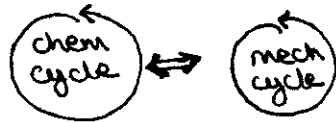
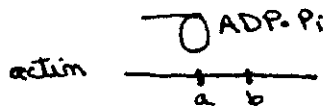
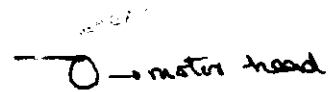


Molecular motors and mechanochemistry

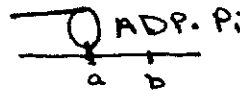
The challenge in mechanochemistry is to connect the chemical & mechanical cycles of a motor in order to understand the transduction of chemical energy to mechanical work.



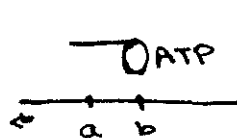
Smooth muscle myosin



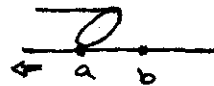
→



↓ power stroke  
product release  
ADP, Pi



← ATP  
detachment  
↑  
recovery

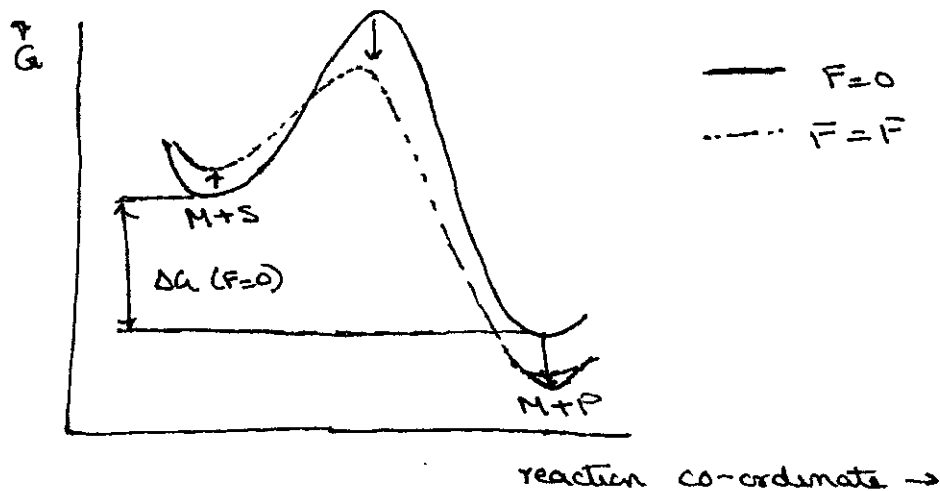
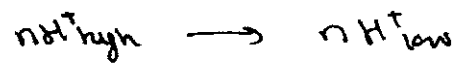
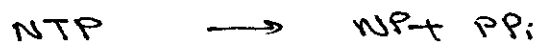
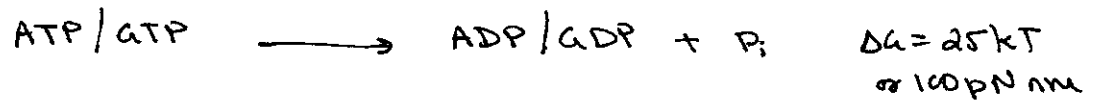


- There are coupling rules between the chemical & mechanical cycle that prevent the motor from running backwards.

## Landscape of a motor

- motors are coupled to an energetically favorable reaction. Therefore motor reactions are energetically downhill.

- energy sources:



## Mechanical properties of motors

- Step size: distance moved per catalytic cycle
- Stall force: - The force at which the velocity of the motor reduces to zero.  
Hence, this is the maximum force that a functional motor can generate
- Stall force of a motor may have some relevance to its biological functions.

eg RNAP ( $F_{\text{stall}} \approx 25 \text{ pN}$ ) has to knock off obstacles encountered while travelling on DNA

$\phi 29$  ( $F_{\text{stall}} \approx 57 \text{ pN}$ ) has a higher stall force presumably because of the high pressure build-up as it packages DNA inside the viral capsid.

- Mechanical efficiency  $\Rightarrow \eta = \frac{F \cdot \Delta x}{\Delta G} = \frac{\text{output work}}{\text{input energy}}$

- max efficiency is achieved at stall

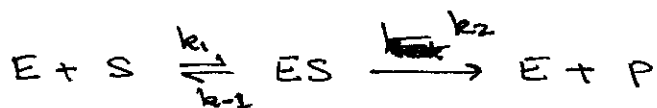
-  $\eta < 100\%$   $\rightarrow$  energy dissipated as heat

pg 3.

$\rightarrow$  work along an orthogonal axis

## Mechanochemistry

- chemical reaction  $\implies$  force-generation
- How does inhibitory or assisting force affect the output of the chemical reaction?
- Measurable outputs for a motor are quantities like velocity, stepping duration etc. Mechanochem is understood best by looking at the motor output at various chemical variables such as substrate concentrations, inhibitors etc & simultaneously at various forces
- A quick review of enzyme kinetics



Under initial velocity conditions

$$v = \frac{V_{max} [S]}{[S] + K_m} \quad ; \quad K_m = \text{Michaelis constant} = \frac{k_{-1} + k_2}{k_1}$$

$v$  = motor velocity

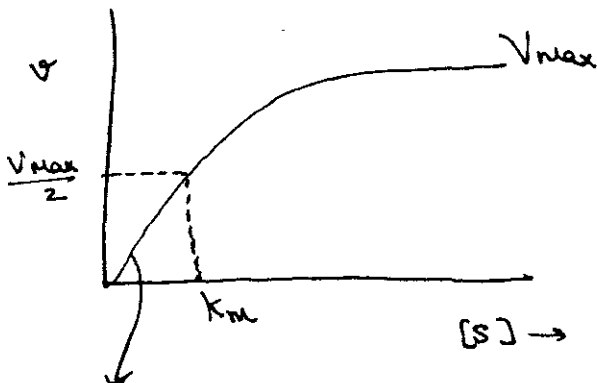
$V_{max}$  = maximum velocity obtainable when substrate binding is not limiting  
 $\equiv k_{cat}$  for a single motor enzyme

$[S]$  = substrate concentration

$K_m$  =  $[S]$  at which  $v = V_{max}/2$

At saturating substrate concentrations  $v = V_{max}$

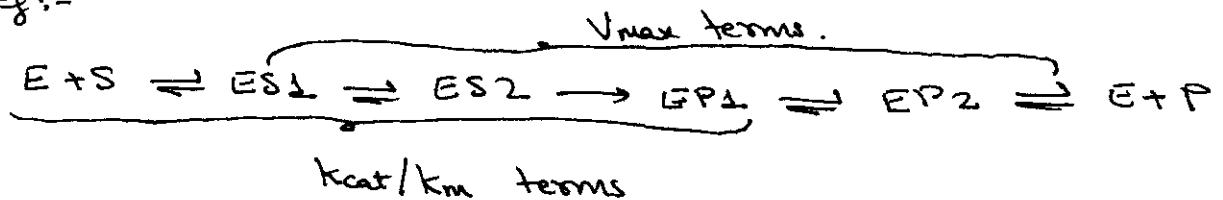
At  $S \ll K_m$ ,  $v = \frac{V_{max} [S]}{K_m}$



initial slope =  $\frac{k_{cat}}{K_m}$  = effective 2nd order binding constant

Enzyme reactions are generally more complex than the one discussed earlier.

eg:-



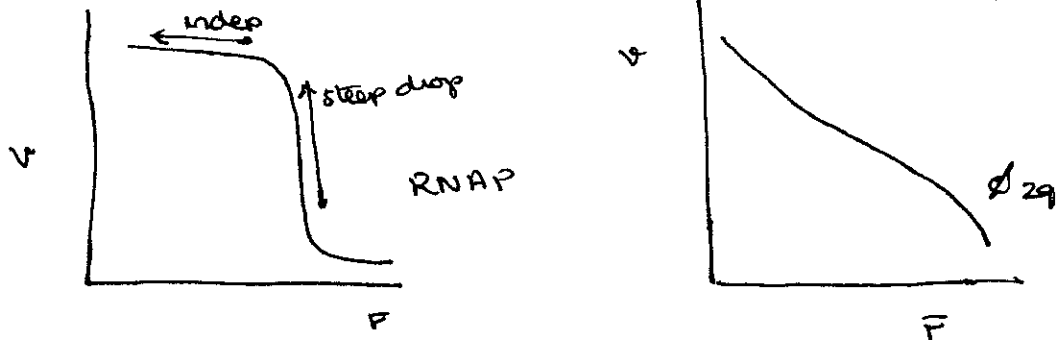
In general:-

$k_{cat}/K_m$  includes all steps from substrate binding to the first irreversible step

$V_{max}$  includes all steps other than substrate binding

## Force-velocity curves & mechanism

eg: F-V curves at saturating substrate concentrations have different shapes for different enzymes



in RNAP, force dependent step is not rate limiting until stall

in  $\sigma 29$ , force dependent step is rate limiting even at low forces.

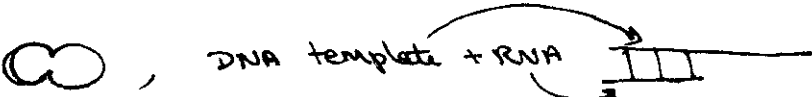
There are two major classes of models to describe motor mechanism

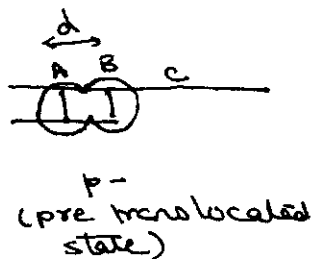
↳ Brownian ratchet

↳ Power stroke

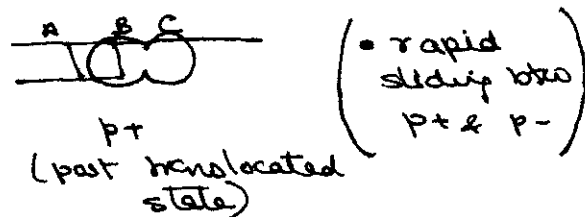
### ① Brownian ratchet

In this model, the motor can thermally visit two adjacent binding sites on its track via brownian motion & this can be rectified by coupling it to another stochastic process which is energetically downhill such as ATP hydrolysis.

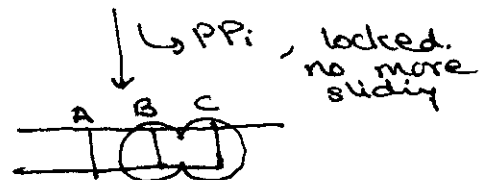
eg T7 RNAP  , DNA template + RNA



$$K(F) \rightleftharpoons \frac{k_1}{k_{-1}}$$



1 NTP



In such a mechanism:

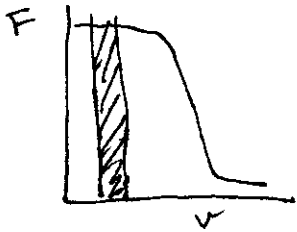
- chemical energy favors forward motion rather than doing the mechanical work directly on the motor
- rate of the reaction is proportional to the relative occupancies of P+ & P-

$$\frac{P_+}{P_-} = K \exp(-Fd/k_B T)$$

where d is the distance btw P- & P+

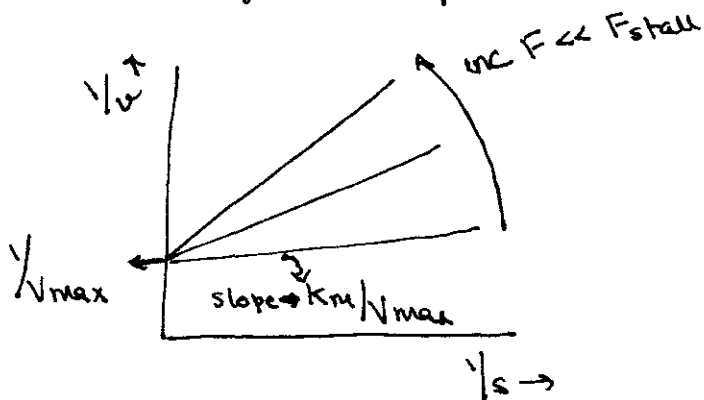
- The effect of force will be to push the  $k_1/k_{-1}$  equilibrium towards the "P-" state. This will result in a higher  $K_m$  at higher forces. In other words, more NTP will be required to reach  $V_{max}$

Thomen et al, PRL 94, 128102 (2005).



experiments conducted away from stall force to look at inhibition of T7 RNAP by force at various substrate concentrations

Plot of  $1/v$  vs  $1/s$  at various forces



Plot resembles a simple competitive inhibition plot

- $v_{max}$  is the same at all forces
- $k_{cat}/K_m$  decreases with increasing force
- $K_m$  increases with increasing force

The data is fully consistent with a brownian ratchet model for T7 RNAP.



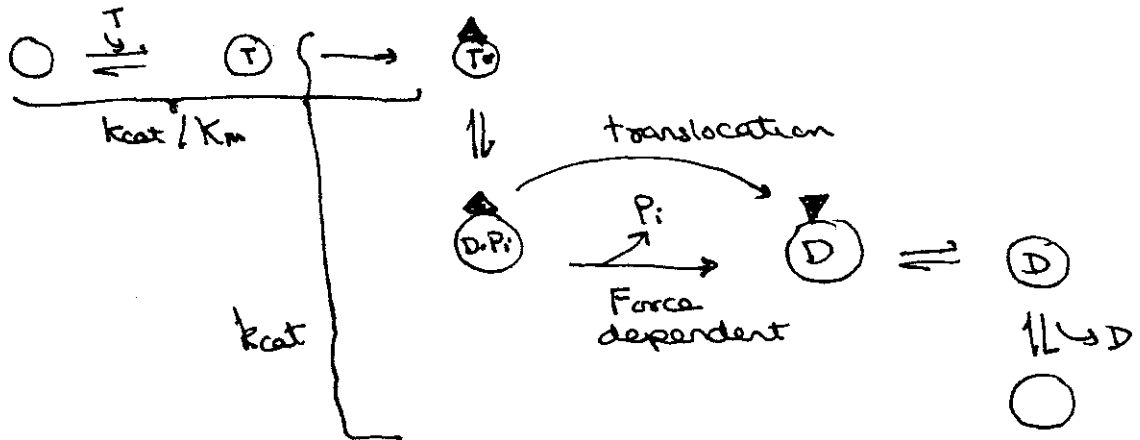
② Power stroke model (myosin,  $\phi 29$ )

In this model, the chemical reaction is mechanically coupled to movement by some part of the molecule

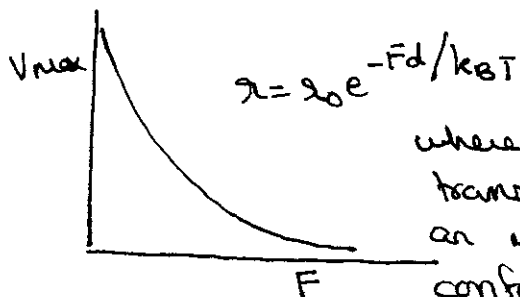
Archives

eg  $\phi 29$ ; Chemla, Y. et al Cell, Vol 122 (5), 683-692

minimal motor mechanism:-  $\circ$  motor  
 $\Delta$  attachment to DNA



This model is consistent with the observation that  $V_{max}$  decreases with force, where as  $k_{cat}/K_m$  is independent of force.



where  $d$  is the distance to the transition state & can be an indication of the size & the conformational change in the force dependent step.

Some points to keep in mind while looking at force-vel curves

- Not all cases can be clearly identified as brownian ratchet or power stroke.
- The dependence of  $v_{max}$  &  $v_{max}/k_m$  on force will be dictated by the details of the biochemical mechanism (ie first reversible step location, two-step versus one step substrate binding mechanisms etc). Hence interpretation of results depends on prior knowledge about the biochemical mechanism.
- Force could induce off-pathway inhibition introducing effects on  $v_{max}$  &  $k_{cat}/k_m$  that have nothing to do with the on-pathway mechanism of the enzyme.

Reference :-

Bustamante et al Annual Rev Biochem 2004. 73 : 705-748