



ELSEVIER

Available online at www.sciencedirect.com

SCIENCE @ DIRECT®

Physica A 334 (2004) 583–590

PHYSICA A

www.elsevier.com/locate/physa

Topological phase transitions of random networks

Imre Derényi*, Illés Farkas, Gergely Palla, Tamás Vicsek

*Department of Biological Physics, Biological Physics Research Group of HAS, Eötvös University,
Pázmány P. stny. 1A, Budapest H-1117, Hungary*

Received 20 October 2003

Abstract

To provide a phenomenological theory for the various interesting transitions in restructuring networks we employ a statistical mechanical approach with detailed balance satisfied for the transitions between topological states. This enables us to establish an equivalence between the equilibrium rewiring problem we consider and the dynamics of a lattice gas on the edge-dual graph of a fully connected network. By assigning energies to the different network topologies and defining the appropriate order parameters, we find a rich variety of topological phase transitions, defined as singular changes in the essential feature(s) of the global connectivity as a function of a parameter playing the role of the temperature. In the “critical point” scale-free networks can be recovered.

© 2003 Elsevier B.V. All rights reserved.

PACS: 89.75.Hc; 05.70.Fh; 64.60.Cn; 87.23.Ge

One of the most fruitful recent approaches to the description of complex systems consisting of many similar units has been the analysis of the underlying network of interactions. It has turned out that seemingly very different systems can be characterized by a few major classes of stochastic graphs representing the overall features of the structure of connections among the units [1,2]. These developments have greatly advanced our potential to interpret the fundamental common features of diverse systems including social groups, technological, biological and other networks. The effects of both the restructuring [3] and the growth [4] of the associated graphs have been considered leading to a number of exciting discoveries about the laws concerning their diameter, clustering and degree distribution.

Although the conceptual basis of most of the related works has largely relied on analogies (scaling, percolation, etc.) with statistical mechanics, only very few recent

* Corresponding author.

works [5–7] have been devoted to the problem of directly connecting the graph theoretical aspects of networks to statistical mechanics or thermodynamics. On the other hand, using a thermodynamic formalism for the changes in graphs being in an equilibrium-like state is expected to provide a significantly deeper insight into the processes taking place in systems being in a saturated state and, as such, dominated by the fluctuating rearrangements of links between their units.

As an example, let us take a given number of units interacting in a “noisy” environment. These units can be people, firms, genes, etc. The probability for establishing a new or ceasing an existing interaction/connection between two units depends on both the noise and the advantage/disadvantage gained/lost when adopting the new configuration. In this picture, a global transition in the connectivity properties can occur as a function of the level of perturbations. For instance, if the conditions are such that the interactions between the partners become more “conservative” (a relative, short-term gain is more highly valued), as we show later, a transition from a less ordered to a more ordered network configuration can take place. In particular, it has been argued [8] that depending on the level of given types of uncertainties (expected fluctuations) business networks reorganize from a star-like topology to a system of more cohesive, highly clustered ties.

In this paper we use temperature to represent noise and differences in an energy (potential) type quantity to account for advantage or loss during the rearrangement of a network. This enables us to treat the associated microscopic dynamics within the framework of *canonical ensembles* and map the problem onto a *lattice gas* model. After defining an order parameter and a variety of possible potentials, interesting *topological phase transitions (first and second order)* are found both analytically and by numerical simulations. Scale-free networks are recovered at the transition point between a system consisting of many nodes with few links to a system dominated by a single large hub.

We define the partition function

$$Z(T) = \sum_{\{g_a\}} e^{-E_a/T} \quad (1)$$

for an ensemble $\{g_a\}$ of undirected graphs containing N nodes and M links, where E_a denotes the energy of the graph g_a . Our choice for Z is in analogy with that proposed by Berg and Lässig [6] and is motivated by the following physical picture behind the graph restructuring process: the basic event of the rearrangement is the relocation of a randomly selected edge (link) to a new position either by “diffusion” (keeping one end of the edge fixed and connecting the other one with a new node) or by removing the given edge and connecting two randomly selected nodes. Then, the energy difference $\Delta E_{ab} = E_b - E_a$ between the original g_a and the new g_b configurations is calculated and the relocation is carried out following the Metropolis algorithm. The resulting dynamics, by construction, satisfies the detailed balance condition.

This network rearrangement is formally equivalent to a *Kawasaki type lattice gas* dynamics with conserved number of particles moving on a special lattice, which is the edge-dual graph of the fully connected network [9,10]. The sites of this lattice are the

possible $N(N - 1)/2$ connections between the vertices, and the particles wandering on the sites are the M edges.

To be able to monitor topological phase transitions a suitable *order parameter* Φ has to be introduced. As we are primarily interested in the transitions between dispersed and compact states, a natural choice can be either $\Phi = \Phi_s = s_{\max}/M$, the number of edges of the largest connected component of the graph s_{\max} normalized by the total number of edges M , or $\Phi = \Phi_k = k_{\max}/M$, the highest degree in the graph k_{\max} divided by M . We also introduce the corresponding conditional free energy $F(\Phi, T)$ via

$$e^{-F(\Phi, T)/T} = Z(\Phi, T) = \sum_{\{g_a\}_\Phi} e^{-E_a/T}, \tag{2}$$

where $\{g_a\}_\Phi$ is a subset of $\{g_a\}$ and contains the graphs g_a with the same order parameter Φ .

In the $T \rightarrow \infty$ limit the dynamics converges to a totally random rewiring process, and thus, the classical Erdős–Rényi (ER) [11] random graphs are recovered. On the other hand, at low temperatures the topologies with lowest energy occur with enhanced probability. If the energy function is chosen such that compact configurations (which occur exceptionally rarely in the ER graphs) have low energies, then in an intermediate temperature range we expect to see a competition between the entropically favorable dispersed configurations and the energetically favorable compact ones. This is manifested in the shift of the minimum of the conditional free energy $F(\Phi, T)$ from $\Phi = 0$ towards higher values of Φ as the temperature T is decreased from infinity to zero. A sudden change in the position of the global minimum signals a discontinuous (first order) phase transition, whereas a gradual shift indicates either a cross-over or a continuous (second order) phase transition.

In the classical random graph model [9] (corresponding to $T \rightarrow \infty$) by varying the average degree of vertices $\langle k \rangle = 2M/N$, a percolation phase transition occurs at $\langle k \rangle = 1$. For $\langle k \rangle < 1$ the graphs falls apart into small pieces, while for $\langle k \rangle \geq 1$ a giant connected component emerges. Near the critical point the size of the giant component scales as $(\langle k \rangle - 1)M$.

Based on the lattice gas analogy we expect that if $\langle k \rangle < 1$, then for a suitable choice of the energy (one that rewards clustering) a similar dispersed–compact phase transition occurs at a finite temperature $T(\langle k \rangle)$. Such a transition can be best monitored by the order parameter $\Phi_s = s_{\max}/M$ [2,11].

The most obvious energy satisfying the above requirement is a monotonically decreasing function $E = f(s_{\max})$. It can be shown (by counting the number of configurations) that the conditional free energy up to second order in Φ_s (and after omitting the Φ_s independent terms) can be approximated by

$$\frac{F(\Phi_s, T)}{MT} \approx \frac{f(\Phi_s M)}{MT} + [\langle k \rangle - 1 - \ln(\langle k \rangle)]\Phi_s + [\langle k \rangle^2 - 3\langle k \rangle + 2] \frac{\Phi_s^2}{4} \tag{3}$$

in the $N, M \rightarrow \infty$ limit for fixed $\langle k \rangle = 2M/N$.

The simplest choice for the energy function is $f(s_{\max}) = -s_{\max}$. In this case it can be clearly seen from Eq. (3) that as long as $1/T < 1/T_c(\langle k \rangle) = \langle k \rangle - 1 - \ln(\langle k \rangle)$, the free energy has a minimum at $\Phi_s = \Phi_s^*(T) = 0$, i.e., the configuration is dispersed (see

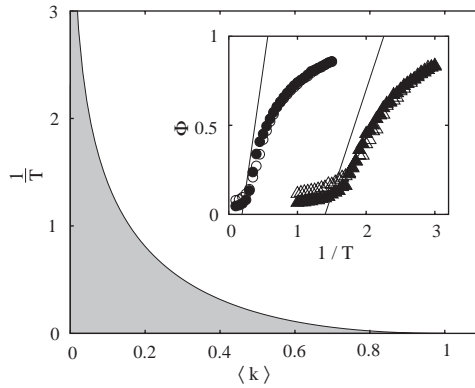


Fig. 1. The phase diagram and the order parameter for the $E = -s_{\max}$ energy. Main panel: the white and shaded areas correspond to the ordered phase (containing a giant component) and the disordered phase, respectively. Inset: The order parameter $\Phi = \Phi_s = s_{\max}/M$ obtained from MC simulations as a function of the inverse temperature for $\langle k \rangle = 0.1$ (triangles) and $\langle k \rangle = 0.5$ (circles). Each data point is an ensemble average of 10 runs, time averaged between $t = 100N$ and $500N$ MC steps. The open and closed symbols represent $N = 500$ and 1000 vertices, respectively. The critical exponent, in agreement with the analytical approximations (solid lines), was found to be 1.

main panel of Fig. 1). When the temperature drops below $T_c(\langle k \rangle)$, the minimum moves away from $\Phi_s = 0$ and a giant component appears. Near the critical temperature $T_c(\langle k \rangle)$ the order parameter at the minimum of the free energy can be estimated from Eq. (3) as $\Phi_s^*(T) = 2[1/T - 1/T_c(\langle k \rangle)]/[\langle k \rangle^2 - 3\langle k \rangle + 2]$, indicating that we are dealing with a *second-order topological phase transition* (see inset of Fig. 1).

For other forms of $f(s_{\max})$, such as $-s_{\max}^2$ or $-s_{\max} \ln(s_{\max})$, first-order topological phase transitions are also expected to occur. We have found such transitions numerically (not shown). In addition, similar results are expected for any (non-global) energy $E = \sum_j f(s_j)$, where the summation goes over each connected component and s_j denotes the number of edges in the j th one. However, because the total number of edges, $M = \sum_j s_j$, is conserved by the dynamics, $f(s_j)$ must decrease faster than $-s_j$ in this case.

Next we turn to another important class of the energy functions, where the energies are assigned to the vertices rather than to the connected components of the graph: $E = \sum_{i=1}^N f(k_i)$, where k_i denotes the degree (number of neighbors) of vertex i . This energy is consistent with a dynamics, in which the change of the degree of a vertex depends only on the structure of the graph in its vicinity. The fitness of an individual vertex depends on its connectivity. The most suitable order parameter for this class of graph energy is $\Phi_k = k_{\max}/M$. Again, due to the conservation of the number of edges, $M = \sum_i k_i$, the single vertex energy $f(k_i)$ should decrease faster than $-k_i$, if aggregation is to be favored.

If we express the single vertex energy in the form $f(k_i) = k_i g(k_i)$, then the total energy of the graph can also be written as $E = \sum_{i=1}^N f(k_i) = \sum_{i=1}^N \sum_{i'} g(k_{i'})$, where i' runs over all vertices that are neighbors of vertex i . Therefore, the graph energy can

also be interpreted in such a way, that every vertex i collects an energy $g(k_i)$ from each of its neighbors. In this interpretation, the fitness of an individual vertex depends on the connectivities of its neighbors.

A relevant case is when $f(k_i) = -(J/2)k_i^2$, or equivalently, $g(k_i) = -(J/2)k_i$, corresponding to a linear preference of the number of pairs that can be chosen from the edges of a vertex, or the number of edges owned by the neighbors of a vertex. Furthermore, this form is in full analogy with the usual definition of the energy $E = -J \sum_{\langle i,j \rangle} n_i n_j$ of a lattice gas on the edge-dual graph of the fully connected network with nearest-neighbor attraction. The summation here runs over all adjacent pairs of lattice sites, and $n_i = 1$ if site i is occupied and 0 otherwise. When this energy is applied to normal cubic lattices, we recover the standard lattice gas model of nucleation of vapors. The negative energy unit $-J$ associated with a pair of edges sharing a vertex in the original graph is equivalent to the binding energy between the corresponding occupied nearest-neighbor sites on the edge-dual graph. By measuring the energies (and temperature) in units of J we can set $J = 1$, without losing generality. Thus, from now on J will be omitted.

For $f(k_i) = -k_i^2/2$, the topology with the lowest overall energy is a “star”, where all the M edges are connected to a single node. Both Monte–Carlo (MC) simulations (see Fig. 2) and a simple theoretical approximation indicate that a first-order phase transition occurs between a dispersed configuration and a star as the temperature is varied. For large enough systems, a sudden change of the order parameter between zero and one can be observed. The hysteresis appearing between cooling and heating is consistent with a first-order transition.

The numerical results are supported by the following *simple theoretical analysis* of the free energy. The idea is to focus on the regime, where the most highly connected vertex has already accumulated the majority of the edges ($\Phi_k > \frac{1}{2}$) and, thus, the energy of the entire graph can be approximated by $f(k_{\max}) = -k_{\max}^2/2$. Then, in the

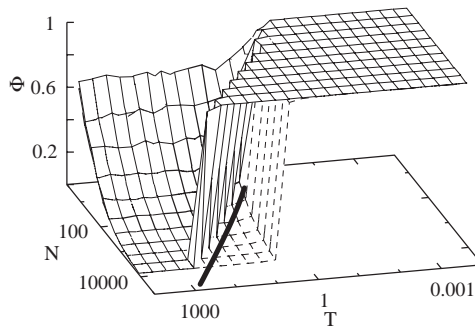


Fig. 2. The order parameter $\Phi = \Phi_k = k_{\max}/M$ as a function of the temperature and the system size for $E = \sum_i -k_i^2/2$ and $\langle k \rangle = 0.5$. The simulations started either from a star (corresponding to $T = 0$, solid line) or a classical random graph ($T = \infty$, dashed line). Each data point represents a single run, time averaged between $t = 100N$ and $200N$ MC steps. The thick solid line shows the analytically calculated spinodal $T_1 = M/\ln(N)$.

thermodynamic limit, the free energy to leading order can be written as

$$\frac{F(\Phi_k, T)}{MT} \approx \frac{f(\Phi_k M)}{MT} + \Phi_k \ln(N), \quad (4)$$

where the Φ_k independent terms have been dropped. Note that this approximation would be valid even for $\Phi_k < \frac{1}{2}$, if the energy of the graph was simply defined as $E = f(k_{\max})$.

For $f(\Phi_k M) = -(\Phi_k M)^2/2$ the parabola given by Eq. (4) has a maximum at $\Phi_k = (T/M) \ln(N)$. When $T \rightarrow 0$, this maximum also shifts towards zero and $F(\Phi_k, T)$ becomes a descending parabola on the $[0, 1]$ interval. This means that the minimum of the free energy is at $\Phi_k = 1$, the star configuration. In contrast, when the temperature goes above the $T_1 = M/\ln(N)$ spinodal point (thick solid line in Fig. 2), the maximum leaves the $[0, 1]$ interval and the free energy becomes an ascending parabola, resulting in a minimum at a low value of Φ_k (corresponding to an ER random graph). However, this value cannot be deduced from Eq. (4), because it is a valid approximation only for $\Phi_k > \frac{1}{2}$. For intermediate temperatures the maximum of the parabola separates the two extreme topologies (the dispersed random graph and the star), among which one is a metastable configuration and the other one is absolutely stable.

Another application-motivated choice for the single vertex energy is $f(k_i) = -k_i \ln(k_i)$, or equivalently, $g(k_i) = -\ln(k_i)$, inspired, in part, by the logarithmic law of sensation. It is the logarithm of the degree of a vertex that its neighbors can sense and benefit from. For this energy the configuration of lowest energy is a fully connected subgraph [or almost fully connected if M cannot be expressed as $n(n-1)/2$]. However, the star configuration is also quite favorable. The order parameter $\Phi_k = k_{\max}/M$ can easily distinguish between these two configurations, because $k_{\max} \approx \sqrt{2M}$ for a fully connected subgraph and $k_{\max} \approx M$ for a star. Our MC simulations demonstrate (Fig. 3) that as we cool down the system, the dispersed random graph first assembles to a configuration with a few large stars (sharing most of their neighbors), and then at lower temperatures it reorganizes into an almost fully connected subgraph. The hysteresis near this latter transition suggests that it is a first-order phase transition. On the other hand, the former transition is accompanied by a singularity in the heat capacity and no hysteresis is observed, indicating that it is a second-order phase transition.

For $\Phi_k > \frac{1}{2}$ Eq. (4) can be used again as a good approximation for the free energy of the graph. By plugging $f(\Phi_k M) = -(\Phi_k M) \ln(\Phi_k M)$ into that expression, we get $F(\Phi_k, T)/(MT) \approx (1 - 1/T) \ln(N) \Phi_k$ to leading order, which is linear in Φ_k . In agreement with our observations above, this formula predicts that for $T < 1$ the star is a stable configuration ($\Phi_k=1$ is a minimum of the free energy), and for $T > 1$ it becomes unstable. The transition at $T = T_c = 1$ is thus step-like with no hysteresis, indicating a second-order phase transition with an infinitely large critical exponent. The observed deviation of T_c from 1 is a finite size effect.

A remarkable feature of the MC dynamics is that by crossing T_c from above, a scale-free graph (with a degree distribution $\sim k^{-\gamma}$ with $\gamma \simeq 3$) appears at some point of the evolution of the graph from the random configuration towards the star. This supports the notion that scale-free graphs are typically non-equilibrium (dynamical) configurations. The MC dynamics is governed by the change of the energy associated

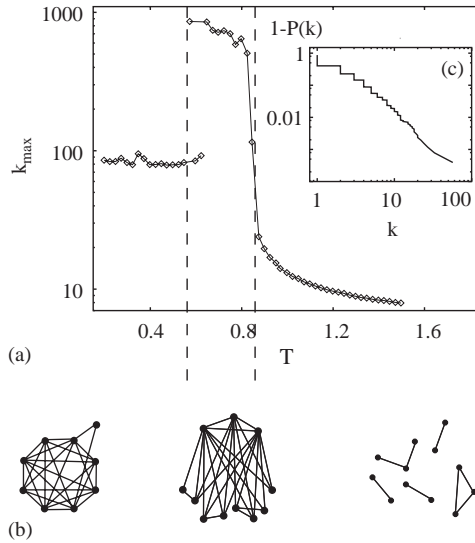


Fig. 3. Phases of the graph when the energy is $E = -\sum_i k_i \ln(k_i)$. (a) The largest degree k_{\max} for $N = 10,224$ vertices and $M = 2556$ edges. Each data point represents a single run, time averaged between $t = 5000N$ and $20,000N$ MC steps. The data points are connected to guide the eye. There is a sharp, continuous transition near $T = 0.85$ and a first-order transition (with a hysteresis) around $T = 0.5 - 0.6$. (b) The three different plateaus in (a) correspond to distinct topological phases: $k_{\max} = \mathcal{O}(1)$ to the classical random graph, $k_{\max} = \mathcal{O}(M)$ to the star phase (a small number of stars sharing most of their neighbors) and $k_{\max} = \mathcal{O}(\sqrt{M})$ to the fully connected subgraph. (c) The (cumulative) degree distribution at $T = 0.84$ and $t = 600N$ follows a power law.

with the relocation of an edge. Estimating the energy change of a vertex by the derivative of the single vertex energy $f(k_i) = -k_i \ln(k_i)$, we get $\Delta E = 1 - \ln(k_i)$. Plugging this into the Boltzmann factor, $\exp[-\Delta E/T]$, at $T = T_c = 1$ we get a quantity proportional to k_i for the acceptance/rejection ratio of a randomly selected move. Since the preferential attachment in the Barabási–Albert model [4] is proportional to k_i , it is *natural that our dynamics also produces scale-free graphs*.

Due to the macroscopic number of edges of the most highly connected vertices in the compact configurations, the graph energy is a non-extensive quantity. More precisely, the energy in different topological states scales differently with the size of the system. For instance, when $f(k_i) = -k_i \ln(k_i)$, the energy of the star and the fully connected subgraph scales as $N \ln(N)$, whereas that of the dispersed state scales as N . Thus (unlike in the mean-field Ising model), there is *no way to choose an appropriate coupling constant that could render the energy extensive in all topological states simultaneously*.

However, the dispersed state (having an extensive graph energy) can equally be studied in the grand canonical ensemble. There, the degree distribution can be expressed as [6] $P_k = C \exp[-f(k)/T - \mu k]/k!$, where C is a normalization factor and the chemical potential μ is adjusted to give the correct $\langle k \rangle$. For $f(k) = -k \ln(k)$, using Stirling’s

formula, the distribution takes the form $P_k = C \exp[-(\mu - 1)k]k^{(1/T-1)k}/\sqrt{2\pi k}$. When $T > 1$, this has a tail, which decays faster than exponential, consequently, each vertex has a small degree. For $T < 1$, on the other hand, the tail becomes divergent, signaling a phase transition at $T = T_c = 1$. However, note that in the $T < 1$ temperature range, due to the non-extensive contribution of the diverging degrees, the ensembles are not equivalent, and the grand canonical description loses its validity.

At the critical temperature, the grand canonical description might still be valid. Thereby choosing a more general single vertex energy $f(k_i) = -(k_i - \alpha) \ln(k_i)$ and setting $\langle k \rangle$ such that $\mu = 1$, the degree distribution acquires a power-law tail ($P_k \sim k^{-(\alpha+1/2)}$) and the network becomes scale-free. We have to stress though that the scale-free network at T_c is not general: for $\mu > 1$ the tail decays exponentially, and for $\mu < 1$ the tail diverges.

Although in this paper we assumed that $\langle k \rangle \leq 1$, this is not a necessary requirement, when the energy is assigned to single vertices. For large average degree ($\langle k \rangle > 2$) the only difference is that one vertex cannot collect all the edges, and thus, several stars appear in the “star” configuration. Finally, further interesting directions in the context of the above study include the investigation of additional relevant forms for the energy [e.g., $E = (k - n)^2$ with $n > 1.5$] and the joint effects of restructuring and growth.

The authors are grateful to Gábor Tuszányi for many valuable discussions. This research has been supported by Hungarian National Science Foundation, Grant no: OTKA 034995.

References

- [1] A.-L. Barabási, R. Albert, *Rev. Mod. Phys.* 74 (2002) 47.
- [2] J.F.F. Mendes, S.N. Dorogovtsev, *Evolution of Networks: from Biological Nets to the Internet and WWW*, Oxford University Press, Oxford, 2003.
- [3] D.J. Watts, S.H. Strogatz, *Nature* 393 (1998) 440–442.
- [4] A.-L. Barabási, R. Albert, *Science* 286 (1999) 509.
- [5] Z. Burda, J.D. Correia, A. Krzywicki, *Phys. Rev. E* 64 (2001) 046 118.
- [6] J. Berg, M. Lässig, *Phys. Rev. Lett.* 89 (2002) 228701.
- [7] Z. Burda, A. Krzywicki, *Phys. Rev. E* 67 (2003) 046 118.
- [8] D. Stark, B. Vedres, Santa Fe Institute, Working Paper, 2003.
- [9] B. Bollobás, *Graph Theory, An Introductory Course*, Springer, Berlin, 1979.
- [10] A. Ramezanpour, V. Karimipour, A. Mashaghi, *Phys. Rev. E* 67 (2003) 046 107.
- [11] P. Erdős, A. Rényi, *Publ. Math. Inst. Hung. Acad. Sci.* 5 (1960) 17–61.