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Efficient numerical methods for the random-field Ising model: Finite-size scaling, reweighting extrapolation, and computation of response functions

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It was recently shown [Phys. Rev. Lett. **110**, 227201 (2013)] that the critical behavior of the random-field Ising model in three dimensions is ruled by a single universality class. This conclusion was reached only after a proper taming of the large scaling corrections of the model by applying a combined approach of various techniques, coming from the zero- and positive-temperature toolboxes of statistical physics. In the present contribution we provide a detailed description of this combined scheme, explaining in detail the zero-temperature numerical scheme and developing the generalized fluctuation-dissipation formula that allowed us to compute connected and disconnected correlation functions of the model. We discuss the error evolution of our method and we illustrate the infinite limit-size extrapolation of several observables within phenomenological renormalization. We present an extension of the quotients method that allows us to obtain estimates of the critical exponent α of the specific heat of the model via the scaling of the bond energy and we discuss the self-averaging properties of the system and the algorithmic aspects of the maximum-flow algorithm used.

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I. INTRODUCTION

The random-field Ising model (RFIM) is one of the archetypal disordered systems [1–11], extensively studied due to its theoretical interest, as well as its close connection to experiments in condensed matter physics [12–17]. In particular, several important systems can be studied through the RFIM: diluted antiferromagnets in a field [15], colloid-polymer mixtures [17,18], colossal magnetoresistance oxides [19,20], phase coexistence in the presence of quenched disorder [21–23], nonequilibrium phenomena such as the Barkhausen noise in magnetic hysteresis [24,25] or the design of switchable magnetic domains [26], etc.

The existence of an ordered ferromagnetic phase for the RFIM, at low temperature and weak disorder, followed from the seminal discussion of Imry and Ma [1], when the space dimension is greater than two ($D > 2$) [27–31]. This has provided us with a general qualitative agreement on the sketch of the phase boundary, separating the ordered ferromagnetic phase from the high-temperature paramagnetic one. The phase-diagram line separates the two phases of the model and intersects the randomness axis at the critical value of the disorder strength. Such qualitative sketching has been commonly used in most papers for the RFIM [32–37] and closed form quantitative expressions are also known from the early mean-field calculations [37]. However, it is generally true that the quantitative aspects of phase diagrams produced by mean-field treatments are very poor approximations.

On the theoretical side, a scaling picture is available [27–29]. The paramagnetic-ferromagnetic phase transition is ruled by a fixed point [in the renormalization-group (RG) sense] at temperature $T = 0$ [14]. The spatial dimension D is replaced by $D - \theta$, in hyperscaling relations ($\theta \approx D/2$). Nevertheless, one expects only two independent exponents [2,8,9,14], as in standard phase transitions [38].

Unfortunately, establishing the scaling picture is far from trivial. Perturbation theory predicts that, in $D = 3$, the ferromagnetic phase disappears upon applying the tiniest random field [3]. Even if the statement holds at all orders in perturbation theory [5], the ferromagnetic phase is stable in $D = 3$ [31]. Nonperturbative phenomena are obviously at play [39,40]. Indeed, it has been suggested that the scaling picture breaks down because of spontaneous supersymmetry breaking, implying that more than two critical exponents are needed to describe the phase transition [41].

On the experimental side, a particularly well researched realization of the RFIM is the diluted antiferromagnet in an applied magnetic field [15]. Yet, there are inconsistencies in the determination of critical exponents. In neutron scattering, different parametrizations of the scattering line shape yield mutually incompatible estimates of the thermal critical exponent, namely $\nu = 0.87(7)$ [42] and $\nu = 1.20(5)$ [43]. Moreover, the anomalous dimension $\eta = 0.16(6)$ [42] violates hyperscaling bounds, at least if one believes experimental claims of a divergent specific heat [16,44]. Clearly, a reliable parametrization of the line shape would be welcome. This program has been carried out for simpler, better understood problems [45]. Unfortunately, it is a common belief that we do not have such a strong command over the RFIM universality class.

The model has been also investigated by means of numerical simulations [34,46,47]. However, typical Monte Carlo schemes get trapped into local minima with escape time exponential in ξ^θ , where ξ denotes the correlation length. Although sophisticated improvements have appeared [48–52], these simulations produced low-accuracy data because they were limited to linear sizes of the order of $L_{\max} \leq 32$. Larger sizes can be achieved at $T = 0$, through the well-known mapping of the ground state to the maximum-flow optimization problem [53–66]. Yet, $T = 0$ simulations lack

many tools, standard at $T > 0$. In fact, the numerical data at $T = 0$ and their finite-size scaling analysis mostly resulted in strong violations of universality [55,57,58,60].

The criteria for determining the order of the low temperature phase transition and its dependence on the form of the field distribution have been discussed throughout the years [37,67–73]. In fact, different results have been proposed for different field distributions, like the existence of a tricritical point at the strong disorder regime of the system, present only in the bimodal distribution [37,69]. Currently, despite the huge efforts recorded in the literature, a clear picture of the model's critical behavior is still lacking. Although the view that the phase transition of the RFIM is of second order is well established [48–50,64], the extremely small value of the exponent β continues to cast some doubts. Moreover, a rather strong debate exists with regards to the role of disorder: the available simulations are not able to settle the question of whether the critical exponents depend on the particular choice of the distribution for the random fields, analogous to the mean-field theory predictions [37]. Thus the whole issue of the model's critical behavior is under intense investigation [41,48,49,51,52,74–78].

Recently, progress has been made towards this direction by the present authors [79]. In particular, using a combined approach of state of the art techniques from the pool of statistical physics and graph theory, it was shown that the universality class of the RFIM is independent of the form of the implemented random-field distribution. This, somehow unexpected, according to the current literature, result, was reached only after a proper taming of the large scaling corrections, a fact that, although emphasized many years ago [53], was overlooked in numerous subsequent relevant investigations of the model. In the current paper we present the full technical details of our numerical implementation, originally outlined in Ref. [79] and we provide some further numerical results relevant to the scaling behavior of the specific heat and the self-averaging aspects of the model in terms of the magnetic susceptibility and the bond energy. We also discuss the scaling aspects of the implemented maximum-flow algorithm.

The methods that we shall explain in the present paper will be useful way beyond the context of the 3D RFIM. The most obvious generalization is of course the RFIM in higher dimensions (see, e.g., [80]). However, similar ideas can be applied to many disordered systems and should be useful when one needs to take derivatives, or to perform reweighting extrapolations, with respect to the disorder-distribution parameters. The ability to obtain these derivatives is most important when the relevant RG fixed point lies at zero temperature (thus parameters other than temperature should be varied to cross the phase boundaries). For instance, for 2D Ising spin glasses several RG fixed-points appear at $T = 0$ depending on the nature of the couplings distribution [81]. It should be possible then to study the corresponding phase boundaries and RG flows using our formalism. Another difficult problem that can be tackled with the current prescription is the diluted antiferromagnet in a uniform external field [15]. The ground state of this model is degenerate, and it is thus difficult to sample with uniform probability from the set of all ground states [59]. Even if in experiments the external field is uniform, in simulations it is desirable to add a small, local random noise to the

magnetic field [52]. The small random magnetic fields make it possible to employ the full formalism that we derive in the following sections. Furthermore, the fluctuation-dissipation formulas elucidated below are also valid when working at finite (rather than zero) temperature, which is necessary for some algorithms [48,51].

The outline of paper is as follows: In the following Sec. II we define the model and the random-field distributions under study. In Sec. III we outline the $T = 0$ maximum-flow algorithm, and in Sec. IV we define the set of useful physical observables that will be mainly analyzed. However, a complication arises: the sought observables cannot be straightforwardly computed, as we explain in Sec. V. The problems are overcome in Sec. VI, where we derive explicitly a fluctuation-dissipation formalism that allowed us to compute connected and disconnected correlation functions from the $T = 0$ data for each field distribution distinctively. The use of a reweighting method with respect to the disorder strength consists another asset at hand of our combinatorial scheme. In Sec. VII we give a brief description of our finite-size scaling vehicle, the quotients method [82]. In Sec. VIII and on the basis of our main physical result of a single universality class [79], we illustrate the size evolution of several effective critical exponents and we present a finite-size scaling analysis of additional numerical data for the bond energy. For this latter task, we adopt an extension of the quotients method, necessary for monitoring the scaling of the effective exponent α of the specific heat. Furthermore, we discuss the self-averaging aspects of the model, by implementing a proper noise to signal ratio for the magnetic susceptibility and the bond energy, and we estimate the critical slowing-down exponent z of the zero-temperature algorithm used to generate the ground states of the model. Our contribution ends with a summary in Sec. IX.

II. MODEL AND RANDOM-FIELD DISTRIBUTIONS

Our $S_x = \pm 1$ spins are placed on a cubic lattice with size L and periodic boundary conditions. The Hamiltonian of the RFIM in a general form may be written as

$$\mathcal{H} = -J \sum_{\langle x,y \rangle} S_x S_y - \sum_x h_x S_x, \quad (1)$$

where in the above equation J is the nearest-neighbors' ferromagnetic interaction, which is set to be $J = 1$. With h_x we denote the set of independent quenched random fields. Common field distributions considered in the literature are the Gaussian and bimodal distributions [12,14,83], for which marginally distinct results have been proposed [55,57,58,60].

In the current work the quenched random fields h_x are extracted from one of the following double Gaussian (dG) or Poissonian (P) distributions (with parameters h_R, σ):

$$dG^{(\sigma)}(h_x; h_R) = \frac{1}{\sqrt{8\pi\sigma^2}} \left[e^{-\frac{(h_x - h_R)^2}{2\sigma^2}} + e^{-\frac{(h_x + h_R)^2}{2\sigma^2}} \right], \quad (2)$$

$$P(h_x; \sigma) = \frac{1}{2|\sigma|} e^{-|h_x|/\sigma}. \quad (3)$$

The limiting cases $\sigma = 0$ and $h_R = 0$ of Eq. (2) correspond to the well-known bimodal (b) and Gaussian (G) distributions, respectively. In the Poissonian and Gaussian cases the strength

of the random fields is parametrized by σ , while in the double Gaussian case we shall take $\sigma = 1$ and 2, and vary h_R .

As we are only interested in a $T = 0$ study of the model by estimating ground states via the use of efficient optimization methods that will be discussed below, a proper choice of the random-field distributions is of major importance in our task. In particular, the main advantage of considering the double Gaussian distribution of Eq. (2) is that one can mimic for certain values of σ the double-peak structure of the bimodal distribution, capturing its effects and at the same time escaping the implication of nondegenerate ground states. As is well known, for cases of discrete distributions, like the bimodal, degeneracy complicates the numerical solution of the system at $T = 0$, since one has to sweep over all the possible ground states of the system [56,59]. On the other hand, for the cases of the above distributions (2) and (3), the ground state of the system is nondegenerate, so it is sufficient to calculate just one ground state in order to get the necessary information.

III. ZERO-TEMPERATURE ALGORITHM

As already discussed extensively in the literature (see Refs. [84,85] and references therein), the RFIM captures essential features of models in statistical physics that are controlled by disorder and have frustration. Such systems show complex energy landscapes due to the presence of large barriers that separate several metastable states. If such models are studied using simulations mimicking the local dynamics of physical processes, it takes an extremely long time to encounter the exact ground state. However, there are cases where efficient methods for finding the ground state can be utilized and, fortunately, the RFIM is one such clear case. These methods escape from the typical direct physical representation of the system, in a way that extra degrees of freedom are introduced and an expanded problem is finally solved. By expanding the configuration space and choosing proper dynamics, the algorithm practically avoids the need of overcoming large barriers that exist in the original physical configuration space. An attractor state in the extended space is found in time polynomial in the size of the system and when the algorithm terminates, the relevant auxiliary fields can be projected onto a physical configuration, which is the guaranteed ground state.

The random field is a relevant perturbation at the pure fixed point, and the random-field fixed point is at $T = 0$ [27,86,87]. Hence the critical behavior is the same everywhere along the phase boundary and we can predict it simply by staying at $T = 0$ and crossing the phase boundary at the critical field point. This is a convenient approach because we can determine the ground states of the system exactly using efficient optimization algorithms [53–66,79,88–93] through an existing mapping of the ground state to the maximum-flow optimization problem [94–96]. A clear advantage of this approach is the ability to simulate large system sizes and disorder ensembles in rather moderate computational times. We should underline here that even the most efficient $T > 0$ Monte Carlo schemes exhibit extremely slow dynamics in the low-temperature phase of these systems [84,85]. Further assets in the $T = 0$ approach are the absence of statistical errors and equilibration problems, which, on the contrary, are the

two major drawbacks encountered in the $T > 0$ simulation of systems with rough free-energy landscapes [84,85].

The application of maximum-flow algorithms to the RFIM is nowadays well established [88]. The most efficient network flow algorithm used to solve the RFIM is the push-relabel algorithm of Tarjan and Goldberg [97]. For the interested reader, general proofs and theorems on the push-relabel algorithm can be found in standard textbooks [95,96]. In the present study we prepared our own C version of the algorithm that involves a modification proposed by Middleton *et al.* [63,64,98] that removes the source and sink nodes, reducing memory usage and also clarifying the physical connection [64,98]. For the sake of completeness, we recall here the algorithm we use, which is *exactly* the algorithm proposed in Refs. [63,64,98].

The algorithm starts by assigning an excess x_i to each lattice site i , with $x_i = h_i$. Residual capacity variables r_{ij} between neighboring sites are initially set to J . A height variable u_i is then assigned to each node via a global update step. In this global update, the value of u_i at each site in the set $\mathcal{T} = \{j | x_j < 0\}$ of negative excess sites is set to zero. Sites with $x_i \geq 0$ have u_i set to the length of the shortest path, via edges with positive capacity, from i to \mathcal{T} . The ground state is found by successively rearranging the excesses x_i , via push operations, and updating the heights, via relabel operations. When no more pushes or relabels are possible, a final global update determines the ground state, so that sites which are path connected by bonds with $r_{ij} > 0$ to \mathcal{T} have $\sigma_i = -1$, while those which are disconnected from \mathcal{T} have $\sigma_i = 1$. A push operation moves excess from a site i to a lower height neighbor j , if possible, that is, whenever $x_i > 0$, $r_{ij} > 0$, and $u_j = u_i - 1$. In a push, the working variables are modified according to $x_i \rightarrow x_i - \delta$, $x_j \rightarrow x_j + \delta$, $r_{ij} \rightarrow r_{ij} - \delta$, and $r_{ji} \rightarrow r_{ji} + \delta$, with $\delta = \min(x_i, r_{ij})$. Push operations tend to move the positive excess towards sites in \mathcal{T} . When $x_i > 0$ and no further push is possible, the site is relabeled, with u_i increased to $1 + \min_{\{j | r_{ij} > 0\}} u_j$. This is defined as a single push-relabel step; the number of such steps will be our measure of algorithmic time. In addition, if a set of highest sites \mathcal{U} becomes isolated, with $u_i > u_j + 1$, for all $i \in \mathcal{U}$ and all $j \notin \mathcal{U}$, the height u_i for all $i \in \mathcal{U}$ is increased to its maximum value, L^3 , as these sites will always be isolated from the negative excess nodes. The order in which sites are considered is given by a queue. In this paper, we have used the first-in-first-out (FIFO) queue [98]. The FIFO structure executes a push-relabel step for the site i at the front of a list. If any neighboring site is made active by the push-relabel step, it is added to the end of the list. If i is still active after the push-relabel step, it is also added to the end of the list. This structure maintains and cycles through the set of active sites. Last but not least, the computational efficiency of the algorithm has been increased via the use of periodic global updates every L^3 relabels [64,98].

Using the above version of the push-relabel algorithm, we performed large-scale simulations of the RFIM defined above in Eqs. (1)–(3) for a wide range of the simulation parameters. Our tactic included three steps: originally, we performed preliminary runs with $\mathcal{N}_{\text{samples}} = 10^6$, where $\mathcal{N}_{\text{samples}}$ counts the number of independent disorder realizations, to locate the h_R or σ values (depending on the parametrization) of

TABLE I. Summary of simulation details.

Distribution	L_{\min}	L_{\max}	$\mathcal{N}_{\text{samples}} (\times 10^6)$
G	8	192	10
dG $^{(\sigma=1)}$	8	128	50
dG $^{(\sigma=2)}$	8	128	10
P	8	192	10

the crossing points of the connected correlation length of the system for pairs of lattice sizes of the form $(L, 2L)$, as this is indicated in the main heart of the scaling method used (see below). Subsequently, the main part of the simulations took place in these estimated crossing points, with details, in terms of linear system sizes and disorder-averaged ensembles, summarized in Table I. In Table I $L_{\min}(L_{\max})$ denotes the minimum(maximum) linear size considered within the sequence of size points $L \in \{8, 12, 16, 24, 32, 48, 64, 96, 128, 192\}$. Finally, we performed an additional set of simulations for triplets of systems sizes as shown in Table II in order to compute the critical exponent of the specific heat via the scaling of the bond energy. This will be exemplified in Sec. VIII.

IV. OBSERVABLES

An instance of the random fields $\{h_x\}$ is named a sample. Thermal mean values are denoted as $\langle \dots \rangle$, while the subsequent average over samples is indicated by an overline. The two most basic quantities are the bond energy and the order-parameter density:

$$E_J = -J \sum_{\langle x,y \rangle} S_x S_y, \quad m = \frac{1}{L^D} \sum_x S_x. \quad (4)$$

A crucial feature of the RFIM is that we have to deal with *two* different correlation functions, namely the *disconnected* and the connected propagators.

 TABLE II. Effective critical exponent ratio $(\alpha - 1)/\nu$ using a three lattice-size variant $(L_1, L_2, L_3) = (L, 2L, 4L)$, see Eq. (71), of the original quotients method.

Distribution	(L_1, L_2, L_3)	$(\alpha - 1)/\nu$
G	(12,24,48)	-0.758(11)
	(16,32,64)	-0.793(17)
	(24,48,96)	-0.860(30)
	(32,64,128)	-0.881(75)
dG $^{(\sigma=1)}$	(16,32,64)	0.954(66)
	(24,48,96)	-0.036(23)
	(32,64,128)	-0.309(23)
dG $^{(\sigma=2)}$	(12,24,48)	-0.735(16)
	(16,32,64)	-0.766(16)
	(24,48,96)	-0.882(60)
	(32,64,128)	-0.867(56)
P	(12,24,48)	-1.120(6)
	(16,32,64)	-1.089(10)
	(24,48,96)	-1.071(42)
	(32,64,128)	-0.970(37)

The disconnected propagator, is straightforward to compute both in real, $G_{xy}^{(\text{dis})}$, and Fourier space, $\chi_k^{(\text{dis})}$:

$$G_{xy}^{(\text{dis})} = \overline{\langle S_x S_y \rangle}, \quad \chi_k^{(\text{dis})} = L^D \overline{\langle |m_k|^2 \rangle}_k, \quad (5)$$

where

$$m_k = \frac{1}{L^D} \sum_x e^{ik \cdot x} S_x. \quad (6)$$

In particular, special notations are standard for vanishing wave vector: $m_{k=(0,0,0)} = m$ (i.e., the order-parameter density) and $\chi_{k=(0,0,0)}^{(\text{dis})} = \chi^{(\text{dis})}$ (i.e., the disconnected susceptibility).

On the other hand, we have the connected propagator:

$$G_{xy} = \frac{\partial \langle S_x \rangle}{\partial h_y}. \quad (7)$$

At finite temperature, one could compute G_{xy} from the fluctuation-dissipation theorem

$$G_{xy} = \frac{1}{T} \overline{\langle S_x S_y \rangle - \langle S_x \rangle \langle S_y \rangle}. \quad (8)$$

However, we work directly at $T = 0$, as explained in Sec. III. Therefore, Eq. (8) is clearly unsuitable for us, and the methods of Sec. VI will be needed (see also Ref. [8]). For later use, we note the symmetry

$$G_{xy} = G_{yx} = \frac{G_{xy} + G_{yx}}{2}. \quad (9)$$

In fact, our numerical data will *never* verify this symmetry (because of statistical fluctuations); hence we prefer to use the symmetrized propagator $(G_{xy} + G_{yx})/2$. Now, the connected propagator in Fourier space is

$$\chi_k = \frac{1}{L^D} \sum_{x,y} e^{ik \cdot (x-y)} \frac{G_{xy} + G_{yx}}{2}. \quad (10)$$

Again, the case of vanishing wave vector deserves a special naming: $\chi_{k=(0,0,0)} = \chi$ is the connected susceptibility.

From both propagators, we compute the connected, ξ , and disconnected, $\xi^{(\text{dis})}$, second-moment correlation lengths [38,99]. Let $k_{\min} = (2\pi/L, 0, 0)$; then

$$\xi^{\#} = \frac{1}{2 \sin(\pi/L)} \sqrt{\frac{\chi^{\#}}{\chi_{k_{\min}}^{\#}} - 1}, \quad (11)$$

where the superscript $\#$ stands both for the connected or the disconnected case [100]. Of course, we improve our statistics by computing $\chi_{k_{\min}}^{\#} = \frac{1}{3} [\chi_{k=(2\pi/L, 0, 0)}^{\#} + \chi_{k=(0, 2\pi/L, 0)}^{\#} + \chi_{k=(0, 0, 2\pi/L)}^{\#}]$.

Other important quantities are the well-known universal Binder ratio

$$U_4 = \frac{\overline{\langle m^4 \rangle}}{\langle m^2 \rangle^2}, \quad (12)$$

and the susceptibilities ratio

$$U_{22} = \frac{\chi^{(\text{dis})}}{\chi^2} \quad (13)$$

that we use as a platform for investigating the validity of the so-called two-exponent scaling scenario; see Sec. VIII.

V. PROBLEMS WITH THE STRAIGHTFORWARD APPROACH

Computing response functions is very important. Unfortunately, the traditional approach for disordered systems (see, e.g., [101]) is not feasible at zero temperature. The problem is easily understood by considering the example of the Monte Carlo computation of the magnetic susceptibility.

The traditional approach would start by generating $\mathcal{N}_{\text{samples}}$ of the random fields according to the appropriate probability density $w(\{h_x\})$. Then, one would add to each random field a uniform external field

$$h_x \rightarrow h_x + H, \quad (14)$$

and the magnetic susceptibility would be estimated as

$$\chi^{\text{naive}} = \frac{1}{\mathcal{N}_{\text{samples}}} \sum_{s=1}^{\mathcal{N}_{\text{samples}}} \left. \frac{\partial \langle m_s \rangle_H}{\partial H} \right|_{H=0}, \quad (15)$$

where $\langle m_s \rangle_H$ is the thermal expectation value of instance s under the displaced magnetic fields in Eq. (14). Yet, as we explain below, the naive Monte Carlo estimator (15) yields $\chi^{\text{naive}} = 0$ with probability one for any smooth random-field probability density $w(\{h_x\})$ such as ours; recall Eqs. (2) and (3).

The approach outlined in Eq. (15) fails because, at zero temperature, the only spin assignment with a nonvanishing statistical weight is the ground state for the Hamiltonian (1). The crucial point is that the ground state is unique, excepting a zero-measure set in the L^D -dimensional space spanned by the random fields. Indeed, consider two arbitrary but fixed spin assignments, $\{S_x^{(1)}\}$ and $\{S_x^{(2)}\}$. The condition of equal energy

$$\mathcal{H}(\{S_x^{(1)}\}) = \mathcal{H}(\{S_x^{(2)}\}) \quad (16)$$

defines a hyperplane in the random-fields space. There are $2^{L^D} (2^{L^D} - 1)/2$ such space-dividing hyperplanes. For random fields $\{h_x\}$ not in these hyperplanes, each of the 2^{L^D} possible spin assignments has a distinct energy, and thus the ground state is unique. Furthermore, the ordering of the 2^{L^D} energy levels is fixed away from the hyperplanes (which are the locus in random-fields space where level crossings happen).

Now, let us suppose that none of the $\mathcal{N}_{\text{samples}}$ instances in Eq. (15) lies *exactly* in one of the dividing hyperplanes [this happens with probability one for any smooth $w(\{h_x\})$]. Then, for H small enough, the fields displacement in Eq. (14) will not cross any of the hyperplanes and thus adding the field H will leave the ground state unvaried. In other words, $d\langle m_s \rangle_H / dH|_{H=0} = 0$, with probability one.

However, the connected susceptibility is *not* zero. The way out of the paradox is simple: the H derivatives in Eq. (15) are actually a sum of Dirac δ functions, centered at the precise H values that cause the displaced fields (14) to cross some of the dividing hyperplanes (16). It is the integral over the random fields of these Dirac δ functions which produces a finite susceptibility $\chi > 0$:

$$\chi = \int \prod_x dh_x w(\{h_x\}) \left. \frac{\partial \langle m \rangle}{\partial H} \right|_{H=0}. \quad (17)$$

We see the heart of the problem: naive Monte Carlo estimations such as Eq. (15) cannot correctly reproduce integrals such as

Eq. (17) when the integrand is such a singular object as a sum of Dirac's δ functions.

Nevertheless, people have tried to overcome the zero-measure problem. For instance, one could keep H finite and compute the Monte Carlo (MC) average

$$[\langle m \rangle_H]^{(\text{MC})} = \frac{1}{\mathcal{N}_{\text{samples}}} \sum_{s=1}^{\mathcal{N}_{\text{samples}}} \langle m_s \rangle_H, \quad (18)$$

and then try to extrapolate to $H \rightarrow 0$ the slope $d[\langle m \rangle_H]^{(\text{MC})} / dH$. Of course, the smaller is H the larger is the number $\mathcal{N}_{\text{samples}}$ needed to observe some H dependency. Yet, reasonable tradeoffs between number of instances and size of the applied field could be empirically found [61].

In Sec. VI we explain a completely different approach that (i) allows one to work directly at $H = 0$ and (ii) avoids Dirac's δ functions. How this is possible can be easily understood by considering the following one-dimensional toy model.

Toy model

Imagine we have a single random field h . In analogy with the general case, let us also assume that the magnetization, regarded as a function of h , is constant but for a set of R discontinuities:

$$\langle m \rangle_h = -1 + \sum_{i=1}^R [m_{i+1} - m_i] \theta(h - h_i). \quad (19)$$

In the above expression $\theta(x)$ is the Heaviside step function, $\theta(x > 0) = 1$ and $\theta(x < 0) = 0$, and the magnetization plateaus are monotonically increasing, $m_{i+1} > m_i$, with $m_1 = -1$ and $m_{R+1} = 1$.

Now, if we displace the field, $h \rightarrow h + H$, and take the H derivative in Eq. (19), a sum of Dirac δ functions will arise, making unfeasible the Monte Carlo method.

However, it is useful to take one step back and recall how the susceptibility is defined. First, we consider the average magnetization as a function of the displaced field

$$\overline{\langle m \rangle}(H) = \int_{-\infty}^{\infty} dw(h) \langle m \rangle_{h+H}. \quad (20)$$

The derivative with respect to H is taken *only after* computing the integral [the random-field probability density $w(h)$ must decrease fast enough at infinity to make the integral convergent]. Yet, a change of variable $h' = h + H$ yields

$$\overline{\langle m \rangle}(H) = \int_{-\infty}^{\infty} dw(h - H) \langle m \rangle_h. \quad (21)$$

The change of variable is mathematically sound, as it relies only on the translational invariance of the integration measure in Eq. (20). If the probability density $w(h)$ is smooth, one can now interchange derivative and integral obtaining

$$\chi_{\text{toy model}} = \int_{-\infty}^{\infty} dw(h) \frac{-1}{w(h)} \frac{dw}{dh} \langle m \rangle_h. \quad (22)$$

The integrand in Eq. (22) is a regular function, allowing for a Monte Carlo estimation of the form

$$\chi_{\text{toy model}}^{(\text{MC})} = \frac{1}{\mathcal{N}_{\text{samples}}} \sum_{s=1}^{\mathcal{N}_{\text{samples}}} \langle m \rangle_{h_s} \left. \frac{-1}{w(h_s)} \frac{dw}{dh} \right|_{h=h_s}, \quad (23)$$

where the independent random fields h_s are obtained with weight $w(h)$. Note that the summands in Eq. (23) *cannot* be interpreted as the magnetic susceptibility of a given instance (there are no Dirac δ functions). However, $\chi_{\text{toy model}}^{(\text{MC})}$ does converge to $\chi_{\text{toy model}}$ in the limit of large $\mathcal{N}_{\text{samples}}$.

VI. FLUCTUATION-DISSIPATION FORMALISM

Reweighting methods are a major asset for numerical studies of critical phenomena [102,103]: From a *single* simulation at a given temperature we get a *continuous* curve for (say) the disconnected susceptibility, $\chi^{(\text{dis})}(T)$.

However, we will be working at zero temperature. Hence standard reweighting methods are not useful for us. In fact, we shall explain here our extension of reweighting methods originally devised for percolation studies [101,104,105]. From a single simulation, we extrapolate the mean value of observables to nearby parameters of the disorder distribution. We varied σ for the Poissonian and Gaussian distributions, see panel (a) in Fig. 1 below for an illustrative flavor, and h_R for the double Gaussian distribution. These reweighting methods were instrumental for our previous work [79].

As we discuss below, a closely related problem is the computation of the connected correlation functions (recall also Sec. V). Our solution for the case of the Gaussian distribution, in Sec. VIA, will turn out to be identical to the one in Ref. [8]. However, modifications are needed for the Poissonian or double Gaussian distributions, which are explained in Secs. VIB and VIC, respectively.

For all three distributions, we shall compute the connected propagator by adding a *source* \tilde{h}_x to the random fields:

$$h_x \rightarrow h_x + \epsilon \tilde{h}_x, \quad (24)$$

where ϵ is a small parameter. At variance with the random fields $\{h_x\}$, the sources $\{\tilde{h}_x\}$ will be arbitrary but *fixed*: the overline will indicate and average only with respect to the $\{h_x\}$. Then, the connected propagator $G_{xy}(= G_{yx})$ follows from the Taylor expansion

$$\overline{\langle S_y \rangle_{\{h_x + \epsilon \tilde{h}_x\}}} = \overline{\langle S_y \rangle} + \epsilon \sum_x G_{yx} \tilde{h}_x + \mathcal{O}(\epsilon^2). \quad (25)$$

In the above expression, $\langle S_y \rangle_{\{h_x + \epsilon \tilde{h}_x\}}$ is the thermal expectation value obtained when plugging $\{h_x + \epsilon \tilde{h}_x\}$ as the random magnetic fields in the Hamiltonian, Eq. (1).

The formalism will be explained in the same way, for all three random-field distributions. We start from the general observation that computing thermal expectation values at $T = 0$ is trivial: one just needs to evaluate the function of interest, see Sec. IV, on the ground-state spin assignment corresponding to a given sample (recall that a sample is characterized by a set of random fields $\{h_x\}$). In this sense, thermal mean values are mere functions of the $\{h_x\}$. Next, we observe that a special function of the random fields, when averaged over the $\{h_x\}$, is equal to the connected propagator. Finally, we show how to

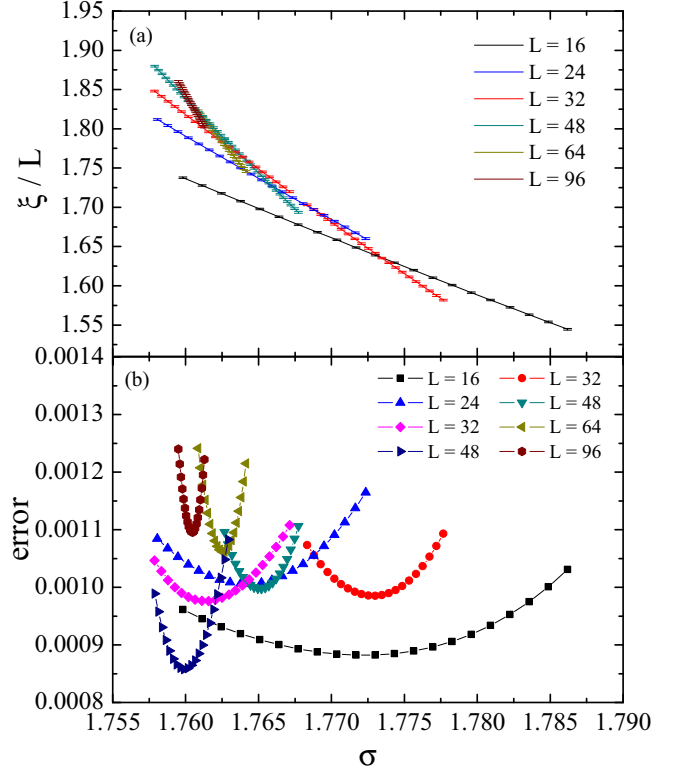


FIG. 1. (a) For several system sizes, we show ξ/L as a function of the strength of the Poissonian random field σ . Lines join data obtained from reweighting extrapolation (discontinuous lines of the same color come from independent simulations). In the large- L limit, ξ/L is L independent at the critical point $\sigma^{(c)}$. In the quotients method, we consider the ξ/L curves for pair of lattices $(L, 2L)$ and seek the σ where they cross. This crossing is employed for computing effective, L -dependent critical exponents with Eq. (68). (b) Illustration of statistical errors in the universal ratio ξ/L for the pairs of the system sizes shown in panel (a).

perform reweighting extrapolations for a generic function of the random fields $\mathcal{F}(\{h_x\})$.

Before we start, let us mention that a practical consideration had an important impact in the designing of our fluctuation-dissipation formalism. We simulated a large number of samples ($\sim 10^7$) on large system sizes ($L = 192$); see Table I. Clearly, storing in the hard drive all the corresponding ground-state assignments is out of the question. Therefore, we need to select *beforehand* a small set of quantities to be computed on the ground-state spin assignment and stored on the hard drive. This small set of observables includes E_J , m , and $m_{k_{\min}}$, recall Sec. IV, but also the quantities needed to compute the connected propagators and the reweighting extrapolations [in all cases, we restricted the wave vectors to a bare minimum: $k = (0, 0, 0)$ and $k = k_{\min}$].

A. Gaussian distribution

The combined probability density for our $N=L^D$ Gaussian random fields with width parameter σ is

$$w^G(\{h_x\}, \sigma) = \frac{1}{(2\pi\sigma^2)^{\frac{N}{2}}} e^{-\frac{1}{2\sigma^2} \sum_x h_x^2}. \quad (26)$$

Our computation starts from Eq. (25):

$$\overline{\langle S_y \rangle_{\{h_x + \epsilon \tilde{h}_x\}}} = \int \prod_x dh_x w^G(\{h_x\}, \sigma) \langle S_y \rangle_{\{h_x + \epsilon \tilde{h}_x\}} \quad (27)$$

$$= \int \prod_x dh'_x w^G(\{h'_x - \epsilon \tilde{h}_x\}, \sigma) \langle S_y \rangle_{\{h'_x\}}. \quad (28)$$

In the above expressions the N integrals extend from $-\infty$ to $+\infty$. We went from (27) to (28) by changing integration variables as $h'_x = h_x + \epsilon \tilde{h}_x$ [we shall drop the prime for the dummy integration variables, h'_x , in Eq. (28)]. Now, one just needs to Taylor expand in the small parameter ϵ in Eq. (28). A direct comparison with Eq. (25) yields

$$G_{zy} = \int \prod_x dh_x w^G(\{h_x\}, \sigma) \frac{h_z \langle S_y \rangle_{\{h_x\}}}{\sigma^2} \quad (29)$$

$$= \frac{\overline{h_z \langle S_y \rangle}}{\sigma^2}. \quad (30)$$

We now use Eq. (10) to compute the propagator in the Fourier space

$$\chi_k = L^D \frac{\overline{h_{-k}^G m_k + h_k^G m_{-k}}}{2\sigma^2}, \quad (31)$$

where m_k was defined in Eq. (6) and

$$h_k^G = \frac{1}{L^D} \sum_x e^{ik \cdot x} h_x. \quad (32)$$

The reader will note that Eq. (31) was obtained in Ref. [8] (yet, our argument is sound as well when one starts directly at $T = 0$, which is exactly our case). Our rationale for recalling this fluctuation-dissipation argument here is that the derivation of the new formulas in Secs. VIB and VIC is completely analogous.

At this point it should be obvious that, for all the observables of interest, we are after the computation of multidimensional integrals of the form

$$\overline{\mathcal{F}}|_\sigma = \int \prod_x dh_x w^G(\{h_x\}, \sigma) \mathcal{F}(\{h_x\}), \quad (33)$$

where $\mathcal{F}(\{h_x\})$ could be $\mathcal{F} = \langle S_z S_y \rangle_{\{h_x\}}$, or $\mathcal{F} = h_z \langle S_y \rangle_{\{h_x\}}$, etc. Now, we need to solve the following three problems.

(1) Compute derivatives with respect to σ , $D_\sigma \overline{\mathcal{F}}$. Recall that σ is the width for the Gaussian weight in Eq. (33).

(2) Extrapolate the expectation values at $\sigma + \delta\sigma$ from integrals at σ such as Eq. (33).

(3) Estimate how large the extrapolation window $\delta\sigma$ may be in a numerical simulation.

Fortunately, we can solve all three problems with a single trick. The starting point is

$$\overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \int \prod_x dh_x w^G(\{h_x\}, \sigma + \delta\sigma) \mathcal{F}(\{h_x\}) \quad (34)$$

$$= \int \prod_x dh_x w^G(\{h_x\}, \sigma) \mathcal{F}(\{h_x\}) \mathcal{R}(\{h_x\}, \sigma, \delta\sigma), \quad (35)$$

where the reweighting factor \mathcal{R} is just the ratio of probability densities:

$$\mathcal{R}(\{h_x\}, \sigma, \delta\sigma) = \frac{w^G(\{h_x\}, \sigma + \delta\sigma)}{w^G(\{h_x\}, \sigma)} \quad (36)$$

$$= \left(\frac{\sigma}{\sigma + \delta\sigma} \right)^N e^{\frac{1}{2}[\sigma^{-2} - (\sigma + \delta\sigma)^{-2}] \sum_x h_x^2}. \quad (37)$$

The computation of σ derivatives follows straightforwardly by Taylor expanding the reweighting factor in $\delta\sigma$:

$$\mathcal{R}(\{h_x\}, \sigma, \delta\sigma + \epsilon) = \mathcal{R}(\{h_x\}, \sigma, \delta\sigma) (1 + \epsilon \mathcal{D} + \mathcal{O}(\epsilon^2)), \quad (38)$$

where

$$\mathcal{D}(\{h_x\}, \sigma, \delta\sigma) = \frac{1}{\sigma + \delta\sigma} \left[\frac{\sum_x h_x^2}{(\sigma + \delta\sigma)^2} - N \right]. \quad (39)$$

Our reweighting formulas can be cast in a more aesthetically appealing form

$$\overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \overline{\mathcal{F} \mathcal{R}_{\sigma, \delta\sigma}}|_\sigma, \quad D_\sigma \overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \overline{\mathcal{F} \mathcal{D}_{\sigma, \delta\sigma}}|_\sigma. \quad (40)$$

Note that Eq. (40) refers to a function \mathcal{F} of the random fields *only*. Explicit dependency on σ , like in $G_{zy} = \overline{h_z \langle S_y \rangle} / \sigma^2$, is not included but can be taken care of straightforwardly.

In summary, in order to perform a complete reweighting study for each sample we need to store on the hard drive only E_J , m_k , $h_{-k}^G m_k + h_k^G m_{-k}$ [restricting ourselves to $k = (0, 0, 0)$ and $k = k_{\min}$], as well as $\sum_x h_x^2$.

The final question we need to address is the following: how large can $\delta\sigma$ reasonably be in a Monte Carlo simulation? Of course, the question is ill posed, because the answer depends on how many samples are simulated. In the limit of infinite statistics, one could have arbitrarily large $\delta\sigma$. However, this ideal situation is never reached in practice. As a rule of thumb one may use many different criteria, but all of them boil down to requiring that the *typical* set of random fields for $\sigma + \delta\sigma$ could also be typical at σ (or, at least, not too unusual). A particularly simple such criterium requires the absolute value of

$$\overline{\sum_x h_x^2}|_{\sigma+\delta\sigma} - \overline{\sum_x h_x^2}|_\sigma = N[(\sigma + \delta\sigma)^2 - \sigma^2] \quad (41)$$

to be no larger than the dispersion of $\sum_x h_x^2$ at σ , namely $\sqrt{2N}\sigma^2$. The resulting bound is

$$|\delta\sigma| \leq \sqrt{\frac{\sigma^2}{2N}}. \quad (42)$$

B. Poissonian distribution

This case is a straightforward translation of the results in Sec. VIA. Since there is not any new idea involved, let us just give the main results.

The connected propagator in real space is

$$G_{zy} = \frac{\overline{h_z \langle S_y \rangle}}{|h_z| \sigma}. \quad (43)$$

Note the small, but crucial, difference with Eq. (30): we correlate $\langle S_y \rangle$ with the *sign* of h_z . In the Fourier space, Eq. (30) translates to

$$\chi_k = L^D \frac{\overline{h_{-k}^P m_k + h_k^P m_{-k}}}{2\sigma}, \quad (44)$$

where m_k was defined in Eq. (6) and

$$h_k^p = \frac{1}{L^D} \sum_x e^{ik \cdot x} \frac{h_x}{|h_x|}. \quad (45)$$

Note, again, that we Fourier transform the sign of the Poissonian random fields.

The reweighting factor is again the ratio of probability densities for the Poisson fields:

$$\mathcal{R}(\{h_x\}, \sigma, \delta\sigma) = \left(\frac{\sigma}{\sigma + \delta\sigma} \right)^N e^{[\sigma^{-1} - (\sigma + \delta\sigma)^{-1}] \sum_x |h_x|}, \quad (46)$$

and the derivative operator follows from a Taylor expansion with respect to $\delta\sigma$:

$$\mathcal{R}(\{h_x\}, \sigma, \delta\sigma + \epsilon) = \mathcal{R}(\{h_x\}, \sigma, \delta\sigma) (1 + \epsilon \mathcal{D} + \mathcal{O}(\epsilon^2)), \quad (47)$$

where

$$\mathcal{D}(\{h_x\}, \sigma, \delta\sigma) = \frac{1}{\sigma + \delta\sigma} \left[\frac{\sum_x |h_x|}{(\sigma + \delta\sigma)} - N \right]. \quad (48)$$

The final reweighting formulas can be cast in exactly the same form that we found for the Gaussian random fields

$$\overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \overline{\mathcal{F}} \mathcal{R}_{\sigma, \delta\sigma} |_{\sigma}, \quad D_{\sigma} \overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \overline{\mathcal{F}} \mathcal{R}_{\sigma, \delta\sigma} D_{\sigma} |_{\sigma}. \quad (49)$$

As for the maximum reasonable reweighting extrapolation, we also use an analogous criterium: the absolute value of the difference

$$\sum_x |h_x| |_{\sigma+\delta\sigma} - \sum_x |h_x| |_{\sigma} = N(\sigma + \delta\sigma) - N\sigma \quad (50)$$

should be no larger than the dispersion of $\sum_x |h_x|$ at σ , namely $\sqrt{N}\sigma$. The resulting bound is

$$|\delta\sigma| \leq \sqrt{\frac{\sigma^2}{N}}. \quad (51)$$

C. Double Gaussian distribution

Our formalism for this distribution is slightly more complicated. Let us start by explaining how we obtain a random field distributed as prescribed in Eq. (2). For each h_x we extract two independent random variables. One of them is discrete, $\eta_x = \pm 1$, with 50% probability. The other variable, φ_x , is Gaussian distributed with zero mean and unit variance. Then, we set [the width σ is given in Eq. (2)]

$$h_x = h_R \eta_x + \sigma \varphi_x. \quad (52)$$

The combined probability density for our $2N$ variables is

$$w^{\text{dG}}(\{\eta_x, \varphi_x\}) = \frac{e^{-\frac{1}{2} \sum_x \varphi_x^2}}{2^N (2\pi)^{N/2}}. \quad (53)$$

In order to understand the origin of the additional complications for this distribution, let us add a source (i.e., $h_x \rightarrow h_x + \epsilon \tilde{h}_x$), while we simultaneously modify the position of the peaks (i.e., $h_R \rightarrow h_R + \delta h_R$) [106].

Under our circumstances, Eq. (25) reads

$$\overline{\langle S_y \rangle_{\{h_x + \epsilon \tilde{h}_x\}} |_{h_R + \delta h_R}} = \sum_{\{\eta_x\}} \int \prod_x d\varphi_x w^{\text{dG}}(\{\eta_x, \varphi_x\}) \langle S_y \rangle_{\{\tilde{h}_x\}}, \quad (54)$$

$$\hat{h}_x = h_R \eta_x + \delta h_R \eta_x + \sigma \varphi_x + \epsilon \tilde{h}_x. \quad (55)$$

The sum in Eq. (54) extends to the 2^N possible values of the discrete variables η_x . The problem now is apparent from Eq. (55). For each site, we have a single integration variable, namely φ_x . We need to carry out a change of variable that brings Eq. (55) to the form in Eq. (52):

$$\varphi'_x = \varphi_x + \frac{\delta h_R \eta_x + \epsilon \tilde{h}_x}{\sigma}. \quad (56)$$

In other words, if $\delta h_R \neq 0$, there is no way of distinguishing $\delta h_R \eta_x$ from the source term $\epsilon \tilde{h}_x$.

With this caveat in mind, and dropping the prime in the dummy integration variables, Eq. (54) now reads

$$\begin{aligned} \overline{\langle S_y \rangle_{\{h_x + \epsilon \tilde{h}_x\}} |_{h_R + \delta h_R}} &= \sum_{\{\eta_x\}} \int \prod_x d\varphi_x w^{\text{dG}} \\ &\times \left(\left\{ \eta_x, \varphi_x - \frac{\delta h_R \eta_x + \epsilon \tilde{h}_x}{\sigma} \right\} \right) \langle S_y \rangle_{\{\tilde{h}_x\}}, \end{aligned} \quad (57)$$

$$\hat{h}_x = h_R \eta_x + \sigma \varphi_x. \quad (58)$$

Now,

$$\begin{aligned} w^{\text{dG}} \left(\left\{ \eta_x, \varphi_x - \frac{\delta h_R \eta_x + \epsilon \tilde{h}_x}{\sigma} \right\} \right) \\ &= w^{\text{dG}}(\{\eta_x, \varphi_x\}) \mathcal{R}(\{\eta_x, \varphi_x\}, \delta h_R) \\ &\times \exp \left[\frac{\epsilon}{\sigma} \sum_x \tilde{h}_x \left(\varphi_x - \frac{\delta h_R \eta_x}{\sigma} \right) \right] \exp \left[-\frac{\epsilon^2}{2\sigma^2} \sum_x \tilde{h}_x^2 \right], \end{aligned} \quad (59)$$

where the reweighting factor appropriate for our implementation of the double Gaussian distribution is

$$\mathcal{R}(\{\eta_x, \varphi_x\}, \delta h_R) = \exp \left[\frac{\delta h_R}{\sigma} \sum_x \eta_x \varphi_x \right] \exp \left[-\frac{\delta h_R^2 N}{2\sigma^2} \right]. \quad (60)$$

Taylor expanding with respect to ϵ in Eq. (59) we finally get the connected propagator

$$G_{zy} |_{h_R + \delta h_R} = \frac{1}{\sigma} \overline{\mathcal{R}_{\delta h_R} \left(\varphi_z - \frac{\delta h_R \eta_z}{\sigma} \right) \langle S_y \rangle} \Big|_{h_R}. \quad (61)$$

In particular, the correction term $-\delta h_R \eta_z$ was absent for the Poissonian and Gaussian distributions. Similarly, one can get the connected propagator in the Fourier space

$$\begin{aligned} \chi_k |_{h_R + \delta h_R} &= \frac{L^D}{\sigma} \overline{\mathcal{R}_{\delta h_R} \left(\hat{\varphi}_{-k} - \frac{\delta h_R \hat{\eta}_{-k}}{\sigma} \right) m_k} \\ &+ \overline{\left(\hat{\varphi}_k - \frac{\delta h_R \hat{\eta}_k}{\sigma} \right) m_{-k}} \Big|_{h_R}, \end{aligned} \quad (62)$$

where m_k was defined in Eq. (6) and

$$\hat{\varphi}_k = \frac{1}{L^D} \sum_x e^{ik \cdot x} \varphi_x, \quad \hat{\eta}_k = \frac{1}{L^D} \sum_x e^{ik \cdot x} \eta_x. \quad (63)$$

Instead, for disconnected observables (e.g., E_J , connected propagators, etc.), the reweighting formulas take the standard form

$$\overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \overline{\mathcal{F}\mathcal{R}_{\delta h_R}}|_{\sigma}, \quad D_{\sigma}\overline{\mathcal{F}}|_{\sigma+\delta\sigma} = \overline{\mathcal{F}\mathcal{R}_{\delta h_R}\mathcal{D}_{\delta h_R}}|_{\sigma}, \quad (64)$$

where, in this case, the derivative operator is

$$\mathcal{D}(\{\eta_x, \varphi_x\}, \delta h_R) = \frac{1}{\sigma} \sum_x \left(\eta_x \phi_x - \frac{\delta h_R}{\sigma} \right). \quad (65)$$

Finally, we need to assess the maximum sensible size for δh_R . The simplest way to proceed is to compute the moments of the reweighting factor

$$\overline{\mathcal{R}_{\delta h_R}^k} = \exp \left[N \frac{k^2 - k}{2} \frac{\delta h_R^2}{\sigma^2} \right]. \quad (66)$$

If we now demand the dispersion (i.e., square root of variance) to be as large as the mean value, we get

$$|\delta h_R| \leq \sigma \sqrt{\frac{\log 2}{N}}. \quad (67)$$

VII. QUOTIENTS METHOD

To extract the values of critical points, critical exponents, and dimensionless universal quantities, we employed the quotients method, also known as phenomenological renormalization [38,82,107]. This method allows for a particularly transparent study of corrections to scaling, that up to now have been considered as the Achilles' heel in the study of the $D \geq 3$ random-field problem. We should note that previous applications of the method include diluted antiferromagnets [48] and the spin-glass problem; see Ref. [108] and references therein.

We compare observables computed in a pair of lattices ($L, 2L$). We start imposing scale invariance by seeking the L -dependent critical point: the value of σ (h_R for the dG), such that $\xi_{2L}/\xi_L = 2$ [i.e., the crossing point for ξ_L/L , see Fig. 1(a)]. Now, for dimensionful quantities O , scaling in the thermodynamic limit as $\xi^{x_O/\nu}$, we consider the quotient $Q_O = O_{2L}/O_L$ at the crossing. For dimensionless magnitudes g , we focus on g_{2L} . In either case, one has

$$Q_O^{(\text{cross})} = 2^{x_O/\nu} + \mathcal{O}(L^{-\omega}), \quad g_{(2L)}^{(\text{cross})} = g^* + \mathcal{O}(L^{-\omega}), \quad (68)$$

where x_O/ν , g^* , and the scaling-corrections exponent ω are universal.

Examples of dimensionless quantities are the connected and disconnected correlation lengths over the system size, i.e., ξ/L and $\xi^{(\text{dis})}/L$, and the Binder ratio U_4 . Instances of dimensionful quantities are then the derivatives of ξ , $\xi^{(\text{dis})}$ ($x_{\xi} = 1 + \nu$), the connected and disconnected susceptibilities χ and $\chi^{(\text{dis})}$ [$x_{\chi} = \nu(2 - \eta)$, $x_{\chi^{(\text{dis})}} = \nu(4 - \bar{\eta})$], and the ratio U_{22} [$x_{U_{22}} = \nu(2\eta - \bar{\eta})$] (see also Sec. IV), which as already noted above will serve as an alternative platform for investigating the validity of the so-called two-exponent scaling scenario [8,9].

Let us point out here that an extension of the quotients method using the sequence of three lattice-size points ($L, 2L, 4L$) will be presented below in Sec. VIII. This generalization is necessary for the scaling study of the bond energy of the RFIM, which is governed by a nondiverging background term.

VIII. RESULTS AND DISCUSSION

Let us start with a few illustrations on the main heart of the scaling method applied, that is the crossing of the universal ratio ξ/L together with the error evolution of the presented numerical scheme. As already mentioned above, we varied σ for the Poissonian and Gaussian distributions, see panel (a) in Fig. 1, and h_R for the double Gaussian distribution. Panel (b) in Fig. 1 shows the statistical errors of the universal ratio corresponding to the pairs of system sizes shown in panel (a) of the same figure. As expected, the error is minimal exactly at the simulation point denoted hereafter as $\sigma^{(s)}$ or $h_R^{(s)}$, respectively, and increases further away from it. Furthermore, a comparative illustration with respect to the errors induced by the reweighting method and the disorder averaging process is shown in Fig. 2 again for the universal ratio ξ/L of an $L = 64$ Poissonian RFIM and for three sets of simulations, as outlined in the figure. Clearly, this latter accuracy test serves in favor of the proposed scheme.

The full application of Eq. (68) to our four random-field distributions has been summarized in Table II of Ref. [79], where all the estimates of critical points, universal ratios, and critical exponents are given, together with the corrections-to-scaling exponent $\omega = 0.52(10)$ (see also Fig. 4 in Ref. [79]). In particular, the computation of the corrections-to-scaling exponent ω has been performed by means of a joint fit, third-order polynomial in $L^{-\omega}$, for the dimensionless quantities ξ/L , $\xi^{(\text{dis})}/L$, and U_4 using data for $L \geq 24$. We should point out that joint fits share the value of some fitting parameters such as the $L \rightarrow \infty$ extrapolation (which is the same for all random-field distributions), or the corrections-to-scaling exponent ω (which is common to all magnitudes). Here, we provide some

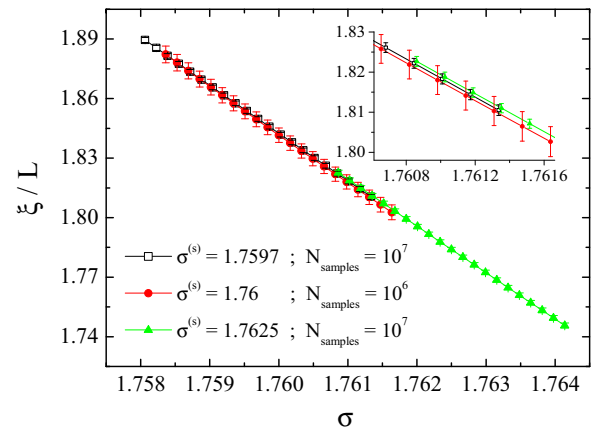


FIG. 2. Universal ratio ξ/L of an $L = 64$ Poissonian RFIM for three different sets of simulations, corresponding to different simulation values $\sigma^{(s)}$ and different sets of random realizations. The inset is a mere enlargement of the intermediate regime of σ values.

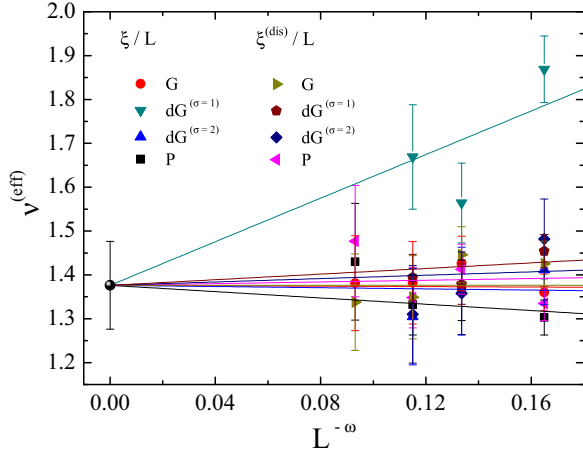


FIG. 3. Infinite limit-size extrapolation of the effective critical exponent ν .

complementary illustrations, showing the infinite limit-size extrapolation of the effective exponents of the correlation length ν , the anomalous dimension η , and the two-exponent difference $2\eta - \bar{\eta}$, the latter serving as an independent test of the two-exponent scaling scenario in the theory of the random-field problem [8].

Figures 3 and 4 illustrate the infinite limit-size extrapolation of the effective exponents ν and η , respectively, for all the four types of distributions studied. The solid lines are joint polynomial fits of first order in $L^{-\omega}$ including data points for $L \geq 32$, extrapolating to $L^{-\omega} = 0$, as shown by the filled circle in each figure. We remind the reader that for the effective exponent ν we have two sets of data for each of the four distributions coming from the connected and disconnected correlation lengths [79]. Let us comment here that our estimation $\nu = 1.38(10)[3]$ is similar to the most modern computations that suggest on average a value of $1.35(7)$ [61,63,65,66]. For the anomalous dimension estimate $\eta = 0.5153(9)[2]$, we note also $\eta = 0.50(3)$ from Ref. [61] as a comparison. Obviously, our errors for ν are

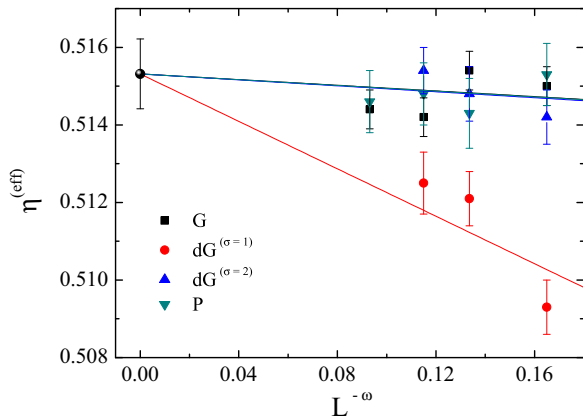


FIG. 4. Infinite limit-size extrapolation of the effective critical exponent η . Four solid lines are shown corresponding to the four random-field distributions as in Fig. 3, although they are not easily discerned due to very close values of their linear coefficient terms.

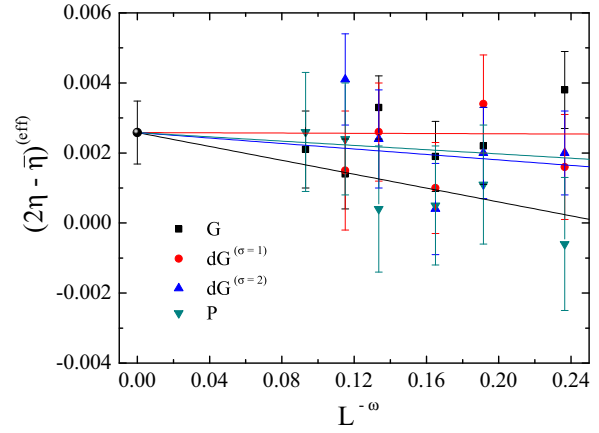


FIG. 5. Infinite limit-size extrapolation of the effective exponent $2\eta - \bar{\eta}$.

larger than those for η because we compute derivatives as connected correlations [109] (see also the discussion in Sec. VI).

Subsequently in Fig. 5 we perform an extrapolation of the effective exponent difference $2\eta - \bar{\eta}$, corresponding to the dimensional quantity U_{22} , Eq. (13), in order to discuss the two-exponent scaling scenario. The solid lines in this figure illustrate a joint polynomial fit, first-order in $L^{-\omega}$, including data points for $L \geq 16$, giving $2\eta - \bar{\eta} = 0.0026(9)[1]$ as shown by the filled black circle at $L^{-\omega} = 0$. However, we should note here that if one fixes the infinite limit-size point $(2\eta - \bar{\eta})|_{L=\infty}$ to zero, the fit becomes of better quality in terms of the merit χ^2/DOF [79]. Unfortunately, in the present $D = 3$ case, we can not draw a definite conclusion on the validity of the two-exponent scaling scenario. Additional work is under way to tackle this problem at higher dimensions ($D > 3$) [80].

We turn our discussion now to the most controversial issue of the specific heat of the RFIM. The specific heat of the RFIM can be experimentally measured [16] and is, for sure, of great theoretical importance. Yet, it is well known that it is one of the most intricate thermodynamic quantities to deal with in numerical simulations, even when it comes to pure systems. For the RFIM, Monte Carlo methods at $T > 0$ have been used to estimate the value of its critical exponent α , but were restricted to rather small systems sizes and have also revealed many serious problems, i.e., severe violations of self-averaging [40,110]. A better picture emerged throughout the years from $T = 0$ computations, proposing estimates of $\alpha \approx 0$. However, even by using the same numerical techniques, but different scaling approaches, some inconsistencies have been recorded in the literature. The most prominent was that of Ref. [61], where a strongly negative value of the critical exponent α was estimated. On the other hand, experiments on random field and diluted antiferromagnetic systems suggest a clear logarithmic divergence of the specific heat [16].

The specific heat can be estimated using ground-state calculations and applying thermodynamic relations employed by Hartmann and Young [61] and Middleton and Fisher [64]. The method relies on studying the singularities in the bond-energy density E_J [111]. This bond energy density is the first

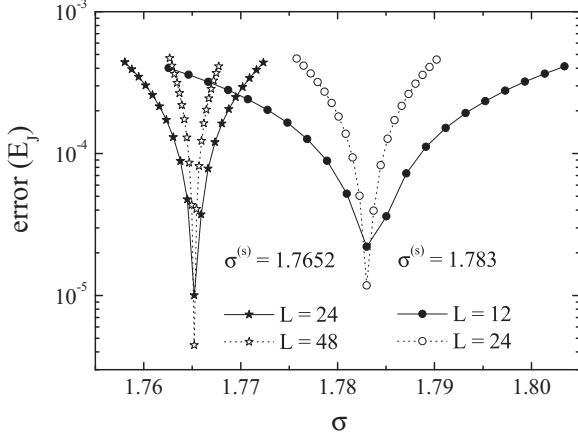


FIG. 6. Semilogarithmic illustration of statistical errors appearing in the three lattice-size variant of the quotients method. We show results for the Poissonian RFIM and $L_1 = 12$, $L_2 = 24$, and $L_3 = 48$. The values of the field strength where the simulations were performed for both pairs of system sizes are also given in the figure.

derivative $\partial E/\partial J$ of the ground-state energy with respect to the random-field strength, say σ [61,64]. The derivative of the sample averaged quantity \overline{E}_J with respect to σ then gives the second derivative with respect to σ of the total energy and thus the sample-averaged specific heat C . The singularities in C can also be studied by computing the singular part of \overline{E}_J , as \overline{E}_J is just the integral of C with respect to σ . The general finite-size scaling form assumed is that the singular part of the specific heat behaves as

$$C_s \sim L^{\alpha/\nu} \tilde{C}[(\sigma - \sigma^{(c)})L^{1/\nu}]. \quad (69)$$

Thus one may estimate α by studying the behavior of \overline{E}_J at $\sigma = \sigma^{(c)}$ [64]. The computation from the behavior of \overline{E}_J is based on integrating the above scaling equation up to $\sigma^{(c)}$, which gives a dependence of the form

$$\overline{E}_J(L, \sigma = \sigma^{(c)}) = A + BL^{(\alpha-1)/\nu}, \quad (70)$$

with A and B nonuniversal constants.

Since $\alpha - 1$ is negative, Eq. (70) is dominated by the nondivergent background A , forcing us to modify the standard phenomenological renormalization. We get rid of A by considering three lattice sizes in the following sequence: $(L_1, L_2, L_3) = (L, 2L, 4L)$. We generalize Eq. (68) by taking now the quotient of the differences $Q_O = (O_{4L} - O_{2L})/(O_{2L} - O_L)$ at the crossings of the pairs $(L, 2L)$ and $(2L, 4L)$, respectively. Applying this formula to the bond energy we obtain

$$Q_{\overline{E}_J}^{(\text{cross})} = 2^{(\alpha-1)/\nu} + \mathcal{O}(L^{-\omega}). \quad (71)$$

Of course, at variance with the standard two lattice-size phenomenological renormalization, statistical errors are significantly amplified by the reweighting extrapolation, as it can be clearly seen in Fig. 6. Hence we have preferred to carry out an independent set of simulations for parameters corresponding to the crossing points identified in the main analysis of the quotients method. Our results for the effective exponent ratio $(\alpha - 1)/\nu$ are given in Table II and their extrapolation is shown in Fig. 7. We have excluded from the fitting procedure the data of the double Gaussian distribution

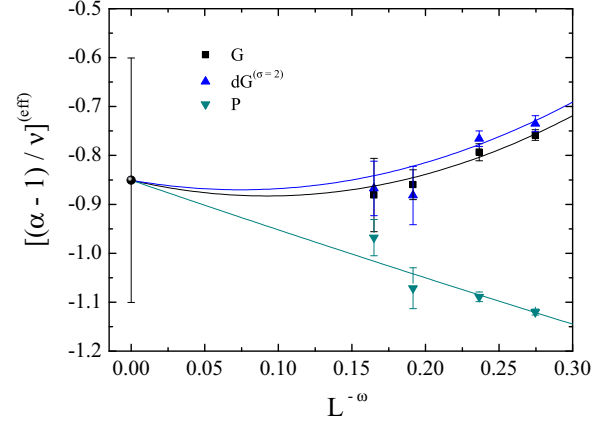


FIG. 7. Infinite limit-size extrapolation of the effective exponent ratio $(\alpha - 1)/\nu$.

with $\sigma = 1$ as their inclusion destabilized the fit. The solid lines in Fig. 7 show a joint polynomial fit, second order in $L^{-\omega}$. The extrapolated value for the exponent ratio is $(\alpha - 1)/\nu = -0.85(25)$ and is marked by the filled circle in the figure at $L^{-\omega} = 0$. Using now our estimate $\nu = 1.38(10)$ for the critical exponent of the correlation length, simple algebra (and error propagation) gives the value $\alpha = -0.16(35)$ for the critical exponent of the specific heat. Let us point out here that also Middleton and Fisher, using the scaling of the bond energy at the candidate critical field value $\sigma^{(c)} = 2.27$, proposed a value of $\alpha = -0.12(16)$ [64], compatible with our result. Although the error proposed by the latter authors is much smaller than ours, we have to note that their method implies an *a priori* knowledge of the “exact” value of the critical field. Obviously, as we have no command over this value, what is usually done is to use some candidate values of the critical field around the best known estimate and then repeat the simulations for all those candidate values. However, even in this case the results are ambiguous, as a change in the value of $\sigma^{(c)}$ by a factor of $\delta\sigma^{(c)} = 10^{-3}$ results, on an average, in a change of the order of $\delta\alpha \approx 0.04$ in the value of α [112].

Following the discussion above, our numerical studies of disordered systems are carried out near their critical points using finite samples; each sample is a particular random realization of the quenched disorder. This makes it then crucial to study the dependence of some observables with the disorder, the so-called self-averaging properties of the system. The study of these properties in disordered systems has generated in the past years a large amount of work [40,113–119], still mostly focused on the bond- and site-diluted versions of the Ising model in $D = 2$ and 3.

A typical measure of the self-averageness of a random system via a physical quantity A is given from $R_A = [\overline{(A)^2} - \overline{A}^2]/\overline{A}^2$. Aharony and Harris [113] predicted that the size evolution of $R_A(L)$ for the random system depends on whether the system is controlled by the pure or the random fixed point, i.e., $R_A(L) \propto L^{(\alpha/\nu)_{\text{pure}}}$ for pure fixed point, and $R_A(L) \propto \text{const} \neq 0$ for random fixed point, respectively, as $L \rightarrow \infty$. In the case of the random fixed point, the system has no self-averaging. On the other hand, the system exhibits weak self-averaging in the case of the pure fixed point. Clearly

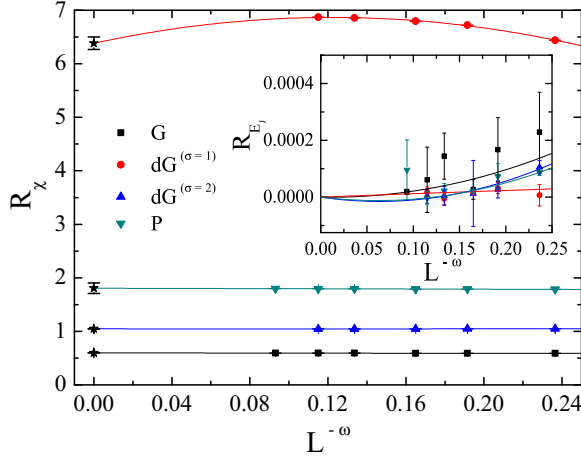


FIG. 8. Infinite limit-size extrapolation of the effective ratios R_χ (main panel) and R_{E_J} (inset).

enough, the system is expected to be self-averaging if $R_A \rightarrow 0$, as $L \rightarrow \infty$.

The RFIM is a nice candidate to check the analytical predictions of Aharony and Harris on self-averaging [113], monitoring the infinite limit-size extrapolation of R_A . In particular, we investigated here the behavior of the ratio for two observables, the connected susceptibility, R_χ , and the bond energy, R_{E_J} . In Fig. 8 we plot the effective values of the ratios R_χ and R_{E_J} in the main panel and inset, respectively, estimated at the crossing points of ξ/L as usual, for all our four distributions studied, as indicated by the different colors and symbols. In both cases, the solid lines show a joint, second-order in $L^{-\omega}$ polynomial fit, using as a lower cutoff the lattice size $L_{\min} = 16$. For the case of R_χ , the extrapolated values shown by black stars are dependent on the field distribution and are clearly nonzero, indicating violation of self-averaging.

In particular, we quote the following limiting values: $R_\chi = \{0.600(2), 6.38(11), 1.047(4), 1.81(9)\}$, for the cases of the Gaussian, double Gaussian with $\sigma = 1$, double Gaussian with $\sigma = 2$, and Poissonian distributions, respectively. The above results verify the prediction of Ref. [113], according to which the susceptibility at the critical point is not self-averaging for models where the disorder is relevant, relevant meaning that the critical exponent of the specific heat of the corresponding pure model is positive ($\alpha_{\text{pure}} > 0$) [120]. As for the self-averaging ratio for the bond energy, shown in Fig. 8, inset, it goes to zero with increasing system size, indicating self-averaging in the thermodynamic limit.

Finally, we present some computational aspects of the implemented push-relabel algorithm and its performance on the study of the RFIM. Although its generic implementation has a polynomial time bound, its actual performance depends on the order in which operations are performed and which heuristics are used to maintain auxiliary fields for the algorithm. Even within this polynomial time bound, there is a power-law critical slowing down of the push-relabel algorithm at the zero-temperature transition [53,98]. This critical slowing down is certainly reminiscent of the slowing down seen in local algorithms of statistical mechanics at finite temperature, such

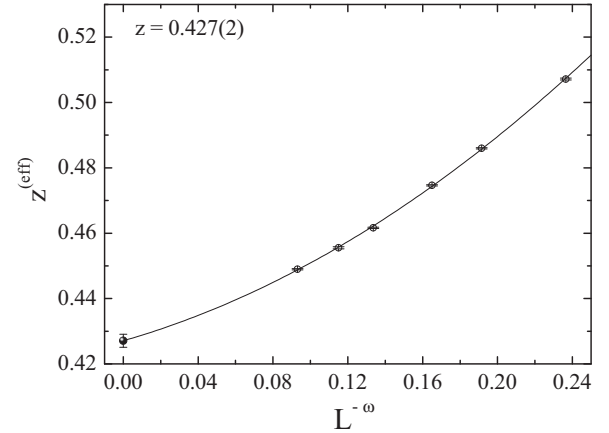


FIG. 9. Infinite limit-size extrapolation of the effective exponent z of the push-relabel algorithm.

as the Metropolis algorithm, and even for cluster algorithms. In fact, Ogielski [53] was the first to note that the push-relabel algorithm takes more time to find the ground state near the transition in three dimensions from the ferromagnetic to paramagnetic phase.

A direct way to measure the dynamics of the algorithm is to examine the dependence of the running time, measured by the number of push-relabel operations, on system size L . Such an analysis has already been performed in Ref. [121] for the Gaussian $D = 3$ RFIM and a FIFO queue implementation, as in the current paper, finding a dynamic exponent $z = 0.43(6)$, using the data collapse technique and fixing the values $\sigma^{(c)} = 2.27$ and $\nu = 1.37$ in the scaling ansatz. Here, we present a complementary analysis based on the numerical data also for Gaussian RFIM, using our scaling approach within the quotients method and without assuming prior knowledge of the critical field and correlation length exponent. Our fitting attempt is shown in Fig. 9, where the solid line is a second order in $L^{-\omega}$ polynomial for system sizes $L \geq 16$ and the obtained estimate for the dynamic critical exponent z is $0.427(2)$, very close to the estimate of Ref. [121].

IX. SUMMARY AND OUTLOOK

To summarize, we have presented in the current paper a fluctuation-dissipation approach to the study of the random-field Ising model, using as a platform the three-dimensional version of the model. We combined several efficient numerical methods, from zero-temperature optimization algorithms to generalized fluctuation-dissipation formulas and reweighting extrapolations that allowed the computation of response functions, as well as advanced finite-size scaling techniques that offered us the possibility to tackle some of the hardest open problems in the random-field literature, like the existence and role of scaling corrections and the universality principle of the model. We hope that this contribution gives a clear overview of all the technical details of our implementation, paving the way to even more sophisticated studies in the field of disordered systems. Currently, using the prescription outlined above, we are dealing with the random-field problem at higher dimensions and we expect to provide clearcut results regarding

the validity of the two-exponent scaling scenario, one of the building blocks in the scaling theory of the random-field Ising model.

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