Exchange Monte Carlo Method and Application to Spin Glass Simulations

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Abstract

We propose an efficient Monte Carlo algorithm for simulating a “hardly-relaxing” system, in which many replicas with different temperatures are simultaneously simulated and a virtual process exchanging configurations of these replica is introduced. This exchange process is expected to let the system at low temperatures escape from a local minimum. By using this algorithm the three-dimensional $\pm J$ Ising spin glass model is studied. The ergodicity time in this method is found much smaller than that of the multi-canonical method. In particular the time correlation function almost follows an exponential decay whose relaxation time is comparable to the ergodicity time at low temperatures. It suggests that the system relaxes very rapidly through the exchange process even in the low temperature phase.
I. INTRODUCTION

The low temperature phase of spin glasses (SG) and other complex systems generally have numerous local minima which are separated to each other by energy barriers. In studying such systems, we have to take account of each configuration for these local minima and the fluctuation around it. The characteristic time in which the system escapes from a local minimum, however, increases rapidly as temperature decreases. This situation causes “hardly-relaxing” problem in using conventional Monte Carlo (MC) simulations based on a local updating.

So far, various improvements have been (and is being) made on MC algorithms to overcome this problem. There are mainly two categories in such algorithms, i.e., the cluster algorithm and the extended ensemble method. The Swendsen-Wang [1] and the Wolff [2] algorithms are in the first category in which non-local updating is employed to let the system easily change over energy barriers. The second one is based on the idea that an extension of the canonical ensemble can be introduced so as to perform MC sampling efficiently at low temperatures. The multi-canonical method [3, 4] and the simulated tempering [5] are two such algorithms. Some applications of the SG system have been attempted for the multi-canonical method [3, 6, 7] and for the simulated tempering [8, 9, 10]. In this work we propose another efficient algorithm belonging to the second category. It can be regarded as an parallelized version of the simulated tempering, although the extended ensemble used is very different.

Our method treats a compound system which consists of many replicas of the system concerned. The temperature attributed to each replica is distributed in a range including both high and low temperature phases. To this system we perform a standard MC simulation in the following way: (1) Each replica is simulated simultaneously and independently as canonical ensemble for a few MC steps (MCS). (2) Exchange of the configurations for a pair of replicas is tried by referring to the energy cost of the whole system. In effect, state traverses over the replicas, due to the exchange process, being cooled down and warmed up, and thus replica at low temperatures can escape from a local minimum very easily. Since the present method never violates the detailed balance in the canonical sense, each replica is guaranteed to be equilibrated to the canonical distribution with its own temperature. An advantage of our method to the simulated tempering is that we need not to estimate weighting factor (or free energy) to equidistribute the probability of visiting temperatures.

We then apply the exchange MC method to a three-dimensional $\pm J$ Ising spin glass (SG) model and show that it works well even in the SG phase. To characterize the time scale of relaxation in our method, we observe the ergodicity time, which is defined as the average MCS per one travel over the whole temperature range, and a modified autocorrelation function. It is found that the autocorrelation function follows the exponential decay and the relaxation time at low temperatures takes a moderate value which is comparable with the ergodicity time. The order parameter distribution $P(q)$ is an even function in the whole temperature range, which provide us an evidence for reaching the equilibrium.

This paper is organized as follows: In Sec. 2 we describe the exchange MC algorithm in detail. In Sec. 3 we show how to determine the temperature mesh in order that the exchange process occurs properly. In Sec. 4 we report the relaxational behavior of our method by applying it to the SG model. The last section is devoted to discussion and summary.
II. EXCHANGE MC METHOD

In our method we treat a compound system which consists of non-interacting $M$ replicas of the system concerned. The $m$-th replica, described by a common Hamiltonian $H(X)$, is associated with inverse temperature $\beta_m$, i.e., each replica is in contact with its own heat bath having different temperature (for convenience we take $\beta_m < \beta_{m+1}$). A state of this extended ensemble is specified by $\{X\} = \{X_1, X_2, \ldots, X_M\}$, and the partition function is given as

$$Z = \text{Tr}_{\{X\}} \exp(-\sum_{m=1}^{M} \beta_m H(X_m)) = \prod_{m=1}^{M} Z(\beta_m), \quad (2.1)$$

where $Z(\beta)$ is the one for the original system. For a set of temperature $\{\beta\}$ given, the probability distribution of finding $\{X\}$ becomes

$$P(\{X, \beta\}) = \prod_{m} P_{\text{eq}}(X_m, \beta_m), \quad (2.2)$$

where

$$P_{\text{eq}}(X, \beta) = Z^{-1}(\beta) \exp(-\beta H(X)) \quad (2.3)$$

In constructing a Markov process for exchange MC we introduce a transition matrix $W(X, \beta_m|X', \beta_n)$ which is a probability of exchanging configurations of the $n$-th and $m$-th replicas. In order that the system remains at equilibrium, it is sufficient to impose the detailed balance condition on the transition matrix:

$$P(\cdots; X, \beta_m; \cdots; X', \beta_n; \cdots)W(X, \beta_m|X', \beta_n) = P(\cdots; X', \beta_m; \cdots; X, \beta_n; \cdots)W(X', \beta_m|X, \beta_n). \quad (2.4)$$

From eq.(2.3) we obtain

$$\frac{W(X, \beta_m|X', \beta_n)}{W(X', \beta_m|X, \beta_n)} = \exp(-\Delta), \quad (2.5)$$

where

$$\Delta = (\beta_n - \beta_m)(H(X) - H(X')). \quad (2.6)$$

Therefore the replica-exchange part of transition probability can be expressed as

$$W(X, \beta_m|X', \beta_n) = \begin{cases} 1, & \text{for } \Delta < 0 \\ \exp(-\Delta), & \text{for } \Delta > 0 \end{cases} \quad (2.7)$$

if one adopts the Metropolis method.

For the actual MC procedure, the following two steps are performed alternately:

1. Each replica is simulated *simultaneously* and *independently* as canonical ensemble for a few MCS by using a standard MC method.
2. Exchange of two configurations $X_m$ and $X_{m+1}$, is tried and accepted with the probability $W(X_m, \beta_m | X_{m+1}, \beta_{m+1})$.

Here we restrict the replica-exchange to the case $n = m + 1$ because the acceptance ratio of the exchange trial decreases exponentially with the difference $\beta_m - \beta_n$.

The canonical expectation value of a physical quantity $A$ is measured in usual way as follows:

$$
\langle A \rangle_{\beta_m} = \frac{1}{N_{\text{mcs}}} \sum_{t=1}^{N_{\text{mcs}}} A(X_m(t)).
$$

(2.8)

Another expression can be obtained when the exchange procedure mentioned above is regarded as for temperature, i.e., temperatures, instead of configurations, of a pair of replicas are to be exchanged. Then the above quantity is expressed as

$$
\langle A \rangle_{\beta} = \frac{1}{N_{\text{mcs}}} \sum_{t=1}^{N_{\text{mcs}}} \sum_{m=1}^{M} A(\tilde{X}_m(t)) \delta_{\beta, \beta_m(t)},
$$

(2.9)

where we introduce the time-dependent inverse temperature $\beta_m(t)$ and the configuration $\tilde{X}_m$ in this temperature-exchange scheme. Note that both schemes are completely equivalent to one another. One can choose either of the two schemes in actual implementation of the present method.

### III. DETERMINATION OF TEMPERATURE

The parameters we have to determine are only the set of (inverse) temperatures $\{\beta_m\}$. The highest temperature should be set in the high temperature phase where relaxation (correlation) time is expected to be very short and there exists only one minimum in the free energy space, otherwise the system would not completely forget where it was trapped before even if it visits to the highest temperature. On the other hand, the lowest temperature is somewhere in the low temperature phase whose properties we are interested in. In this sense the temperature range is considered to be given. Then we estimate the number of temperatures required in the range as follows. In order that each replica wanders over the whole temperature region the acceptance probabilities of the exchange process for every pair of replicas at different temperatures have to be of order of one and nearly constant. The logarithm of the probability $e^{-\Delta}$ of exchanging $\beta_n$ and $\beta_{n+1} = \beta_n + \delta$ is to order $\delta^2$

$$
\Delta = \delta(\mathcal{H}(X_{n+1}) - \mathcal{H}(X_n)) \sim \delta^2 \frac{d}{d\beta} E,
$$

(3.1)

where the instantaneous value of the energy $\mathcal{H}$ is approximated by the thermal expectation value $E$. Since the energy $E$ is an extensive variable, $\delta$ should be of order of $\frac{1}{\sqrt{N}}$ to satisfy the condition that $\Delta \sim O(1)$. In other words, the number of temperatures in the range we have to simulate is of the order of $\sqrt{N}$. In the case where a second-order phase transition exists, the number is modified as $\sqrt{N^{1+\alpha \nu}}$, where $\alpha$ and $\nu$ are the exponent of the specific heat and the correlation length. This means that we need more replicas if the specific heat
diverges at the transition temperature. In spin glass and other glassy systems, the specific heat has usually no singularity at the transition temperature.

A set of temperatures \( \{ \beta_m \} \) can be obtained by the iteration procedure which is first introduced to the simulated tempering. [10] For given \( \{ \beta_m \} \), the acceptance ratios \( \{ p_m \} \) are evaluated by simulating an appropriate MCS. Then a new set \( \{ \beta'_m \} \) is constructed by using the old set \( \{ \beta_m \} \):

\[
\beta'_1 = \beta_1, \\
\beta'_m = \beta'_{m-1} + (\beta_m - \beta_{m-1}) \frac{p_m}{c}, \quad (m = 1, 2, \cdots, M), \\
c = \frac{1}{M-1} \sum_{m=1}^{M} p_m.
\]

(3.2)

In each iteration the MCS required to estimate \( p_m \) are found not so long. We will show a concrete examination in the next section.

In closing this section, we propose some necessary conditions to check the efficiency and the equilibration.

(i) The exchange happens with a non-negligible probability for all adjacent pairs of replicas.

(ii) Each replica moves around the whole temperature range in suitable MCS.

(iii) In moving the temperature space the system forgets where it was trapped.

We can obtain the temperatures \( \{ \beta_m \} \) satisfying the condition (i) by the iteration procedure mentioned above. The ergodicity time defined in the following section gives a criterion for the condition (ii). Unfortunately, even if the condition (i) and (ii) are satisfied, it is possible that the condition (iii) is violated. One trivial possibility for this is that the largest temperature is not high enough. If some of these conditions are broken down, we may not be able to improve the present method due to the absence of any other controlling parameters.

**IV. THE MODEL AND SIMULATIONS**

We consider the three dimensional \( \pm J \) Ising SG model on the simple cubic lattice. The real-replica Hamiltonian is defined as

\[
\mathcal{H}(\{ \sigma, \tau \}) = -\sum_{ij} J_{ij}(\sigma_i \sigma_j + \tau_i \tau_j),
\]

(4.1)

where \( \{ \sigma \} \) and \( \{ \tau \} \) take the values \( \pm 1 \). The interactions \( \{ J_{ij} \} \) are quenched random variables taking \( \pm 1 \) with equal probability. To be more accurate, \( \{ J_{ij} \} \) are distributed so that the number of the ferromagnetic bonds, \( J_{ij} = +1 \), is exactly a half of the total bonds. Then the Edwards-Anderson SG order parameter [12] is computed as the overlap between the two copies,

\[
q = \frac{1}{N} [\sum_i \langle \sigma_i \tau_i \rangle],
\]

(4.2)

where \( \langle \cdots \rangle \) and \( [\cdots] \) denote thermal average for the Hamiltonian (4.1) and random average, respectively. We have simulated on lattices of linear size \( L = 6, 8, 12, \) and 16 with periodic
boundary condition. For local updating we have adopted the conventional sublattice-flipping with heat bath method. For the exchange process the replica pairs \((\beta_m, \beta_{m+1})\) are divided into two subgroups, \(i.e.,\) odd-\(m\) and even-\(m\) groups. In actual updating the exchange trial is performed for one of these subgroups after each local updating. It means that our one MC step consists of one local updating MC step and a half exchange-trial per replica-pair.

A. temperature setting and the ergodicity time

The actual local updating have been performed with the multi-spin coding technique \([15–17]\) which simulates 32 different physical systems at once. This fixes the total number of temperatures to be \(M = 32\) in all lattice sizes. To determine the temperatures \(\{\beta_m\}\) a typical sample of size \(L = 12\) has been used. The initial condition we used is

\[
\beta_m = \beta_1 + (\beta_M - \beta_1) \frac{m - 1}{M - 1},
\]

with \(\beta_1 = 0.4\) and \(\beta_M = 2\). We then performed 400 MC steps of simulation for each iteration of eq.(3.2) to evaluate the acceptance probabilities \(\{p_m\}\), and found that the temperatures converge after several iterations. Finally we smoothed \(\{\beta_m\}\) by spline interpolation and obtained the temperature range \((0.86 \leq T/J \leq 2.39)\) which includes both the high and low temperature phases. These temperatures so obtained was used for all samples and sizes. In fig. 1 we show the acceptance ratios for various sizes. These values are found to be of order of one and do not depend on bond realization so sensitively.

In order to confirm that the obtained temperature set satisfies the condition (ii) mentioned in the previous section, we investigated the ergodicity time \(\tau_E\) which is defined as the average MC step for a specific replica to move from the lowest to the highest temperature. The dependence of our observed ergodicity time on the lattice size is shown in Fig. 2, with that of the multi-canonical method by Berg et al. \([9]\) If the total number of temperature points we simulate is fixed as in the present analysis, the acceptance ratio, \(e^{-\Delta}\), of exchange process between replicas reduces exponentially as a function of system size \(L\) and the \(\tau_E\) grows rapidly with increasing \(L\). As shown in Fig. 3, however, it is clear that our ergodicity time is much shorter than the multi-canonical ergodicity time in a lattice size suitable for our actual simulation.

B. relaxation

Because each replica wanders over the temperature space, the (equilibrium) time correlation function can not be argued in the ordinary sense. Here to investigate relaxation dynamics of the present method, we study an autocorrelation function in the temperature-exchange scheme,

\[
q(t, \beta_m) = \frac{1}{N} \sum_i \langle \tilde{\sigma}_i(0) \tilde{\sigma}_i(t) \rangle^{(m)}_{\text{path}},
\]

where \(\langle \cdots \rangle^{(m)}_{\text{path}}\) means an average along trajectories in temperature space evolving from the initial state \(\{\tilde{\sigma}_i(0)\}\) with temperature \(\beta_m\). The function \(q(t, \beta)\) is expected to include
the slowest relaxation mode of the present method. In Fig. 3 we show \( q(t, \beta) \) at various temperatures for \( L = 12 \). The data are averaged over 10 samples. As indicated by the straight lines, the function \( q(t, \beta) \) follows nearly exponential decay for large time \( t \). We evaluate the largest relaxation time from least square fit to a single exponential

\[
q(t, \beta) \sim q_0 \exp(-t/\tau) \quad t \gg 1.
\]

As shown in Fig. 3(a) the relaxation time \( \tau \) has a crossover temperature \( T_{cr} \sim 1.5 \), above which \( \tau \) behaves like the relaxation time obtained by a conventional local spin flip dynamics. Below \( T_{cr} \), on the other hand, \( \tau \) seems to almost become saturated and to be comparable with \( \tau_E \), and the relaxation amplitude \( q_0 \) grows rapidly (see Fig. 3(b)). It suggests that the largest relaxation mode of the exchange dynamics dominates the relaxation at low temperatures. In any case we can see that the system relaxes within a reasonable MCS and thus the condition (iii) is satisfied.

C. order parameter distribution

In this subsection we show the obtained distribution function \( P_J(q) \) of overlap function for a sample, which is defined by

\[
P_J(q) = \langle \delta(q - \frac{1}{N} \sum_i \sigma_i \tau_i) \rangle.
\]

The distribution function \( P_J(q) \) is important to SG study because all physical quantities of our interest can be obtained from this function. The distribution is expected to be an even function from the invariance of the Hamiltonian under the transformation \( \sigma_i \rightarrow -\sigma_i \).

We demonstrate in Fig. 4(a) that for \( L = 16 \) the distribution \( P_J(q) \) is symmetric even below the SG transition temperature. Note that only \( 3 \times 10^5 \) MCS, which is about 10 times larger than \( \tau_E \) are used to obtain Fig. 4. The distribution of the same sample obtained by the conventional MC method is also shown in Fig. 4 with the broken line. Obviously, it is far from symmetric and even the peak positions differ with each other. It strongly suggests that the system does not reach the thermal equilibrium yet by the conventional MC method.

V. DISCUSSION AND SUMMARY

One may consider that, for random systems, \( \{\beta_m\} \) should be determined for each sample separately. It is, however, not the case because the exchange probability \( \{p_m\} \) is insensitive to bond realization as long as \( \beta_{m+1} - \beta_m \) is sufficiently small. It is contrasted to the case of the multi-canonical method, where the energy density should be estimated for each sample because it strongly depends on the structure of local minima in a sample.

An advantage of the present method to the simulated tempering is that we need not estimate weighting factor (or free energy) to equidistribute the probability of visiting temperatures. This is because the phase space of the present method is a direct product of the original one in contrast with that of the simulated tempering, which is given by a direct sum. Since the weight factor in the simulated tempering is an extensive parameter, the
probability is very sensitive to it, and temperature traversing would be easily broken if the estimation is not so good.

A more important merit of the present method appears in evaluating $P(q)$ of the SG system, where two copies of a system are independently simulated. We have $M$ replicas for each copy, so that we can take $M(M - 1)/2$ un-correlated samplings of overlap in one simulation as compared with $M$ sampling is by $M$ independent runs of the simulated tempering.

As shown in the previous section, the model to be simulated is not supposed to have any specific aspect, so that we can, in principle, apply the present method to various models without modification. Unfortunately the present method seems not to work well in the systems exhibiting the first order phase transition. Since the energy has a finite gap at transition point, the exchange between replicas below and above transition temperature hardly occurs in a large lattice, meaning that the condition (i) in section 3 is not satisfied.

In summary we have proposed a new MC method for simulating hardly relaxing systems. We have applied this method to the three dimensional $\pm J$ Ising spin glass system, and found that the system really traverses over wide temperature space and the largest relaxation time in this dynamics is given by the ergodicity time, which is much smaller than the conventional one. As a result, the order parameter distribution $P(q)$ can be obtained below $T_c$ up to $L = 16$ within much shorter than the conventional MCS. Numerical results of physical quantities and discussion on the SG phase transitions will be discussed in detail elsewhere. [20]

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FIGURES

FIG. 1. Temperature dependence of the acceptance probability of exchange process. The sample-to-sample errors are smaller than 1/10 of the size of symbols.

FIG. 2. Ergodicity time $\tau_E$ as a function of the system size $V = L^3$. The circles correspond to our data, and squares to multi-canonical method by Berg et al. The unit of relaxation time is MC step per spins.

FIG. 3. The modified autocorrelation function $q(t, \beta)$ for $1/\beta = T = 0.862, 1.088, 1.256, 1.399, \text{and} 1.609$ (top to down).

FIG. 4. Temperature dependence of the relaxation time $\tau$ (a) and its amplitude $q_0$ (b) of the modified autocorrelation function (Fig. 3).

FIG. 5. Distribution function $P_J(q)$ of the overlap for a sample with $L = 16$ at $T = 0.924$. The solid and dashed line represent $P(q)$ by using the exchange MC and by a conventional MC method, respectively. The same MCS are used in two methods.
Acceptance ratio

\[ T \]

\[ L=6 \]
\[ L=8 \]
\[ L=12 \]
\[ L=16 \]
This work
Berg et al
$q(t, \beta)$
$P(q)$

- exchange MC method
- conventional MC method