Approaching equilibrium and the distribution of clusters

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We investigate the approach to stable and metastable equilibrium in both nearest-neighbor and long-range Ising models using a cluster representation. The distribution of nucleation times is determined using the Metropolis algorithm and the corresponding ϕ^4 model using Langevin dynamics. We find that the nucleation rate is suppressed at early times even after global variables such as the magnetization and energy have apparently reached their time independent values. The mean number of clusters whose size is comparable to the size of the nucleating droplet becomes time independent at about the same time that the nucleation rate reaches its constant value. We also find subtle structural differences between the nucleating droplets formed before and after apparent metastable equilibrium has been established.

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I. INTRODUCTION

Understanding nucleation is important in fields as diverse as materials science, biological physics, and meteorology [1–9]. Fundamental progress was made when Gibbs assumed that the droplet can be considered to be a fluctuation about metastable equilibrium, and hence the probability of a droplet is independent of time [10]. Langer [11] has shown that the probability of a droplet can be related to the analytic continuation of the stable state free energy in the limit that the metastable state lifetime approaches infinity. Hence the assumption by Gibbs is valid in this limit. It has also been shown that the Gibbs assumption is applicable in systems for which the interaction range $R \rightarrow \infty$ [12,13].

For metastable states with finite lifetimes equilibrium is never reached because a large enough fluctuation would initiate the transformation to the stable state. However, if the probability of such a fluctuation is sufficiently small, it is possible that systems investigated by simulations and experiments can be well approximated as being in equilibrium. Hence, for metastable lifetimes that are very long, we expect the Gibbs assumption to be a good approximation.

In practice, nucleation is not usually observed when the lifetime of the metastable state is very long. Processes such as alloy formation, decay of the false vacuum, and protein crystallization generally occur during a continuous quench of a control parameter such as the temperature. It is natural to ask if the nucleation process that is observed occurs when the system can be reasonably approximated by treating it as if it were in metastable equilibrium.

It is frequently assumed that metastable equilibrium is a good approximation when the mean value of the order parameter and various global quantities are no longer changing with time. As an example, we consider the nearest-neighbor Ising model on a square lattice and equilibrate the system at temperature $T=4T_c/9$ in a magnetic field h=0.44. The relatively small value of the linear dimension L=200 was chosen in order to avoid nucleation occurring too quickly. At time t=0 the sign of the magnetic field is reversed. In Fig. 1 we plot the evolution of the magnetization m(t) and the energy e(t) per spin using the Metropolis algorithm. The solid lines

are the fits to an exponential function with the relaxation time $\tau_g \approx 1.5$. In the following we will measure the time in terms of Monte Carlo steps per spin.

A major goal of this paper is to address the question: Can the system be treated as being in metastable equilibrium for $t \gtrsim \tau_g$? If so, the nucleation rate would be time independent. We will address this question for both the short-range and long-range Ising models.

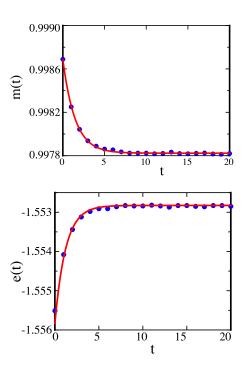


FIG. 1. (Color online) The evolution of the magnetization m(t) and the energy e(t) per spin of the nearest-neighbor Ising model on a square lattice with linear dimension L=200 using the Metropolis algorithm. The system was prepared at temperature T=4 T_c /9 in the external magnetic field h=0.44. At time t=0 the sign of the magnetic field is reversed. The solid lines are fits to an exponential function with relaxation time τ_g =1.5 and 1.2, respectively. (Time is measured in Monte Carlo steps per spin.) The data is averaged over 5000 runs.

If the nucleation rate is independent of time, the probability of a droplet occurring at time t after the change of magnetic field is an exponentially decreasing function of time. To understand this dependence we divide the time into intervals Δt and write the probability that the system nucleates in a time interval Δt as $\lambda \Delta t$, where the nucleation rate λ is a constant. The probability that nucleation occurs in the time interval (N+1) is given by

$$P_N = (1 - \lambda \Delta t)^N \lambda \Delta t. \tag{1}$$

If we assume that $\lambda \Delta t$ is small and write $N=t/\Delta t$, we can write

$$P(t)\Delta t = (1 - \lambda \Delta t)^{t/\Delta t} \lambda \Delta t \to e^{-\lambda t} \lambda \Delta t, \tag{2}$$

where $P(t)\Delta t$ is the probability that the system nucleates at a time between t and $t+\Delta t$ after the change of the magnetic field. In the following we ask if the nucleation rate and the mean values of the order parameter and other thermodynamic quantities become time independent at approximately the same time after a quench or is the approach to metastable equilibrium more complicated?

In Sec. II we determine the probability distribution of the nucleation times and find that the nucleation rate becomes a constant only after a time τ_{nequil} that is much longer than the relaxation time τ_g of m(t) and e(t). In Sec. III we study the microscopic behavior of the system and determine the relaxation time τ_s for n_s , the mean number of clusters of size s, to approach its equilibrium value [14]. Our main result is that τ_s is an increasing function of s, and the time required for n_s to reach its equilibrium value is the same order of magnitude as τ_{nequil} for values of s comparable to the nucleating droplet. That is, the time for the number of clusters that are the size of the droplet to reach its equilibrium value is considerably longer than the time for the mean value of the order parameter to become independent of time within the accuracy that we can determine.

In Secs. IV and V we show that there are subtle differences between the structure of the nucleating droplets that occur before and after the nucleation rate has become time independent. This difference suggests the possibility of finding even greater differences in the nucleating droplets in systems of physical and technological importance. We summarize and discuss our results in Sec. VI. In the Appendix we study the evolution of the clusters after a quench to the critical temperature of the Ising model and again find that that the clusters equilibrate in size order, with the smaller clusters equilibrating first. Hence in principle, an infinite system will never equilibrate. How close to equilibrium a system needs to be and on what spatial scale so that it can be treated by equilibrium methods depends on the physical process of interest.

II. DISTRIBUTION OF NUCLEATION TIMES

We simulate the Ising model on a square lattice with interaction range R with the Hamiltonian

$$H = -J\sum_{\langle i,j\rangle} s_i s_j - h\sum_i s_i, \tag{3}$$

where h is the external field. The notation $\langle i,j \rangle$ in the first sum means that the distance between spins i and j is within the interaction range R. We studied both nearest-neighbor (R=1) and long-range interactions $(R \ge 20)$. The interaction strength J is scaled as J=4/q, where q=2R(R+1) is the number of interaction neighbors per spin. The external field h and the temperature are measured in terms of J in the simulations. All of our simulations are at temperature $T=4T_c/9$, where T_c is the critical temperature. For R=1 the critical temperature is $T_c\approx 2.269$. For $R \ge 20$ the mean-field result $T_c=4$ is a good approximation to the critical temperature [15]. As discussed in Sec. I the system is equilibrated in a magnetic field h. The time t=0 corresponds to the time immediately after the magnetic field is reversed.

The clusters in the Ising model are defined rigorously by a mapping of the Ising critical point onto the percolation transition of a properly chosen percolation model [9,16,17]. Two parallel spins that are within the interaction range R are connected only if there is a bond between them. The bonds are assigned with the probability $p_b = 1 - e^{-2\beta J}$ for R = 1 and $p_b = 1 - e^{-2\beta J(1-\rho)}$ near the spinodal, where ρ is the density of the stable spins, and β is the inverse temperature [18]. Spins that are connected by bonds form a cluster.

Because the intervention method [19] of identifying the nucleating droplet is time consuming (see Sec. IV), we use a simpler criterion in this section to estimate the nucleation time. We monitor the size of the largest cluster (averaged over 20 bond realizations) and estimate the nucleation time as the time when the largest cluster first reaches a threshold size s^* . The threshold size s^* is chosen so that the largest cluster begins to grow rapidly once its size is greater than or equal to s^* . Because s^* is larger than the actual size of the nucleating droplet, the nucleation time that we estimate by this criterion will be 1 to 2 Monte Carlo steps per spin later than the nucleation time determined by the intervention method. Although the distribution function P(t) is shifted to slightly later times, the nucleation rate is found to be insensitive to the choice of the threshold.

Figure 2 shows P(t) for R=1 and h=0.44, where $P(t)\Delta t$ is the probability that nucleation has occurred between time t and $t+\Delta t$. The results for P(t) were averaged over 5000 runs. Note that P(t) is an increasing function of t for early times, reaches a maximum at $t = \tau_{\text{nequil}} \approx 60$, and fits to the expected exponential form for $t \gtrsim \tau_{\text{nequil}}$. The fact that P(t) falls below the expected exponential for $t < \tau_{\rm nequil}$ indicates that the nucleation rate is reduced from its equilibrium value and that system is not in metastable equilibrium. Similar nonequilibrium effects have been studied and observed in Ising-type [20–23] and continuous systems [24]. We conclude that the time for the nucleation rate to become time independent after the change of magnetic field is much longer than the relaxation time $\tau_{o} \simeq 1.5$ of the magnetization and energy. We will refer to nucleation that occurs before the nucleation rate has become time independent as transient nucleation.

The mean size of the droplet was estimated by finding the maximum cluster size that failed to grow (instead of using

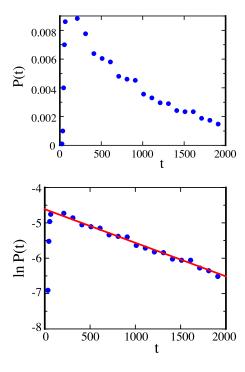


FIG. 2. (Color online) The distribution of nucleation times P(t) averaged over 5000 runs for the same system as in Fig. 1. The threshold size was chosen to be $s^* = 30$. (The mean size of the droplet is ≈ 25 spins.) (a) P(t) begins to decay exponentially at $\tau_{\text{nequil}} \approx 60$. The nucleation rate after equilibrium has been established is determined from the log-linear plot in (b) and is $\lambda \approx 9 \times 10^{-4}$ [see Eq. (2)].

the intervention method). This size represents an upper bound to the size of the nucleating droplet. For h=0.44 this size was found to be approximately 25 spins, which is consistent with our observation that a cluster starts to grow rapidly once its size reaches 20–25 spins.

In order to see if the same qualitative behavior holds near the pseudospinodal [25], we simulated the long-range Ising model with R=20 and h=1.258. In the mean-field limit $R\to\infty$ the spinodal field is at $h_s=1.2704$ [12] (for $T=4T_c/9$). A plot of m(t) for this system is shown in Fig. 3(a) and is seen to have the same qualitative behavior as in Fig. 2 for R=1; the relaxation time $\tau_g\approx 4.5$. In Fig. 3(b) the distribution of nucleation times is shown, and we see that P(t) does not decay exponentially until $t \gtrsim \tau_{\rm nequil} = 40$. According to Ref. [26], $\tau_{\rm nequil}$ should become comparable to τ_g in the limit $R\to\infty$ because the free energy is described only by the magnetization in the mean-field limit. We find that the difference between $\tau_{\rm nequil}$ and τ_g is smaller for R=20 than for R=1, consistent with Ref. [26].

III. RELAXATION OF CLUSTERS TO METASTABLE EQUILIBRIUM

Given that there is a significant time delay between the relaxation of the magnetization and the energy and the equilibration of the system as measured by the nucleation rate, it is interesting to monitor the time dependence of the

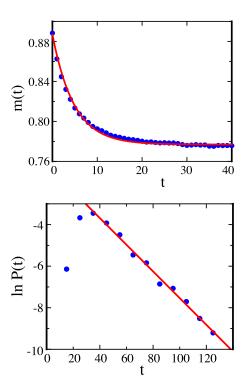


FIG. 3. (Color online) (a) The evolution of m(t) for the long-range Ising model on a square lattice with R=20, h=1.258, and L=500. The solid line is an exponential fit with the relaxation time $\tau_g \approx 4.5$. The data is averaged over 2000 runs. (b) The distribution of nucleation times P(t) for the same system and number of runs. P(t) decays exponentially for $t \approx \tau_{\text{nequil}} \approx 40$. The nucleation rate once equilibrium has been established is $\lambda = 6.4 \times 10^{-2}$. The mean size of the nucleating droplet is determined to be 300 spins using the same method as for R=1.

cluster-size distribution after the reverse of the magnetic field. After the change the system gradually relaxes to metastable equilibrium by forming clusters of spins in the stable direction. How long is required for the number of clusters of size *s* to reach equilibrium? In particular, we are interested in the time required for clusters that are comparable in size to the nucleating droplet.

We first consider R=1 and monitor the number of clusters n_s of size s at time t. To obtain good statistics we chose L=200 and averaged over 5000 runs. Figure 4 shows the evolution of $n_6(t)$, which can be fitted to the exponential form

$$n_s(t) = n_s \,_{\infty} [1 - e^{-t/\tau_s}].$$
 (4)

We find that $\tau_s \approx 8.1$ for s=6. By doing similar fits for a range of s, we find that the time τ_s for the mean number of clusters of size s to become time independent increases linearly with s over the range of s that we can simulate (see Fig. 5). The linear dependence of τ_s on s is consistent with the theoretical prediction in Ref. [27]. The extrapolated value of τ_s corresponding to the mean size of the droplet (≈ 25 spins by direct simulation) is $\tau_{\rm extrap} \approx 34$. That is, it takes a time of $\tau_{\rm extrap} \approx 34$ for the mean number of clusters whose size is the order of the nucleating droplets to become time independent. The time $\tau_{\rm extrap}$ is much longer than the

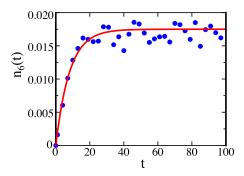


FIG. 4. (Color online) The evolution of the number of clusters of size s=6 averaged over 5000 runs for R=1 and the same conditions as in Fig. 1. The fit is to the exponential form in Eq. (4) with $\tau_s \approx 8.1$ and $n_{s,\infty} = 0.0175$.

relaxation time $\tau_g \approx 1.5$ of the macroscopic quantities m(t) and e(t) and is comparable to the time $\tau_{\text{nequil}} \approx 60$ for the nucleation rate to become independent of time.

Because the number of clusters in the nucleating droplet is relatively small for R=1 except very close to coexistence

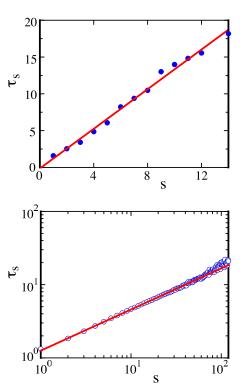


FIG. 5. (Color online) (a) The equilibration time τ_s as a function of the cluster size s for R=1 and h=0.44 and the same conditions as in Fig. 1. The s dependence of τ_s is approximately linear. The extrapolated value of τ_s corresponding to the mean size of the droplet (≈ 25 spins) is $\tau_{\rm extrap} \approx 34$, which is the same order of magnitude as time $\tau_{\rm nequil} \approx 60$ for the nucleation rate to become independent of time. (b) Log-log plot of the equilibration time τ_s versus s for R=20 and h=1.258 and the same conditions as in Fig. 3(b). We find that $\tau_s \sim s^x$ with the exponent $x \approx 0.56$. The extrapolated value of τ_s corresponding to the mean size of the droplet (≈ 300 spins) is $\tau_{\rm extrap} \approx 30$, which is comparable to the time $\tau_{\rm nequil} \approx 40$ for the nucleation rate to become independent of time.

(small h), we also consider a long-range Ising model with R=20 and h=1.258 (as in Fig. 3). The relaxation time τ_s of the clusters near the pseudospinodal fits to a power law $\tau_s \sim s^x$ with $x\approx 0.56$ [see Fig. 5(b)]. If we extrapolate τ_s to s=300, the approximate size of the nucleating droplet, we find that the equilibration time for clusters of the size of the droplet is $\tau_{\text{extrap}} \approx 30$, which is comparable to the time $\tau_{\text{nequil}} \approx 40$ for the nucleation rate to become independent of time.

To determine if our results are affected by finite size effects, we compared the equilibration time of the clusters for lattices with linear dimension L=2000 and L=5000. The equilibration times of the clusters were found to be unaffected.

IV. STRUCTURE OF THE NUCLEATING DROPLET

Because nucleation can occur both before and after the nucleation rate has become independent of time, we ask if there are any structural differences between the nucleating droplets formed in these two cases. To answer this question, we determine the nature of the nucleating droplets for the one-dimensional (1D) Ising model where we can make R (and hence the size of the nucleating droplets) large enough so that the structure of the nucleating droplets is well defined. In the following we take $R=2^{12}=4096$, h=1.265, and $L=2^{18}$. The relaxation time for m(t) is $\tau_g \approx 40$, and the time for the distribution of nucleation times to reach equilibrium is $\tau_{\text{nequil}} \approx 90$.

We use the intervention method to identify nucleation [19]. To implement this method, we choose a time at which a droplet might exist and make many copies of the system. Each copy is restarted using a different random number seed. The idea is to determine if the largest cluster in each of the copies grows in approximately the same place at about the same time. If the percentage of copies that grow is greater than 50%, the nucleating droplet is already in the growth phase; if it is less than 50%, the time chosen is earlier than nucleation. We used a total of 20 trials to make this determination

Our procedure is to observe the system for a time $t_{\rm obs}$ after the intervention and determine if the size of the largest cluster exceeds the threshold size s^* at approximately the same location. To ensure that the largest cluster at $t_{\rm obs}$ is the same cluster as the original one, we require that the center of mass of the largest cluster be within a distance r^* of the largest cluster in the original configuration. If these conditions are satisfied, the droplet is said to grow. We choose $t_{\rm obs}=6$, $r^*=2R$, and $s^*=2000$. (In comparison, the size of the droplet for the particular run that we will discuss is ≈ 1080 spins.)

There is some ambiguity in our identification of the nucleation time because the saddle point parameter is large but finite [9]. This ambiguity manifests itself in the somewhat arbitrary choices of the parameters $t_{\rm obs}$, r^* , and s^* . We tried different values for $t_{\rm obs}$, r^* , and s^* and found that our results depend more strongly on the value of the parameter r^* than on the values of $t_{\rm obs}$ and s^* . If we take $r^* = R/2$, the nucleating droplets almost always occur one to two Monte Carlo steps per spin later than for $r^* = 2R$. The reason is that the linear size of the droplet is typically 6-8R, and its center of

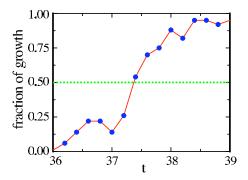


FIG. 6. (Color online) The fraction of copies for which the largest cluster grows for a particular run for a 1D Ising model with $R=2^{12}$, h=1.265, and $L=2^{18}$. The time for 50% growth is ≈ 37.4 . The largest cluster at this time corresponds to the nucleating droplet and has ≈ 1080 spins. For this intervention 100 copies were considered; 20 copies were considered for all other runs.

mass might shift more than R/2 during the time $t_{\rm obs}$. If such a shift occurs, a cluster that would be said to grow for $r^* = 2R$ would not be counted as such because it did not satisfy the center of mass criterion. This shift causes an overestimate of the time of the nucleating droplet. A reasonable choice of r^* is 20%-40% of the linear size of the nucleating droplet. The choice of parameters is particularly important here because the rate of growth of the transient nucleating droplets is slower than the growth rate of droplets formed after the nucleation rate has become independent of time. Hence, we have to identify the nucleating droplet as carefully as possible.

Because nucleation studies are computationally intensive, we used an algorithm for simulating Ising models with a uniform long-range interaction [28]. The algorithm uses a hierarchical data structure to store the magnetization at many length scales, and can find the energy cost of flipping a spin in time $O[(\ln R)^d]$, rather than the usual time $O(R^d)$, where d is the spatial dimension.

Figure 6 shows the fraction of copies for which the largest cluster grows as a function of the intervention time. For this particular run the nucleating droplet is found to occur at $t \approx 37.4$.

We simulated 100 systems in which nucleation occurred before global quantities such as m(t) became independent of time, $t < \tau_g \approx 40$, and 100 systems for which nucleation occurred after the nucleation rate became time independent $(t > \tau_{\rm nequil} \approx 90)$. We found that the mean size of the nucleating droplet for $t < \tau_g$ is ≈ 1200 with a standard deviation of $\sigma \approx 150$ in comparison to the mean size of the nucleating droplet for $t > \tau_{\rm nequil}$ of ≈ 1270 and $\sigma \approx 200$. That is, the nucleating droplets formed before the nucleation rate has become independent of time are somewhat smaller.

We introduce the *cluster profile* ρ_{cl} to characterize the shape of the largest cluster at the time of nucleation. For a particular bond realization a spin that is in the stable direction might or might not be a part of the largest cluster due to the probabilistic nature of the bonds. For this reason bond averaging is implemented by placing 100 independent sets of bonds between spins with probability $p_b = 1 - e^{-2\beta J(1-\rho)}$ in the

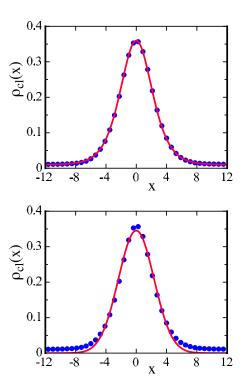


FIG. 7. (Color online) Comparison of the mean cluster profile (•) in the 1D Ising model after the nucleation rate has become independent of time with (a) the form in Eq. (5) and (b) a Gaussian. Note that Eq. (5) gives a better fit than the Gaussian, which underestimates the peak at x=0 and the wings. The x axis is measured in units of R.

stable direction. The clusters are identified for each set of bonds, and the probability p_i that spin i is in the largest cluster is determined. The values of p_i for the spins in a particular bin are then averaged using a bin width equal to R/4. This mean value of p_i is associated with $\rho_{\rm cl}$. Note that the spins that point in the unstable direction are omitted in this procedure. The mean cluster profile is found by translating the peak position of each droplet to the origin.

Figure 7(a) shows the mean cluster profile formed after the nucleation rate has become independent of time $(t > \tau_{\text{nequil}} \approx 90)$. The position x is measured in units of R. For comparison we fit ρ_{cl} to the form [12]

$$\rho(x) = A \operatorname{sech}^{2}(x/w) + \rho_{0}, \tag{5}$$

with A_{cl} =0.36, w_{cl} =2.95, and ρ_0 =0 by construction. In Fig. 7(b) we show a comparison of ρ_{cl} to the Gaussian form $A_g \exp[-(x/w_g)^2]$ with A_g =0.35 and w_g =3.31. Note that Eq. (5) gives a better fit than a Gaussian, which underestimates the peak at x=0 and the wings. Although Unger and Klein [12] derived Eq. (5) for the magnetization saddlepoint profile, we see that this form also provides a good description of the cluster profile.

A comparison of the cluster profiles formed before and after the nucleation rate has become independent of time is shown in Fig. 8. Although both profiles are consistent with the form in Eq. (5), the transient nucleating droplets are more compact, in agreement with the predictions in Ref. [26].

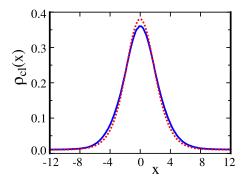


FIG. 8. (Color online) The cluster profiles of the nucleating droplets formed before (dashed line) and after (solid line) the nucleation rate has become independent of time. Both profiles are consistent with the form given in Eq. (5), but the transient nucleating droplets are slightly more compact. The fitting parameters are A = 0.38 and w = 2.67 for the transient droplets and A = 0.35 and w = 2.95 for the droplets formed after the nucleation rate has become independent of time.

We also directly coarse grained the spins at the time of nucleation to obtain the density profile of the coarse-grained magnetization $\rho_m(x)$ [see Fig. 9(a)]. The agreement between the simulation and analytical results [29] are impressive, especially considering that the analytical form is valid only in the limit $R \rightarrow \infty$. The same qualitative differences between the nucleating droplets that occur before and after the nucleation rate has become independent of time is found [see Fig. 9(b)], although the magnetization density profile is much noisier than that based on the cluster analysis.

V. LANGEVIN SIMULATIONS

It is interesting to compare the results for the Ising model and the Langevin dynamics of the ϕ^4 model. One advantage of studying the Langevin dynamics of the ϕ^4 theory is that it enables the efficient simulation of systems with a very large interaction range R. If all lengths are scaled by a large value of R, the effective magnitude of the noise decreases, making faster simulations possible.

The coarse-grained Hamiltonian analogous to the 1D ferromagnetic Ising model with long-range interactions in an external field h can be expressed as

$$H[\phi(x)] = \int \left[-\frac{1}{2} \left(R \frac{d\phi(x)}{dx} \right)^2 + \epsilon \phi^2(x) + u \phi^4(x) - h \phi(x) \right] d\phi(x), \tag{6}$$

where $\phi(x)$ is the coarse-grained magnetization. A dynamics consistent with this Hamiltonian is given by

$$\frac{\partial \phi}{\partial t} = -M \frac{\delta H}{\delta \phi} + \eta = -M \left[-R^2 \frac{\partial^2 \phi}{\partial x^2} + 2\varepsilon \phi + 4u \phi^3 - h \right] + \eta, \tag{7}$$

where *M* is the mobility and $\eta(x,t)$ represents zero-mean Gaussian noise with $\langle \eta(x,t) \eta(x',t') \rangle = 2kTM \delta(x-x') \delta(t-t')$.

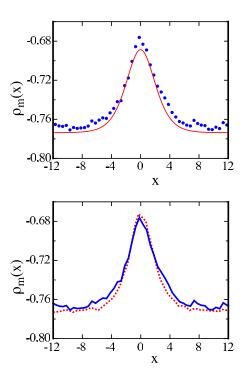


FIG. 9. (Color online) (a) The magnetization density profile of the nucleating droplets formed after the nucleation rate has become independent of time. The solid line is the analytical solution [29], which has the form in Eq. (5) with the calculated values A = 0.085, w = 2.65, and $\rho_0 = -0.774$. (b) Comparison of the density profile of nucleating droplets formed before (dashed line) and after (solid line) the nucleation rate has become independent of time by coarse graining the magnetization. The same qualitative differences between the nucleating droplets that occur before and after the nucleation rate has become independent of time are observed as in Fig. 8, although the magnetization density profile is much noisier than the cluster density profile.

For nucleation near the spinodal the potential $V = \varepsilon \phi^2 + u \phi^4 - h \phi$ has a metastable well only for $\varepsilon < 0$. The magnitude of ϕ and h at the spinodal are given by $h_s = \sqrt{(8|\varepsilon|^3/27u)}$ and $\phi_s = \sqrt{(|\varepsilon|/6u)}$, and are found by setting

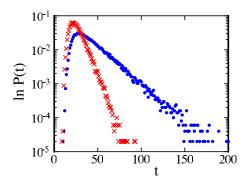


FIG. 10. (Color online) Log-linear plot of the distribution P(t) of nucleation times for the one-dimensional Langevin equation with $R=2000~(\times)$ and $R=2500~(\bullet)$ averaged over 50 000 runs. The distribution is not exponential for early times, indicating that the system is not in metastable equilibrium. Note that the nucleation rate is a rapidly decreasing function of R.

V'=V''=0. The distance from the spinodal is characterized by the parameter $\Delta h=|h_s-h|$. For $\Delta h/h_s\ll 1$, the bottom of the metastable well ϕ_{\min} is near ϕ_s , specifically $\phi_{\min}=-\phi_s(1+\sqrt{2\Delta h/3h_s})$.

The stationary solutions of the dynamics are found by setting $\delta H/\delta \phi = 0$. Besides the two uniform solutions corresponding to the minima in V, there is a single nonuniform solution, which approximates the nucleating droplet profile when the nucleation barrier is large. When $\Delta h/h_s \ll 1$, the profile of the droplet is described by Eq. (5) with $A = \sqrt{h_s/6\Delta h}/\phi_s$, $w = (8h_s\Delta h\phi_s^2/3)^{-1/4}$, and $\rho_0 = \phi_{\min}$ [12].

The dynamics (7) is numerically integrated using the scheme [30]

$$\phi(t + \Delta t) = \phi(t) - \Delta t M \left[-R^2 \frac{\partial^2 \phi}{\partial x^2} + 2\varepsilon \phi + 4u \phi^3 - h \right] + \sqrt{\frac{\Delta t}{\Delta x}} \eta, \tag{8}$$

where $d^2\phi/dx^2$ is replaced by its central difference approximation. Numerical stability requires that $\Delta t < (\Delta x/R)^2$, but it is often desirable to choose Δt even smaller for accuracy.

As for the Ising simulations, we first prepare an equilibrated system with ϕ in the stable well corresponding to the direction of the external field h. At t=0 the external field is reversed so that the system is no longer in equilibrium. We choose M=1, T=1, $\varepsilon=-1$, u=1, and $\Delta h=0.005$. The scaled length of the system is chosen to be L/R=300. We choose R to be large so that, on length scales of R, ϕ fluctuates near its equilibrium value $\phi_{\min} \approx -0.44$.

After nucleation occurs ϕ will rapidly grow toward the stable well. To determine the distribution of nucleation times, we assume that when the value of the field ϕ in any bin reaches 0, nucleation has occurred. This relatively crude criterion is sufficient for determining the distribution of nucleation times if we assume that the time difference between the nucleation event and its later detection takes a consistent value between runs.

Figure 10 compares the distribution of 50 000 nucleation times for systems with R=2000 and R=2500 with $\Delta x/R$ =1 and Δt =0.1. The distribution shows the same qualitative behavior as found in the Metropolis simulations of the Ising model (see Fig. 2). For example, the distribution of nucleation times is not exponential for early times after the quench. As expected, the nucleation rate decreases as R increases. Smaller values of Δx and Δt give similar results for the distribution.

To find the droplet profiles, we need to identify the time of nucleation more precisely. The intervention criterion, which was applied in Sec. IV, is one possible method. In the Langevin context we can employ a simpler criterion: nucleation is considered to have occurred if ϕ decays to the saddle-point profile [given by Eq. (5) for $\Delta h/h_s \ll 1$] when ϕ is evolved using noiseless dynamics [26,31]. For fixed Δh these two criteria agree in the $R \to \infty$ limit, but can give different results for finite R [32].

In Fig. 11 we plot the average of 1000 density profiles of the nucleating droplets formed after the nucleation rate has

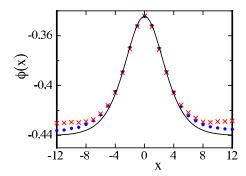


FIG. 11. (Color online) Comparison of the density profile $\phi(x)$ of the nucleating droplets found by numerically solving the Langevin equation after the nucleation rate has become independent of time for R=2000 (×) and R=4000 (\bullet) to the theoretical prediction (solid line) from Eq. (5) using the calculated values A=0.096, w=3.58, and ρ_0 =-0.44. The numerical solutions are averaged over 1000 profiles. The results suggest that as R increases, the observed nucleation profiles approach the mean-field theory prediction.

become independent of time for R=2000 and R=4000. Note that there are noticeable deviations of the averaged profiles from the theoretical prediction in Eq. (5), but the deviation is less for R=4000. The deviation is due to the fact that the bottom of the free energy well in the metastable state is skewed; a similar deviation was also observed in the Ising model. We also note that the individual nucleating droplets look much different from their average. It is expected that as R increases, the profiles of the individual nucleating droplets will converge to the form given by Eq. (5).

In Fig. 12 we compare the average of 1000 density profiles of nucleating droplets before and after the nucleation rate has become independent of time. As for the Ising model, there are subtle differences consistent with the predictions of Ref. [26]. The transient droplets have slightly lower background magnetization and compensate by being denser and more compact.

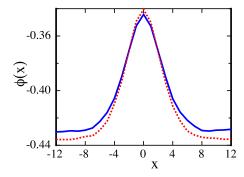


FIG. 12. (Color online) The density profile of the nucleating droplets found from numerical solutions of the Langevin equation formed before (dotted line) and after (solid line) the nucleation rate has become independent of time. Nucleation events occurring before t=15 are transient, and events occurring for $t \ge 30$ can be treated assuming that the system is in metastable equilibrium. Both plots are the result of 1000 averaged profiles with an interaction range R=2000.

VI. SUMMARY

Although the time independence of the mean values of macroscopic quantities such as the magnetization and the energy is often used as an indicator of metastable equilibrium, we find that the observed relaxation time of the clusters is much longer for sizes comparable to the size of the nucleating droplet. This longer relaxation time is consistent with the measured nonconstant nucleation rate even when global quantities such as the magnetization and energy appear to be stationary. By identifying the nucleating droplets, in the onedimensional long-range Ising model and the Langevin equation, we find structural differences between the nucleating droplets, which occur before and after the nucleation rate has become independent of time. Our results suggest that using global quantities as indicators for metastable equilibrium may not be appropriate in general, and distinguishing between equilibrium and transient nucleation is important in studying the structure of nucleating droplets. Further studies of transient nucleation in continuous models of more realistic systems would be of interesting and practical importance.

Finally, we note a subtle implication of our results. For a system to be truly in equilibrium would require that the mean number of clusters of all sizes be independent of time. The larger the cluster, the longer the time that would be required for the mean number to become time independent. Hence, the bigger the system, the longer the time that would be required for the system to reach equilibrium. Given that the system is never truly in metastable equilibrium so that the ideas of Gibbs, Langer, and others are never exactly applicable, when is the system close enough to metastable equilibrium so that any possible simulation or experiment cannot detect the difference? We have found that the magnetization and energy are not sufficient indicators for nucleation and that the answer depends on the process being studied. For nucleation the equilibration of the number of clusters whose size is comparable to the size of the nucleating droplet is the relevant indicator.

ACKNOWLEDGMENTS

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APPENDIX: RELAXATION OF CLUSTERS AT THE CRITICAL TEMPERATURE

Accurate determinations of the dynamical critical exponent z have been found from the relaxation of the magnetization and energy at the critical temperature. In the following we take a closer look at the relaxation of the Ising model by studying the approach to equilibrium of the distribution of clusters of various sizes.

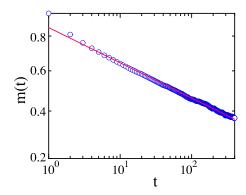


FIG. 13. (Color online) The relaxation of the magnetization m(t) of the nearest-neighbor 2D Ising model at $T=T_c$ starting from T_0 =0; L=5000. The straight line is the fit to a power law with slope ≈ 0.057 .

We consider the R=1 Ising model on a square lattice with L=5000. The system is initially equilibrated at either zero temperature $T_0=0$ (all spins up) or at $T_0=\infty$, and then instantaneously quenched to the critical temperature T_c . The Metropolis algorithm is used.

As a check on our results we first determine m(t) starting from T_0 =0. Scaling arguments suggest that m(t) approaches its equilibrium value as [33,34]

$$f(t) = Bt^{-\beta/\nu z} + f_{\infty},\tag{A1}$$

where the static critical exponents are $\beta=1/8$ and $\nu=1$. The fit of our results in Fig. 13 to Eq. (A1) yields the estimate $z\approx 2.19$, which is consistent with previous results [35]. Note that no time scale is associated with the evolution of m(t).

We next determined $n_s(t)$, the number of clusters of size s at time t after the temperature quench. Because all the spins are up at t=0, the number of (down) clusters of size s begins at zero and increases to its (apparent) equilibrium value $n_{s,\infty}$. The value of the latter depends on the size of the system.

Figure 14 shows the evolution of clusters of size s=100 for one run. Because we know of no argument for the time dependence of $n_s(t)-n_{s,\infty}$ except in the mean-field limit [34], we have to rely on empirical fits. We find that the time de-

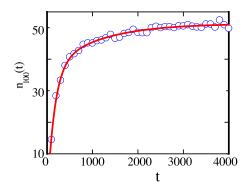


FIG. 14. (Color online) The evolution of the number of clusters of size s=100 at $T=T_c$ starting from $T_0=0$. The fit to Eq. (A2) gives $n_{s,\infty}=51.3$, $C_1=-42$, $C_2=-15$, $\tau_1=156$, and $\tau_2=1070$.

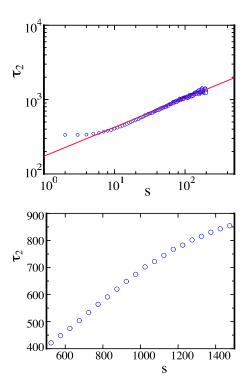


FIG. 15. (Color online) The relaxation time τ_2 versus the cluster size s at $T=T_c$ for R=1 starting from (a) $T_0=0$ and (b) $T_0=\infty$. The log-log plot in (a) yields $\tau_2 \sim s^{0.4}$.

pendence of $n_s(t)$ can be fitted to the sum of two exponentials,

$$n_s(t) - n_{s,\infty} = C_1 e^{-t/\tau_1} + C_2 e^{-t/\tau_2},$$
 (A2)

where C_1 , C_2 , τ_1 , and τ_2 are parameters to be fitted with $\tau_2 > \tau_1$.

Figure 15(a) shows the relaxation time τ_2 as a function of s for R=1 at $T=T_c$ starting from $T_0=0$. Note that the bigger the cluster, the longer it takes to reach its equilibrium distribution. That is, small clusters form first, and larger clusters are formed by the merging of smaller ones. The s dependence of τ_2 can be approximately fitted to a power law with the exponent 0.4.

To prepare a configuration at $T_0=\infty$, the system is randomized with approximately half of the spins up and half of the spins down. The temperature is instantaneously changed to $T=T_c$. As before, we focus on the relaxation of down spin clusters. In contrast to the $T_0=0$ case, the evolution of the clusters falls into three classes (see Fig. 16). For small clus-

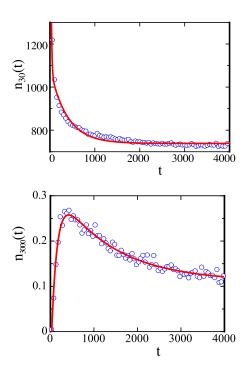


FIG. 16. (Color online) The time dependence of the number of clusters of size s=30 and s=3000 at T= T_c for R=1 starting from T_0 = ∞ . Note that n_{s =30 monotonically decreases to its equilibrium value and n_{s =1000 overshoots its equilibrium value. (a) C_1 =2367, C_2 =332, n_{s =30, ∞ =738, τ_1 =16, and τ_2 =403. (b) C_1 =-0.42, C_2 =0.22, n_{s =3000, ∞ =0.11, τ_1 =130, and τ_2 =1290.

ters $(1 \le s \le 40)$, n_s monotonically decreases to its equilibrium value. This behavior occurs because the initial random configuration has an abundance of small clusters so that lowering the temperature causes the small clusters to merge to form bigger ones. For intermediate size clusters (40 < s<4000), n_s first increases and then decreases to its equilibrium value. The initial growth is due to the rapid coalescence of smaller clusters to form intermediate ones. After there are enough intermediate clusters, they slowly coalesce to form bigger clusters. For clusters with s > 4000, n_s slowly increases to its equilibrium value. The range of sizes for these different classes of behavior depends on the system size. In all three cases $n_s(t)$ can be fitted to the sum of two exponentials. One of the two coefficients is negative for 40 < s<4000 for which $n_s(t)$ overshoots its equilibrium value. The relaxation time τ_2 is plotted in Fig. 15(b) as a function of s and does not appear to be a simple function of s for the range of values of s studied.

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