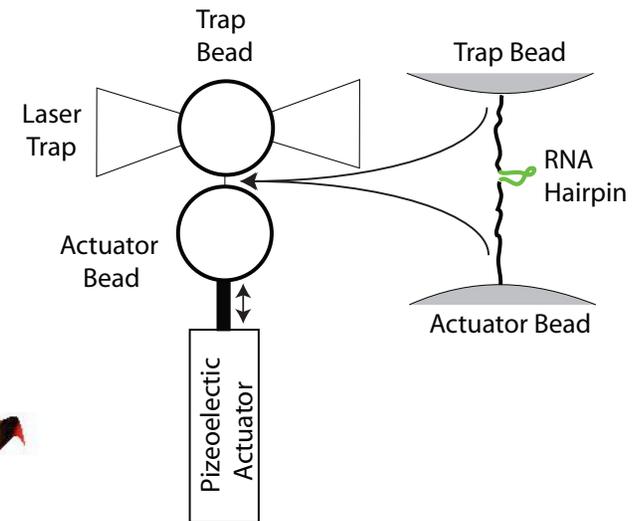
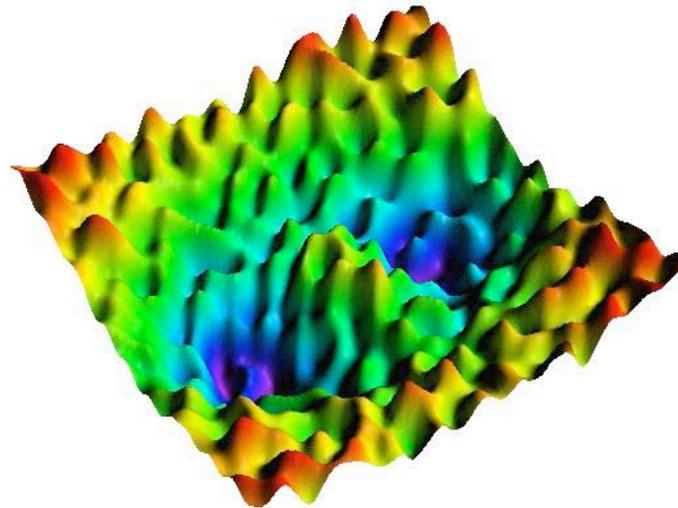
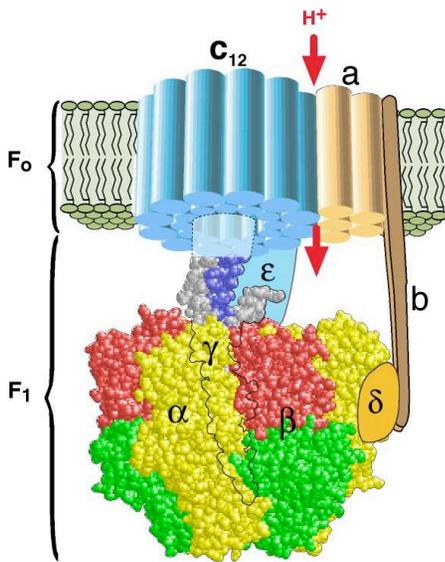


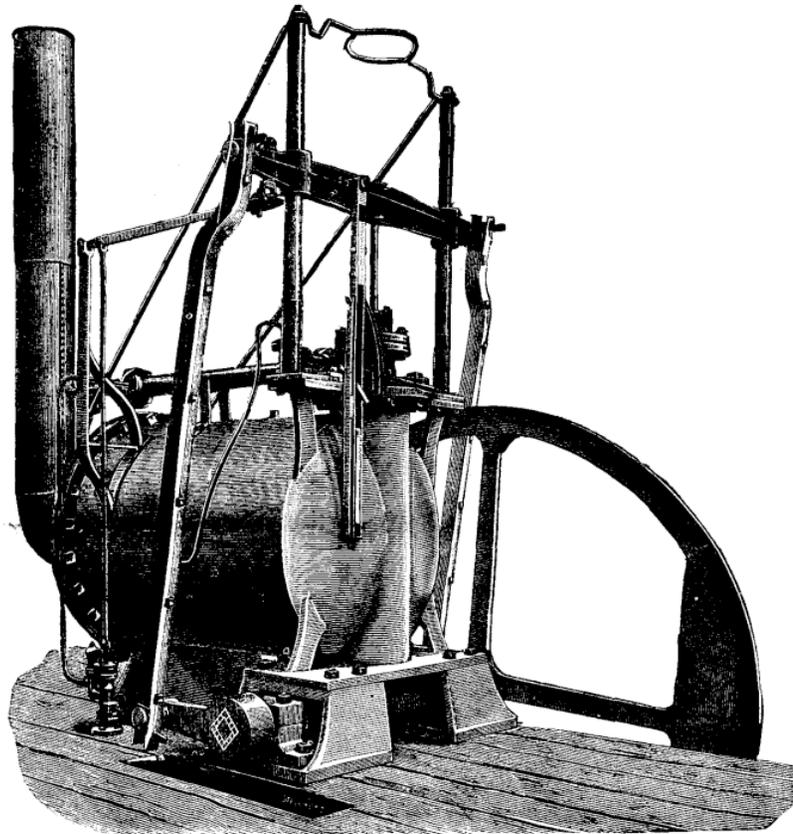
There and Back Again: The statistical dynamics of trajectories

Gavin E. Crooks
Physical Biosciences
Lawrence Berkeley Natl. Lab



Steam Engines

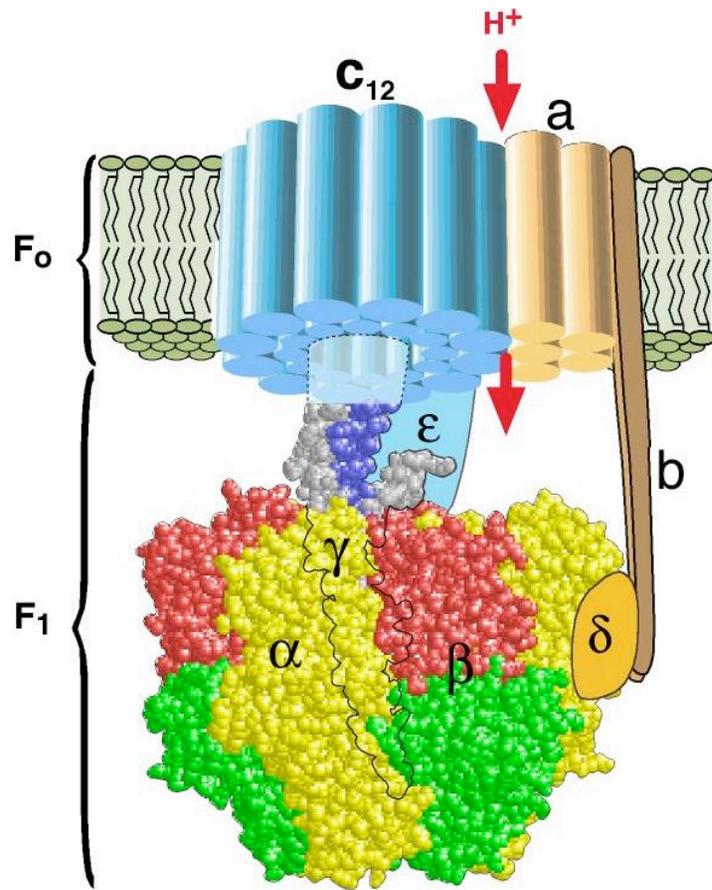
19th Century Thermodynamics



meters
kilojoules/cycle

21st Century Thermodynamics

Molecular Engines



ATP Synthase

Nanometers
Out-of-equilibrium
Large fluctuations
Dissipation $\sim kT$ per cycle

Equilibrium Statistical Mechanics

Probability of state i

$$\rho_i = \frac{1}{Z} e^{-E_i/k_B T} = e^{\beta F - \beta E_i}$$

Partition Function

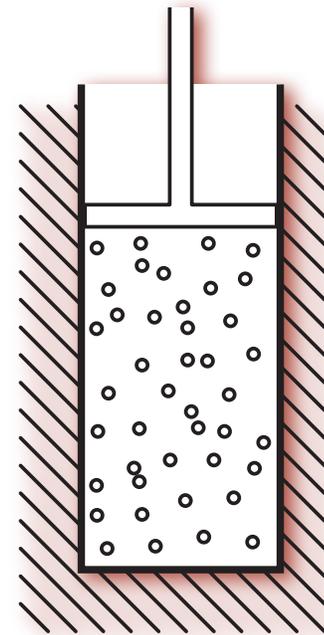
Boltzmann's Constant

Temperature

Free Energy

$\beta = 1/kT$

Energy of state i



This fundamental law is the summit of statistical mechanics, and the entire subject is either the slide-down from this summit, as the principle is applied to various cases, or the climb-up to where the fundamental law is derived... — Feynman

Out-of-Equilibrium

Probability of state i

$$\rho_i \neq \frac{1}{Z} e^{-E_i/k_B T} = e^{\beta F - \beta E_i}$$

Partition Function

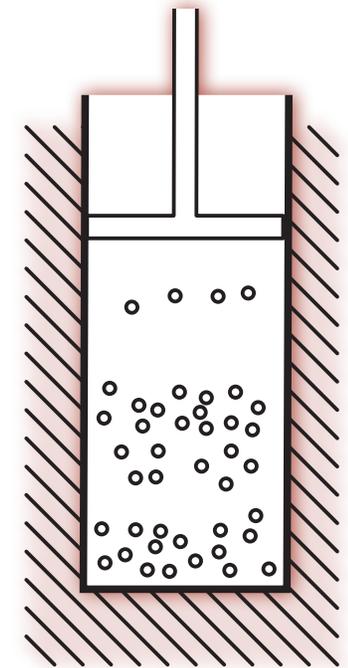
Boltzmann's Constant

Temperature

Free Energy

Energy of state i

$\beta = 1/kT$



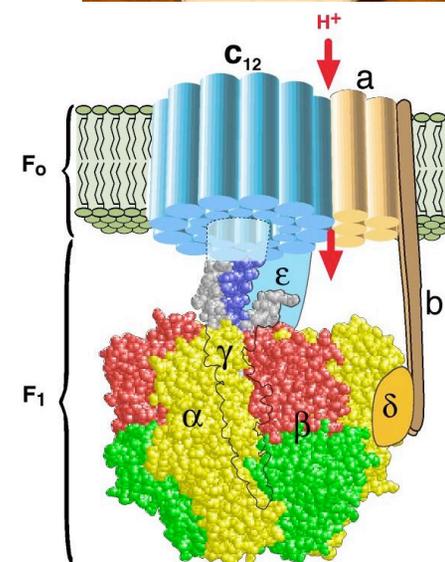
Dynamics around equilibrium is always more interesting than equilibrium itself
- John Doyle 2007

Information is physical — Rolf Landauer

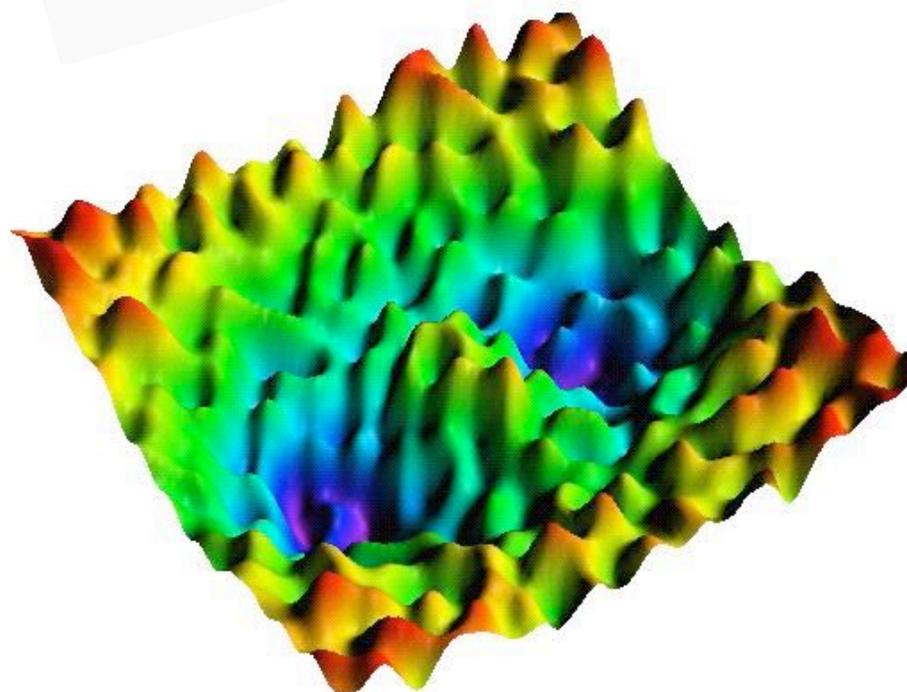
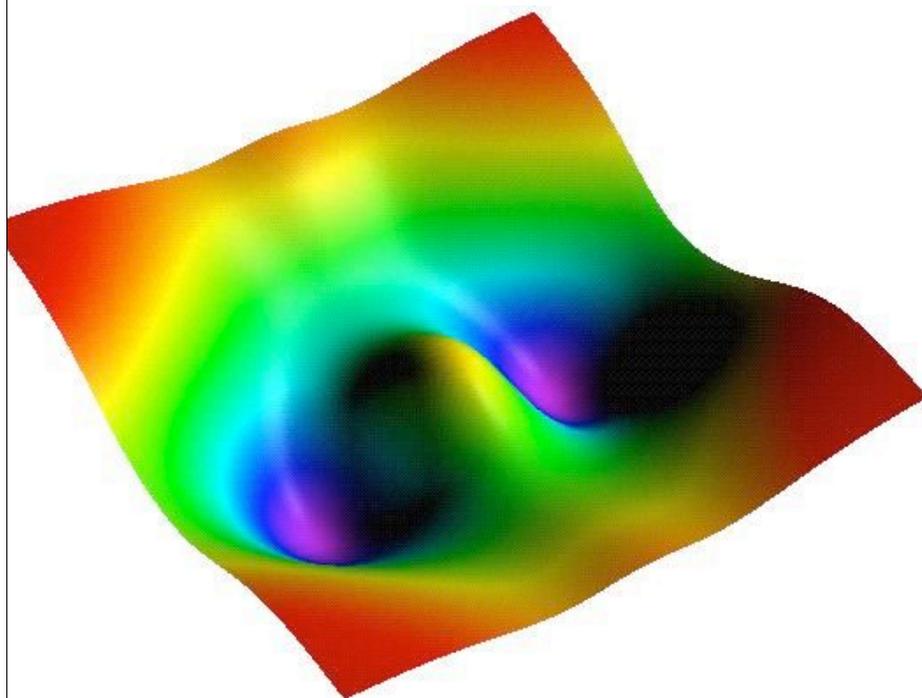
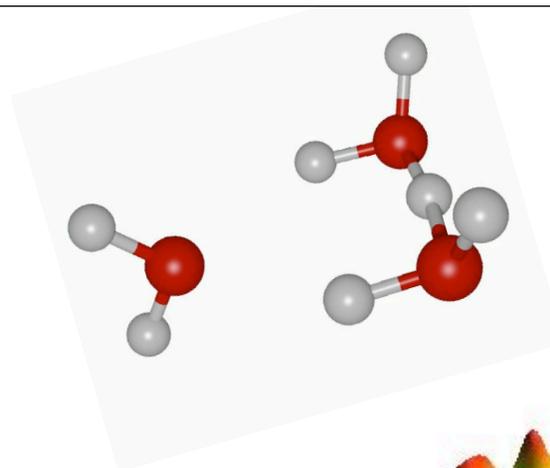
Information is bits

$$\begin{aligned}\Delta S_{\text{Melting}} &= 22 \text{ J / (mol K)} \\ &\sim 2.5 \text{ nats / molecule} \\ &\sim 4 \text{ bits / molecule}\end{aligned}$$

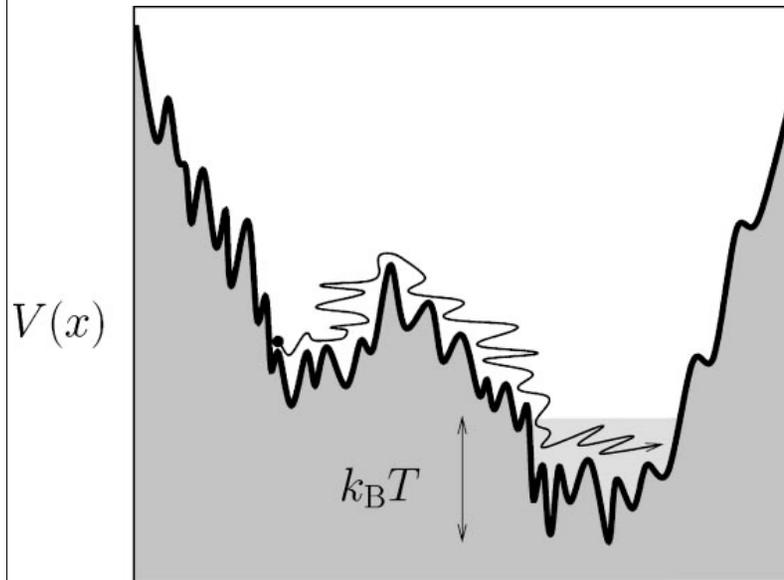
$$\begin{aligned}\Delta F_{\text{ATP Hydro.}} &= 20 \text{ kJ / mol} \\ \beta \Delta F_{\text{ATP Hydro.}} &\sim 8 \text{ nats / molecule} \\ &\sim 12 \text{ bits / molecule}\end{aligned}$$



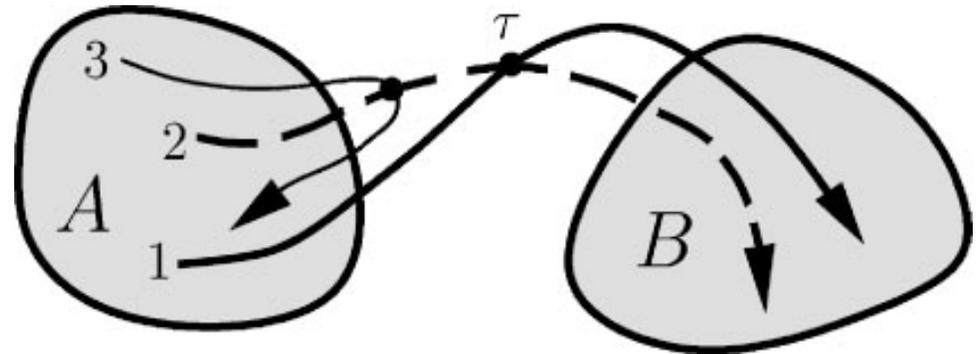
Transition Path Sampling



Transition Path Sampling

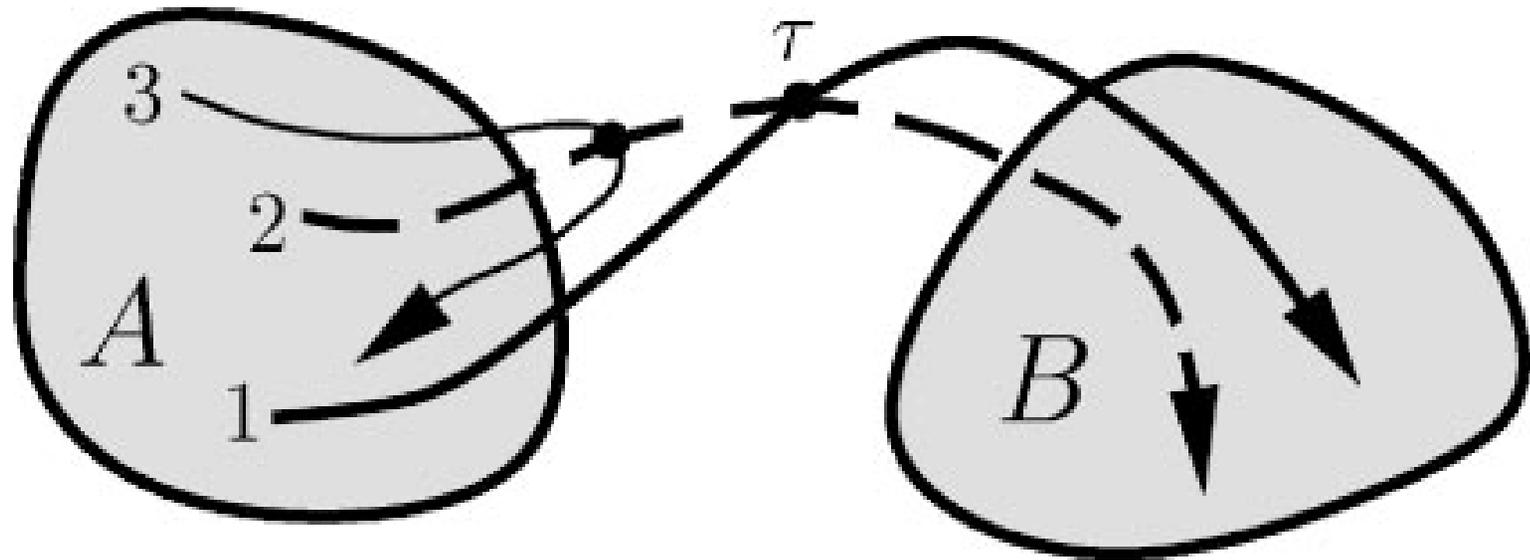


Configurational
Monte Carlo

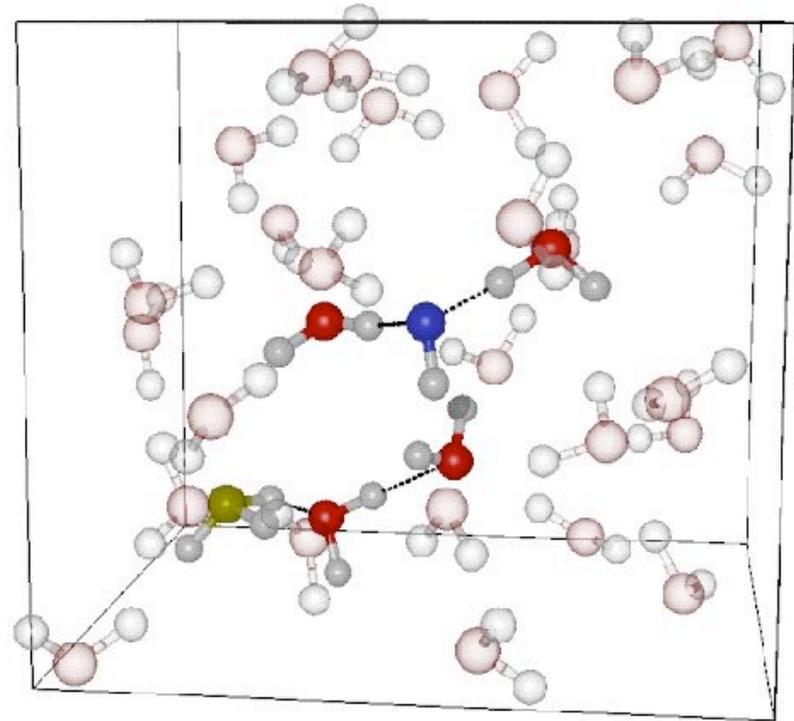
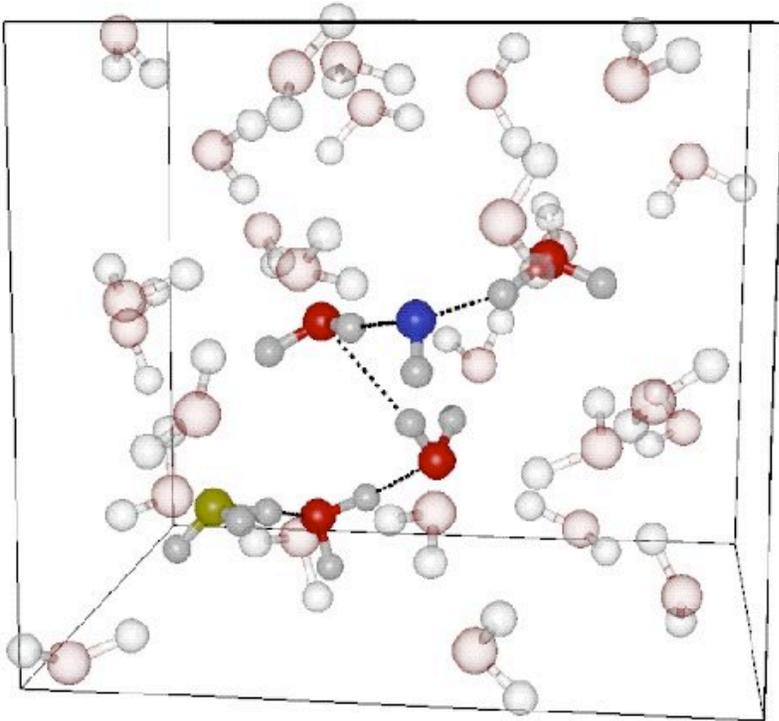


Transition Path
Monte Carlo

Transition Path Sampling



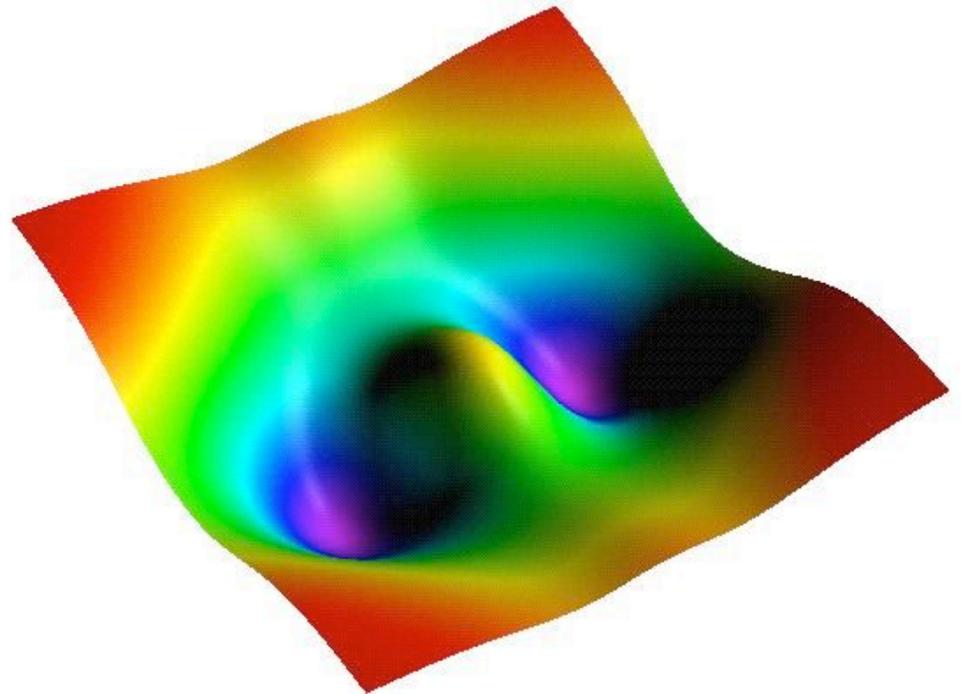
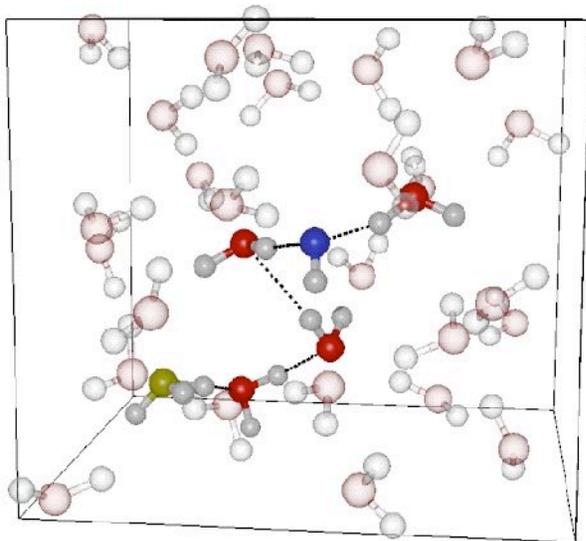
Water Autoionization



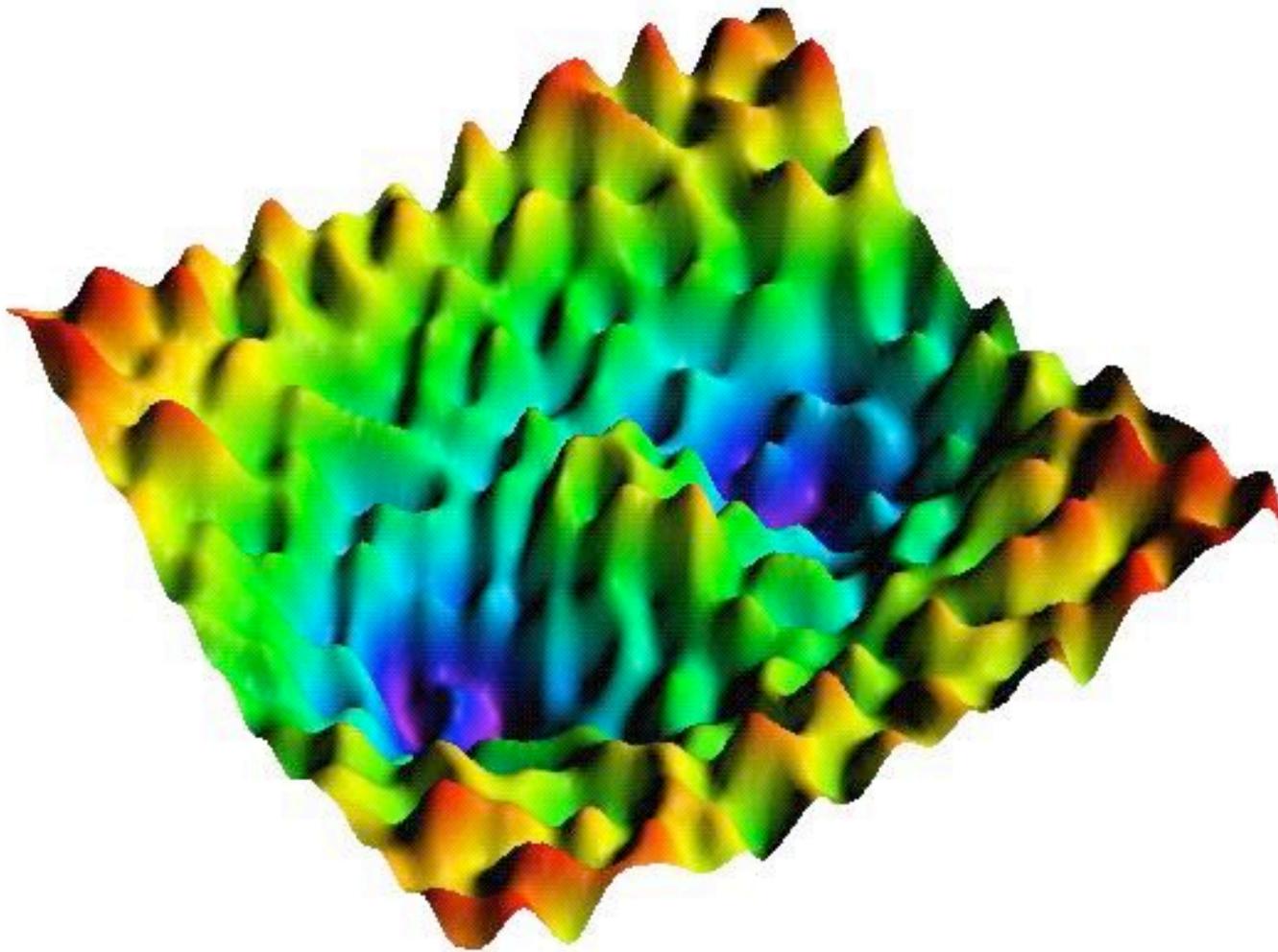
Geissler et al 2000

Open Problem:

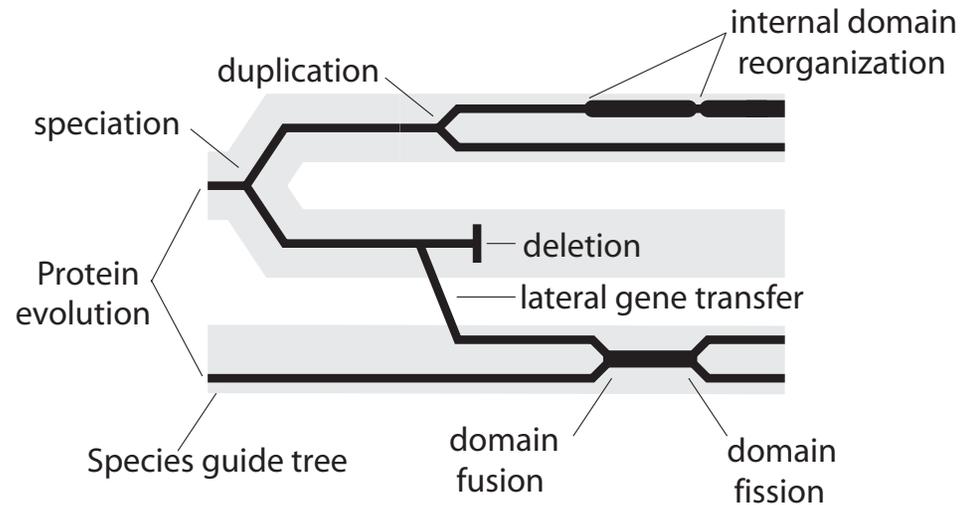
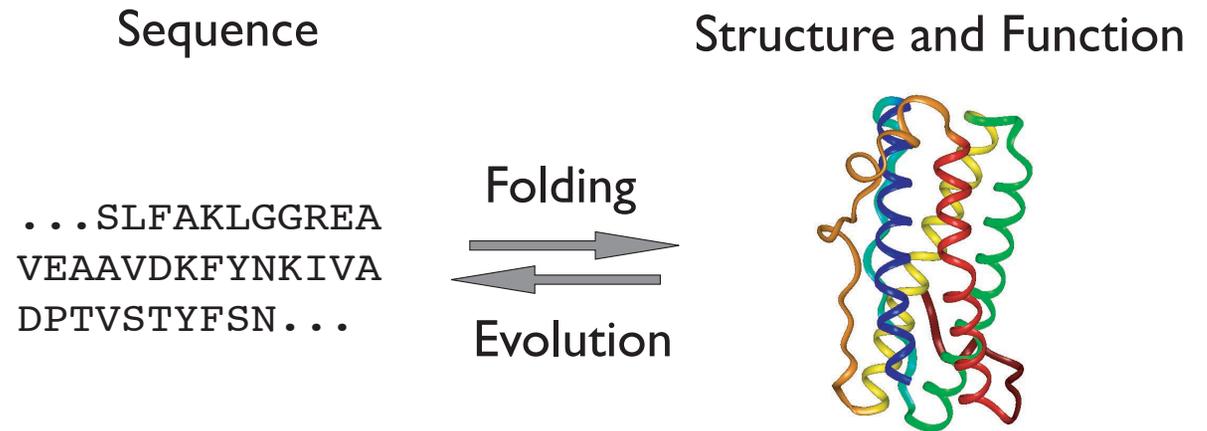
Assume we obtained a transition path ensemble between two conformations of a protein, how do we write a paper describing what happens? -Peter Bolhuis



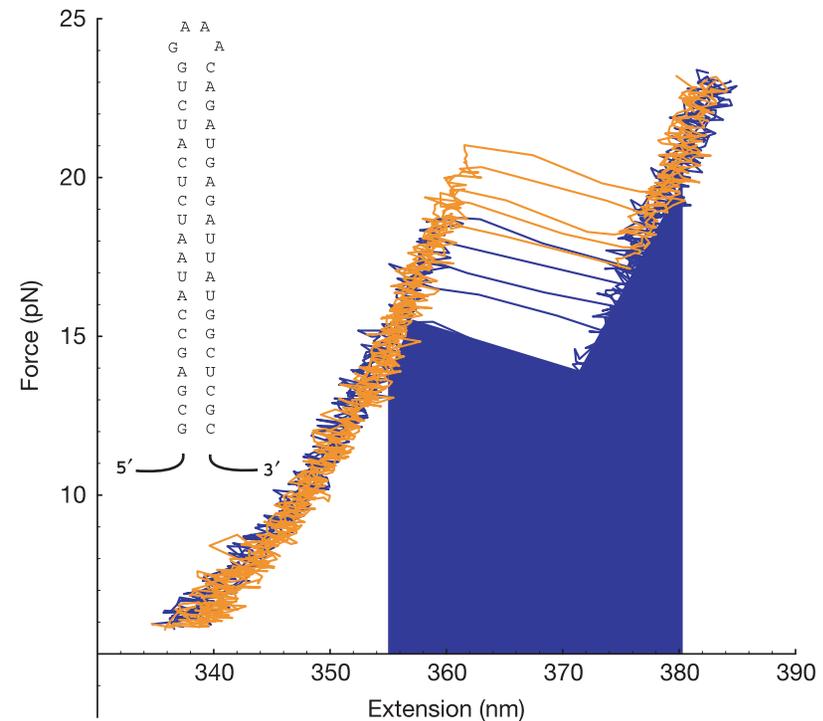
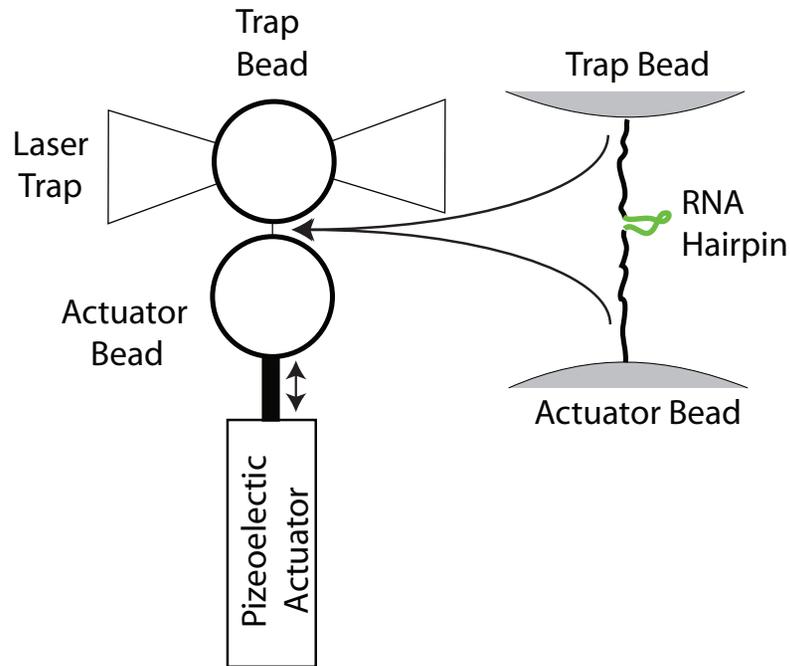
Open Problem : Intermediate Metastable states



Protein Evolution



RNA hairpin unfolding (& refolding)



J. Liphardt, S. Dumont, S.B. Smith, I. Tinoco, C. Bustamante (2002)
D. Collin, F. Ritort, C. Jarzynski, S.B. Smith, I. Tinoco, C. Bustamante (2005)

Clausius Inequality

(1865)

$$dS \geq \frac{1}{T} dQ$$

$$\Delta F \leq \langle W \rangle$$

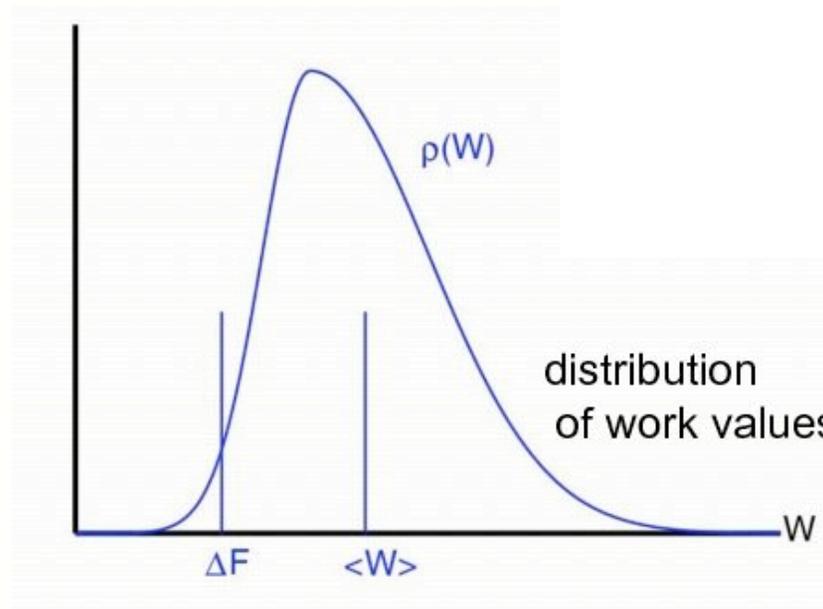
Free
Energy

Work

Jarzynski Equality

(1997)

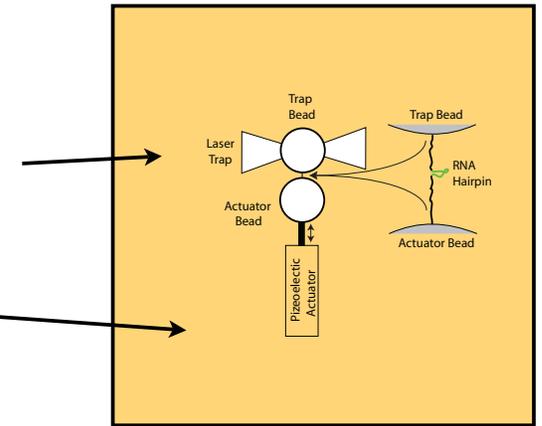
$$\langle e^{-\beta W} \rangle = e^{-\beta \Delta F}$$



Heat & Work

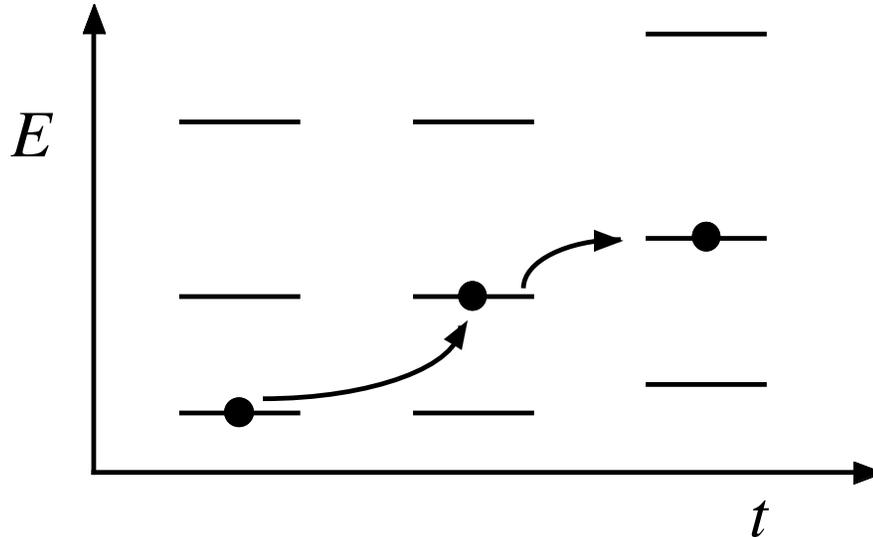
System of Interest

Thermal Environment



$$\Delta E = Q + W$$

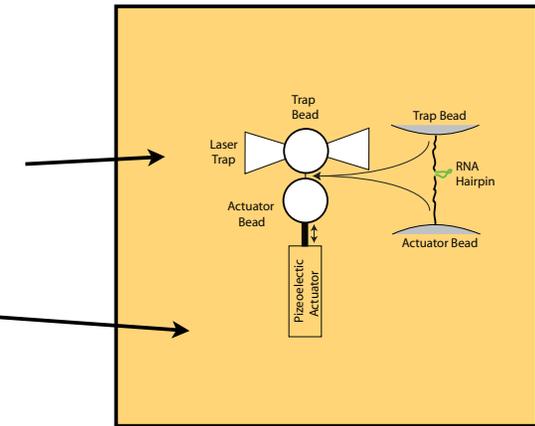
Internal Energy Heat Work



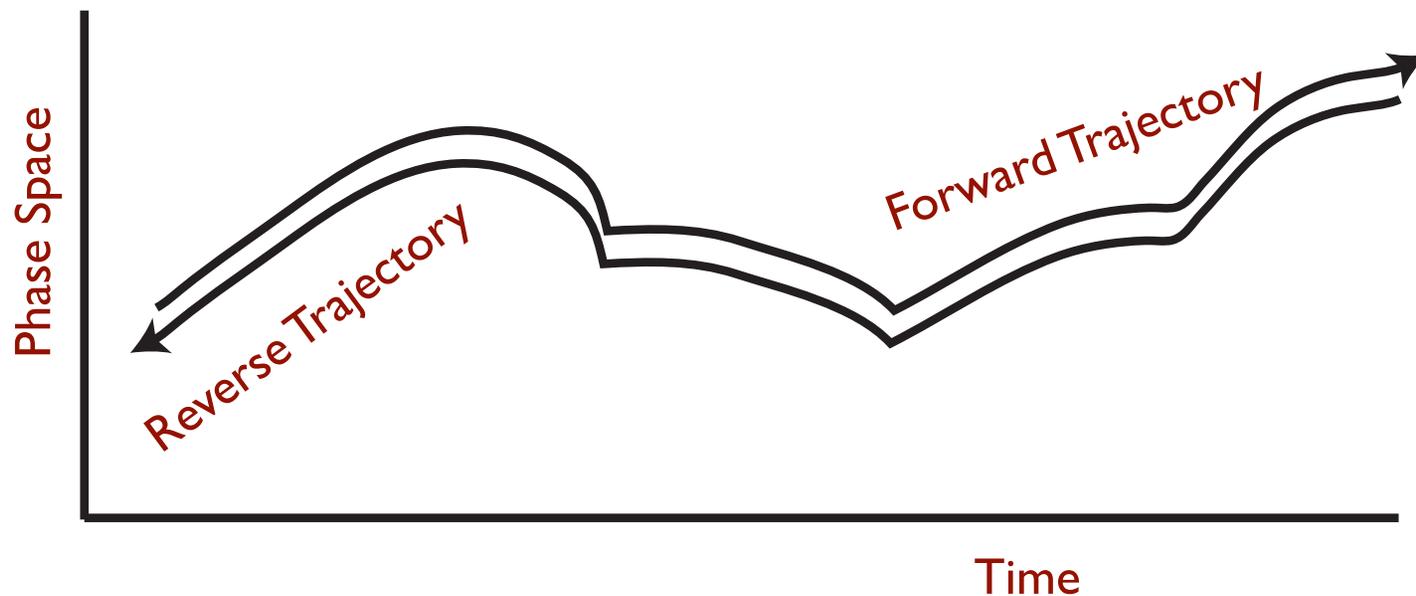
Time Reversibility

System of Interest

Thermal Environment



$$\frac{P(\text{Forward})}{P(\text{Reverse})} = e^{+\beta Q} \text{ Heat}$$



Work Fluctuation Theorem

Assumptions:

- 1) Equilibrium boundaries
- 2) Time-reversal symmetry

Forward protocol
(e.g. compression,
unfolding)

Work probability
distribution

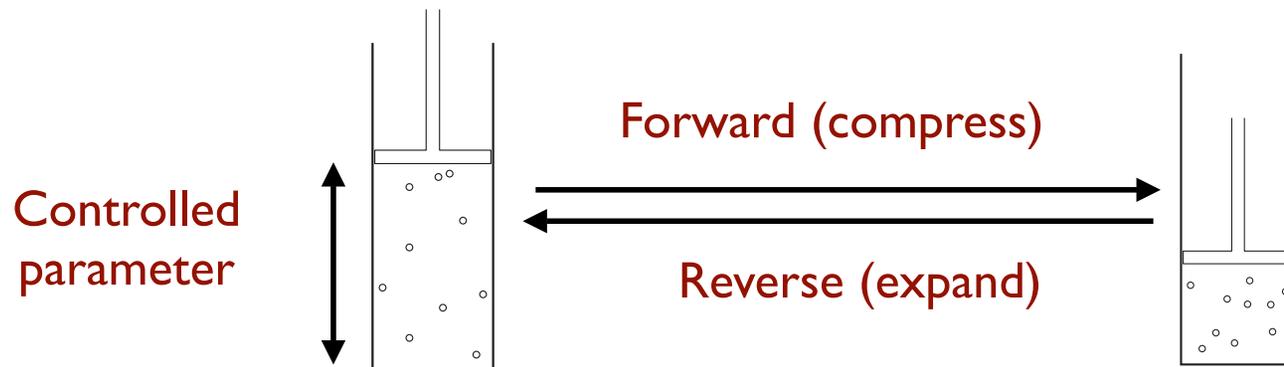
$$\frac{\rho_F(+W)}{\rho_R(-W)} = e^{\beta(W - \Delta F)}$$

Conjugate reverse protocol
(e.g., expansion, refolding)

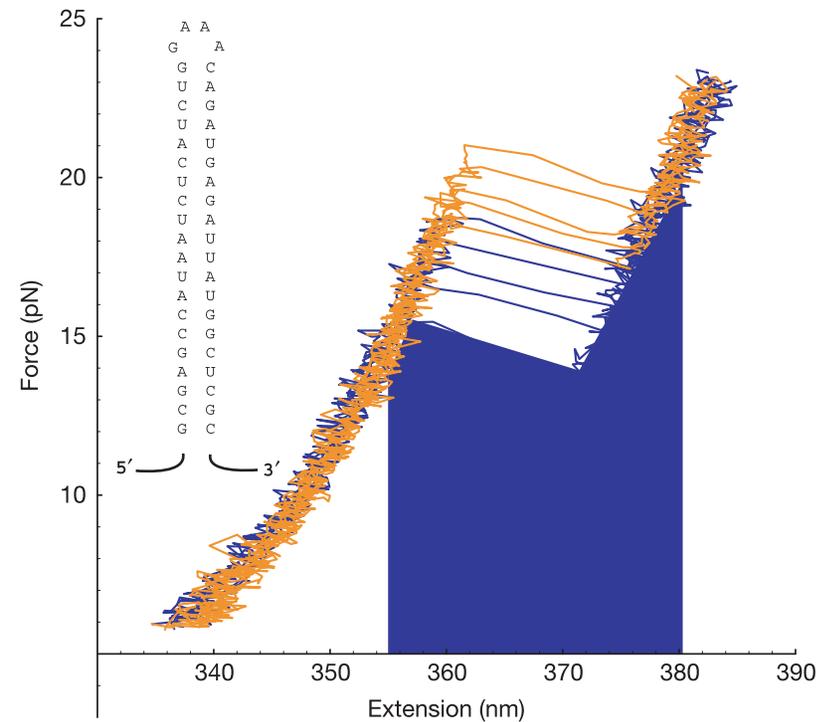
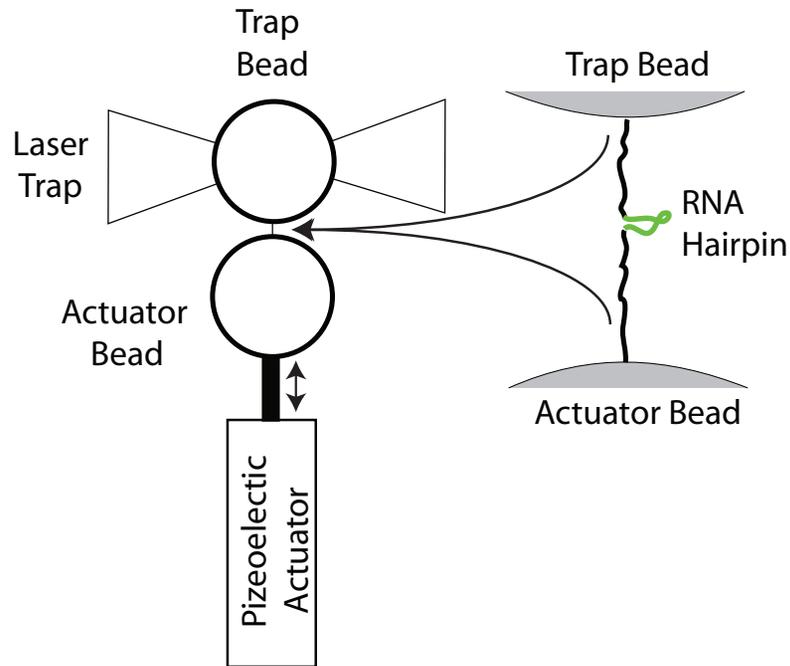
Work

Free Energy
Change

Inverse
Temperature



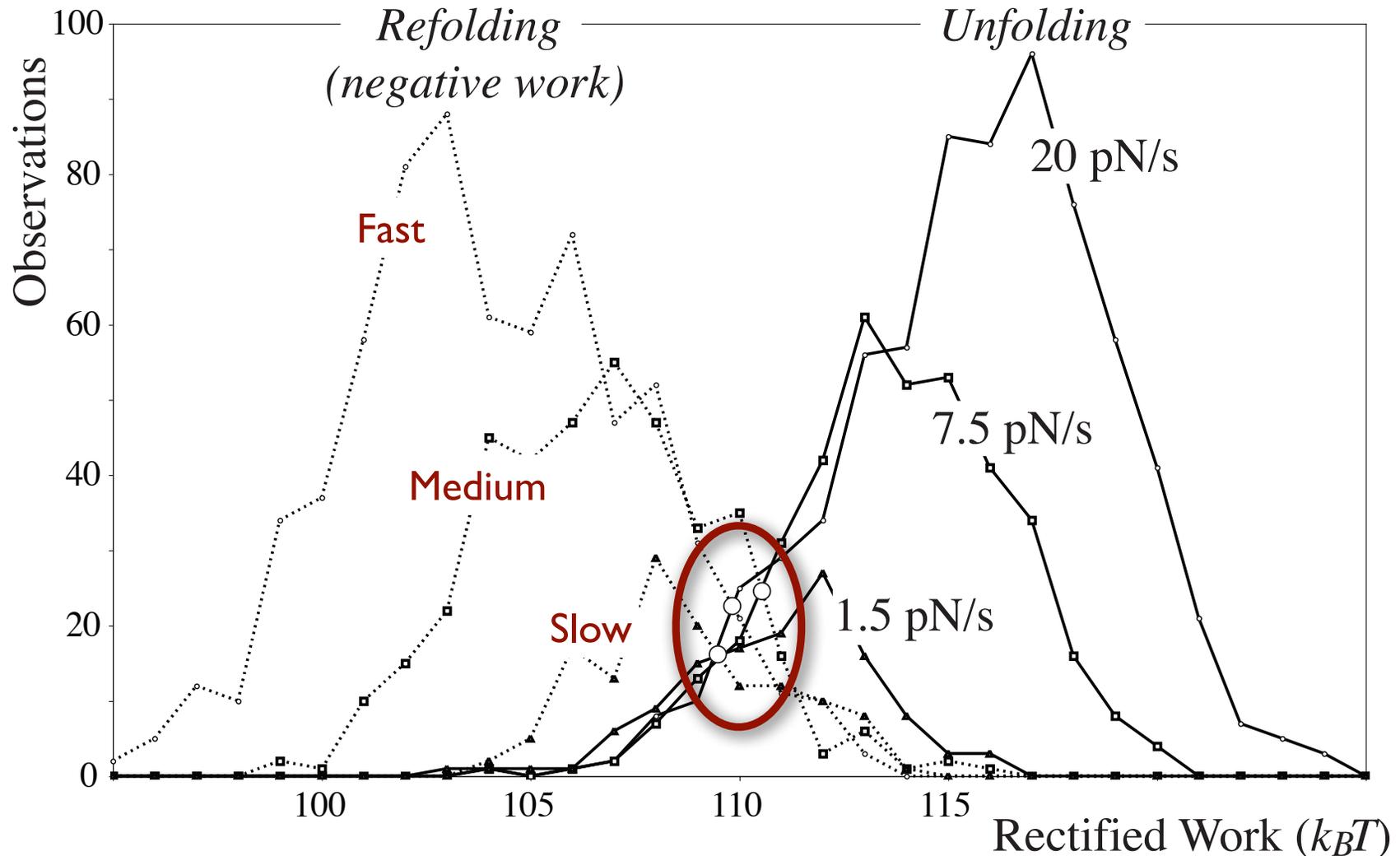
RNA hairpin unfolding (& refolding)



J. Liphardt, S. Dumont, S.B. Smith, I. Tinoco, C. Bustamante (2002)
D. Collin, F. Ritort, C. Jarzynski, S.B. Smith, I. Tinoco, C. Bustamante (2005)

Work of Unfolding

$$\frac{\rho_F(+W)}{\rho_R(-W)} = e^{\beta(W - \Delta F)}$$



Equilibrium Statistical Mechanics

Probability of state i

$$\rho_i = \frac{1}{Z} e^{-E_i/k_B T} = e^{\beta F - \beta E_i}$$

Partition Function

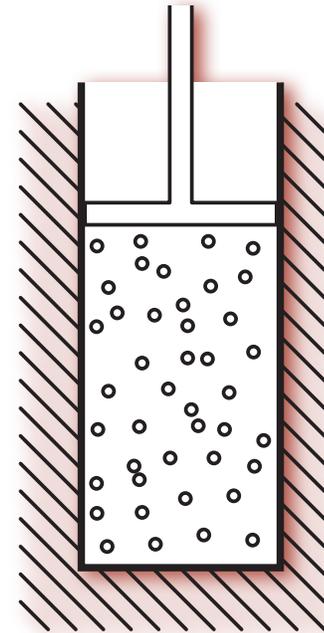
Boltzmann's Constant

Temperature

Free Energy

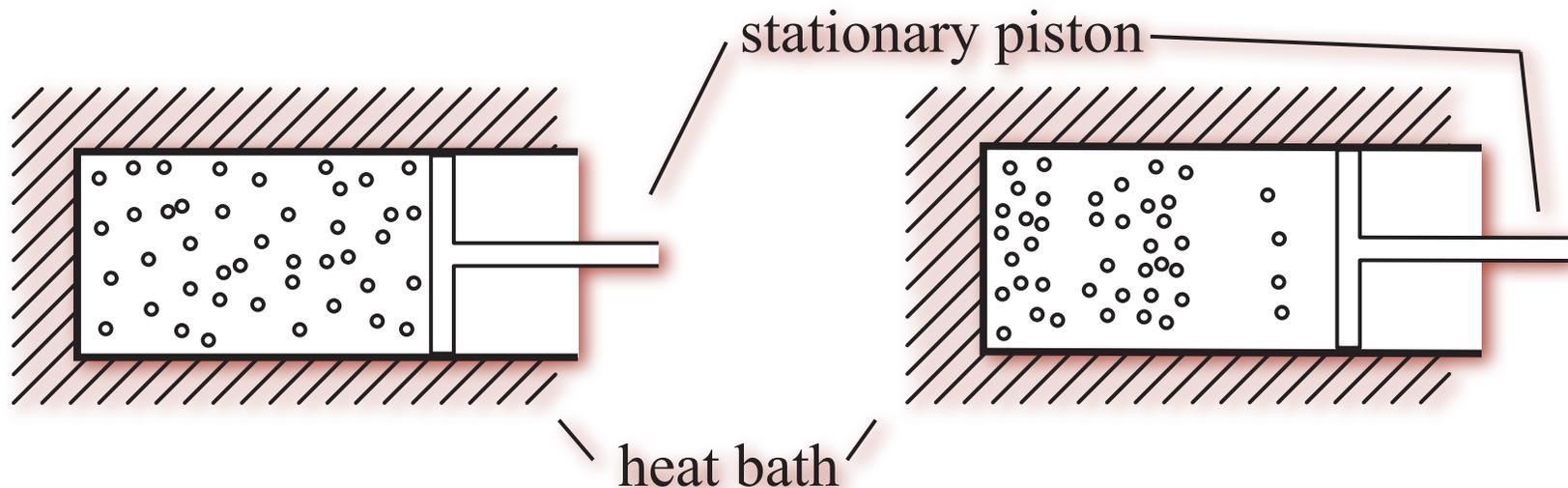
$\beta = 1/kT$

Energy of state i



Non-Equilibrium Statistical Ensembles

What is the **best description** that we can construct of a thermodynamic system that is **not in equilibrium**, given only **one** (or a few) extra parameters over and above those needed for a description of the same system at equilibrium?

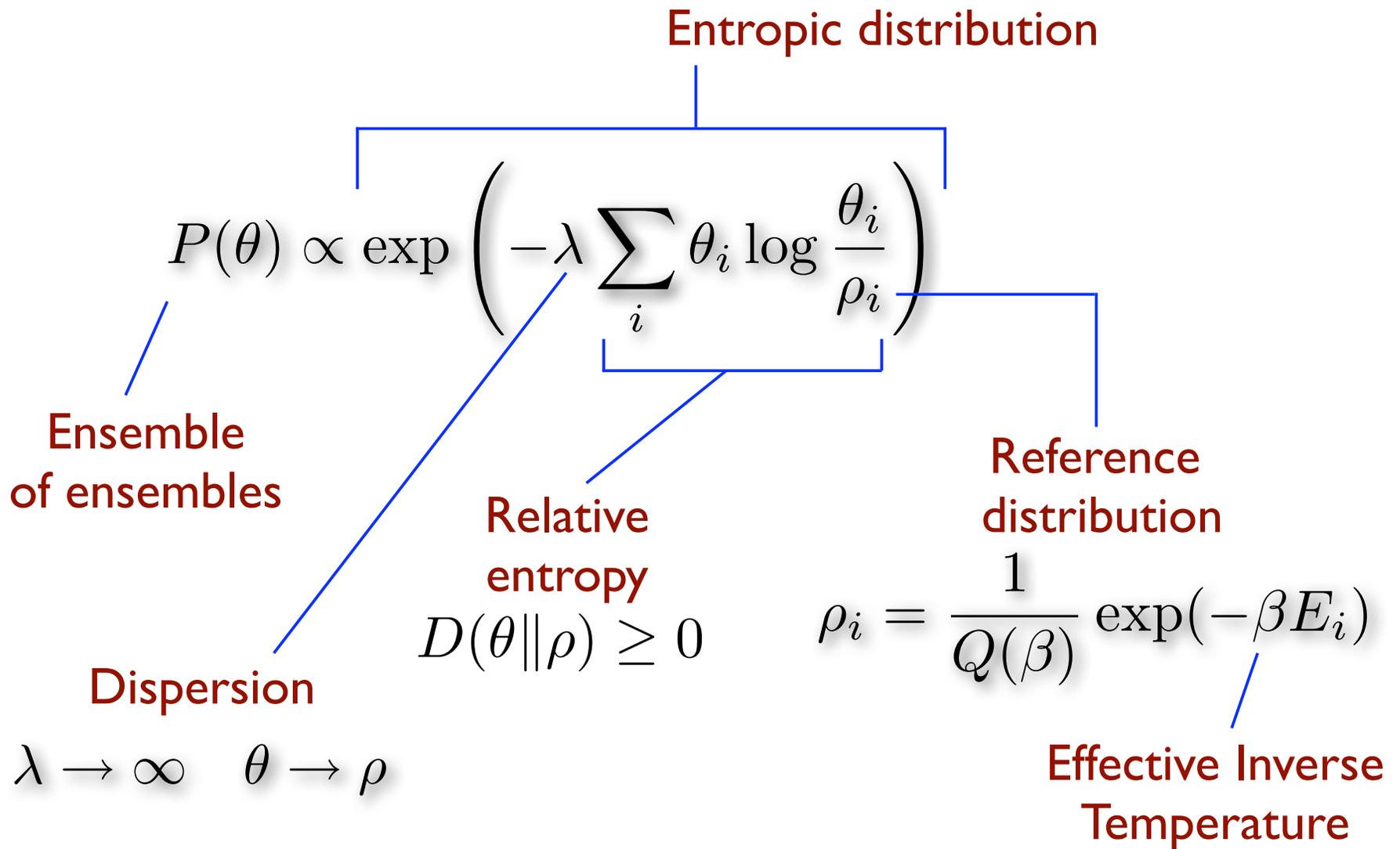


Non-Equilibrium Statistical Mechanics

What is the **best description** that we can construct of a thermodynamic system that is **not in equilibrium**, given only **one** (or a few) extra parameters over and above those needed for a description of the same system at equilibrium?

- Tsallis Statistics (Tsallis 1988) $S_q(\rho) = \frac{1}{q-1} (1 - \sum_i \rho_i^q)$
Maximize generalized q-entropy
- Superstatistics (Beck & Cohen 2002)

Canonical Hyperensemble



Open Problems

- Design principles of molecular machines.
- Transition path sampling with meta-stable intermediate states.
- Extracting meaningful reaction coordinates.
- Statistical ensembles out-of-equilibrium.
- Measuring non-equilibrium entropy.