Percolation transitions in the survival of interdependent agents on multiplex networks, catastrophic cascades, and SOS

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The “SOS” in the title does not refer to the international distress signal, but to “solid-on-solid” (SOS) surface growth. The catastrophic cascades are those observed by Buldyrev \textit{et al.} in interdependent networks, which we re-interpret as multiplex networks with agents that can only survive if they mutually support each other, and whose survival struggle we map onto an SOS type growth model. This mapping not only reveals non-trivial structures in the phase space of the model, but also leads to a new and extremely efficient simulation algorithm. We use this algorithm to study interdependent agents on duplex Erdős-Rényi (ER) networks and on lattices with dimensions 2, 3, 4, and 5. We obtain new and surprising results in all these cases, and we correct statements in the literature for ER networks and for 2-d lattices. In particular, we find that \(d = 4\) is the upper critical dimension, that the percolation transition is continuous for \(d \leq 4\) but – at least for \(d \neq 3\) – not in the universality class of ordinary percolation. For ER networks we verify that the cluster statistics is exactly described by mean field theory, but find evidence that the cascade process is not. For \(d = 5\) we find a first order transition as for ER networks, but we find also that small clusters have a nontrivial mass distribution that scales at the transition point. Finally, for \(d = 2\) with intermediate range dependency links we propose a scenario different from that proposed in W. Li \textit{et al.}, PRL 108, 228702 (2012).

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I. INTRODUCTION

While percolation and related epidemic processes had appeared until recently as a mature subject that holds few surprises, this has changed dramatically during the last few years \cite{1}. A host of new models have been proposed, such as percolation on growing networks \cite{2}, percolation on hierarchical structures \cite{3}, agglomerative percolation \cite{4,5}, explosive percolation \cite{6}, percolation on interdependent networks \cite{7}, percolation where nodes cooperate to infect their neighbors \cite{8,10}, and percolation where spreading agents cooperate \cite{11,12}. Indeed, some of these models are not entirely new and are related to models studied since the 1970’s. The model of \cite{3}, e.g., can be viewed as a version of percolation on lattices with long range contacts \cite{13,14}, while cooperative percolation \cite{8,10} can be viewed as a variant of bootstrap percolation \cite{15,17,18} with heterogeneous nodes \cite{19}. But these models had previously been widely considered as curiosities, while only the recent developments have shown their wide range and wide spread applicability.

The range of behaviors found in these models is bewildering. Instead of the continuous (“second order”) transition with standard finite size scaling (FSS) observed in ordinary percolation (OP), one finds everything from infinite order transitions with Kosterlitz-Thouless (KT) type scaling \cite{2} to first order transitions with KT type scaling \cite{14} to first order hybrid transitions \cite{10,12,17,19}, and – last but not least – second order transitions with completely different FSS behavior \cite{20}.

Some of these results were obtained by simulations, others were derived analytically. With very few exceptions \cite{2,3}, the latter rely on the fact that mean field theory becomes exact on random locally tree-like networks \cite{21,23}. The latter allows very elegant treatments based on self-consistency equations derived using message passing arguments, as e.g. demonstrated in \cite{3,17,19} for cooperative percolation and in \cite{24} for several interdependent network models \cite{7,25,26,31}. But these results can be very deceptive – as seen by simulations \cite{12,27} – when applied uncritically to networks which are either not random or not (locally) tree-like. For such networks dynamic message passing \cite{25,31} and the cavity method \cite{31,32} have been applied recently with very promising results, but their applicability to most of the above models is still far from obvious. The fact that network topology can change the critical behavior entirely was demonstrated recently \cite{12} for cooperative spreading agents (so-called syndemics or co-infections). Another warning should be the fact that interdependent two- and three-dimensional lattices do not show the first order transition found on random networks, but show second order transitions \cite{27}, if all links are short range.

All this shows that efficient methods to simulate such models are badly needed. While there exist very efficient methods for OP and for all versions of cooperative percolation, this is not the case for the class of models introduced in \cite{7} and developed further in \cite{24,27,32,43}, where the algorithms used so far in large simulations are extremely slow. After the present work was done, we learned of a recent algorithm \cite{44} that is fast but highly non-trivial, and which has so far not yet been applied to any real problem \cite{45}. It is the purpose of the present paper to present such an algorithm and to use it for several
different network topologies. As we shall see, the results are most disturbing, as the behavior is different for each case. This shows that one should be extremely cautious in applying results obtained mathematically for random locally tree-like models to real world situations.

In deriving the algorithm we use a mapping of the problem onto a solid-on-solid (SOS) type growth model. This mapping is also of interest by itself, as it shows that the model has a number of non-trivial features that might become useful in future analytic treatments.

II. THE MODEL, ITS MAPPING ONTO SOS TYPE GROWTH, AND THE RESULTING ALGORITHM

The models studied in [7] [24, 27, 32, 33] were originally presented as interdependent networks showing failure cascades subsequent to random removal of nodes, but as noted in [24, 27] they are more easily interpreted as single multiplex networks. More precisely, while in general interdependent networks some nodes depend on each other, these dependencies always were assumed in most of these papers to be pairwise and mutual. In this case each pair of interdependent nodes can be identified into a single node, and the network becomes just a duplex network, i.e. one set of nodes connected by two sets of (undirected) links. In some other papers [24, 26, 34] this was generalized to $m$ networks with mutual dependencies among all nodes in clusters of sizes $m \leq m$, where each cluster contains at most one node per network. In this case identification of all nodes in such clusters leads to $m$-plex networks. In the following we shall only consider the case $m = 2$, i.e. duplex networks or pairwise mutual dependencies.

Another slight but important shift was made by [24, 27] who noted that the model can be formulated as a self-consistency problem without any reference to cascades triggered by node removal. Rather, as pointed out in [48], it describes (for $m = 2$) the case where each node needs two essential supplies for being active. More precisely, we assume that there exist a source node that supplies both resources, and that the resources can be transported only on active nodes. It is this latter interpretation which we adopt: We usually consider a cluster $C$ of nodes as viable or, if any two of its nodes are connected by paths on both sets of links, where we also demand that both paths are entirely confined to $C$. In that case, any node in $C$ can be supplied with both resources, if the source node is also in $C$. Conversely, if any node in $C$ can obtain both resources, i.e. if any node in $C$ is doubly connected to the source, then (by the assumed undirectedness of the links) also two arbitrary nodes in $C$ must be doubly connected. Notice that in this way we do not consider the network(s) itself as failed or intact, but we consider only the activities on the networks as present or absent. To use the main example used in [7]: If there is a power failure, the electricity network itself might still be perfectly intact, it is just the activities on the power grid and on the associated computer network that have gone down.

The next difference with the bulk literature on interdependent networks is that we do not primarily delete nodes, but we consider the fate of viable clusters as we change the densities of links. On ER lattices, decimating nodes is just equivalent to renormalizing the connectivity and thus equivalent to decimating links [27]. On regular lattices this is not true, but keeping all nodes just reduces the number of control parameters by one. For two-dimensional (2-d) lattices we checked, however, that decimating both sites and links leads to the same universality class.

A last difference with [7] is that we consider not only the largest viable cluster on the entire network, but we study all viable clusters, in agreement with [44]. As we shall see, this comes with no additional effort. Indeed, only by studying all viable clusters we can verify with our algorithm which one is the largest one. Again we consider this as more realistic. To stay with the electric breakdown example: Assume we have a power failure in central Italy which disconnects the north from the south. This would be considered by [7] as catastrophic, since there would no longer exist a giant viable cluster. But as long as there local viable clusters around Milano and Napoli, people there would be perfectly happy.

In spite of all the differences with [7] mentioned in this section, notice that we are mathematically still dealing with precisely the same class of models, and expect the same type(s) of phase transitions.

Our algorithm works for any type of duplex network with undirected links, although the mapping onto an SOS growth model requires of course strictly spoken that we deal with a planar lattice (since SOS models are models for 2-d surfaces). We shall use therefore a language adequate for this special case, although it should be understood that everything in the following applies also the general case. We thus have a set of $N$ points which are partially connected by two sets of bonds ("red" and "green", say) between neighboring sites. Bonds of either color are placed randomly and independently, with probability $q \in [0, 1]$.

Our algorithm has two ingredients. In the first part we pick a seed (or "source") site and find the largest viable cluster connected to this seed. In the second part we repeat this for all possible seeds.

(a) Finding the largest viable cluster $C$ attached to point $i$: This is simply done by alternatively performing "epidemic" or "Leath-type" spreading processes on the red and green bonds, each starting from site $i$ (this can be done breadth or depth first; we actually used breadth first. We also assume for definiteness that we start with the red bonds). We do not fix bond occupancies "on the run", but we rather determine them before we start with the first epidemic. We follow the spreading until it dies due to the finiteness of the lattice, which gives us a first cluster $C_1$. Since $C_1$ is connected to the seed site but not necessarily doubly connected,
we know that the largest viable cluster attached to site \( i \) must be contained in it, \( C \subseteq C_1 \). Therefore, when we generate the second epidemic using the green bonds, we restrict ourselves to \( C_1 \), generating thereby a cluster \( C_2 \) with \( C \subseteq C_2 \subseteq C_1 \). As we proceed alternatingly, we generate thus a chain of nested sets

\[
C \ldots \subseteq C_h \subseteq C_{h-1} \ldots \subseteq C_1.
\]  

Since the lattice is finite, this must stop at a finite value of \( h \) which we call \( h_i \). It is easy to see that \( C_{h_i} \) is equal to \( C \). If not, then \( C_{h_i} \) would contain at least one point which is not connected to the seed by paths of either color, but in that case \( C_{h_i+1} \) would be strictly larger than \( C_{h_i} \), which is in conflict to our assumption that the iteration stops at \( h_i \).

In the following, we shall call each epidemic starting from \( i \) a “wave”, and \( h \) its height. Since the supports \( C_h \) of successive waves are nested, we obtain in this way a landscape with a single mountain of peak height \( h_i \), and “terraces” of heights \( h < h_i \). Each terrace is just

\[
\Delta C_h \equiv C_h \setminus C_{h+1}. 
\]  

(2)

The part of the lattice which was not even touched by the first wave is called the zeroth terrace with height \( h = 0 \).

(b) Finding viable clusters attached to other points:

After we have obtained the largest cluster attached to \( i \) we could pick another site \( j \) (either randomly or by going systematically through the lattice), and repeat the same process. To find the largest viable cluster on the entire lattice \( C_{\text{max}} \), we would then have to repeat this for all sites. But this would be extremely time-consuming. We are interested in the phase transition where \( C_{\text{max}} \) becomes macroscopic in the limit \( N \to \infty \). This happens at a critical value \( q_c \) of \( q \) which is far above the critical value for single epidemics. Thus for all seeds the first few waves will most likely be huge, and most of the CPU time is spent for finding out clusters which overlap so much that they are nearly identical and cover nearly the entire lattice. The resulting CPU time would then be roughly \( O(N^2) \).

Fortunately, there is no need to use this brute force algorithm. Let us consider the system of waves and terraces for the second seed \( j \). Since the entire lattice is covered by terraces from the first seed \( i \), we can assume that \( j \) is on a terrace of some height \( h \in 0, 1, \ldots h_i \). If the heights were equal to \( h_i \), \( j \) would be part of the first cluster. Thus we can assume that \( h < h_i \). If \( h = 0 \), then \( j \) is not connected to \( i \) by the red bonds, and thus any viable cluster attached to \( j \) must be contained in the zeroth terrace \( \Delta C_0 \). Otherwise (if \( 0 < h < h_i \)) we can use

**Lemma 1:** If \( h > 0 \), the first \( h \) waves starting from \( j \) cover exactly the same clusters \( C_{h'} \) (with \( h' \leq h \)) as the waves starting from \( i \).

**Proof:** The proof is by induction and uses the fact that each wave just covers all sites on a given network that can be reached from the seed, where each of these networks is the subgraph of the original network reached by the previous wave. First of all, it is clear that the lemma holds for \( h = 1 \), because \( C_1 \) is the set of all points on the original network that are connected to the seed by red bonds. Since the two seeds are connected by red bonds due to the assumption that \( h > 0 \), any point \( k \) connected to \( i \) must also be connected to \( j \) and vice versa. Let us now assume that for some \( h' < h \) the clusters attached to \( i \) and \( j \) are the same. Then we can use exactly the same argument, just replacing the original network by \( C_{h'} \) and using the appropriate color of the bonds.

Thus we do not need to follow the first \( h \) waves starting from \( j \), and we can immediately start with wave \( h + 1 \). In doing so, we can use also another important simplification because of

**Lemma 2:** Assume that point \( j \) is on a terrace of height \( h < h_i \). Then the entire next wave is confined to this terrace.

**Proof:** For the proof we have to show that the next wave neither spills over to the lower terrace, nor to the higher.

For the first part we can assume \( h > 0 \). Assume now that the wave actually does spill over, i.e. there exists a point \( k \) which is connected to \( j \) but is on a lower terrace. This point would not be connected to \( i \) on any terrace with \( h' \geq h \), while \( j \) is. This cannot be.

For the second part we assume for the moment that the wave spills over to a higher terrace, i.e. there exists a point \( k \) which is connected to both \( k \) and \( i \). But since \( j \) is not on this higher terrace (i.e. is not connected to \( i \)), this cannot be either.

Thus when building the terraces for site \( j \) we can restrict ourselves to waves for which all boundaries of the terraces generated by the first seed form obstacles which cannot be crossed. It is easy to see that this generalizes also to all later seeds, when the landscape is made up by any number of terraces and boundaries between them:

**Proposition 1:** If a seed \( j \) happens to be on any terrace with a height \( h \) which is not a local maximum (i.e., \( j \) is not in a locally maximal viable cluster), then all previous wave boundaries form obstacles for all avalanches starting at \( j \) which have to be followed in order to get the viable cluster attached to \( j \).

The proof uses the same arguments as above. We just have to realize that successive waves that started from later and later seeds are just distinguished by more and more restricted subsets of the original network to which they are confined. Whatever these subsets are, the above proofs go through without modification.

It should now be clear why we call this procedure an SOS type growth process. As in any SOS model, we have a rough landscape and the growth of this landscape proceeds by localized events, each of which builds a hierarchy of nested terraces. As in surface diffusion problems with strong Schwoebel barriers, these events cannot spill over the boundaries set by previous events.

Notice that neighboring terraces belonging to different seeds can have the same height. Also in this case, their
boundaries cannot be crossed by later waves from other seeds. To implement this restriction we either use an additional marker for each site which tells us the seed to which it belongs; or, alternatively, when each wave is finished, we cut all bonds connecting the wave with its complement. Both methods allow us to prevent the waves from spilling over the terrace boundary, without modifying the dynamics inside any terrace. Also, when we start with a new seed \( j \), we choose the color of the first wave according to the landscape height \( h \) at this seed, and we count subsequent heights by adding to this \( h \). Since now every site \( i \) is “infected” precisely by \( h_i \) waves, it follows that the algorithm has time complexity \( N(h) \), where \( (h) \) is the average height. This estimate holds for arbitrary graphs, provided all node degrees are bounded.

Before we show numerical results, we present also a second proposition which we did not find useful for numerics, but which could be very useful for mathematical treatments.

**Proposition 2:** The final landscape and the system of all terrace boundaries is independent of the sequence by which the seed points are chosen.

Thus the landscape is not a property of the realization of the algorithm (which involves an arbitrary choice of going through all seeds), but is an inherent property of the network. In essence, it says that the growth of the surface is Abelian in a similar way as the Bak-Tang-Wiesenfeld sand pile model is Abelian \[49\].

For the proof we can use the fact that any permutation of sites can be written as a product of pairwise transpositions. We thus need to show only that it does not matter whether we first take seed \( i \) and then seed \( j \) or the inverse. But this follows from the arguments in the proofs of Lemma 1 and 2.

We should add that we checked both propositions also numerically, finding perfect agreement.

Finally, we should point out how our sequences of waves are related to the failure cascades of \[7\] and to other algorithms proposed for them. One essential difference is that we do not demand that the cluster surviving the cascade is always the maximal one. Rather, the maximal cluster is determined at the end, when all viable clusters are known. This simplifies the algorithm of course considerably. Also the complexity of the algorithm of \[13\] is related to the fact that they always follow the largest viable cluster. This precaution is not taken in the algorithm of \[92\], where it is assumed that a cluster which starts large at the beginning of the cascade is not overtaken later by one which starts smaller. This is true for sparse tree-like graphs (whence their algorithm gives correct results), but it is not true in general. Related to this is the absence of the notion of ‘seed’ or ‘source’ in these algorithms. While in our algorithm each ‘cascade’ gives just one cluster arbitrarily picked by the seed, the cascades studied in \[7\] are constructed so as to lead always to the largest cluster. As a consequence, the average cascade length in \[7\] is in our interpretation essentially the average height of the largest viable cluster, which is in general (but not always?) an upper bound for the average surface height.

A last difference between our algorithm and that of \[7\] is that we always start our ‘cascades’ from the full (undamaged) system, while the cascades of \[7\] are supposed to be triggered by (small and successive?) damages, starting from an already partially damaged system. On the other hand, our algorithm shares with that of \[7\] that it does not deal – in contrast to what is suggested in \[7\] – with any real dynamics. Both algorithms deal with pseudodynamics in a fictitious time, and should not be confused with actual cascading processes going on in real time.

**III. APPLICATIONS**

### A. Erdős-Rényi Networks

We first apply our algorithm to duplex ER networks where both layers have the same average degree \( z = \langle k \rangle \). In this case it is known from \[24, 27\] that the generating function formalism of \[21, 22\] gives the exact threshold and the exact dependency of the order parameter on \( \langle k \rangle \). Thus it can be used to test the accuracy of the algorithm. On the other hand, we shall also use it to find the average height of the landscape and the average height of its maxima (i.e., of the largest clusters). The latter is just the average cascade life time in the interpretation of \[7\]. Since during the cascades the clusters are supercritical, it is not a priori clear whether the mean field theory arguments used in \[7\] to derive a scaling law for this life time are exact. To make a short summary, we will find that the algorithm works perfectly, but the mean field arguments of \[7\] for the life time seem to be to be only approximate. This implies that also the arguments of \[7\] in favor of the percolation threshold value have to be taken with care – although they give the correct result – because they rely on the assumed mean field cascade dynamics.

We consider networks with \( N = 2^m \) nodes, with \( m = 4, \ldots, 26 \). The average degree was varied in the range 1.9 to 3.2. The critical point is known in this case to be at \( z_c = 2.45540748 \ldots \). The number of realizations simulated at \( z_c \) varied from \( \approx 10^8 \) for \( N = 1024 \) to \( > 600 \) for \( N = 2^{26} \).

The density of the giant viable cluster on an infinite ER network is given by the largest solution of the equation \[27\]

\[
F(S) = S - \left(1 - e^{-z S}\right)^2 = 0. \tag{3}
\]

It vanishes for \( z < z_c \), and has both a jump and a square root singularity at \( z = z_c \),

\[
S = 0.511700 + a \sqrt{z - z_c} + \ldots \quad \text{for} \quad z > z_c. \tag{4}
\]
For finite $N$ there are of course corrections, the analytic form of which is not known.

Numerical results for $S_N$, the average relative size of the giant cluster on a graph of size $N$, are given in Figs. 1 to 3. Actually, in these plots we show (in contrast to analogous plots in the next subsections) $S_N$ conditioned on realizations which do have a giant cluster. Notice that all other clusters are extremely small (for $N > 2^{20}$ we found only clusters of sizes 1 or 2 apart from the giant one, and even for $N = 1024$ the largest non-giant clusters had sizes < 8), thus it is straightforward to identify giant clusters, except for $N < 10^3$. In Fig. 1 we show $S_N$ versus $z$. Since this gives a perfect agreement with Eq. (3), except for $z$ extremely close to $z_c$, we plot in Figs. 2 and 3 differences between analytical and numerical results for a more sensitive comparison. From Fig. 2 we see that the finite-$N$ corrections are indeed not monotonic, as one might have suggested from Fig. 1, but they decrease fast with $N$. For $N = 2^{26}$, the deviations from the theory for $N = \infty$ are $< 10^{-4}$ for all $z$ except extremely close to $z_c$, which is just the level of statistical errors. Thus we verified with very high precision that our algorithm gives results in agreement with Eq. (3) not only near the critical point, but also for all $z > z_c$. The agreement at $z = z_c$ is checked in Fig. 3, where the differences $\Delta S_N = S_N - S_\infty$ are plotted against $N$. We see a perfect scaling law

$$\Delta S_N \sim N^{-\alpha} \quad \text{with} \quad \alpha = 0.23(1). \quad (5)$$

To our knowledge there is in theoretical prediction for $\alpha$.

There also seems to exist no prediction for the behavior of $P_N(z)$, the chance that there exists a viable giant cluster on a network of size $N$. In analogy to ordinary percolation we expect that this is a step function in the limit $N = \infty$, i.e. $P_\infty(z) = \Theta(z - z_c)$, but the behavior for finite $N$ is non-trivial. In Fig. 4 we plot $P_N(z)$ against $(z - z_c)N^{1/2}$. We see that all curves become essentially parallel, i.e. we have a data collapse apart from a correction to scaling that leads to a shift of the effective transition point towards higher $z$ as $N$ increases.

“Surface” heights (i.e. average cascade life times) measured in these simulations are discussed in the next figures. In Fig. 5 we show for three different network sizes how the average surface heights, averaged over all nodes, depend on $z$. As expected, they are small for $z < z_c$ and become quickly independent of $N$: For small $z$ there

FIG. 1. (color online) Order parameter (density of giant viable clusters) for ER networks plotted against the average degree $z$. The continuous curve is the prediction of Eq. (3), while the points are from simulations with $N$ ranging from $2^{14}$ to $2^{26}$. For $S < 0.511700$... the theoretical curve is strictly vertical, while it has square-root behavior above.

FIG. 2. (color online) Differences between the data points and the theoretical curve in Fig. 1. To enhance the region $z \approx z_c$, the data are plotted against $z - z_c$ on a logarithmic axis.

FIG. 3. (color online) Differences between the data points for $z = z_c$ and the theoretical value 0.511700 towards which they should converge for $N \to \infty$. We see indeed a straight line on a log-log plot, which leads to Eq. (5).
The most conspicuous feature of Fig. 4 is, however, the sharp peaks developing at $z = z_c$. These peaks are only small clusters to start with, and the cascades terminate quickly. For $z \gg z_c$ there are only small holes in the viable clusters and they also are quickly identified, whence the cascades terminate soon also in this regime. It is less obvious why the heights do not seem to be independent of $N$ as $z$ becomes very large. The most conspicuous feature of Fig. 4 is, however, the sharp peaks developing at $z = z_c$ as $N$ increases. Figures 6 to 8 deal with several aspects of these peaks.

Heights of the peaks (more precisely the average surfaces heights at $z = z_c$) are shown in Fig. 6. In the same figure we also show, in addition to the average surface heights, the (averages of the) surface peaks, i.e. the highest points in the “landscapes”. At least for ER networks (where we have no large viable clusters apart from the giant ones) we expect these peak heights to be precisely the average durations of the cascades studied in [7] (see also [13]). We see from Fig. 6 that the average heights show a perfect scaling,

$$\langle h \rangle \sim N^\gamma, \quad \gamma = 0.280(1).$$

In contrast, the peak heights in the landscape seem to increase with the same asymptotic power law, but with large finite-$N$ corrections. The latter are presumably the usual bias corrections in extremal properties evaluated over finite domains. One has similar (logarithmic?) corrections e.g. in the average size of the largest cluster in subcritical (ordinary) percolation on finite lattices.

Eq. (6) is in clear contradiction to the result $h_{\text{peak}} \sim N^{1/4}$ obtained in [7] by mean field arguments which do give also the correct Eq. (3) for the cluster size and the exact value of $z_c$. This is rather puzzling. One might try to explain it by noting that the value of $z_c$ depends only on clusters which are barely viable, and thus the local tree-likeness of sparse ER networks should be sufficient – while the cascade dynamics deals always with supercritical clusters. But then it is not clear why Eq. (3) should be correct also for $z > z_c$, which we took great pain to verify. Notice that the violation of the mean field prediction seen in Fig. 5 is not related to the observations in [43]. In that paper it was shown that an apparent violation of mean field theory is observed, if one does not use the true critical value of $p_c$ in the analysis, but an effective one that changes from realization to realization. In our analysis, $p_c$ is always the true critical point. In any case, our value of $\gamma$ disagrees both with the mean field exponent $1/4$ obtained in [7] and with the exponent $1/3$ proposed in [43], which was based on simulations of much smaller systems with much lower statistics.

The “wings” of the peaks seen in Fig. 5 are studied more closely in Fig. 7, where we plotted the heights (on a log scale) against $\log|z - z_c|$. For the subcritical regime $z < z_c$ we clearly see a power law with exponent $1/2$, in perfect agreement with [43] (notice, however, that we could not confirm the other scaling laws suggested in [43]). We did not try to derive this analytically, but doing this should not be difficult. For $z > z_c$ the situation is less clear, but it seems that for large $N$ a power law with exponent $1/3$ emerges, even if corrections are large for finite $N$.

The two panels of Fig. 8 show finally two attempts to produce a data collapse for the entire critical region. In panel (a) we plotted $\langle h \rangle/N^\gamma$ against $(z - z_c)N^{3/2}$. This gives us a perfect data collapse subcritically. In the critical and supercritical regions the plot does not look too bad either, but a closer inspection shows that it is indeed unacceptable. In contrast, in panel (b) we plotted the data against $(z - z_c)N^{0.5}$. This time the collapse is very poor in the wings, but we get a very good collapse in the critical region – apart from the same shift of the
ER networks, \( <k> = k_c \)

average max. peak height
average surface height

FIG. 6. (color online) Average “surface” heights at \( z = z_c \) plotted against networks network sizes on a log-log plot, and average values of the peaks in each surface. Both curves clearly disagree with the scaling \( \propto N^{1/4} \) proposed in [7], but give rather an exponent 0.280(1).

\[ \langle h \rangle \propto N^{\gamma} \]

\[ |<k> - k_c| N^{\gamma} \]

B. 2-dimensional Lattices

We next study 2-d square lattices of size \( L \times L \), with \( 32 \leq L \leq 32768 \) and with helical boundary conditions. The two sets of links are different (randomly chosen) subsets of nearest neighbor bonds. For each one of them we include a bond with probability \( p \) and exclude it with probability \( 1 - p \). In addition, we can also make a fraction \( q \) of sites inaccessible. For \( p = 1 \) and \( 0 < q < 1 \) the red and green links coincide, and we simply deal with site percolation with control parameter \( q \). On the other hand, if \( p < 1 \) and \( q = 1 \), the sets of red and green links overlap. Each one of them would define a bond percolation problem, and the viable clusters are subsets of intersections of red and green percolation clusters.

For \( p < 1 \) and \( q < 1 \), the single color problems would correspond to mixed site-bond percolation. We studied also that case, because the problem was originally formulated in [7] as a robustness problem for network breakdown under successive node removal. In that interpretation one needs \( p < 1 \) for having a multiplex network, and \( q < 1 \) due to node removal. Actually, however, the problem just becomes more complicated by having both \( p \) and \( q \), without adding any conceptual advantage. If one wants to interpret results obtained with \( q = 1 \) (i.e. effective critical point also seen in Fig. 4. Notice that the variables on the x-axis in Figs. 4 and 8 are the same.

FIG. 7. (color online) Log-log plots of \( \langle h \rangle \) against \( |z - z_c| \). The straight lines indicate power laws which we claim to hold in the limit \( N \to \infty \).

FIG. 8. (color online) Data collapse plots of re-scaled surface heights \( \langle h \rangle / N^\gamma \) against \( (z - z_c) N^{2\gamma} \) with \( 2\gamma = 0.56 \) (panel a) and against \( (z - z_c) N^{\gamma} \) (panel b). Neither plot is fully satisfactory, but in spite of the bigger overall discrepancies we claim that panel (b) gives the correct collapse in the critical region.
with an undiluted lattice) in terms of robustness against damage, then one simply has to implement the damage not as site removal, but as bond removal.

In the following we shall present results for \( q = 1 \) and varying \( p \) (because this is conceptually the simplest), and for \( p = 0.6 \) and varying \( q \). The latter was done in order to compare with the results of [27, 50, 51]. It leads to \( q_c \approx 0.96 \). In neither case the exact critical point is known. The total number of realizations were in both cases between \( > 10^8 \) for the smallest, and several thousand for the largest lattices.

Results for \( p = 0.6 \) are shown in Figs. 9 to 11. In Fig. 9 we show a data collapse similar to the corresponding plots in [27, 50]. We see a perfect collapse, if we use \( q_c = 0.960512(4) \) and \( \nu = 1.2 \). These values agree roughly with those proposed in [27], but are very different from those in [50]. In particular, we would obtain a very bad collapse (for any value of \( \nu \)), if we would use \( q_c = 0.9609 \) as proposed in [50]. Also, we would obtain a very bad collapse (for any value of \( q_c \)), if we would take for \( \nu \) the value 4/3 as in ordinary percolation. We should point out that the largest lattice sizes studied in [50] were only \( 3000 \times 3000 \).

Supercritical data are plotted as \( S \) against \( q - q_c \), on a log-log plot. For large \( L \) and \( q \) not too close to \( q_c \) we expect a power law

\[
S \sim (q - q_c)^\beta.
\]

This is indeed nicely confirmed, with \( \beta = 0.163(2) \). Notice that this is perfectly consistent with the scaling relation \( D = 2 - \beta/\nu \), but is again different from the value for OP. Notice also that the power law holds only for \( q - q_c < 0.01 \). For larger distances from the critical point, corrections would be needed.

Finally, we show in Fig. 11 a log-log plot of \( S \) against \( L \). To make the plot more significant, we actually divide out that the largest lattice sizes studied in [50] were only \( 3000 \times 3000 \).

Essentially the same results were obtained when we kept \( q = 1 \) fixed (i.e., no sites were deleted), and \( p \) was allowed to vary. This time we studied only lattices with sizes up to \( 16384 \times 16384 \), but with higher statistics. The critical point is now \( p_c = 0.576132(5) \). All critical expo-
nents are compatible with these given above, and have roughly the same error bars. The main difference with the case $p = 0.6$ is that now corrections in the far supercritical region do not make the curves in the log-log plot of $S$ against $p - p_c$ turn up (as in Fig. 10), but make then turn slightly down. To avoid duplication we do not show the same plots as for $p = 0.6$, but we do show a data collapse plot for the variance (“susceptibility”) of $S$.

Average landscape heights are shown in Fig. 13. We see a behavior rather different from that for ER graphs. While the latter showed a very sharp peak at the critical point, whose height and sharpness both increased with powers of the system size, we have now a very wide bump far in the supercritical region, whose width does not seem to depend on $L$ at all, and whose height increases logarithmically with $L$. The most conspicuous feature of Fig. 13 is that the slopes at $p = p_c$ become increasingly steeper with $L$, roughly as

$$\frac{d}{dp} \langle h \rangle \sim L^{0.7} \text{ at } p = p_c. \quad (8)$$

In contrast to the ER case, where we could find a decent scaling behavior of $\langle h \rangle$ near the critical point, we were unable to find any convincing scaling in the present case.

Before leaving this subsection, we should mention that all scaling laws and critical exponents found in this subsection were also found in a very different microscopic realization of the 2-d lattice model discussed in subsection III.F.

C. 3-dimensional Lattices

For lattices with $d > 2$ we made only simulations without site decimations, i.e. we always used $q = 1$.

In [27], simulations of very small lattices had suggested that the transition is also for 3-dimensional lattices continuous and not in the OP universality class. This is important, since it could be argued that the transition is continuous for $d = 2$ because there the two sets of links necessarily overlap strongly, and in that case the transition can be continuous even on ER networks [25]. In three dimensions, this overlap is much reduced.

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much smaller than found in [27]. In Fig. 14 we show a log-log plot of rescaled infinite cluster sizes against $L$ for eight values of $p$ close to $p_c$. Here $L$ is defined as $L = N^{1/3}$, where $N$ is the number of lattice sites. Since we used helical b.c., where $N$ was always a power of 2, the so defined $L$ is not always an integer, and the lattices are not always exact cubes. This has however no noticeable effect on the scaling. Since we want to compare with OP, where $L^d S \sim L^{D_{OP}}$ with $D_{OP} = 2.523(1)$ [52, 53], we multiply the data with the corresponding power of $L$.

Thus we expect one curve (the critical one) to be horizontal, if and only if the present model is in the universality class of OP. We see that the most straight curve has a slightly negative slope. This gives $p_c = 0.356707(4)$ and $D_f = 2.475(8)$ (the estimate of [27] was 2.40(1)), which would nominally indicate that the present model is in a different universality class. As we shall see, however, there are important corrections to scaling. Thus the error in this estimate may be severely underestimated.

A reason for it to be not in the OP universality class is, of course, the divergence of the surface height when $L \to \infty$. This is clearly seen from Fig. 16, where we find the same logarithmic increase as in $d = 2$, and a power law for the slope at $p = p_c$ as in Eq. (8), but with exponent $\approx 0.82$.

D. 4-dimensional Lattices

As a first result we show the average surface heights, see Fig. 17. Their behavior is very similar to those in two and three dimensions: For any $N = L^d$ their maxima occur in the supercritical region, and their heights increase logarithmically with $L$. Also, the slopes $d(\bar{h})/dp$ diverge

\[ \beta = (d - D_f)\nu \approx 0.42, \]

in agreement with OP. In summary it seems that corrections to scaling were severely misjudged in [27], and we cannot exclude that the model is in the OP universality class for $d = 3$. 

FIG. 15. (color online). Data collapse plot for the size of the largest viable cluster on 3-d lattices. In contrast to the 2-d case we now have huge corrections to scaling in the supercritical region.

These corrections to scaling are most clearly seen in a data collapse plot, see Fig. 15. In contrast to the analogous plot for $d = 2$ in Fig. 9, we see now huge deviations in the supercritical region. Nevertheless, in the critical and under critical regions the collapse seems to be very good. Notice that we used here the values for $D_f$ and $p_c$ obtained from the previous plot. For the correlation length exponent we obtain $\nu = 0.80(3)$, which is significantly lower than the value 0.8734 obtained from OP [53]. Hyper scaling finally gives $\beta = (d - D_f)\nu \approx 0.42$, in agreement with OP. In summary it seems that corrections to scaling were severely misjudged in [27], and we cannot exclude that the model is in the OP universality class for $d = 3$.

FIG. 16. (color online) Average SOS landscape heights for 3-d lattices, plotted against $p$.

FIG. 17. (color online) Average SOS landscape heights for 4-d lattices, plotted against $p$. 

As a first result we show the average surface heights, see Fig. 17. Their behavior is very similar to those in two and three dimensions: For any $N = L^d$ their maxima occur in the supercritical region, and their heights increase logarithmically with $L$. Also, the slopes $d(\bar{h})/dp$ diverge.
for $L \to \infty$ with a power law, again with power slightly smaller than one (our best estimate is $\approx 0.9$).

In contrast to this, the behavior of the order parameter is completely different. In Fig. 18 we show plots of $S$ versus $p$, for lattice sizes with $N = 2^{16}, 2^{18}, \ldots 2^{28}$, corresponding to $L = 16, \ldots 128$. We see clearly a continuous transition, but with order parameter equal to (or at least very close to) $\beta = 1$. This would indicate that $d = 4$ is the upper critical dimension, in striking contrast to OP, where $d_u = 6$. An attempt to collapse the data according to standard FSS leads to Fig. 19, and to $D_f \approx 8/3$ and $\nu \approx 3/4$. These would not correspond to any known mean field theory, but we should be aware that we should expect logarithmic corrections, if indeed $d_u = 4$. The poor quality of the collapse might indicate that such corrections are indeed present, but with our present data we have no chance to estimate them in detail.

Finally, we show in Fig. 20 the integrated mass distribution on a large lattice for $p \approx p_c$. If usual FSS would hold, we would expect a power law

$$p(m) \sim m^{1-\tau}$$

with

$$\tau = 1 + d/D_f.$$  \hfill (10)

Inserting here our numerical result $D_f = 8/3$ gives $\tau = 2.5$. The data show a very rough power law, with large deviations at intermediate mass values, but the distribution agrees surprisingly well with Eq. (10) for large masses.

\begin{figure}[h]
\centering
\includegraphics[width=0.4\textwidth]{fig18}
\caption{(color online) Average density of the largest viable cluster on 4-d lattices with $N$ sites, plotted against $p$. The sharpness of the kink at $p \approx 0.2715$ increases with $N$.}
\end{figure}

\begin{figure}[h]
\centering
\includegraphics[width=0.4\textwidth]{fig19}
\caption{(color online) The same data as in Fig. 18, but plotted such that the collapse onto a single curve near the critical point.}
\end{figure}

\begin{figure}[h]
\centering
\includegraphics[width=0.4\textwidth]{fig20}
\caption{(color online) Log-log plot of the integrated mass distribution of 4-d viable clusters at $L = 128$ and $p = 0.27145$. The straight line has slope $-3/2$ as predicted from standard scaling laws, accepting the exponents found in Fig. 19.}
\end{figure}

\begin{figure}[h]
\centering
\includegraphics[width=0.4\textwidth]{fig21}
\caption{(color online) Log-log plot of the integrated mass distribution on a large lattice for $p \approx p_c$. If usual FSS would hold, we would expect a power law

$$p(m) \sim m^{1-\tau}$$

with

$$\tau = 1 + d/D_f.$$  \hfill (10)

Inserting here our numerical result $D_f = 8/3$ gives $\tau = 2.5$. The data show a very rough power law, with large deviations at intermediate mass values, but the distribution agrees surprisingly well with Eq. (10) for large masses.}

\subsection*{E. 5-dimensional Lattices}

If the present problem represents indeed a new universality class with upper critical dimension $d_u = 4$, then the behavior in $d = 5$ should be essentially the same as for $d \to \infty$, and in particular as on ER graphs. To verify this we show first (in Fig. 21) the order parameter $S$ (the density of the largest viable cluster) as a function of $p$. We see that the curves for different sizes $N$ intersect indeed in a single point, with $p_c = 0.22614(1)$ and $S_c = 0.31(1)$. This jump of $S$ is a clear sign of a discontinuous transition, although the rise of $S$ for $p > p_c$ indicates that the transition is also, as for ER networks, hybrid.
FIG. 21. (color online) Average density $S$ of the largest viable cluster on 5-d lattices with $N$ sites, plotted against $p$. Notice that we averaged here over all runs (even if they did not have a giant viable cluster), while the averages in subsection III.A (e.g. Fig. 1) were done only over runs that did have a giant cluster.

FIG. 22. (color online) Average SOS landscape heights for 5-d lattices, plotted against $p$.

The same conclusion is reached by looking at surface heights (Fig. 22), which display the same sharp peak as for ER networks. Also, the heights of the peaks increase with roughly the same power of the system size ($h_{\text{max}} \sim N^{0.30(2)}$), not logarithmically as on lattices with $d \leq 4$.

On the other hand, the statistics of finite viable clusters is markedly different from that for ER networks. There, apart from the giant viable cluster all other clusters are of size one in the $N \to \infty$ limit [2], and for the system sizes studied in the present work most of them have sizes one or two. On five-dimensional lattices, however, the distribution of finite viable clusters is non-trivial. The average mass of the second-largest clusters peaks for $p_c - p \sim L^{-2}$, and the peak height increases roughly as $L^2$ (both of these exponents are very crude estimates). Moreover, the mass distribution of the finite clusters shows rough scaling, $p(m) \sim m^{-2.4}$ at $p = p_c$ (see Fig. 23).

F. Cross-over to Mean Field Theory via Long Range Links on 2-d Lattices

1. General Remarks: Long Range Connectivity vs. Long Range Dependency

In general, there are two main ways how the anomalous behavior of critical phenomena seen on low dimensional lattices can cross over to mean field theory. One is by increasing the dimension, the second is by making the theory more and more non-local. In the present model we found that the transition via dimension increase is very non-standard, by showing first a transition to a mean field critical point (at $d = 4$), which is then replaced by a first order transition at $d > 4$. In view of this, it is of interest to check what mechanism(s) are at play when one makes the model more and more non-local.

In the interpretation as multiplex networks (where any interdependent node pairs are joined into a single node) adopted in this paper, the most natural way to do this is by increasing the lengths of the (“connectivity”) links. In analogy with standard critical phenomena one would expect that the model stays in the same universality class as long as the characteristic length of the bonds are finite, and crosses over to mean field theory only when this length becomes infinite, e.g. by having power-behaved link length distributions (see, e.g., [14, 54]). One cannot rule out, however, that there exists a tricritical point at
some finite length.

In the original interpretation of [7] as a model with distinct connectivity and dependency links, one can make either of them long ranged, or both. In [35, 37, 42] it is assumed that nodes are located on 2-d lattices with nearest neighbor connectivity links, while dependency links become long ranged. This is presumably not very realistic in most applications: Since dependency links are much more crucial than connectivity links, one would assume that they are the shorter ones in any realistic natural network. In the following we shall just make some comments on the model of [35, 37, 42], and leave the more natural model of multiplex networks with long range connectivity links to a future publication.

2. Random Dependency Links

As in [35], we first discuss the case where the dependency links are completely random (while, as we said, the connectivity links are the nearest neighbor bonds). The transition is obtained by varying the site occupancy $q$ of one of the two dependency partners (i.e., any pair of dependency partners is present with probability $q$), and keeping all dependency links (i.e., $p = 1$). This is the $r \rightarrow \infty$ limit of the model discussed in the next subsection. Following cascades, it was shown in [35] that the transition is first order in this limit, and closed formulas were given for the threshold $q_c$ and for the density of the viable cluster at threshold.

As in the case of random locally tree-like graphs [24, 27], these formulas are indeed obtained more easily from consistency considerations without reference to any cascade dynamics. Let us denote by $S_{\text{OP}}(p)$ the order parameter (i.e. the density of the infinite cluster) in ordinary site percolation with site occupancy $p$ (the subscript “OP” stands for ‘ordinary percolation’). Let us now consider the percolation of one type of nodes, say of nodes of type A. Since dependencies are completely random, they are not correlated with the (non-trivial) order parameter fluctuations, and they just mean that a finite density of A nodes will be killed by not having a viable dependency partner. This fraction is exactly the same as the fraction of B nodes that are killed because they have no viable A partner. As a result, starting with occupancy $p$ in ordinary site percolation leads to the same order parameter as starting with occupancy $q$ in the dependency model, provided that [35]

$$q = p \times p / S_{\text{OP}}(p),$$

and $S_{\text{OP}}(p)$ is also the density of the viable cluster when starting with $q$. Plotting the r.h.s. of Eq. (12) versus $p$ gives a convex function (see Fig. 24), the minimum of which is the critical value of $q$,

$$q_c = \min_p p^2 / S_{\text{OP}}(p) = 0.682892(5)$$

and $S_c \equiv S_{\text{OP}}(p^*) = 0.6031(2)$ is the density of the giant viable cluster at threshold. The numerical values are in good agreement with the less precise estimates of [35].

![Fig. 24](image)

FIG. 24. (color online) Plot of $p^2 / S_{\text{OP}}(p)$ versus $p$. The points are most precise near the minimum of the curve, where each point is obtained from $\approx 10^5$ lattices of size $32768 \times 32768$. The statistical errors of these points are smaller than their sizes.

![Fig. 25](image)

FIG. 25. (color online) Average density $S$ of the largest viable cluster on 2-d lattices with random dependency links and dependency links between nearest neighbors, plotted against $q$. In the main figure, averages are taken over all runs (even they did not lead to a giant viable cluster, while the inset shows averages restricted to runs with giant viable clusters.

The existence of a first order transition with these parameter values was also confirmed by direct simulations.
In Fig. 25 (main plot) we show the order parameter (the density of the largest cluster, averaged over all runs) plotted against \( q \). Densities obtained by averaging only over those runs which did lead to a giant viable cluster are shown in the inset. As in the ER case, it was trivial to distinguish between runs with and without giant clusters, because all non-giant clusters had sizes 1 or 2 (except on very small lattices, where also clusters of size 3 were seen). Figure 25 confirms not only the values of \( q_c \) and \( S_c \), but it also shows that the transition is hybrid, because the slope at threshold is infinite. The latter is a direct consequence of the fact that the minimum in Fig. 24 is quadratic.

The dependence of the average SOS surface heights on \( q \) and on \( N = L^2 \) is also very similar to that for ER networks. As functions of \( q \) they show peaks that become narrower and higher with increasing \( L \), and the peak values (more precise, the values at \( q = q_c \)) show a power law

\[
\langle h(q_c) \rangle \sim L^{0.556(10)},
\]

(see Fig. 26). This is in qualitative agreement with findings by D. Zhou (quoted in [42]), who obtained however an exponent 0.44, in clear disagreement with our value 0.556. On the other hand, the scaling \( \langle h(q_c) \rangle \sim N^{0.278(5)} \) is the same as that for ER networks (see Eq. (6)).

3. Intermediate-Range Dependency Links

Irrespective of all details, the results of the last subsection show unambiguously that the transition is first order in the limit of infinitely long dependency links and nearest neighbor connectivity links, and the transition occurs roughly at \( q \leq 0.7 \). This should be remembered if we now consider, following [35,37], dependency links with intermediate range.

In [35,37] the dependency links were random vectors \( \mathbf{x} = (x,y) \) with \( ||x|| < r \) in the maximum norm. It was found that the model stays second order, as long as \( r < 8 \). At \( r = r^* \approx 8 \) the transition becomes first order, and for \( r > 8 \) there are claimed to be two first order transitions with a metastable phase in between. The percolation is again implemented as site percolation, i.e. in one of the two lattices only a fraction \( q \) of nodes is open, while all connectivity bonds and all sites in the other lattice are open.

We verified that there is an abrupt change of the transition type at \( r = r^* \), although we found \( r^* \approx 7 \) instead of \( r^* \approx 8 \), see Fig. 27. The most conspicuous feature in the range \( 1 \leq r < r^* \) is a dramatic increase of the surface heights. For all \( r \) the critical points are at those values of \( q \) where \( \langle h \rangle \) changes fastest, which is to the left of the peaks for \( r < 7 \) and to its right for \( r = 8 \). For all \( r < 7 \) we found that the transition is not in the OP universality class, but in the universality class discussed in subsection III.B. This is most clearly seen for \( r = 1 \), where simulations are fastest (due to the smallness of \( \langle h \rangle \)) and where also corrections to scaling are smallest. One might have anticipated that there are large scaling corrections for \( r = 1 \) because of the nearness to \( r = 0 \) where the model would be just ordinary site percolation, but this is not true. Indeed, we found critical exponents in perfect agreement with the estimates of subsection III.B,
and with even somewhat smaller statistical errors. We do not show any plots as they are so similar to those in subsection III.B.

For $r > r^*$ the transitions look very much like first order transitions, in agreement with the claims in \[35, 37, 42\]. If so, then the transition should be tricritical exactly at $r^*$, but we were not able to measure tricritical exponents. As discussed in \[35, 37, 42\], what happens at $r = r^*$ is that fronts become unstable. Consider a scenario where open sites exist only in half-space $x > 0$, while all sites with $x < 0$ are closed. In ordinary percolation this would correspond just to a boundary where all clusters are cut off at $x < 0$. This case of percolation near the boundary is reduced and is governed by a new critical index, but otherwise the boundary has no effect on the bulk behavior.

This is still true for $r < r^*$, but not for $r > r^*$. Although we disagree with \[35, 37, 42\] on details and on the theoretical treatment, we agree with them that nodes near the boundary are removed from the viable cluster by dependencies, which leads then to more removals, etc., so that such boundaries create moving fronts which finally destroy the entire viable cluster – unless $q$ is large and $r$ is sufficiently close to $r^*$. If the front established in this latter case has a sharp profile and the density behind it is finite (which is clearly suggested by the simulations), one has a first order transition.

This gives thus indeed rise to the “upper” of the two claimed transitions for $r > r^*$ (where $q_c(r) > q_c(r^*) \approx 0.74$) \[35, 37, 42\] – with an important caveat discussed below. But we found no indication for the lower transition, where $q_c$ is supposed to be lower than the critical value at $r^*$. In Fig. 28 we show the order parameter for $r = 12$ (i.e., well above $r^*$) for system sizes between $L = 128$ and 8192. According to \[35, 42\], there should be a first order phase transition at $q = 0.74$. We see that there is a rather rapid cross-over at $q = 0.74$, if $L \approx 2000$ (which is presumably the size on which the claim of \[35, 42\] was based), but we can definitely rule out a real phase transition within the range $0.68 < q < 0.765$. Indeed, extrapolating to larger $L$ we can be rather sure that no transition occurs for any $q < 0.8$. Notice that $q_c = 0.753(1)$ for $r = 8$ (data not shown), so the transition point is definitely increasing with increasing $r$. We cannot say numerically whether it agrees with the transition defined via the propagation of fronts (is at $q \approx 0.83$ for $r = 12$), but this seems to be the only plausible option.

Analogous results were obtained for $r = 20$ and $r = 30$ (data not shown). Again we find rapid but nevertheless smooth crossovers at values of $q$ which increase rapidly with $L$. In both cases the cross-over happens at the “transition points” seen in \[35, 42\] when $L \approx 10^3$, but we can rule out real phase transitions anywhere near these points.

![FIG. 28. (color online) Densities of the largest viable cluster for the model of \[35\] with $r = 12$, plotted against $q$ for various system sizes. Here, averages are taken over all runs, including those with no giant viable cluster.](image)

![FIG. 29. (color online) Densities of the largest viable cluster for the model of \[35\] with $r = 12$ (same data as in Fig. 27), but conditioned on clusters which do have a giant viable cluster.](image)

Further insight is obtained by plotting not the order parameter averaged over all runs, but by conditioning onto runs with a giant cluster. Again it is very easy to distinguish between runs with and without giant clusters (although the distinction is not as sharp as for $r = \infty$). The results for $r = 12$ are plotted in Fig. 29. The perfect data collapse shows that the difference between the curves for different $L$ seen in Fig. 28 is entirely due to differences in the probabilities $P_l$ with which a giant cluster is reached. The structures of the giant clusters themselves are completely independent of $L$. In Fig. 30, these data are plotted together with analogous results for $r = 20$ and $r = 30$, and with the results for $r = \infty$ (obtained in the previous sub-subsection; see the inset in Fig. 25) extrapolated to $L \to \infty$. All four curves seem to...
become identical for large \( q \), while they fan out systematically for small \( q \). This figure suggests that there would be very little dependence on \( \mathcal{R} \) (except for very small \( q \)), if there were not a mechanism which would dramatically affect the probability for a giant viable cluster to occur.

\[
1 - P_\infty \sim c(q)L^2,
\]

suggesting that giant cluster formation is prevented by a rare extensive mechanism, i.e. by a mechanism whose probability to arise is \( \propto L^2 \). This is of course true for the formation of voids in the initial configuration. We also see that these probabilities decrease, for fixed \( \mathcal{L} \), roughly exponentially with \( q \). This is again what we would expect for the random formation of voids, with

\[
p(m) \sim \frac{L^2}{m} (1 - q)^m
\]

being roughly the probability to form a void of \( m \) sites.

Numerical estimates of void sizes based on this estimate and on Fig. 31 give void sizes which are typically half as large as those quotes in \[35\]. This is not unreasonable, given the fact that voids were assumed in \[35\] to be circular disks, while they could indeed have different shapes.

Thus we do agree with basic features proposed in \[35, 37, 42\], but we do not agree with the existence of a second phase transition curve. We also would not call “metastable” the states for \( q \) slightly above this supposed transition curve, since metastability applies to systems subject to stochastic (not frozen) noise, and refers to states which are actually unstable on long time scales. In the present case we would rather speak of “conditional stability”, since the viable clusters are absolutely stable, but they arise only conditioned on particular initial configurations.

4. A Final Remark on First Order Transitions and an Alternative Scenario

As we said, the data suggest for \( r > r^* \) a first order transition, and the location of this transition seems to be on the curve given in \[35, 37, 42\]. Although we have no direct numerical evidence to doubt this, there is indirect evidence and theoretical arguments. As suggested in \[35, 37, 42\], the transition for \( r > r^* \) is essentially a percolation transition, where the spreading of voids induced by dependencies becomes critical. In \( d \geq 3 \) there exist indeed tricritical points in generalized percolation models where the phase transition changes from second to first order \[9, 10\]. There is however strong evidence \[10, 57, 58\] that such tricritical points are absent in \( d = 2 \). This seems related to the Aizenman-Wehr theorem \[59\] on absence of first order phase transitions in random 2-d systems, although details are far from clear. Physically, it corresponds to the fact that phase boundaries cannot stay flat in 1+1-dimensional isotropic random systems, but become crumpled on large scales. This is e.g. what is found in the \( T = 0 \) random field Ising model \[57, 58\], and such interfaces seem always to be in the OP universality class, even if they look very smooth on small scales \[59\].

As suggested by Fig. 32, this is not true for simulations on lattices of size \( L_\perp \times L_\parallel \) with \( L_\parallel \gg L_\perp \), and with zero initial densities in the regions \( x < x_- = 20 \) and \( x > x_+ = L_\parallel - 20 \). The latter enforce interfaces which would stay near \( x_\pm \) if \( q > q_c \), but move inwards.

FIG. 30. (color online) Densities of giant viable clusters for \( \mathcal{R} = 12, 20, 30, \) and \( \infty \).

FIG. 31. (color online) Probabilities \( 1 - P_\infty \) that no giant viable cluster is formed, for \( \mathcal{R} = 12 \).
for $q < q_c$. We find, even for $r$ very close to $r^*$, that interfaces at $q \approx q_c$ are very straight, with fluctuations much smaller than $L_\perp$. For larger values of $r$ (data not shown) the effect was much enhanced, and the interfaces were essentially straight lines. This is clearly a finite size effect, which is particularly enhanced by the specific choice of dependency links made in \cite{35,37,42}. By choosing such links to be in a square region with sharp boundaries, fluctuations of the interface are strongly suppressed.

We thus conclude that all simulations of \cite{35,37,42} are far from the true asymptotic region. It might be the fact that the positions of the critical points are nevertheless correct, but they may also be far off. In any case, the fact that the transitions for $r > r^*$ look like first order cannot be taken as evidence that they really are so. Explosive percolation \cite{6} should be a warning that not all transitions that look first order also are so. In the present case, the length scale introduced by $r$ should be enough to confuse the picture.

Indeed, we conjecture that the transition is in the universality class of OP not for $r < r^*$ (as claimed in \cite{35}), but for $r > r^*$ – where it just corresponds to ordinary percolation of voids. We have to admit that we do not have direct numerical evidence to support this claim, nor do we see any possibility to obtain evidence in favor or against it in the near future.

### IV. DISCUSSION

The purpose of this paper was twofold: On the one hand, we proposed an efficient algorithm to simulate viable clusters in multiplex networks, and the percolation transitions related to them. This was motivated by the fact that exact results for this problem are obtainable only in mean field theory (i.e. on random locally loopless graphs), and there exist ample indications that such predictions can be very misleading in real applications. Indeed, the algorithm presented in this paper is not only fast, but by mapping the problem onto a solid-on-solid model we were able also to discuss some non-trivial structures which might become important for themselves.

The other aim of the paper was to apply this algorithm in several simple networks and lattices, in order to test previous claims and to understand better the nature of the percolation transition(s) in this model. We found several surprises:

- While the geometric structure of viable clusters on ER networks is indeed as predicted by mean field theory, it seems that the (pseudo-)dynamics of cascades is not. We speculated that this might be because the clusters are supercritical during the cascades, but more work is needed to understand this.

- On finite-dimensional lattices the model is not in the universality class of ordinary percolation, but represents its own new universality class. This is seen most dramatically in four and five dimensions, where OP would show continuous transitions with non-trivial anomalous exponents. Rather, we found a first order transition in $d = 5$, and a continuous transition with $\beta = 1$ in $d = 4$. Thus it seems as if the model has upper critical dimension $d_\text{uc} = 4$, although the true mean field solution has a discontinuous transition. In $d = 2$ the results are less dramatic, but it seems clear that it is not in the OP universality class. In $d = 3$, finally, we cannot make a clear statement because of very strong corrections to scaling.

- While the order parameter exponent is $\beta = 1$ for $d = 4$ as in other models at the upper critical dimension, other exponents like the fractal cluster dimension and the correlation length are different. They seem to be simple rational numbers, but precise estimates are difficult due to strong (logarithmic?) corrections.

- In general, all first order transitions are hybrid. Thus, the order parameter makes on ER networks and on 5-dimensional lattices not only a jump, but it also shows a power law. More surprisingly, in $d = 5$ there seems also to be a non-trivial scaling mass distribution of finite clusters.

As regards geometric networks embedded in low dimensions with semi-local links, we partly confirmed the
plex networks with behavior is an open question. We do not find the double phase transitions claimed there when the lengths of these links are above a (tri-)critical value. We do find that there exists such a (tri-)critical value above which the transition seems to be first order, but we give strong evidence that this is related to very large finite size corrections. More importantly, we claim that the model studied in [35, 37, 42] is very unrealistic in assuming dependency links to be much longer than connectivity links. Whether a model where this relation is reversed shows similar behavior is an open question.

Other open questions concern the behavior of multiplex networks with > 2 types of links. We have not yet tried to extend our algorithm to this case. Even less clear is the behavior in case of non-mutual (asymmetric) dependencies, which cannot be mapped onto multiplex networks at all. The same is true for directed multiplex networks. Finally, the true asymptotic behavior in the 2-d case with long range dependency links could, eventually, only be solved analytically or by means of a different model realization which avoids the introduction of a new length scale.

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