

Competition between quenched disorder and long-range connections: A numerical study of diffusion

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The problem of random walk is considered in one dimension in the simultaneous presence of a quenched random force field and long-range connections, the probability of which decays with the distance algebraically as $p_l \simeq \beta l^{-s}$. The dynamics are studied mainly by a numerical strong disorder renormalization group method. According to the results, for $s > 2$ the long-range connections are irrelevant, and the mean-square displacement increases as $\langle x^2(t) \rangle \sim (\ln t)^{2/\psi}$ with the barrier exponent $\psi = 1/2$, which is known in one-dimensional random environments. For $s < 2$, instead, the quenched disorder is found to be irrelevant, and the dynamical exponent is $z = 1$ like in a homogeneous environment. At the critical point, $s = 2$, the interplay between quenched disorder and long-range connections results in activated scaling, however, with a nontrivial barrier exponent $\psi(\beta)$, which decays continuously with β but is independent of the form of the quenched disorder. Upper and lower bounds on $\psi(\beta)$ are established, and numerical estimates are given for various values of β . Besides random walks, accurate numerical estimates of the graph dimension and the resistance exponent are given for various values of β at $s = 2$.

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I. INTRODUCTION

Long-range interactions are known to affect the dynamics of various systems such as equilibrium spin models [1], nonequilibrium processes [2], or diffusion [3]. A well-known generalization of normal diffusion is the Lévy flight [3,4], where long jumps are allowed with probabilities decaying algebraically with the distance as $p_l \sim l^{-1-f}$. These long jumps lead to superdiffusion with an anomalous dynamical exponent $z = f$ for $f < 2$, while for fast enough decay ($f > 2$) the diffusion is normal ($z = 2$). Contrary to long-range interactions, local heterogeneities, frequently slow the dynamics, which manifests itself in an increased dynamical exponent compared to that of the pure system. This is also the case for diffusion in various types of random environments [3,5]. Here the most dramatic slowing is induced by the random-force disorder, where the direction of the local force acting on the particle is random. This type of disorder is known to be relevant if the dimension d is less than 2 [6,7]. In one dimension, the mean-square displacement increases in time as $\langle x^2(t) \rangle \sim (\ln t)^4$ [8], where $\langle \cdot \rangle$ denotes average over stochastic histories for a given random environment and the overbar denotes average over random environments. This type of ultra-slow dynamics corresponds formally to zero dynamical exponent $z = 0$.

It is an intriguing question what happens when both long-range jumps and random force disorder are present in a system and whether there is a nontrivial dynamical behavior due to the interplay between them. In case of diffusion, a dynamic renormalization group analysis showed that the dynamical exponent equals the Lévy index f even in the presence of weak disorder for $f < 2$ [9]. Thus occasional long jumps enable the particle to escape from trapping regions of the random environment, and disorder is irrelevant in the above sense.

An alternative form of long-range interactions is realized when the interaction strength is constant but only certain

pairs of sites interact with each other, which are selected randomly with a probability decaying asymptotically with the distance l of the sites as $p_l \simeq \beta l^{-s}$. This paradigm arises, e.g., in the context of magnetic [10] and conducting [11] properties of linear polymers with cross-links between certain monomers or in models of social or communication networks that exhibit the small-world phenomenon [12,13]. The interactions are in this case represented by a random graph, which is composed of short edges between neighboring sites and long edges between certain remote sites. This model carries thus an intrinsic quenched disorder due to the interaction network as opposed to the former paradigm, which can be regarded as an annealed counterpart of the latter. Here even the geometrical properties of the interaction graph are nontrivial [14–20]. Adding long edges to a one-dimensional lattice, the resulting graph is quasi-one-dimensional if $s > 2$, whereas it is formally infinite-dimensional if $s < 2$ [19]. These domains are separated by a critical point at $s = 2$, where the graph dimension is conjectured to be finite and to depend on the prefactor β [15]. The geometrical properties of the underlying interaction network indicate that such models may show nonconventional dynamical behavior. Indeed, the contact process with long-range interactions of this type is conjectured to exhibit activated dynamical scaling at $s = 2$ with β -dependent critical exponents [21,22].

One can pose the question of how the dynamics in general are affected by the simultaneous presence of this type of long-range connections and quenched local disorder. In this paper we shall study this issue by considering a simple dynamical process, the random walk with random-force disorder on graphs containing long edges with algebraically decaying probability. Most efforts will be devoted to the critical point $s = 2$, where the graph dimension is finite and which gained less attention in earlier numerical studies. As the dynamics are expected to be slow here, we shall mainly apply a numerical strong disorder renormalization method [23] carried out in the configuration space [24–27] rather than Monte Carlo simulations. Besides the above investigations, the nontrivial geometrical properties such as the graphs' dimension and the

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random-walk dimension will be estimated at $s = 2$ as well. Our results, which are accurate for moderate β at the critical point $s = 2$, indicate that the dynamics are still activated; i.e., the mean-square displacement increases as $\langle x^2(t) \rangle \sim (\ln t)^{2/\psi(\beta)}$. However, the barrier exponent $\psi(\beta)$ differs from $1/2$ and decreases continuously with the prefactor β .

The rest of the paper is organized as follows. The precise definition of the model is given in Sec. II. In Sec. III numerical estimates of the shortest-path dimension are presented. In Sec. IV bounds on the random-walk dimension are established, as well as numerical estimates are given. In Sec. V the strong disorder renormalization group method is reviewed and applied to the problem of random walks with long-range connections in random environment. Finally, the results are discussed in Sec. VI.

II. THE MODEL

The finite connectivity networks to be studied are defined as follows. Assume that the sites are numbered by the integers $1, 2, \dots, N$ and define the distance between site i and j as $l_{ij} = \min(|i - j|, N - |i - j|)$. In words, the sites are arranged on a ring with unit spacing between them. Then all pairs of sites with a distance $l_{ij} = 1$ (i.e., neighboring sites on the ring) are connected with an edge, and all pairs with $l_{ij} > 1$ are connected independently with the probability [15,28]

$$P(l) = 1 - \exp(-\beta l^{-s}), \quad (1)$$

where β and s are positive constants. For large l , this probability has the asymptotic form $p(l) \simeq \beta l^{-s}$.

On each realization of the above connectivity network a continuous-time random walk is considered, which is specified by the set of i.i.d. quenched random transition rates w_{ij} from site i to site j . Note that the transition rates through long connections ($l_{ij} > 1$) are random just as those through short connections ($l_{ij} = 1$). Note, furthermore, that the source of randomness is twofold in this model. First, the random topology of the connectivity network, and, second, the set of random transition rates.

III. SHORTEST PATHS

From the point of view of dynamical processes like random walk, the relevant metric is not the distance l_{ij} but the length ℓ of shortest paths (or chemical distance) between two sites, which is given by the number of (short or long) edges that constitute the path. If $s > 2$, the expected value of the length of long edges is finite, and therefore the average shortest path length is asymptotically proportional to the distance l ; in other words, the system is quasi-one-dimensional. At the critical point $s = 2$, the mean shortest path length has been conjectured to grow algebraically with the distance:

$$\bar{\ell} \sim l^{1/d_g(\beta)}, \quad (2)$$

where the graph dimension $d_g(\beta)$ depends on β [15]. Indeed, later a lower bound on $d_g(\beta)$ was proven: $d_g(\beta) \geq \ln 3 / \ln(3 - \delta)$, where $\delta = 1 - e^{-\beta/9}$, as well as an upper bound, $d_g(\beta) \leq 1/(1 - \beta)$, for $\beta < 1$ [18]. The latter inequality is related to the presence of so-called cut points, which are defined as follows.

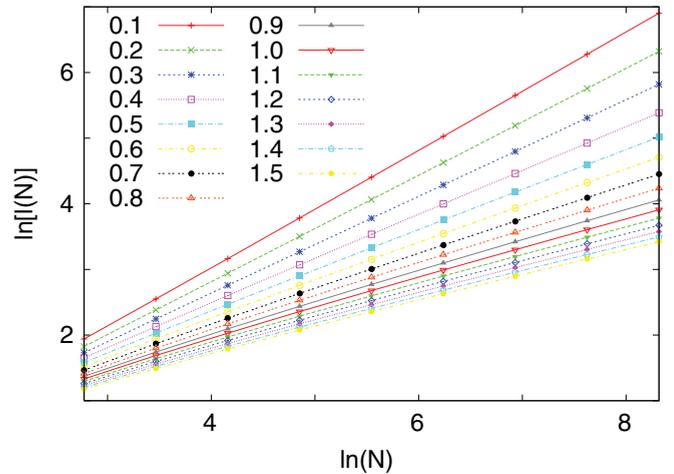


FIG. 1. (Color online) The logarithm of the numerically calculated mean length of shortest paths between sites in a distance $N/2$ plotted against $\ln N$ for different values β at $s = 2$. The statistical errors are smaller than the size of the symbols.

First, instead of the cyclic graphs defined in the previous section, let us consider acyclic ones, which are defined in the same way except that the modified distance $l_{ij} = |i - j|$ is used there. In such acyclic graphs, a cut point is a site the removal of which (and all of its edges) results in that the graph becomes disconnected. The expected number of such points tends to zero for $s < 2$, is $O(N)$ for $s > 2$ and $O(N^{1-\beta})$ for $s = 2, \beta < 1$ [18,28]. Obviously, the shortest path between site 1 and site N must contain all cut points of the graph, which implies the above lower bound on $d_g(\beta)$. For $s < 2$, the graph dimension is formally infinite; for details see Refs. [15,18,19].

We considered networks of size $N = 2^n$ with $n = 4, 5, \dots, 12$ and calculated the length of the shortest path between pairs of sites in a distance $l = N/2$ numerically by a breadth-first search. Here 10^6 independent networks have been generated for each size, and ℓ between $N/8$ pairs of sites has been calculated in each realization. The mean length of shortest paths obtained in this way is plotted against N in Fig. 1.

As can be seen, the data are compatible with the power law in Eq. (2), and the estimated graph dimensions, which are obtained extrapolating the effective two-point exponents to $N \rightarrow \infty$, lie between the rigorous bounds quoted above; see Fig. 2 and Table I.

IV. RANDOM WALK AND ELECTRICAL RESISTANCE

Before discussing the random walk with quenched random transition rates, we investigate the homogeneous model, where the transition rates are equal, e.g., $w_{ij} = 1$. For this model, the random walk is known to be recurrent if $s \geq 2$ and transient if $1 < s < 2$ [29,30]. On the grounds of calculations in similar but analytically tractable random networks [31], superdiffusion is expected for $s = 2$; i.e., the mean-square displacement in an infinite network is expected to behave asymptotically as

$$\overline{\langle x^2(t) \rangle} \sim t^{2/d_w}, \quad (3)$$

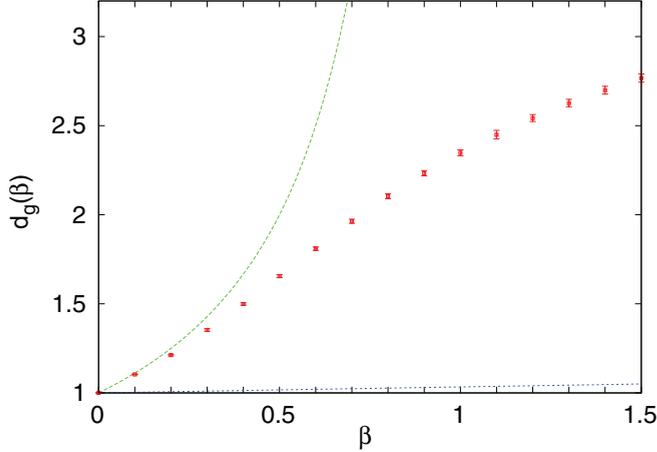


FIG. 2. (Color online) Graph dimensions estimated from the data shown in Fig. 1 for different values of β . The solid lines represent the rigorous bounds on $d_g(\beta)$ [18].

where the random-walk dimension d_w [32] is less than 2. Although the exact value of d_w is unknown, we can establish bounds on it in terms of the graph dimension, and then we will give numerical estimates. In order to do this, we make use of the well-known relationship between diffusion and resistor networks [3,5]. For symmetric random walks (where $w_{ij} = w_{ji}$ holds) on an arbitrary graph, a resistor network can be defined where the edges of the graph are resistors with a resistance $R_{ij} = 1/w_{ij}$. In the case of homogeneous transition rates, the mean first-passage time from one site to another one whose distance is on the order of the graph diameter (e.g., site 1 and site $N/2 + 1$ for cyclic graphs) is related to the effective resistance $R(N)$ of the corresponding resistor network between the two sites as $t \sim NR(N)$ for large N [31,33]. Assuming that Eq. (3) holds, the average effective resistance behaves

TABLE I. The estimated values of $1/d_g$, ζ and ψ for different values of β . The estimate for ψ has been obtained by the finite-size-scaling of $\ln(1/\Omega_N)$ using discrete randomness with $r = 0.1$.

β	$1/d_g$	ζ	ψ
0.1	0.906(2)	0.902(3)	0.452(5)
0.2	0.825(3)	0.809(4)	0.405(5)
0.3	0.739(4)	0.719(4)	0.352(5)
0.4	0.667(3)	0.633(5)	0.315(10)
0.5	0.604(3)	0.550(8)	0.265(10)
0.6	0.552(3)	0.473(10)	0.224(10)
0.7	0.509(3)	0.405(13)	0.185(10)
0.8	0.475(3)	0.343(13)	0.158(10)
0.9	0.448(3)	0.299(14)	0.135(10)
1.0	0.426(3)	0.257(16)	0.115(10)
1.1	0.408(4)	0.220(17)	0.09(1)
1.2	0.393(3)	0.204(19)	0.075(20)
1.3	0.381(3)	0.18(2)	0.07(2)
1.4	0.371(3)	0.16(2)	0.065(20)
1.5	0.361(3)	0.15(2)	0.055(20)

asymptotically as

$$R(N) \sim N^\zeta, \quad (4)$$

where the resistance exponent ζ is related to the random-walk dimension as

$$d_w = 1 + \zeta. \quad (5)$$

Bounds on the random-walk dimension at $s = 2$ can be obtained by simple considerations as follows. In a cyclic graph of size N let us consider two sites in a distance $l = N/2$. The length of the shortest path between them is $O(N^{1/d_g})$ according to Eq. (2). From Eq. (3) we obtain that the mean first-passage time from one site to the other one is $O(N^{d_w})$. Deleting now all edges other than those contained in the shortest path, one obtains a one-dimensional system in which the diffusion is normal. So the expected first-passage time in this one-dimensional chain is $O(N^{2/d_g})$, and it is easy to see that this time must not be greater than that in the original network, implying the inequality

$$d_w \geq 2/d_g. \quad (6)$$

Consider again the above one-dimensional chain between the two sites. The effective resistance of this chain is $O(N^{1/d_g})$, and this must be larger than the effective resistance between the two points in the original network, which is $O(N^\zeta)$, since removing links from the original network does not decrease the effective resistance. So we have the relation $1/d_g \geq \zeta$ or, using Eq. (5),

$$d_w \leq 1 + 1/d_g. \quad (7)$$

Since, for $\beta > 0$, $d_g(\beta) > 1$ holds, we have $d_w < 2$, i.e., superdiffusion when $\beta > 0$. Furthermore, since both the lower and upper bounds decrease with β and for $\beta = 0$ the above relations hold as equalities, we conclude that the random-walk dimension must also depend on β .

In order to estimate $d_w(\beta)$, we have numerically calculated the effective resistance between sites 1 and $N/2 + 1$ in cyclic networks of size N by the method described in the next section. We considered system sizes $N = 2^n$ with $n = 4, 5, \dots, 14$ and generated 10^6 independent networks for $n \leq 12$ and 10^5 for $n = 13, 14$. The average resistance is plotted against N for different values of β in Fig. 3. As can be seen, the data are in agreement with the power law in Eq. (4) with a β -dependent resistance exponent. The estimated values of $\zeta(\beta)$ obtained by extrapolating the two-point effective exponents are plotted for different values of β in Fig. 4 and are given in Table I. The numerical estimates of $d_w(\beta)$ and $d_g(\beta)$ are found to satisfy the inequalities (6) and (7).

V. RANDOM WALK IN RANDOM ENVIRONMENT

A. Finite graph dimension ($s \geq 2$)

After having studied the geometry and homogeneous random walks we turn to the full problem, where the transition rates are quenched random variables. In one dimension, which corresponds to $\beta = 0$ in our model, the mean-square displacement in an infinite system behaves as $\langle x^2(t) \rangle \sim (\ln t)^4$ [8].

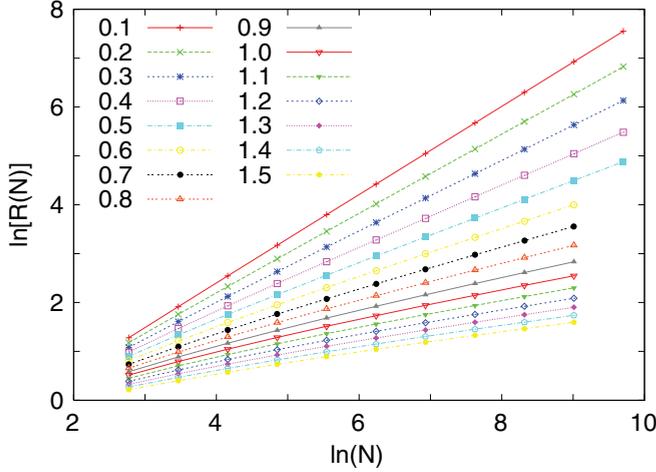


FIG. 3. (Color online) The logarithm of the numerically calculated average resistance between sites in a distance $N/2$ plotted against $\ln N$ for different values β at $s = 2$. The statistical errors are smaller than the size of the symbols.

It is intuitively obvious that adding long edges to the one-dimensional chain results in the mean-square displacement increasing faster with time since considerably shorter paths between far away sites are opened. For $s > 2$, on the other hand, the number of cut points, which must be contained in shortest paths, is $O(N)$. Therefore the logarithm of the typical time to travel through a domain of length N is at least $O(N^{1/2})$. As a consequence, for $s > 2$, the dynamics must obey the asymptotic law

$$\overline{\langle x^2(t) \rangle} \sim (\ln t)^{2/\psi}, \quad (8)$$

with the barrier exponent $\psi = 1/2$ as in one dimension, and only the proportionality constant is expected to be modified by the presence of long edges.

At the critical point $s = 2$, $\beta > 0$, the form in Eq. (8) with $\psi = 1/2$ is a lower bound on $\overline{\langle x^2(t) \rangle}$. An upper bound for

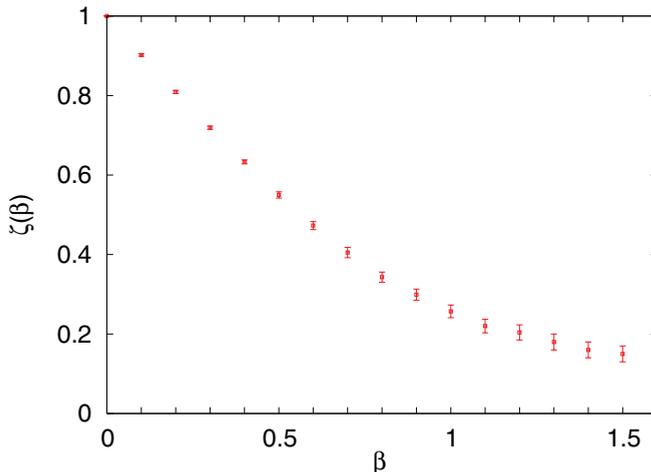


FIG. 4. (Color online) Resistance exponents estimated from the data shown in Fig. 1.

$\beta < 1$ can be easily established since the number of cut points is $O(N^{1-\beta})$; thus the logarithm of mean first-passage time must be typically at least $O(N^{(1-\beta)/2})$. Hence we expect for $s = 2$ and at least for $\beta < 1$ the activated dynamics given in Eq. (8) to hold with some barrier exponent $\psi(\beta)$, which may depend on β and lies in the range

$$(1 - \beta)/2 \leq \psi(\beta) \leq 1/2. \quad (9)$$

This type of ultra-slow dynamics suggests that a strong disorder renormalization group (SDRG) method can be applied to the process. This method has been originally constructed for studying disordered quantum spin chains [34], and later it was adapted for investigating dynamical processes; for a review see Ref. [23]. For random walks in one-dimensional random environments the method has been formulated in terms of the potential barriers [35]. Apart from tree-like graphs, however, potential cannot be defined in general, and the method has to be formulated in terms of transition rates [24–27].

In the SDRG procedure, sites with short waiting time are successively eliminated, and the remaining transition rates are modified such that the long-time dynamical behavior of the system remains unaltered. The detailed scheme of the method is the following. Let us introduce the exit rate from site i as $\Omega_i = \sum_j w_{ij}$, where the summation goes over all sites connected with site i . The inverse of Ω_i gives the expected waiting time at site i : $\tau_i = 1/\Omega_i$. The elementary step of the procedure is that the site that has the smallest waiting time is chosen and removed from the network together with their edges. The transition rates w_{jk} between sites that were connected to the eliminated site i are modified as

$$\tilde{w}_{jk} = w_{jk} + w_{ji}w_{ik}/\Omega_i. \quad (10)$$

If these sites were not connected before the decimation (i.e., $w_{jk} = w_{kj} = 0$), then a new edge is created between them with the rates given above. Through the transition rates, the exit rates and the waiting times of sites neighboring to the decimated one are also renormalized. This decimation step is then iterated starting from a finite system of size N until only one site remains with an effective waiting time $\tau_N = 1/\Omega_N$. The renormalization rule in Eq. (10) can be obtained from the condition that the ratios of steady-state probabilities of all sites not yet decimated (active sites) remain unchanged when a site is eliminated. Concerning the dynamics, the replacement of the original system by the one site smaller effective one is an approximation in the sense that the waiting time in the eliminated site is neglected. In certain systems, however, the distribution of the logarithm of waiting times becomes broader and broader as the ratio of active sites is decreased. In these systems, which are described by an infinite randomness fixed point, the above approximation becomes more and more accurate as the fraction of active sites tends to zero and the SDRG is said to be asymptotically exact [23]. The last effective waiting time obtained by this method τ_N is then on the order of the expected escape time from the system embedded in a larger environment, or in the case of an acyclic network, it is on the order of the mean first-passage time from site 1 to site

N . So, in the case of the activated dynamics given in Eq. (8), the last exit rate must scale with N as

$$\ln(1/\Omega_N) \sim N^\psi \quad (11)$$

for large N . It is easy to show that, in case of symmetric rates $w_{ij} = w_{ji}$, the renormalization rules in Eq. (10) reduce to the rules of calculating the effective resistance in the equivalent resistor network. This method has been used for calculating the effective resistance discussed in the previous section. A simplification in that case is that the sites can be eliminated in an arbitrary order.

We have performed the above SDRG procedure numerically in finite systems and calculated the distribution of the last effective exit rate and the average of its logarithm. In most of the numerical calculations, the random transition rates have been drawn from a discrete distribution, where the rates on the edge (ij) are either $w_{ij} = 1$ and $w_{ji} = r$ with probability $1/2$ or $w_{ij} = r$ and $w_{ji} = 1$ with probability $1/2$. Here r is constant that lies in the range $0 < r < 1$. We also considered a continuous distribution, namely, a uniform one in the domain $(0, 1)$, from which all transition rates are drawn independently. The size of systems was $N = 2^n$ with $n = 6, 7, \dots, 13$. The number of independent realizations was 10^6 for $n \leq 12$ and $\beta < 1$, while for larger β and larger N , where the SDRG procedure is slower, it was less; nevertheless, in all points it was at least a few times 10^4 .

The distributions of $\ln(1/\Omega_N)$ for $\beta = 0.7$ and for different system sizes are shown in Fig. 5. As can be seen, they broaden with increasing N , which justifies the applicability of the SDRG method *a posteriori*. The distributions are qualitatively similar for all considered values of β in the range $[0, 1.5]$. The probability densities have the scaling property

$$\rho[\ln(\Omega_0/\Omega_N), N] = N^{-\psi} \tilde{\rho}[\ln(\Omega_0/\Omega_N)N^\psi], \quad (12)$$

in accordance with the assumption written in Eq. (11). Here, the constants Ω_0 and ψ are found to depend on β .

Instead of the optimal data collapse, we have estimated the barrier exponent ψ systematically by the finite-size scaling of the average $\ln(1/\Omega_N)$. Here the form $\ln(\Omega_0/\Omega_N) = cN^\psi$ has been fitted to triples of data points for sizes N , $2N$, $4N$, and the effective values of ψ have been extrapolated to $N \rightarrow \infty$.

In one dimension it is known that the barrier exponent $\psi = 1/2$ is universal; i.e., it is independent of the distribution of transition rates. In order to test this property for $\beta > 0$ we have studied the point $\beta = 0.7$ with discrete randomness with $r = 0.1, 0.02$ and with the uniform one. The estimated values of ψ for these three cases are in order: 0.185 (10), 0.189 (10), and 0.19 (1). The differences are not significant, which suggests universality with respect to the distribution of transition rates. So the barrier exponent is expected to be a characteristic of the connectivity network. Next, we have determined the barrier exponents for different values of β ; see Figs. 6, 7, and Table I. As can be seen, ψ varies with β , and even for $\beta \geq 1$, where we do not have a lower bound on $\psi(\beta)$ at our disposal, the estimated values differ significantly from zero.

After presenting the numerical results we close this section with a heuristic reasoning, which is based on the SDRG

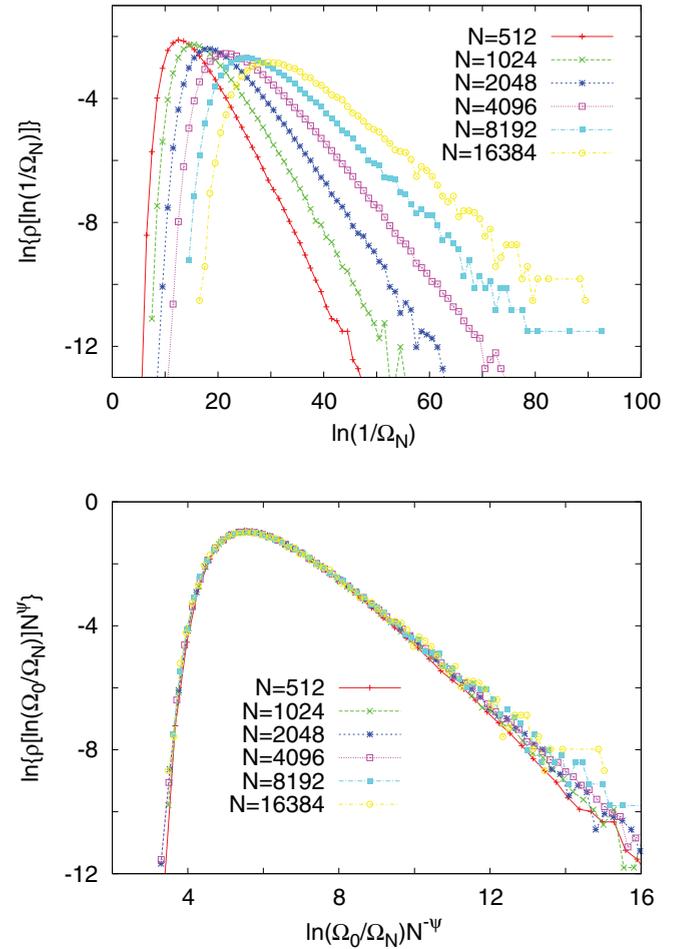


FIG. 5. (Color online) Top: Histogram of the logarithm of the last exit rate obtained by numerical renormalization, for $s = 2$, $\beta = 0.7$, and different system sizes N . The rates were drawn from the discrete distribution with $r = 0.1$ described in the text. Bottom: Scaling plot of the same data with $\psi = 0.19$ and $\ln \Omega_0 = 5.5$.

procedure and yields an upper bound on $\psi(\beta)$. Let us consider an acyclic network of size N and find the last site, which the SDRG ends up with. The last effective waiting time is in the order of the expected first-passage time from site 1 to site N , as we have already mentioned. Let us find now the shortest path that connects site 1 with site N and goes through the last site obtained by the SDRG. Its length is $O(N^{1/d_g})$. Then, following the rules of the SDRG method, we eliminate all sites except those contained in the shortest path. The elimination of the rest of the graph will generate new (long) edges between the sites of the path, but the initial rates on the edges of the path are never decreased. Furthermore, it is obvious that applying the SDRG procedure to this effective system ends up with the same site as when it was applied to the initial network. It is plausible to assume that adding long edges to a one-dimensional system decreases the expected first-passage time. Therefore the last effective waiting time of the system must not be longer than that of the above one-dimensional path (without the generated long edges). In the latter system the logarithm of the mean first-passage time is $O(N^{1/(2d_g)})$, so we obtain

$$\psi(\beta) \leq 1/[2d_g(\beta)]. \quad (13)$$

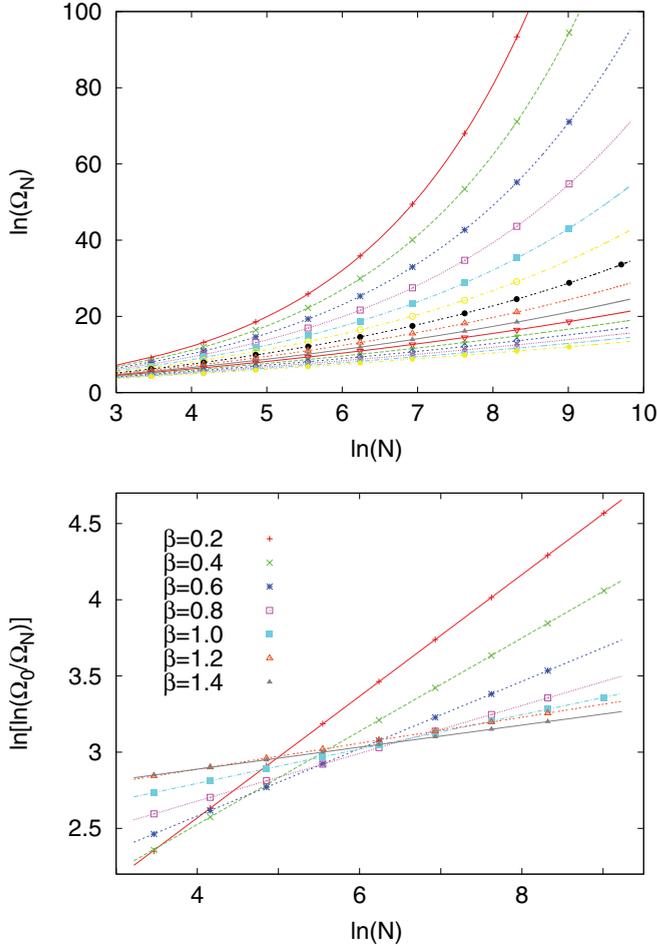


FIG. 6. (Color online) Top: The average of the logarithm of the last exit rate $\ln(1/\Omega_N)$ plotted against $\ln N$ for the same values of β as in Fig. 3. Data have been obtained by the numerical SDRG method using discrete randomness with $r = 0.1$. The lines are fits of the form in Eq. (11) to the data. Bottom: The logarithm of the quantity plotted in the upper figure for certain values of β . The slope of the straight lines is $\psi(\beta)$.

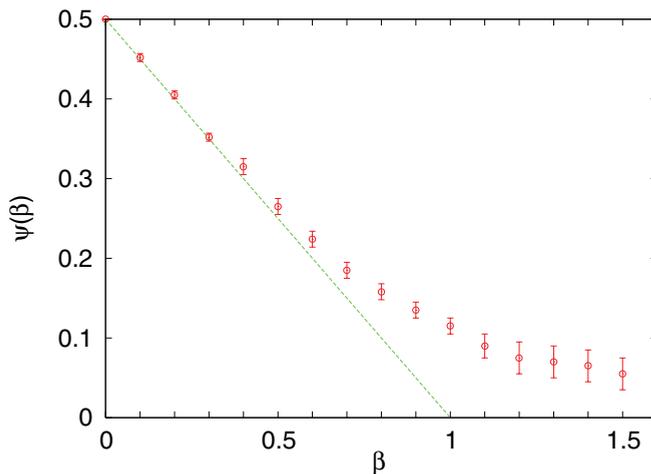


FIG. 7. (Color online) The estimated values of $\psi(\beta)$ given in Table I plotted against β . The straight line is the lower bound $(1 - \beta)/2$.

The numerical estimates are found to satisfy this relation for all β ; see Table I.

B. Infinite graph dimension ($1 < s < 2$)

In the domain $1 < s < 2$, the number of edges of the graph is still $O(N)$, but the shortest path length increases with the distance poly-logarithmically [15,19]. For an even smaller decay exponent, $s < 1$, the number of edges is $O(N^{2-s})$, and the shortest path length is bounded [15]; therefore we have restricted ourselves to the range $1 < s < 2$. Here the graph dimension is formally infinite. According to our numerical results, the average resistance seems to saturate with N , which corresponds to a vanishing resistance exponent, $\zeta = 0$. Using Eq. (5) it follows that the random walk with homogeneous transition rates is ballistic here, i.e., $d_w = 1$.

In case of random transition rates, we have calculated the distribution of the last exit rate by the SDRG method at several points. For $s = 1.5$, $\beta = 1$, the results are shown in Fig. 8. As can be seen, the distributions do not broaden with increasing N , indicating that the system is not described by an infinite randomness fixed point. In this case, the waiting times neglected in the procedure may be considerable, and the last waiting time is not necessarily on the order of the first-passage time through the system, but it may be only a vanishing fraction of the latter. The last waiting time is found to scale algebraically with N as $\tau_N \sim N^a$; see the scaling plot in Fig. 8. It is remarkable that the data fit well to the Fréchet distribution given by the probability density

$$f(\tilde{\tau}) = b\tilde{\tau}^{-1-b}e^{-\tilde{\tau}^{-b}}, \quad (14)$$

where $\tilde{\tau} = c\tau_N N^{-a}$ and b, c are constants. This is well known for the random walk and other processes in one-dimensional random environments in the driven phase [36], where the environment can be divided into roughly independent trapping regions with random trapping times having the asymptotical probability density $\rho(\tau) \sim \tau^{-1-b}$. The number of trapping

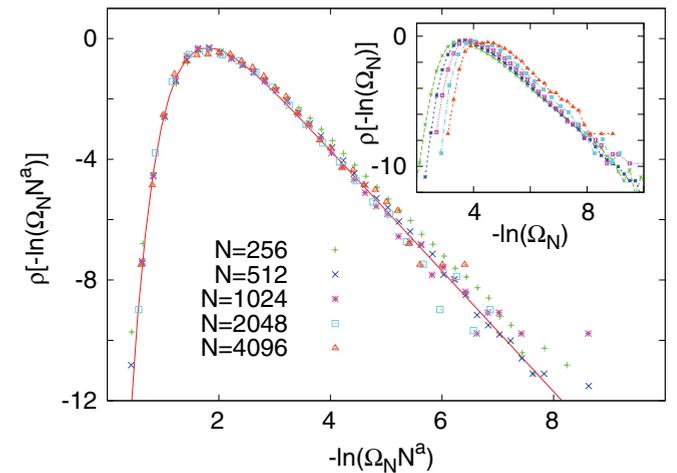


FIG. 8. (Color online) Scaling plot of the distribution of the logarithm of the last exit rate obtained by numerical renormalization, for $s = 1.5$, $\beta = 1$, and different system sizes N . The rates were drawn from the discrete distribution with $r = 0.1$. The scaling exponent is $a = 0.3$. The solid line is the Fréchet distribution given in Eq. (14) with $b = 2$. The unscaled data are shown in the inset.

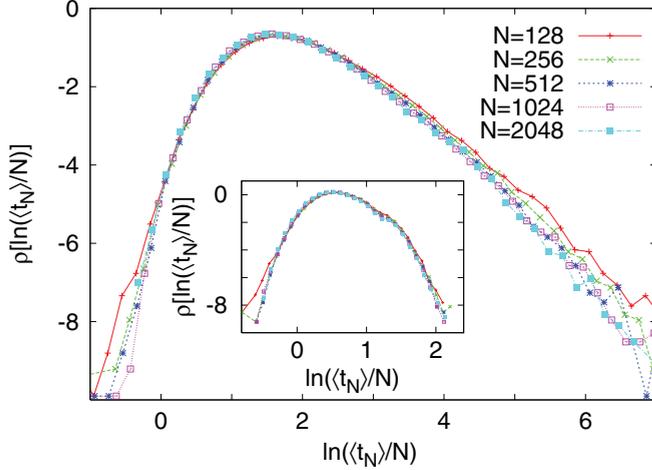


FIG. 9. (Color online) Scaling plot of the distribution of the logarithm of the mean first-passage time $\langle t_N \rangle$ obtained by Monte Carlo simulation in networks with $s = 1.5$, $\beta = 1$ for different system sizes. For the transition rates discrete randomness has been used with $r = 0.1$. The inset shows the same quantity in the same type of networks but with homogeneous transition rates ($r = 1$).

regions is proportional to N in one dimension; thus the distribution of largest waiting time follows Eq. (14) with $a = 1/b$. In our model with $1 < s < 2$, however, the latter relation seems not to be valid. Applying the above assumption on independent trapping regions with a power-law trapping time distribution this implies that the number of effectively independent trapping regions is not proportional to N but is only $O(N^{ab})$.

We have found that the above properties of the distribution of the largest waiting time are generally valid in the region $1 < s < 2$; however, the exponents a and b depend on s and β .

In order to reveal the dynamics we have carried out Monte Carlo simulations in cyclic networks and measured the first-passage time from site 1 to site $1 + N/2$. For each system size, 10^5 independent networks with random transition rates have been generated, and the mean first-passage time has been calculated in each network from 100 measurements. First, we have determined the distribution of the mean first-passage time in networks with homogeneous rates. As can be seen in the inset of Fig. 9, the mean first-passage time scales as $\langle t_N \rangle \sim N$, in accordance with $d_w = 1$.

The same quantity for random transition rates is shown in the same figure. As can be seen, the distributions are broader than in the homogeneous case, but the motion is still ballistic, i.e., $\langle t_N \rangle \sim N$. Thus we conclude that although the steady-state probabilities are inhomogeneous and there are trapping regions with large waiting times in case of random environments, the random-walk dimension is not altered compared to the homogeneous model ($d_w = 1$). So in this sense the disorder in the transition rates is irrelevant for $s < 2$.

VI. DISCUSSION

In this work we have studied one-dimensional random walks in the simultaneous presence of quenched disordered

transition rates and quenched long-range connections. The effects of these two components when they are present separately are known to be antagonistic: The former induces ultra-slow activated dynamics while the latter leads to superdiffusion. According to our results obtained mainly by a strong disorder renormalization group method, the scaling exponents are independent of the distribution of transition rates in general, but they do depend on the decay exponent s and the prefactor β of the probability of long connections. For $s > 2$, the long-range connections are irrelevant, and the dynamics follows activated scaling with a barrier exponent $\psi = 1/2$ like in one-dimensional random environments. For $s < 2$, where the graph dimension is formally infinite, the disorder in the transition rates still leads to the formation of trapping regions with large waiting times; however, the dynamical exponent is not altered compared to the homogeneous environment, so the disorder is irrelevant in this sense. At $s = 2$, the effects of disorder and long-range connections are comparable, and their interplay results in activated scaling with a nontrivial barrier exponent, which varies continuously with β . The barrier exponent is found to decrease with β , and a positive lower bound on $\psi(\beta)$ has been established only for $\beta < 1$. The numerical estimates are found to be above this exact lower bound, and the smooth decrease with β seems to continue also beyond $\beta = 1$ all the way to the largest value ($\beta = 1.5$) studied in this work. The numerical investigations are, unfortunately, limited to moderate β since, for larger β , the number of edges is larger as well, and consequently the SDRG procedure becomes more and more time consuming. In addition to this, the effective strength of disorder decreases with β , which would necessitate the investigation of larger and larger systems in order to see the correct asymptotical behavior. So it is still an open question whether $\psi(\beta)$ remains positive for arbitrarily large β or it becomes zero above some finite critical value.

In the light of the nontrivial results obtained here at the point $s = 2$, one can pose the question whether similar marginal behavior can be observed in the corresponding annealed model, i.e., in the simultaneous presence of quenched disordered transition rates and Lévy-flight-type long-range transitions at the index $f = 2$, which separates the phases where one of the above two components is irrelevant. Furthermore, the marginal behavior found at $s = 2$ suggests that an analogous behavior is possible in the case of more complex interacting systems with quenched disorder and long-range connections.

Throughout this work we have assumed that the average of the random force acting on the particle is zero. On the basis of the results, we expect the interplay between random forces with nonzero average and long-range connections also to result in nontrivial dynamics. The investigation of this issue is left for future work.

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