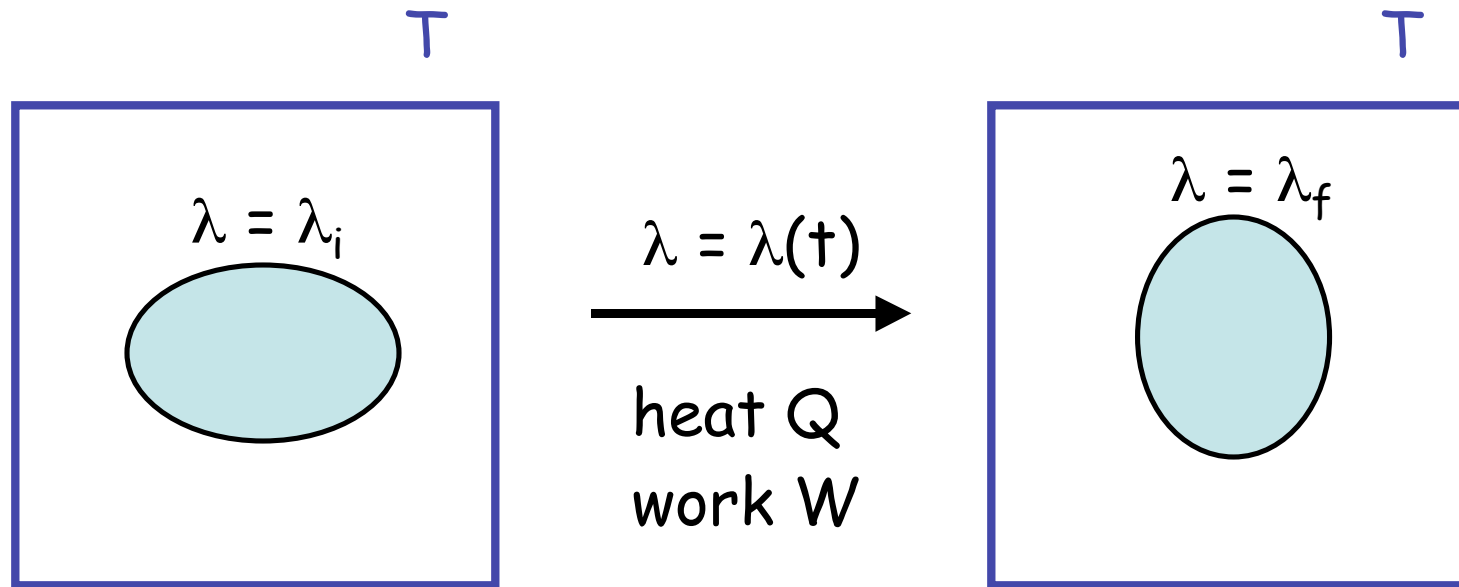


Free Energy Calculations

- Non-equilibrium SMD simulations
using Jarzynski equality
- Equilibrium MD simulations
Umbrella Sampling (WHAM)

Jarzynski Identity



2nd Law: $\langle W \rangle \geq \Delta F = F(\lambda_f) - F(\lambda_i)$

Jarzynski (1997): $\langle \exp(-\beta W) \rangle = \exp(-\beta \Delta F)$

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

$\langle W \rangle \geq \Delta F \quad \text{JE} \Rightarrow \text{2nd law}$

difficult to estimate

Derivation of Jarzynski Identity

- Hamiltonian systems:
energy conservation, Liouville's theorem
- Stochastic systems:
Markov property, balance condition

Stochastic algorithms in MD:

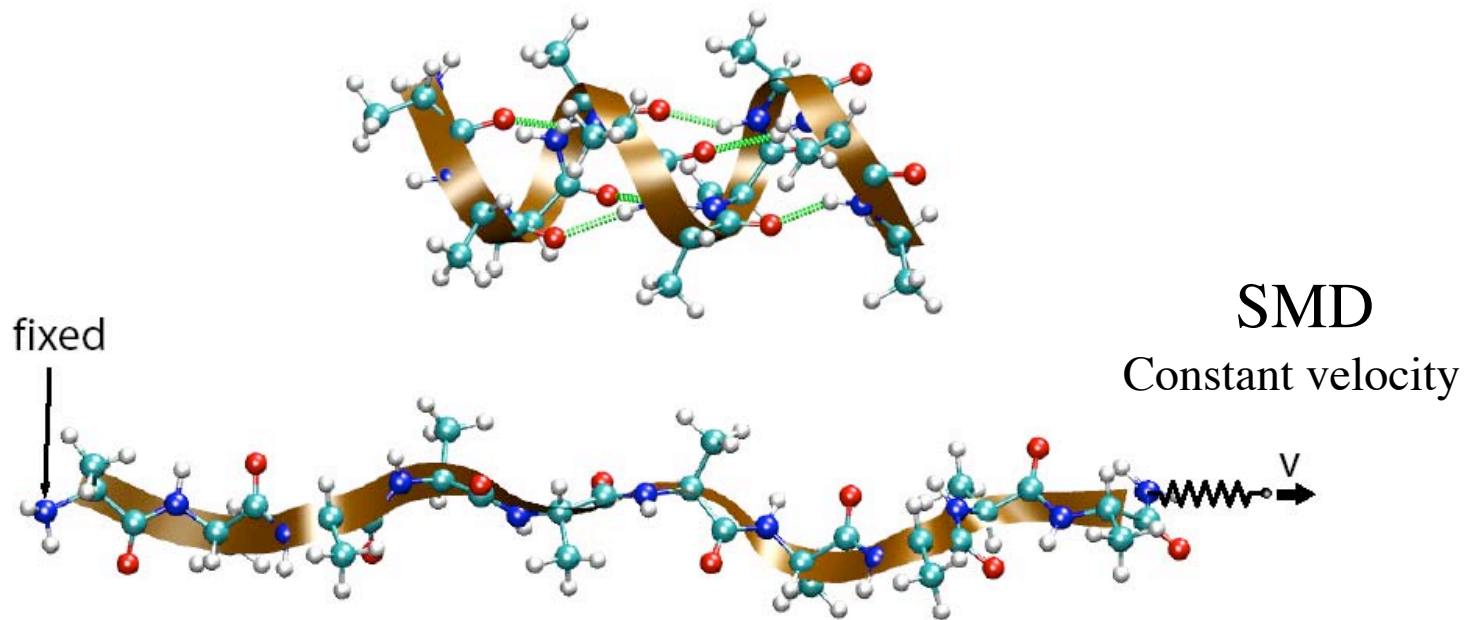
- NVT (Nose-Hoover)
- NpT (Langevin piston) (Gibbs free energy)

markovian (no History)

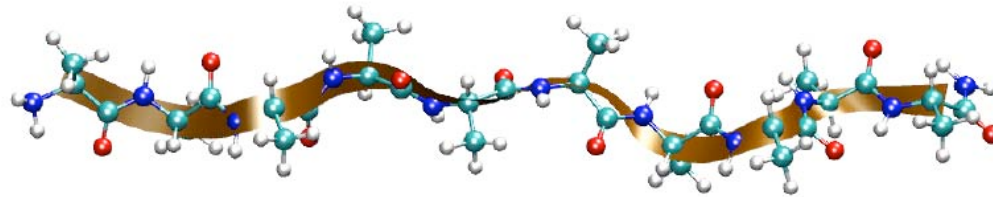
Balance Condition \Rightarrow Jarzynski Identity

Helix-Coil Transition of Deca-alanine in vacuum

- Small, but not too small: 104 atoms
- Short relaxation time \rightarrow reversible pulling \rightarrow exact PMF



A system with Hamiltonian H_0
 ϕ free energy along the reaction pathway



External potential: $h(r) = k(r - \lambda)^2/2$

$$H(r, p) = H_0(r, p) + h(r, \lambda)$$

$$\Delta F = - (1/\beta) \log \langle e^{-\beta W} \rangle$$

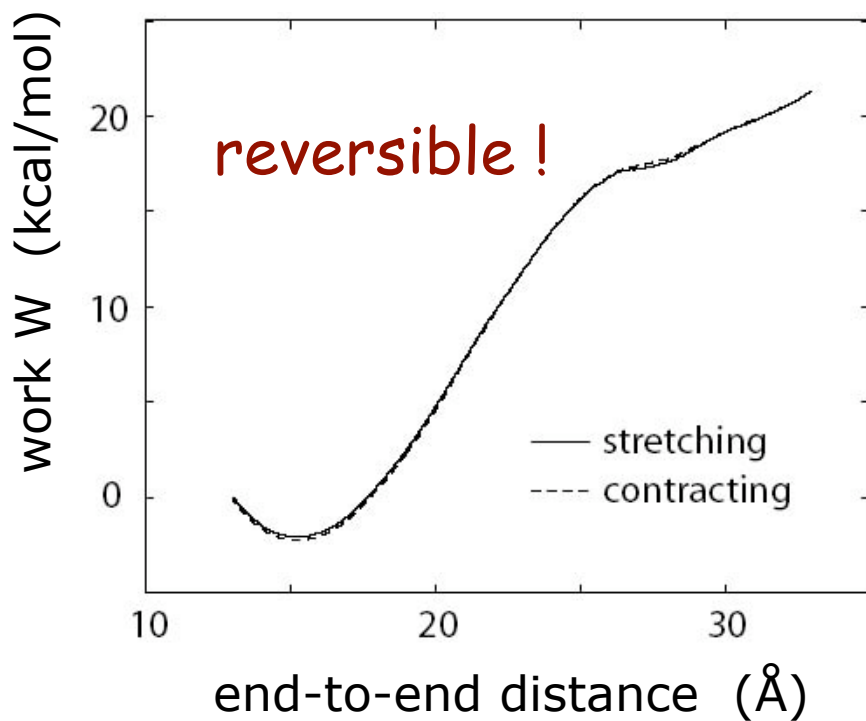
For stiff springs
(large spring constant)

$$\Delta F \approx \Delta \phi$$

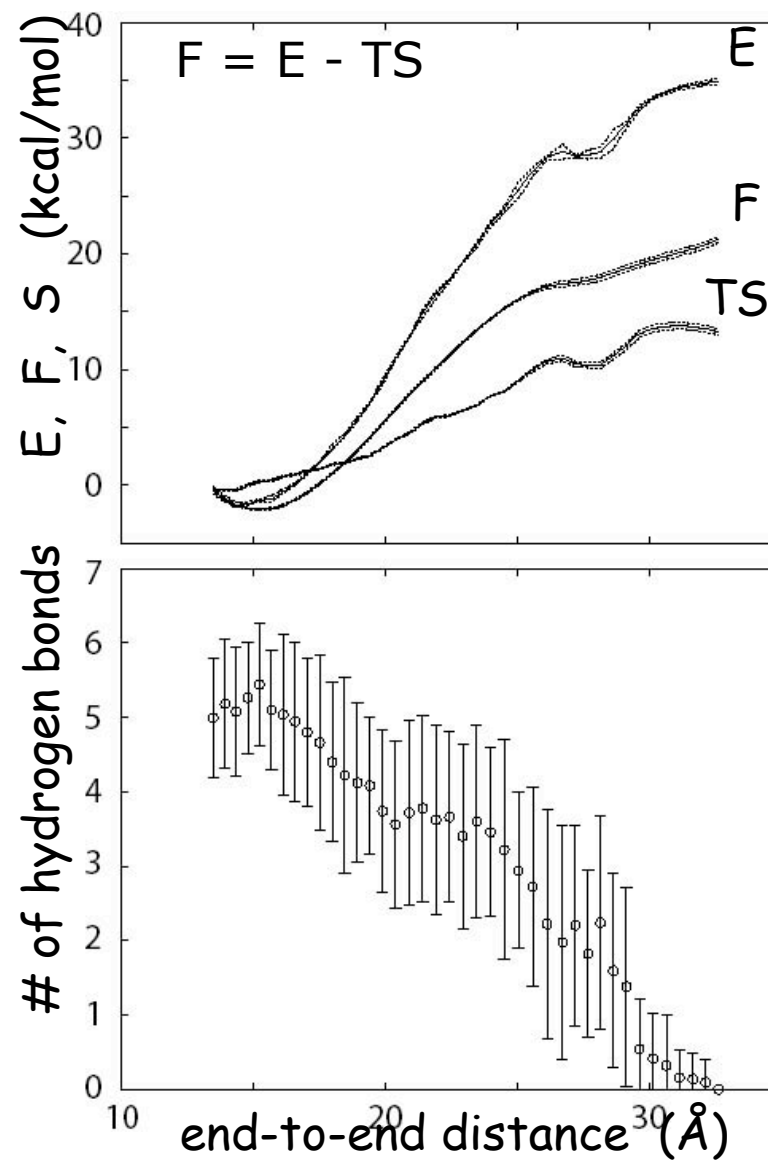
here we have used 500pN/A,
Covalent bond strength 3500pN/A **Stiff!**

Reversible Pulling ($v = 0.1 \text{ \AA/ns}$)

Park, Khalili-Araghi, Tajkhorshid & Schulten, J Chem Phys 119, 3559 (2003)



$$\Delta F = \langle W \rangle$$
$$TS = E - F$$



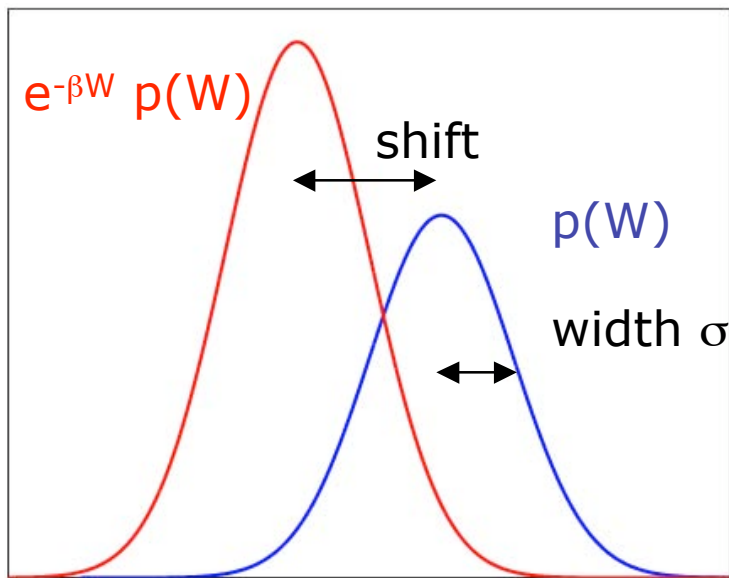
Sampling Error and Truncation Error

$$\Delta F = - (1/\beta) \log \langle e^{-\beta W} \rangle$$

$$= \langle W \rangle - (\beta/2) (\langle W^2 \rangle - \langle W \rangle^2)$$

$$+ (\beta^2/6) (\langle W^3 \rangle - 3\langle W^2 \rangle \langle W \rangle + 2\langle W \rangle^3) + \dots$$

Gaussian distributions



Example: Gaussian
 Shift = $\sigma^2 / k_B T$
 shift/width $\sim \sigma / k_B T$



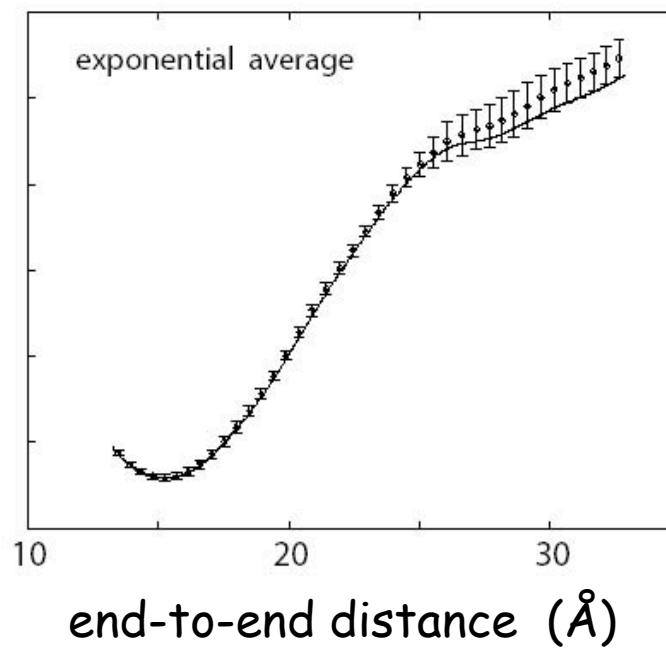
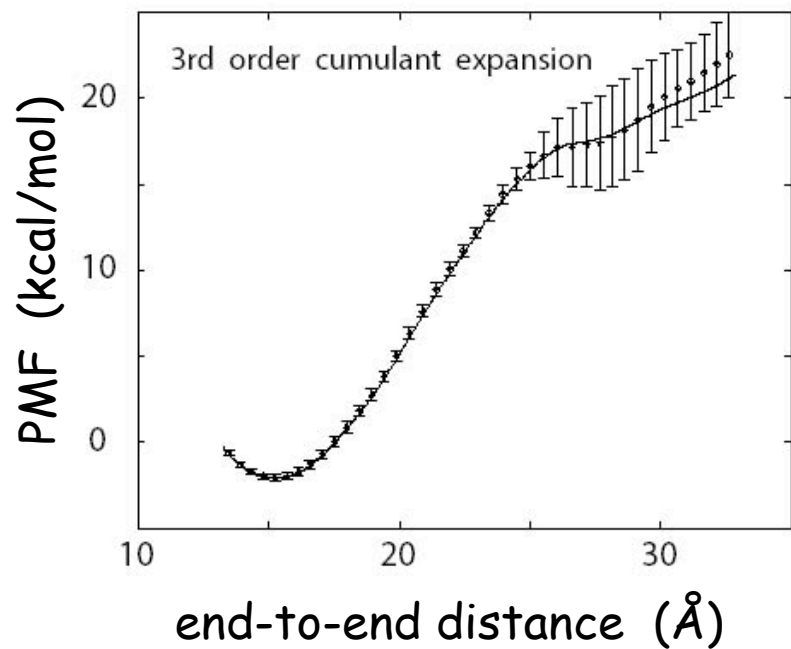
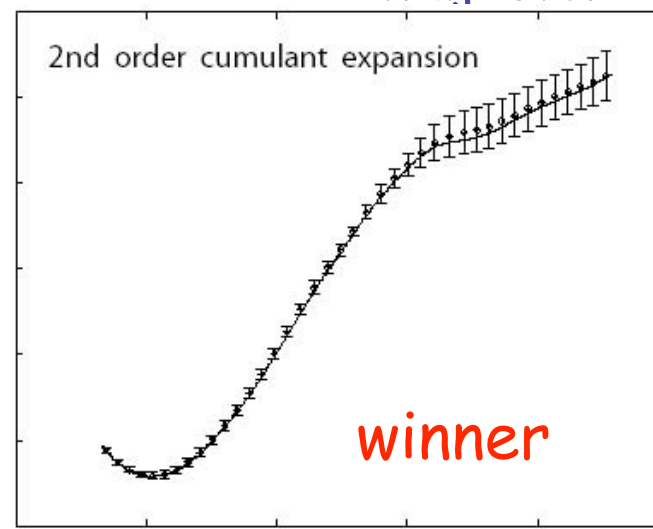
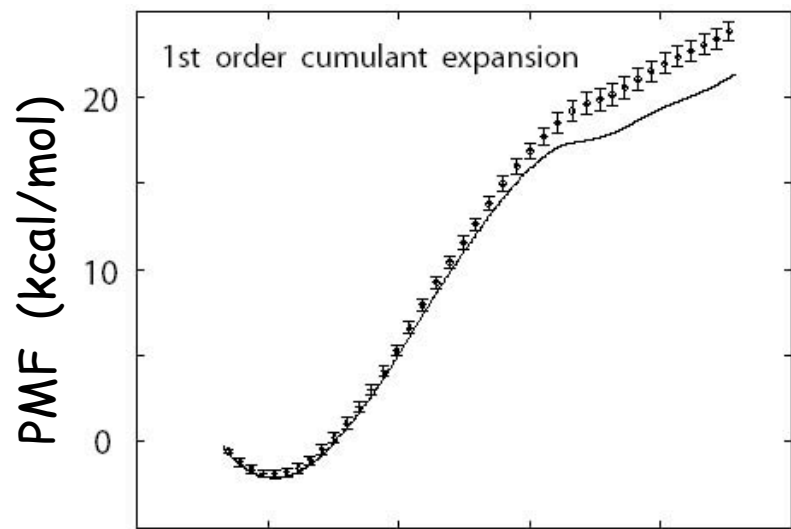
big in strong nonequilibrium

W

Biggest contribution to ΔF comes from small values of work, far from its average value, which requires ample sampling

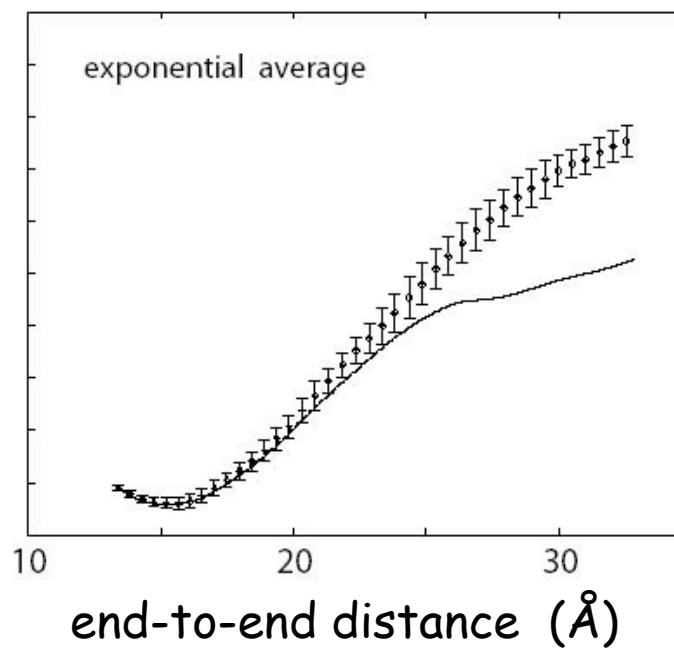
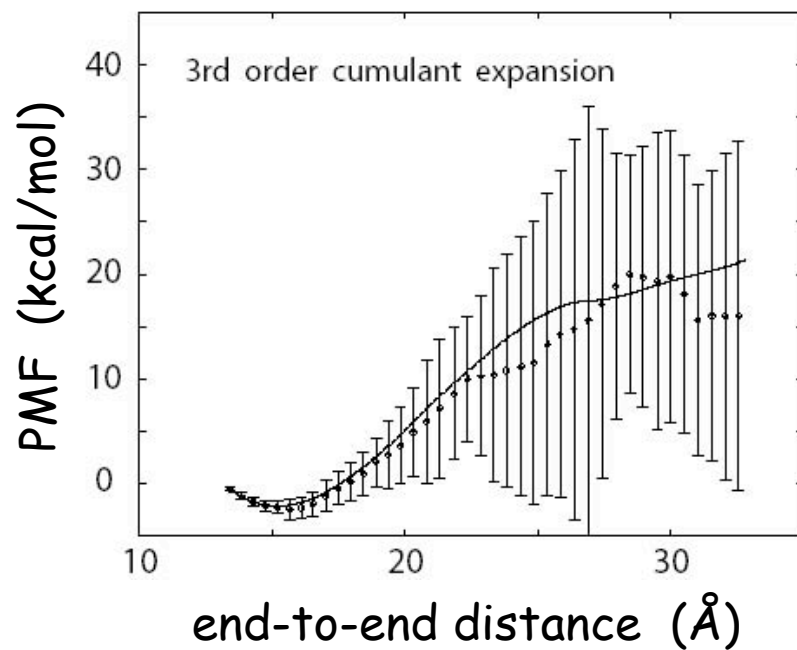
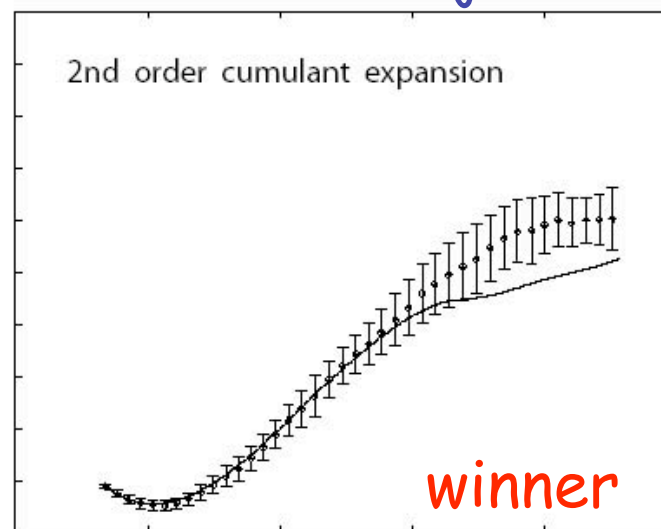
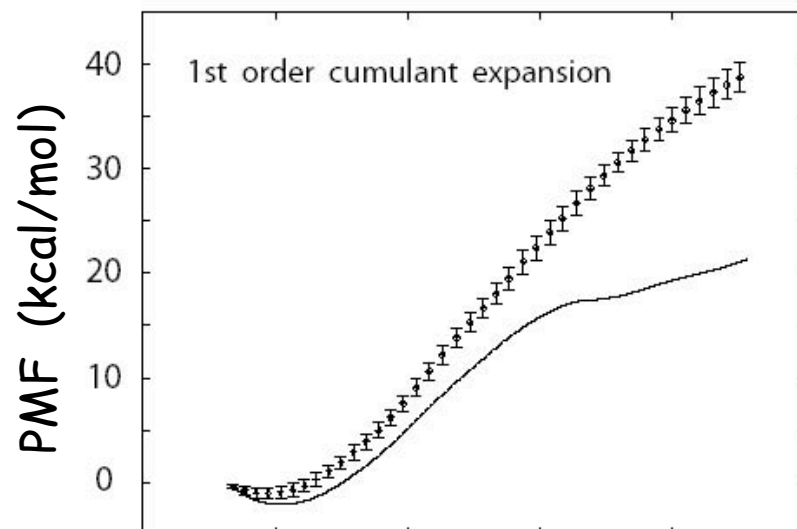
Irreversible Pulling ($v = 10 \text{ \AA/ns}$)

10 blocks of 10 trajectories



Irreversible Pulling ($v = 100 \text{ \AA/ns}$)

10 blocks of 10 trajectories



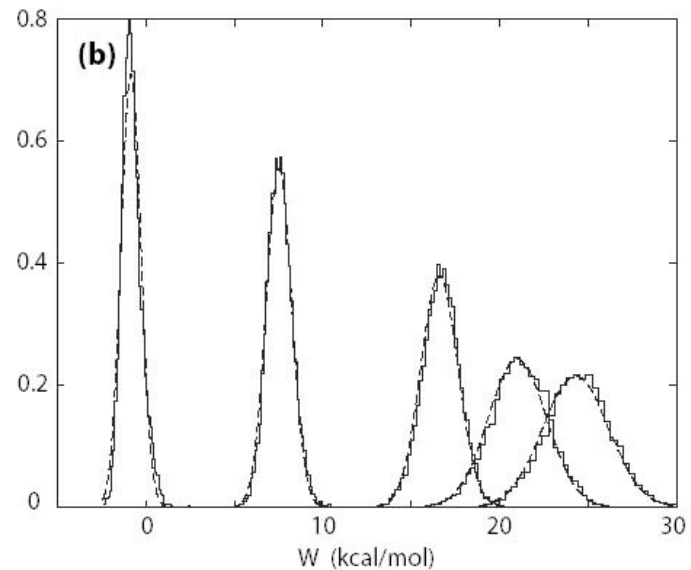
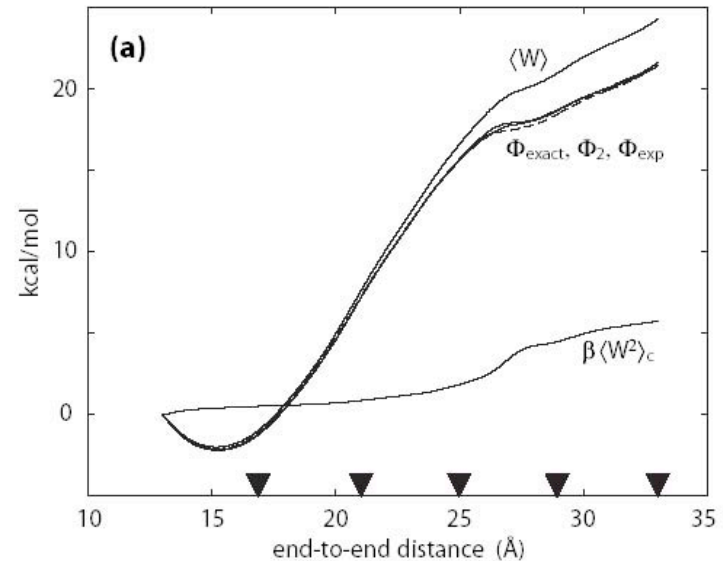
Guassian Work Distribution

- Langevin dynamics
- Harmonic guiding potential
- Stiff spring approximation

Guassian Work Distribution

$$v = 10 \text{ \AA/ns}$$

Park & Schulten, J Chem Phys, in press.



PMF from Umbrella Sampling

Equilibrium sampling of the configuration space.

Takes very long time

Confine the system to a small region, by applying a biasing potential

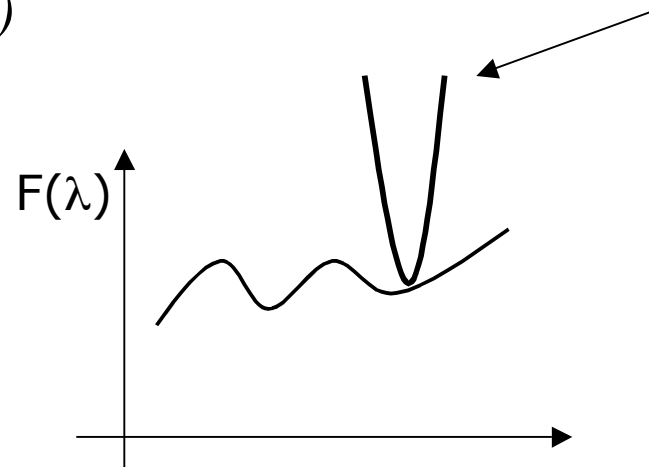
$$E_i = E_0 + V_i(x) \quad P_i(x) \sim e^{\beta V_i(x)} h_i(x) \\ = Z_i e^{-\beta H(x)}$$

h_i : Histograms built from equilibrium MD simulations

$$V_i(x) = k (x - x_i)^2 / 2$$

Choice of biasing harmonic potential

$$k \Delta x^2 \approx k_B T$$



WHAM

Weighted Histogram Analysis Method

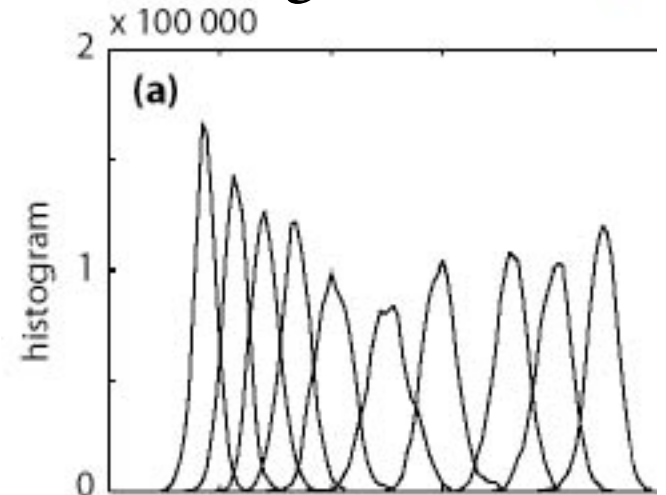
Histogram

Biassing potential => free energy difference in each region

Free energy in the overlapping regions
has to be matched.

$$P_0^{\text{est}} = \sum_i w_i(x) \exp[V_i / (k_B T)] Z_i / Z_0 P_i(x)$$

To minimize the statistical error, a **weight function** is used.



Weight function that minimizes the error:

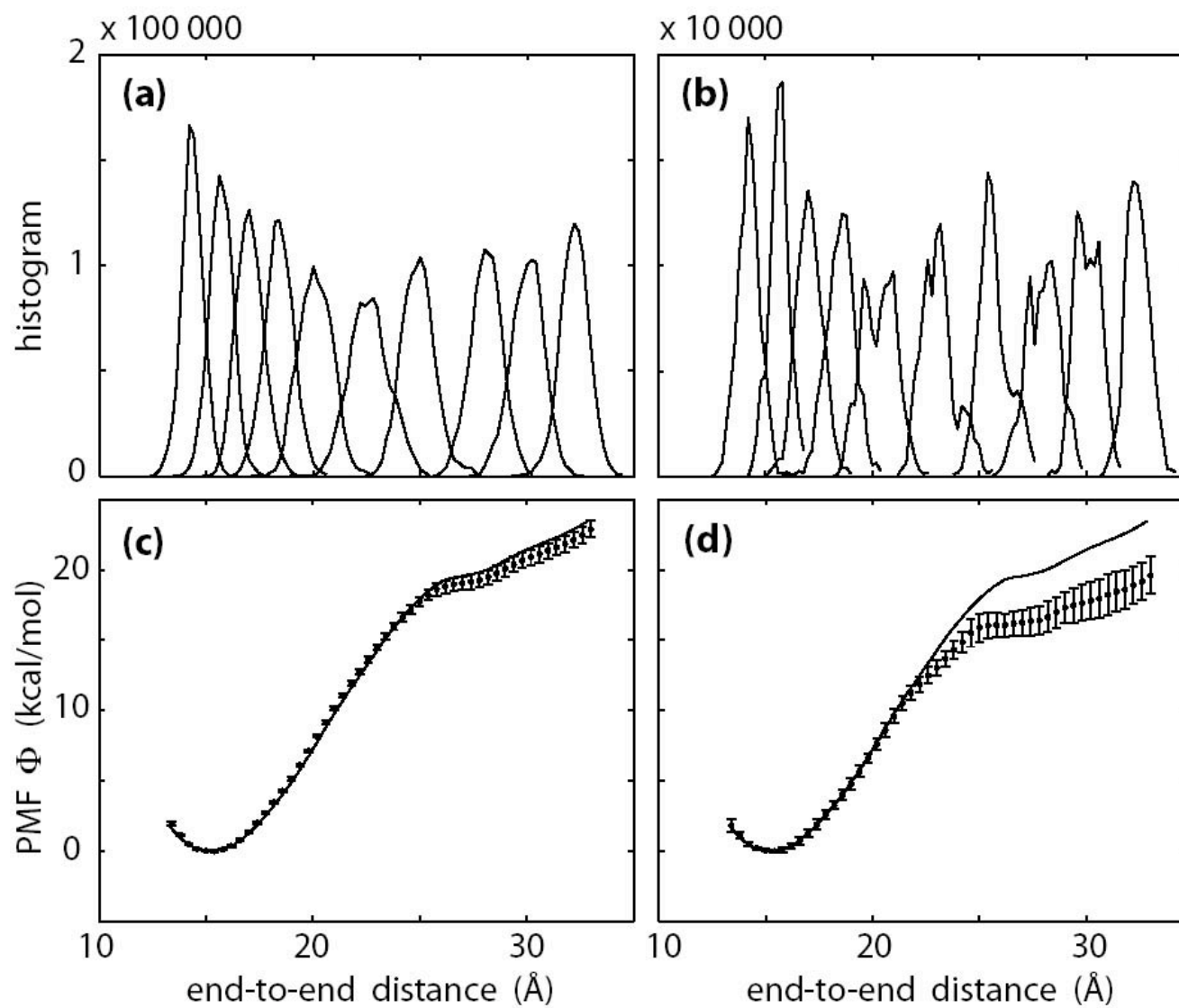
$$w_i(x) = [\exp(-V_i(x)/k_B T) Z_0/Z_i] / \sum_i \exp(-V_i(x)/k_B T) Z_0/Z_i$$

This results in a set of equations that has to be solved self consistently for Z_i

$$P_0^{\text{est}} = \sum_i w_i(x) \exp[V_i / (k_B T)] Z_i / Z_0 P_i(x)$$

$$F(x) = -k_B T \log (P_0^{\text{est}}(x))$$

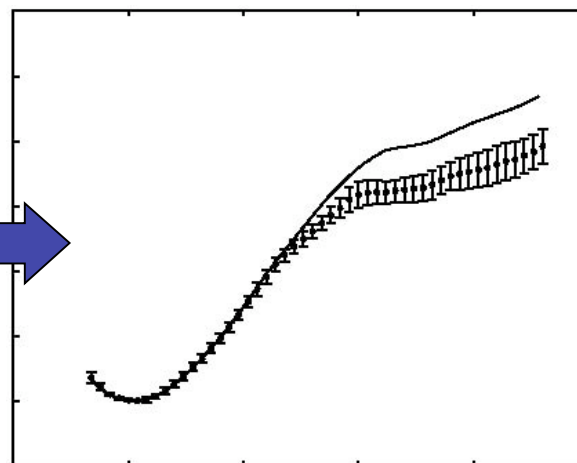
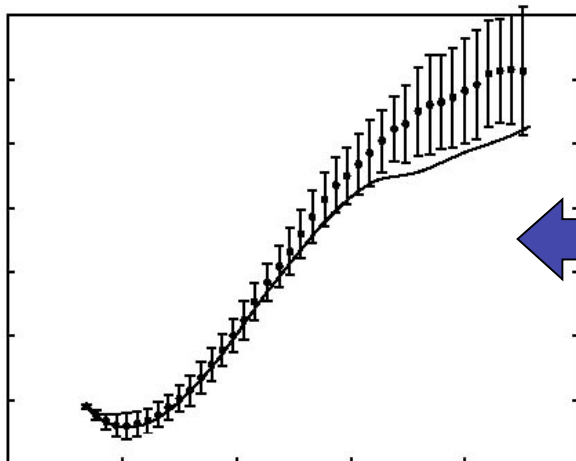
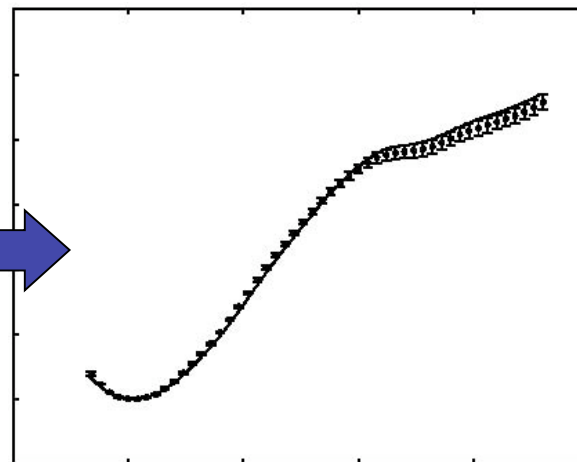
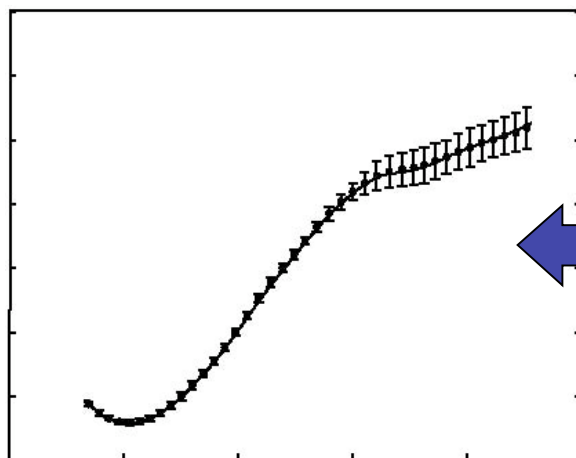
Umbrella Sampling w/ WHAM



SMD-Jarzynski

Umbrella Sampling

(equal amount of simulation time)



simple analysis

coupled nonlinear equations (WHAM)

uniform sampling of the reaction coordinate

nonuniform sampling of the reaction coordinate