Ensemble renormalization group for disordered systems

Maria Chiara Angelini,1 Giorgio Parisi,2 and Federico Ricci-Tersenghi2
1Dipartimento di Fisica, Università “La Sapienza”, Piazzale A. Moro 5, I-00185 Roma, Italy
2Dipartimento di Fisica, INFN – Sezione di Roma 1, CNR – IPCF UOS Roma, Università “La Sapienza”, Piazzale A. Moro 5, I-00185 Roma, Italy

(Received 12 December 2012; published 2 April 2013)

We propose and study a renormalization group transformation that can be used also for models with strong quenched disorder, like spin glasses. The method is based on a mapping between disorder distributions, chosen such as to keep some physical properties (e.g., the ratio of correlations averaged over the ensemble) invariant under the transformation. We validate this ensemble renormalization group by applying it to the hierarchical model (both the diluted ferromagnetic version and the spin glass version), finding results in agreement with Monte Carlo simulations.

DOI: 10.1103/PhysRevB.87.134201

PACS number(s): 05.10.Cc, 75.10.Nr, 64.60.ae

I. INTRODUCTION

The renormalization group (RG) is a fundamental tool in theoretical physics.1 It allows to characterize phase transitions and critical phenomena by computing critical exponents and universality classes. The real-space RG can be viewed as a decimation procedure that takes a system made of N dynamical variables and reduces it to a smaller system, in a way which preserves, or scales appropriately, some important physical observables. Such a decimation induces a RG transformation on the system couplings, and the study of such a transformation allows one to identify critical points and critical exponents.

Real-space RG transformations have been studied in great detail for homogeneous models,1 but much less is known for disordered models that contain quenched randomness in the Hamiltonian (either random fields and/or random couplings). This is especially true for strongly frustrated models, like spin glasses (SG), for which a satisfying RG transformation is still lacking. For example, for the Edwards-Anderson SG model2 on a D-dimensional lattice all the attempts to develop a field theory by performing an $\epsilon$ expansion around the upper critical dimension $D_u = 6$ have proved to be very complicated.3,4 These studies have led to the discovery of fixed points different from the mean-field (MF) ones, however, the implications of that are not completely clear. In particular the existence of replica symmetry breaking (RSB) fixed points in the non-mean-field region $D < D_u$ has been shown only very recently5 and estimates of critical exponents in $D = 3$ are still not reliable.

In this framework the development of a (semi)analytical real-space RG for disordered models would be very welcome. The outcome of such a RG transformation could be well compared with Monte Carlo (MC) simulations that provide accurate estimates of critical temperatures and critical exponents for disordered models (at a much higher computational cost).

Since the first developments of RG transformations for models with quenched disorder it was clear that one has to deal with distributions of couplings.6–8 However, the RG transformation for an entire coupling distribution is highly nontrivial. Previous attempts to develop a real-space RG for disordered systems8 focused on transformations mapping a single sample of size $N$ to a smaller system (without loss of generality we can set the size of the smaller system to $N/2$). In formulas, we can write the mapping \( \{J_{ij}\} \rightarrow \{J'_{ij}\} \) as the one solving a set of equations like

\[
\langle O_k(\{J_{ij}\}) \rangle = \langle O_k'(\{J'_{ij}\}) \rangle ,
\]

where the angular brackets are thermal averages with respect to the Gibbs-Boltzmann distribution, the primed quantities refer to the smaller system, and the number of observables $O_k$ is enough to determine the new couplings $\{J'_{ij}\}$. Given an ensemble of systems of size $N$, the above transformation can be applied to each of them to obtain an ensemble of systems of size $N/2$.

However, we believe that such a mapping is suboptimal for models with quenched disorder and a better RG transformation should consider explicitly the average over the quenched disorder (as was done, e.g., in Ref.9). What we are proposing is a mapping between the probability distributions of couplings $P(J_{ij}) \rightarrow P'(J'_{ij})$ such that the following equations hold:

\[
\langle O_k(\{J_{ij}\}) \rangle = \langle O_k'(\{J'_{ij}\}) \rangle .
\]

The overbar represents the average over the quenched disorder (i.e., the couplings in the present case). The rationale behind this choice is that in models with strong disorder (like SG) sample-to-sample fluctuations may dominate thermal ones.

It is worth noticing that standard RG transformations working on coupling distributions typically use Eq. (1). For example, the simplest RG transformation for the bond percolation problem in two dimensions ($p' = 2p^2 - p^3$) or even the more complicated one of Ref.7 can be viewed as single sample transformations, where $p$ is the bond density by which the larger samples are generated and $p'$ is the bond density of the renormalized samples (one-by-one). Instead, the new approach in Eq. (2) would prescribe to compute some physical quantity (e.g., a correlation or a probability of being connected) in the ensemble of larger systems with bond density $p$, and to repeat the same computation in the ensemble of smaller systems with several bond densities to find the best value $p'$ such that the physical observables match in the two ensembles. The new approach we are proposing resembles in some way the finite size scaling analysis which is performed...
on Monte Carlo data measured in systems of different sizes. Obviously, for models where the RG transformation is exact (like, e.g., the diamond hierarchical lattices\textsuperscript{10}) the old and the new approaches provide the same answer.

Two simple examples may help elucidate even more the limits of the RG transformation working sample by sample, Eq. (1), and thus justify the use of the one in Eq. (2) that we will call ensemble RG (ERG). In a diluted ferromagnet with long-range interactions, where the couplings are positive with probability $p$ and null with probability $1-p$, a single step of the decimation procedure induced by Eq. (1) typically generates all nonzero couplings (i.e., $p = 1$). This is clearly not very useful if one is willing to follow the RG flow in the $p-T$ plane. Moreover, in frustrated models the decimated system is typically much less frustrated than the original one: The extreme case is the transformation of a four-spins system to lower critical dimensions (and thus limits of the RG transformation working sample by sample, Eq.(1), and thus justify the use of the one in Eq.(2) that generates all nonzero couplings (i.e., $p = 1$). This tendency to reduce frustration makes the RG using Eq.(1) clearly unfit to describe SG fixed points.

In principle, our ERG scheme can be applied to any disordered system. We choose here to apply it to the hierarchical model (HM), which is a particular one-dimensional long-range model, whose Hamiltonian for $N = 2^n$ spins can be constructed iteratively in the following way:\textsuperscript{11}

$$H_n(s_1, \ldots, s_{2^n}) = H_{n-1}(s_1, \ldots, s_{2^{n-1}}) + H_{n-1}(s_{2^{n-1}+1}, \ldots, s_{2^n}) - c^n \sum_{i<j=1}^{2^n} J_{ij}s_i s_j .$$ \tag{3}

In practice $H_n$ is the sum of interactions at $n$ different levels. We have studied three versions of this model: The ferromagnet (FM), where $J_{ij} = 1$; the diluted ferromagnet (DFM), where a random fraction $1 - p$ of FM couplings are set to zero; and the SG version,\textsuperscript{12} with Gaussian couplings $P(J) \propto e^{-J^2/2}$.

There are many reasons to test a new RG transformation on the HM. First, by properly tuning the topological factor $c$ that controls how fast the couplings’ intensity decays with distance, the HM can emulate a $D$-dimensional short-range (SR) model: $c \simeq 2^{1-\frac{2}{D}}$ for DFM and $c \simeq 2^{1-\frac{2}{D}+1/2}$ for SG.\textsuperscript{13} These relations are exact around the upper critical dimensions because the long-range HM and the SR $D$-dimensional model have the same field theory at leading order. To have a phase transition at a finite temperature, the $c$ parameter must satisfy $c_L^{\text{FM}} = \frac{1}{2} < c < \frac{1}{2} = c_\infty^{\text{FM}}$ for the DFM and $c_L^{\text{SG}} = \frac{1}{2} < c < \frac{1}{2} = c_\infty^{\text{SG}}$ for the SG. The lower bound values $c_L$ correspond to lower critical dimensions (and thus $T_c = 0$), while when $c = c_\infty$ the energy is no longer extensive (thus $T_c = \infty$). For $c > 2^{-\frac{2}{D}} = c_L^{\text{FM}}$ in DFM and $c > 2^{-\frac{2}{D}} = c_L^{\text{SG}}$ in SG, the model shows mean-field critical properties (like for $D > D_U$ in the SR models). So, tuning a single parameter in the HM, we can move from the MF region to a non-MF one.

The second reason to choose the HM is that, if the system is decimated by a standard block-spin transformation, the new Hamiltonian does not contain any multispin terms (at variance to what happens on finite dimensional lattices\textsuperscript{1}). So, considering only pairwise interactions in the RG is not an approximation for the HM. Moreover, the FM version can be exactly solved in a time growing only polynomially with $N$ since the probability distribution of the magnetization

\begin{figure}[h]
\centering
\includegraphics[width=\textwidth]{erg.png}
\caption{Schematic representation of the ERG for a SG HM with $n = 3$. Variances ($\sigma_i$) and renormalized variances ($\sigma'_i$) at each level $k$ are indicated.}
\end{figure}

\begin{equation}
p_n(m) \propto e^{\frac{\mu}{n} m + \frac{\sigma^2}{2} n} \sum_{m_L,m_R} p_{n-1}(m_L) p_{n-1}(m_R) \delta_{m_L+m_R,m} , \tag{4}
\end{equation}

where $m_L$ and $m_R$ are the magnetizations of the half systems. It is worth stressing that Eq. (4) is no more valid for the SG version of the HM, which does not admit a polynomial time solution. In Ref.14, a generalization of Eq. (4) has been proposed to describe the distribution of the overlap $q$ in the SG version of the HM; however, we will not make use of this relation due to the difficulties in its treatment.

\section{II. ENSEMBLE RENORMALIZATION GROUP FOR HM}

We describe now in detail how to apply the ERG to the HM. We assume couplings to remain independent during the RG, but we allow couplings to have a different probability distribution $P_k(J)$ (or a different value $J_k$ in the FM version) at each level $k \in \{1, 2, \ldots, n\}$: In the original HM all couplings have the same probability law, but we have seen that the RG iteration produces different couplings at different levels. In the analyzed versions, each coupling distribution is parametrized by $K \in \{1, 2\}$ numbers (that is, the mean for FM, the variance for SG, the fraction of nonzero couplings, and the mean for DFM), otherwise the search for a solution to Eq. (2) would become too difficult. We start from an ensemble of systems with $n$ levels that we want to reduce to an “equivalent” ensemble of smaller systems of $n - 1$ levels. The whole procedure is described in Fig. 1 for a SG HM with $n = 3$.

(1) First we compute $(n - 1)K$ observables $\langle O_j \rangle_j, j \in \{K+1, \ldots, Kn\}$ in the larger systems extracted from the original coupling distribution (in the FM the overbar can be omitted).

(2) Then we identify the new distributions of couplings $P'_k(J'), k \in \{1, \ldots, n - 1\}$ for the ensemble of smaller systems, i.e., we determine the new $(n - 1)K$ parameters of the $P'$ distributions by requiring that $\langle O'_i \rangle'_{P'} = \langle O_{i+K} \rangle_P$ for any $i \in \{1, 2, \ldots, (n - 1)K\}$. 

\begin{verbatim}
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\end{verbatim}
Finally we build a new ensemble of systems of the original size. They are constructed joining with random couplings extracted from the original distribution $P_0(J)$ two smaller systems with couplings extracted from $P'_k(J')$, $k \in \{1, 2, \ldots, (n - 1)\}$ found at step 2.

Primed quantities refer to the smaller systems. In this case the parameters to be determined are the variances of the Gaussian distributions at each level. The first two steps are the true renormalization steps, while the last step is required to obtain a final system size that will allow us to iterate the method until convergence. In the disordered versions, the assumption that the renormalized coupling distributions are of the same type as the original ones is an approximation. However, it can be easily improved adding extra terms in the distributions. The thermal averages are computed exactly: This is easy to do in the FM, thanks to Eq. (4), while in the DFM and in the SG we do it by exhaustive enumeration, thus limiting us to a small number of levels in the disordered cases. The average over the disorder is not exact but taken over $\sim 10^5$ samples. Step 2 is actually accomplished by minimizing $\sum_k (\langle O_k \rangle - \langle O_k \rangle)^2$, and we have checked that the reached minimum is always very close to zero. Since the couplings’ distributions are different at each level, we do not see any better option than extracting the new couplings at level $n$ in step 3 from the original distribution $P_0(J)$.

The FM version of the HM, for which exact RG equations can be written, is a benchmark for our numerical implementation of the ERG. For $c = 2^{-1}$, which corresponds to $D \simeq 3$, the critical temperature is $T_c = 0.848154717$ (Ref. 16), and the critical exponents are $\eta = 4/3$, $\gamma = 1.299140730159(1)$ (Ref. 17), leading to $\nu = 1.948711095$ using the scaling relations. To compare this critical exponent $\nu$ with the one for a SR three-dimensional (3D) Ising FM, it can be shown that the following equation should be used with a proper $c$-$D$ relation:

$$\nu_L(c) = D\nu_{SR}(D).$$

In the present case $\nu_{SR}(c = 2^{-1/3})/3 = 0.649570365$ is close to $\nu_{SR}(3) = 0.6301(4)$ (Ref. 18).

The observables $O_k$ that we use in the RG equations are the correlation of the magnetization at level $k + 1$, normalized by those at level $k$, with $k \in \{1, \ldots, n - 1\}$:

$$\langle O_k \rangle = \frac{\langle m_{L_k} m_{R_k} \rangle}{\langle m_{L_k} m_{R_k} \rangle}.$$

The denominator is needed to reduce finite size effects and to ensure that a solution to the RG equations always exists (this is not true in general for other observables, like the magnetization).

Applying the previously described procedure, the flux of couplings and correlations can be followed, see Fig. 2. If $T \simeq T_c$, the renormalized couplings stay for a while close to the critical fixed point (FP) and then go towards the high temperature (HT) FP if $T > T_c$ or the low temperature (LT) FP if $T < T_c$. We estimate the critical temperature as the temperature dividing the flows towards the two different FP. Please note that the HT and LT fixed points are not characterized by the usual $J = 0$ and $J = \infty$ coupling values: The reason for this is that in step 3 of our procedure we used a new coupling of the original intensity. Nonetheless the couplings’ flows clearly differentiate HT and LT behaviors.

To extract critical exponents from the RG equations, we focus on the early regime, when the coupling flows leave the critical FP. From the Wilson relation an equation relating the renormalized coupling after $x$ RG steps can be obtained:

$$\frac{J_1(x)}{T_1} - \frac{J_2(x)}{T_2} = \left(\frac{1}{T_1} - \frac{1}{T_2}\right)b^x,$$

where $b = 2$ is the scaling factor in our case. Thus, the $\nu$ exponent can be estimated from a fit like the one in Fig. 3. The values obtained for the critical temperature and the critical exponents in $D \simeq 3$ are well comparable with the known ones: $T_c$ extrapolates to 0.8478(1) in the large $n$ limit and the measured $\nu$ exponent is 2.076(6) for $n = 13$ [although the extrapolation of $\nu$ to the $n \to \infty$ limit is difficult due to strong finite size effects, e.g., $\nu = 2.79(12)$ for $n = 4$]. Moreover, we have checked that our numerical RG recovers the right bounds on $c$, namely, $T_c \to 0$ for $c_L = 1/4$ and $T_c \to \infty$ for $c_\infty = 1/2$.

We consider now the DFM. The Hamiltonian of the model is always the one in Eq. (3), but the couplings at level $k$ are independent random variables extracted from the distribution $P_k(J) = p_k \delta(J - J_k) + (1 - p_k) \delta(J)$.

At the beginning $p_k = p$ and $J_k = 1$ for any $k$, while under the RG they will differentiate. The number of parameters to be determined in the ERG is $2(n - 1)$, and we use the following observables, with $k \in \{1, \ldots, n - 1\}$, to find them:

$$\left(\frac{\langle m_{L_k} m_{R_k} \rangle}{\langle m_{L_k} m_{R_k} \rangle}\right)^2.$$
phase diagram found with the ERG is confirmed by a set of MC simulations (explained later in the text) whose $T_c$ estimates are also shown in Fig. 4. The only disappointment about this phase diagram is that we do not find an unstable FP along the critical line as expected for a $D = 3$ SR model. However, this can be explained by noticing that the $\alpha$ exponent of this model is very small, $\alpha = 0.051288905$, and so the crossover from the pure behavior can be extremely long.

Finally we study the SG version. The Hamiltonian is always the one in Eq. (3), and the couplings at level $k$ are distributed with a Gaussian law of zero mean and variance $\sigma^2_k$ (at the beginning $\sigma^2_k = 1$ for any $k$). The assumptions that the renormalized couplings are independent and normally distributed could be released by adding extra terms in the coupling distributions, but we leave these generalizations for future work. In the SG case the observables used to fix the $n - 1$ variances are normalized SG correlations at different levels:

$$\langle O_n \rangle = \frac{\sum_{i \in L_n, j \in K_n} \langle s_i s_j \rangle^2}{\sqrt{\sum_{i \in L_n} \langle s_i \rangle^2 \sum_{j \in K_n} \langle s_j \rangle^2}}. $$

Because of the computational costs we use $n \leq 4$, so the early regime leaving the critical FP is rather short, and the stationary regime is soon reached (with respect to the FM case). This effect is also enhanced by the disorder: Indeed, even exactly at criticality, the SG ensemble contains many samples which are not critical, and the couplings of these samples flow away from the critical values very quickly. So, it seems unavoidable that disorder increases the instability of critical FP and consequently the uncertainty on the estimates of the critical exponents. Nonetheless we can distinguish two temperature regions separated by a critical temperature $T_{SG}$ [see Fig. 5 for an effective dimension $D \simeq 3$ and $T_{SG} = 0.58(1)$], such that above $T_{SG}$ the correlations and couplings decay towards zero, while below $T_{SG}$ the correlations and couplings variances grow, suggesting that the system is in a SG phase. In Fig. 5 we have plotted only the couplings’ and correlations’ variances measured at the lowest level $k = 1$, but (as in the FM, see Fig. 2) the renormalized variances at the other levels are related to those at $k = 1$: For example, $\sigma_2 > \sigma_1$ if $T > T_{SG}$ and $\sigma_2 < \sigma_1$ if $T < T_{SG}$ (remember that the parameters at the lowest level are those which are less influenced by the coupling of the original intensity at level $n$ because they have been renormalized more times).

Also in the SG case we are able to estimate the $\nu$ exponent from the flux of the couplings at early times. The procedure used is the same as that in the FM case and typical fits are shown in Fig. 6 for $D \simeq 8.2$ in the mean-field region and for $D \simeq 3$ below the upper critical dimension. We obtain $\nu = 4.15(10)$ in $D \simeq 8.2$ and $\nu = 4.34(6)$ in $D \simeq 3$. In Fig. 7 we report (with the label HM ERG) the estimates of $\nu$ as a function of the exponent $\sigma$ defined as $c = 2^{-(1+\sigma)/2}$: We see that in the mean-field region ($\sigma < 1/4$) the results are compatible with the expected mean-field behavior ($\nu = D/2$ that corresponds to $\nu = 1/3$) and, more interestingly, the critical exponent $\nu$ has a minimum around the upper critical dimension $D_U = 6$, $\nu_U = 1/3$, as confirmed by our MC simulations (label HM MC in Fig. 7) and those in Ref. 12. This minimum was not observed in previous RG studies, while it is present in SR models: In Fig. 7 we report the $D_{VSR}$ estimates for the EA model in $D = 3, 4, 5$ (Refs. 15, 20, and 21) following the relation introduced previously: $D = \frac{5}{2}$. The same nonmonotonic behavior for the $\nu$ exponent has been also seen in a 1d SG model with long-range (LR) power-law decaying interactions ($J_{ij} \propto |r_{ij}|^{-(1+\sigma)/2}$) (Refs. 15 and 22), also shown in Fig. 7.
we study the dimensionless Binder parameter, and we also show results from the ε expansion and from MC simulations of the HM. For comparison, we use the parallel tempering algorithm, running simultaneously at 20 different temperatures. Two replicas have been in parallel to measure the overlap between them. Equilibration has been checked by the standard method of observing the convergence of the measured observables (e.g., energy and overlap moments) to their asymptotic values. We have found that the equilibration time is \( \tau \simeq 10^3 - 10^6 \) MC steps for the largest sizes at smaller temperatures and we have acquired data for \( 5 \cdot \tau \) MC steps. For the model with Gaussian couplings we have used \( n = 6, 7, 8, 9, 10 \), while for the one with \( \pm J \) couplings, which is easier to simulate, \( n = 6, 7, 8, 9, 10, 11 \). Averages were performed over 400 samples for the larger systems up to 2000 samples for the smaller ones.

We have run also standard MC simulations (with nonparallel tempering) for the DFM at several bond concentrations (\( \rho = 0.3, 0.5, 0.7, 0.8 \)).

Critical temperatures have been estimated from the crossing points of the scale-invariant observables: These crossing points should approach the critical temperature for large sizes. In particular we study the dimensionless Binder parameter, defined as \( B = \frac{1}{2} \left[ 3 - \frac{\langle m^4 \rangle}{\langle m^2 \rangle^2} \right] \) for the DFM and \( B = \frac{1}{2} \left[ 3 - \frac{\langle q^4 \rangle}{\langle q^2 \rangle^2} \right] \) for the SG. We can construct also another scale-invariant observable from the susceptibilities, \( \chi_{\text{FM}} = m_2 L \) and \( \chi_{\text{SG}} = q_2 L \), knowing analytically its dimension \( 2 - \eta \). In fact the \( \eta \) exponent is not renormalized in LR systems. Defining \( c = 2 - (1 + \sigma) \) for the FM and \( c = 2 - (1 + \sigma)/2 \) for the SG, \( \eta = 2 - \sigma \) in the whole region \( c_L < c < c_\infty \). Thus if we divide \( \chi \) by its dimension, the resulting \( m_2 L^{-1+\eta} \) or \( q_2 L^{-1+\eta} \) should remain finite at the critical temperature. We can apply the same argument to the fourth-moment susceptibility, obtaining another scale-invariant observable \( m_4 L^{-2+2\eta} \) for DFM and \( q_4 L^{-2+2\eta} \) for SG.

In Fig. 8, we plot the three scale-invariant observables, \( B, q_2 L^{-1+\eta} \) and \( q_4 L^{-2+2\eta} \) for different sizes \( L = 2^n \) as a function of the temperature for a SG HM with \( c = 2^{-\frac{2}{3}} \) and a Gaussian distribution of the couplings. The crosses of the curves should approach the critical temperature, which we estimate to be \( T_c = 0.55(1) \). Unfortunately the data do not allow us to estimate the correction to scaling exponent \( \omega \). Nonetheless we can say that the Binder parameter is the observable that has the largest finite size effects. The critical temperature is compatible with the one obtained from the ERG \( T_c = 0.58(1) \). Please consider that the estimate from ERG is obtained with a small number of levels \( n = 4 \) (and in a much faster time). In Fig. 4 we report the critical temperatures of the DFM obtained by an analysis very similar to the one just described.

To measure the critical exponent \( \nu \) we have used the more accurate data for the model with \( \pm J \) interactions, which should be in the same universality class. We use the same observables

![Figure 8](https://example.com/figure8.png)

**FIG. 8.** (Color online) Scale-invariant observables, \( B, q_2 L^{-1+\eta} \) and \( q_4 L^{-2+2\eta} \) for different sizes \( L = 2^n \) as a function of the temperature for a SG HM with \( c = 2^{-\frac{2}{3}} \) and a Gaussian distribution of the couplings. The crosses of the curves should approach the critical temperature, which we estimate to be \( T_c = 0.55(1) \).
The method has been applied to the hierarchical model and is able to find a SG transition also for effective dimension $D \simeq 3$ in the non-mean-field region. The reliability of the method has been tested comparing the values of critical temperatures and critical exponents with those obtained in MC simulations. The agreement for the critical temperatures is optimal for all the versions we have studied and the ERG method allows to compute them in a time much smaller than MC simulations. The critical exponents obtained with the ERG method are more affected by finite size effects than critical temperatures. In fact the error in the ERG estimate for the SG exponents is around 20%. However, such a large error is not unexpected since we find it also in the ferromagnetic case, where the correct exponent is recovered only in the $n \to \infty$ limit.

The ERG method is able to reproduce the correct behavior of the $\nu$ exponent, which shows a minimum at the upper critical value of $\nu = c_1$ (or $\sigma = \sigma_1$). Such a behavior is predicted by the $\epsilon$ expansion of the SG version and found in our MC simulations. This accurate prediction by the ERG makes us confident that the method is reliable. The present results by the ERG solve an apparent inconsistency problem between the $\epsilon$ expansion and another real-space RG approach recently proposed by the authors of Ref. 8, where the $\nu D$ exponent was found to decrease linearly with $D$, with no minimum at all around the upper critical dimension. In this work we have shown that such a minimum in $\nu D$ exists if a better RG transformation is used. In particular we have shown that the right way to do renormalization when disorder is present is to consider RG on an ensemble of systems and not on single samples.

This statement can be confirmed also from a comparison with the method of analysis used in MC simulations. When exponents are determined using finite size scaling (FSS), we look at the crossing point of some dimensionless quantities (like the Binder parameter) averaged over disorder for different sizes. Indeed FSS is an approximate RG where only a parameter, the temperature, is varied. In this perspective, our RG is a generalization of FSS, where we vary more parameters, and for this reason we can look at more observables. However, like in FSS, the useful observables are those averaged over the disorder.

ACKNOWLEDGMENTS

This research has received financial support from the European Research Council (ERC) through grant agreement No. 247328 and from the Italian Research Minister through the FIRB project No. RBFR086NN1.
13 We are aware that more accurate $c$-$D$ relations can be defined, as discussed exhaustively by the authors of Ref. 15. However, the scope of our comparison with short-range models in the present work is only qualitative and does not pretend to be quantitative. For this reason the simplest relation is sufficient for our purpose.