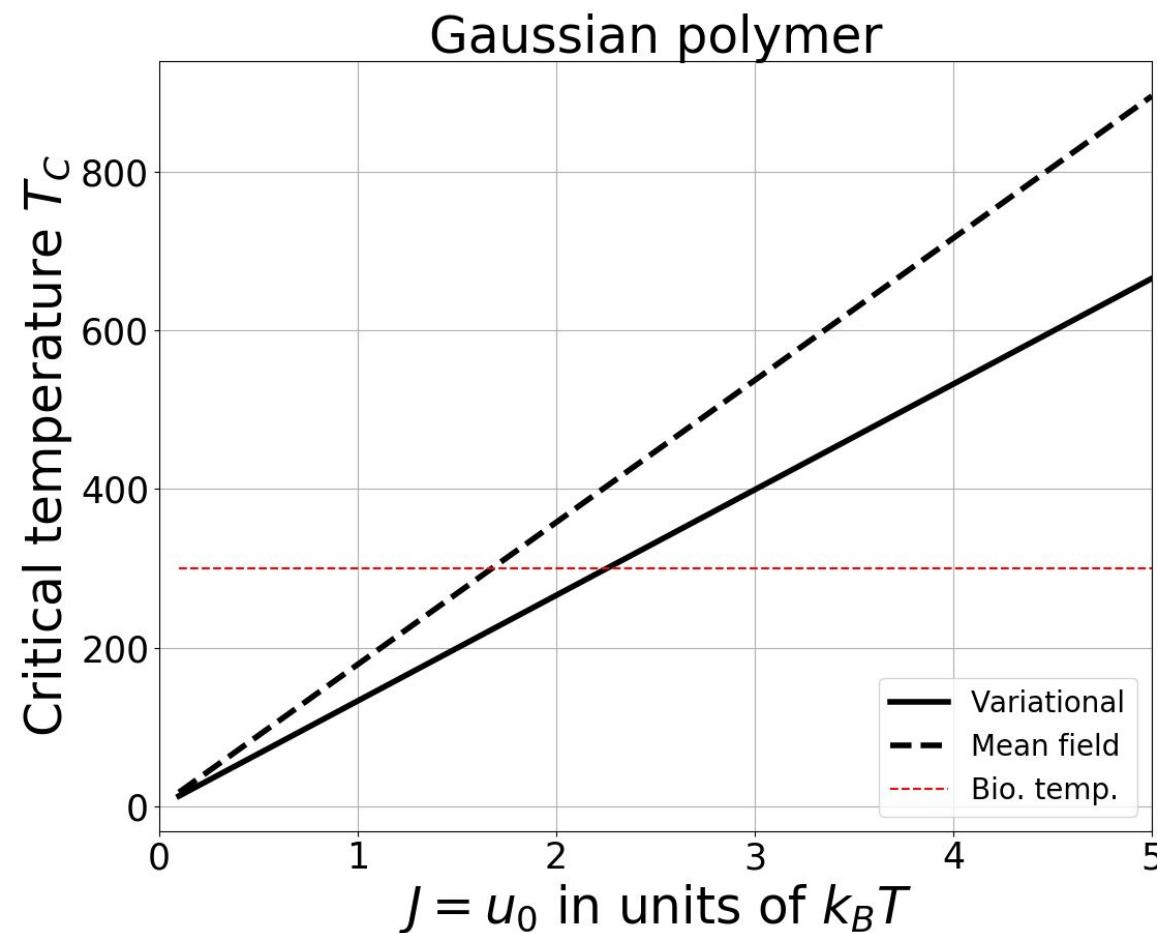
$$\Omega_{\text{LRLG}} \approx \Omega_V^*$$

Analytical results !

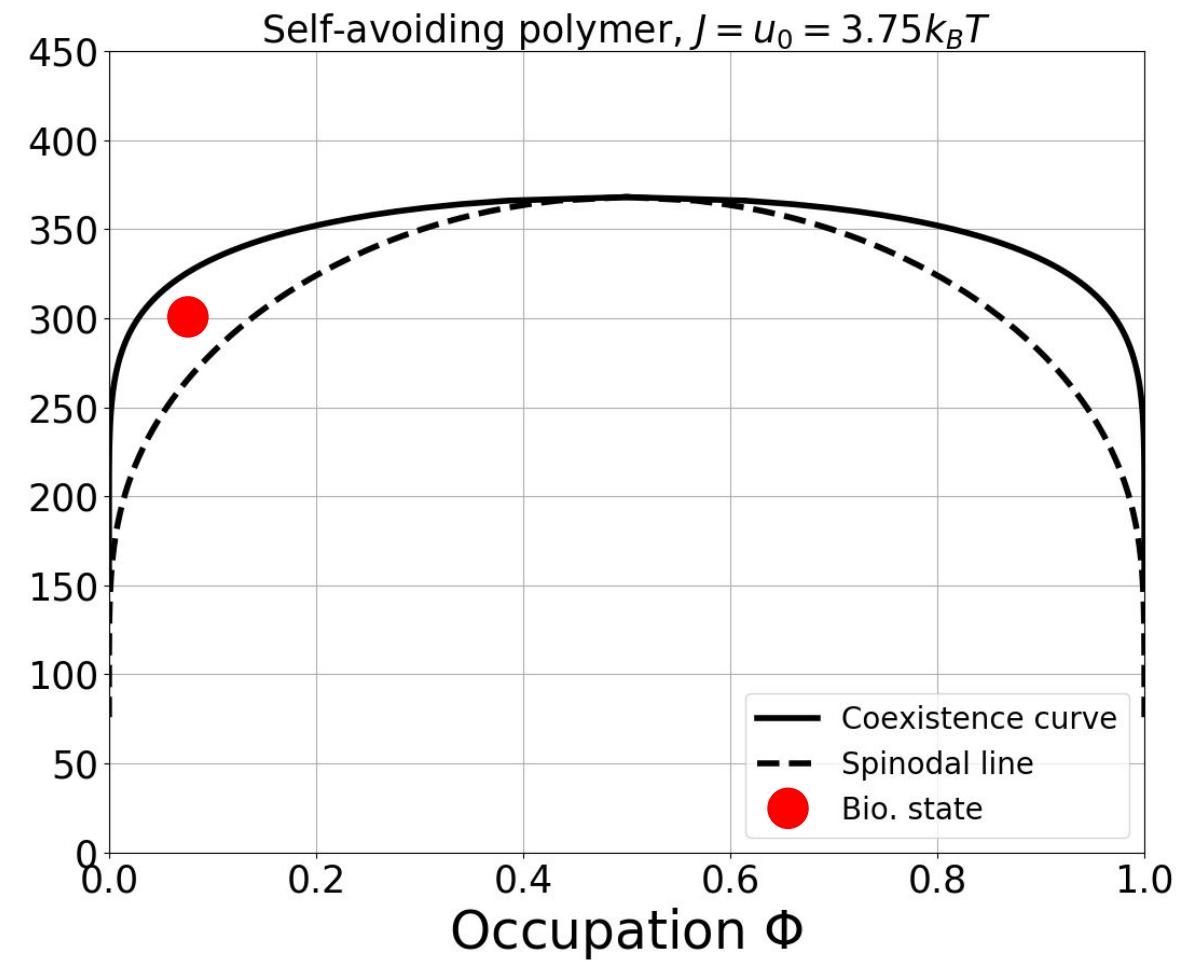
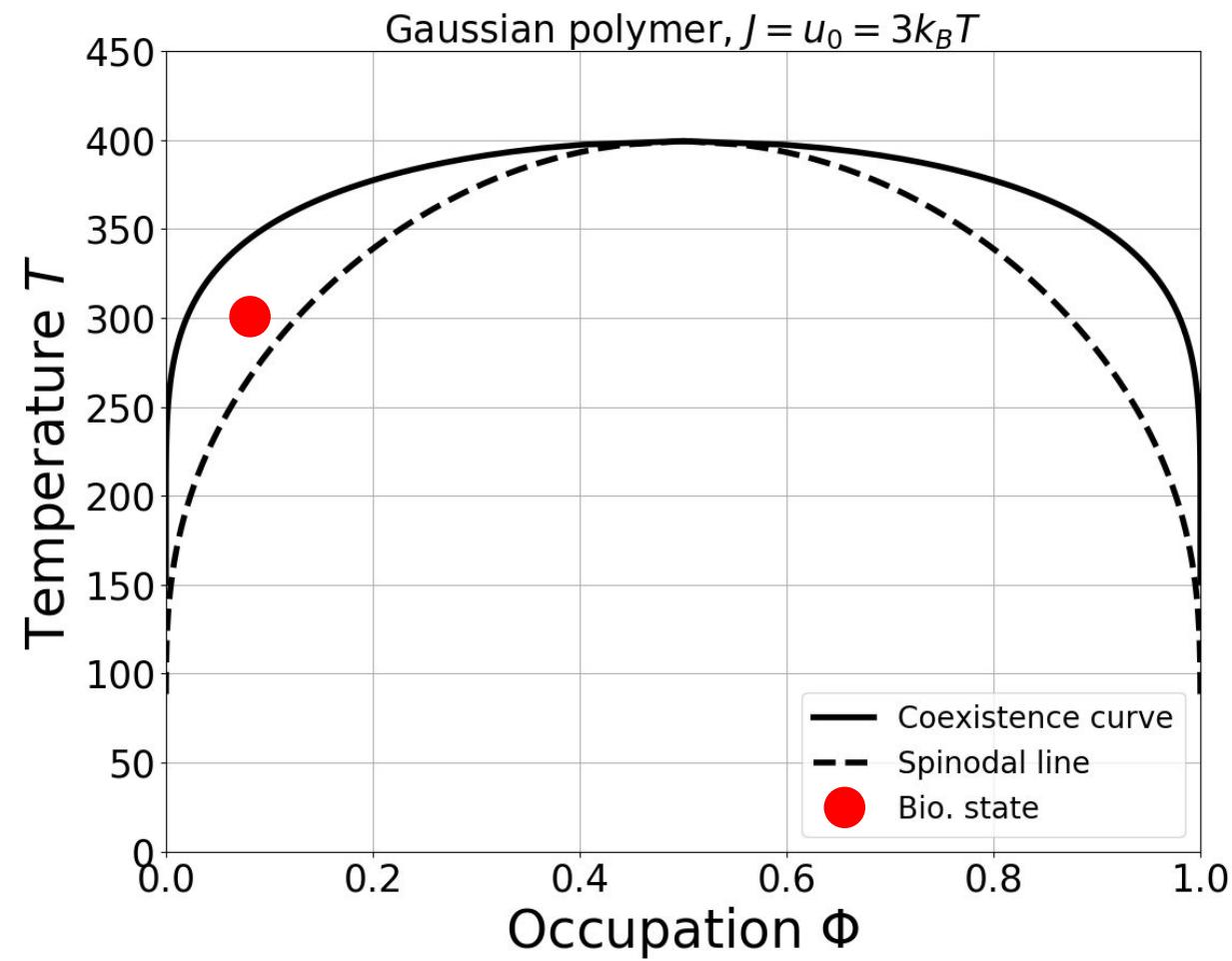
Parameters overview

	Parameter	Symbol	Value
Polymer parameters	Physiol. Temperature	T_{bio}	300 K
	Monomer length	l_m	16 bp (~ 5.4 nm)
	DNA persistence length	l_p	150 bp (~ 50 nm)
	Kuhn length	b	23.6 nm
Spatial potential $U(R)$	PDF LR behavior exponent	g	SAP: 0.27, G: 0
	Flory exponent	ν	SAP: 0.588, G: 0.5
	Hard-core int. range	σ	l_m
	Spatial int. range	a	2σ (~ 10.8 nm)
Work parameters	Spatial int. intensity	u_0	$u_0 = J$
	Spreading intensity	J	0 to $5k_B T$
	Mean occupation	Φ	0 to 1

1D Long Range Lattice Gas: critical temperature



1D Long Range Lattice Gas: phase diagrams

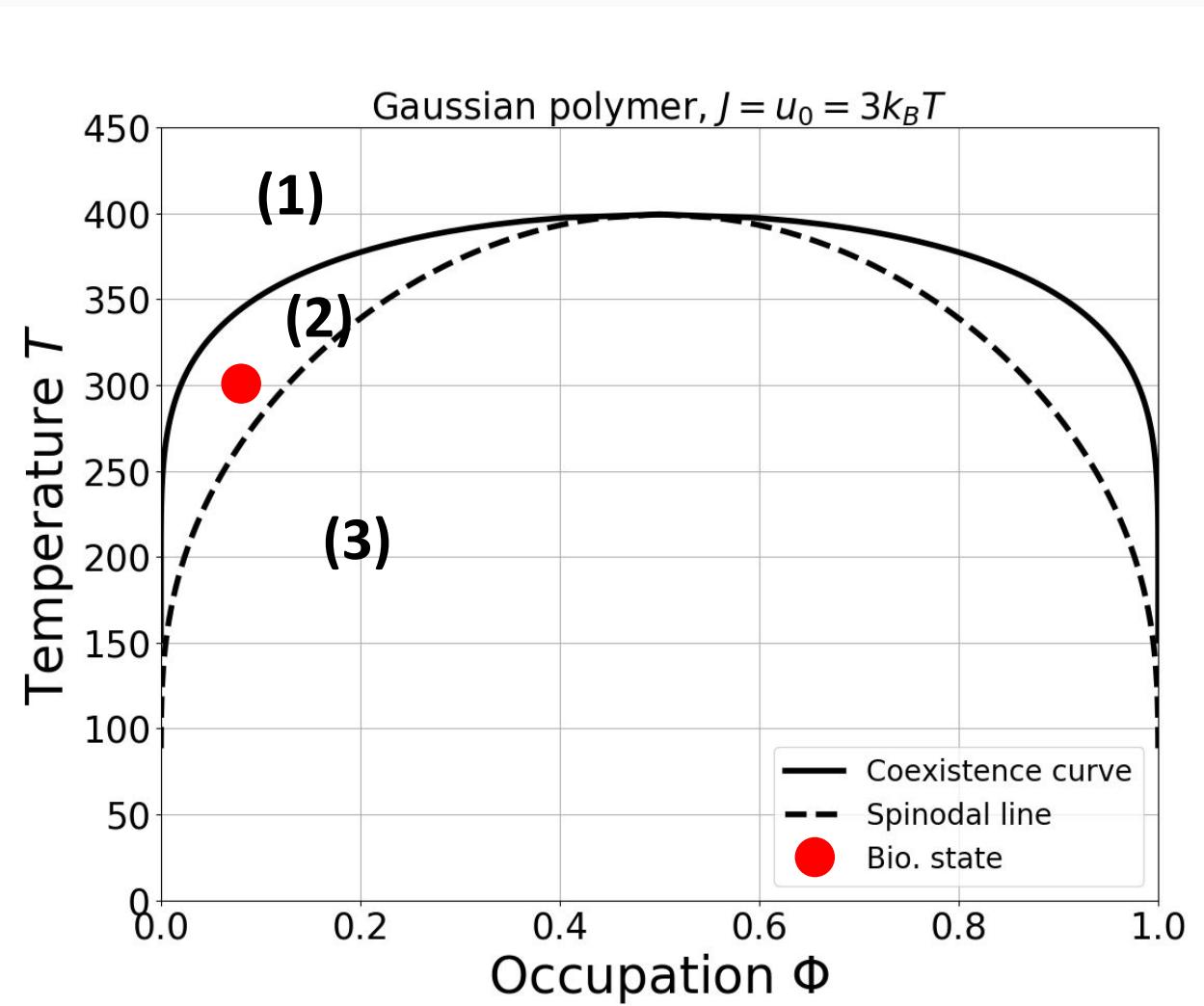


Biological regime for F-plasmid:



$\Phi = 300 \text{ ParB} / 3750 \text{ sites} \approx 0.08 \text{ (upper bound)}$
 $T = 300 \text{ K}$

Low occupation metastable regime

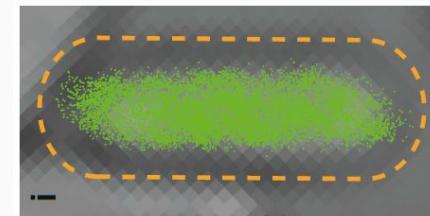


Expected ParB distribution...

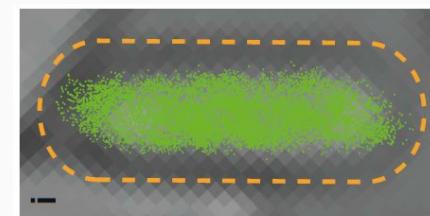
...without *parS*

...with *parS*

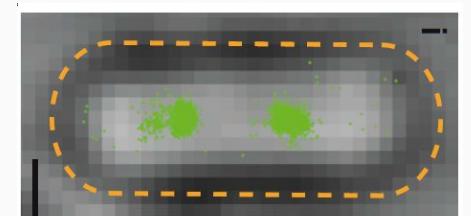
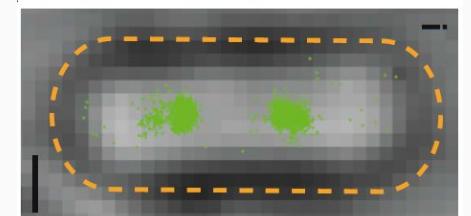
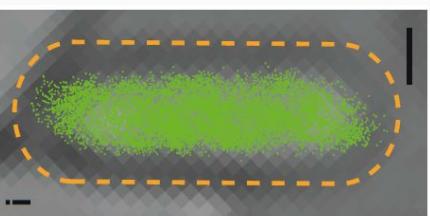
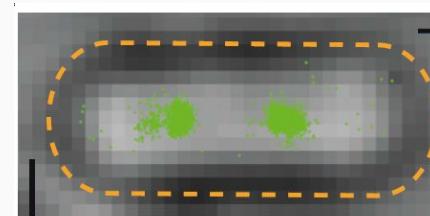
(1)



(2)



(3)



In short

Sum up and perspectives

1. Polymer fluctuations \Rightarrow effective long-range interactions between bound-particles \Rightarrow phase separation of bound-proteins on the polymer.
 2. Variational method \Rightarrow whole mean-occupation/temperature phase diagram.
 3. The low occupation metastable coexistence region explains the experimental data for ParB distribution.
-
- What is the consequence for the polymer distribution?
 - Occupation/density profile?
 - Quantitative test? Accurate prediction of J or u_o ?

Team and collaborators in France

Physical modeling (Montpellier)

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