

estimates:  $k_{\text{dif}} = K_s c$   $K_s = 4\pi DR \approx 10^8 M^{-1} s^{-1}$

$$c \sim 10 \text{ nM} - 0.1 \text{ mM} \quad \begin{matrix} \text{Smoluchowski} \\ \text{coeff.} \end{matrix}$$

$$\Rightarrow k_{\text{dif}} \sim 1 \text{ s}^{-1} \sim 10^4 \text{ s}^{-1}$$

$k_{\text{dri}} \Rightarrow$  a molecule diffusing 1 nm away from target

$$\text{time} \sim \frac{R^2}{6D} \quad R = 1 \text{ nm} \\ D = 20 \mu\text{m}^2/\text{s}$$

$$\sim 8 \text{ ns}$$

$$k_{\text{dri}} \sim \frac{1}{8 \text{ ns}} = 10^8 \text{ s}^{-1}$$

To find  $\tau_{1 \rightarrow 3}$  (avg. time from FAR to BOUND)

use example from earlier lecture:

$$\sum \Omega_{ij} \tau_i = -1$$

$$\tau_{1 \rightarrow 3} = \frac{k_{\text{dif}} + k_{\text{dri}} + k_{\text{att}}}{k_{\text{att}} + k_{\text{dif}}}$$

effective rate from  $1 \rightarrow 3$

$$k_b \equiv \frac{1}{\tau_{1 \rightarrow 3}} = k_{\text{dif}} \left[ \frac{k_{\text{att}}}{k_{\text{dif}} + k_{\text{dri}} + k_{\text{att}}} \right]$$

$\uparrow$  binding rate

$\equiv \alpha_b$  coefficient

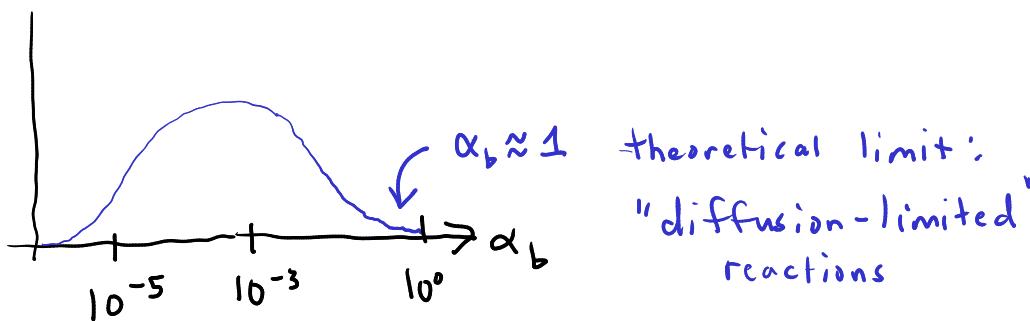
$$0 \leq \alpha_b \leq 1$$

$$\Rightarrow k_b = \alpha_b k_{dif} = \alpha_b K_s C \leq k_{dif}$$

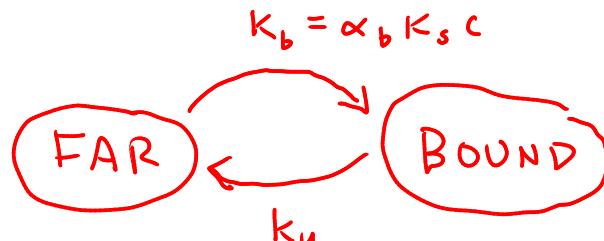
"Speed limit"

since  $k_{dif} \ll k_{dri} \Rightarrow \alpha_b \approx \frac{k_{att}}{k_{dri} + k_{att}}$  approx.  
indep. of  $C$

histogram of  $\alpha_b$  over all biochemical reactions



effective 2 state approx.



eff. unbinding rate

$$= \frac{1}{\tau_{3 \rightarrow 1}} = \frac{k_{det} k_{dri}}{k_{det} + k_{dri} + k_{att}}$$

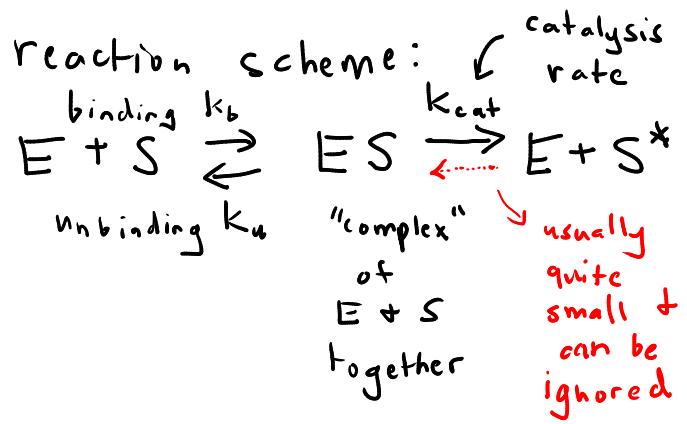
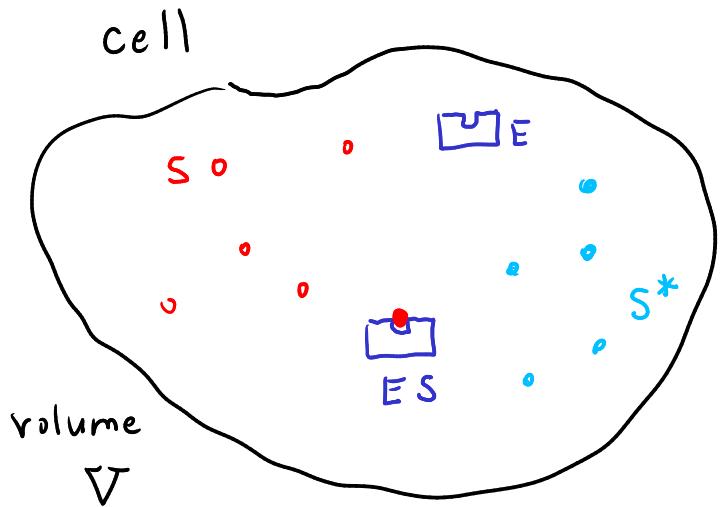
const. indep. of  $C$

focus on a full chemical reaction:

enzyme  $E$  (one type of protein)

substrate  $S$  (another protein or small molecule)

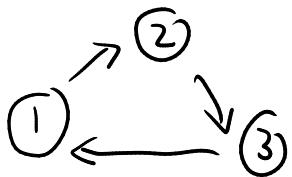
reaction  $S \rightarrow S^*$  (modified version of substrate)



chemical state:

$$\vec{n} = (n_S, n_E, n_{ES}, n_{S^*})$$

$n_\alpha$  = # of type  $\alpha$  in cell



$\Omega_{nm}$   $\rightsquigarrow$   $\Omega_{\vec{n}\vec{m}}$   $\equiv$  trans. rate  
 $m \rightarrow n$  generalize from state

$\vec{m}$  to  $\vec{n}$   
(for  $\vec{m} \neq \vec{n}$ )

probability  $p_{\vec{n}}(t)$  = prob. to be in state  $\vec{n}$  at time t

$$\sum_{\vec{n}} p_{\vec{n}}(t) = 1$$

$\left[ \sum_{n_S=0}^{\infty} \sum_{n_E=0}^{\infty} \dots \right]$

columns of  $\Omega_{\vec{n}\vec{m}}$   
sum to zero

$$\sum_{\vec{n}} \Omega_{\vec{n}\vec{m}} = 0$$

$$\frac{d}{dt} p_{\vec{n}}(t) = \sum_{\vec{m}} \Omega_{\vec{n}\vec{m}} p_{\vec{m}}(t)$$

chemical master eqn.

(accurate, but hard way of doing chemistry)

conservation laws (stoichiometry)

$$n_S + n_{ES} + n_{S^*} = \text{const.} \equiv M_S$$

$$n_E + n_{ES} = \text{const.} \equiv M_E$$

example:  $M_S = 2$ ,  $M_E = 2$

allowed  
states

$$\vec{n} = \begin{pmatrix} n_s \\ 2 \\ 2 \\ 0 \\ 0 \end{pmatrix}$$

binding

catalysis

$$\begin{pmatrix} 1 & 1 & 1 & 0 \end{pmatrix}$$
$$\begin{pmatrix} 1 & 2 & 0 & 1 \end{pmatrix}$$

...  
.

catalysis

$$\begin{pmatrix} 0 & 0 & 2 & 0 \end{pmatrix}$$
$$\begin{pmatrix} 0 & 1 & 1 & 1 \end{pmatrix}$$

binding