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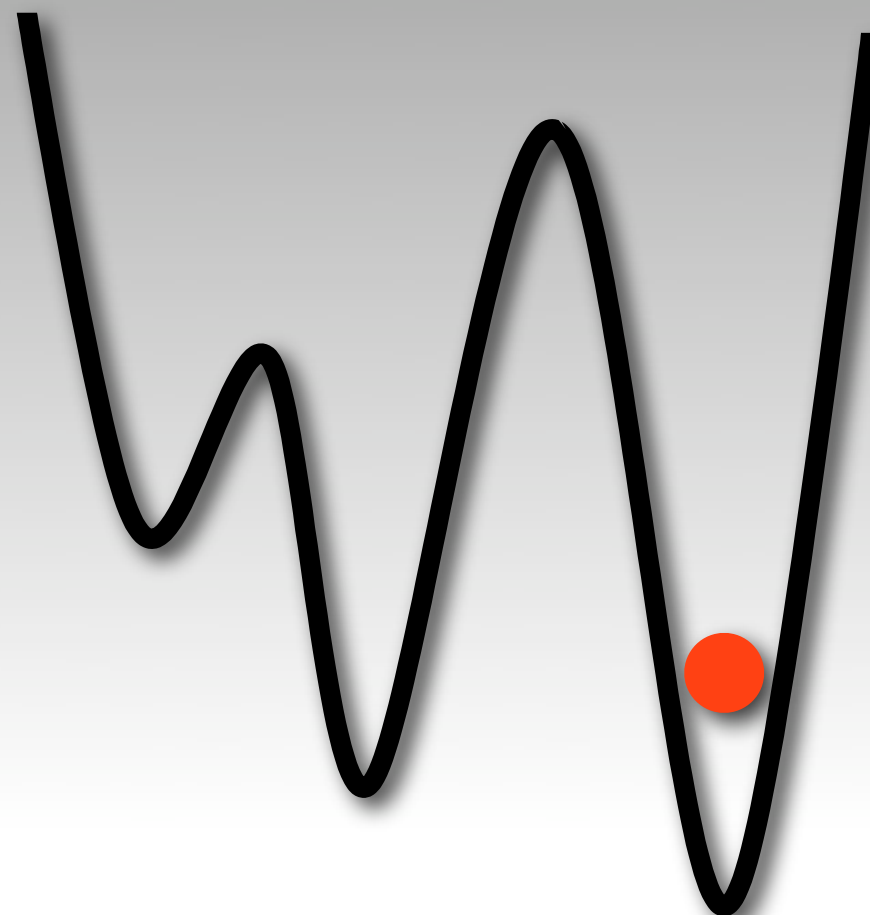
Helmut G. Katzgraber



<http://katzgraber.org>

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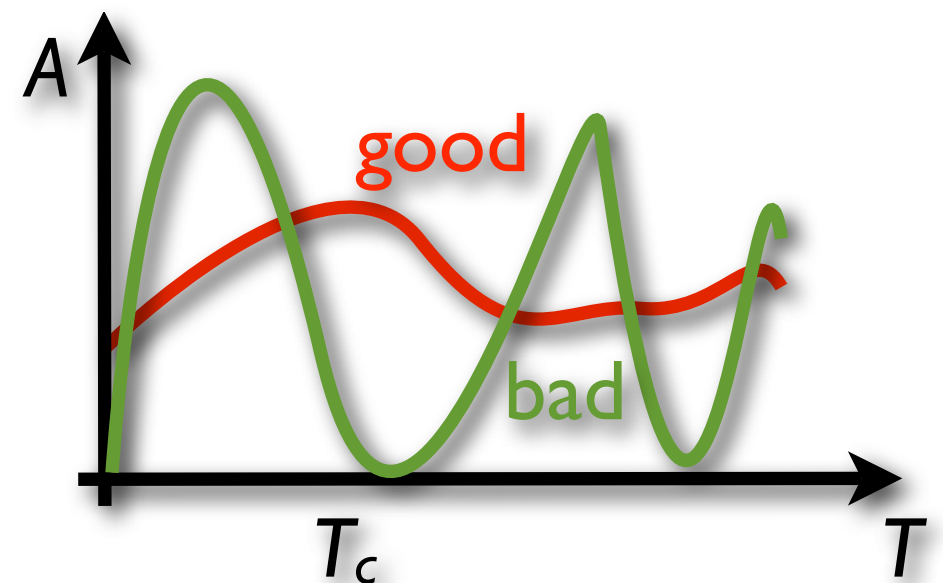
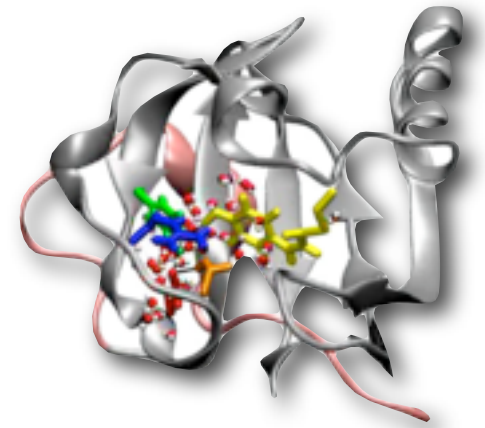
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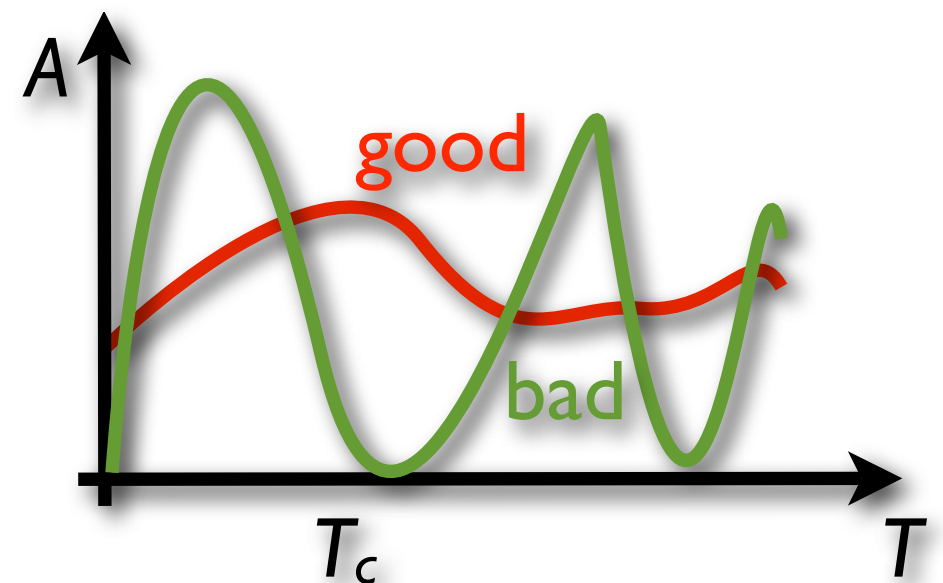
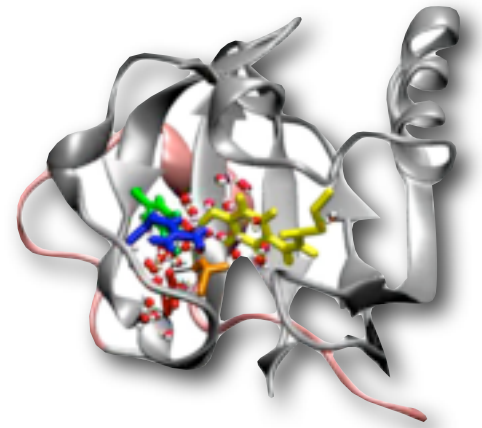
# Outline

- **Brief overview of optimization problems:**
  - Why do we need finite-temperature algorithms?
- **Exchange (parallel tempering) Monte Carlo:**
  - Advantages over simple Monte Carlo.
  - Algorithm description.
- **Tuning the method:**
  - Traditional vs feedback-optimized.
  - Some extensions and applications.
- Presented research done in collaboration with D. Huse, S. Trebst and M. Troyer.



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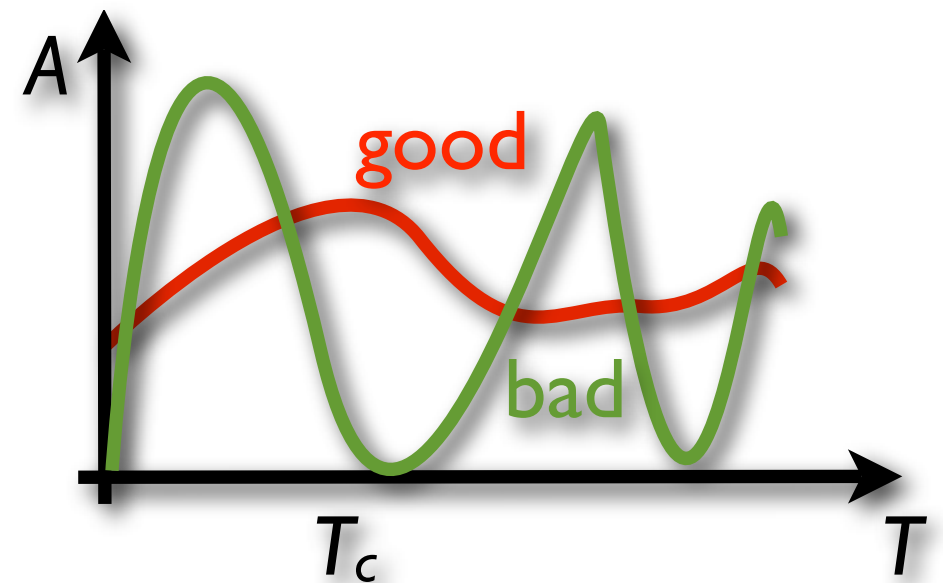
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# Exchange vs parallel tempering

- **History:**

- The method was developed in 1996 by Hukushima and Nemoto. They called it “**exchange** Monte Carlo.”
- Marinari & Parisi developed a similar method called simulated tempering and suggested a variation they dubbed “**parallel tempering** Monte Carlo”.
- Recently, I discovered that Geyer presented another version in 91 in a proceedings book. So far, I could not get my hands on it...

- **Conundrum:**

- What should we call it?
- Traditionally it is called “parallel tempering.”
- Politically correct probably “exchange Monte Carlo.”

# Optimization problems

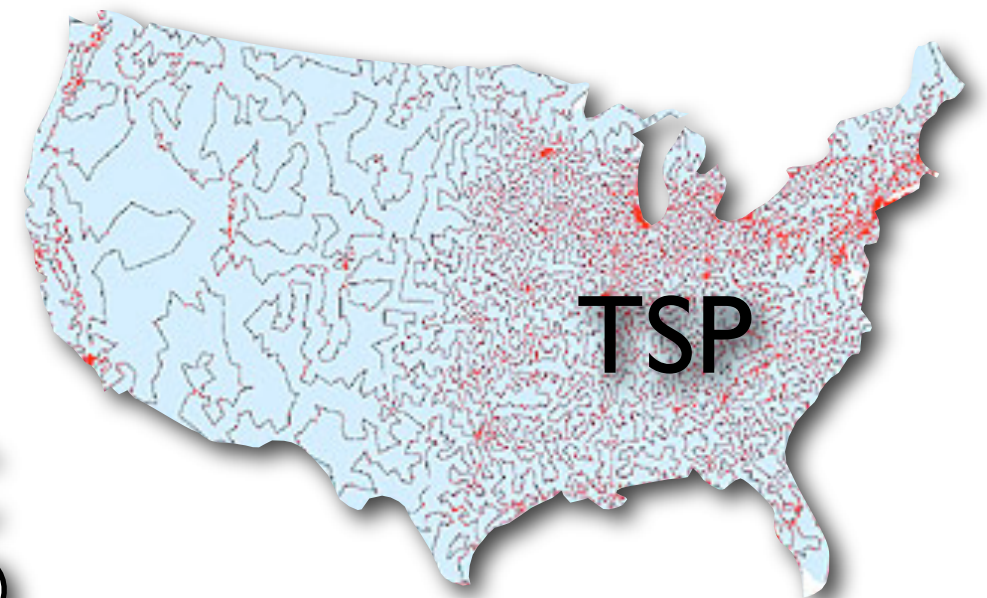
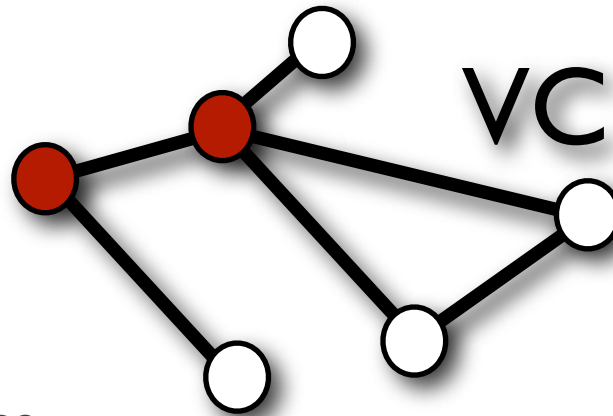


# Overview of optimization problems

- **Typical goals one usually wants to tackle:**
  - Minimize a cost function of a problem (e.g., calculate the ground-state energy of a system).
  - Compute an observable (e.g., energy, magnetization, ...) at low temperature.

- **Typical optimization problems:**

- Physical, biological, chemical systems
- $k$ -SAT
- NPP
- Vertex cover problem
- TSP, ...



**NPP**  $\sum_{i \in A_1} a_i = \sum_{i \in A_2} a_i$

$$A = \{a_1, a_2, \dots, a_n\} \rightarrow A_1 \cup A_2$$

**3-SAT**

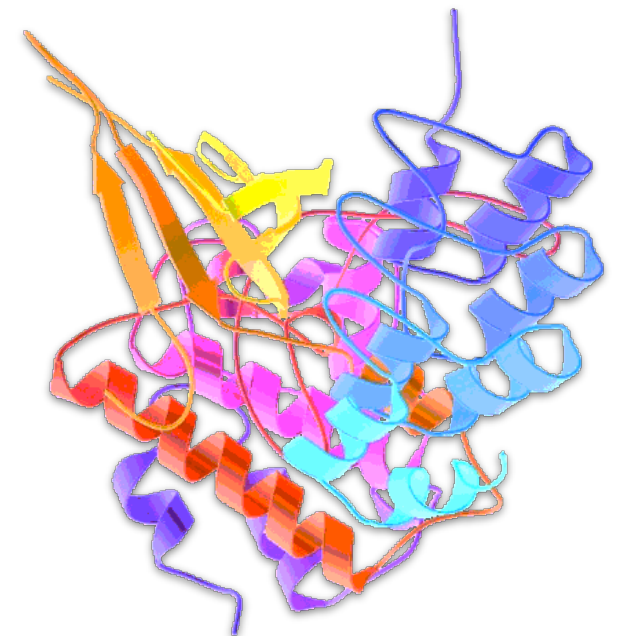
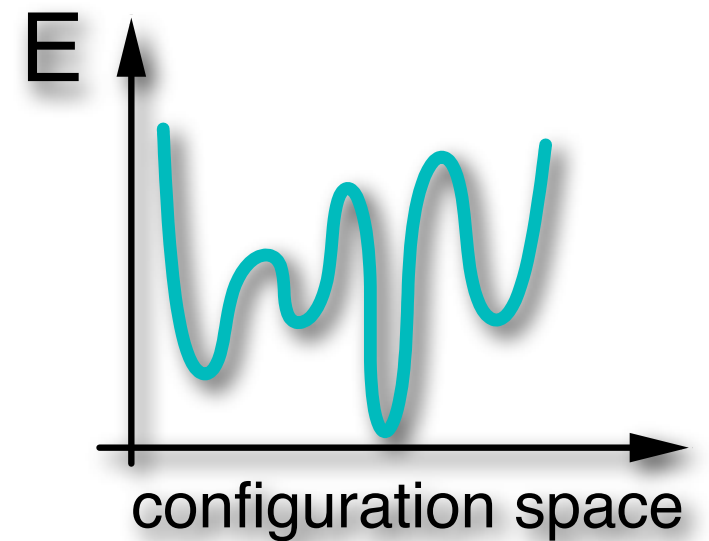
$(x_{11} \text{ OR } x_{12} \text{ OR } x_{13}) \text{ AND } (x_{21} \text{ OR } x_{22} \text{ OR } x_{23}) \text{ AND } \dots$

# Optimization in physical systems

- Several physical problems which contain randomness can be classified as (often hard) optimization problems.
- Due to the randomness, one obtains competing interactions and thus a complex energy landscape.

- **Examples:**

- Spin glasses:  $\mathcal{H} = - \sum_{ij} J_{ij} S_i S_j$   
 $\mathcal{P}(J_{ij})$  random
- Structural glasses
- Polymers in random media (interfaces)
- Biomolecules (proteins, ...)
- Quantum wave functions
- Reconstruction of geological structures from seismic measurements, ...



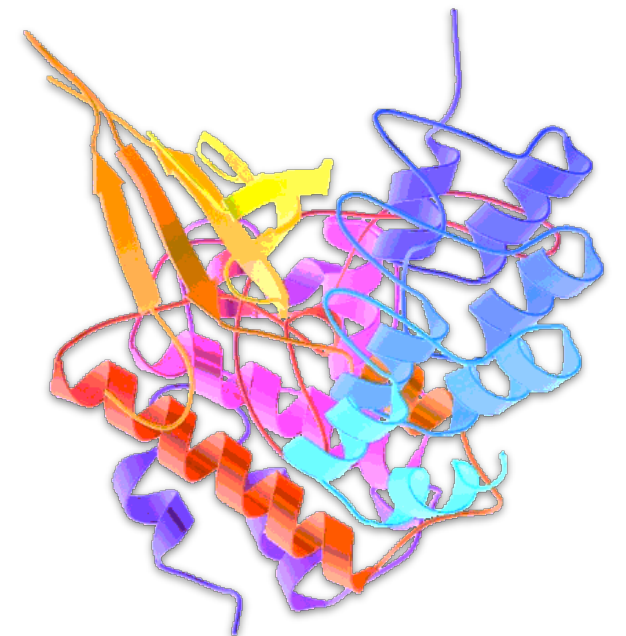
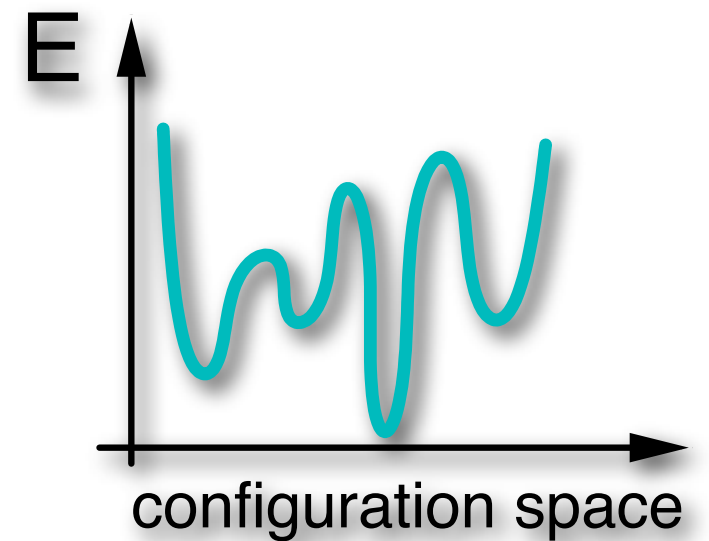
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mañana



# What will be discussed here?

- **So far:**
  - Methods to compute ground states of complex systems (zero-temperature methods).
  - Methods to study phase transitions and overcome critical slowing down (for example cluster algorithms).
  - Methods to tackle systems which undergo first-order transitions (for example Wang-Landau algorithm).
  - Methods for infinite temperature (series expansions).

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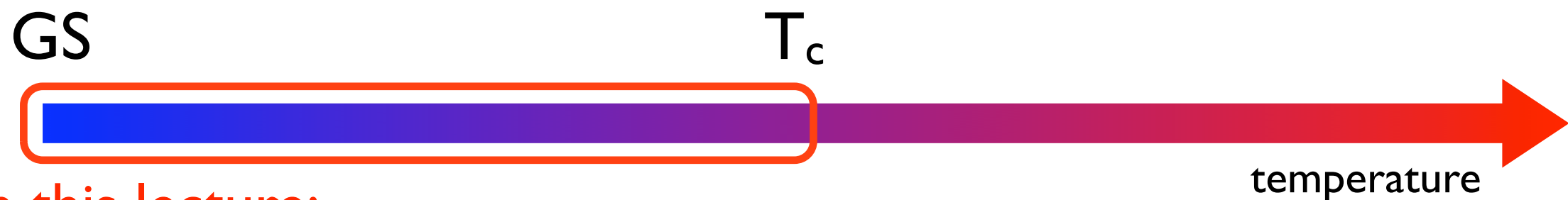
$T_c$



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  - Study systems with rough energy landscapes at low temperatures.
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# Simple benchmark model family



# Prototype for a magnet: the Ising model

- **Hamiltonian:**

$$\mathcal{H} = - \sum_{\langle ij \rangle} J_{ij} S_i S_j - H \sum_i S_i$$

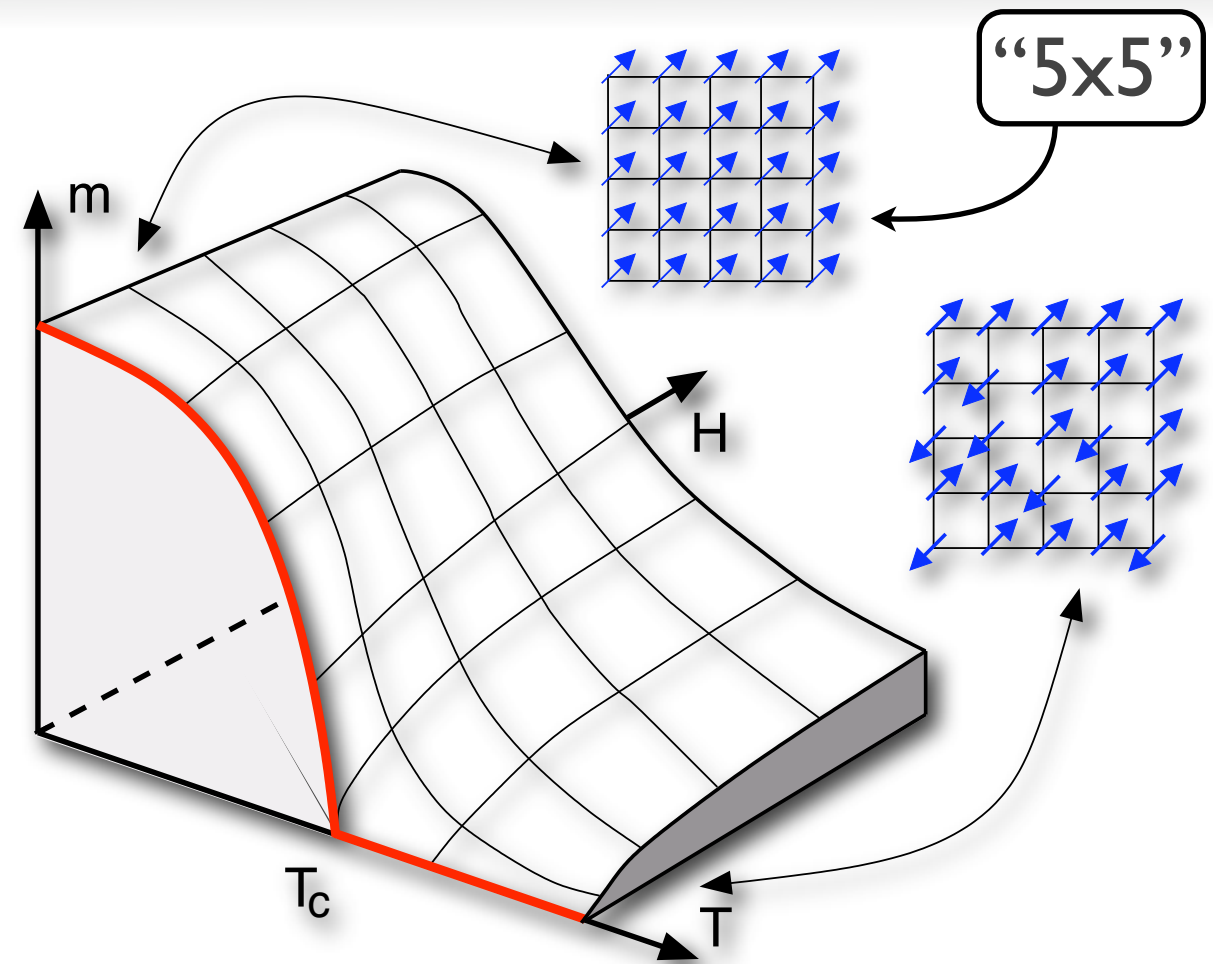
$$J_{ij} = 1 \quad \forall i, j \quad i \neq j$$

- **Order parameter:**

$$m = \frac{1}{N} \sum_i S_i \quad (\text{magnetization})$$

- **Some properties:**

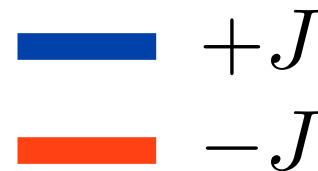
- Describes many magnetic systems (also other spin symmetries).
- Exact solution in 2 space dimensions (Onsager).
- Nearest-neighbor interactions, in 2D  $T_c \sim 2.27$ .



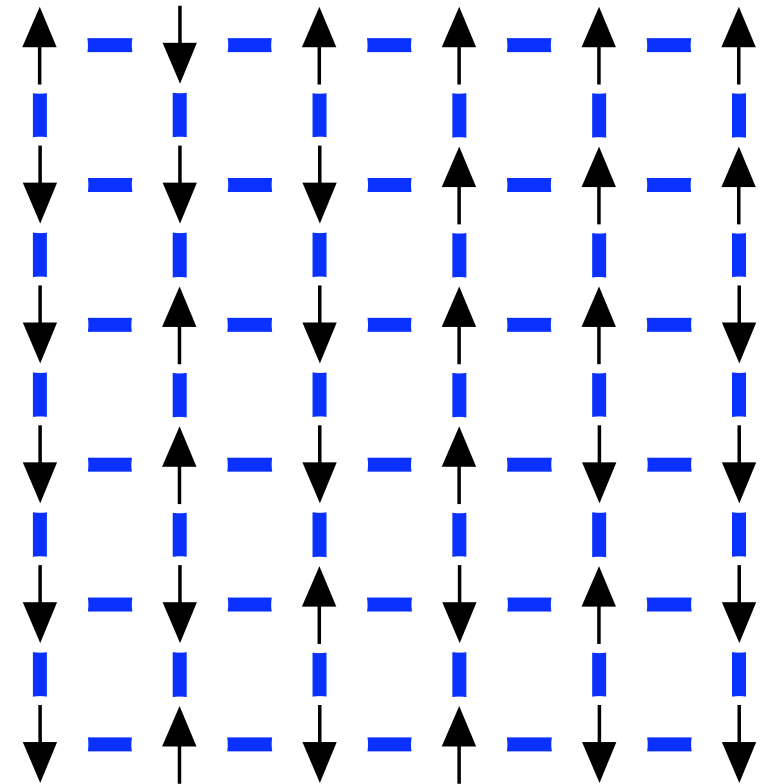
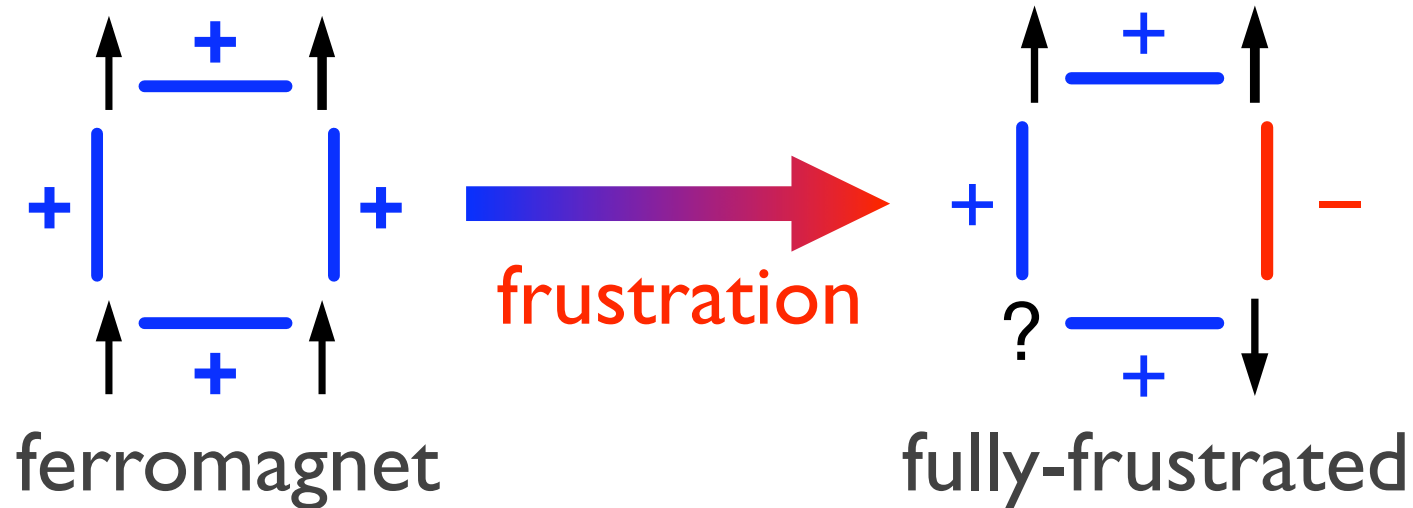
# Adding frustration...

- General Hamiltonian:

$$\mathcal{H} = - \sum_{\langle ij \rangle} J_{ij} S_i S_j$$



- Introduce frustration between the spins:



- Properties of the fully-frustrated Ising model:

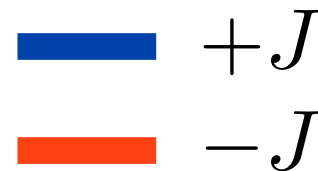
- Huge ground-state degeneracy and complex energy landscape.
- $T_c = 0$  in 2D.
- $\prod_{\square} J_{ij} < 0 \quad \forall i, j$

- What happens if we add randomness to the frustration?

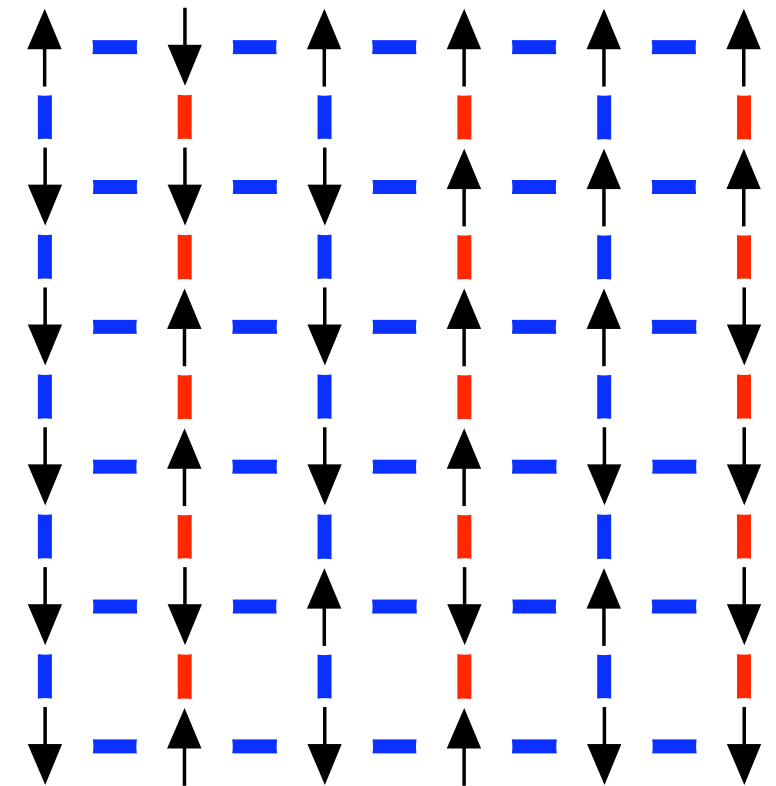
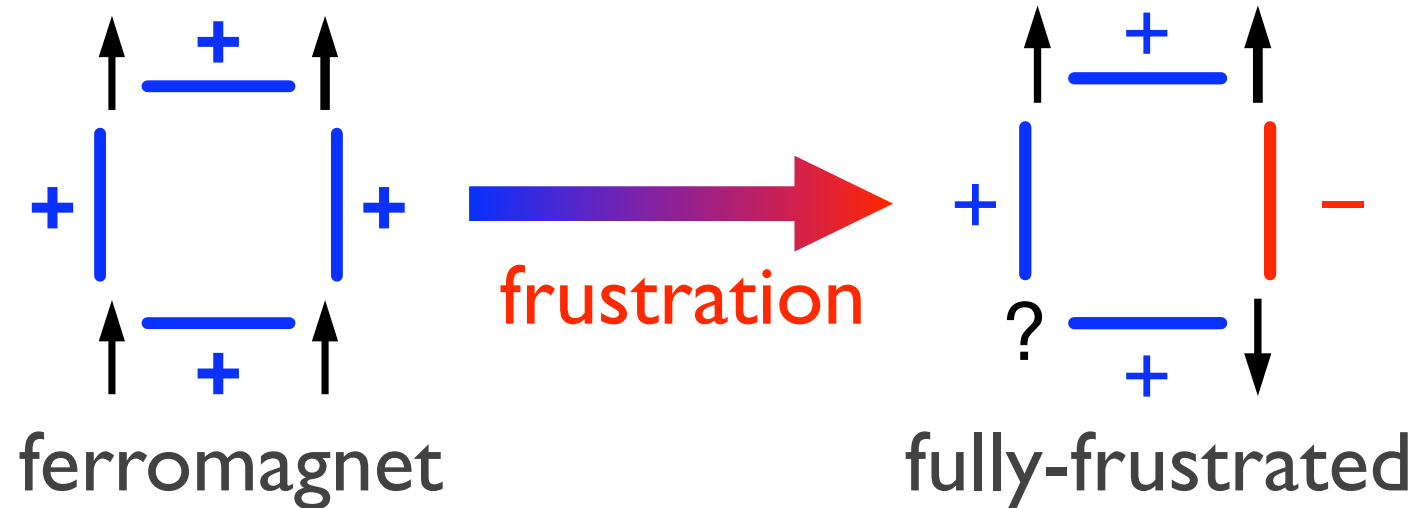
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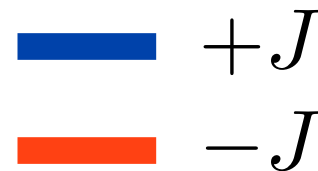
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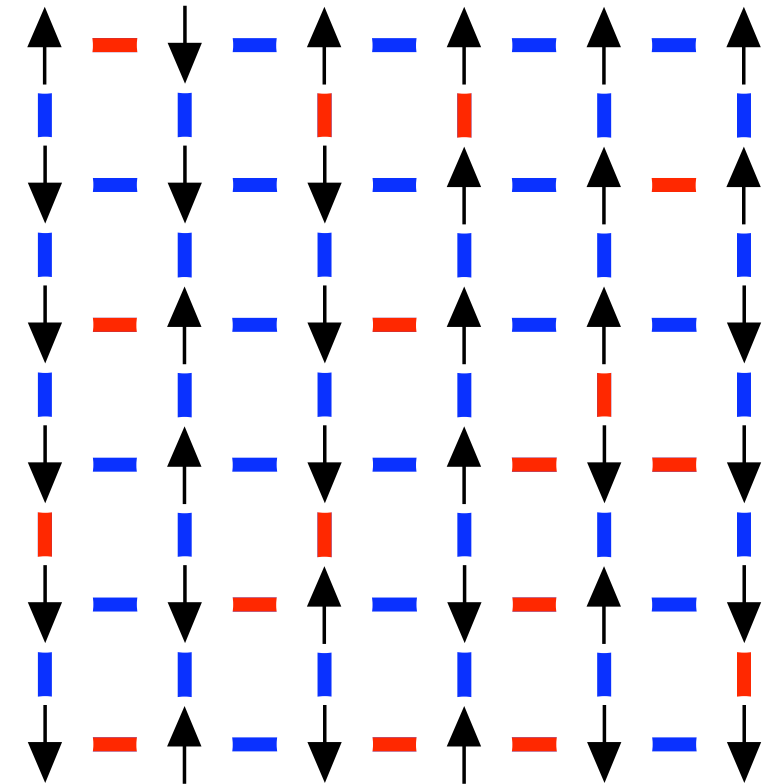
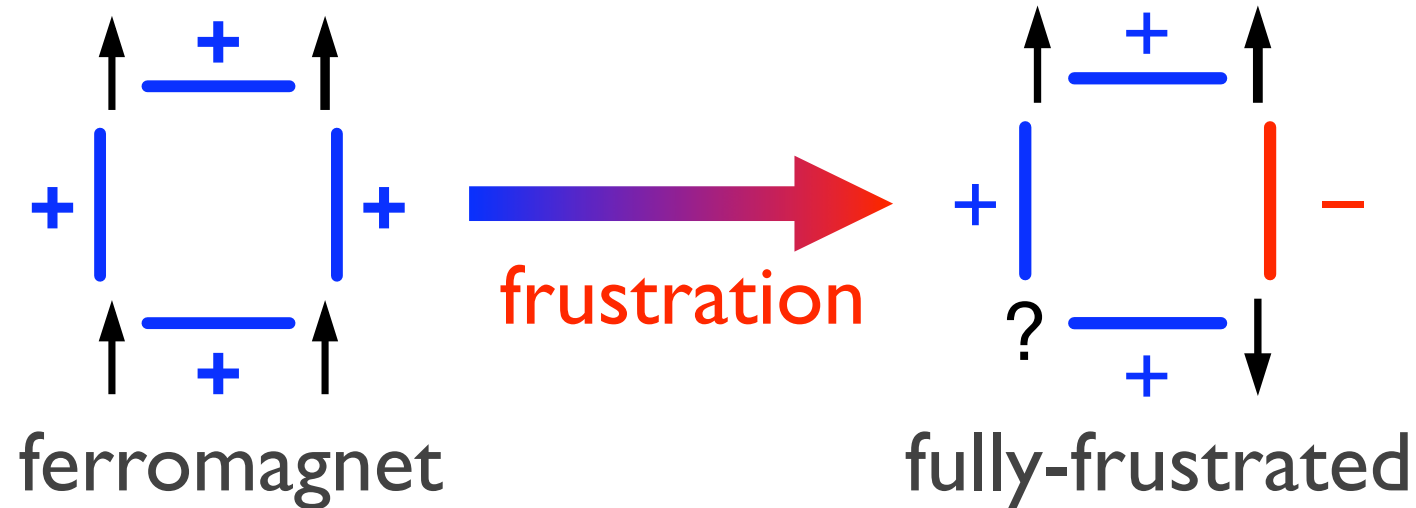
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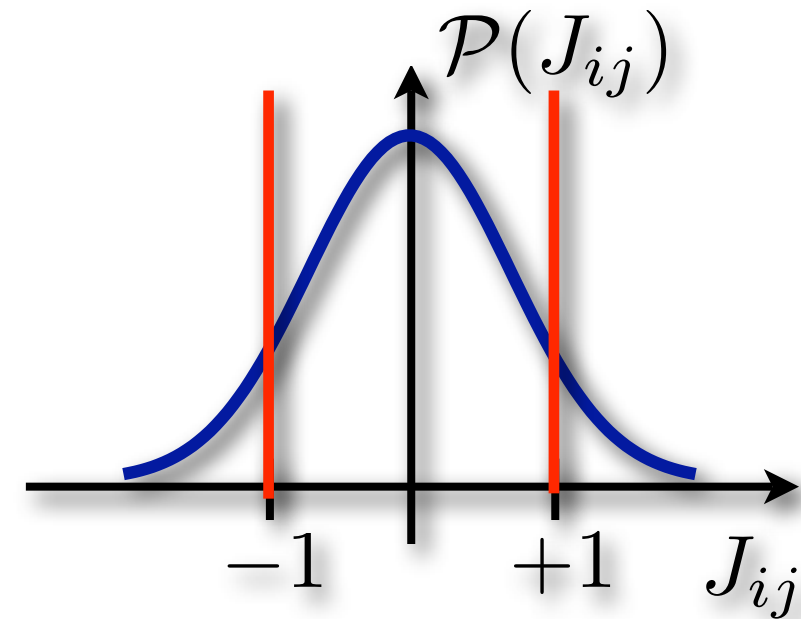
# Adding frustration and disorder: spin glass

- **Spin-glass Hamiltonian:**

$$\mathcal{H} = - \sum_{\langle i,j \rangle} J_{ij} S_i S_j \quad S_i \in \{\pm 1\}$$

- **Details about the model:**

- Unconventional “order.”
- Only mean-field solution.
- Bimodal random bonds: high degeneracy.
- Gaussian random bonds: unique ground state.
- 2D  $T_c = 0$
- 3D  $T_c = 0.951(2)$
- 4D  $T_c = 1.805(10)$



- Will be discussed in detail tomorrow and not discussed here further.

# Reminder: Simple Monte Carlo



# The original Metropolis paper (1953)...

THE JOURNAL OF CHEMICAL PHYSICS

VOLUME 21, NUMBER 6

JUNE, 1953

## Equation of State Calculations by Fast Computing Machines

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(Received March 6, 1953)

A general method, suitable for fast computing machines, for investigating such properties as equations of state for substances consisting of interacting individual molecules is described. The method consists of a modified Monte Carlo integration over configuration space. Results for the two-dimensional rigid-sphere system have been obtained on the Los Alamos MANIAC and are presented here. These results are compared to the free volume equation of state and to a four-term virial coefficient expansion.

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### II. THE GENERAL METHOD FOR AN ARBITRARY POTENTIAL BETWEEN THE PARTICLES

In order to reduce the problem to a feasible size for numerical work, we can, of course, consider only a finite number of particles. This number  $N$  may be as high as several hundred. Our system consists of a square† con-

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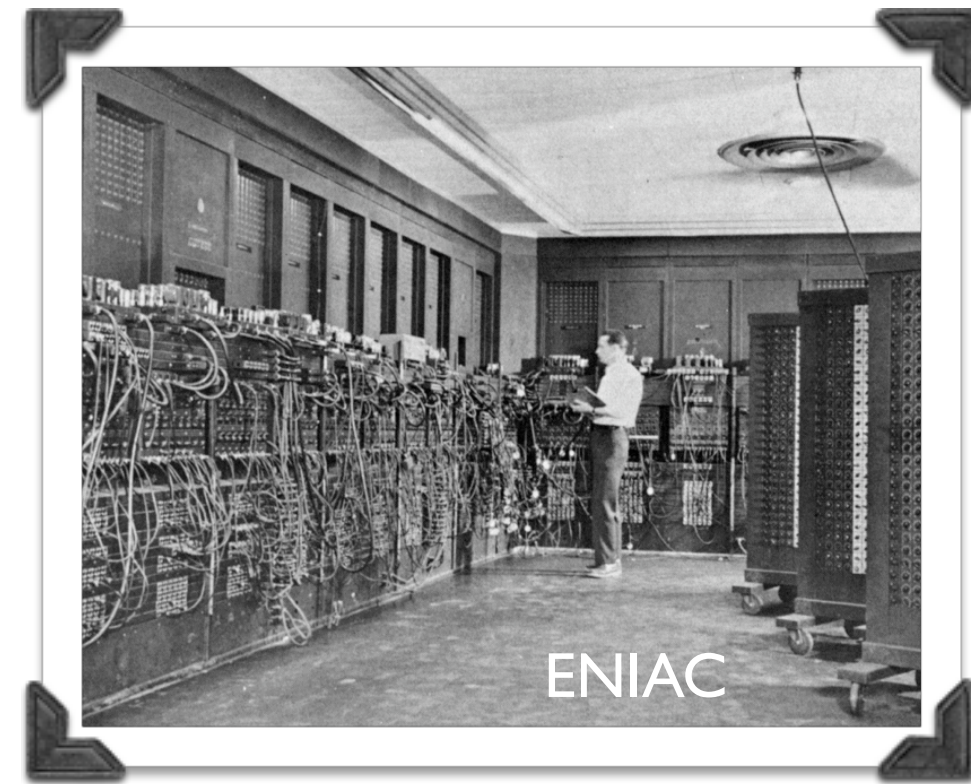
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# The original Metropolis paper: some facts

Phys. Plasmas 12, 057303 (05)

- 50 years later at a Los Alamos meeting:
  - Only M. Rosenbluth attended, although with terminal cancer.
  - Metropolis mainly contributed CPU time on MANIAC.
  - von Neumann and Ulam invented the Monte Carlo method in 1946 and pointed out that it could be used for simulations.
  - Teller: Statistical averages can be made as ensemble averages.
  - Interesting author list: two couples. How often does this happen?
- Why Los Alamos?
  - The US was bulding the atomic bomb. At least one good thing came out.





# Reminder: simple Monte Carlo

- In statistical mechanics we want to compute the average of an observable  $O$ :

$$\langle O \rangle = \sum_n P_n^{\text{eq}} O_n \quad P_n^{\text{eq}} = \frac{e^{\beta E_n}}{\sum_n e^{-\beta E_n}}$$

- **Problem:** The number of states is exponentially large.
- **Solution:** Statistically sample a few (smartly chosen) states to obtain an estimate of  $O$  but with a statistical error.
  - If we chose the states according to  $P_n^{\text{eq}}$  and ensure detailed balance we obtain a Markov chain for  $\langle O \rangle_{\text{est}}$

$$\langle O \rangle_{\text{est}} = \frac{1}{M} \sum_i^M O_i$$

where  $M$  is the number of states sampled.

- **Metropolis algorithm:** accept a new configuration if  $(e^{-\Delta E/T} > \text{rand}())$

$$P_{\text{accept}} = \min(1, e^{-\Delta E/T})$$

# Why does simple Monte Carlo fail here?

- The systems we are interested in have rugged energy landscapes.
- At low temperature, when  $\Delta E$  is large

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is “never” accepted.

- **How can we resolve the problem?**
  - Tunnel through barrier.
  - Heat up the system to overcome the barrier.
- **Note:**
  - Simulated annealing has similar problems (stuck in valleys).



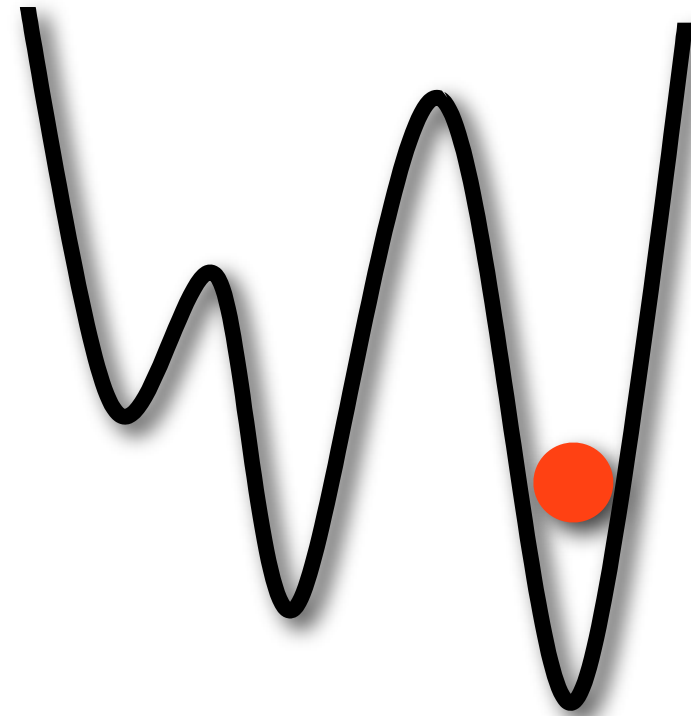


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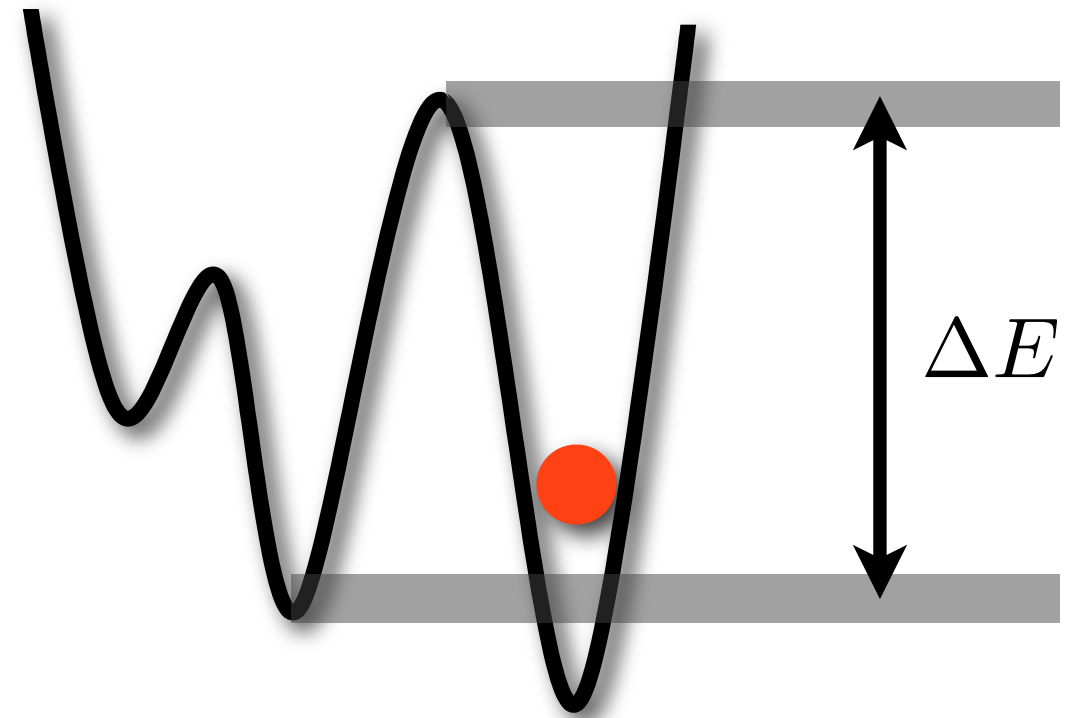
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# Exchange (parallel tempering) Monte Carlo

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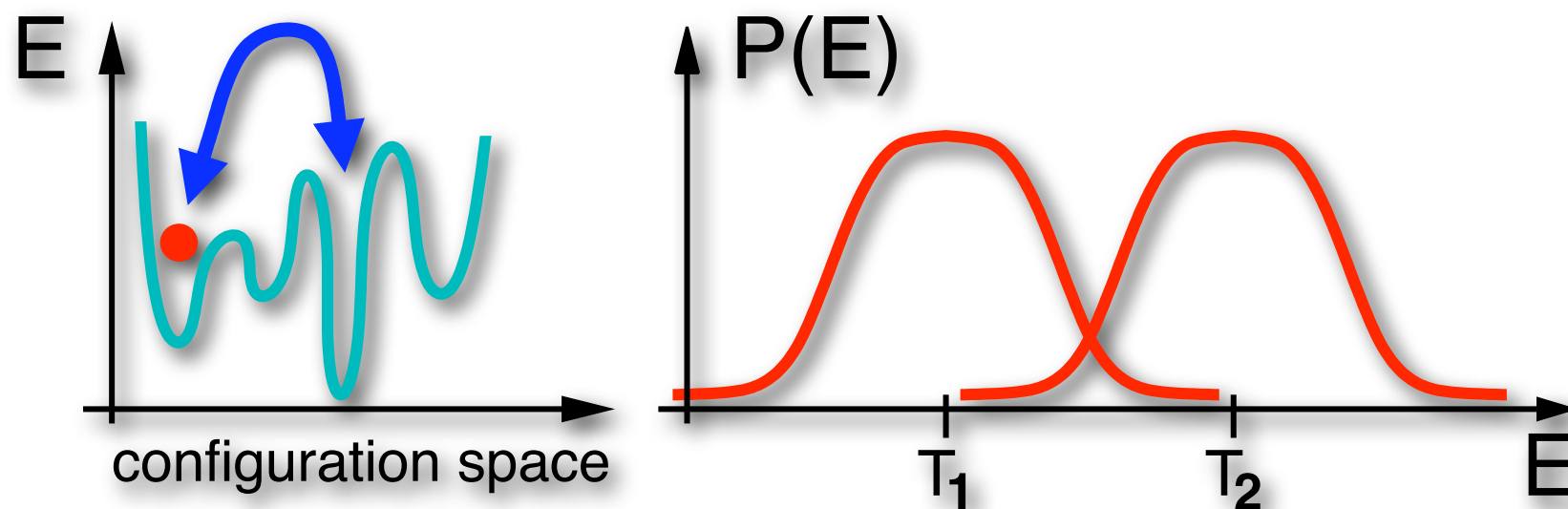
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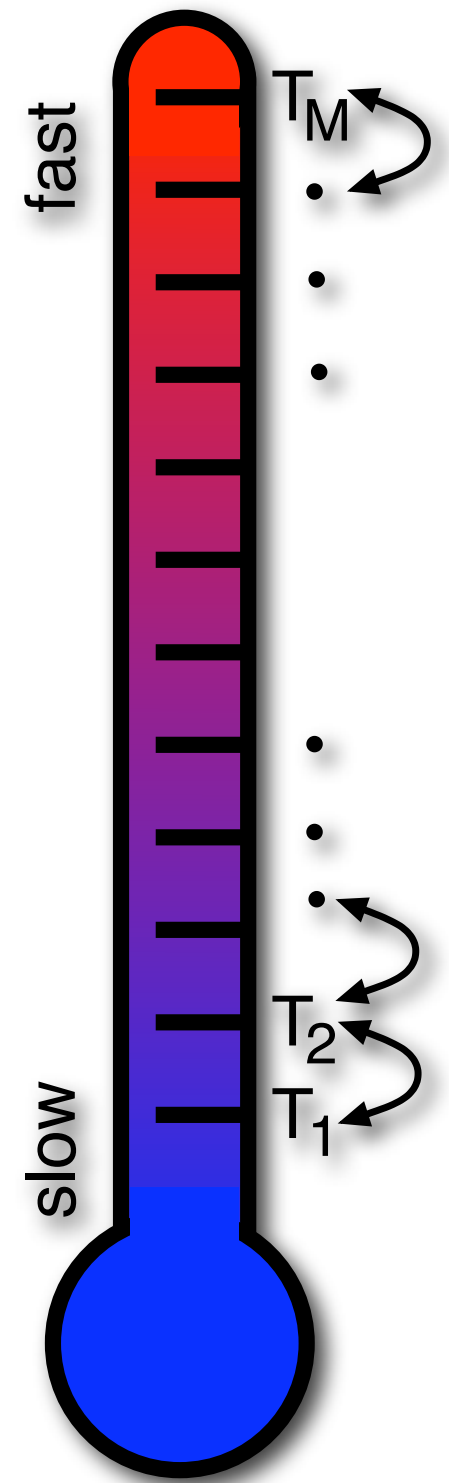
# Exchange (parallel tempering) Monte Carlo

Hukushima & Nemoto (96)

- **Idea:**
  - Simulate  $M$  copies of the system at different temperatures with  $T_{\max} > T_c$  (typically  $T_{\max} \sim 2T_c^{\text{MF}}$ ).
  - Allow swapping of neighboring temperatures: easy crossing of barriers.



- **What has to be tuned?**
  - Number of temperatures.
  - Position of temperatures.



# Parallel tempering: algorithm and details

- **Brief outline of the algorithm:**

- Perform a Monte Carlo update between *neighboring* replicas.
- Best to keep temperatures and change pointers to configurations.

$$P(S_{m+1} \leftrightarrow S_m, \beta_{m+1} \leftrightarrow \beta_m) = e^{-\Delta} \quad : \quad \Delta > 0$$

$$P(S_{m+1} \leftrightarrow S_m, \beta_{m+1} \leftrightarrow \beta_m) = 1 \quad : \quad \Delta \leq 0$$

$$\Delta = (\beta_{m+1} - \beta_m)(E_m - E_{m+1})$$

$$\beta = 1/T \quad \text{[obeys detailed balance]}$$

- **How often do we call the swap routine?**

- Optimal ratio of lattice sweeps ( $N$  spin updates) and swaps is 1:1, i.e., after each lattice sweep, perform an attempted parallel tempering swap [see Katzgraber *et al.*, JSTAT P03018 (2006)].

# How fast is fast? Example: 3D spin glass

- Equilibration times:

$$\tau_{\text{eq}}^{\text{PT}} \approx 300 \text{ MCS}$$

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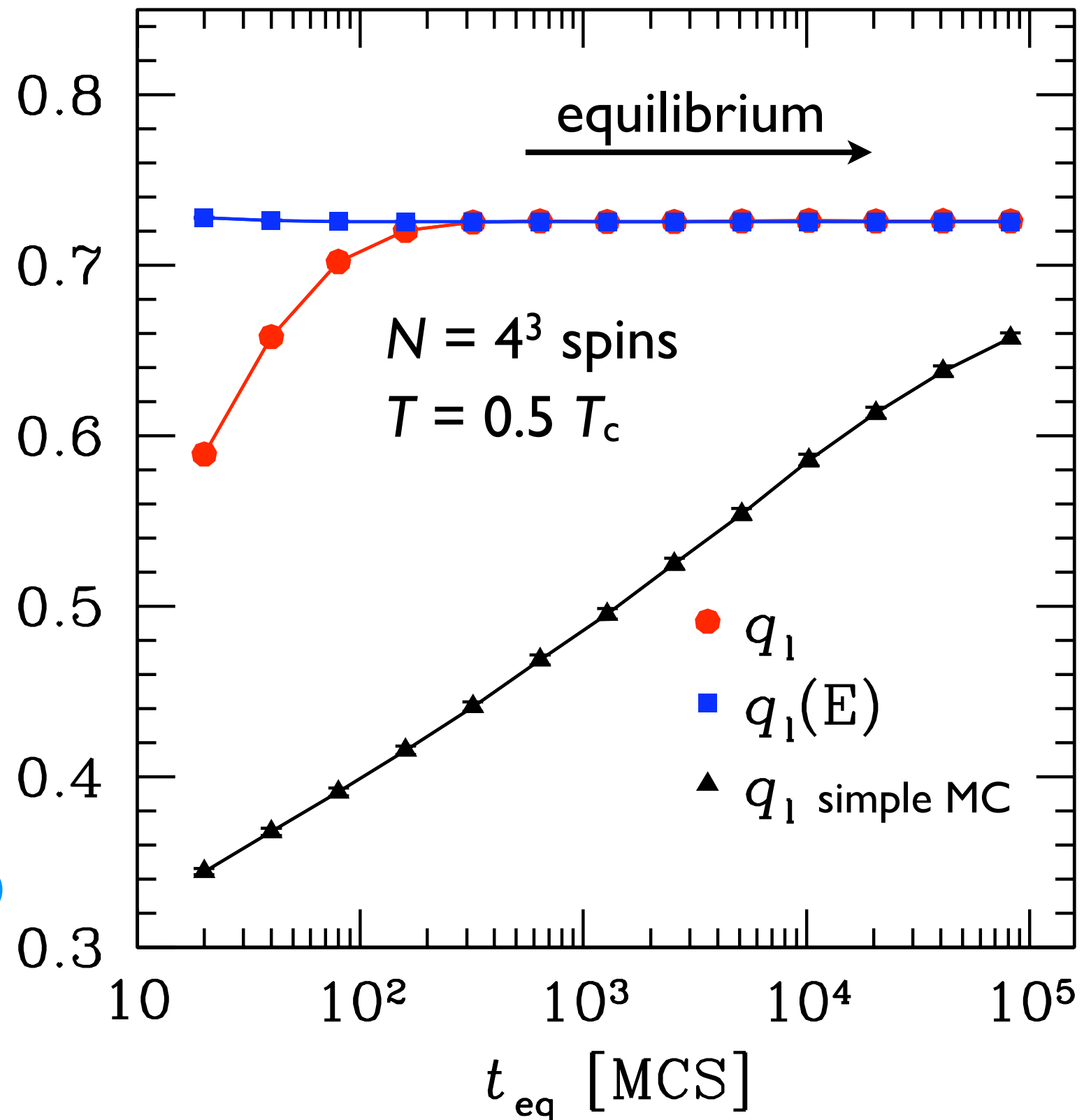
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$$q_l(E) = \frac{2T|E|}{z} + 1$$

$$z = 2D$$

Once both agree, the system is in equilibrium (only Gaussian disorder).

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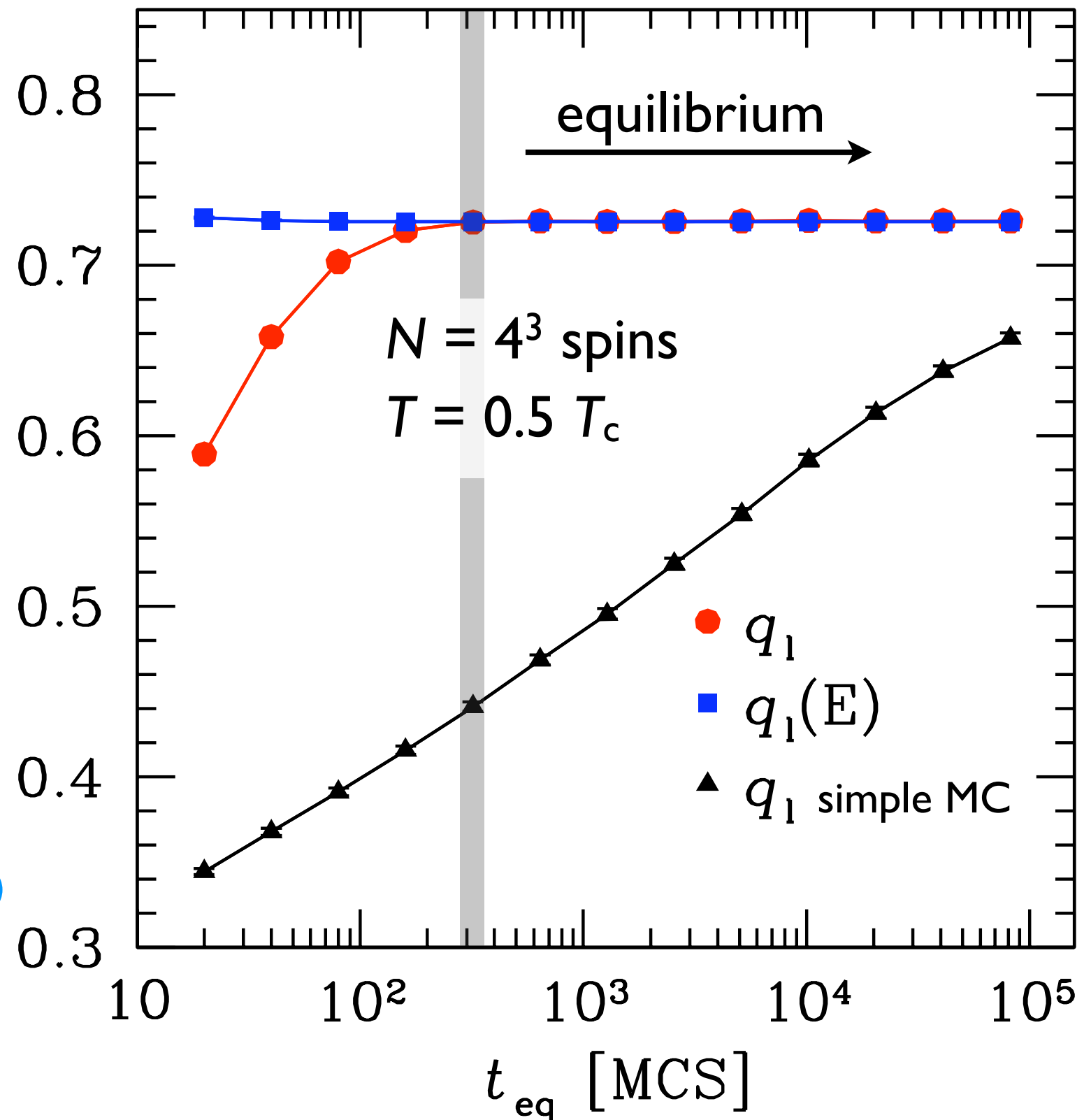
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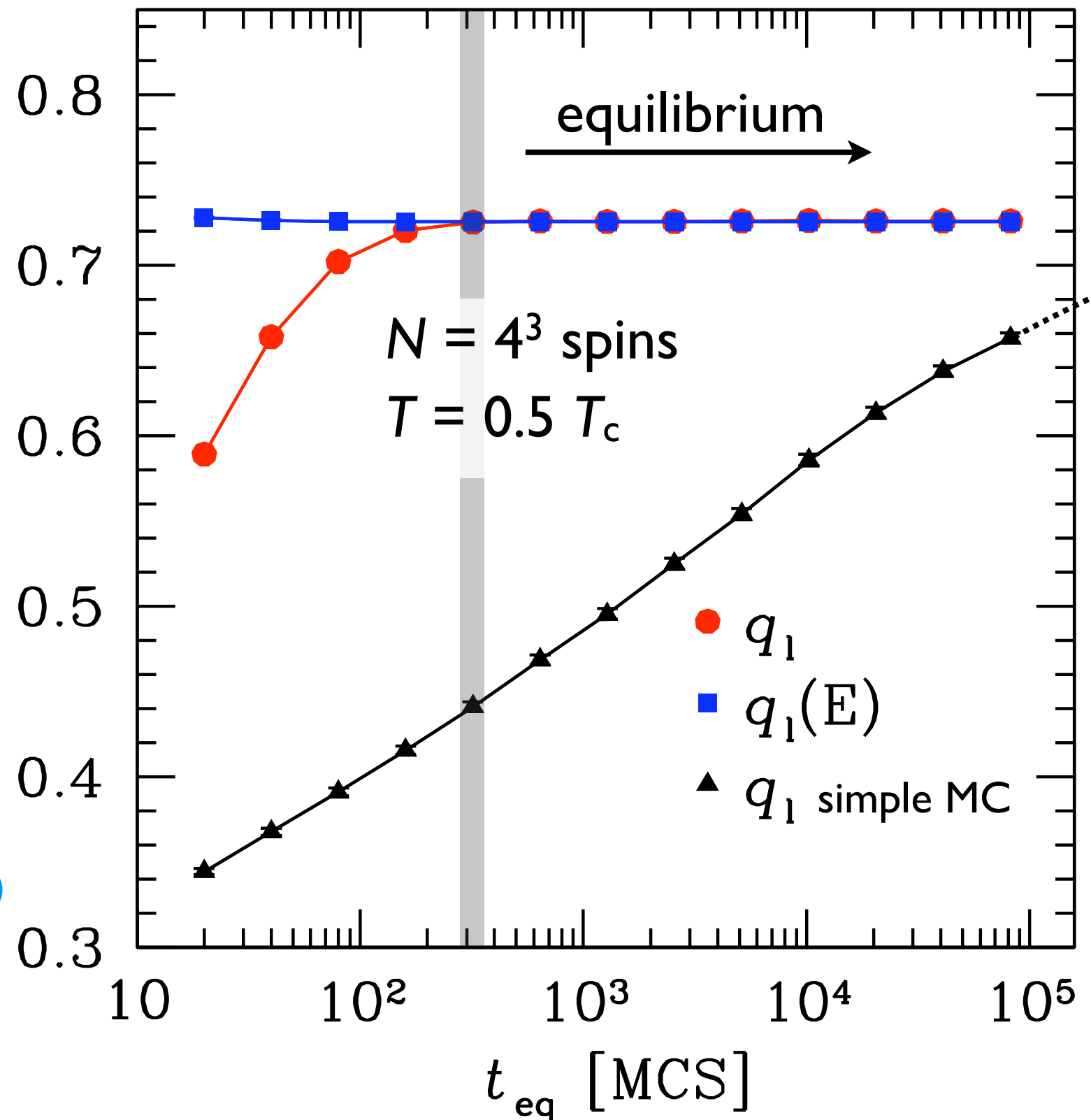
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Katzgraber *et al.* PRB (01)

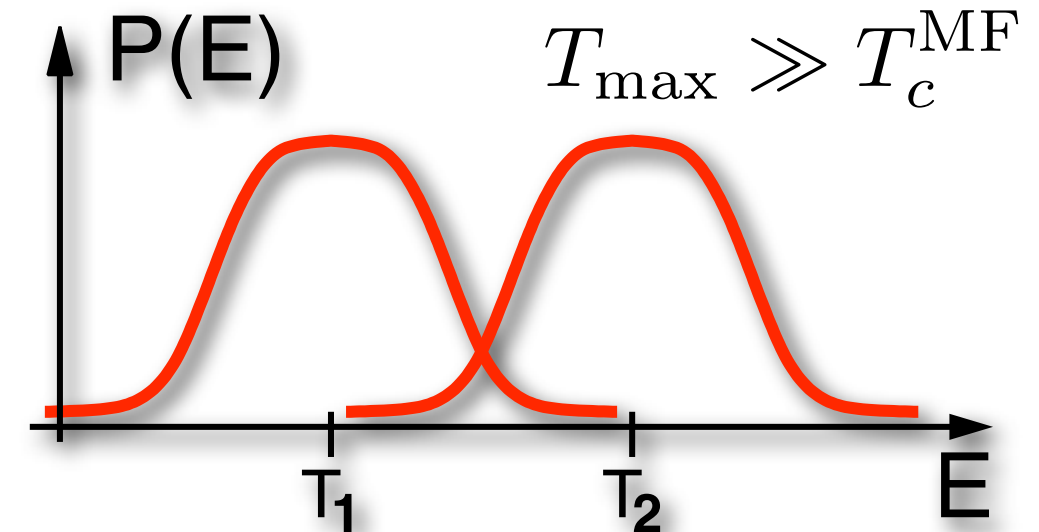


# How do we choose the temperatures?

# How many temperatures do we need?

- **Two possible scenarios:**

- Temperatures too far apart: parallel simple Monte Carlo chains.
- Temperatures too close: overhead.



- **What determines the number  $M$  of temperatures?**

- The energy distributions of the system at  $T_1$  and  $T_2$  have to overlap.
- Because  $\Delta E \sim C_V \longrightarrow M \sim \sqrt{N^{1+\alpha/d\nu}}$
- Note: Systems for which  $C_V|_{T \rightarrow 0} \rightarrow 0$  require many temperatures.

- In principle, we need as many temperatures such that the method works. Measure? Acceptance rates.

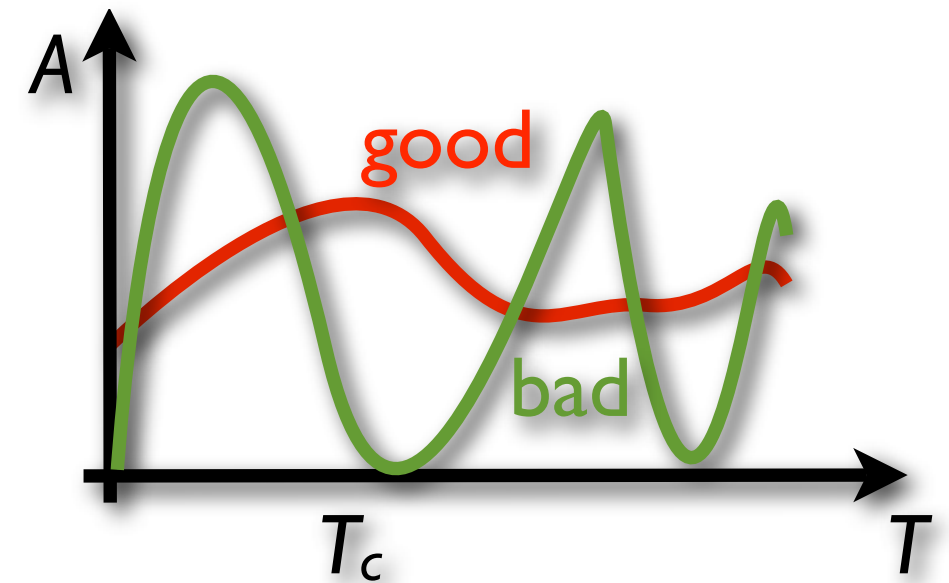
# Measuring acceptance rates

- **Definition:**

$$A = \frac{N_{\text{accept}}}{N_{\text{trial}}}$$

- **Traditional wisdom:** Tune the temperature set such that...

- ...  $0.2 \leq A \leq 0.9$ .
- ...  $A$  is approximately independent of temperature.



- Detailed implementation which gives flat acceptance rates: Incomplete beta function law [uses  $A = f(C_V)$ ].

- **Notes:**

*Predescu et al., JSTAT (03)*

- A quick run (no need to equilibrate) will immediately produce stable acceptance rates (easy tuning by hand).
- It has been claimed that  $A \sim 0.3$  is optimal.

*Rathore et al., J. Chem. Phys. (05)*

# Practical/traditional approaches

- **Geometric progression:**

- Works well when  $C_V \sim \text{const}$  (like for spin glasses).
- Iteratively construct a temperature set and tune  $M$  with  $\lambda$ .

$$\frac{1}{T_i} = \lambda R^{i-1} \frac{1}{T_{\min}} \quad R = \left[ \frac{T_{\min}}{T_{\max}} \right]^{1/(M-1)}$$

- **By hand:**

- If  $C_V$  diverges (phase transition) start from a geometric progression.
- Interlace extra temperatures by hand.
- Tedious, but after a while you get a feeling for it.

- **But... are temperature-independent acceptance rates optimal?**

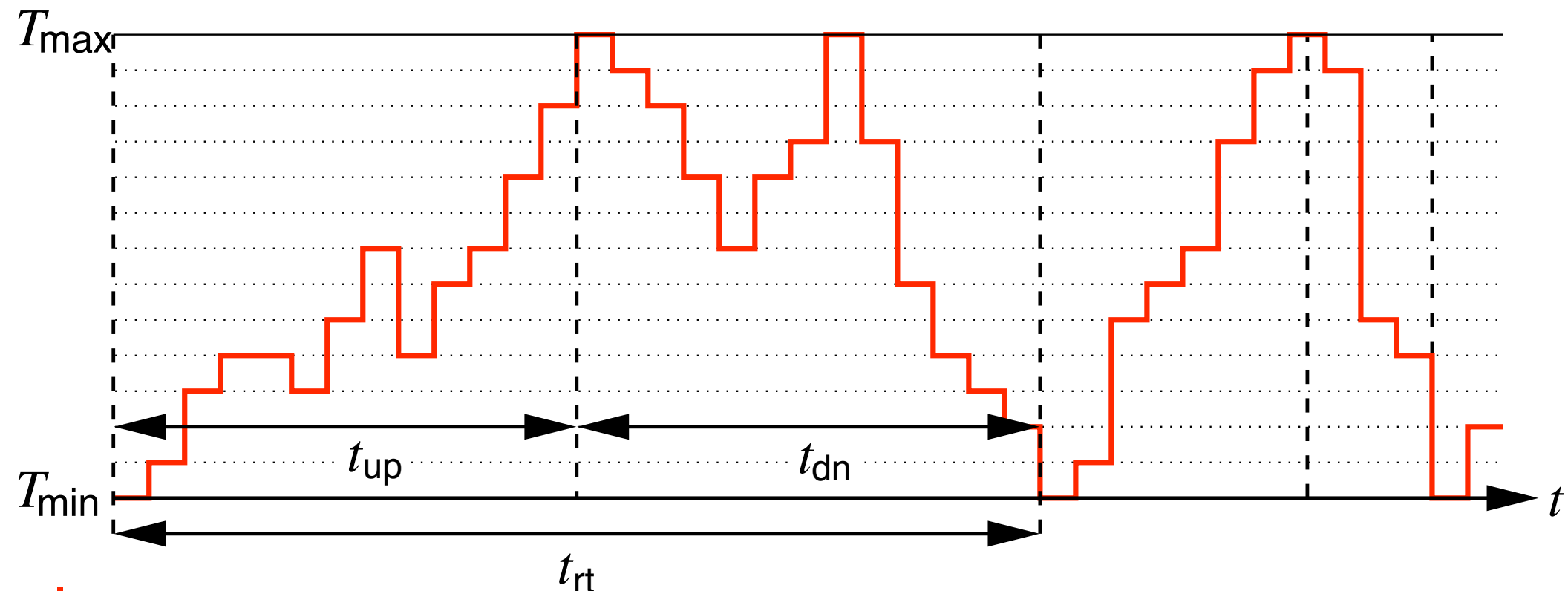
- Replicas do a random walk in temperature space.



# Optimizing the ensemble $\{T_i\}$

Katzgraber *et al.*, JSTAT (06)

- Track random walk of the replicas in temperature space...



- Goal:**
  - Minimize the round-trip time  $t_{\text{rt}}$  and ensure that  $t_{\text{up}} \sim t_{\text{dn}}$ .
  - This shall ensure an efficient sampling of temperature space.
- How?**
  - Tune the ensemble  $\{T_i\}$  at a fixed number of temperatures  $M$ .
  - Quick run at the beginning of the simulation.

# Conventional temperature sets...

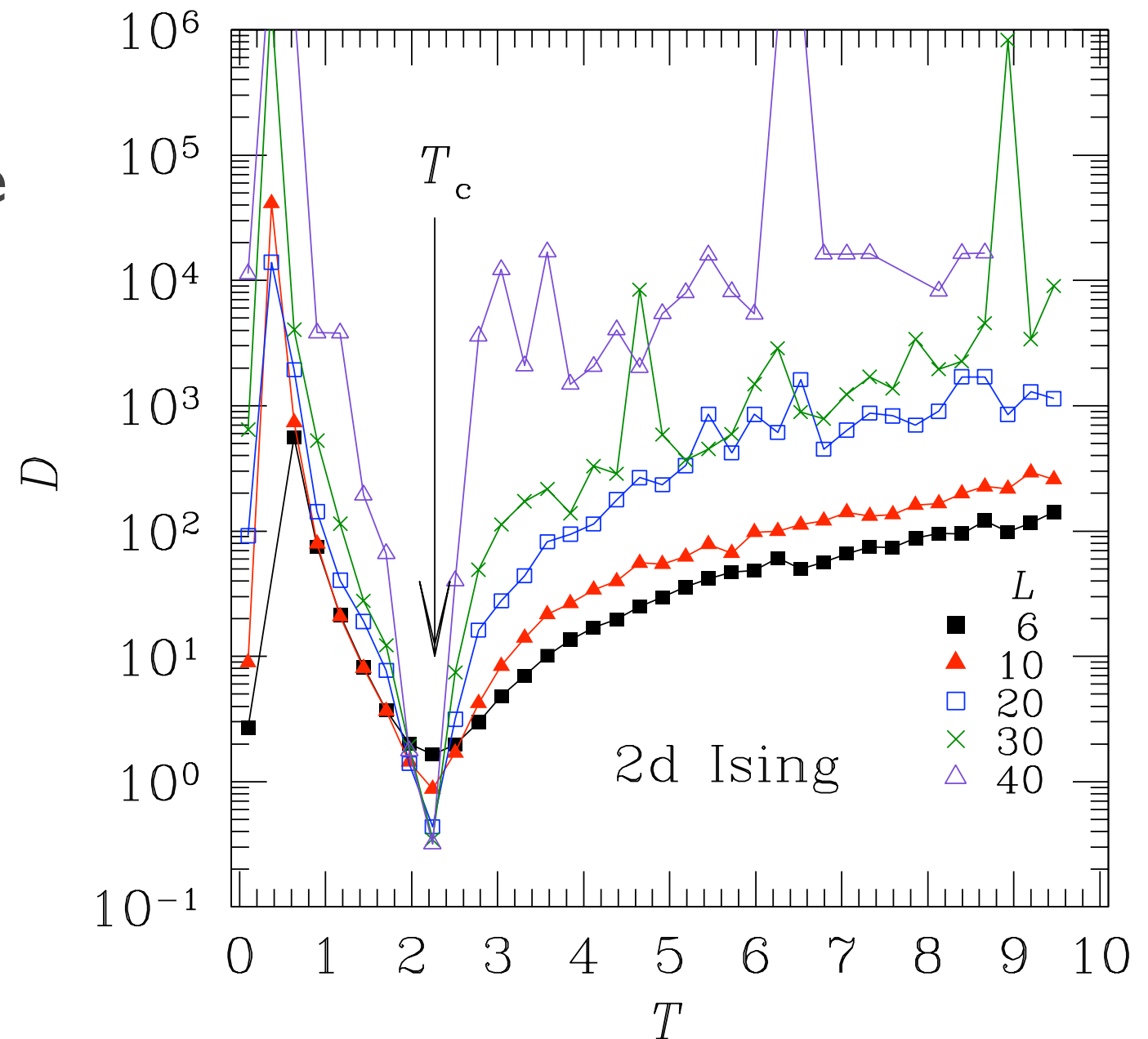
- Start from a temperature set with  $A(T) \sim \text{const.}$
- Track one replica and measure the local diffusivity  $D$  in the ensemble  $\{T_i\}$ .

- Ising model:

- Bottleneck at  $T_c$ !

- Fully-frustrated Ising model:

- Bottleneck at  $T = 0$ .



- Goal: change  $\{T_i\}$  so that  $D \sim \text{const.}$  for each  $\Delta T_i = T_{i+1} - T_i$ .

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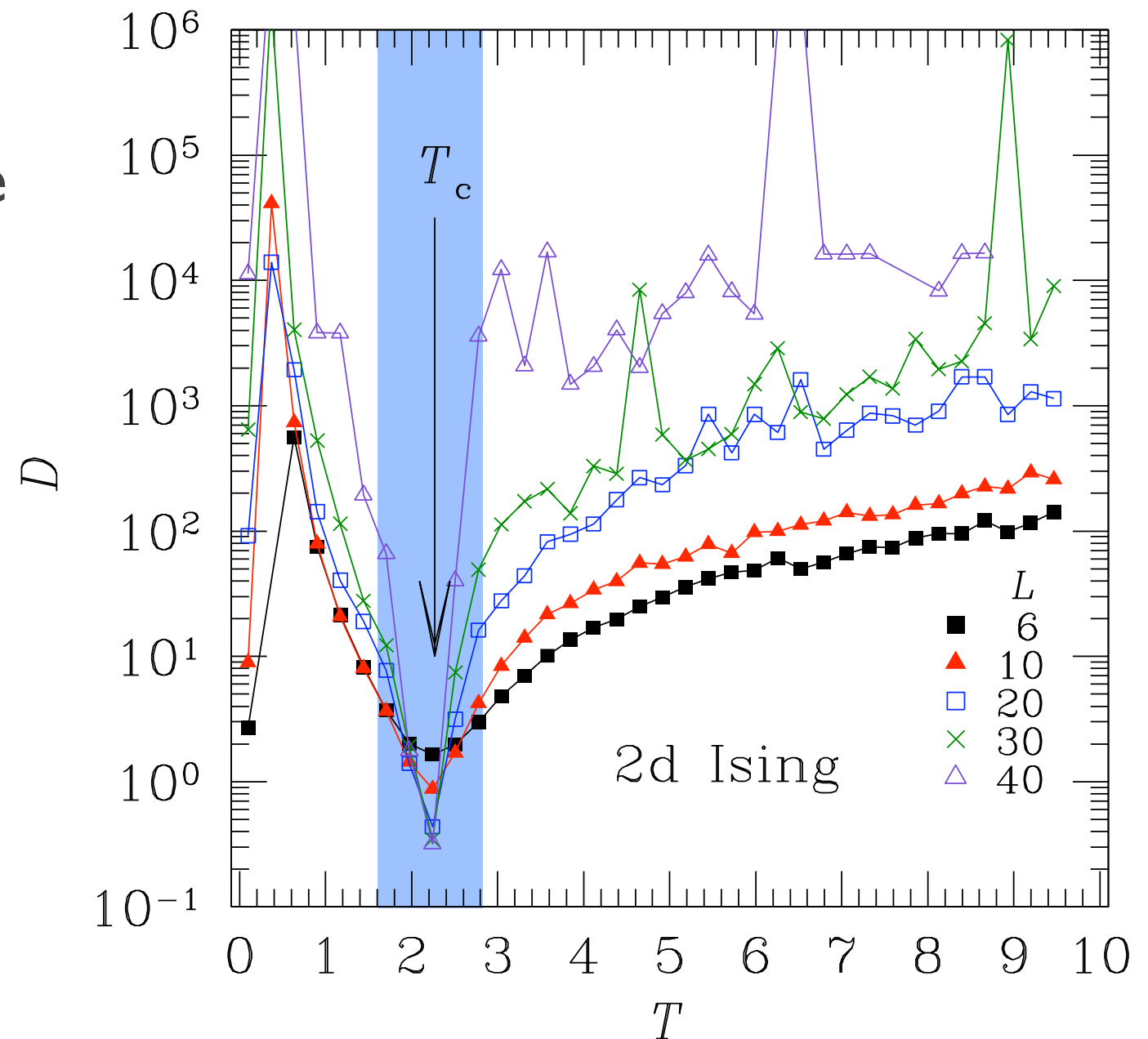
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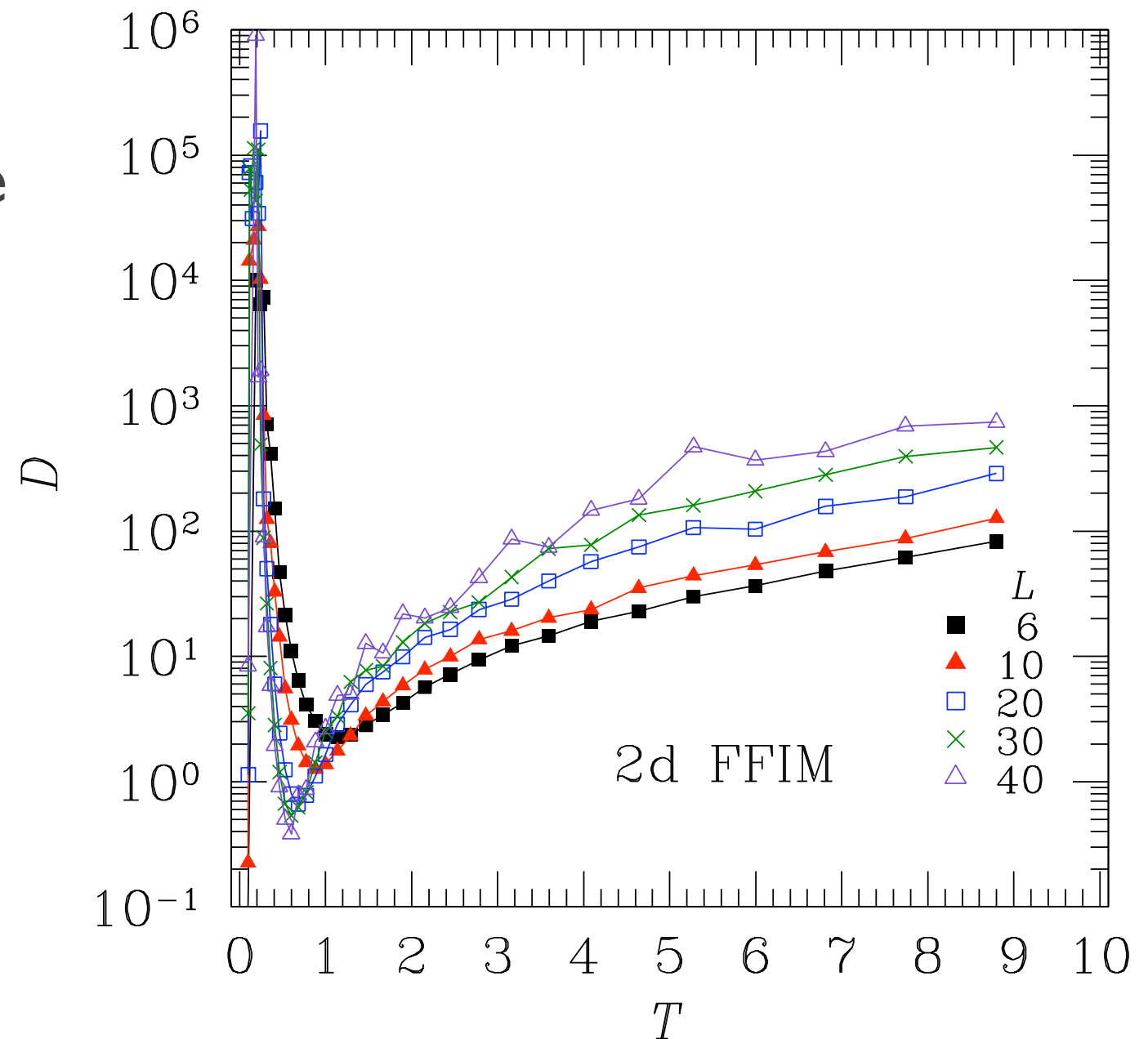
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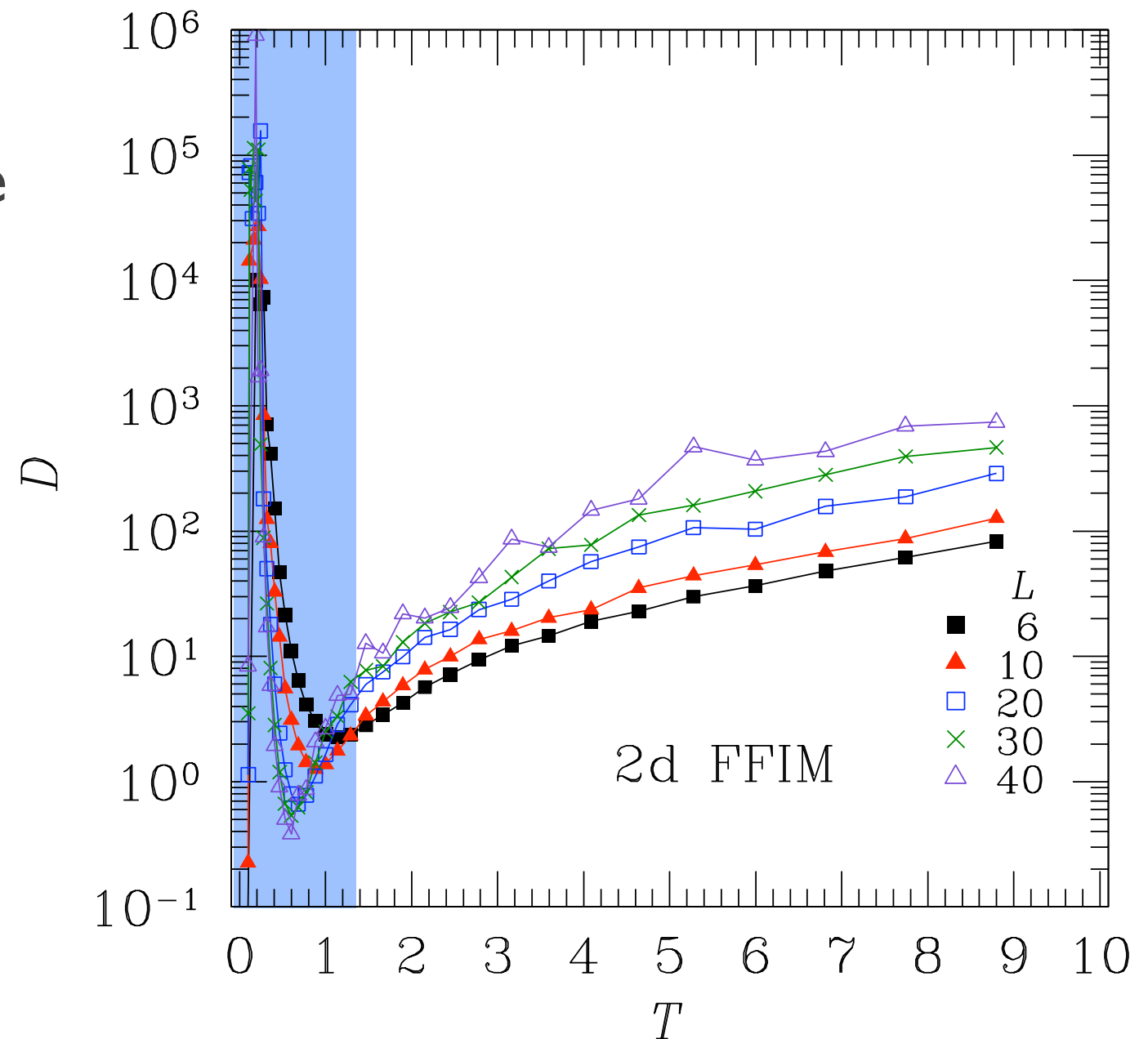


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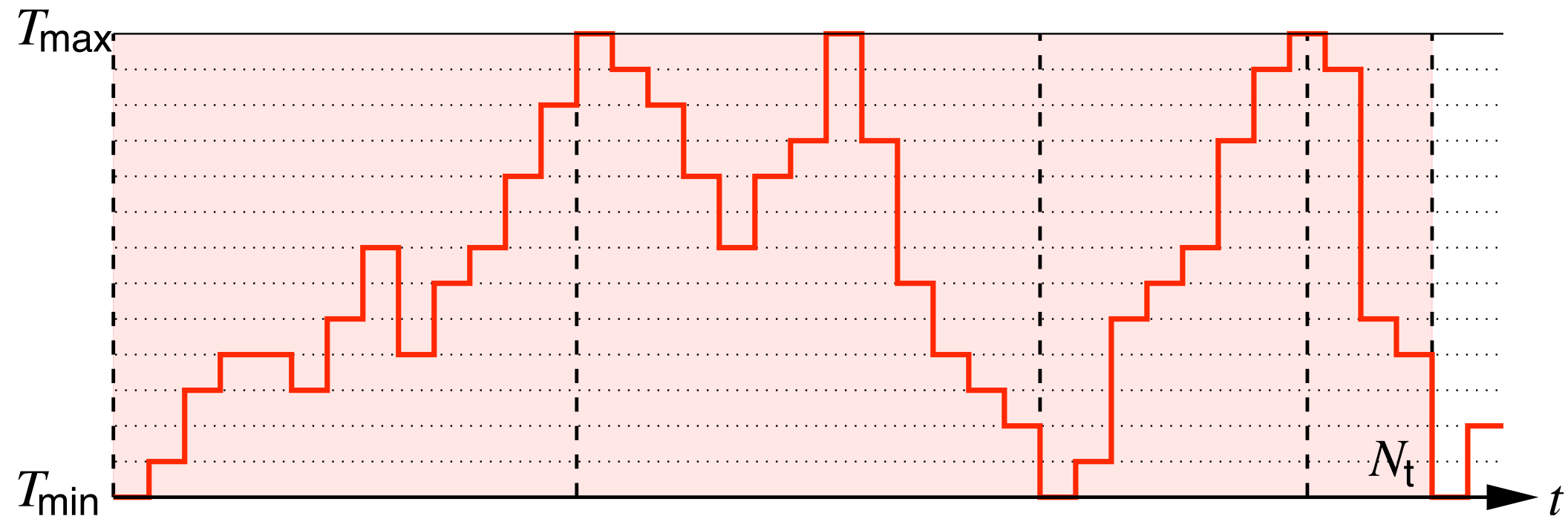


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# Feedback method

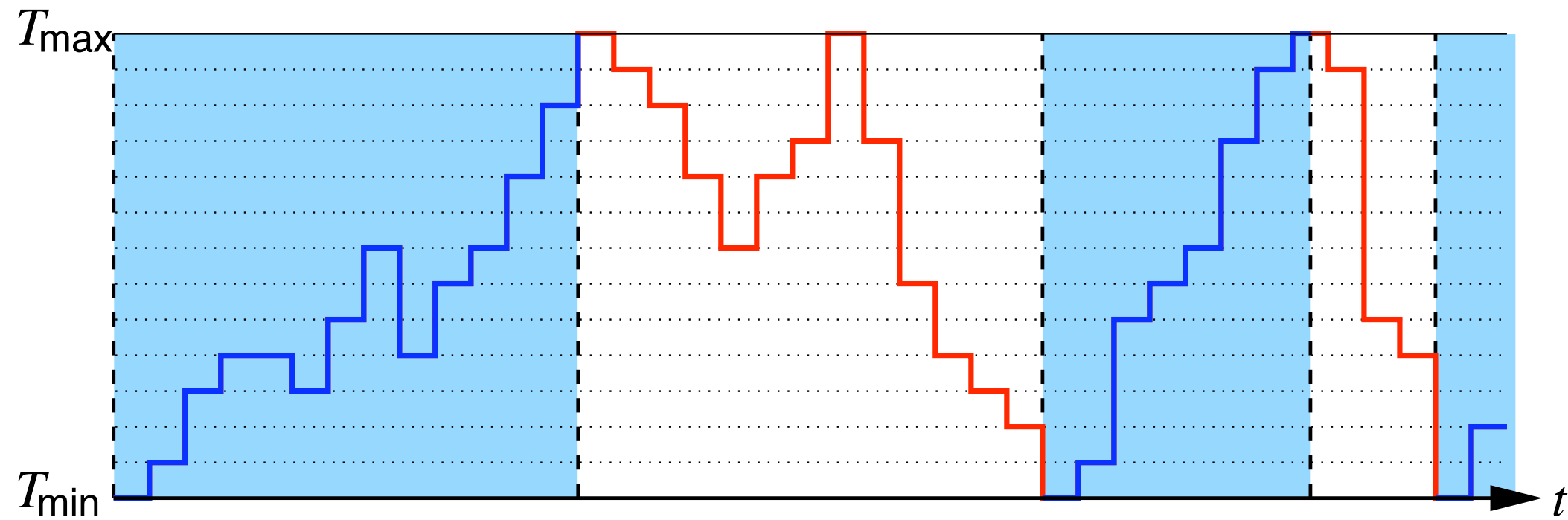
Katzgraber et al., JSTAT (06)



- Track replicas for  $N_t$  MCS and compute a **histogram  $n(T)$**  of the number of times a replica hits a given  $T$ .
- Compute a **histogram of directed walkers  $n_{\text{up}}(T)$** .
- Calculate the fraction  $f(T) = n_{\text{up}}(T)/n(T)$  [ $D(T) = (df/dT)^{-1}$ ].
- Calculate  $\{T'_i\}$  from  $\{T_i\}$ :
 
$$\frac{1}{\Delta T'_i} = C \sqrt{\frac{1}{\Delta T_i} \frac{df}{dT_i}} \sim \frac{1}{\sqrt{\Delta T_i D(T_i)}}$$
- Choose  $C$  such that  $[T'_{\min}, T'_{\max}]$  maps back to  $[T_{\min}, T_{\max}]$ . **Iterate!**

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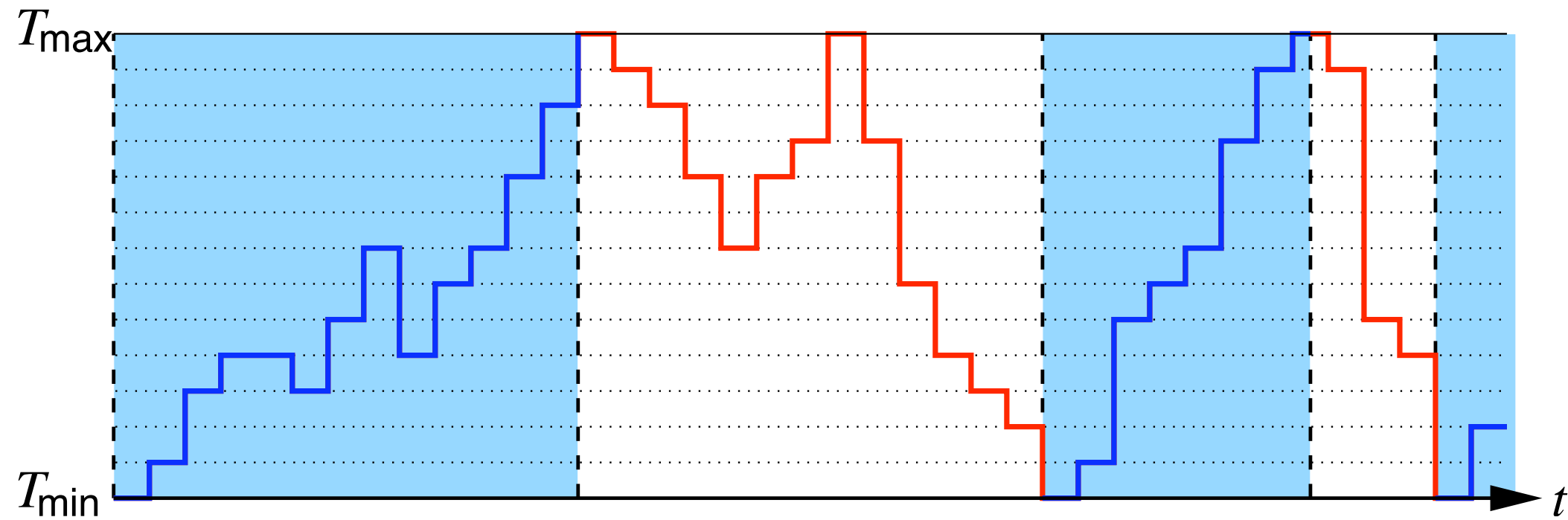
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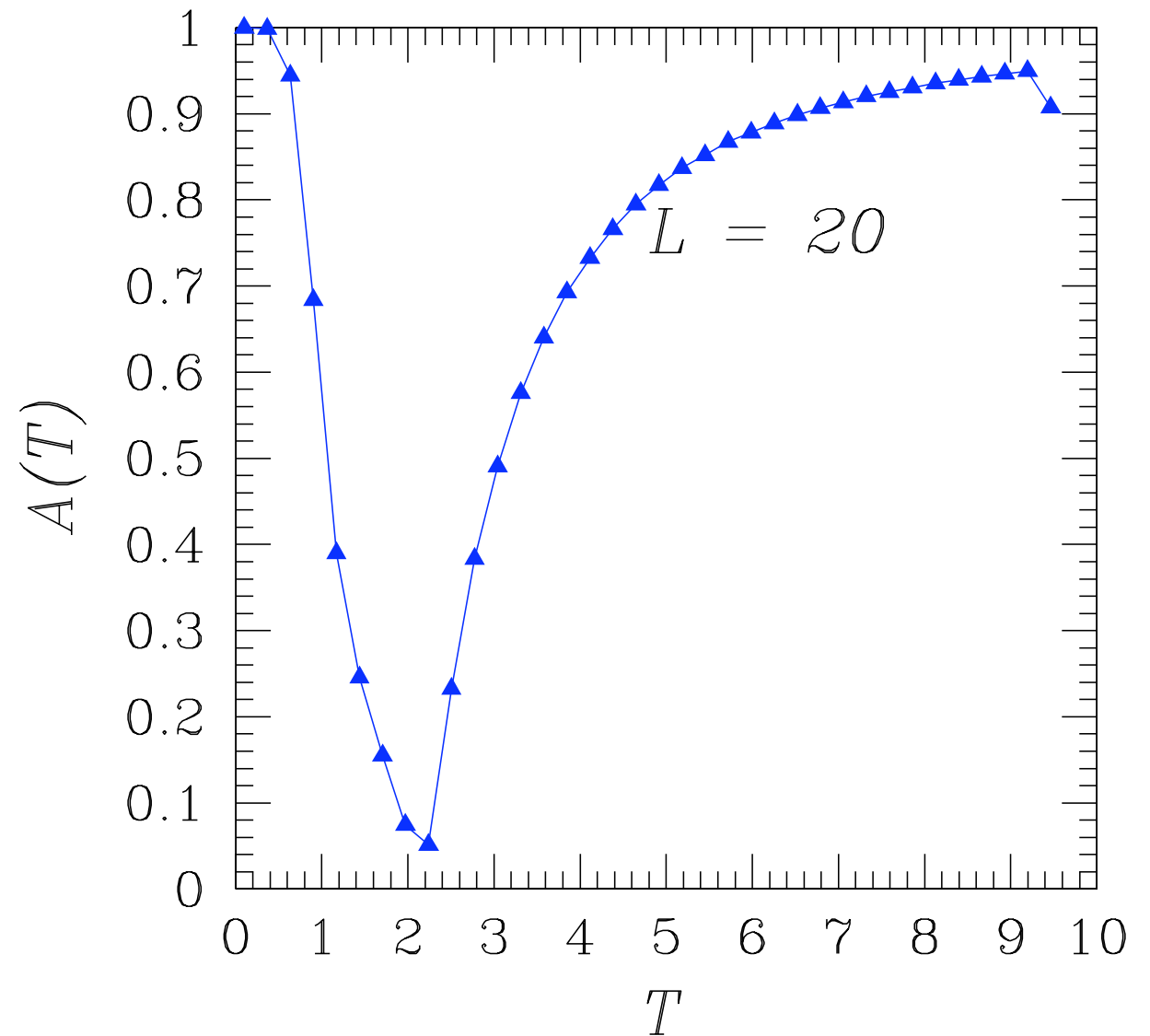
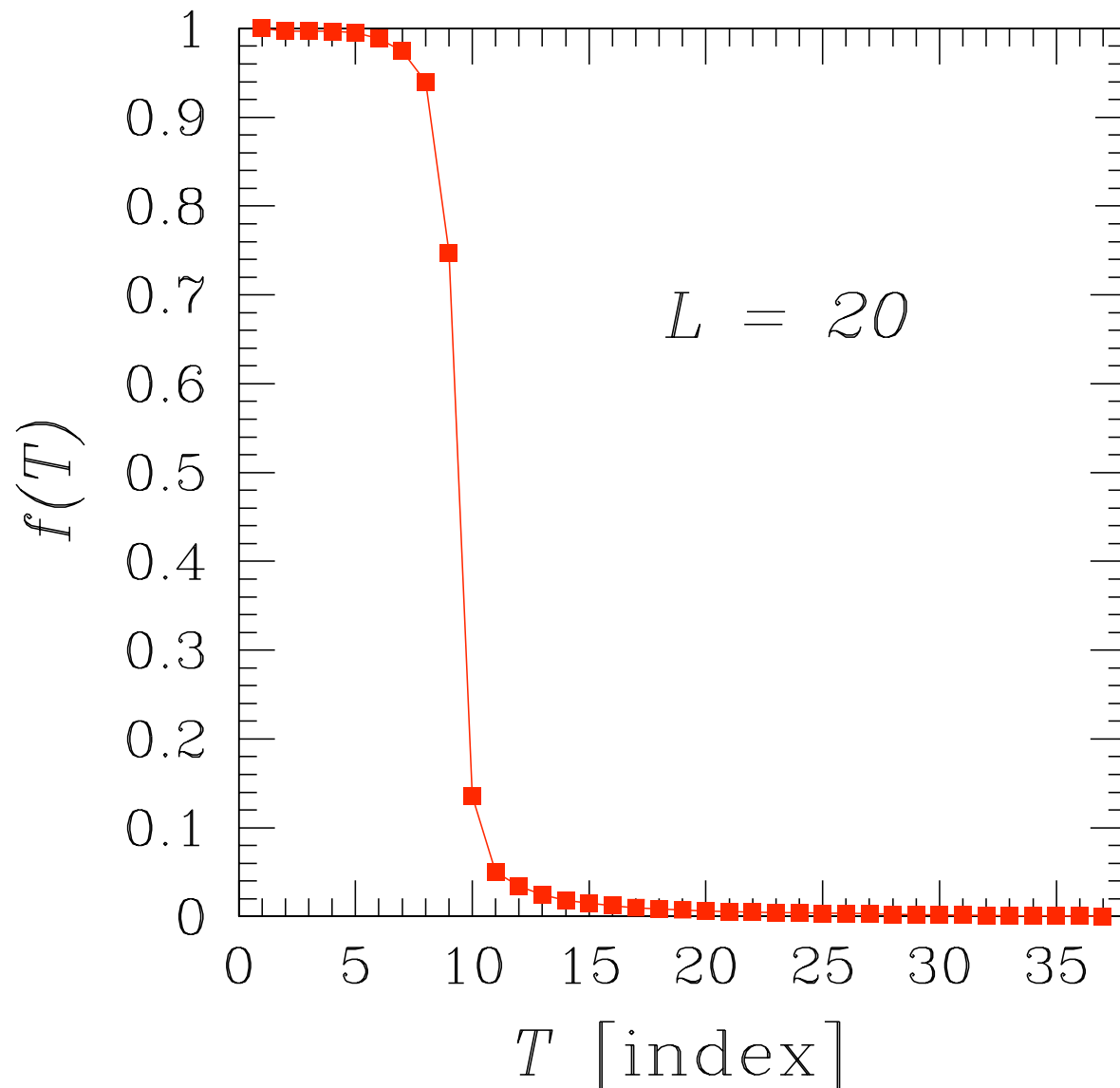
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Katzgraber et al., JSTAT (06)



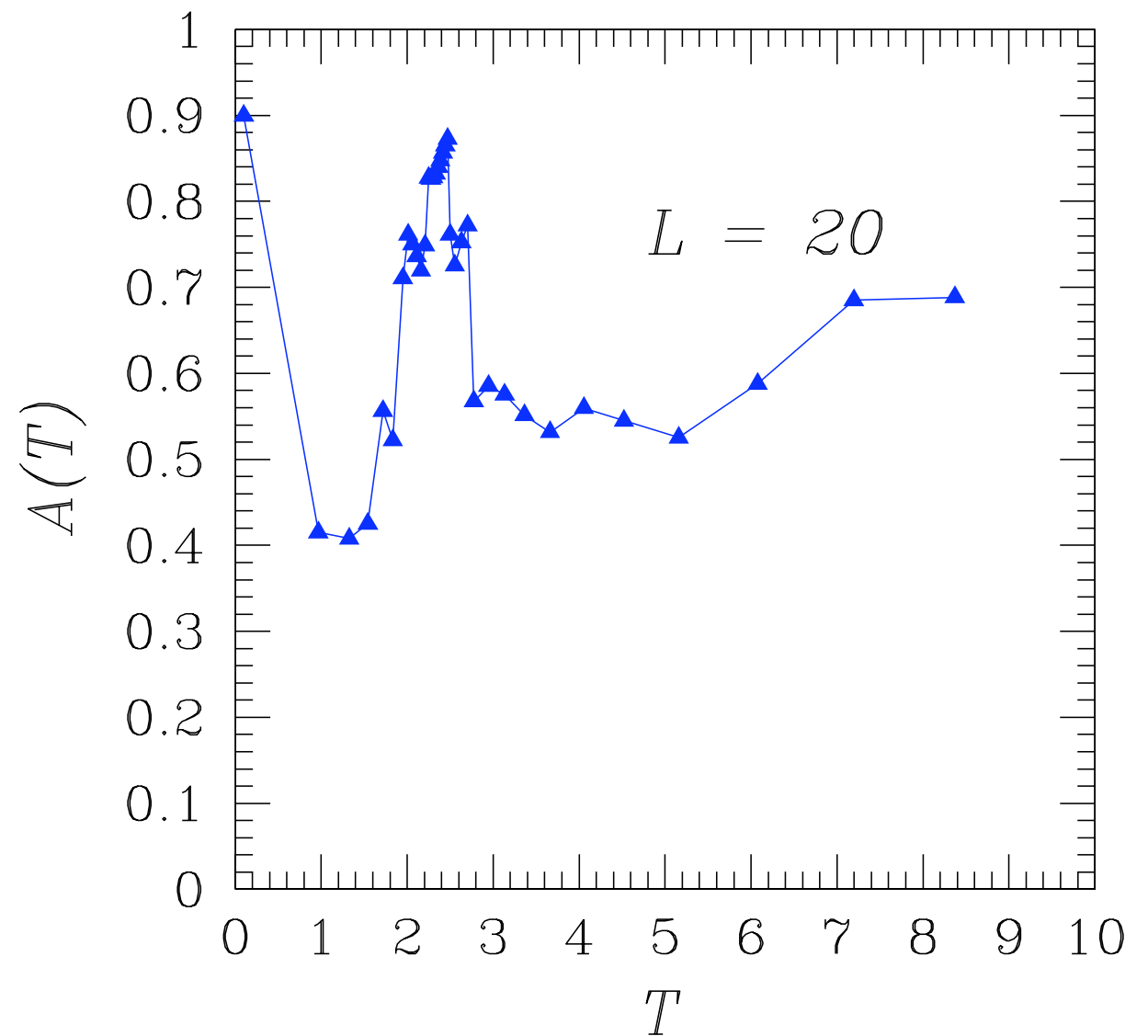
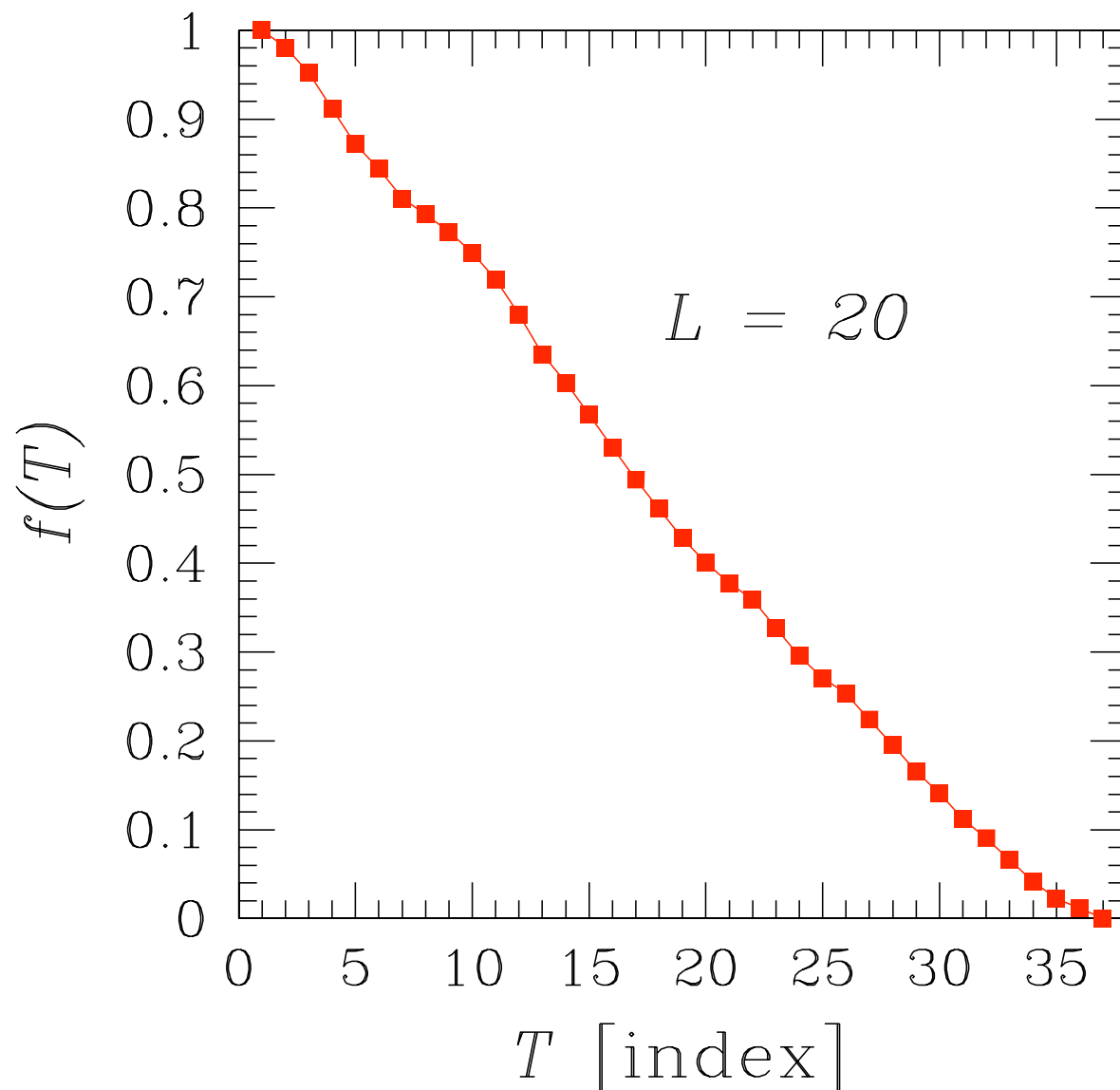
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# Feedback method: Ising model



- Start from a geometric progression temperature set (not good!).
- Feedback maximizes the **rate of round trips**.
- Feedback reallocates resources where needed (critical point):  
**Acceptance rates not constant.**

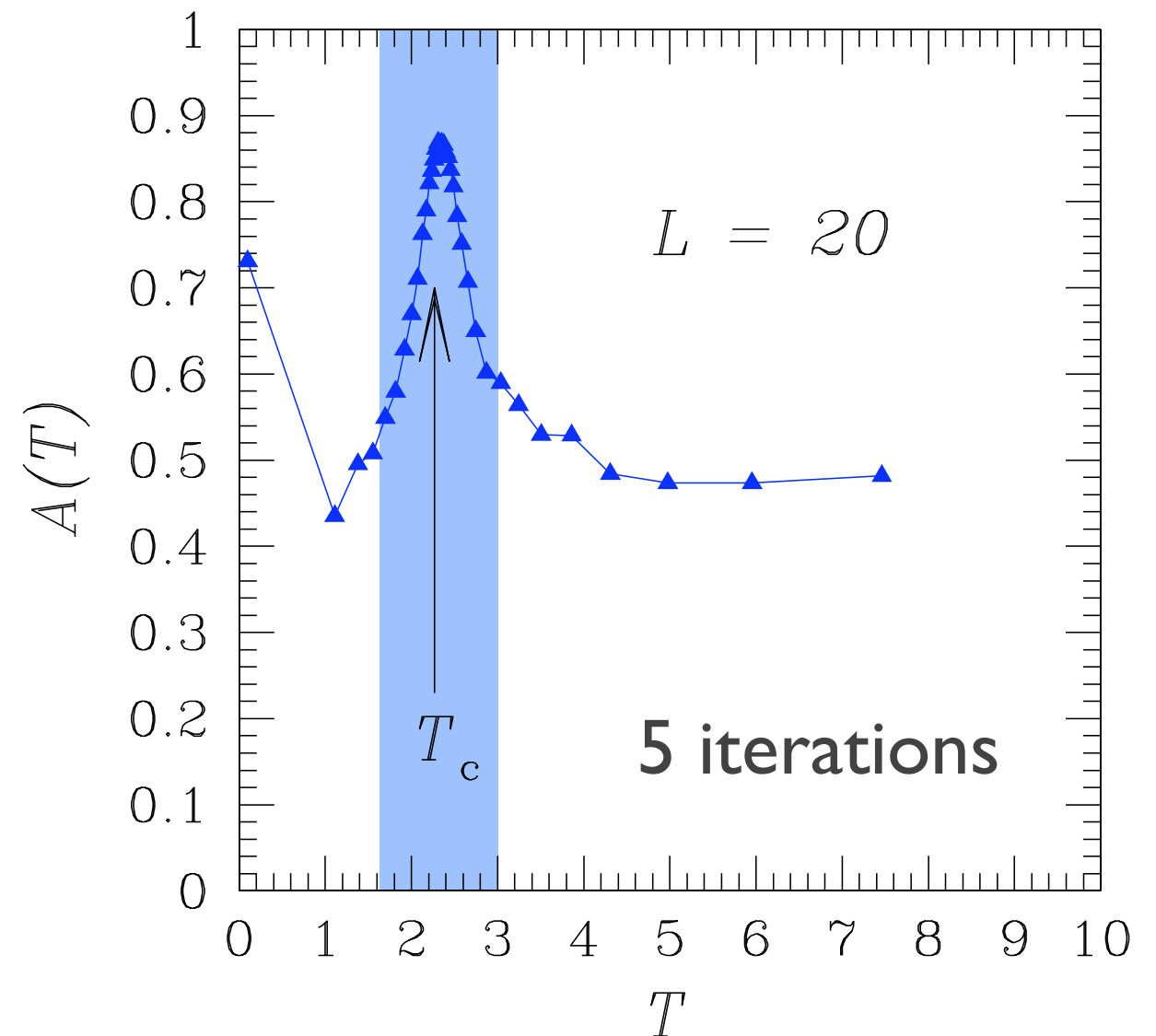
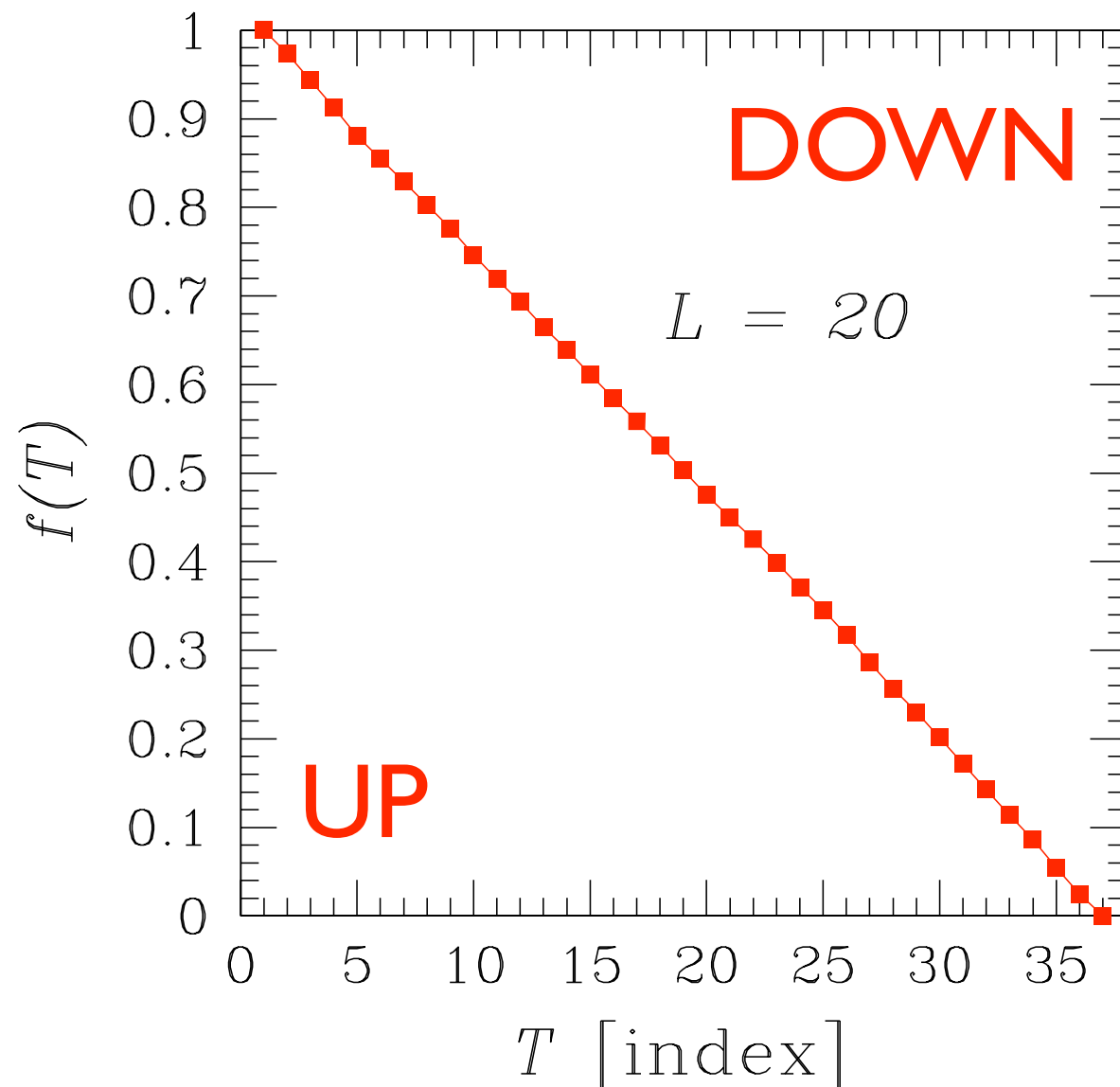
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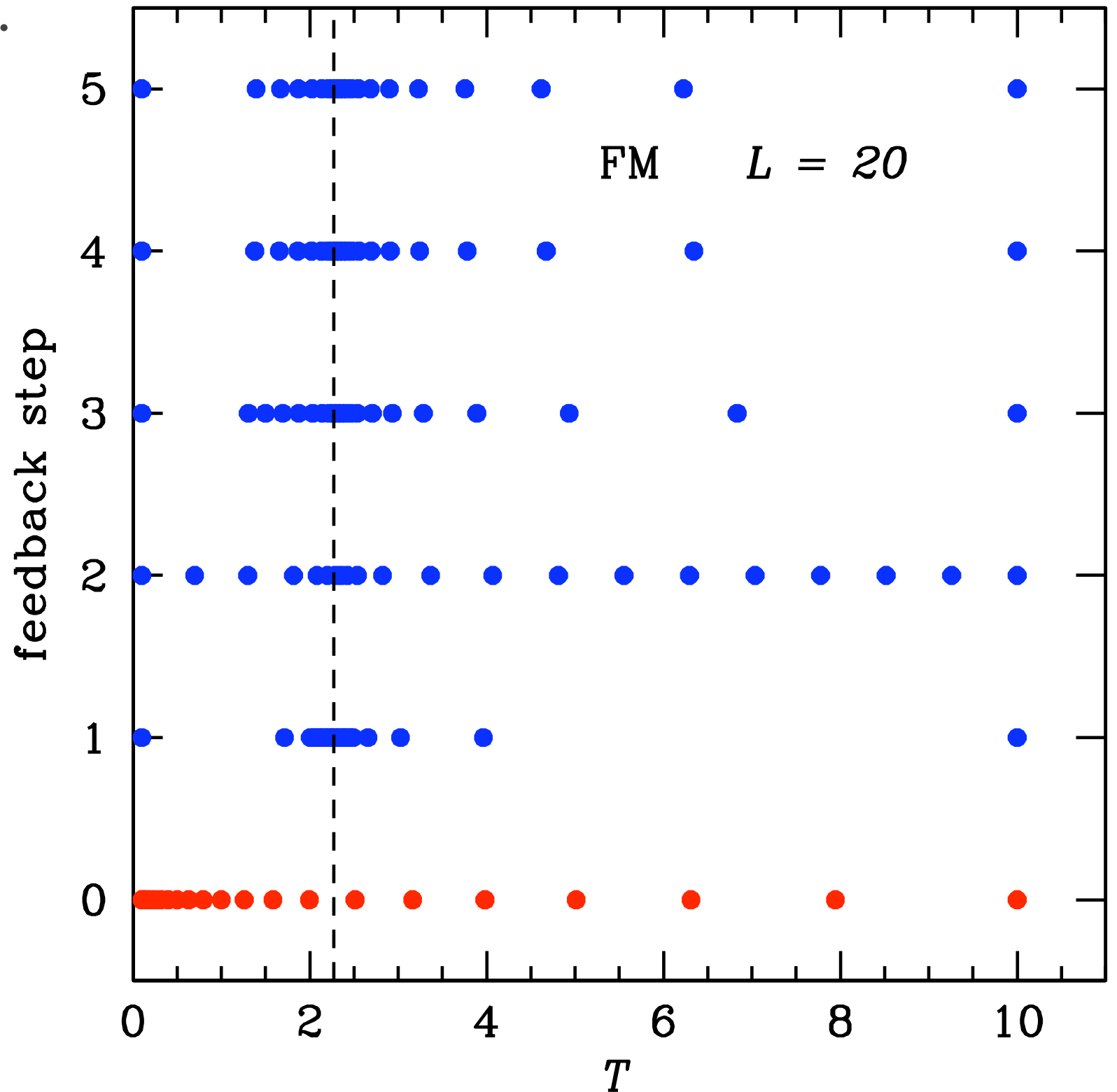
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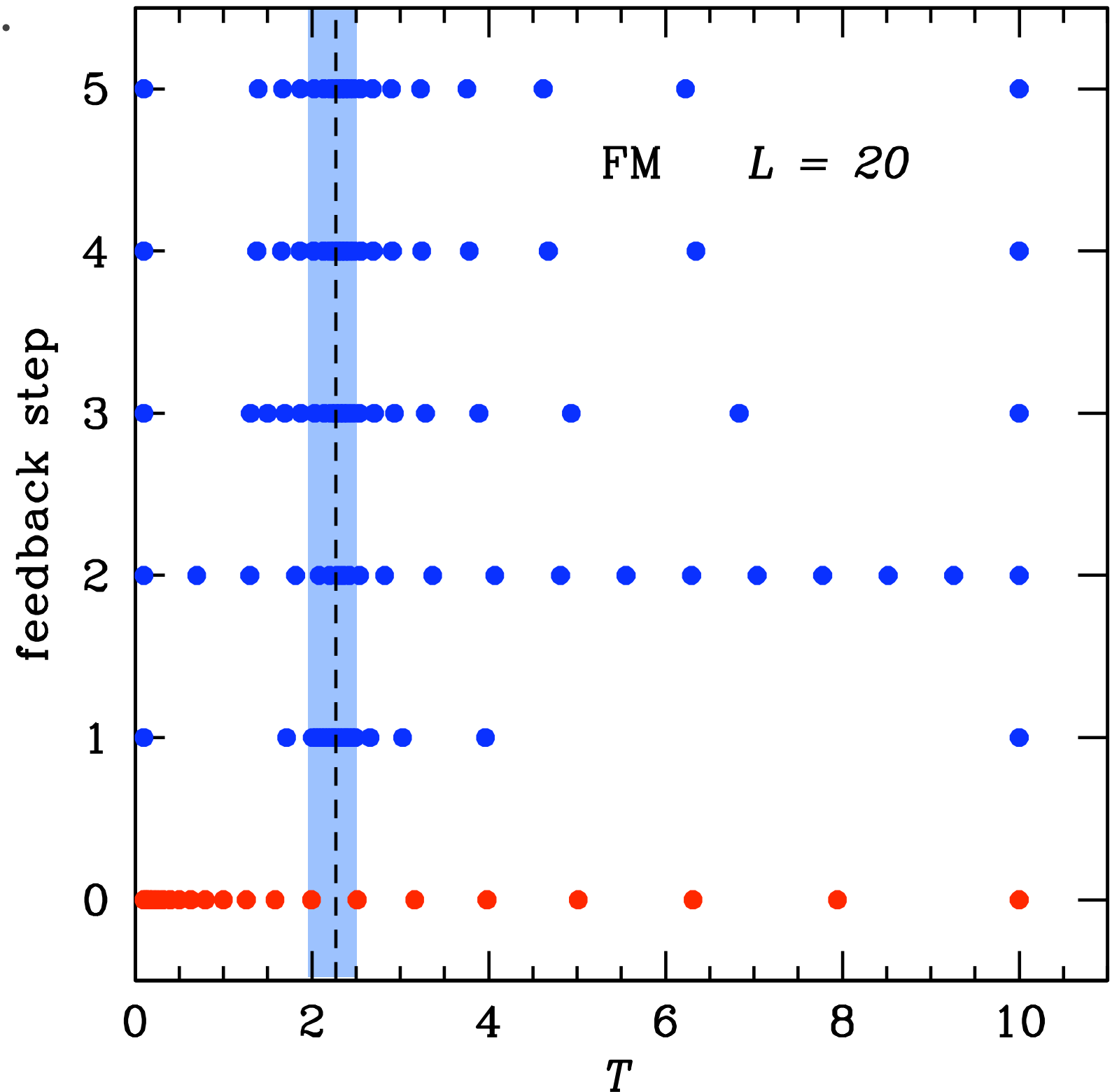
# Evolution of the temperature set

- Data for the Ising model.
- After few iterations the temperature set converges.
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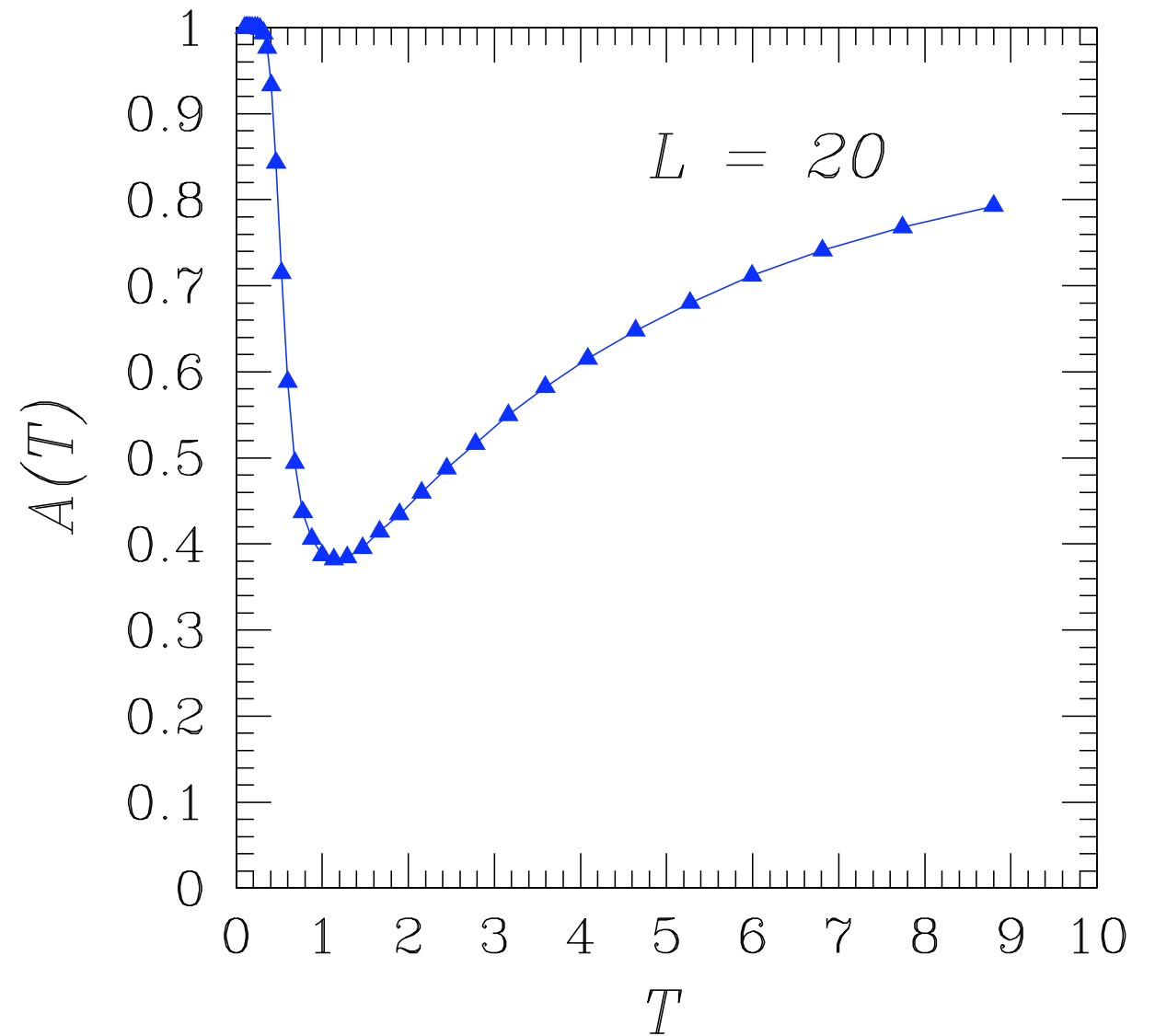
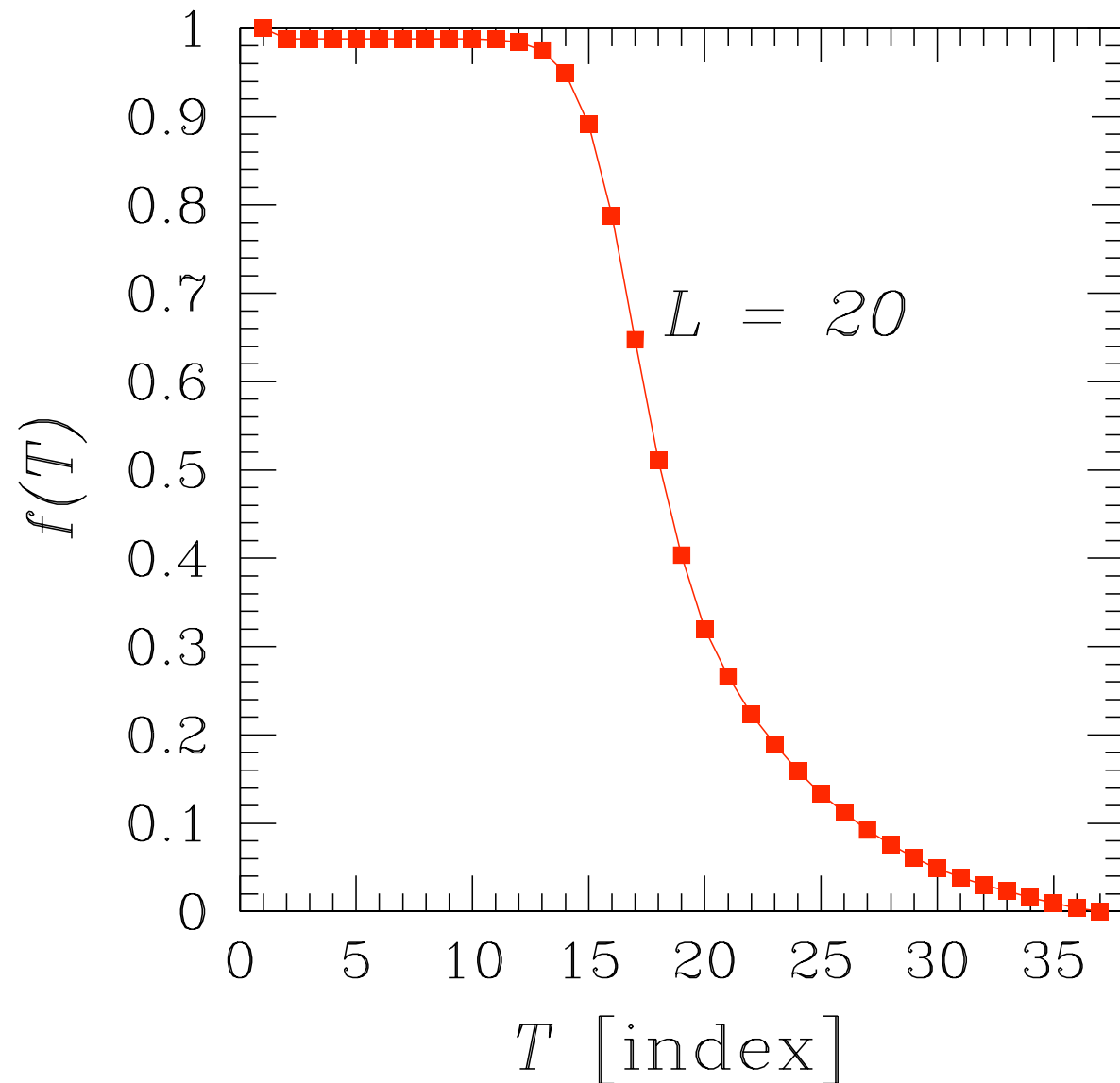


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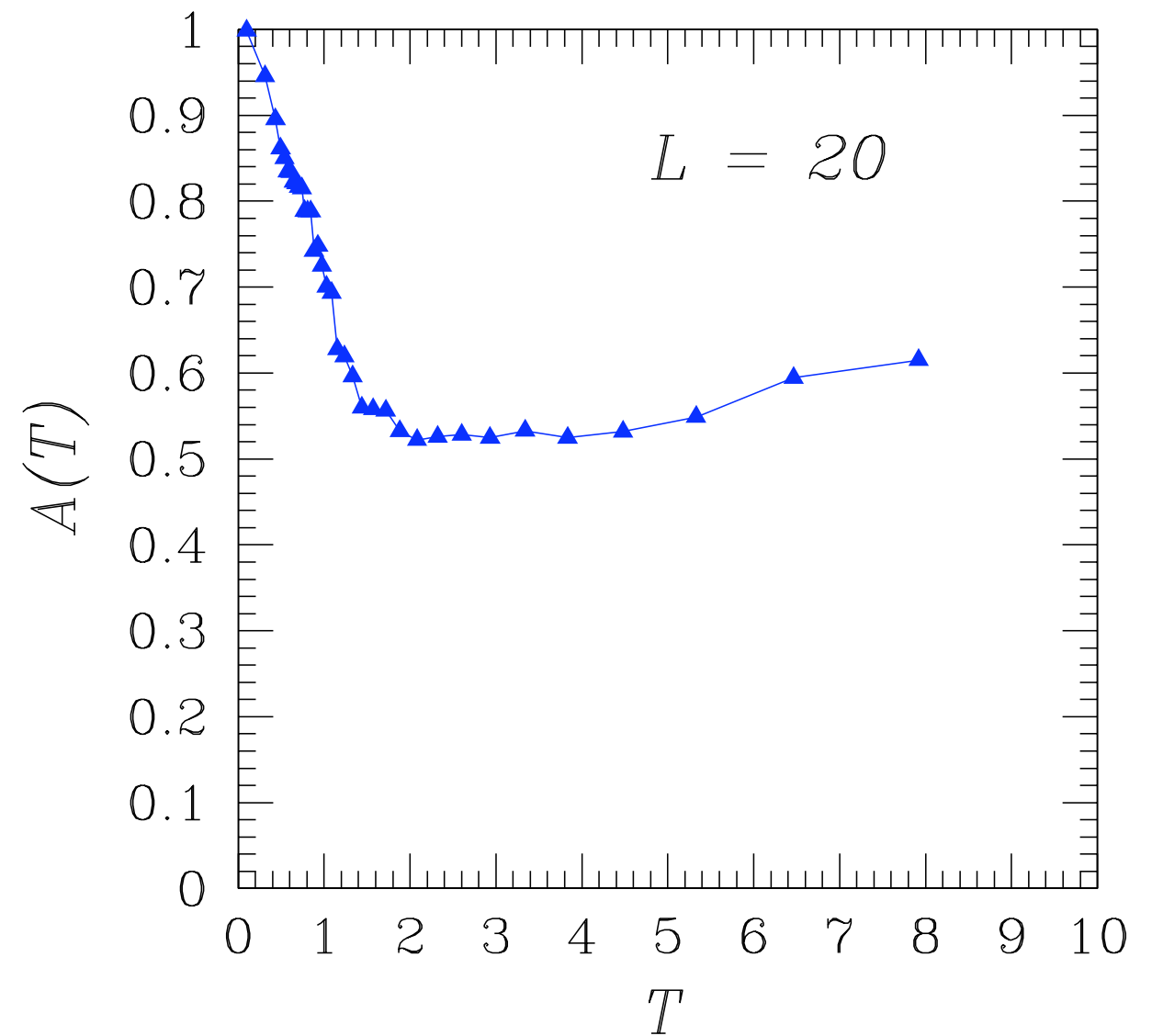
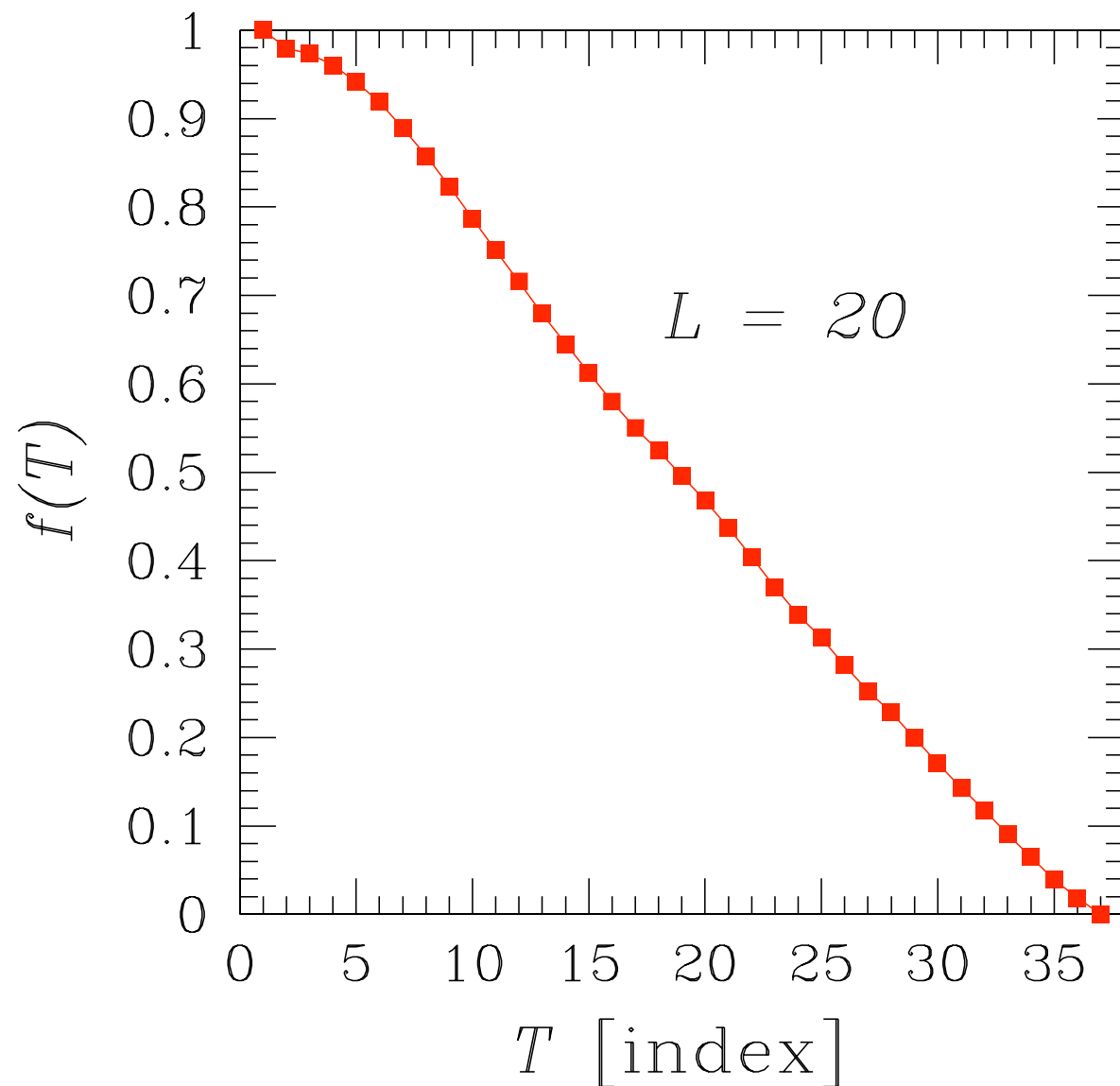


# Feedback method: FFIM



- Start from a geometric progression temperature set (not good!).
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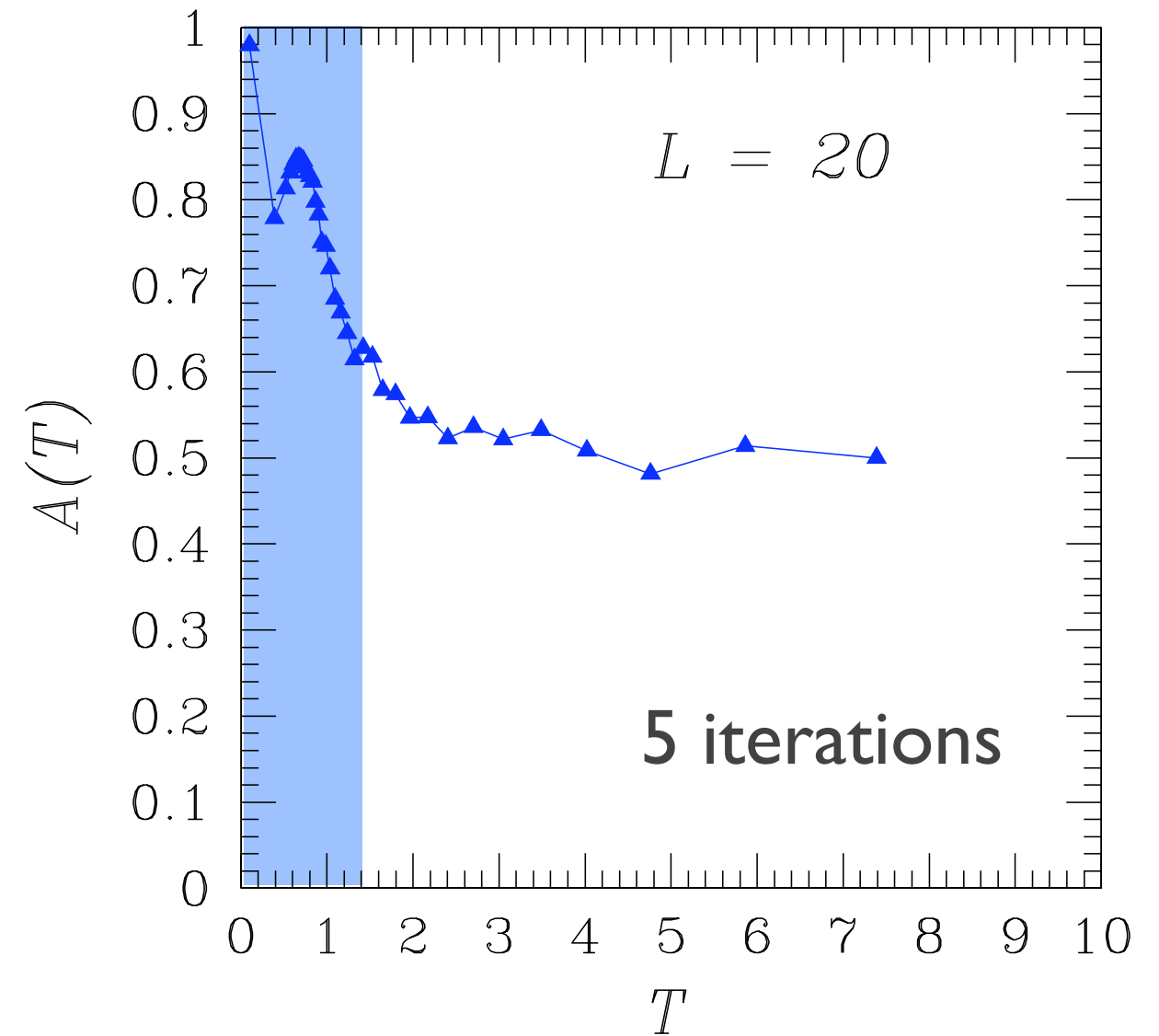
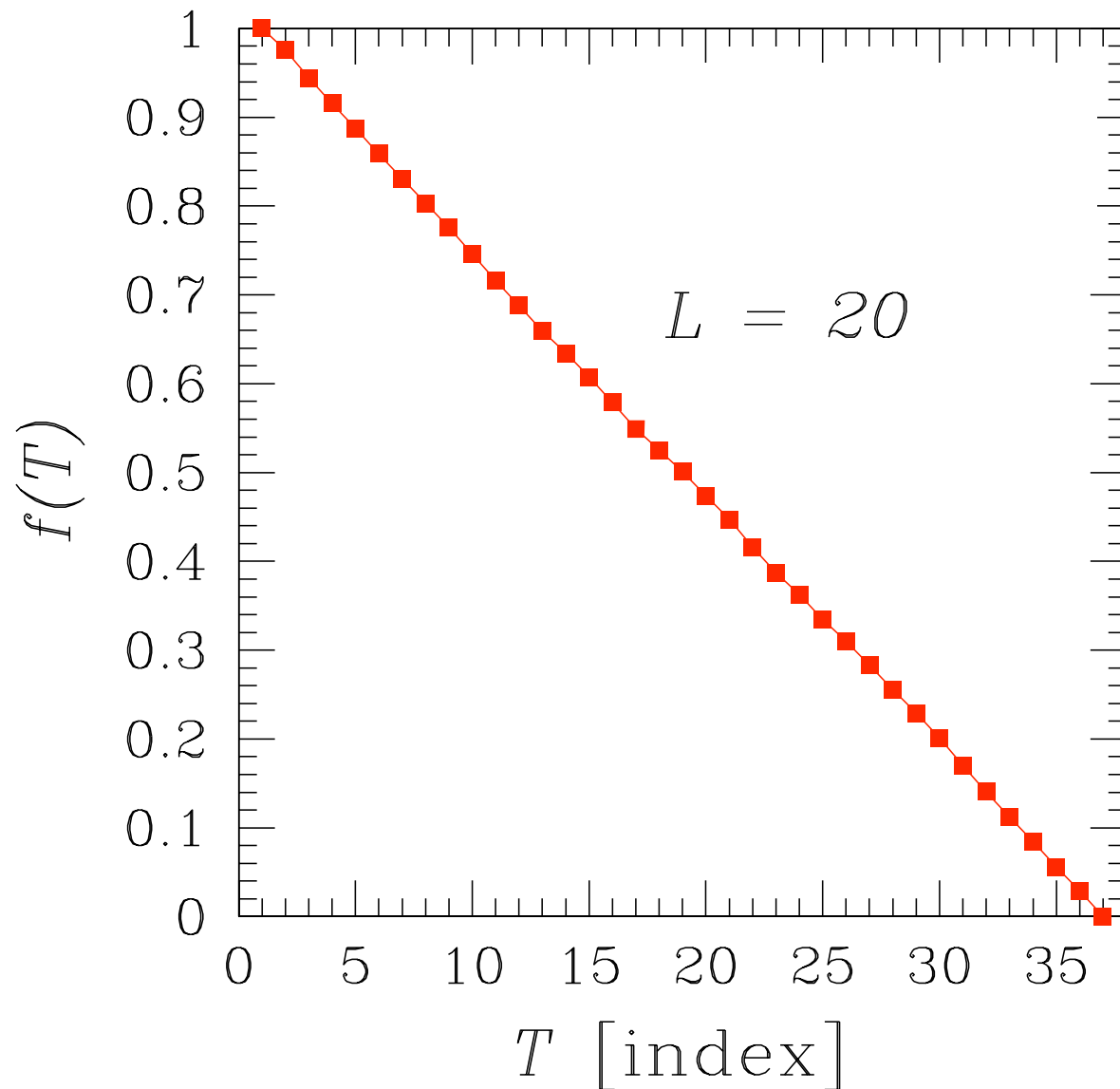
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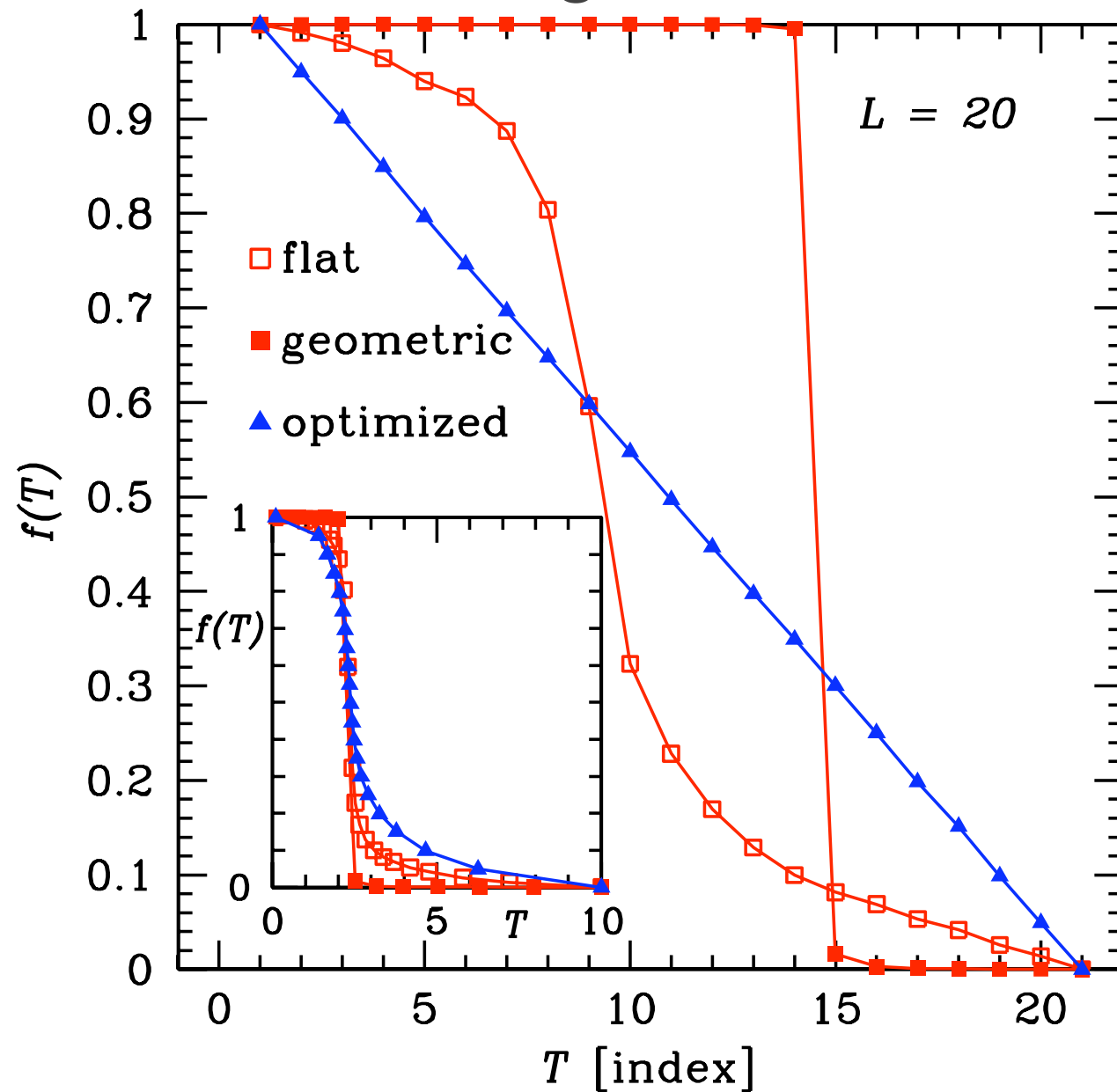
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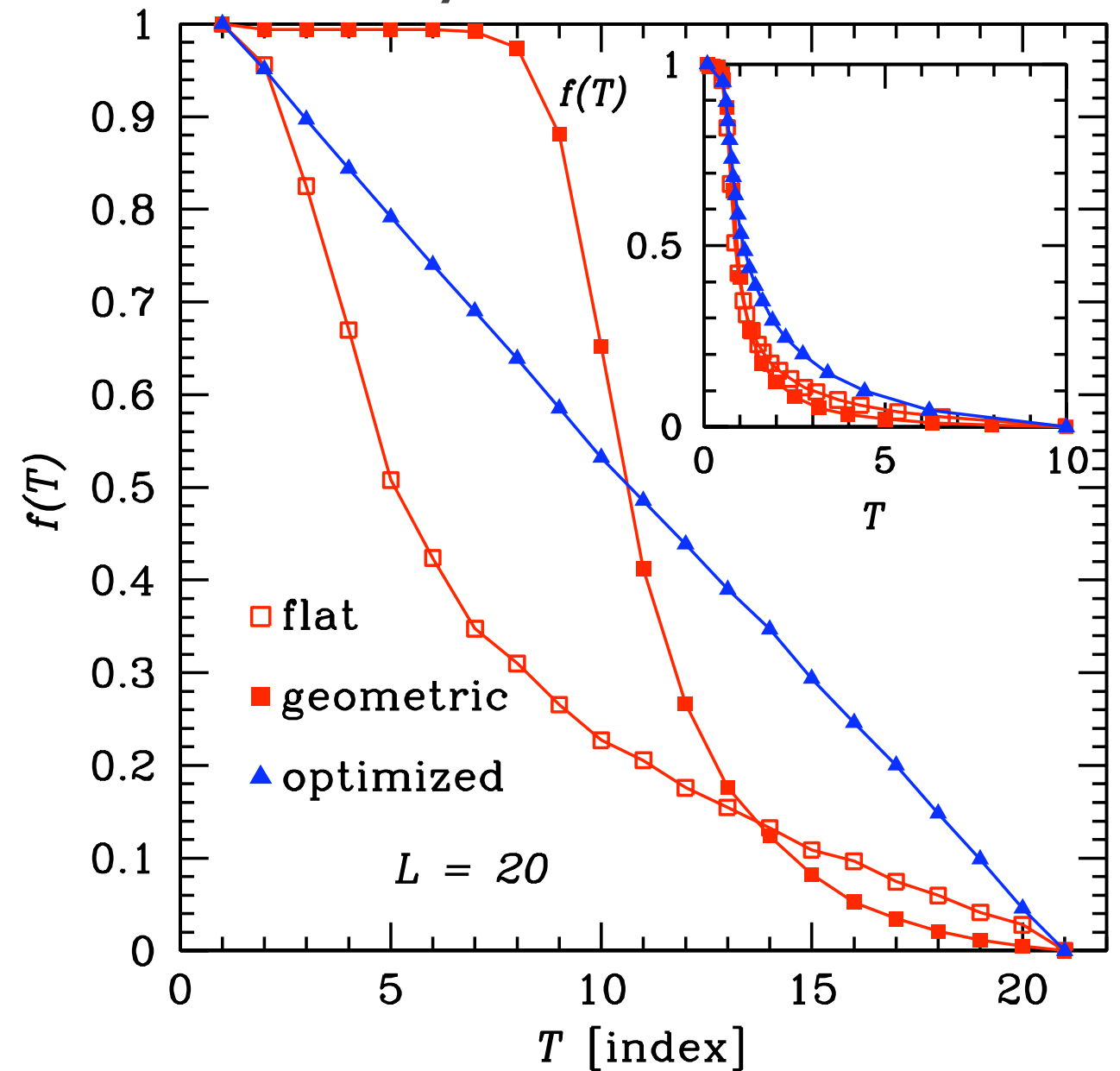
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# Comparison of fractions

Ising model



Fully frustrated model

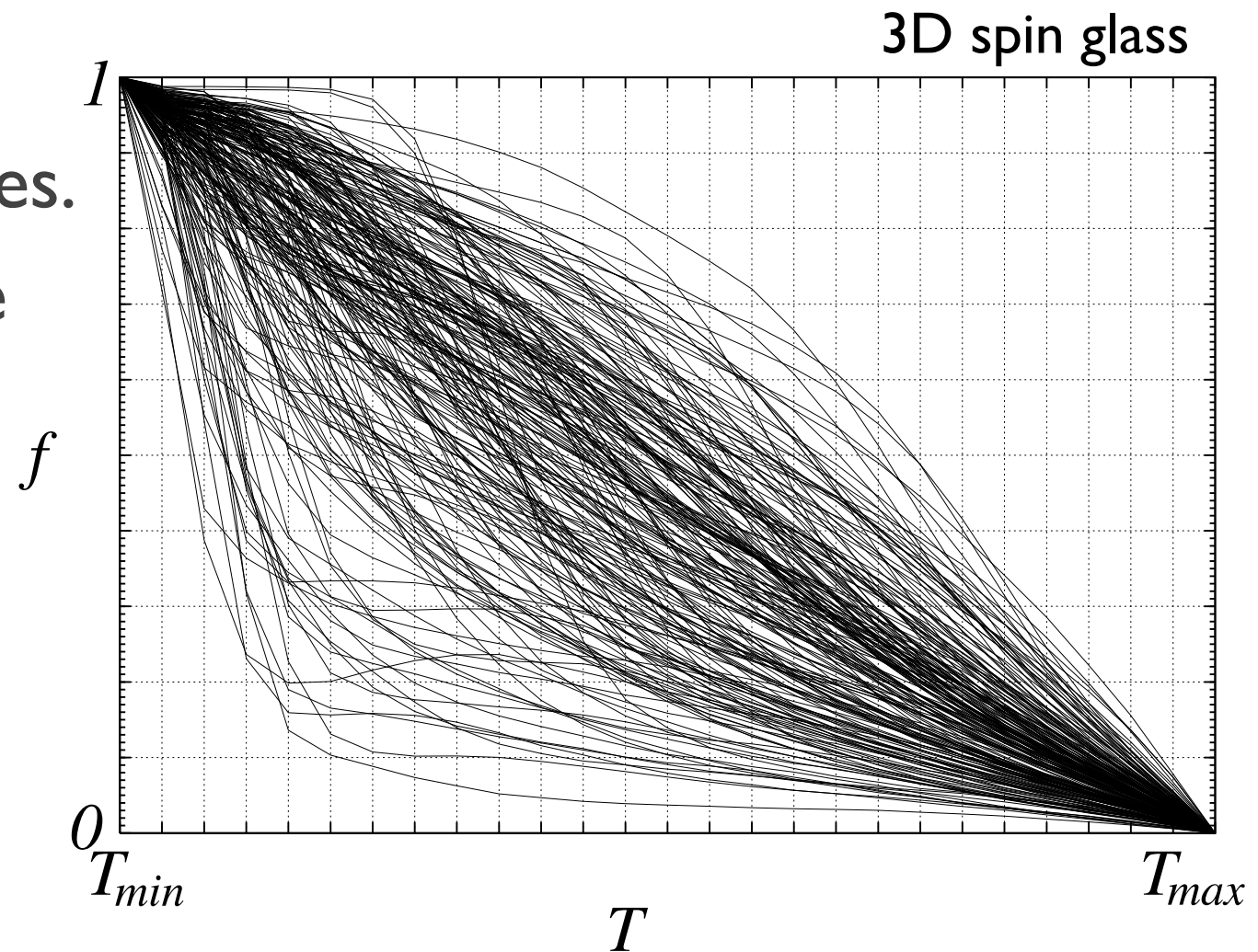


- In general, a “flat” [ $A(T) \sim \text{const}$ ] temperature set is not too bad, but not optimal.
- If  $C_v$  diverges, a geometrical progression is bad.

# What happens for systems with disorder?

- **Problems:**

- We need to deal with averages.
- Each fraction depends on the given disorder configuration.
- Each disorder configurations converges independently...
- The round-trip times are fat tail distributed.
- Each disorder realization has its own temperature set.



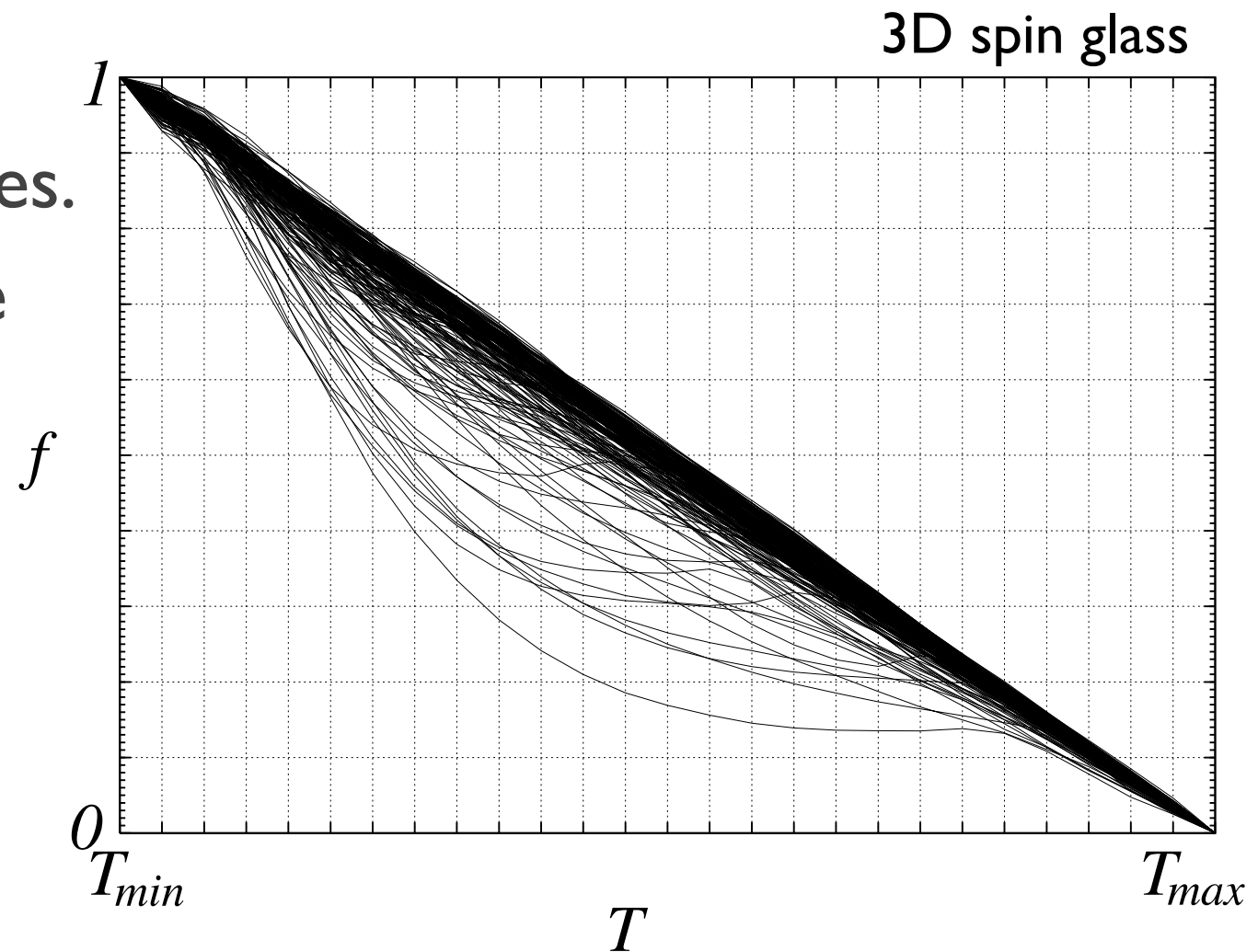
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- Fortunately  $C_v \sim \text{const.}$  and a “flat” temperature set is close to optimal “on average.”

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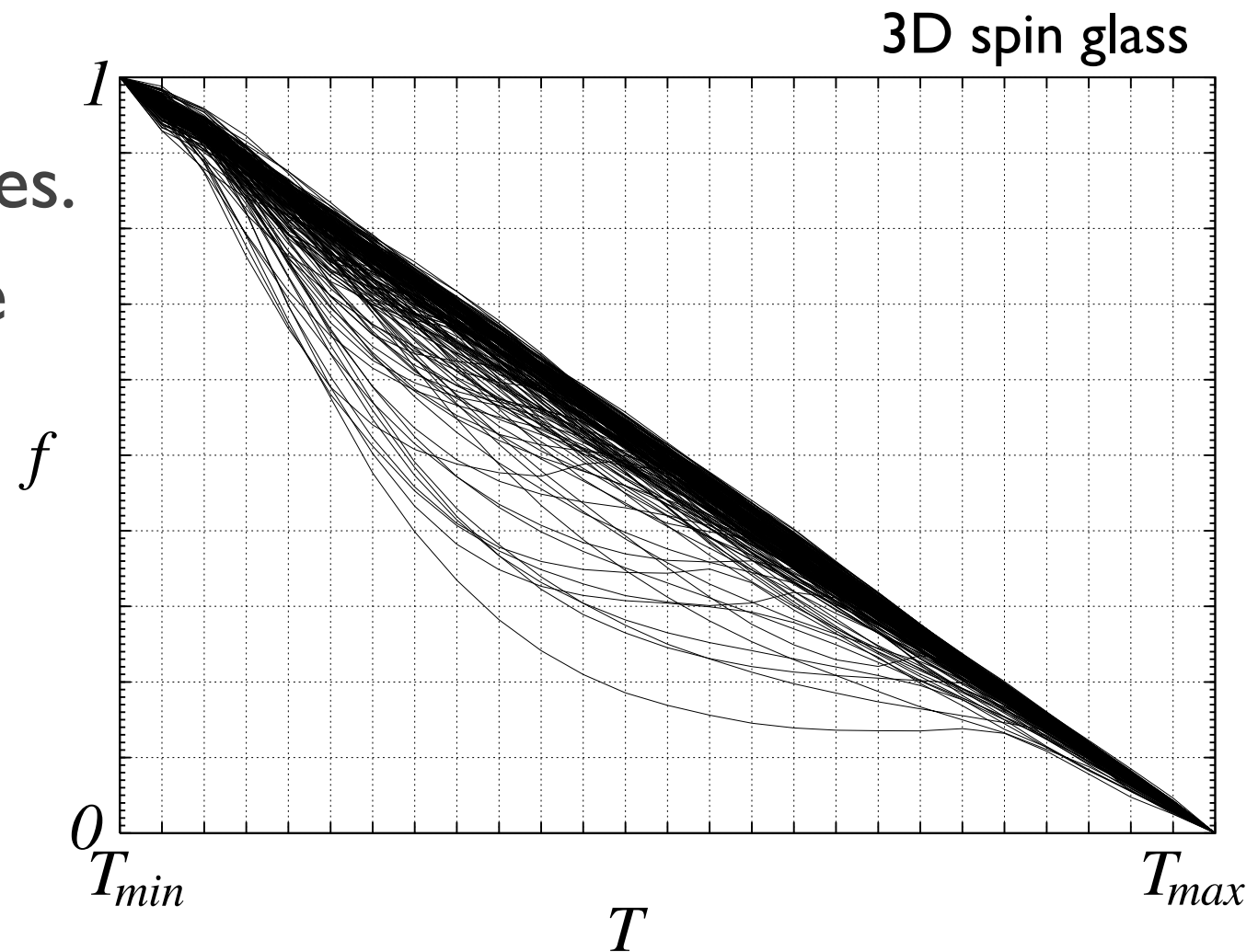
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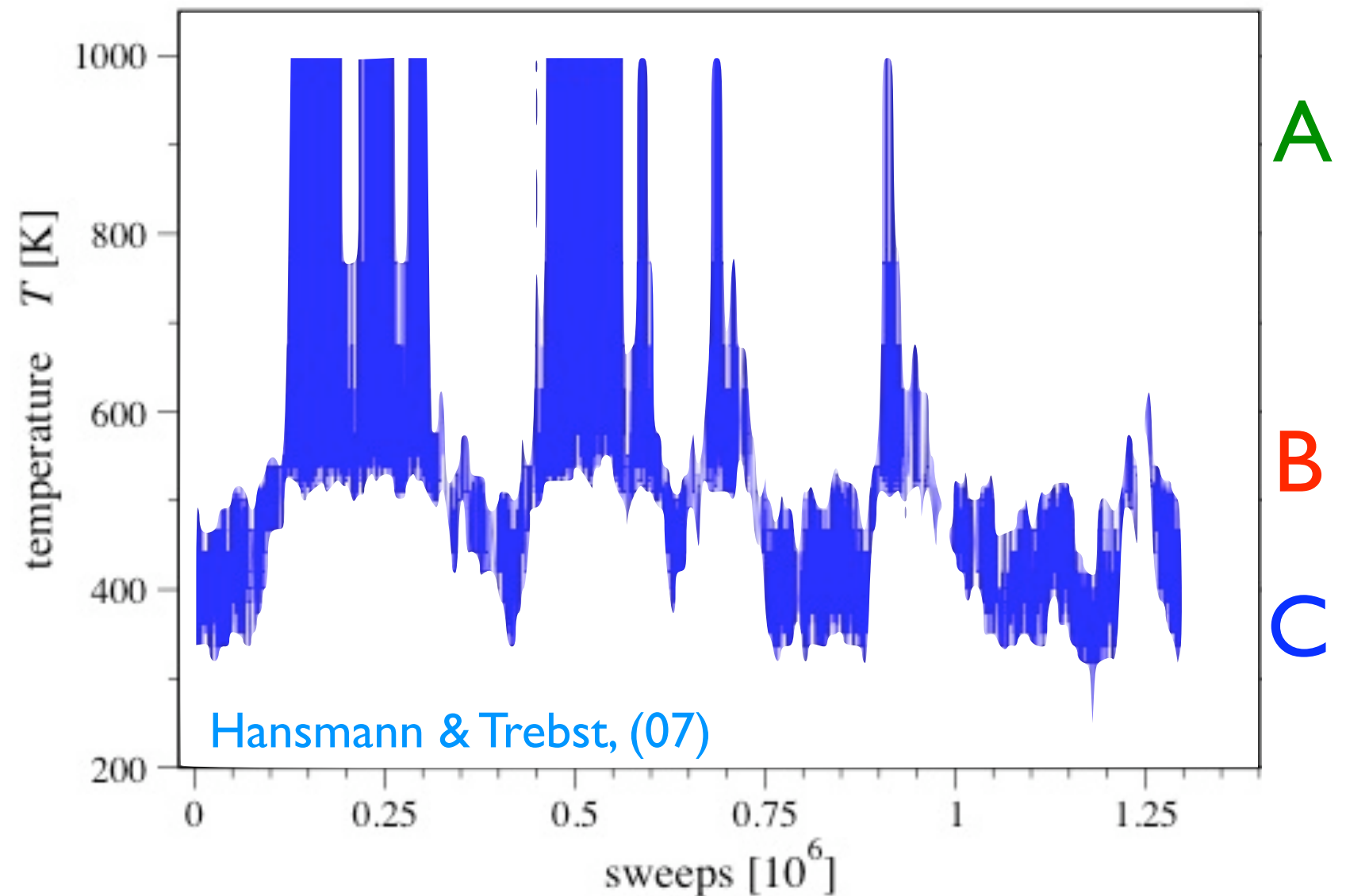
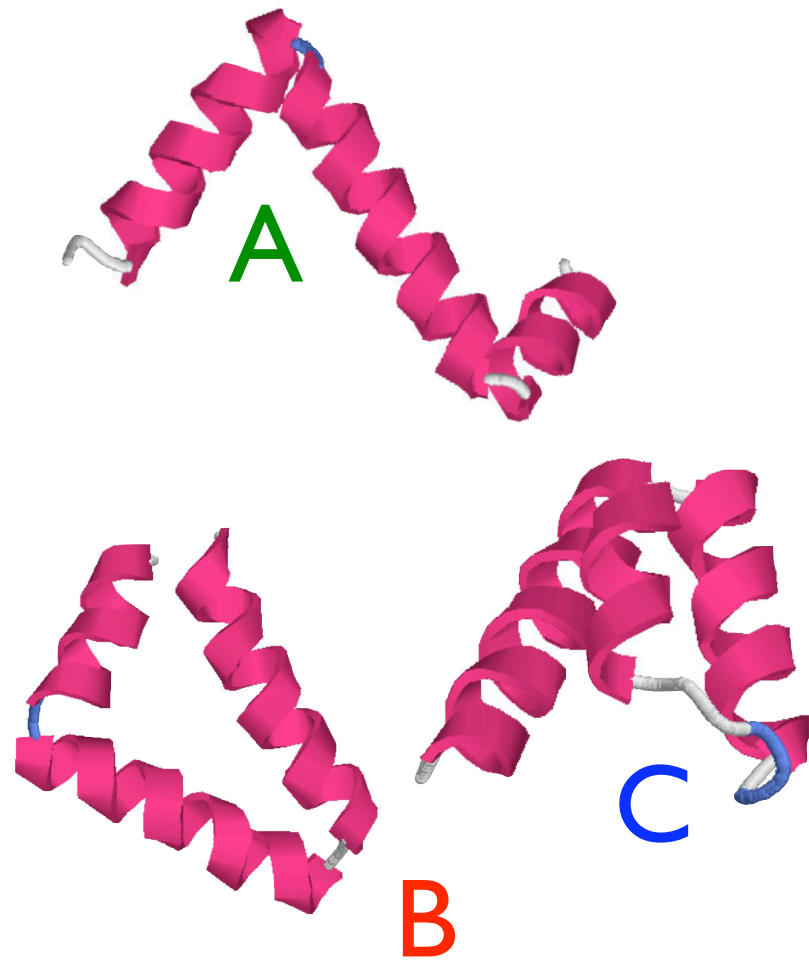
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Do not use for glasses...



# Simulations in (bio)chemistry

- **Example:** Protein A

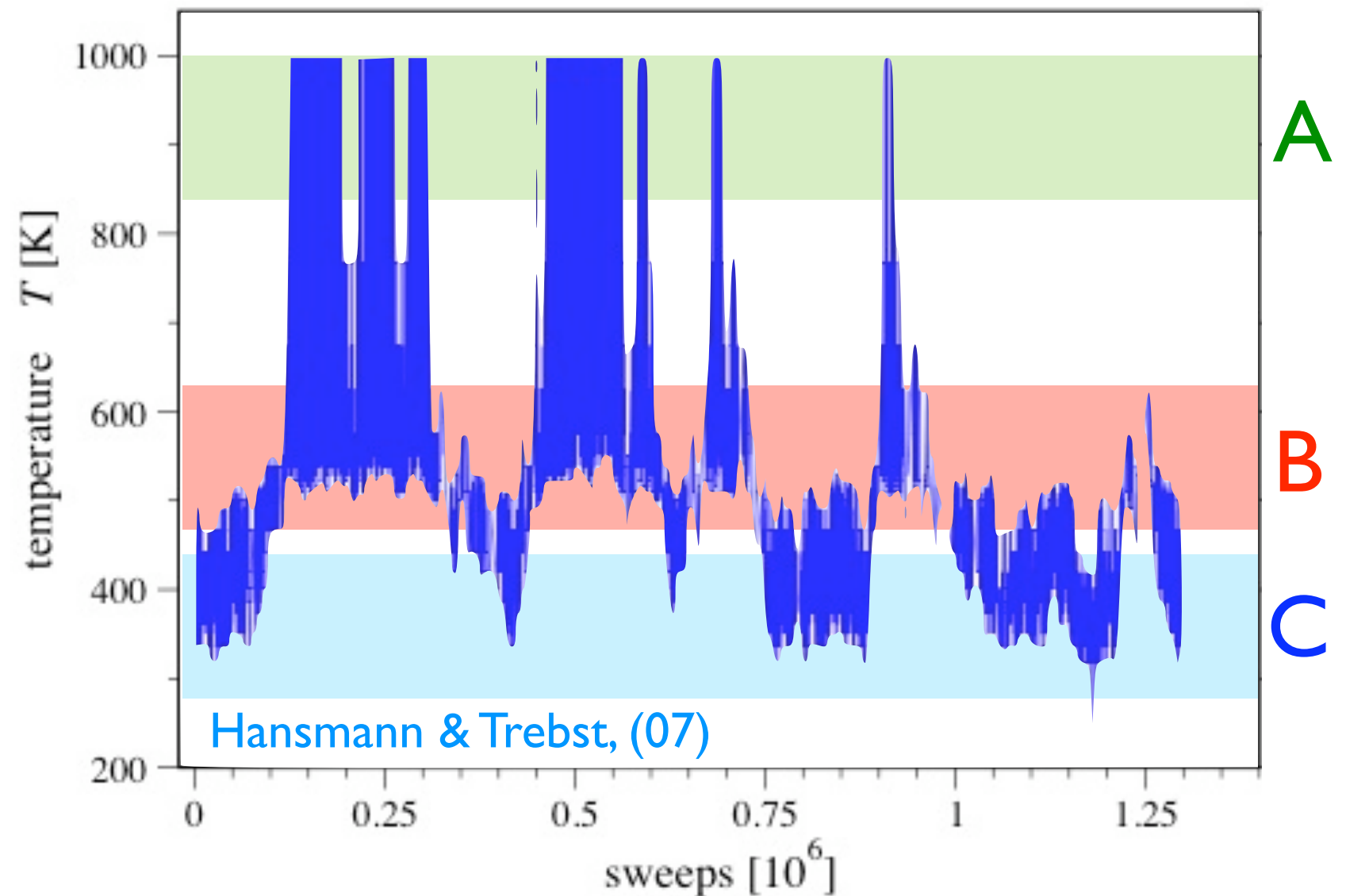
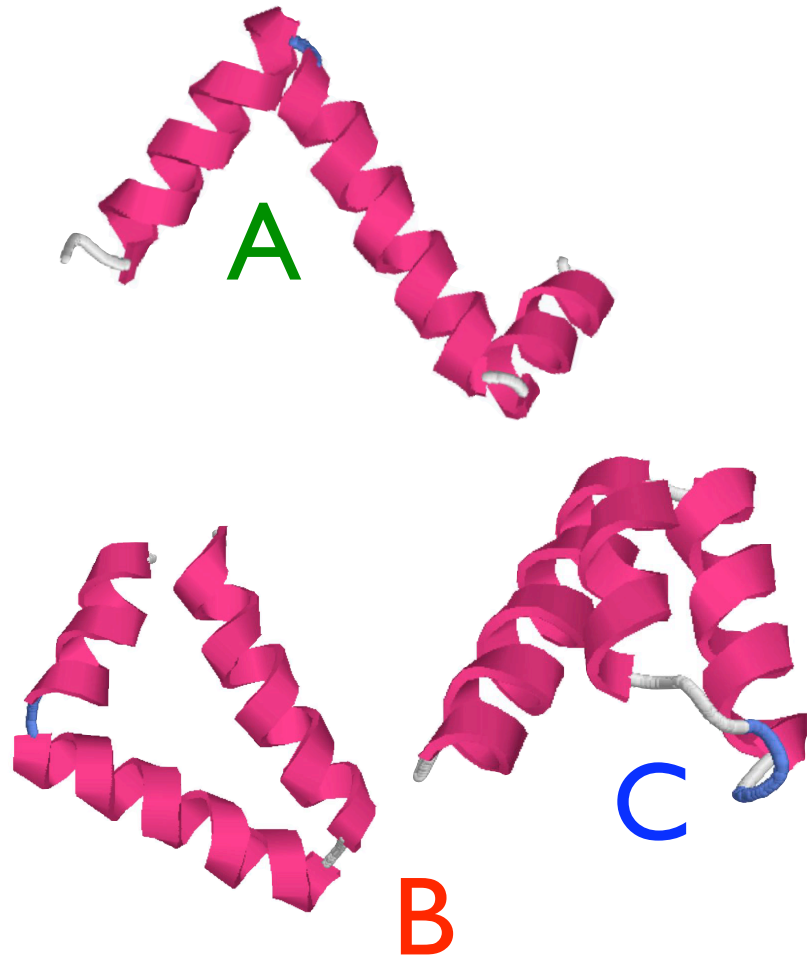


- It can happen that the replicas will only move in subspaces of the phase space (A, B, C).
- Feedback optimization helps overcome these problems easily. Simulations otherwise impossible.



# Simulations in (bio)chemistry

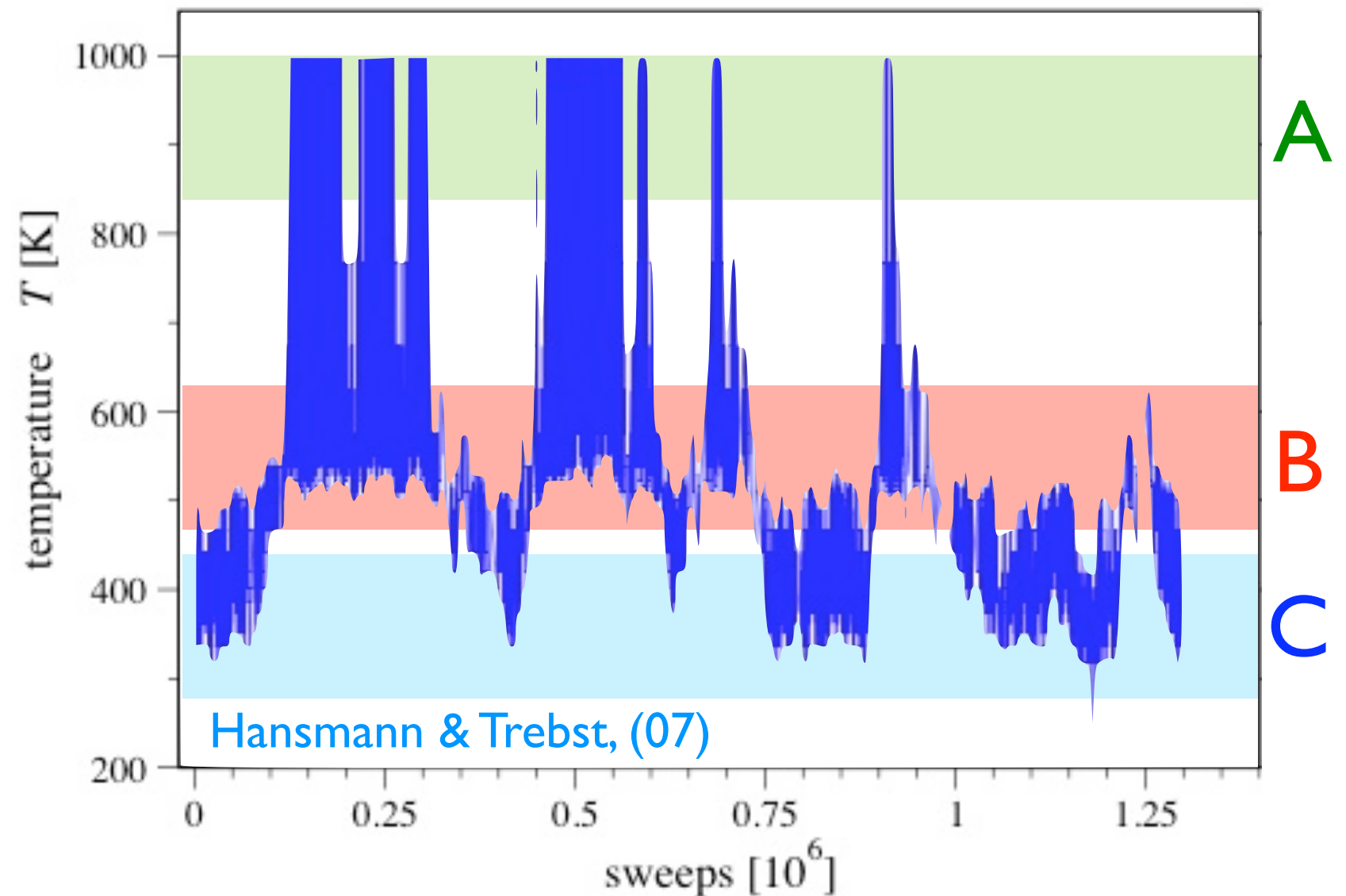
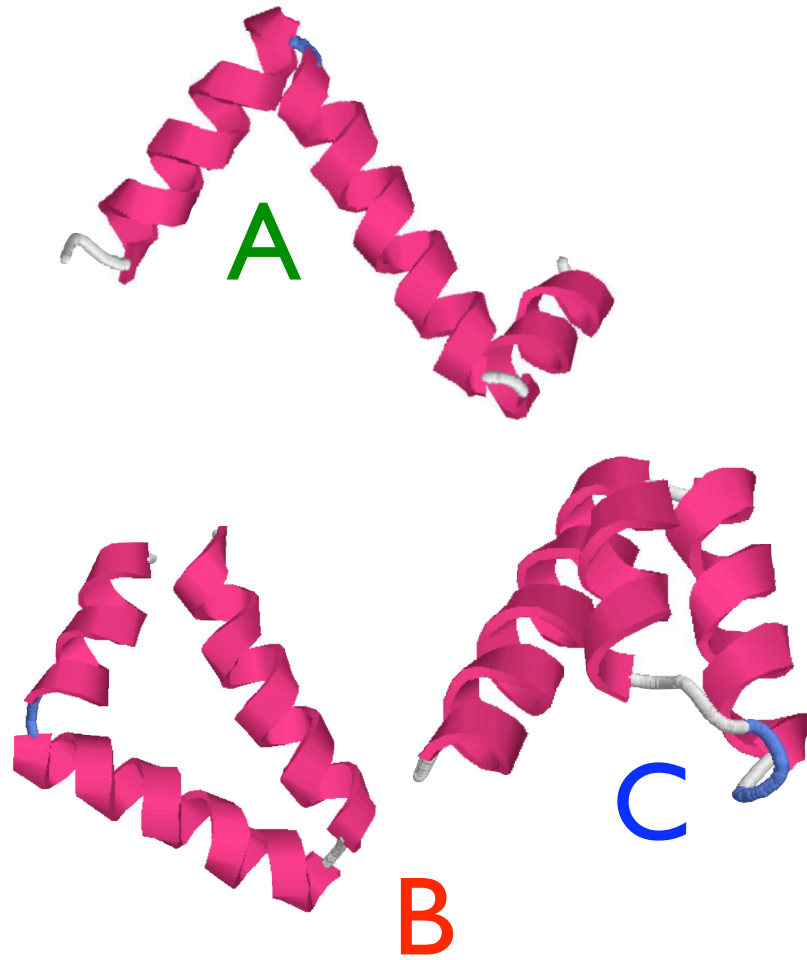
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Use for biology applications...

# Some final thoughts (and extensions)...

# Heuristic ground state search

- Outline of the algorithm:
  - Perform an exchange MC run with  $T_{\min}$  close to zero.
  - Simulate two copies of the system with different Markov chains.
  - The system is in thermal equilibrium:
    - after each lattice sweep record the spin configuration and energy for the lowest  $T$  if these match in both copies.
    - If a lower energy is found, replace the previously recorded energy.
  - Measure for  $1/4t_{\text{eq}}$ .
- For small to intermediate system sizes where other algorithms do not work well the method provides heuristic ground states with high probability (98% up).

# Extensions and combinations

- Any control variable can be used:
  - Field
  - Temperature and field
  - Coupling constants in QCD
  - Frequencies (e.g., in a Holstein model)
  - ...
- Combinations with other algorithms possible:
  - Exchange Monte Carlo molecular dynamics (biomolecules).
  - Exchange quantum Monte Carlo (quantum spin glasses).
  - Bayesian periodogram (planet search in star systems).
  - Iterative search methods (combinatorial problems).
  - Cluster exchange Monte Carlo (diluted spin glasses).
  - ...





Thank you.