Hierarchical aggregation in percolation model

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June 13, 2005

Contents

1 Introduction 2

2 Model 4
   2.1 Dynamics .................................. 4
   2.2 Tree representation of clusters .................. 4
   2.3 Horton-Strahler ranking ........................ 5

3 Mass-rank relationship 6

4 Cluster size distributions 7
   4.1 Observed dynamics of cluster population .......... 7
   4.2 Time-dependent laws for cluster size distribution .... 8
   4.3 Averaged scaling ................................ 9
   4.4 Correction to simple scaling ..................... 9

5 Properties of percolation cluster 10
   5.1 Mass dynamics of percolation cluster ............. 11
   5.2 Rank dynamics of percolation cluster ............. 11
   5.3 Mass-rank relation for percolation cluster ........ 12
   5.4 Rank distribution within percolation cluster ...... 13

6 Tokunaga scaling for side branching 13

7 Discussion 15

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Abstract

There is a growing belief that the complex dynamics of seismicity can be better understood by studying the collective behavior of numerous lithosphere instability sources rather than focusing on the details of each of them. Classical site-percolation is a simple and tractable model which exhibits such important general features of complex systems as criticality and phase transitions of second kind. It also illustrates the mechanism of hierarchical aggregation, which is very important for explaining collective phenomena in material fracture and earthquake nucleation processes. We study the dynamics of a 2D site percolation model on a square lattice using the hierarchical approach introduced by Gabrielov et al., Phys. Rev. E, 60, 5293-5300, 1999. The key elements of the approach are the tree representation of clusters and the Horton-Strahler scheme for cluster ranking. Accordingly, the evolution of percolation model is considered as a hierarchical inverse cascade of cluster aggregation. We analyzed the growth of the percolation cluster and established the time-dependent rank distribution of its subclusters, as well as corresponding laws for its mass, rank, and their relationship. We report several phenomena premonitory to the onset of percolation that complement the traditional power-law increase of the model’s observables. In addition, we have shown that the Tokunaga side-branching constraint uniquely determines the mass-rank relationship for a general aggregation process (not necessarily originated from the percolation model). The results can be used for development and improvement of earthquake prediction techniques.

1 Introduction

Earthquakes pose an intolerable threat to society; yet their complex dynamics can hardly be described by a constitutive set of a few fundamental equations. There is an increasing belief that predictive understanding of the behavior of the lithosphere, which generates earthquakes, might result if the research focus shifts from an increasing number of specific sources of lithospheric instability and deformation to their essential collective behavior [1, 20, 21, 22, 48, 50, 37]. During the last two decades, several general concepts and approaches for studying collective phenomena have been successfully applied to various geophysical problems, particularly to describing the dynamics of seismicity. Among them are the concepts of phase transition of second kind [19, 37] and self-organized criticality (SOC) [4]. The latter applies to systems that are maintained near a critical point: a state with no natural length scale typically reflected in a power-law statistics of a system’s observables (think of the power-law distribution of seismic moments).

Classical site-percolation model presents probably the simplest and best studied system that exhibits criticality and experiences (geometrical) phase transition of second kind [44]. It is widely used as a toy model for spatially distributed stochastic processes, such as diffusion in disordered media, forest fires, gelation, semiconduction, etc. Importantly for our study, percolation model presents a very transparent mechanism of the process of hierarchical aggregation, which is very relevant for the dynamics of seismicity. An
early idea of using hierarchical aggregation for describing essential properties of material fracture and earthquake nucleation appeared in the seminal works of Allegre et al. [2] and Newman and Knopoff [30, 31, 23] and has been actively employed since then [6, 7, 12, 27, 29, 35, 39, 55]. A recent general review of the theory and models of kinetics of irreversible aggregation is given by Leyvraz [24].

The principal novelty facilitating the recent efforts in the hierarchical aggregation research is the use of the Horton-Strahler scheme [16, 45, 34] to assign ranks to the elements of hierarchical structures (cascades), as well as the Tokunaga classification [46, 34], which is used to describe the complex self-similar branching within those structures. The rediscovery of the Horton-Strahler scheme (see Sect. 2.3), which originated in hydrology and has not been well known in physical applications, allowed Gabrielov et al. [14] to formulate an exactly solvable model for steady-state approximation to the aggregation process. Their analytical results were confirmed by extensive simulations of Morein et al. [26], who focused on a steady-state regime in a modified forest-fire model; and Zaliapin et al. [56], who applied the hierarchical description to the classical percolation model. The Horton-Strahler ranking scheme is shown essential for building an empirical description and developing numerical models in various geophysical, biological, and computational applications [3, 14, 26, 34, 47, 53]. Notably, hierarchical ranks are more feasible than masses for observations in practice; they present a convenient “macroscopic” measure of cluster size, and provide a natural alternative to the logarithmic binning commonly used in analysis of modeled and observed processes.

In this study we describe the evolution of a percolation model (and, particularly, the percolation cluster) in terms of hierarchical aggregation of smaller clusters into larger ones using the Horton-Strahler ranking scheme. Our goal is three-fold: First, this contributes to a novel understanding of the percolation phenomenon as a time-dependent hierarchical inverse cascade process [51]. Second, this further validates the analytical modeling approach introduced by Gabrielov et al. [14] and developed by Morein et al. [26] for a steady-state approximation to a general aggregation process. Third, this sets a basis for developing earthquake prediction techniques based on dynamical properties of the inverse cascades. A simple model considered in this work naturally suggests several patterns premonitory to the percolation onset; they could be further tested in this and other models and observations.

Specifically, we describe the evolution of percolation model in terms of an inverse cascade of hierarchical cluster aggregation. We focus on the evolution of the first spanning cluster in the classical site-percolation model, and describe it by tracing the consecutive hierarchical fusion of smaller clusters into larger ones. Noteworthy, we are interested not in a final solution of a percolation state, but in an evolutionary path leading from the juvenile single-particle clusters to a self-similar population of clusters of arbitrary large size (limited by the finiteness of the lattice), the percolation cluster included. Thus we depart from the steady-state assumption of [14, 26, 51] as well as from the asymptotic focus on the percolation onset typical for the classical percolation studies [44].

Many phenomena encountered in the percolation model mimic what we see when
the phase transitions of second kind occur. Note however that these phenomena are of purely geometrical and statistical rather than physical nature. Indeed, the physical percolation theory is largely predicated in this geometrical model and there are many empirical links between them; this is why the percolation model is said to be an example of the geometrical phase transition of the second kind, and why its nomenclature emerges from that of the physical critical phenomena.

2 Model

2.1 Dynamics

We consider the classical 2D site-percolation model [44]. The model dynamics starts with an empty \( L \times L \) square lattice. At each step a particle is dropped into a randomly chosen unoccupied site; thus each site can be either occupied by one and only one particle or empty. Two sites are considered neighbors if they share one side; each site on a square lattice has four neighbors. Cluster is defined as a group of occupied neighbor sites [44]. Time refers to the steps at which particles drop onto the lattice. Since we do not have annihilation of particles, time is formally equivalent to the number of particles on the lattice. It is convenient to normalize time by the lattice size \( L^2 \) so it varies from \( \rho = 0 \) at the start to \( \rho = 1 \) when all sites are occupied. During the system evolution, occupied sites start to aggregate and clusters begin to form. Once a sufficient number of particles is accumulated, a percolation cluster is formed connecting the opposite sides of the lattice vertically and/or horizontally (Fig. 1). In the limit of an infinite lattice, the percolation cluster is formed at \( \rho_c \approx 0.59274606 \) [33], while for a finite lattice the critical time \( \rho_c(L) \) is smaller [44]:

\[
\rho_c(L) = \rho_c - cL^{-3}.
\]  

(1)

2.2 Tree representation of clusters

We represent each cluster by a tree that reflects the time-dependent formation of a cluster (its history), and is a subject for quantitative analysis. Specifically, each one-particle cluster is represented by a trivial tree consisting of a single node. When two clusters merge their trees are also merged by adding a new node (parent) for which they become children (and siblings to each other.) In our model, the coalescence of two or more clusters can only be materialized by adding to the lattice a new particle which will be a neighbor to one or more existing clusters. Figure 2a illustrates the four possible types of coalescence; we call \( k \)-coalescence a situation when a newly dropped particle (marked \( N \) in the figure) is a neighbor to exactly \( k = 1, 2, 3 \) or 4 existing clusters (gray numbered sites). Numerical simulations on a square lattice with \( L = 2000 \) suggest the following relative frequencies \( Q_k \) of \( k \)-coalescences: \( Q_1 \approx 0.628, Q_2 \approx 0.318, Q_3 \approx 0.052, Q_4 \approx 0.002 \). Figures 2b,c illustrate how a tree is formed for different coalescence types. There are two basic situations: When a new particle is a neighbor to only one existing
cluster, it is considered as an individual one-particle cluster that is connected to the existing one. The connecting node of the tree in this case does not correspond to a particle on the lattice (panel b). When a new particle is a neighbor to two, three, or four existing clusters, it is not considered as an individual cluster. Instead, it corresponds to the connecting node in the tree (panel c). Thus, the connecting node in a tree may or may not correspond to a lattice particle depending on the coalescence type. The branching parameter (number of children for a given parent) of a tree for any cluster varies between 2 and 4. Note that both 1- and 2-coalescences result in merging only two clusters; accordingly, most of the observed coalescences (about 95% according to our simulations on L=2000 lattice) involve only two clusters while coalescence of three or four clusters is a rare event.

The consecutive process of tree formation for a simple four-particle cluster is illustrated in Fig. 3. To construct the tree one needs to consider all consecutive coalescences that have formed the cluster, not only its final shape. Therefore, it is clear that the same tree may correspond to clusters of different shape: Figure 4a shows two 11-particle clusters that both correspond to the same tree shown in panel b. Accordingly, working with trees, we unavoidably narrow the information about the cluster population. Notice however that trees capture an excessively larger amount of information than mere cluster masses. Summing up, the time evolution of a cluster is necessary and sufficient to uniquely determine the corresponding tree, while the inverse is not true. Generally, clusters of different shape may correspond to the same tree (see Fig. 5); as well as clusters of the same final shape but with different formation history may correspond to different trees (see Fig. 4). The non-uniqueness of the tree representation of clusters does not affect our results, and it is beyond the scope of this paper. Next, we describe the ranking of clusters, presenting a conventional alternative to the logarithmic binning of cluster masses.

2.3 Horton-Strahler ranking

The appropriate ordering of trees (clusters) is very important for meaningful description and analysis of the model dynamics. The problem of such an ordering is not trivial since the clusters may grow and coalesce in a variety of peculiar ways. An advantageous solution to this problem is given by the Horton-Strahler topological classification of ramified patterns [16, 45, 3, 34] illustrated in Fig. 4b: One assigns ranks to the nodes of a tree, starting from \( r = 1 \) at leaves (clusters consisting of one particle.) When two or more clusters with ranks \( r_i, i = 1, \ldots, n \geq 2 \), merge together, a new cluster is formed with the rank [3]:

\[
r = \begin{cases} 
  r_1 + 1, & \text{if } r_i = r_1 \ \forall \ i = 1, \ldots, n \\
  \max (r_i), & \text{otherwise.}
\end{cases}
\]

The rank of a cluster is that of the root of the corresponding tree. It is also possible to consider an alternative definition of ranks [34]: When at least two clusters with rank \( r \) coalesce, and other coalescing clusters have a lower rank, the rank of a new cluster
becomes \( r + 1 \). Clearly, the two definitions coincide when only two clusters coalesce. The results reported in this paper are independent of a particular definition, since coalescence of more than two clusters (especially of high ranks) is a rare event.

3 Mass-rank relationship

In this section we analyze the average mass \( m_r \) of clusters with a given rank \( r \); this is important for connecting various mass and rank scaling laws. The observed mass-rank distribution of clusters at percolation is shown in Fig. 6; it obeys the exponential relation

\[ m_r = 10^{\gamma(r-1)} = c^{r-1}, \]

with \( \gamma \approx 0.625 \), \( c = 10^\gamma \approx 4.2 \). Our simulations suggest that the mass distribution within a given rank is approximately lognormal (not shown) with the mean given by Eq. (2) and a rank-independent standard deviation. The steady-state simulations of [26] suggest \( c = 4.325 \), which is reasonably close to \( c = 4.2 \) that we observe at percolation.

The exponential relation of Eq. (2) happens to be valid over the entire time interval \( 0 < \rho < \rho_c \); the corresponding dynamics of \( c(\rho) \) is shown in Fig. 7. It grows with time from about 2.0 at the earliest stages to 4.2 at percolation. This growth reflects the fact that clusters become more weighty with time due to coupling with the clusters of lower ranks (which does not change the rank but increases the mass). The growth is not monotonous; it is accompanied by pronounced log-periodic oscillations which are associated with creation of new ranks. The log-periodic oscillations that accompany general power-law increase of observed parameters have been found in many systems including hierarchical models of defect development [35], biased diffusion on random lattices [43], diffusion-limited aggregation (DLA) [40], and others. Log-periodic oscillations can be naturally explained by the Discrete Scale Invariance (DSI) [41], which occurs in a system whose observables scale only for a discrete set of values. A famous example of DSI is given by the Cantor set that possesses a discrete scale symmetry: In order to superimpose its scaled image onto the original, one has to stretch it by the discrete factors \( 3^n, n = 1, 2, \ldots \), not a continuous set of values. The Cantor set and percolation belong to systems with built-in geometrical hierarchy, leading to DSI. In our particular system, ranks take only a countable set of values. Creation of new ranks necessarily disrupt the system in a discontinuous way resulting in the log-periodicity.

Recall that the models of [14, 26] use the “fractal correction” \( \epsilon \) to the cluster shape; this correction affects the rate \( r_{ij} \) of cluster coalescence:

\[ r_{ij} \approx \epsilon^{-|j-i|} L_i L_j, \]

where \( L_i \) is the total boundary size of the clusters of rank \( i \). The correction \( \epsilon \) can be expressed as [14]

\[ \epsilon = \frac{1}{\sqrt{c}} \frac{c - 1}{c - 2}. \]
which, together with results of Fig. 7, shows that in the percolation model the effective value of $\epsilon$ decreases in time passing the Euclidean limit $\epsilon = 1$ [14] at $(\rho_c - \rho) \approx 0.14$ and approaching the steady-state “fractal” $\epsilon \approx 0.68$ [26] at $\rho = \rho_c$. The interval $2 < c \leq 4.2$ observed during $0 < \rho \leq \rho_c$ corresponds to $\infty > \epsilon \geq 0.68$.

4 Cluster size distributions

Time-dependent rank distribution of clusters provides essential information about the model dynamics. Also it presents a direct counterpart of the Gutenberg-Richter law for magnitude distribution in observed seismicity. The study [56] demonstrated that simple substitution of the mass-rank relation (2) into well-known laws for the dynamics of cluster masses [44] does not provide a proper description of the time-dependent rank distribution, suggesting an original non-trivial character of a hierarchical description. In this section we summarize time-dependent laws for the mass- and rank-distributions [44, 56], starting with a look at empirical observations.

4.1 Observed dynamics of cluster population

The dynamics of the total number ($n_r \cdot L^2$) of the clusters of a given rank $r$ is illustrated in Fig. 8 for $r = 5, 6, 7$. The population follows a characteristic bell-shaped trajectory with the peak prior to percolation. One does not observe a steady-state behavior in the cluster dynamics: The population of each rank steadily develops to its peak as a result of merging of the clusters of lower ranks; then it starts decreasing, giving birth to the clusters of higher ranks. As naturally follows from the model definition, the peak of the population of a higher rank comes after the peak of a lower rank. Figure 9 shows the population dynamics for the ranks $1 \leq r \leq 11$ in semilogarithmic scale; here one clearly sees the similarity in the dynamics of different ranks. Note that this figure is remarkably similar to Fig. 7 from [51] that shows the dynamics of clusters with logarithmically binned masses.

Figures 10,11 illustrate the mass and rank distribution for a system with $L = 2000$ at $\rho = 0.29$ (dashed line), and at percolation (dash-dotted line). To smooth out statistical fluctuations, Fig. 10 shows the number of clusters with mass equal to or larger than $m$: $\sum_{m' \geq m} n_m(\rho_c)$. The cluster size distribution is prominently transformed during the model development: its downward bend at the early times is substituted at percolation with an almost linear (in bi- or semi-log scale) shape with a slight upward bend for the largest clusters.
4.2 Time-dependent laws for cluster size distribution

It is known that the time-dependent mass distribution is described by Stauffer’s two-exponent scaling law [42, 44, 25]:

\[ n_m(\rho) \sim m^{-\tau} f_0(z), \quad z = (\rho_c - \rho)m^{1/2} + z_0, \]  

(3)

where \( n_m(\rho) \) is the number of clusters of mass \( m \) per lattice site at instant \( \rho \). The Fisher exponent \( \tau = 187/91 \approx 2.05 \) is universal for 2D systems [11, 44]; the function \( f_0 \) has a bell-shaped form with maximum to the left of percolation; and the shift \( z_0 \) is independent of \( m \).

Considered as a function of \( m \), the two-exponent scaling explains the power law mass distribution at percolation (with \( q_0 = f_0(z_0) \)):

\[ n_m(\rho_c) \sim q_0 m^{-\tau}, \]  

(4)

it also explains the downward bend for \( \rho < \rho_c \), clearly observed in Fig. 10 (dashed line). At the same time, as a function of \( \rho \) it describes the bell-shaped dynamics of clusters with given mass \( m \). Equation (4) suggests the slope \( \tau - 1 \approx 1.05 \) for the cumulative mass distribution, while the observed slope 0.96 is somewhat less than that (see Fig. 10). This is due to the impact of two concurrent phenomena: so-called “deviation from scaling” at small \( m \) [17] and finite-size effects at large \( m \) [25, 17] (see Sect. 4.4).

When we switch to the rank description, the two-exponential scaling is no longer sufficient. As shown in [56], the number \( n_r(\rho) \) of clusters of rank \( r \) at instant \( \rho \) is obeying the three-exponent scaling

\[ n_r(\rho) \sim g_0(z) 10^{-br}, \quad z = (\rho_c - \rho)h(r) + z'_0, \]  

(5)

\[ h(r) = a_1 10^{\sigma_1 r}, \quad z'_0(r) = a_2 10^{-\sigma_2 r} \]  

(6)

with

\[ \sigma_1 \approx 0.23, \quad \sigma_2 \approx 0.03, \quad a_1 \approx 1.54, \quad a_2 \approx 1.43, \]

and function \( g_0 \) being approximately Gaussian:

\[ g_0(z) = \exp \left(-z^2/2\right). \]

The three-exponent scaling (5) is consistent with an exponential approximation to the rank distribution at percolation (with \( p_0 = g_0(z'_0) \))

\[ n_r(\rho_c) \sim p_0 10^{-br}, \quad b \approx 0.62. \]  

(7)

Possible deviations from the pure exponential law at \( \rho < \rho_c \) (clearly observed in Fig. 11) and dynamics of a given rank (see Figs. 8,9) are described by specific form of the functions \( g_0(\cdot), h(\cdot), \) and \( z'_0(\cdot) \). The function \( g_0(z) \) is shown in Fig. 12 where different symbols depict clusters of different ranks. The collapse is obvious, confirming the validity of the three-exponent scaling (5), (6).
In the Stauffer’s scaling (3) for cluster masses, the time renormalization \((\rho_c - \rho)m^{1/2}\) collapses the dynamics of mass \(m\) clusters onto the master curve \(f_0(z - z_0)\) with its only peak shifted by \(z_0\) leftward from percolation; the shift \(z_0\) is mass independent. Similarly, in the scaling (5), (6) for ranks the time renormalization \((\rho_c - \rho)10^{m^r}\) collapses the dynamics of rank \(r\) clusters onto the master curve \(g_0(z - z_0')\), although the shift now is rank dependent and is given by \(10^{\sigma_2r}\).

4.3 Averaged scaling

In applications, it is often impossible to measure the size distribution of system elements at a given time instant. Moreover, sometimes the instantaneous size distribution does not exist at all; this is indeed the case for the systems described by marked point processes widely used to model seismicity, volcano activity, starquakes, etc. [10]. In such situations one uses the averaged measurements. For instance, the famed Gutenberg-Richter law [15, 48, 5] that gives exponential approximation to the size distribution of earthquakes (via their magnitudes) is valid only after appropriate averaging over a suitable spatio-temporal domain. This explains the importance of the question: How do the distributions of Eq. (3), (5) change after temporal averaging?

The integration of (3) over the interval \(0 \leq \rho \leq \rho_c\) gives [56]:

\[
\hat{n}_m \propto m^{-5/2}.
\]

(8)

The validity of (8) is confirmed by the observed averaged mass distribution shown by the solid line in Fig. 10: it retains the power-law form (4) of the distribution at percolation while the slope is increased by \(1/2\) due to averaging.

Similarly, for ranks one obtains [56]:

\[
\hat{n}_r \propto 10^{-\alpha r}, \quad \alpha \approx 0.87.
\]

(9)

Again, the averaged rank distribution retains the exponential form (7) of the distribution at percolation; while its index has increased due to averaging (see Fig. 11).

4.4 Correction to simple scaling

Due to finiteness of the lattice, the results of previous sections require some corrections to exactly match the simulated rank distributions. The pure power and exponential laws in Figs. 10, 11 are just first-order approximations to the observed cluster distributions at percolation. In both cases one sees the downward bending for small clusters and upward bending for large clusters. These are not due to statistical fluctuations. The downward bending for small clusters is explained by “deviations from scaling” [17]: it can be shown analytically that the small clusters do not yet obey the general scaling law of Eqs. (4), (7) which holds only for large enough masses (ranks). The upward bend at large clusters is due to finite-size effects [17, 25]: each large cluster that reaches outside the lattice
boundary is “seen” as a number of smaller clusters, thus creating the upward deviation from the pure power (exponential) law. This phenomenon is especially important when the system is close to percolation and clusters of arbitrary large sizes have already been formed. The appropriate scale corrections for the mass distribution were studied by Hoshen et al. [17] and Margolina et al. [25].

In case of the mass distribution, the corrections to scaling are given by [25]:

\[ n_m(\rho_c) \simeq m^{-\tau} \left( q_0 + q_1 m^{-\Omega} + q_L m^{1/D} L^{-1} \right), \]

where \( \Omega \approx 0.75 \), \( 1/D = 48/91 \) is the universal mean cluster radius exponent, and \( q_0, q_1, q_L \) are independent of \( s \) and \( L \). The first additional term describes the deviation from scaling for small clusters, while the second one is responsible for finite-size effects.

For the rank distribution, the “deviations from scaling” at lower clusters are only observed for \( r = 1 \); while the finite-size effects at large clusters are clearly present for many ranks. Zaliapin et al. [56] proposed the following correction to scaling for the rank distribution:

\[ n_r(\rho_c) \simeq 10^{-br} \left( p_0 + p_L 10^{dr} L^{-1} \right), \quad r > 1. \]

with

\[ d = \frac{1}{D} \log_{10} c \approx 0.33 \]

and showed that the observed ranks \( 4 \leq r \leq 9 \) nicely follow the predicted scaling.

Importantly, the corrections to scaling (11) act at all cluster sizes, so they can not be neglected even for the intermediate clusters, not only for the largest ones. Indeed, their effect decreases with \( L \), but this decrease is very slow. Notably, as shown by Morein et al. [26] (their Fig. 5) even for lattices as large as \( L = 30,000 \) during the process when clusters as large as 2% of the lattice size are removed, the cluster size distribution clearly exhibits the upward deviations at large ranks (\( r = 11, 12, 13 \)). For smaller systems these deviations become dominant and may lead to an artificial decrease of the observed slope of cluster size distribution; this is demonstrated in Fig. 10,11 and is also seen in the analysis of Turcotte et al. [51] (their Fig. 9).

5 Properties of percolation cluster

This section analyzes the growth of the percolation cluster; that is, we consider the time-dependent properties of the cluster that will first span the lattice (unavoidably using the information from the “future”). Such properties, being compared to the average statistics over the entire cluster population, might be useful for general description of the model as well as for improving the observer’s knowledge about the moment of the approaching percolation. Studying the percolation cluster separately from the rest of the model can be considered an analog to concentrating on the region surrounding the epicenter of a large regional earthquake versus the entire region; the latter approach being traditionally and effectively used in earthquake prediction research. We emphasize that studying the
percolation cluster may differ from studying the largest cluster on the grid. Indeed, for large lattices the percolation cluster is almost surely the largest one at the moment of percolation; this equivalence though may not hold at the earlier stages of the model dynamics as well as for intermediate size lattices. Below we use the term “percolation cluster” to denote the cluster that will percolate first in a given model run.

5.1 Mass dynamics of percolation cluster

The time-dependent mass $m_p(\rho)$ of the percolation cluster, averaged over 1,000 runs of the model on $L = 2,000$ grid, is shown in Fig. 13. One can easily distinguish the following three major stages of the mass development: Intermediate dynamics, $0.3 > (\rho_c - \rho) > 0.01$, is well described by the scaling law

$$m_p \approx 5.0 \times (\rho_c - \rho)^{-2.05}.$$  \hspace{1cm} (12)

However, at early times, $(\rho_c - \rho) > 0.3$, as well as in the immediate vicinity of percolation, $(\rho_c - \rho) < 0.01$, the mass dynamics deviates downward from the scaling of Eq. 12. At early times this is due to the “deviations from scaling” [17], while at later times to the finite size effects [17, 25] (See Sect. 4.4). The downward deviation from the power scaling at later times seems a nice signal of the approaching percolation. However such a signal might be undetectable in a particular model run, as illustrated in the insert of Fig. 13: While the overall scaling with exponent 2.05 is clearly observed, the downward deviation prior to percolation is not seen for this particular realization. This problem is very similar to the one typically faced in earthquake prediction studies: many premonitory seismicity patterns (the Benioff strain release [18] being the most popular one) are easily detected by the stacking data from many large earthquakes, but almost unseen in the observed seismicity prior to a single earthquake.

It should be noted though that the direct analog of a power law Benioff strain release—a power law mass increase of the percolation cluster—seems to be nicely detectable even in a single model run (see insert in Fig. 13), and may serve as a basis for predicting the onset of percolation. We leave a detailed treatment of such a prediction technique for a further study, and proceed now with the dynamics of the rank of the percolation cluster.

5.2 Rank dynamics of percolation cluster

We argued above that the ranks are more feasible for observations than masses. This advocates the study of rank dynamics for the percolation cluster. Figure 14 shows the time-dependent rank $r_p(\rho)$ of the percolation cluster, averaged over 1,000 runs of the model on an $L = 2,000$ grid. The following exponential increase is clearly observed at intermediate times to percolation, $0.3 > (\rho_c - \rho) > 0.01$:

$$r_p \approx 2.2 - 3.35 \times \log_{10}(\rho_c - \rho).$$  \hspace{1cm} (13)
As in the mass dynamics, the downward deviations from this law at early times, \((\rho_c - \rho) > 0.3\), and close to percolation, \((\rho_c - \rho) < 0.01\), can be explained by “deviations from scaling” and finite size effects (see Sect. 5.1). Again, the delay in creation of the consecutive rank may serve as a signal of approaching percolation, but this delay would be hard (if possible at all) to detect in a single model run.

Striking is the following observation: The building of the percolation rank is monotonous, it lacks the characteristic log-periodic oscillations observed for the process of rank creation on the entire grid (see Fig. 7 and its discussion).

5.3 Mass-rank relation for percolation cluster

Having established the scaling laws for mass (Eq. 12) and rank (Eq. 13) of the percolation cluster, we now proceed with connecting these laws. One obviously may rewrite them as

\[
(\rho_c - \rho) \approx \left( \frac{m_p}{5.0} \right)^{-1/2.05}, \quad (\rho_c - \rho) \approx 10^{-(r_p - 2.2)/3.35},
\]

which leads to

\[
\left( \frac{m_p}{5.0} \right)^{1/2.05} \approx 10^{(r_p - 2.2)/3.35}
\]

or

\[
\log_{10} m_p \approx \frac{2.05}{3.35} - \left( \log_{10} 5.0 - \frac{2.05 \times 2.2}{3.35} \right) \approx 0.61 \times r_p - 0.65.
\]

The index 0.61 is fairly close to that of the mass-rank relation (2) (with \(\gamma \approx 0.625\)) estimated for the entire cluster population at percolation (see Fig. 6). This prompts one to conjecture that the index of the mass-rank scaling for the percolation cluster matches that of the total cluster population at percolation. This observation, schematically illustrated in Fig. 15, depicts a very special character of the percolation cluster: at any time instant it possesses the information about the index of the mass-rank scaling law that will be observed for the entire cluster population at percolation. Thus, comparison of the mass-rank scalings for the entire cluster population with that of the percolation cluster may serve as another precursor for the percolation onset. Indeed, we do not know in advance which cluster will percolate first, so one may use instead the largest observed cluster. Detailed analysis of the power and usefulness of such a premonitory pattern is beyond the scope of the present study.

We also observe from Fig. 15 that the mass \(m_p\) of the cluster that will percolate first is always less than the average mass of clusters of the same rank at percolation. This observation is supported by Fig. 16 that shows the dynamics of the average mass \(m_r\) of clusters of a given rank. Strikingly, the average mass continues to increase when both the total number \(n_r\) of clusters and their total mass \(M_r\) significantly decreased having passed their respective peak values. In other words, the “fat” members of a rank population still survive when their “thin” companions already have merged together turning into clusters of a higher rank.
5.4 Rank distribution within percolation cluster

Recall that each cluster in our model is an aggregate of smaller ones that merged together at previous time steps. Thus, it is natural to consider the rank distribution of subclusters within the percolation cluster at different stages of its development. This distribution for four time instants: $\rho = 0.29, 0.5, 0.54$, and at percolation is shown in Fig. 17. The rank distribution within the percolation cluster is always approximately exponential (modulo the largest ranks that deviate significantly from the overall pattern):

$$\tilde{m}_r \propto 10^{-0.3r},$$

(14)

where $\tilde{m}_r$ is the number of clusters of rank $r$ within the percolation cluster. Notable is the upward bend that reflects the “critical” character of the percolation cluster (see Sect. 7 for details). Importantly, the distribution is stable in time; there is no transformation from convex to almost linear shape observed for the entire cluster population (see Sect. 4, Fig. 11). The index of the exponential law (14) is twice less than that of the rank distribution for the entire cluster population (see Sect. 4, Eq. 7, Fig. 11). Thus the relative number of larger clusters within the percolation one is higher comparing to that for the entire cluster population.

6 Tokunaga scaling for side branching

In this section we show how the index $c$ of the mass-rank relationship (2) is expressed via the parameters of another important scaling law (Tokunaga scaling [46]) that describes self-similar side branching of clusters. First, following [34, 46] we define the branching ratio $T_{ij}$ for a given cluster (tree) as the number $N_{ij}$ of its subclusters (nodes) of rank $i$ that joined subcluster (node) of rank $j$, averaged over subclusters (nodes) of rank $j$:

$$T_{ij} = \frac{N_{ij}}{N_j}.$$

(15)

Many natural hierarchies are shown to exhibit self-similar side branching, obeying the Tokunaga scaling constraint [34, 46]:

$$T_{i+i+k} = T_k = s_0 s^{k-1}.$$

(15)

Particularly, it was predicted by Gabrielov et al. [14] and later confirmed by simulations [26] that clusters produced by aggregation dynamics obey the Tokunaga scaling (15) asymptotically in $k$.

Moreover, Gabrielov et al. [14] established a relationship between parameters $s_0$, $s$ of the Tokunaga scaling (15) and the mass-rank distribution (2). Specifically, if one assumes clusters of a regular non-fractal shape (Euclidean limit of the model), then

$$s_0 \approx 0.55495813, \quad s = 1/s_0 \approx 1.80193774, \quad c = 1/s_0^2 \approx 3.24697602.$$
Observed systems though are expected to produce fractal clusters (non-Euclidean case), which might change the values of the parameters and their interplay; for example, the steady-state simulations of Morein et al. [26] suggest

\[ s \approx 3.0253, \quad s_0 \approx 0.6993, \quad c \approx 4.325. \]  \hspace{1cm} (17)

The general connection between \( s_0, s \) and \( c \) can be established by noticing that the Tokunaga scaling uniquely determines the mass of rank \( r \) clusters. To express the average mass \( m_r \) via the parameters \( s_0, s \) we notice that the mass of a rank \( r \) cluster is the sum of two \( r - 1 \) cluster masses that formed the cluster (we ignore the possibility for three or more clusters to coalesce at the same step), plus a unit mass of a joining particle, plus the mass of all the lower-rank clusters that joined the considered cluster, hence:

\[
\begin{align*}
m_1 &= 1 \\
m_2 &= (2m_1 + P) + T_{12}(m_1 + P) \\
m_3 &= (2m_2 + 1) + T_{23}(m_2 + 1) + T_{13}(m_1 + P) \\
&\quad \ldots \\
m_k &= (2m_{k-1} + 1) + \sum_{i=1}^{k-1} T_{k-i,k}(m_{k-i} + 1) - (1 - P)T_{1,k}, \ k \geq 3. \hspace{1cm} (18)
\end{align*}
\]

Here the coefficient \( P \) addresses the possibility for a one-particle cluster to join another cluster in two ways: via a one-particle connector (with probability \( P \)) or directly (with probability \( 1 - P \)); the clusters with \( r > 2 \) can only join other clusters using a one-particle connector.

Assuming the Tokunaga scaling (15) we rewrite Eq. (18) for \( k \geq 3 \) as

\[
m_k = (2m_{k-1} + 1) + \sum_{i=1}^{k-1} T_{i}(m_{k-i} + 1) - (1 - P)T_{1,k},
\]

and using the mass-rank relation (2) with \( c > 1 \) obtain

\[
\begin{align*}
c^{k-1} &= 2c^{k-2} + 1 + \sum_{i=1}^{k-1} s_0 s^{i-1} \left[ (c^{k-i-1} + 1) - (1 - P)s_0 s^{k-1} \right] \\
&= c^{k-2} \left[ 2 + \frac{1}{c^{k-2}} + s_0 \frac{1 - (s/c)^{k-1}}{1 - s/c} + \frac{s_0}{c^{k-2}} \frac{s^{k-1} - 1}{s - 1} - (1 - P)s_0 (s/c)^{k-2} \right].
\end{align*}
\]

It is easily checked that this equation has a solution only if \( c > s \); thus \( s/c < 1 \) and for large \( k \) then follows

\[
c^{k-1} = c^{k-2} \left[ 2 + \frac{s_0}{1 - s/c} \right]
\]

leading to the final equation

\[
c^2 - c(2 + s + s_0) + 2s = 0.
\]
A simple analysis shows that this equation always has at least one solution for \( s_0 > 0 \), and the only solution that ensures \( c > s \) is given by

\[
c = \frac{2 + s + s_0 + \sqrt{(2 + s + s_0)^2 - 8s}}{2},
\]  

(19)

while \( c > 1 \) implies \( s > 1 + s_0 \).

For the Euclidean values of \( s_0, s \) (16) the above formula gives \( c(s_0, s) = 3.24697960 \), which is remarkably close (6 digits) to the result of [14]. Furthermore, the non-Euclidean values of (17) solve Eq. (19) exactly. We found it quite convincing that our complimentary set of assumptions used to derive (19) lead to the same numerical results as an analytical study [14] and simulations of [26]. This suggests an internal stability of a system that follows an aggregation dynamics with respect to the details of its particular realization. The relationship (19) provides a striking constraint for the scaling of aggregation processes, and may be useful in working with observed systems.

7 Discussion

This work continues a line of research initiated recently by Turcotte et al. [51], Gabrielov et al. [14], Morein et al. [26], and Zaliapin et al. [56], who have introduced and studied hierarchical inverse cascade description of aggregation (coagulation) phenomena. Notably, the inverse cascade approach is shown to explain self-organized critical behavior [51], while the aggregation phenomena are proven important for evolution of many natural hazardous processes such as earthquakes, landslides, and forest fires [52, 26]; they are also commonly employed for description and modeling of material fracture.

Following [14] we described clusters in 2D site-percolation model on a square lattice by hierarchical trees that reflect the history of cluster formation; the Horton-Strahler scheme was used to rank the trees and thus the corresponding clusters. We concentrated on the development of the first percolation cluster, thus working with a system that does not exhibit the steady-state dynamics, contrary to the studies [14, 26] that have developed mean-field steady-state approximations to the system.

The most important results of percolation theory are traditionally given in terms of cluster masses [44]. Accordingly, in order to bridge the classical results to the novel rank description it is vitally important to establish the mass-rank scaling laws, which was done in [14, 26]. In this study we have analytically expressed the index of the mass-rank exponential relationship (2) via the parameters of the Tokunaga side-branching constraint (15). The predictions and simulations of [14, 26] are found to be in a perfect agreement with this relationship, suggesting the universality of the essential characteristics of the aggregation process, independently of its particular (physical or analytical) realization.

We studied in detail the transition of the system from earlier stages to the vicinity of percolation and reported several characteristic phenomena observed as \( \rho \to \rho_c \). They include transformation of the cluster size distribution not unlike that observed in seismicity, steel samples fracture, and previous models of hierarchical fractures [27, 36, 12, 55].
In our simple model these phenomena are partly explained (qualitatively as well as quantitatively) by finite-size effects; nevertheless we believe that they should not be neglected as irrelevant side-effects of numerical simulation. In fact, in practice we often work with systems that are described by intermediate depth hierarchies (in other words they have an intermediate number of degrees of freedom). The percolation results related to small and intermediate lattices might be of high relevance in describing such systems. In addition, simulations on large lattices ($L = 30,000$) performed by Morein et al. [26] show that finite-size effects are not negligible even for large $L$.

We have studied the growth of the percolation cluster tracing its evolution from the earliest times to the onset of percolation and established scaling laws for mass and rank of the percolation cluster. We also studied the frequency-size distribution of subclusters within the percolation cluster. We have reported several features that may be used for prediction of the onset of percolation: power-law increase of the mass of the percolation cluster (this property is known from the classical studies [44]); exponential increase of the rank of the percolation cluster; downward deviations of the mass- and rank- scalings prior to percolation; and convergence of the index of the mass-rank distribution for the entire cluster population to that within the percolation cluster. Indeed, if we may count precisely the occupied sites, then the onset of percolation is determined by the critical time $\rho_c$, and there is no need in additional premonitory patterns. At the same time, it is not uncommon in prediction research that premonitory patterns whose emergence is obvious in a simple model can be effectively applied for prediction of critical events in more complex systems as well as in observations (e.g., see [13, 27, 28, 36, 38, 55]), so the further analysis of the above patterns seems promising.

Our closing remark is on the index $\tau$ of cluster mass distribution at percolation (Eq. (4)). The studies of Gabrielov et al. [14] and Morein et al. [26] predict $\tau = 2$; which slightly deviates from the well established theoretical value of the Fisher exponent $\tau = 187/91 \approx 2.05$. The index of the mass distribution is an essential characteristic of a system, thus even this slight difference of 2.5% might seem disappointing, implying the intrinsically approximate character of the modeling of [14, 26]. In fact, this implication is not true. To validate the approach of [14, 26] we notice that the Fisher exponent is tightly connected to the precise count of cluster particles on a site-level, hardly feasible in practice. At the same time, the studies [54, 8] have demonstrated that when we “characterize the size distribution of clusters in a way that circumvents the site-level description” considering any “macroscopic measure of the length scale of the cluster”, the exponent of the corresponding scaling law becomes 2, universally for all 2D systems. An example of a “macroscopic measure” is the linear size in an arbitrary direction, the radius of gyration, the diameter of the covering disk, etc. Clearly, the modeling of [14, 26] deals with such a macroscopic measure of cluster size, and hence predicts the correct scaling exponent.

Acknowledgments. We are sincerely grateful to Bill Newman for numerous focused discussions that have helped significantly in organizing and presenting the results. We
thank Kristy Tiampo and two anonymous reviewers for their comments that helped to improve the paper; and Vladimir Keilis-Borok, Gleb Morein, and Donald Turcotte for their continuous interest in the work. This study was partly supported by NSF, grant ATM 0327558.
References


21
Figure 1: Sample percolation model. The percolation cluster is shown in black: it connects the top and bottom sides of the grid. Coincident smaller clusters are shown in gray, empty sites are white.
Figure 2: Multiple coalescence of clusters. a) Coalescence of clusters is materialized by adding to the lattice a new particle $N$ (black) that is a neighbor to one, two, three, or four existing clusters (numbered gray sites). The relative frequencies $Q_k$, $k = 1, 2, 3, 4$ of $k$-coalescences based on simulations with $L = 2,000$ are shown in the figure. The corresponding tree is constructed as shown in panel b) (for $k = 1$) and c) (for $k = 2$). The cases $k = 3, 4$ are analogous to $k = 2$. Note that about 95% of coalescences result in merging two clusters. See text for details.
Figure 3: Tree representation of clusters: scheme. The dynamics is from left to right. At first step particle A is dropped onto the lattice and a one-particle cluster is formed; it is represented by a one-node tree. At second step another one-particle cluster B is formed; it is represented by another one-node tree. At third step new particle C coalesces with cluster A to form two-particle cluster AC. This cluster is represented by a three-node tree; note that the connecting node of the tree does not correspond to any particle. At fourth step new particle D connects existing clusters AC and B to form four-particle cluster ABCD. This cluster correspond to a five-node tree.

Figure 4: a) Non-uniqueness of tree representation. Two different 11-particle clusters that correspond to the same tree shown in panel b). Particles have been dropped according to their alphabet marks; so first was the particle A, then B, etc. b) Horton-Strahler ranking: illustration. The ranks are shown next to the tree nodes.
Figure 5: Non-uniqueness of tree representation. Three clusters with the same final shape but different history of formation are represented by different hierarchical trees. Particles have been dropped according to their alphabet marks; so first was the particle A, then B, etc. The Horton-Strahler ranks are shown next to the tree nodes.
Figure 6: Mass-rank distribution observed on a 2,000×2,000 lattice at percolation. Dots – individual clusters, balls – average mass $m_r$ within a given rank. Line shows the relation $m_r = [10^{0.625}]^{r-1} \approx 4.2^{r-1}$. 

slope = 0.625
Figure 7: Parameter $c$ of the mass-rank relation $m_r = c^{r-1}$ as a function of time. At percolation $c(\rho_c) \approx 4.2$; the Euclidean limit of [14] corresponds to $c = 3.25$, it is reached at $\rho_c - \rho \approx 0.14$.

Figure 8: Dynamics of population $n_r \cdot L^2$ of a given rank, $r = 5, 6, 7$ for $L = 2,000$. Moment of percolation is depicted by a vertical dashed line.
Figure 9: Dynamics of population $n_r$ of a given rank, $1 \leq r \leq 11$ in semilogarithmic scale. Moment of percolation is depicted by a vertical dashed line. (Cf. Fig. 7 from [51]).

Figure 10: Mass distribution of clusters observed on a $2,000 \times 2,000$ lattice at percolation $\rho = \rho_c$ (dash-dotted line), $\rho = 0.48$ (dashed line), and averaged over $0 < \rho < \rho_c$ (solid line). To smooth out statistical fluctuations we show the cumulative distribution: $\propto \sum_{m'>m} n_{m'}$. For comparison, all curves are normalized to unity at $m = 1$. 
Figure 11: Rank distribution of clusters observed for $2,000 \times 2,000$ lattice at percolation $\rho = \rho_c$ (dash-dotted line), $\rho = 0.29$ (dashed line), and averaged over the percolation cycle $0 < \rho < \rho_c$ (solid line). For comparison, all curves are normalized to unity at $r = 1$.

Figure 12: Three-exponent scaling for rank dynamics. The master Gaussians $g_0(z)$ for different ranks collapse when using the renormalization given by Eqs. (5),(6).
Figure 13: Dynamics of the mass of the cluster which will percolate first. Main panel shows the average over 1,000 runs of a model with $L = 2,000$, insert shows the dynamics of a single model run.
Figure 14: Dynamics of the rank of the cluster which will percolate first; average over 1,000 runs of a model with $L = 2,000$. 

slope = 3.35
Figure 15: Mass-rank relationship for the entire cluster population at percolation (top solid line), entire cluster population prior to percolation (dashed line), and the cluster that will percolate first (bottom solid line).

Figure 16: Dynamics of number of clusters $n_r$ (solid line), total mass $M_r$ (dashed line), and average mass $m_r$ (dash-dotted line) for clusters of rank $r = 5$. 
Figure 17: Rank distribution of subclusters within the cluster that will percolate first. For comparison, all curves are normalized to unity at $r = 1$ and arbitrarily shifted downward.