

microstates obey

LDB:

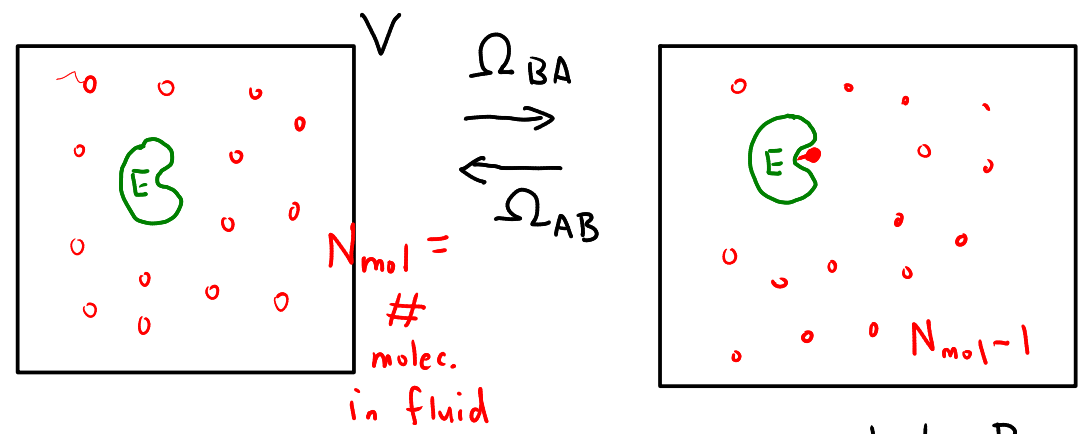
$$\frac{k_{mn}}{k_{nm}} = e^{-\beta(H_m - H_n - W_{mn})}$$

↑
other work

$$\frac{\Omega_{nm}}{\Omega_{mn}} = \frac{G_n k_{nm}}{G_m k_{mn}} = \frac{G_n}{G_m} e^{-\beta(H_n - H_m - W_{nm})}$$

$$= e^{-\beta(H_n - H_m - k_B T \ln \frac{G_n}{G_m} - W_{nm})}$$

example:
enzyme E
w/ fixed position



macrostate A
⇒ includes all configs of molec. in fluid

macrostate B
⇒ molec. bound to enzyme

binding

unbinding

$$\frac{\Omega_{BA}}{\Omega_{AB}} = e^{-\beta(H_B - H_A - k_B T \ln \frac{G_B}{G_A})}$$

$\underbrace{H_B - H_A}_{= 0 \text{ for molec. forming a bond}}$

$W_{BA} = 0$
 (no other work terms)

(if binding thermodyn. favorable)

$G_A = \#$ of configs of N_{mol} molecules in a box of volume V (ignore vol. of $E \ll V$)

divide up volume V into small volumes $V_{mol} =$
 vol. of molecule (big enough to fit one)

$$\# \text{ boxes} = N_{pos} = \frac{V}{V_{mol}}$$

how many ways to arrange N_{mol} molec. into
 N_{pos} possible positions?

$$\Omega_A = \binom{N_{pos}}{N_{mol}} = \frac{N_{pos}!}{N_{mol}! (N_{pos} - N_{mol})!}$$

$$\Omega_B = \binom{N_{pos}}{N_{mol}-1} = \frac{N_{pos}!}{(N_{mol}-1)! (N_{pos} - N_{mol} + 1)!}$$

$$\frac{\Omega_B}{\Omega_A} = \frac{N_{mol}}{N_{pos} - N_{mol} + 1} \approx \frac{N_{mol}}{N_{pos}} \quad \text{when } N_{pos} \gg N_{mol}$$

(dilute sol'n)

$$k_B T \ln \frac{\Omega_B}{\Omega_A} \approx k_B T \ln \frac{N_{mol}}{N_{pos}}$$

$$= k_B T \ln c V_{mol}$$

$$= k_B T \ln V_{mol} + k_B T \ln c$$

$$\underbrace{\quad}_{\mu_0 = \text{const.}}$$

= "standard" chem. potential

$$N_{pos} = \frac{V}{V_{mol}}$$

$$c = \frac{N_{mol}}{V}$$

= concentration

chemical potential $\equiv \mu = k_B T \ln \frac{\delta_B}{\delta_A} = \mu_0 + k_B T \ln c$

binding
unbinding

$$\frac{\Omega_{BA}}{\Omega_{AB}} = e^{-\beta(H_B - H_A - \mu)} \quad (\text{Eq. 1})$$

When $N_{\text{pos}} \gg N_{\text{mol}} \Rightarrow \mu < 0$

increasing $c \Rightarrow \mu$ less negative

\Rightarrow make Ω_{BA} larger
relative to Ω_{AB}

\Rightarrow larger conc. favor
more binding

compare this to the FAR-NEAR-BOUND
model from earlier in the course: $0 \leq \alpha_b \leq 1$

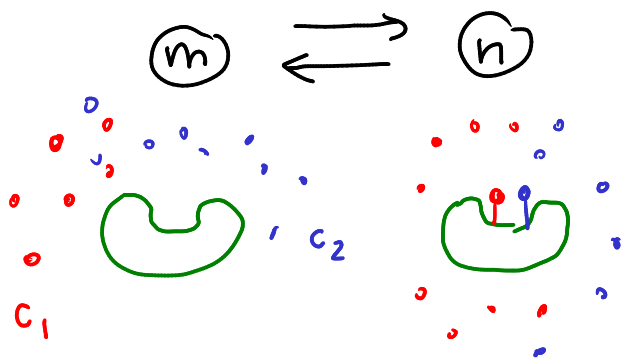
$$\text{Eq. 2: } \frac{\Omega_{BA}}{\Omega_{AB}} = \frac{\text{binding}}{\text{unbinding}} = \frac{k_b}{k_u} = \frac{\alpha_b K_s c}{K_u}$$

both Eq. 1 + Eq. 2 are $\propto c \Rightarrow$ consistent!

generalize to many types of particles
in fluid around system

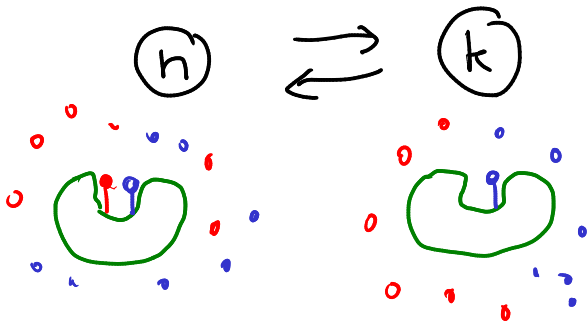
type j molec: $\mu_j = \mu_0^j + k_B T \ln c_j$

\hookrightarrow conc. of
type j



$$\frac{\Omega_{nm}}{\Omega_{mn}} = e^{-\beta(H_n - H_m - \mu_1 - \mu_2)}$$

for every molec. that binds we get a μ_j term w/ minus sign



numerator

$$\frac{\Omega_{nm}}{\Omega_{mn}} = e^{-\beta(H_n - H_m + \mu_1)}$$

numerator trans.

unbinding of type j : $+\mu_j$ term

example:

