Statistical Mechanics of Proteins

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Equilibrium and non-equilibrium properties of proteins

Free diffusion of proteins

- Coherent motion in proteins: temperature echoes
- Simulated cooling of proteins

Temperature Echoes in Proteins

- Coherent motion in proteins: Echoes
- Generation of echoes in *ubiquitin* via velocity reassignments
 - 1) Temperature quench echoes
 - 2) Constant velocity reassignment echoes
 - 3) Velocity reassignment echoes

temperature \Leftrightarrow velocities

kinetic temperature:

$$T(t) = \frac{2}{(3N-6)k_B} \sum_{n=1}^{3N-6} \frac{m_n v_n^2(t)}{2}$$

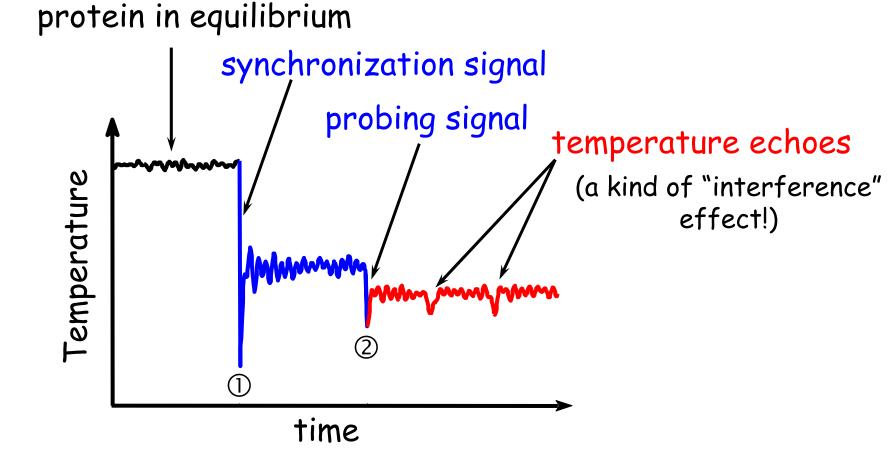
Coherent Dynamics of Proteins

- the internal dynamics of globular proteins comprises a wide range of time scales (10⁻¹⁵ – 1 sec)
- motions of ps (10⁻¹²s) time scale have some coherence, related to concerted motions of many atoms in different parts of the portein
- the coherence of proteins internal dynamics can be investigated via MD simulations by employing the temperature echo technique

<u>Questions</u>: What are, and how do we generate temperature echoes ?

Temperature Echoes

- are sharp, resonance-like features in the time evolution of the protein's temperature
- can be produced through 2 consecutive velocity reassignments



Velocity Reassignments

▶protein ≈ collection of weakly interacting harmonic oscillators having different frequencies

> at $t_1=0$ the 1st velocity reassignment: $v_i(0)=\lambda_1 u_i$ synchronizes the oscillators (i.e., make them oscillate in phase)

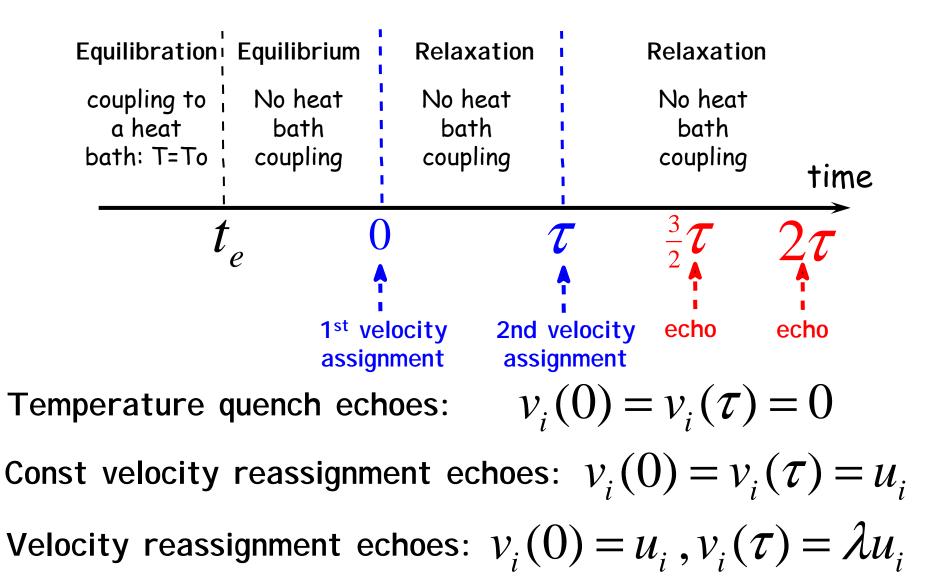
▶ at $t_2 = \tau$ (delay time) the 2nd velocity reassignment: $v_i(\tau) = \lambda_2 u_i$ probes the degree of coherence of the system at that moment

- degree of coherence is characterized by:
- the time(s) of the echo(es)

- the depth of the echo(es)

$$\begin{array}{l} \lambda_1 = \lambda_2 = 0 \implies \text{temperature quench} \\ \lambda_1 = \lambda_2 = 1 \implies \text{constant velocity reassignment} \\ \lambda_1 \neq \lambda_2 \neq 1 \implies \text{velocity reassignment} \end{array}$$

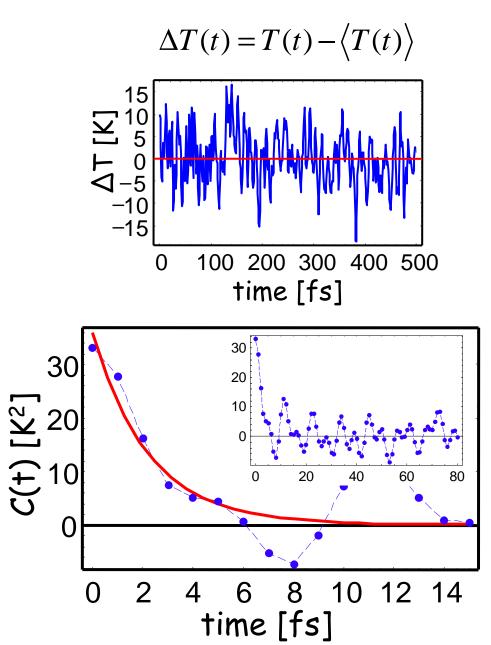
Producing Temperature Echoes by Velocity Reassignments in Proteins



Generating T-Quench Echo: Step1

	 your system is ubiquitin (1UBQ) in vacuum, pre-equilibrated at T₀=300K run all simulations in the <i>microcanonical</i> (NVE) ensemble psf, pdb and starting binary coordinate and velocity files are available in "common/" use NAMD2 configuration file "equil.conf" located in "01_equil_NVE/" to complete a 500 fs (# simulation steps) run extract the temperature time series T(t) from the NAMD2 log (output) file plot T(t) calculate: ⟨T⟩, √⟨T²⟩, C_{TT} = ⟨δT(t) δT(0)⟩
C	

Temperature Autocorrelation Function



$$C(t) = \left\langle \Delta T(t) \, \Delta T(0) \right\rangle$$
$$\rightarrow C(t_i) \approx \frac{1}{N-i} \sum_{n=1}^{N-i} \Delta T(t_{n+i}) \, \Delta T(t_n)$$

$$C(t) = C(0) \exp\left(-t/\tau_0\right)$$

Temperature relaxation time:

 $\tau_0 \approx 2.2 \, fs$

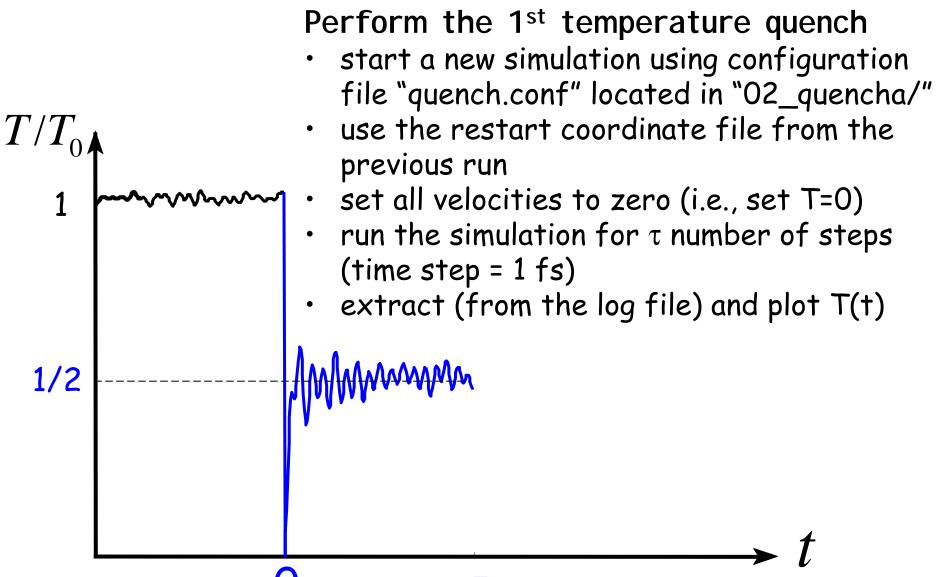
Mean temperature:

 $\langle T \rangle = 299 \, K$

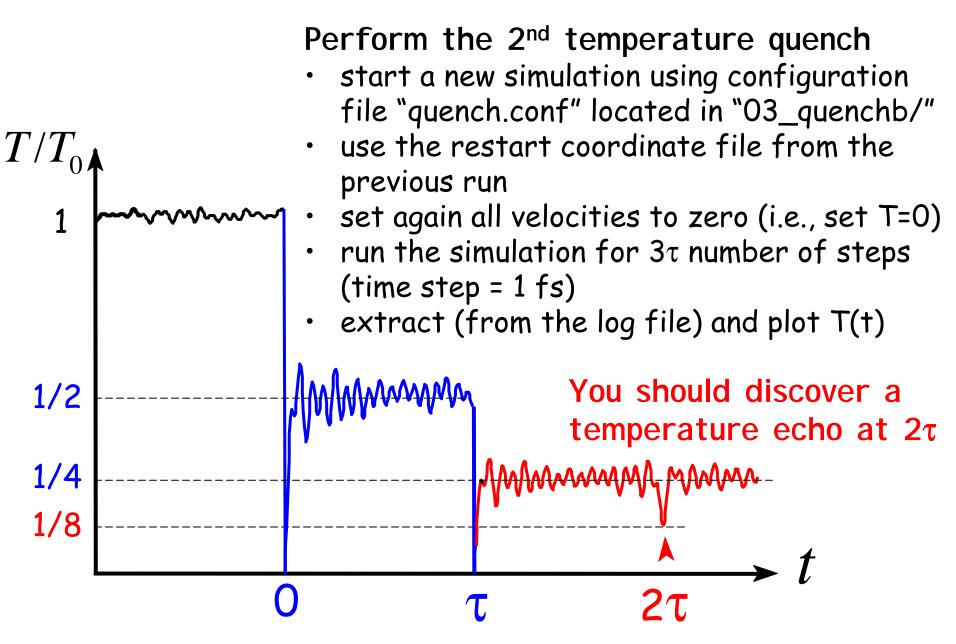
RMS temperature:

$$\sqrt{\left\langle \Delta T^2 \right\rangle} = \sqrt{C(0)} = 6 \, K$$

Generating T-Quench Echo: Step2



Generating T-Quench Echo: Step3



Explanation of the T-Quench Echo

<u>Assumption</u>: protein \approx collection of weakly interacting harmonic oscillators with dispersion $\omega = \omega_{\alpha}$, $\alpha = 1, ..., 3N - 6$

Step1:
$$t < 0$$
 $x(t) = A_0 \cos(\omega t + \theta_0)$
 $v(t) = -\omega A_0 \sin(\omega t + \theta_0)$

Step2:
$$0 < t < \tau$$

 $x_1(t) = A_1 \cos(\omega t + \theta_1)$
 $v_1(t) = -\omega A_1 \sin(\omega t + \theta_1)$
 $\xrightarrow{v_1(0)=0}$
 $\begin{cases} A_1 = A_0 \cos \theta_0 \\ \theta_1 = 0 \end{cases}$

Step3: *t* > *τ*

$$\begin{array}{c} x_2(t) = A_2 \cos\left(\omega t + \theta_2\right) \\ v_2(t) = -\omega A_2 \sin\left(\omega t + \theta_2\right) \end{array} \xrightarrow{v_2(\tau)=0} \begin{cases} A_2 = A_1 \cos \omega \tau \\ \theta_2 = -\omega \tau \end{cases}$$

T-Quench Echo: Harmonic Approximation

for
$$t > \tau$$
: $T(t) \propto \langle v_2^2 \rangle = \langle \omega^2 A_0^2 \cos^2 \theta_0 \cos^2 (\omega \tau) \sin^2 (\omega (t - \tau)) \rangle$

The average must be taken over the distribution of initial phases θ_0 , amplitudes A_0 and angular velocities w

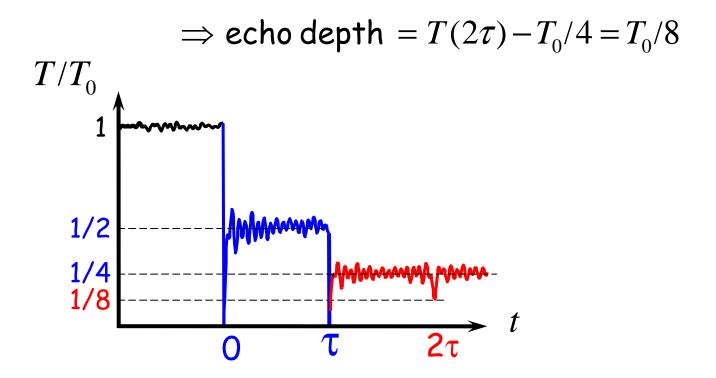
equipartition theorem
$$\Rightarrow \langle A_0^2 \cos^2 \theta_0 \rangle = \frac{1}{2} \langle A_0^2 \rangle = \frac{k_B T_0}{2 m \omega^2}$$

$$T(t) = T_0 \left\langle \cos^2 \left(\omega \tau \right) \sin^2 \left(\omega (t - \tau) \right) \right\rangle = \dots$$
$$= \frac{T_0}{4} \left[1 + \left\langle \cos \left(2\omega \tau \right) \right\rangle - \left\langle \cos \left(2\omega (t - \tau) \right) \right\rangle$$
$$- \frac{1}{2} \left\langle \cos \left(2\omega t \right) \right\rangle - \frac{1}{2} \left\langle \cos \left(2\omega (t - 2\tau) \right) \right\rangle \right]$$

Since: $\left< \cos(\omega?) \right>_{\omega} \approx 0$ unless $? = 0 \Rightarrow$

T-Quench Echo: Harmonic Approximation

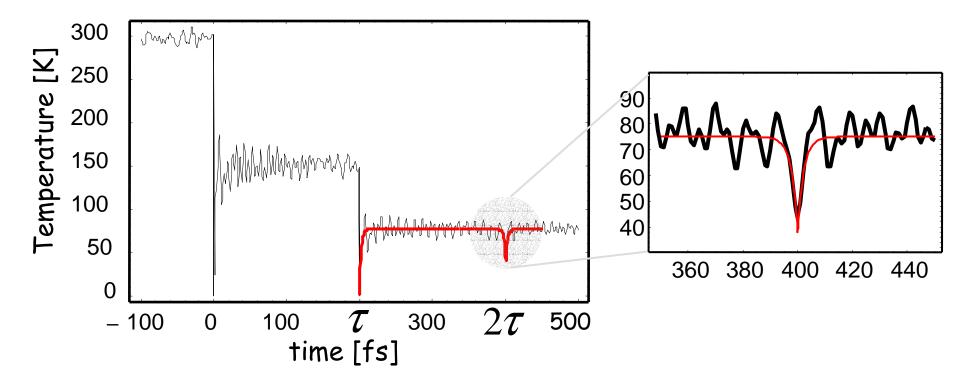
$$T(t) \approx \frac{T_0}{4} \left[1 - \left\langle \cos\left(2\omega(t-\tau)\right) \right\rangle - \frac{1}{2} \left\langle \cos\left(2\omega(t-2\tau)\right) \right\rangle \right]$$
$$\approx \begin{cases} 0 \quad for \ t = \tau \\ T_0/8 \quad for \ t = 2\tau \\ T_0/4 \quad otherwise \end{cases}$$



T(t) and $C_{TT}(t)$

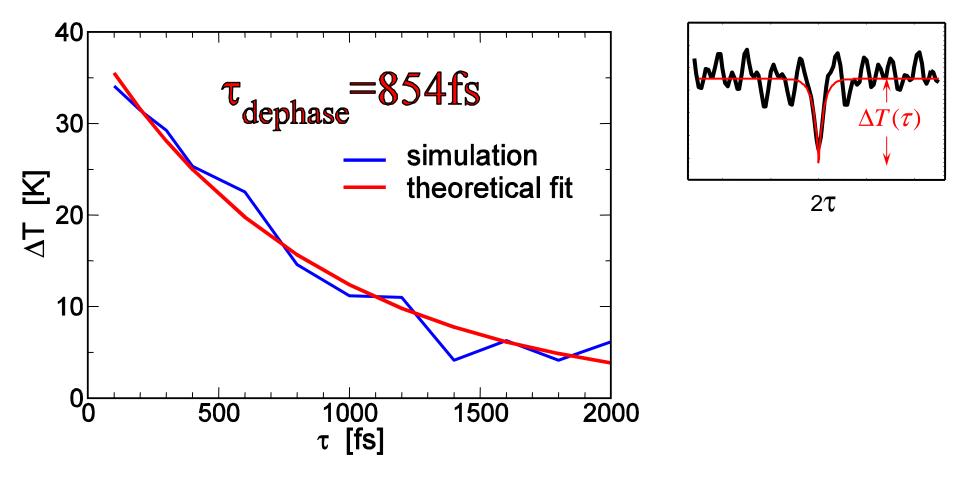
It can be shown that:

T-Quench Echo: Harmonic Approximation



$$T(t) \approx \frac{T_0}{2} \left(1 - C_{TT}(t - \tau) - \frac{1}{2} C_{TT} \left(\left| t - 2\tau \right| \right) \right)$$
$$C_{TT}(t) = \exp\left(-t / \tau_0 \right), \qquad \tau_0 \approx 2.2 \, fs$$

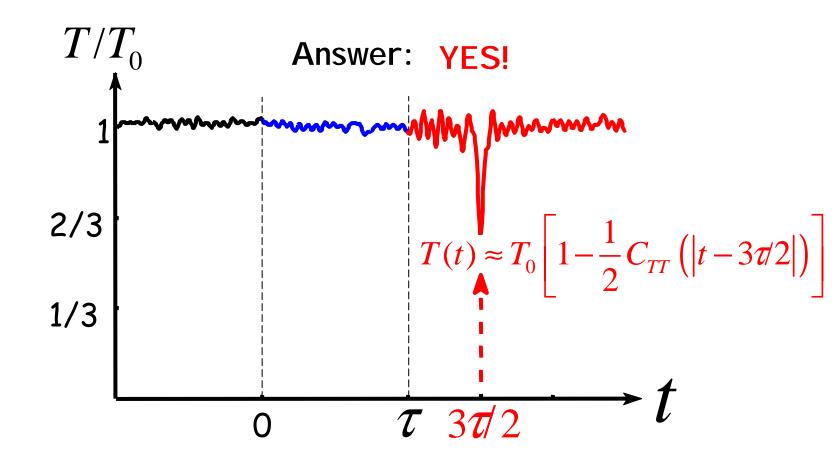
Dephasing Time of T-Quench Echoes



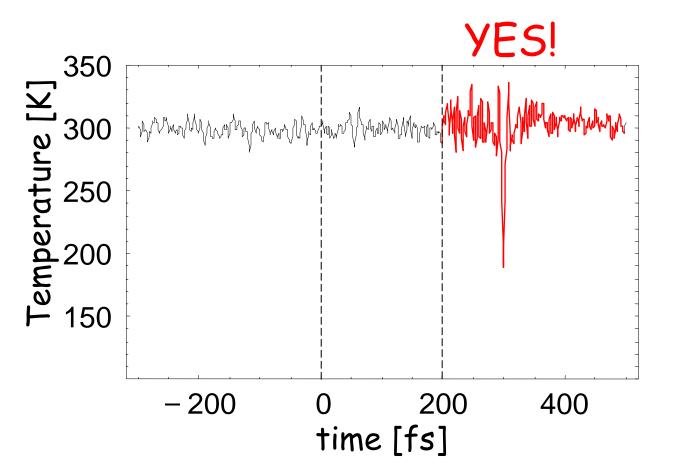
 $\Delta T(\tau) = \Delta T(0) \exp[-\tau / \tau_{dephase}]$

Constant Velocity Reassignment Echo ?

Can we get temperature echo(es) by reassigning the same set of atomic velocities (corresponding to T_0 !) at t = 0 and $t = \tau$? $v_i(0^+) = v_i(\tau^+) = u_i$, i = 1, ..., 3N - 6



Is it possible to produce temperature echo with a single velocity reassignment ?



Reset all velocities at time τ to the values at a previous instant of time, i.e., t=0