Reentrant random quantum Ising antiferromagnet

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We consider the quantum Ising chain with uniformly distributed random antiferromagnetic couplings (1 ≲ J ≲ 2) and uniformly distributed random transverse fields (Γ0 ≲ Γ ≲ 2Γ0) in the presence of a homogeneous longitudinal field, h. Using different numerical techniques (density-matrix renormalization group, combinatorial optimization, and strong disorder renormalization group methods) we explore the phase diagram, which consists of an ordered and a disordered phase. At one end of the transition line (h = 0, Γ0 = 1) there is an infinite disorder quantum fixed point, while at the other end (h = 2, Γ0 = 0) there is a classical random first-order transition point. Close to this fixed point, for h > 2 and Γ0 > 0 there is a reentrant ordered phase, which is the result of quantum fluctuations by means of an order through disorder phenomenon.

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I. INTRODUCTION

Quantum phase transitions are among the fundamental problems of modern physics, the properties of which are studied in different disciplines: solid state physics, quantum field theory, quantum information, and statistical mechanics [1]. Quantum phase transitions take place at T = 0 temperature and these are indicated by singularities in the ground-state expectation values of some observables by varying a control parameter, such as the strength of a transverse field. One basic question in this field of research is how quenched disorder influences the properties of quantum phases and the singularities associated with the quantum phase transitions. This latter problem is theoretically very challenging, since the corresponding quantum state is the result of an interplay between quantum and disorder fluctuations, strong correlations, and frustration.

Many results in this field are known on the random transverse-field Ising chain with short-range interactions. It was Fisher [2], who used a strong disorder renormalization group (SDRG) method [3] and obtained several asymptotically exact results. The phase transition is shown to be controlled by a so-called infinite disorder fixed point [4] (IDFP), at which the distribution of the parameters (couplings and random transverse fields) increase without limit during renormalization. Outside the quantum critical point dynamical observables (susceptibility, autocorrelation functions, etc.) are still singular, due to the presence of strong Griffiths singularities, which are the result of rare regions in the disordered samples [5,6].

IDFP properties are found also for random Heisenberg chains [7–9] as well as in the random singlet phases of SU(2)k anyonic chains [10]. In higher-dimensional systems with discrete symmetry the presence of IDFPs has been demonstrated by numerical application of the SDRG method [11–13], as well as through quantum Monte Carlo (MC) simulations [14,15]. On the contrary the phase transition of the random transverse-field Ising model with long-range interactions, the strength of which decays as a power with the distance, is shown to have a conventional random quantum fixed point [16,17]. The RG flow in this case is similar to that of disordered bosons [18,19].

A quantum system is often described by several parameters and in their space the phase transition takes place at a line or at a (higher-dimensional) surface. The phase transition of the clean system in this case is generally governed by a few fixed points, or in exceptional cases by a line of fixed points. In some cases (such as in the random quantum Ashkin-Teller chain [20,21]) the clean fixed points are found to turn to the same type of IDFPs, mainly due to symmetry reasons. However, there are no detailed studies in the general cases, when the different clean fixed points transform differently due to disorder.

In this paper we are going to study such a more complex system, the antiferromagnetic Ising chain in a mixed transverse and longitudinal field. The clean model has two fixed points. The quantum fixed point governs the critical behavior at any nonzero transverse field, while at zero transverse field there is a classical fixed point, which describes a first-order transition. In the present study we are going to keep the longitudinal fields nonrandom, but at the same time the couplings and the transverse fields disordered and investigate the phase-transition properties of this random system. We explore
the phase diagram numerically by the density-matrix renormalization group (DMRG) method for finite values of the transverse field, while in the zero transverse-field limit, when the system is classical we use combinatorial optimization methods to find the true ground-state configuration. We also use approximate SDRG calculations, as well as perturbation calculations, to see the stability of the random fixed points.

The rest of the paper is organized in the following way. In Sec. II the model is introduced and its properties are described for nonrandom parameters. The disordered model is studied in Sec. III. The two end points of the transition line (zero longitudinal field and zero transverse field) are described asymptotically exactly, while the complete phase diagram is explored numerically through DMRG and SDRG methods. Our results are discussed in Sec. IV and some detailed calculations are presented in the Appendices.

II. THE MODEL

We consider the antiferromagnetic Ising chain with mixed transverse and longitudinal fields. The longitudinal fields are constant, $h \geq 0$, while the nearest-neighbor couplings, $J_i > 0$, and the transverse fields, $\Gamma_i > 0$, may be random, so that the Hamiltonian is defined as

$$\hat{H} = \sum_{i=1}^{L} J_i \sigma_i^z \sigma_{i+1}^z - \sum_{i=1}^{L} \Gamma_i \sigma_i^x - h \sum_{i=1}^{L} \sigma_i^z \tag{1}$$

in terms of the $\sigma_{i}^{x,z}$ Pauli matrices at site $i$. For finite chains we have $L = 4L$ lattice sites and periodic boundary conditions (PBCs).

To the best of our knowledge (the clean version of) this model was studied in the second part of the 1980s. A more general model, with an $m$-spin product interaction term, i.e., $\prod_{j=0}^{m-1} \sigma_{i+1}^{x_j}$ instead of $\sigma_{i}^{z} \sigma_{i+1}^{z}$, has been introduced and studied for $m = 3$ by Penson et al. [22] and for $h > 0$ a phase transition of the three-state Potts universality class has been found. Soon after this model is considered in the vicinity of the classical limit $\Gamma \to 0$ and $h \to m$ (which is called quantum hard rods) and has been studied by finite-size exact diagonalization [23]. For dimers with $m = 2$ the transition is shown to be in the quantum Ising universality class, while for $m = 3$ the transition is in the three-state Potts universality class, which has been shown more precisely by MC simulations on the classical version of the model [24,25]. Finally a detailed study of the clean model in Eq. (1) has been performed in [26] after a preliminary investigation in [27].

Clean model

First we consider the clean model with $J_i = 1$ and magnetic fields $\Gamma_i = \Gamma$ and $h$, and the phase diagram is shown in Fig. 1 which has been calculated by the DMRG method. This phase diagram agrees with the previous calculations obtained by DMRG [26], by experimental realization in an optical lattice [28], by quantum MC [29], and by the fidelity susceptibility method [30]. The phase diagram in Fig. 1 for finite $\Gamma > 0$ contains an ordered antiferromagnetic phase (AFM) and a paramagnetic phase. The transition between them is controlled by a quantum Ising fixed point at $(\Gamma = 1, h = 0)$, the properties of which are known exactly [31].

![FIG. 1. The zero-temperature phase diagram of the clean antiferromagnetic Ising chain with $J = 1$ in transverse ($\Gamma$) and longitudinal ($h$) magnetic fields calculated by the DMRG method. The transition between a quantum antiferromagnetic (AFM) phase and a quantum paramagnetic (PM) phase is controlled by the fixed point of the transverse Ising model (TIM) at $(\Gamma = 1, h = 0)$. The classical multicritical point (CMP) at $(\Gamma = 0, h = 2)$ corresponds to a first-order transition between an AFM phase and a ferromagnetic (FM) phase in the absence of quantum fluctuations. Near the classical multicritical point the phase boundary is linear [23]: $\Gamma^c \approx 1.526492(2 - h)$.](image)

The transition line ends at $(\Gamma = 0, h = 2)$, where there is a classical multicritical point (CMP). In the limiting case $\Gamma \to 0$ and $h \to 2$ the system reduces to the quantum hard dimer model, having the transition at $[23]$ $\Gamma = 1.526492(2 - h)$. This is to be compared with direct calculations $\Gamma \approx 1.5(2 - h)$ in [26] and $\Gamma \approx 1.4(2 - h)$ in [29]. The quantum Ising nature of the transition in the hard dimer limit has been performed with large numerical precision, as well as the conformal properties have been determined. At $\Gamma = 0$ the system is classical and there are no quantum fluctuations. Here the transition takes place at the CMP which is of first order between the AFM phase and a ferromagnetic (FM) phase. At the CMP the ground state is infinitely degenerate, and the entropy per site is finite [32].

III. DISORDERED MODEL

Here we consider random variables in Eq. (1), so that the antiferromagnetic couplings and the transverse fields are independent random numbers, which are taken from some distributions. In the numerical calculations we used boxlike distributions:

$$\pi_1(J) = \begin{cases} 1 & \text{for } J_0 < J \leq J_0 + 1, \\ 0 & \text{otherwise}. \end{cases}$$

$$\pi_2(\Gamma) = \begin{cases} 1/\Gamma_0 & \text{for } J_0\Gamma_0 < \Gamma \leq (J_0 + 1)\Gamma_0, \\ 0 & \text{otherwise}. \end{cases} \tag{2}$$

In the following we argue that the phase diagram of the random system is different, if the smallest coupling is $J_{\min} = J_0 > 0$ (when there is an extended ordered region in $h$) or $J_{\min} = J_0 = 0$ (when the ordered region is restricted to $h = 0$). Indeed, in the classical limit with $\Gamma_0 = 0$ the ground state is
strictly AFM ordered, if \( h < 2J_{\min} \) and for \( h > 2J_{\min} \) domain-wall excitations destroy the AFM order (see more details in Sec. III B). In the following we shall investigate the region with \( J_0 > 0 \) and in the numerical work we keep \( J_0 = 1 \). We note that the case \( J_0 = 0 \) has been studied earlier by Lin et al. [29], so that we are going to study an unexplored system.

Now let us have a look at the phase diagram of the clean system in Fig. 1 having two fixed points, which are located at the two ends of the phase-transition line. In the following we study their stability with respect to disorder in Eq. (2). The TIM fixed point at \( h = 0 \) is transformed to the random transverse Ising model (RTIM), for which several asymptotically exact results are known through SDRG calculations, which are shortly collected in Sec. III A. The other fixed point, the classical multicritical point at \( \Gamma = 0 \), is transformed to a random classical multicritical point (RCMP), the properties of which are studied in Sec. III B.

### A. RTIM at \( h = 0 \)

For zero longitudinal field, \( h = 0 \), the model is equivalent to the random transverse Ising model, for which many asymptotically exact results are known due to SDRG calculations [2], which we shortly recapitulate here. The critical point is located at

\[
\ln J = \ln \Gamma ,
\]

where \( \bar{x} \) stands for the average of the variable \( x \) over disorder, thus with the distribution in Eq. (2) we have \( \Gamma h(h = 0) = 1 \).

The energy scale in the system is the excitation energy (smallest gap) \( \varepsilon \) and its relation with the length scale, \( L \), at the critical point is given by

\[
\varepsilon \sim L^\psi
\]

with a critical exponent \( \psi = 1/2 \).

The spin-spin correlations are defined as

\[
\mathcal{C}(r) = \langle \sigma_i^x \sigma_{r+i}^x \rangle \tag{5}
\]

and their average decay is given as a power of \( r \) at the critical point,

\[
\mathcal{C}(r) \sim \frac{1}{r^{2-\phi}},
\]

where \( \phi = (1 + \sqrt{5})/2 \) is the golden mean.

The deviation from criticality is parametrized by [2]

\[
\delta = \frac{\ln \Gamma - \ln \bar{J}}{\text{var}(\ln h) + \text{var}(\ln \bar{J})} ,
\]

where \( \text{var}(x) \) stands for the variance of the random variable \( x \).

In the disordered phase \( \delta > 0 \), the average correlations decay exponentially with the true correlation length \( \xi \sim 1/\delta^2 \), implying the correlation length exponent \( \nu = 2 \) for the random chain.

Outside the critical point the relation between the energy and the length scale is given by

\[
\varepsilon \sim L^{-\zeta} ,
\]

where the dynamical exponent, \( \zeta \), is given by the positive root of the equation:

\[
\left( \frac{J}{\Gamma} \right)^{\zeta} = 1 ,
\]

which is an exact expression in the entire Griffiths region [5,33,34]. In the vicinity of the critical point, the dynamical exponent to the leading order is given by [2,34]

\[
\zeta \approx \frac{1}{2|\delta|} \quad |\delta| \ll 1 .
\]

### B. The classical limit: \( \Gamma_0 = 0 \)

To explore the ground state of the system in the classical limit let us first sort the random couplings in increasing order, separately at odd \( J_i < J_i < J_i < J_i < J_i < J_i < J_i < J_i < J_i < J_i < J_i \) and even \( 1 < J_1 < J_1 < J_1 < J_1 < J_1 < J_1 < J_1 < J_1 < J_1 < J_1 < J_1 \) positions and let us increase gradually the strength of the longitudinal field from \( h = 0 \). For \( h < J_i + J_i \), the ground state is fully antiferromagnetic, since the longitudinal field is too weak to create a turned domain. This is illustrated in the first row of Fig. 2. For \( h > J_i + J_i \), however, it is energetically favorable to create domain walls at \( i_1 \) and \( i_2 \) and thus turn the spins in the domain between \( i_1 \) and \( i_2 \). At the boundary of the domains the spins are in \( \uparrow \uparrow \) positions. This is illustrated in the second row of Fig. 2. By increasing the value of \( h \) further more and more domains and thus domain walls are created and consequently the FM order monotonously increases. Passing \( h = 4 \) the system is fully FM ordered.

Now let us concentrate on the properties of the system close to \( h = 2 \) where the AFM order is lost. The AFM long-range order in the system is characterized by the average correlation function in Eq. (5), and we choose the two points of reference to be separated by the maximal distance, \( r = L/2 - 1 \). We have calculated \( C(L/2 - 1, \bar{h}) \) numerically by an algorithm, which is described in Appendix A. We obtained

**FIG. 2.** Illustration of the AFM ground state for \( h < J_i + J_i \) (first row) and the ground state with two AFM domains for \( h > (J_i + J_i) \) (second row). Here \( J_i \) (\( J_i \)) is the smallest random coupling at odd (even) positions. The boundaries of the domains are denoted by vertical lines, at \( i_1 \) and \( i_2 \), which are in different parity positions.
FIG. 3. The average correlation function as a function of $h$ for different sizes of the system. From right to left, $L = 2^5, 2^6, 2^7, \ldots, 2^{17}$. The effective transition point, $h^*_L$, is defined as the position of the zero point. Inset: scaling plot in terms of $u = (h - 2)L/2$. The results for $L = 2^7, 2^8, 2^9, \ldots, 2^{17}$ are indistinguishable. The red line represents the theoretical estimate obtained in the case when the ground state contains only one or two domains.

FIG. 4. The average correlation function at $\Gamma_0 = 0.2$ as a function of $h$ for different finite systems. The position of its zero point is used to define effective, size-dependent transition points. These are depicted in Fig. 5 for different values of $\Gamma_0$.

The complete phase diagram of the model has been studied numerically by the DMRG method. In this investigation we used finite samples of length $L = 4, 8, 16, 32, 64$ and their ground state and the first few excited states are calculated. Here the original version of the finite system DMRG scheme was utilized with PBC [39,40]. For the correlations it was systematically and carefully checked that their average value is independent of the basis size $m$ within the error bars. The accuracy of the ground-state energy calculations was in the range of $10^{-6} - 10^{-8}$ and this was in full agreement with the truncation error, the largest basis size being $m = 100-200$ for the different systems.

Averages are performed typically over a few ten thousands of independent samples in which the random couplings are obtained by the transformation $\tilde{J}_i = J_i - J + 3/2$, where $J = \sum_i J_i$ is the average value of the original set of couplings, $1 < J_i < 2, i = 1, 2, \ldots, L$. Evidently the transformed set of couplings, $\tilde{J}_i$, have the same average value, $3/2$, for all random samples. We use a similar rule to define the transformed set of random transverse fields. In the thermodynamical limit the original and the transformed ensembles lead to identical averages. For finite systems, however, the transformed sets usually have smaller sample to sample fluctuations.

We have studied the behavior of the average correlation function, $\langle C(L/2 - 1, h, \Gamma_0) \rangle$, which we have considered also in the classical limit in Sec. III B. We have checked that the DMRG results at $\Gamma_0 = 0$ agree with those calculated previously by combinatorial optimization methods. We remark one that at $\Gamma_0 = 0$ an effective, size-dependent transition point, $h^*_L(0)$, has been defined through the position of the zero value of the average correlation function and we use the same criterion in the quantum regime, for $\Gamma_0 > 0$, too.

As shown in the inset of Fig. 3, this function describes very well the average correlation function, even for not too small values of $u$. In the thermodynamic limit the transition takes place at $h = 2$, where the average correlation function exhibits a jump: $\lim_{m \rightarrow 2^\infty} \lim_{m \rightarrow 1} C(L/2 - 1, h) = 1$ and $\lim_{m \rightarrow 2^\infty} \lim_{m \rightarrow 1} C(L/2 - 1, h) = 0$, thus we have a random first-order transition. We should note, however, that for $h > h^* = 2$ there is a divergent length scale, $\xi \sim (h - h^*)^{-1}$, which measures the typical size of the AFM domains in the system. In this respect the transition is of mixed order [38].
This is illustrated in Fig. 4, where $C(L/2 - 1, h, \Gamma_0)$ is shown for $\Gamma_0 = 0.2$ as a function of $h$ and for different values of $L$. The effective, size-dependent transition points calculated in this way are presented in Fig. 5. One can notice in this figure, that close to the classical limit the effective transition points start to increase with $\Gamma_0 > 0$, have their maximum value around $\Gamma_0 = 0.2$, and then monotonously decrease as $\Gamma_0$ increases further. Before performing a more detailed analysis we note that at $\Gamma_0$ we know the true transition point, $h^t(0) = 2.0$, which is formally the extrapolated value of the series $h^t_\infty(0)$. We have checked that in the regime $L \leq 64$ there are considerable corrections to scaling contributions and similar behavior is expected to happen by extrapolating the data for $\Gamma_0 > 0$, too. Therefore in the vicinity of the classical limit we analyze the difference, $\Delta h^t_\infty(\Gamma_0) = h^t(\Gamma_0) - h^t_\infty(0)$, which is plotted in the inset of Fig. 5. For smaller sizes, $L = 8$ and 16, there are large corrections, in particular for small values of $\Gamma_0$. At the larger sizes ($L = 32$ and 64), however, the differences, $\Delta h^t_\infty(\Gamma_0)$, are stable, and these are positive for $0 < \Gamma_0 \leq 0.2$. We expect that this trend remains in the thermodynamic limit, too.

At the other end of the phase-transition line, close to the RTIM fixed point at $h = 0$, the analysis of the average correlation function leads to less accurate results. In this regime we estimated the location of the transition line in the following way. We calculated the distribution of the excitation gaps at finite chains and compared those with the same type of distributions at the RTIM. Having the gap distributions at $h > 0$ for different values of $\Gamma_0$ the transition point is identified, with that value of $\Gamma_0$, where the (small-gap part of the) gap distributions were closer to that at the RTIM at the same size. We illustrate this procedure in the lower inset of Fig. 6. The distributions are found very similar to that in the RTIM, thus we conclude that for $h > 0$ there is infinite disorder scaling. Indeed the scaling behavior of the small gaps is well described by the relation in Eq. (6). For $h = 0$ there is infinite disorder scaling. The scaling behavior of the small gaps is well described by the relation in Eq. (6) and the scaling exponent is (very close to) $\psi = 1/2$, as illustrated in the upper inset of Fig. 6.

In the next step we calculate the critical average correlation functions, where the transition points have been estimated in the previous paragraph from the scaling of the gaps. From a log-log plot of $C(L/2 - 1, \Gamma_0^c)$ vs $L$ in Fig. 6 one can notice an asymptotic linear trend, which corresponds to a power-law dependence. At $h = 0$ the slope of the lines is compatible with the exact result of the decay exponent in Eq. (6). For somewhat larger longitudinal fields, $h = 0.05$ and 0.1, the slopes of the line segments are somewhat larger than at $h = 0.0$, but due to errors of the calculation one cannot exclude the possibility that the asymptotic exponents agree with that at $h = 0.0$.

We shall come back to study the scaling behavior of the system for small values of $h$ by the SDRG method in the following Sec. III D and postpone the analysis of the phase diagram to the discussion in Sec. IV.
D. SDRG calculations for \( h \ll 1 \)

Here we are going to extend the SDRG method [2], which has been developed for \( h = 0 \) by including small longitudinal fields which are treated as a perturbation. For the sake of simplicity we use here the transformation \( \sigma_j^z = (-1)^j\sigma_j^z \), when the model is described by random ferromagnetic couplings (and transverse fields) in the presence of a staggered longitudinal field. This last part of the Hamiltonian is generally written as

\[
\hat{H}_{\text{longi}} = - \sum_{j=1}^{L} h_j \sigma_j^z,
\]

and in the starting situation \( h_j = (-1)^j h \).

In the SDRG procedure we consider the separated local degrees of freedom, say at position \( i \). These are couplings or sites having the value of the largest gaps: \( 2J_i \) and \( 2\sqrt{\Gamma_i^2 + h_i^2} \), respectively. These gaps are sorted in descending order and the largest one, denoted by \( \Omega \), which sets the energy scale in the problem, is eliminated and between remaining degrees of freedom new terms in the Hamiltonian are generated through perturbation calculation. This procedure is successively iterated, during which \( \Omega \) is monotonously decreasing. At the fixed point, with \( \Omega^* = 0 \) one makes an analysis of the distribution of the different parameters and calculates the scaling properties. In the following we describe the elementary decimation steps.

1. Strong-coupling decimation

In this case the largest local term in the Hamiltonian is a coupling, say \( \Omega = J_i \), connecting sites \( i \) and \( i + 1 \) and the two-site Hamiltonian is given by

\[
\hat{H}_{CP} = -J_i \sigma_i^z \sigma_{i+1}^z - \Gamma_i \sigma_i^x - \Gamma_{i+1} \sigma_{i+1}^x - (h_i \sigma_i^z + h_{i+1} \sigma_{i+1}^z).
\]

(13)

The spectrum of \( \hat{H}_{CP} \) contains four levels, the lower two being separated from the higher two by a gap \( 2J_i \). We omit the higher two levels, which is equivalent to merging the two strongly coupled sites into a spin cluster in the presence of a (renormalized) transverse field \( \Gamma \) and a longitudinal field \( \bar{h} \).

The magnetic moment of the cluster is given by \( \bar{\mu} = \mu_i + \mu_{i+1} \), where the magnetic moments in the starting situation are \( \mu_i = \mu_{i+1} = 1 \). The remaining two lowest energy levels are given by second-order degenerate perturbation method as the eigenvalues of the matrix:

\[
\begin{bmatrix}
-J_i - h_i - h_{i+1} & -\Gamma_i \Gamma_{i+1} / J_i \\
-\Gamma_i \Gamma_{i+1} / J_i & -J_i + h_i + h_{i+1}
\end{bmatrix}
\]

(14)

having a gap

\[
\Delta E_{CP} = 2\sqrt{(\Gamma_i \Gamma_{i+1} / J_i)^2 + (h_i + h_{i+1})^2}.
\]

(15)

Comparing it with the gap of the spin cluster, we obtain for the renormalized values of the parameters

\[
\bar{\Gamma} = \frac{\Gamma_i \Gamma_{i+1}}{J_i}, \quad \bar{h} = h_i + h_{i+1}.
\]

(16)

Note that in the starting situation with \( h_i = -h_{i+1} \) after decimating a strong coupling the longitudinal field is eliminated at the effective composite spin.

2. Strong-transverse-field decimation

In this case the largest local term is a transverse field, say \( \Gamma_i \), and the corresponding energy gap of the one-site Hamiltonian is \( 2\sqrt{\Gamma_i^2 + h_i^2} \). Due to the large \( \Gamma_i \) this site does not contribute to the longitudinal magnetization and therefore it is eliminated. The longitudinal magnetic field, however, should be transformed at the remaining neighboring sites. To calculate the new renormalized coupling between the remaining sites \( i - 1 \) and \( i + 1 \), we calculate energy levels with fixed spins at these sites. Denoting by \( s_{i+1} = \pm (-) \) a \( \uparrow (\downarrow) \) boundary state, the eigenvalue problem with different boundary conditions has the lowest energy as

\[
E_{s_{i+1}} = -\sqrt{\Gamma_i^2 + h_i^2}.
\]

(17)

The renormalized coupling between the remaining sites is given by

\[
J = (E_{\uparrow \downarrow} + E_{\downarrow \uparrow} - E_{\uparrow \uparrow} - E_{\downarrow \downarrow}) / 4 \approx \frac{J_i - J_{i+1}}{\sqrt{\Gamma_i^2 + h_i^2}}.
\]

(18)

where the last relation is calculated perturbatively.

Concerning renormalization of the longitudinal magnetic fields, we require that the sum of these fields is locally conserved, in agreement with the original Hamiltonian. This is obtained by adding \( h_i / 2 \) to the longitudinal fields at the neighboring sites, \( \bar{h}_{i+1} = h_{i+1} + h_i / 2 \), and in this way we try to minimize random-field effects [41].

3. Numerical iteration of the SDRG equations

We have iterated the decimation equations presented in the previous sections for finite periodic chains of length \( L = 256, 512, \) and 1024 up to the last pair of spins and the energy gap of this dimer is identified as the gap of the given sample. We have considered 10 000 independent samples for each case. We have also calculated the total magnetic moment of the samples, and calculated their average value, \( \mu_L \), which scales differently in the different phases. In the ordered phase it is extensive, \( \mu_L \sim L \), while in the disordered phase it approaches a finite limiting value. At the transition point there is a power-law dependence \( \mu_L \sim L^{d_f} \), with a fractal dimension \( 0 < d_f < 1 \). This is related to the decay exponent of the correlation function, since \( C(L) \sim L^{-2(1-d_f)} \), for large \( L \).

First, we have checked that at the RTIM fixed point with \( h = 0 \) the critical point is at \( \Gamma^* \sim 1 \) and the magnetic fractal dimension is \( d_f = 0.81 \), which is in good agreement with the analytical result \( d_f = \phi / 2 \) [see in Eq. (5)]. The distribution of the gaps, as shown in the main panel of Fig. 7, is also in agreement with the scaling relation in Eq. (4).

Next switch on a small longitudinal field, \( h = 0.05 \), when the transition point is moved to \( \Gamma^* \approx 0.9975 \). The estimate for the magnetic fractal dimension \( d_f \approx 0.83 \) is somewhat larger than at \( h = 0 \). The calculated gaps show infinite disorder.
In the vicinity of the RCMP by including quantum fluctuations, $\Gamma_0 > 0$, the system shows reentrant behavior: for $h > 2$ the system with increasing value of $\Gamma_0$ first moves from the disordered phase to the AFM phase and then back to the disordered one. Reentrance in random quantum systems has been observed only in a few cases: in random dimer networks [42,43] and in randomly dimerized AF $S = 1/2$ chains [44]. In classical systems it is usually the result of competing interactions and/or frustration [45]. In our system at the clean classical multicritical point, the ground state is infinitely degenerate. This degeneracy is lifted through disorder [46] at several steps, but the new, nonordered states are typically less favored by quantum fluctuations than the ordered state, which leads to the reentrant behavior.

Between the infinite disorder scaling regime and the random first-order transition region there must be a repulsive, multicritical fixed point, which separates the two parts of the transition line. This fixed point is expected to be the result of the competition between random couplings, random transverse fields, and the homogeneous longitudinal field. With our present investigations we could not explore the properties of this hypothetical fixed point; it will be the aim of further studies.

Outside the transition line the system exhibits singular dynamical behavior due to Griffiths singularities. Near the RTIM these are similar to that mentioned in Sec. III A [see in Eqs. (8) and (9)]. These have been studied in more detail in [29].

The model can be extended and generalized in different directions. In higher dimensions one should consider bipartite lattices, which can accommodate AFM order. Here, at $h = 0$ there is a higher-dimensional RTIM fixed point, which is known to be infinite disorder type [11–13]. Here one should study first the behavior in the classical limit and then the complete phase-transition line. Finally, one can also extend our model with random longitudinal fields. In this case, however, the disorder fluctuations are so strong that no ordered phase exists, even with vanishing quantum fluctuations.

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**APPENDIX A: NUMERICAL ALGORITHM TO CALCULATE THE GROUND STATE IN THE CLASSICAL LIMIT**

It is possible to map the problem of finding the ground state onto a max-flow problem [47]. However, in the one-dimensional problem it is more efficient to use the following simpler algorithm: (i) consider the set $S$ of all pairs of bonds separated by an even number of bonds; (ii) assign to each pair of $S$ the sum of the two couplings, $J_i + J_j$; and sort $S$ according to this sum; and (iii) take each pair $(i, j)$ of $S$ in increasing order and choose the state with the lowest energy between the three following cases. Starting with an
AFM state, (1) flip all spins between i and j; (2) flip all spins between j and i, or (3) flip no spin. Note that we assume L even, and periodic boundary conditions. This algorithm produces all the \( \frac{1}{2} \) ground states when \( h \) varies.

There are \( n = (\frac{L}{2} - 1) \frac{L}{2} \) pair of bonds at even distance, and the sort algorithm has a complexity \( n \ln n \). However, in practice one is interested only in the critical region, which means that only the pair of bonds \((J_i, J_j)\) with \( J_i + J_j \) small enough have to be considered. This greatly accelerates the algorithm.

**APPENDIX B: AVERAGE CORRELATION FUNCTION IN THE CLASSICAL LIMIT**

Let us consider a large finite chain \( L \gg 1 \) in the vicinity of the random classical multicritical point \( h - 2 < 1 \). Let us assume that we are in the transition regime, thus the random samples are either fully AFM ordered, thus \( h < J_{i_1} + J_{i_2} \) and being a fraction \( p_1 \), or contain just two AFM domains \( h \geq J_{i_1} + J_{i_2} \) being a fraction \( p_2 \), and we omit those, which contain more AFM domains, thus \( p_1 + p_2 = 1 \). In the AFM ordered samples the average correlation function is 1. In samples with two domains for a given sampled average correlation function is \( (1 - 4i/L) \), where \( i \leq L/2 \) is the size of the smaller AFM domain: \( i = \min(|i_1 - i_2|, L - |i_1 - i_2|) \). The position of the smallest couplings, \( i_1 \) and \( i_2 \), are random, therefore the distribution of the domain sizes \( 1 \leq i \leq L/2 - 1 \) is uniform. Consequently for samples with two AFM domains the ensemble-averaged correlation function is 0. Then the total average of the correlation function over fully AFM ordered and two domain samples is given by \( C(L/2 - 1, h) = p_1 = 1 - p_2 \).

The fraction of samples with two AFM domains can be calculated through extreme-value statistics [36,37]. For this we should note that with a sample with two domains appears, if \( h - 2 = (J_{i_1} + J_{i_2} - 1) \), where \( \epsilon_1 = J_{i_1} - 1 \) as well as \( \epsilon_2 = J_{i_2} - 1 \) are the smallest values out of the \( L/2 \) ones \((i_1 \) and \( i_2 \) being of different parity) having a parent distribution, which is uniform in \([0,1]\). According to extreme-value statistics [37], the asymptotic form of the distribution of \( \epsilon_1 \), and that of \( \epsilon_2 \), depends on the asymptotic behavior of the parent distribution for small argument. If it is in the form \( P(\epsilon) \sim \epsilon^{\omega} \), the scaling combination reads as \( u_1 = u_0(L/2)^{\gamma} \epsilon_1 \), with \( 1/\gamma = 1 + \omega \) and \( u_0 \) is a constant. For the uniform distribution we have \( \omega = 0 \), thus \( \gamma = 1 \). The distribution of \( u_1 \) is given by the Fréchet distribution:

\[
P(u_1) = \frac{1}{z} u_1^{1/z-1} \exp\left[-u_1^{1/z}\right],
\]

and similarly for \( u_2 \). Then for the longitudinal field the appropriate scaling variable is \( u = u_0(L/2)^{\gamma} (h - 2) = u_1 + u_2 \) and its distribution is given as the convolution of \( P(u_1) \) and \( P(u_2) \), which for \( z = 1 \) is given by

\[
P(u) = u \exp(-u).
\]

The fraction of samples with two domains is given by the accumulated distribution:

\[
p_2(u) = \int_0^u P(u)du = 1 - (1 + u) \exp(-u),
\]

from which the result in Eq. (11) follows.

**APPENDIX C: QUANTUM CORRECTIONS TO THE CLASSICAL LIMIT AT THE RCPM POINT**

Let us consider the \( h > 2 \) part of the phase diagram for a large but finite value of \( L \), at such point where in the ground state of the classical model there is exactly one reversed domain, the domain boundaries being at \( i = a = 2 \alpha + 1 \) and \( i = L \). The couplings at the boundaries \( J_0 \) and \( J_L \) are the smallest at the odd and even positions, respectively, and \( J_0 + J_L - h < 0 \). The energy of the classical ground state is

\[
E_0 = E_{AF} + 2(J_a + J_L - h)
\]

where the classical AFM state has the energy \( E_{AF} = -\sum_{i=1}^L J_i \).

Now let us switch on the transverse fields, and for simplicity let us consider a position-independent strength, \( \Gamma \ll 1 \). The first nonvanishing correction to the AFM state is given by

\[
\epsilon_2^{AF} = -\sum_{j=1}^{L/2} \left[ \frac{\Gamma^2}{2(J_{2j-1} + J_{2j} - h)} \right]^{1/2} \left[ \frac{\Gamma^2}{2(J_{2j} + J_{2j+1} + h)} \right]^{1/2}.
\]

The same type of corrections to the classical ground state are

\[
\epsilon_2^0 = -\sum_{j=1}^{\alpha-1} \left[ \frac{\Gamma^2}{2(J_{2j-1} + J_{2j} - h)} \right]^{1/2} \left[ \frac{\Gamma^2}{2(J_{2j} + J_{2j+1} + h)} \right]^{1/2} - \frac{\Gamma^2}{2(J_{2j-2} + J_{2j-1} - h)} - \frac{\Gamma^2}{2(J_{2j-1} - J_a + h)} - \frac{\Gamma^2}{2(-J_a + J_{2j+1} + h)} + \sum_{j=0}^{L/2-2} \left[ \frac{\Gamma^2}{2(J_{2j} + J_{2j+1} - h)} \right]^{1/2} \left[ \frac{\Gamma^2}{2(J_{2j+1} + J_{2j+2} + h)} \right]^{1/2} - \frac{\Gamma^2}{2(J_{2j} + J_{2j+1} - h)} - \frac{\Gamma^2}{2(J_{2j-2} + J_{2j-1} - h)} - \frac{\Gamma^2}{2(J_{2j-1} - J_a + h)} - \frac{\Gamma^2}{2(-J_a + J_{2j+1} + h)}.
\]

Large contributions to the sums in Eqs. (C2) and (C3) are due to such terms, in which \( h \) in the denominator has a minus sign. In Eq. (C2) there are \( L/2 \) such large terms, while in Eq. (C3) there are just \( L/2 - 1 \). Consequently, on average \( \epsilon_2^0 > \epsilon_2^{AF} \) and \( \epsilon_2^0 - \epsilon_2^{AF} \sim \Gamma^2 > 0 \), thus with increasing \( \Gamma \) the quantum correction is more and more favorable for the AFM ordered state, which at a given critical value can overcome the difference in the classical energy terms in Eq. (C1). This fact is in agreement with reentrance observed numerically in Fig. 5.
[35] If the distribution of the small random couplings is in the form $\pi(J) \sim (J - J_0)^\omega$, $(J - J_0) \ll 1$, then the appropriate scaling combination is $\kappa = u_0/(h - 2)^{1+\omega}$. For the box distribution in Eq. (2), $\omega = 0$.
[41] Using our decimation rules the renormalized longitudinal fields are either zero or a fraction of $h$, but the sum of the longitudinal fields remains zero. If $h$ is small enough we can perform the renormalization up to the last step avoiding the necessity of a longitudinal field decimation.