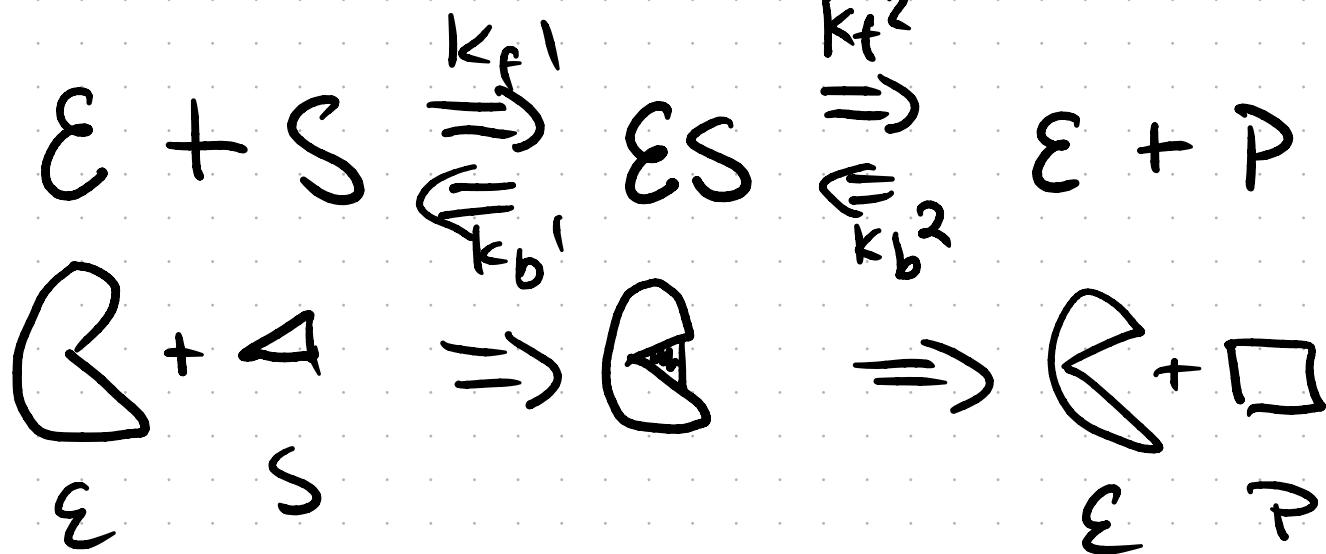
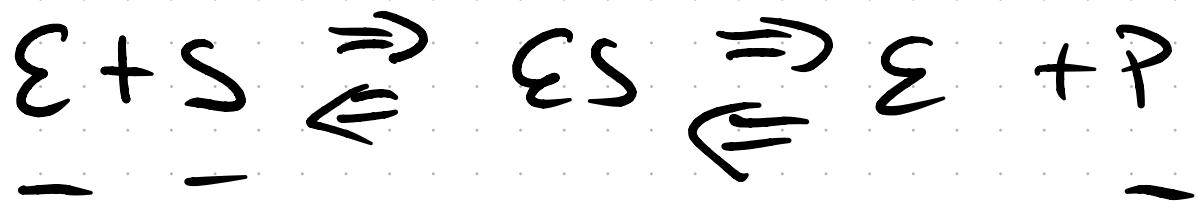


# Enzymes Biological Catalysts

# Michelis-Menten Schene





$$\frac{d[S]}{dt} = -k_f^{-1}[\varepsilon][S] + k_b^{-1}[\varepsilon S]$$

$$\frac{d[P]}{dt} = k_f^2 [ES] - k_b^2 [E][P]$$

$$\frac{d[\text{ES}]}{dt} = k_f' [\text{EJ}[S] + k_b^2 [\text{EJ}[P]] - k_b [\text{ES}] - k_f^2 [\text{ES}]]$$

$$\frac{d[S]}{dt} = -k_f' [\varepsilon][S] + k_b' [\varepsilon S]$$

$$\frac{d[P]}{dt} = k_f^2 [\varepsilon S] - k_b^2 [\varepsilon J][P]$$

$$\begin{aligned}\frac{d[\varepsilon S]}{dt} &= k_f' [\varepsilon J][S] + k_b^2 [\varepsilon J][P] \\ &\quad - k_b' [\varepsilon S] - k_f^2 [\varepsilon S]\end{aligned}$$

Total Enzyme is conserved

$$[\varepsilon]_0 = [\varepsilon S] + [\varepsilon]$$

$$[\varepsilon] = [\varepsilon]_0 - [\varepsilon S]$$

$$\frac{d[\varepsilon s]}{dt} = k_f' [\varepsilon][s] + k_b^2 [\varepsilon][p] - k_b' [\varepsilon s] - k_f^2 [\varepsilon s]$$

$$[\varepsilon] = [\varepsilon]_o - [\varepsilon s] \quad \uparrow$$

$$\frac{d[\varepsilon s]}{dt} = k_f' ([\varepsilon]_o - [\varepsilon s]) [s]$$

$$+ k_b^2 ([\varepsilon]_o - [\varepsilon s]) [p]$$

$$= -[\varepsilon s] \overline{(k_b' + k_f^2) [\varepsilon s]}$$

$$= -[\varepsilon s] \left[ k_f' [s] + k_b^2 [p] + k_b' + k_f^2 \right]$$

$$+ k_f' [\varepsilon]_o [s] + k_b^2 [\varepsilon]_o [p]$$

assumption  
 ①  $[s] \gg [\varepsilon]$

steady state  
 approx

$$[\varepsilon S]_{ss} = \frac{(k_f' [S] + k_b^2 [P]) [\varepsilon]_0}{k_f' [S] + k_b' + k_b^2 [P] + k_f^2}$$



$$\begin{aligned} r &= -\frac{d[S]}{dt} = k_f' [\varepsilon] [S] - k_b' [\varepsilon S] \\ &= k_f' ([\varepsilon]_0 - [\varepsilon S]) [S] - k_b' [\varepsilon S] \end{aligned}$$

$$V = \frac{(k_f' k_f^2 [S] - k_b' k_b^2 [P]) [\varepsilon]_0}{k_f' [S] + k_b' + k_b^2 [P] + k_f^2}$$

approximations

- ① Steady state,  $[\varepsilon] \ll [S]$
- ② Measure initial rate

$$[S] \approx [S]_0, [P] \approx 0$$

$$V_{\text{initial}} \approx \frac{k_f' k_f^2 [S]_0 [\varepsilon]_0}{k_f' [S]_0 + k_b' + k_f^2}$$

$$V_{\text{initial}} \approx \frac{k_f' k_f^2 [S]_0 [\epsilon]_0}{k_f' [S]_0 + k_b' + k_f^2}$$

$$\text{bottom} = k_f' \left( [S]_0 + \frac{k_b' + k_f^2}{k_f'} \right)$$

units of concentration  
↓

$$V_{\text{initial}} = \frac{k_f^2 [S]_0 [\epsilon]_0}{[S]_0 + \frac{k_b' + k_f^2}{k_f'}}$$

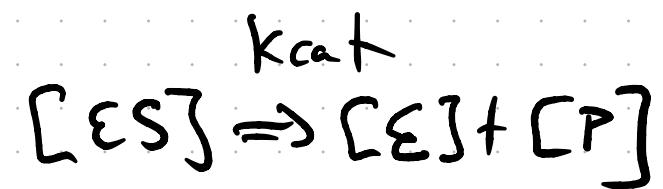
$$K_M = \frac{k_f^2 + k_b'}{k_f'}$$

Michaelis constant

At early times

$$v_{\text{initial}} = \frac{k_2 [S]_0 [\text{E}]_0}{[S]_0 + K_m}$$

$$k_2 = k_{\text{cat}}$$



2 limits

$$[S]_0 \ll K_m$$

$v_{\text{initial}}$  is proportional to  $[S]_0$

$$[S]_0 \gg K_m$$

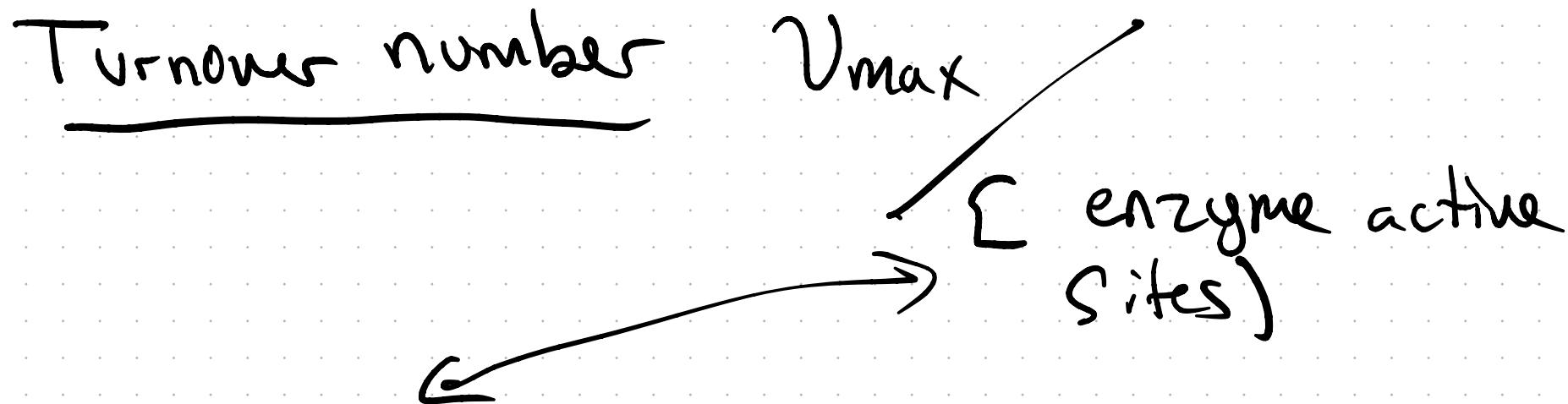
$v_{\text{initial}}$  is 0th order in  $[S]_0$

$$v_{\text{max}} = v_{\text{initial}} = k_{\text{cat}} [\text{E}]_0 \leftarrow \text{maximum rate}$$

$$V_{\text{initial}} = \frac{V_{\text{max}} [S]_0}{[S]_0 + K_m} \quad \text{if } [S]_0 = K_m$$

$$V_{\text{initial}} = V_{\text{max}} / 2 \leftarrow \text{can do this}$$

to find  $K_m$



$$[\text{active sites}] = [\epsilon]_0 \cdot (\# \text{ of sites/enzyme})$$

If # sites/enzyme = 1

then turnover number =  $k_{cat}$

Catalytic Efficiency - how effectively  
an enzyme converts substrate to  
product

$$E = \frac{k_{cat}}{K_m} \text{ — units of } \frac{1}{M} \cdot \frac{1}{s}$$

range of  $E \sim 1 - 10^{10} \frac{1}{ms}$   $\underbrace{10^8 - 10^{10}}$   
diffusion limited

$$G = \frac{k_{cat}}{K_m} = \frac{k_{cat}}{\frac{k_{cat} + k_b'}{k_{f'}}} = \frac{k_{f'}}{1 + \frac{k_b'}{k_{cat}}}$$

Maximum,  $k_{f'}$  or  $k_{cat}$  big

or  $k_b'$  small

# Introduction to Statistical thermodynamics

(Barrick chapter 9)

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$$\frac{k_{on}}{k_{off}} = K_{eq} = e^{-\Delta G^{\circ}/RT}$$

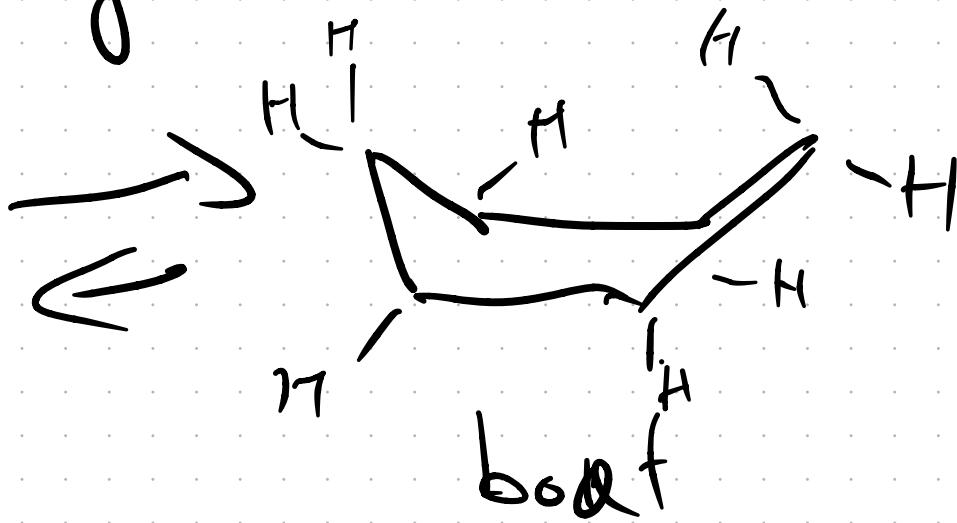
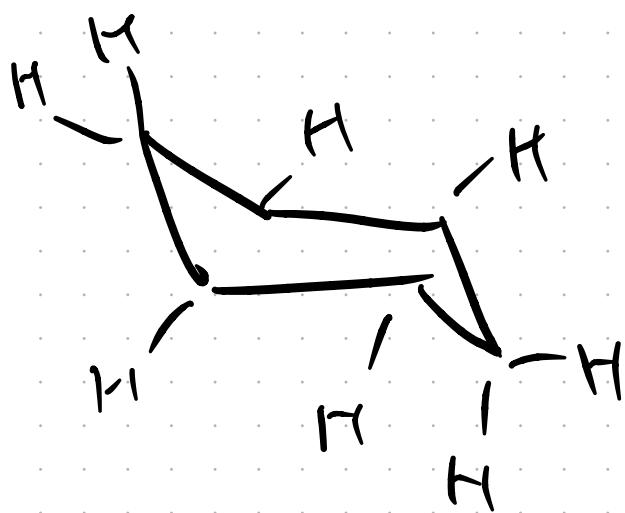
Didn't see how this arises  
for a collection of molecules

Statistical thermodynamics connects  
average properties of sets of molecules  
with Macroscopic thermodynamic properties

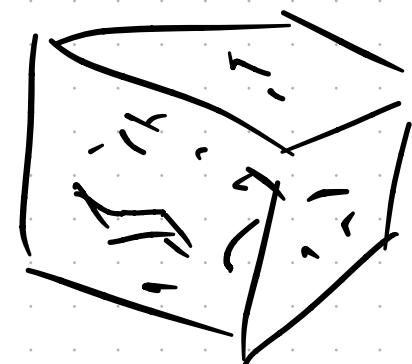
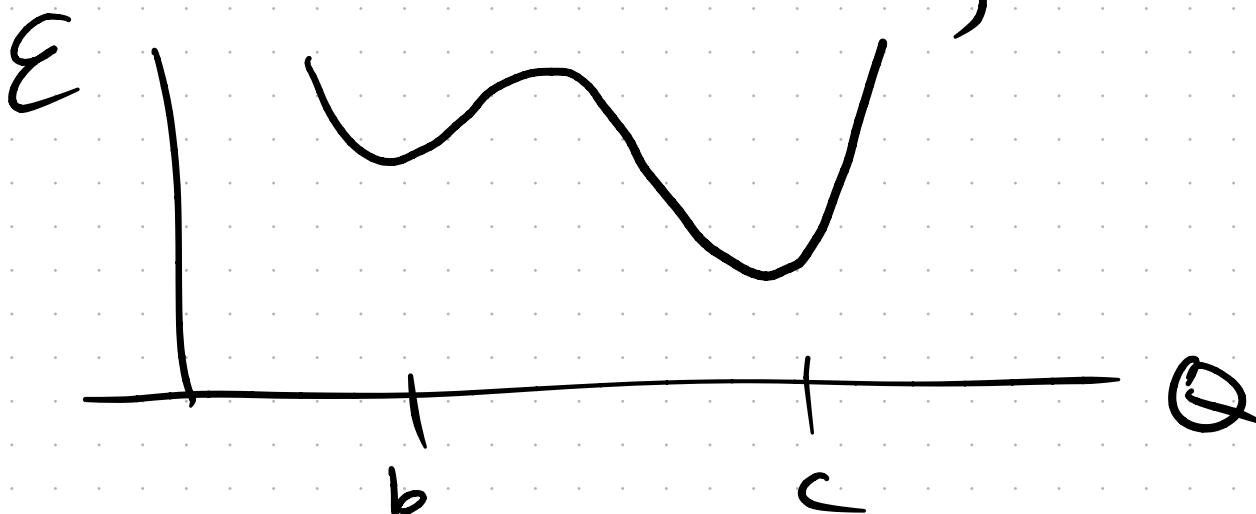
# Molecular Example

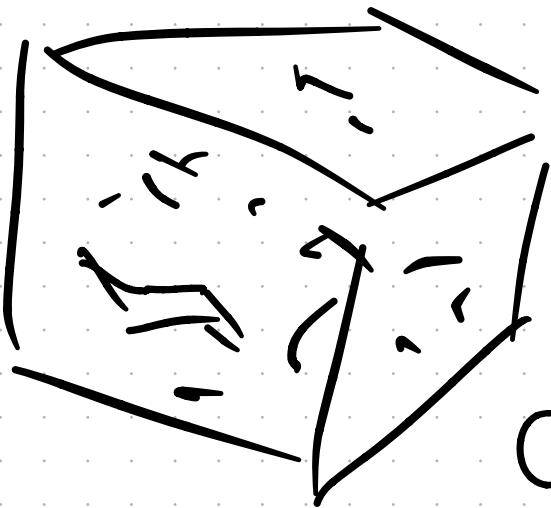


Molecule that goes between 2 states



chair < lower energy





what if we could  
watch this molecule  
and see its cfg. vs time

Can do this with a simulation

Assume all molecules obey Newton's equations

$$F_i = m_i a_i \text{ for every atom}$$

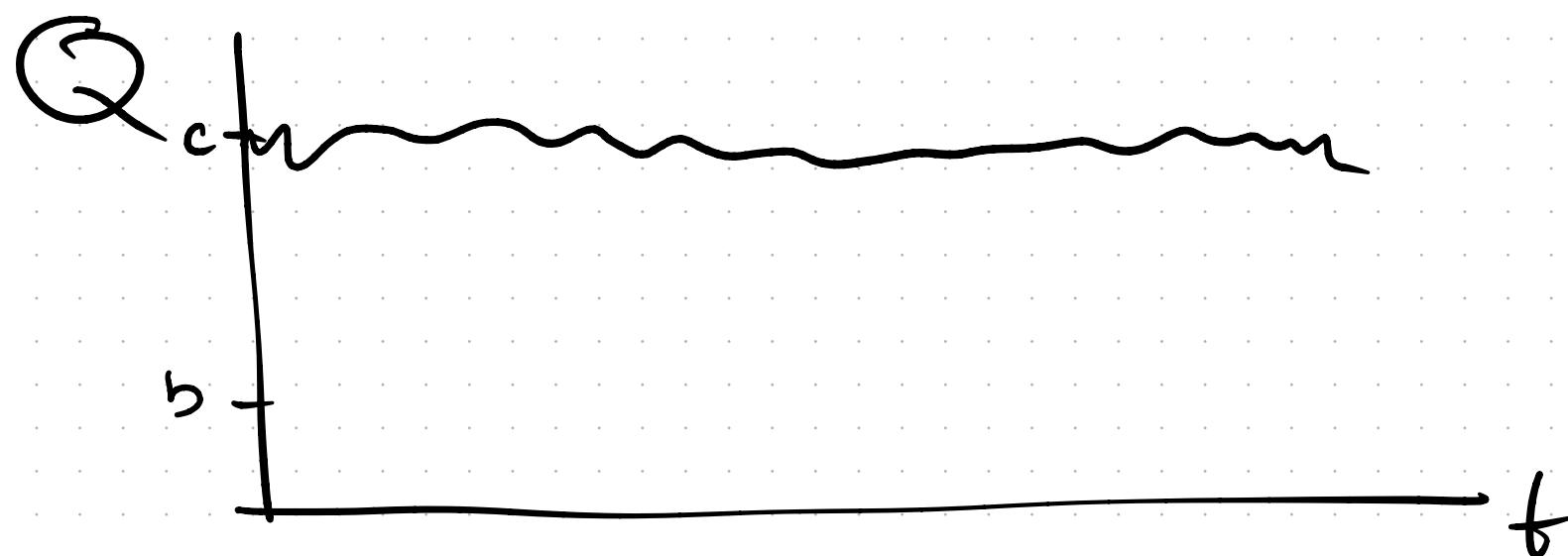
$$F_i = -\frac{\partial U}{\partial \vec{x}_i}$$

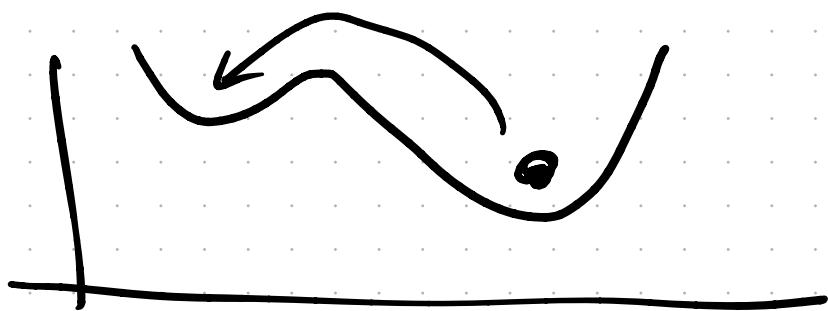
force field  
 $U(\vec{x})$

Problem is that real systems have  $N \sim 10^{23}$

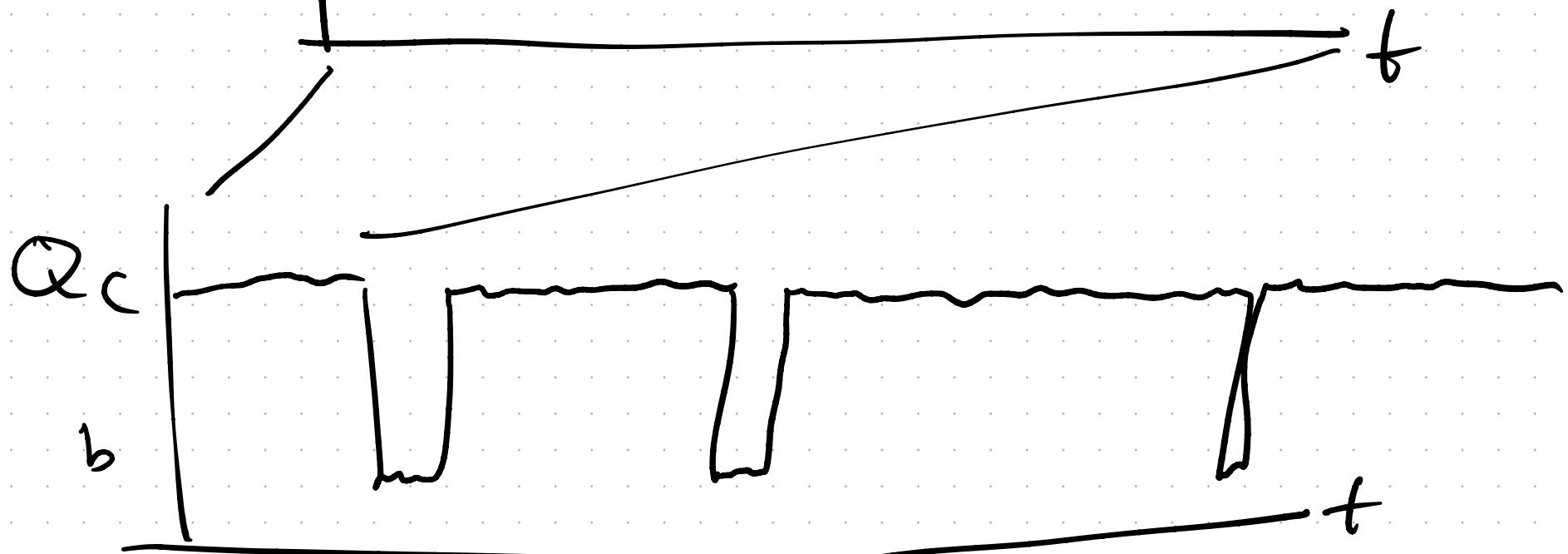
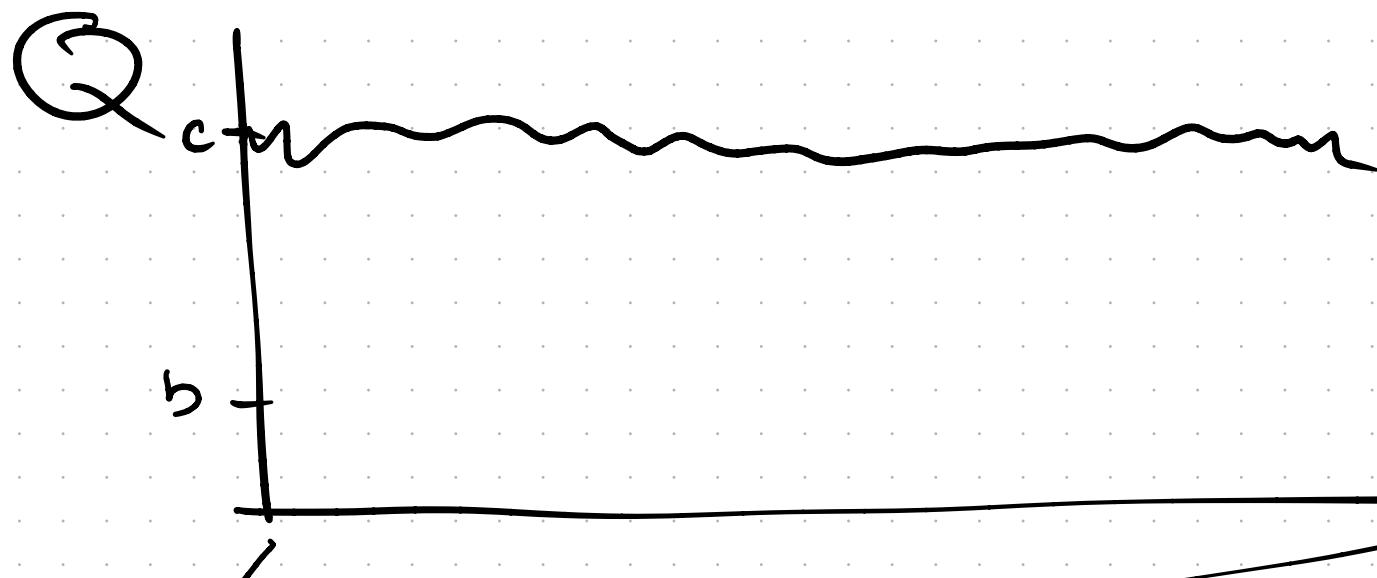
Because real systems are large  
we will treat them statistically

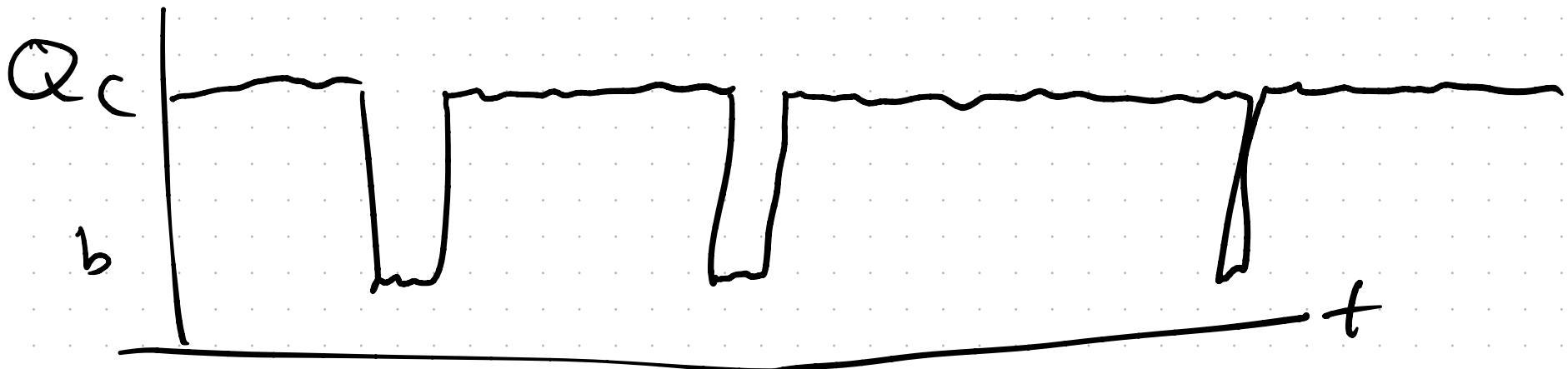
What would happen if we watch  
a 2 state molecule in H<sub>2</sub>O





$$K_{c \rightarrow b} = A e^{-G^*/RT}$$



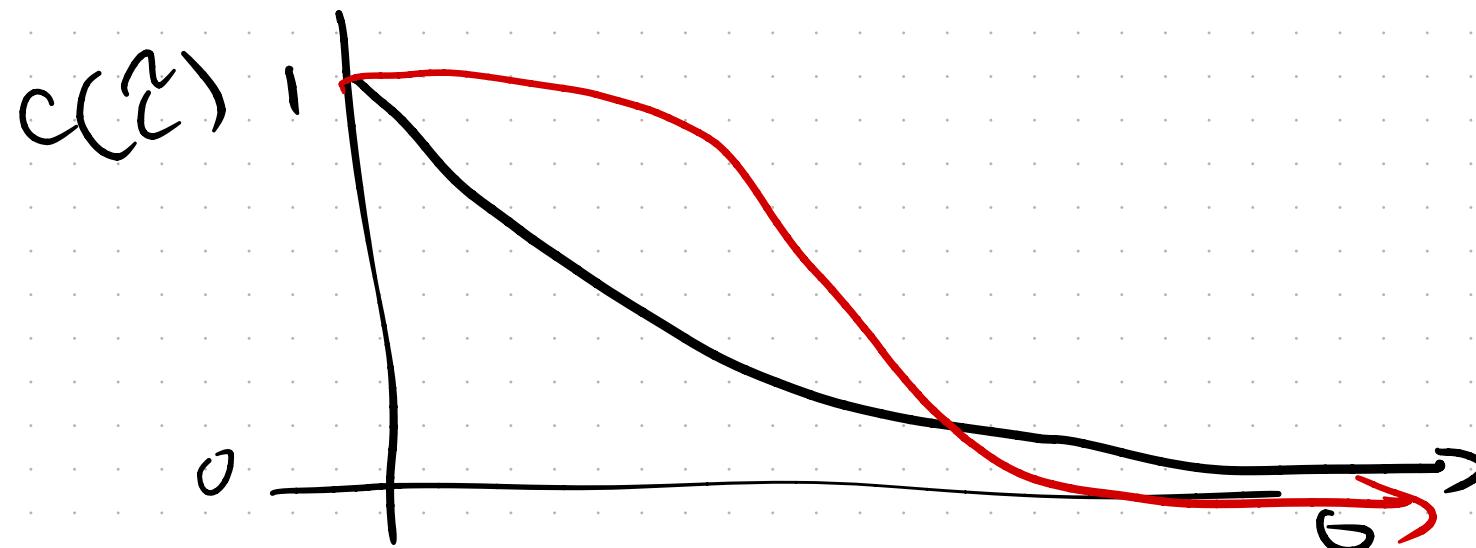


Auto correlation function

tells us over what period of time  
is a molecule's configuration correlated  
with itself

$C(\tau)$  small  $\tau$   $C = 1$

long  $\tau$   $C = 0$



compare  $Q(t)$  to  $Q(t + \Delta)$

calculate  $\langle Q \rangle = \frac{1}{N_t} \sum_{i=1}^{N_t} Q(t_i)$

$$\delta Q(t) = Q(t) - \langle Q \rangle$$

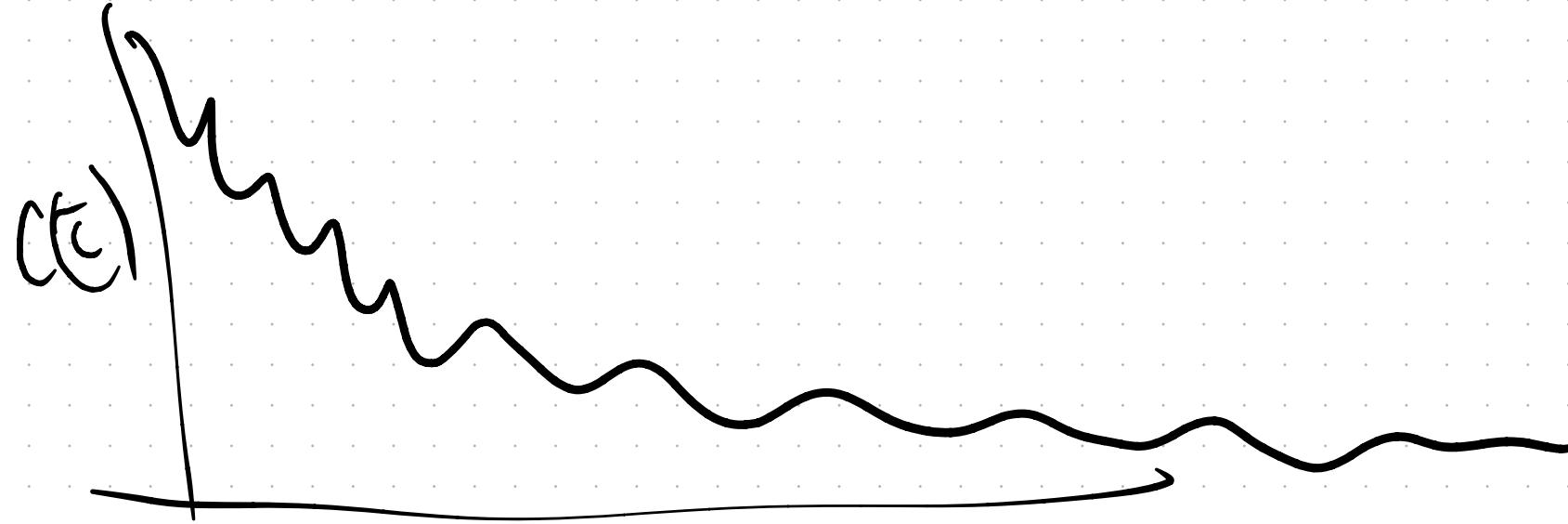
$$\langle \delta Q(t)^2 \rangle = \langle (Q(t) - \langle Q \rangle)^2 \rangle = \text{Var}(Q)$$

$$C(\tau) = \frac{\langle \delta Q(t_i + \tau) \delta Q(t_i) \rangle}{\langle \delta Q(t_i) \delta Q(t_i) \rangle}$$

$$@ \tau = 0 \quad C(0) = 1$$

$$@ \tau \rightarrow \infty \text{ turns} \quad C(\tau \rightarrow \infty) = 0$$

$$C(\tau) = \frac{\frac{1}{N} \sum_{i=1}^N \delta Q(t_i + \tau) \delta Q(t_i)}{\frac{1}{N} \sum_{i=1}^N (\delta Q(t_i))^2}$$



define reaction time

$$\tau_{rxn} = \int_0^{\infty} C(\tau) d\tau \equiv \tau_{rxn}$$

$$C(t) \approx e^{-at} \quad \text{then} \quad a = 1/\tau_{rxn}$$