

# Lecture 6: Proteins; entropy rules

Goal: Introduce Boltzmann distribution, probability of microstate

- Ligand-receptor binding
- Gene regulation
- Cooperativity

PBOC Chapter 6.1.1, 6.1.2, 6.4  
(except 6.4.4)

# Statistical mechanics for biophysics

Numbers:

RNA polymerase molecules in the nucleus

Ligands near cell surface

What can we answer with these models?

Determine the probability of finding the system in a particular (energy) state.

Calculate the average values of observables.

# Statistical mechanics for biophysics

Previously (Lecture 3):

## *Microstates*

Definition: a **microstate** is a microscopic arrangement of the constituents of a system

Example: Ligand binding to a receptor protein

### Lattice model

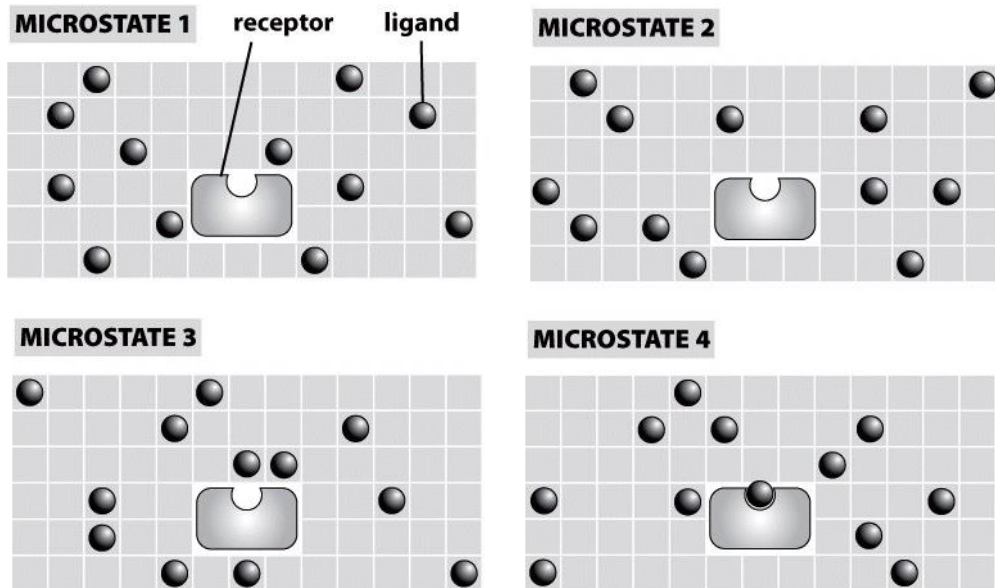
$L$  ligands

$\Omega$  boxes

max. one ligand per box

energy  $\varepsilon_b$  of a bound ligand

energy  $\varepsilon_{sol}$  of a ligand in solution



# Statistical mechanics for biophysics

## *Microstates*

Suppose a system can exist in states with energies  $E_i$ .

What is the probability of finding the system in a given state?

Boltzmann distribution, probability of finding the system in a microstate with energy  $E_i$  (*derivation, Section 6.1.3*)

$$p(E_i) = \frac{1}{Z} e^{-E_i/k_B T}$$

Partition function, normalization factor so that  $\sum_{i=1}^N p(E_i) = 1$

$$Z = \sum_{i=1}^N e^{-E_i/k_B T}$$

# Statistical mechanics for biophysics

## *Microstates*

Suppose a system can exist in states with energies  $E_i$ .  
What is the average energy of the system?

The average energy is the  
(probability) weighted mean of the  
energies of the states:

$$\begin{aligned}\langle E \rangle &= \sum_{i=1}^N E_i p(E_i) = \frac{1}{Z} \sum_{i=1}^N E_i e^{-\beta E_i} \\ &= -\frac{1}{Z} \frac{\partial Z}{\partial \beta} = -\frac{\partial}{\partial \beta} (\ln Z)\end{aligned}$$

# Statistical mechanics for biophysics

Example: Ligand binding

**Lattice model**

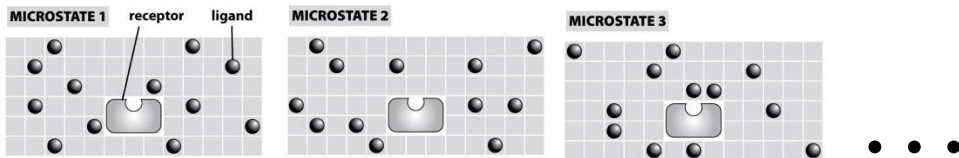
$L$  ligands

$\Omega$  boxes

max. one ligand per box

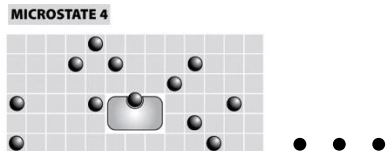
*Microstates: multiplicity* (Lecture 3)

Microstates with receptor unoccupied



$$\text{number of microstates} = \frac{\Omega!}{L!(\Omega-L)!}$$

Microstates with receptor occupied

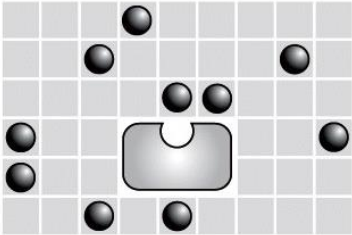
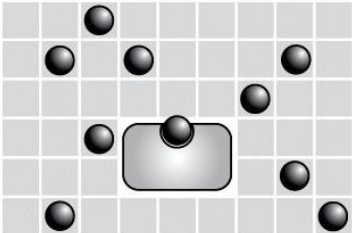


number of microstates?

# Statistical mechanics for biophysics

Example: Ligand binding to receptor protein

*Microstates: energy, weight*

	STATE	ENERGY	MULTIPLICITY	WEIGHT
receptor unbound		$L\varepsilon_{sol}$	$\frac{\Omega!}{L!(\Omega-L)!} \approx \frac{\Omega^L}{L!}$ for $\Omega \gg L$	<p>multiplicity x Boltzmann weight "partial" partition function</p> $\frac{\Omega^L}{L!} e^{-\beta L\varepsilon_{sol}}$
receptor bound		$(L-1)\varepsilon_{sol} + \varepsilon_b$	$\frac{\Omega!}{(L-1)!(\Omega-L+1)!} \approx \frac{\Omega^{L-1}}{(L-1)!}$	$\frac{\Omega^{L-1}}{(L-1)!} e^{-\beta[(L-1)\varepsilon_{sol} + \varepsilon_b]}$

# Statistical mechanics for biophysics

Example: Ligand binding to  
receptor protein

*Microstates: probability*

$$P_{\text{bound}} = \frac{\sum_{\text{states}} \left( \begin{array}{c} \text{diagram of bound state} \end{array} \right)}{\sum_{\text{states}} \left( \begin{array}{c} \text{diagram of unbound state} \end{array} \right) + \sum_{\text{states}} \left( \begin{array}{c} \text{diagram of bound state} \end{array} \right)}$$

$$\begin{aligned} &= \frac{\frac{\Omega^{L-1}}{(L-1)!} e^{-\beta[(L-1)\varepsilon_{\text{sol}} + \varepsilon_b]}}{\frac{\Omega^L}{L!} e^{-\beta L \varepsilon_{\text{sol}}} + \frac{\Omega^{L-1}}{(L-1)!} e^{-\beta[(L-1)\varepsilon_{\text{sol}} + \varepsilon_b]}} \\ &= \frac{\frac{L}{\Omega} e^{-\beta[\varepsilon_b - \varepsilon_{\text{sol}}]}}{1 + \frac{L}{\Omega} e^{-\beta[\varepsilon_b - \varepsilon_{\text{sol}}]}} \end{aligned}$$

Define:  $\Delta\varepsilon = \varepsilon_b - \varepsilon_{\text{sol}}$

$$c(L) = L/V_{\text{box}}$$

$$c(L = \Omega) = c_0 = \Omega/V_{\text{box}}$$

$$= \frac{\frac{c}{c_0} e^{-\beta\Delta\varepsilon}}{1 + \frac{c}{c_0} e^{-\beta\Delta\varepsilon}}$$

Langmuir isotherm

Hill function of coefficient 1



# Statistical mechanics for biophysics

Example: Ligand binding to  
receptor protein

*Microstates: probability*

$$p_{\text{bound}} = \frac{\sum_{\text{states}} \left( \text{diagram of bound state} \right)}{\sum_{\text{states}} \left( \text{diagram of unbound state} \right) + \sum_{\text{states}} \left( \text{diagram of bound state} \right)} = \frac{\frac{c}{c_0} e^{-\beta \Delta \epsilon}}{1 + \frac{c}{c_0} e^{-\beta \Delta \epsilon}}$$

$$L + R \rightleftharpoons LR \quad K_d = \frac{[L][R]}{[LR]} \quad p_{\text{bound}} = \frac{[LR]}{[R] + [LR]}$$

$$p_{\text{bound}} = \frac{\frac{[L]}{K_d}}{1 + \frac{[L]}{K_d}}$$

# Statistical mechanics for biophysics

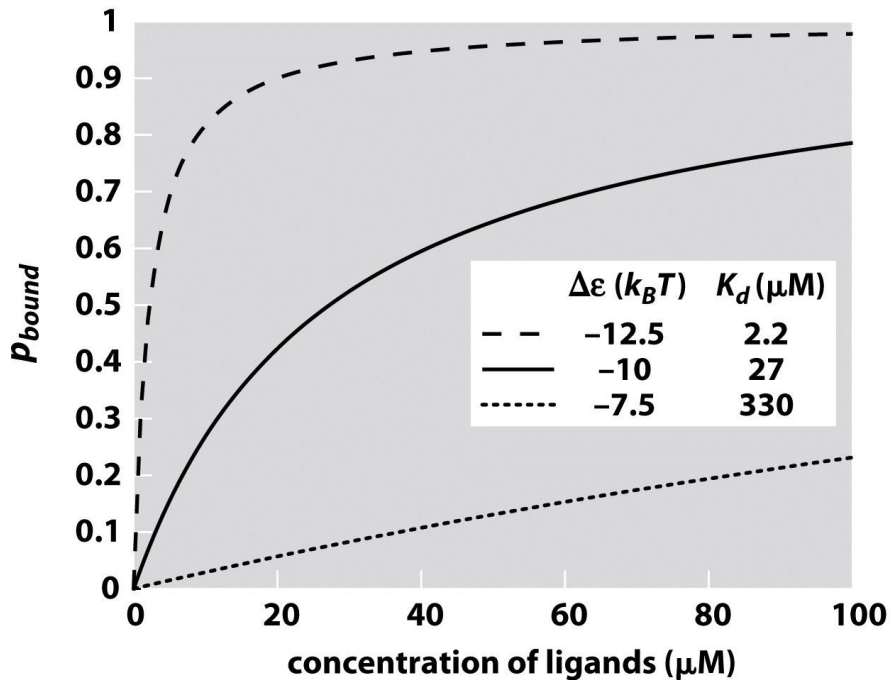
Example: Ligand binding to  
receptor protein

Estimate:  $V_{box} = 1 \text{ nm}^3$

$$\text{then } c_0 = \frac{\Omega}{V_{box}} = \left( \frac{6 \times 10^{23}}{10^{24}} \right)$$

$$= 0.6 \text{ M}$$

*Receptor occupancy*

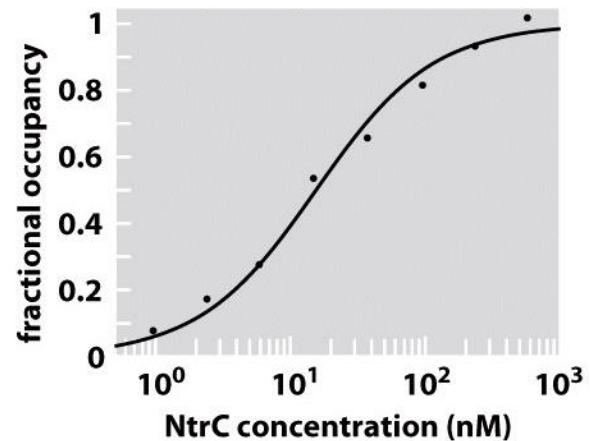
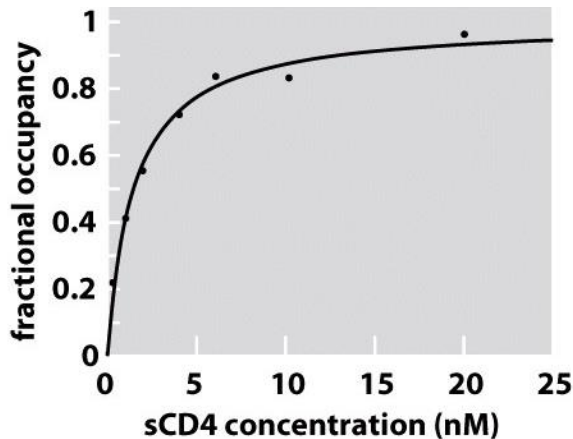
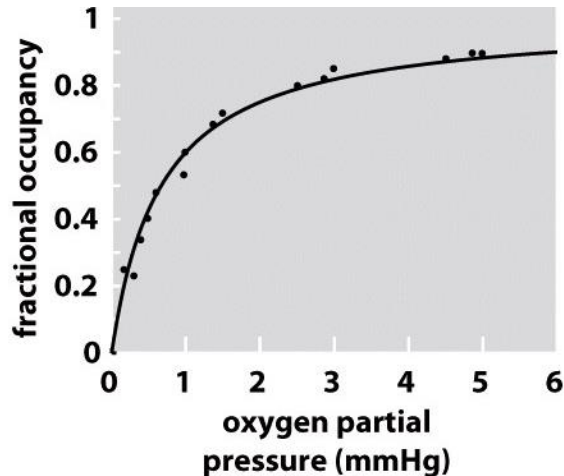


# Statistical mechanics for biophysics

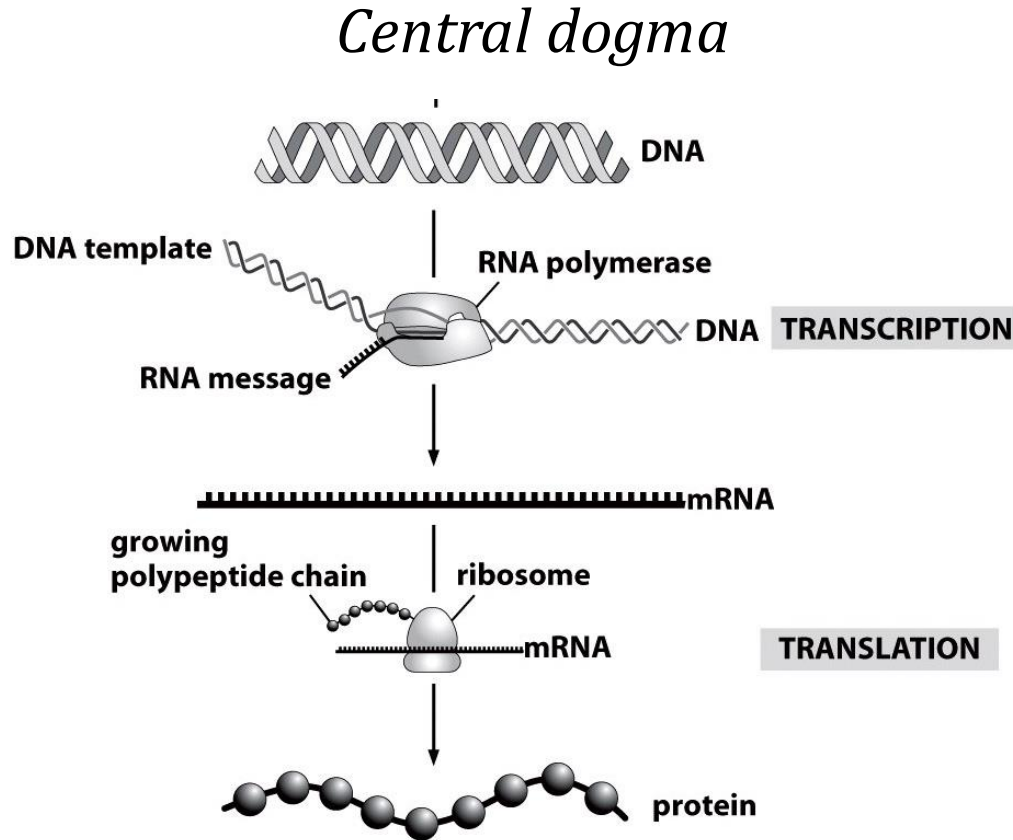
Example: Ligand binding to  
receptor protein

*Receptor occupancy*

Experimental data



# Statistical mechanics for biophysics



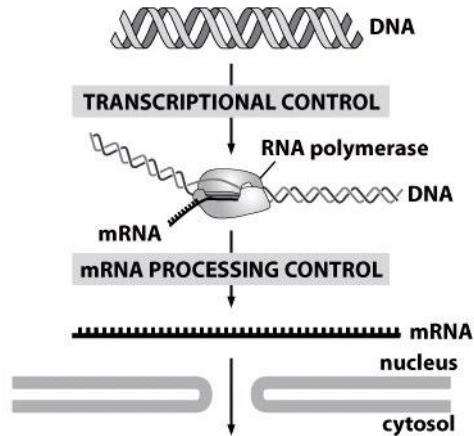
# Statistical mechanics for biophysics

*How do cells make decisions?*

*How can different cells in an organism  
maintain different protein concentrations?*

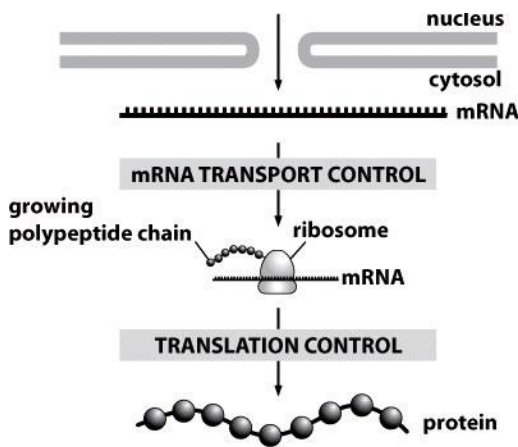
# Statistical mechanics for biophysics

Will the cell make a particular mRNA?



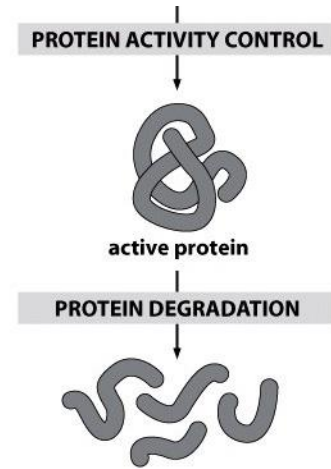
Will an mRNA be processed to become mature?

Will an mRNA be transported to the cytoplasm?



Will an mRNA be translated by the ribosome?

Will a protein exist in an active conformation?

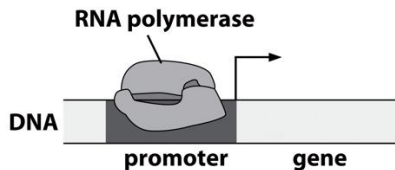


How long will a protein last before being degraded?

# Statistical mechanics for biophysics

Will the cell make a particular mRNA?

Example: Transcriptional control



## Lattice model

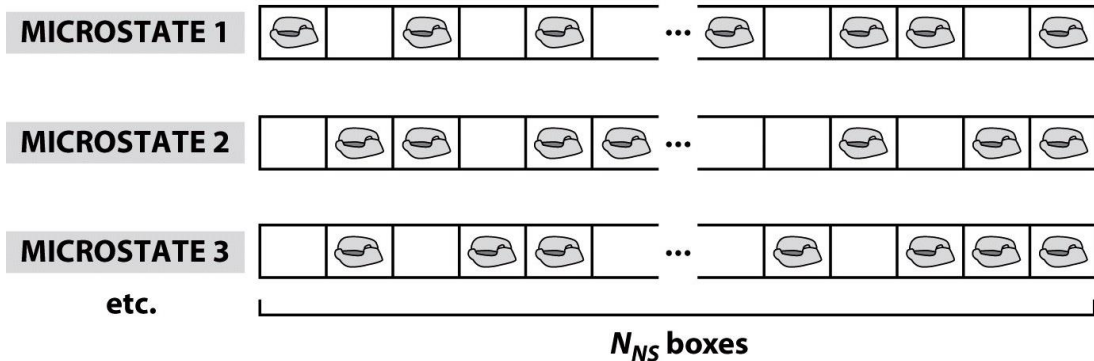
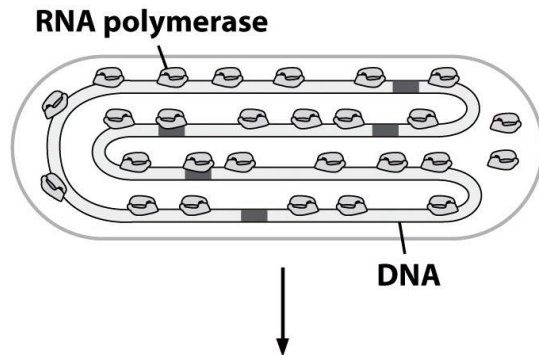
$P$  RNA polymerases

$N_{NS}$  boxes

max. one RNAP per box

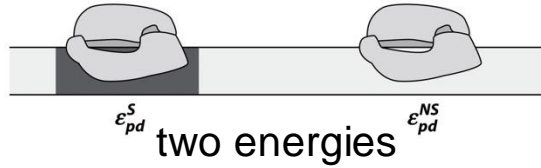
assume all are bound

## Microstates



# Statistical mechanics for biophysics

## Example: Transcriptional control



## Microstates

STATE	ENERGY	MULTIPLICITY	WEIGHT (MULTIPLICITY x BOLTZMANN WEIGHT)
	$P \epsilon_{pd}^{NS}$	$\frac{N_{NS}!}{P! (N_{NS} - P)!} \approx \frac{(N_{NS})^P}{P!}$	$\frac{(N_{NS})^P}{P!} e^{-P \epsilon_{pd}^{NS} / k_B T}$
	$(P-1) \epsilon_{pd}^{NS} + \epsilon_{pd}^S$	$\frac{N_{NS}!}{(P-1)! [N_{NS} - (P-1)]!} \approx \frac{(N_{NS})^{P-1}}{(P-1)!}$	$\frac{(N_{NS})^{P-1}}{(P-1)!} e^{-(P-1) \epsilon_{pd}^{NS} / k_B T} e^{-\epsilon_{pd}^S / k_B T}$



# Statistical mechanics for biophysics

Example: Transcriptional control

*Microstates: probability*

$$\begin{aligned}
 p_{\text{bound}} &= \frac{\sum_{\text{states}} \left( \text{Diagram of DNA with } P \text{ proteins bound to a promoter} \right)}{\sum_{\text{states}} \left( \text{Diagram of DNA with } P \text{ proteins bound to a promoter} \right) + \sum_{\text{states}} \left( \text{Diagram of DNA with } P \text{ proteins bound to a non-specific site} \right)} \\
 &= \frac{\frac{N_{NS}^{P-1}}{(P-1)!} e^{-\beta[(P-1)\epsilon_{NS} + \epsilon_S]}}{\frac{N_{NS}^P}{P!} e^{-\beta P \epsilon_{NS}} + \frac{N_{NS}^{P-1}}{(P-1)!} e^{-\beta[(P-1)\epsilon_{NS} + \epsilon_S]}} \\
 &= \frac{\frac{P}{N_{NS}} e^{-\beta \Delta \epsilon}}{1 + \frac{P}{N_{NS}} e^{-\beta \Delta \epsilon}} \\
 &= \frac{1}{\frac{N_{NS}}{P} e^{\beta \Delta \epsilon} + 1} \quad \begin{array}{l} \text{Langmuir isotherm} \\ \text{Hill function of coefficient 1} \end{array}
 \end{aligned}$$

Define:  $\Delta \epsilon = \epsilon_S - \epsilon_{NS}$

# Statistical mechanics for biophysics

## Example: Transcriptional control

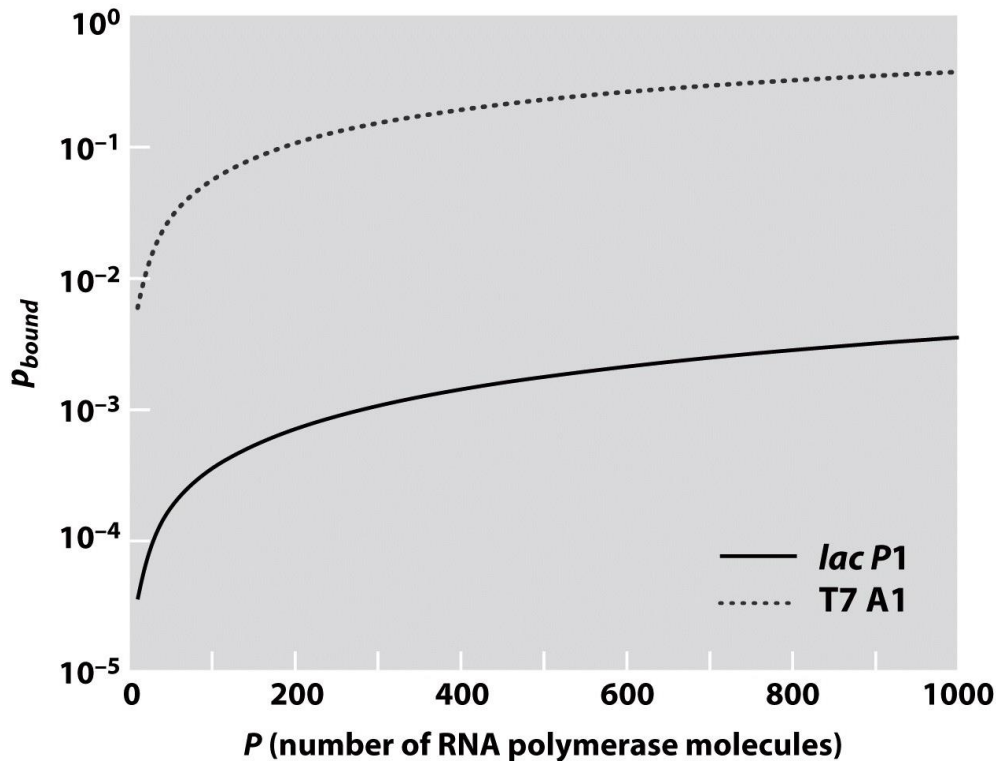
Estimate:

$$\Delta\varepsilon = -2.9 k_B T \text{ E. coli}$$

$$\Delta\varepsilon = -8.1 k_B T \text{ bacteriophage T7}$$

~5000 RNA polymerase in E. coli

## *RNA polymerase occupancy*



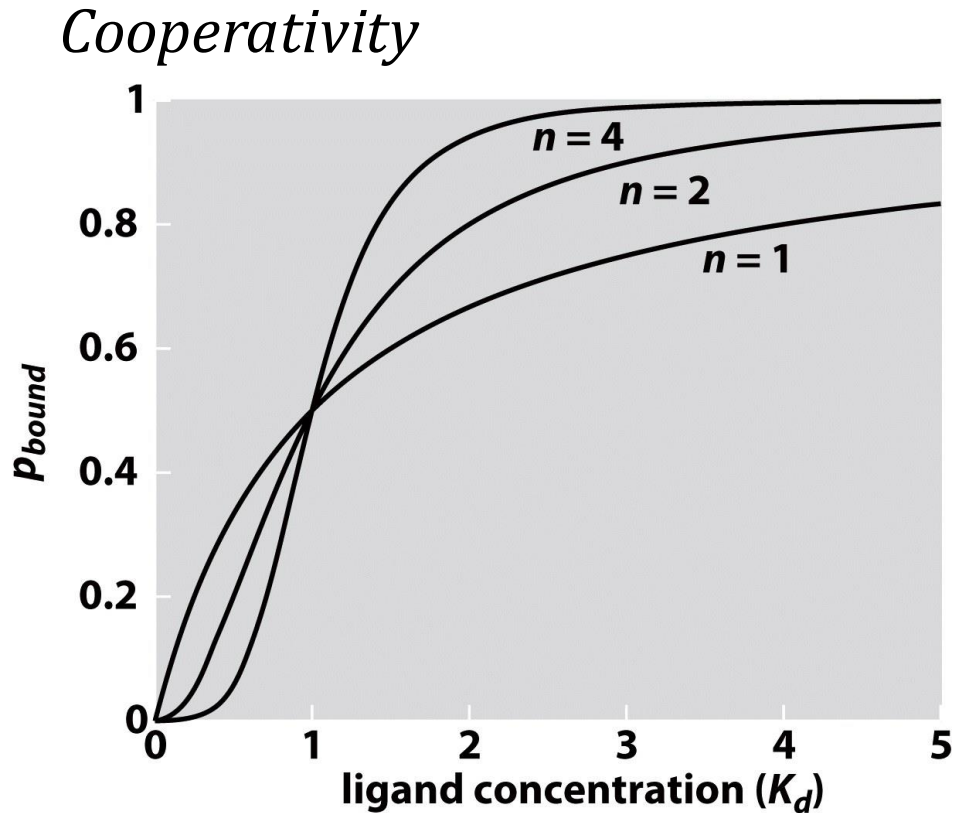
# Statistical mechanics for biophysics

## *Cooperativity*

Analog signal  $\rightarrow$  digital output

# Statistical mechanics for biophysics

Binding curves  
with different Hill  
coefficients



# Lecture 6: Entropy rules

Summary

# Lecture 7: Two-state system, ion channels

Goal: Statistical mechanics modeling. Compute the probability of microstates, including applied forces.

- Two-state system
- Mechanosensitive ion channels

PBOC Chapter 7.1.2, 11.5