Detecting rich-club ordering in complex networks

V. Colizza, A. Flammini, M.A. Serrano, A. Vespignani February 2, 2008

School of Informatics and Department of Physics, Indiana University, Bloomington 47406 IN

Uncovering the hidden regularities and organizational principles of networks arising in physical systems ranging from the molecular level to the scale of large communication infrastructures is the key issue for the understanding of their fabric and dynamical properties ^{1,2,3,4,5}. The "rich-club" phenomenon refers to the tendency of nodes with high centrality, the dominant elements of the system, to form tightly interconnected communities and it is one of the crucial properties accounting for the formation of dominant communities in both computer and social sciences ^{4,5,6,7,8}. Here we provide the analytical expression and the correct null models which allow for a quantitative discussion of the rich-club phenomenon. The presented analysis enables the measurement of the rich-club ordering and its relation with the function and dynamics of networks in examples drawn from the biological, social and technological domains.

Recently, the informatics revolution has made possible the analysis of a wide range of large scale, rapidly evolving networks such as transportation, technological, social and biological networks ^{1,2,3,4,5}. While these networks are extremely different from each other in their function and attributes, the analysis of their fabric provided evidence of several shared regularities, suggesting general and common self-organizing principles beyond the specific details of the individual systems. In this context, the statistical physics approach has been exploited as a very

convenient strategy because of its deep connection with statistical graph theory and because of its power to quantitatively characterize macroscopic phenomena in terms of the microscopic dynamics of the various systems 1,2,3,4,9 . As an initial discriminant of structural ordering, attention has been focused on the networks' degree distribution; i.e., the probability P(k) that any given node in the network shares an edge with k neighboring nodes. This function is, however, only one of the many statistics characterizing the structural and hierarchical ordering of a network; a full account of the connectivity pattern calls for the detailed study of the multi-point degree correlation functions and/or opportune combination of these.

In this paper, we tackle a main structural property of complex networks, the so-called "rich-club" phenomenon. This property has been discussed in several instances in both social and computer sciences and refers to the tendency of high degree nodes, the hubs of the network, to be very well connected to each other. Essentially, nodes with a large number of links - usually referred to as $rich \ nodes$ - are much more likely to form tight and well interconnected subgraphs (clubs) than low degree nodes. A first quantitative definition of the rich-club phenomenon is given by the rich-club coefficient ϕ , introduced by Zhou and Mondragon in the context of the Internet⁷. Denoting by $E_{>k}$ the number of edges among the $N_{>k}$ nodes having degree higher than a given value k, the rich-club coefficient is expressed as:

$$\phi(k) = \frac{2E_{>k}}{N_{>k}(N_{>k} - 1)},\tag{1}$$

where $N_{>k}(N_{>k}-1)/2$ represents the maximum possible number of edges among the $N_{>k}$ nodes. Therefore, $\phi(k)$ measures the fraction of edges actually connecting those nodes out of the maximum number of edges they might possibly share. The rich club coefficient is a novel probe for the topological correlations in a complex network, and it yields important information about its underlying architecture. Structural properties, in turn, have immediate consequences on network's features and tasks, such as e.g. robustness, performance of biological functions, or selection of traffic backbones, depending on the system at hand. In a social context, for example, a strong rich-club phenomenon indicates the dominance of an "oligarchy" of highly connected and mutually communicating individuals, as opposed to a structure comprised of many loosely

connected and relatively independent sub-communities. In the Internet, such a feature would point to an architecture in which important hubs are much more densely interconnected than peripheral nodes in order to provide the transit backbone of the network⁷. It is also worth stressing that the rich club phenomenon is not trivially related to the mixing properties of networks, which enable the distinction between assortative networks, where large degree nodes preferentially attach to large degree nodes, and disassortative networks, showing the opposite tendency ^{10,11,12}. Indeed, the rich club phenomenon is not necessarily associated to assortative mixing. In the top panel of Fig. 1, we sketch a simple construction in which a disassortative network is exhibiting the rich club phenomenon. In other words, the rich club phenomenon and the mixing properties express different features that are not trivially related or derived one from each other (the technical discussion of this point is reported in the methods section).

In Fig. 1, we report the behavior of the rich club coefficient as a function of the degree in a variety of real world networks drawn from the biological, social and technological world. In Table 1, we summarize the basic topological features of these networks and the datasets used. We also consider three standard network models: the Erdös-Rényi (ER) graph 22 , the generalized random network having a heavy-tailed degree distribution obtained with the Molloy-Reed (MR) algorithm 23 , and the Barabasi-Albert (BA) model 24 . In the ER graph, N nodes are connected by E edges randomly chosen with probability p out of the N(N-1)/2 possible pairs of nodes. The MR network is obtained starting from a given degree sequence P(k) (in our case $P(k) \sim k^{-\gamma}$ with $\gamma=3$) by randomly connecting nodes with the constraints of avoiding self-loops and multiple edges. The BA model is generated by using the growing algorithm of Ref. 24 that produces a scale-free graph with power-law degree sequence with exponent $\gamma=3$. In all cases, the generated networks have $N=10^5$ vertices and an average degree $\langle k \rangle=6$.

As is evident from Fig. 1, the monotonic increasing of $\phi(k)$ is a feature shared by all the analyzed datasets. This behavior is claimed to provide evidence of the rich-club phenomenon since $\phi(k)$ progressively increases in vertices with increasing degree (e.g., see Ref.⁷ for the Internet case, where a different representation of the function is adopted with ϕ defined in terms

of the rank r of nodes sorted by decreasing degree values). However, a monotonic increase of $\phi(k)$ does not necessarily implies the presence of the rich-club phenomenon. Indeed, even in the case of the ER graph - a completely random network - we find an increasing rich-club coefficient. This implies that the increase of $\phi(k)$ is a natural consequence of the fact that vertices with large degree have a larger probability of sharing edges than low degree vertices. This feature is therefore imposed by construction and does not represent a signature of any particular organizing principle or structure, as is clear in the ER case. The simple inspection of the $\phi(k)$ trend is therefore potentially misleading in the discrimination of the rich-club phenomenon.

In order to find opportune baselines for the detection of the rich-club phenomenon we focus on the theoretical analysis of $\phi(k)$. In the methods section we derive an expression for the rich club coefficient as a function of the convolution of the two vertices degree correlation function P(k,k'). Interestingly, it is possible to obtain an explicit expression for the rich-club coefficient of random uncorrelated networks. In this case, the two-vertices correlation function is a simple function of the degree distribution, yielding the following behavior for uncorrelated large size networks at large degrees:

$$\phi_{unc}(k) \sim \frac{k^2}{\langle k \rangle N} ,$$
 (2)

where k_{max} is the maximum degree present in the network. Eq.(2) shows unequivocally that the rich-club coefficient is also a monotonically increasing function for uncorrelated networks, so that, in order to assess the presence of rich-club structural ordering, it is necessary to compare it with the one obtained from the appropriate null model with the same degree distribution, thus providing a suitable normalization of $\phi(k)$.

From the previous discussion, a possible choice for the normalization of the rich-club coefficient is provided by the ratio $\rho_{unc}(k) = \phi(k)/\phi_{unc}(k)$, where $\phi_{unc}(k)$ is analytically calculated by inserting in Eq. (4), reported in the methods section, the network's degree distribution P(k). A ratio larger than one is the actual evidence for the presence of a rich-club phenomenon leading to an increase in the interconnectivity of large degree nodes in a more pronounced way than

in the random case. On the contrary, a ratio $\rho_{unc}(k) < 1$ is a signature of an opposite organizing principle that leads to a lack of interconnectivity among large degree nodes. On the other hand, a completely degree-degree uncorrelated network with finite size is not always realizable due to structural constraints. Indeed, any finite size random network presents a structural cut-off value k_s over which the requirement of the lack of dangling edges introduces the presence of multiple and self-connections and/or degree-degree correlations ^{21,25}. Networks with bounded degree distributions and finite second moment $\langle k^2 \rangle$ present a k_{max} that is below the structural one k_s . In this situation, $\phi_{unc}(k)$ is properly defined for all degrees and is representative of the network topology. However, in networks with heavy-tailed degree distribution (e.g., scale-free degree distributions with $2 < \gamma \le 3$, as observed in many real systems), this is no longer the case and k_s is generally smaller than k_{max} . In fact, structural degree-degree correlations and higher order effects, such as the emergence of large cliques²⁶, set in even in completely random networks. The normalization of $\phi(k)$ that takes into account these effects is provided by the expression $\rho_{ran}(k)=\phi(k)/\phi_{ran}(k)$, where $\phi_{ran}(k)$ is the rich-club coefficient of the maximally random network with the same degree distribution P(k) of the network under study. Operatively, the maximally random network can be thought of as the stationary ensemble of networks visited by a process that, at any time step, randomly selects a couple of links of the original network and exchange two of their ending points (automatically preserving the degree distribution). Also in this case an actual rich-club ordering is denoted by a ratio $\rho_{ran}(k) > 1$. Therefore, whereas $\rho_{unc}(k)$ provides information about the overall rich-club ordering in the network with respect to an ideally uncorrelated graph, $\rho_{ran}(k)$ is a normalized measure which discounts the structural correlations due to unavoidable finite size effects, providing a better discrimination of the actual presence of the rich club-phenomenon due to the ordering principles shaping the network.

In Fig. 2, we report the ratios $\rho_{ran}(k)$ for the real world and the simulated networks. The analysis clearly discriminates between networks with or without rich-club ordering. In particular, we identify a strong rich-club ordering in the Scientific Collaboration Network, providing support to the idea that the elite formed by more influential scientists tends to form collaborative

groups within specific domains. This also supports the view that the rich-club phenomenon is a natural tendency in many social networks. We find a clearly opposite result in the decreasing behavior of the rich club spectrum for the Protein Interaction Network and the Internet map at the Autonomous System level. In both cases, this evidence provides interesting information regarding the system structure and function.

The lack of rich-club ordering in the Protein Interaction Network indicates that proteins with large number of interactions are presiding over different functions and thus, in general, are coordinating specific functional modules (whose detailed analysis requires specific tools 27). Figure 3 shows portions of the Protein Interaction Network and the Scientific Collaboration Network including the club of $N_{>k}$ nodes $-N_{>k}=29$ and $N_{>k}=35$ for the Protein Interations, $N_{>k}=30$ and $N_{>k}=36$ for the Scientific Collaboration – and the connections among them. The network representations clearly show the presence of a rich-club phenomenon in the Scientific Collaboration Network, where the majority of rich nodes are highly interconnected forming tight subgraphs, in contrast with the Protein Interaction Network case, where only few links appear to connect rich nodes, the rest linking to lower degree vertices.

In the case of the Internet, the appropriate analysis of the rich-club phenomenon shows that, contrary to previous claims⁷, the structure at the Autonomous System level *lacks* rich-club ordering. This might appear counter-intuitive. It is reasonable to imagine the Internet backbone made of interconnected transit providers which are also local hubs. This is however not the case and an explanation can be easily found in the fact that we are just considering topological properties. Indeed, the backbone hubs are identified more in terms of their bandwidth and traffic capacity than in terms of the sole number of connections. The present result suggests that high degree hubs provide connectivity to local region of the Internet and are not tightly interconnected. The backbone of interconnected transit providers is instead identified by high traffic links which play a crucial role in terms of traffic capacities but whose number might represent a small fraction of the total possible number of interconnections.

The previous discussion points out that, in some cases, the concept of rich-club ordering

should be generalized in order to evaluate the richness of vertices not just in terms of their degree but in terms of the actual traffic or intensity of interactions handled. In this case, we have to consider a weighted network representation of the system where a weight w_{ij} representing the traffic or intensity of interaction is associated to each edge between the vertices i and j. Also in this case, however, the study of the weighted rich-club coefficient alone does not discriminate the actual presence of the rich club effect (see Methods). Given the entanglement of the weight and degree correlations, the appropriate null hypothesis is however more complicated to define and a detailed account of the evaluation of the weighted rich-club effect will be provided elsewhere.

In summary, the presented analysis provides the baseline functions for the detection of the rich-club phenomenon and its effect on the structure of large scale networks. This allows the measurement of this effect in a wide range of systems, finally enabling a quantitative discussion of various claims such as "high centrality" backbones in technological networks and "elitarian" clubs in social systems.

Methods

Analytic expression of the rich club coefficient. The basic analytical understanding of the rich-club phenomenon starts by considering the quantity $E_{kk'}$, representing the total number of edges between vertices of degree k and of degree k' for $k \neq k'$, and twice the number of edges between vertices in the same degree class. We can express the numerator of $\phi(k)$ in Eq. 1 as $2E_{>k} = \int_k^{k_{max}} dk' \int_k^{k_{max}} dk'' E_{k'k''}$, where k_{max} is the maximum degree present in the network and where, for the sake of simplicity, the variable k is thought of as continuous. In turn, the quantity $E_{kk'}$ can be expressed as a function of the joint degree probability distribution 11,21,28,29 via the identity $N\langle k \rangle P(k,k') = E_{kk'}$, yielding

$$\phi(k) = \frac{N\langle k \rangle \int_k^{k_{max}} dk' \int_k^{k_{max}} dk'' P(k', k'')}{\left[N \int_k^{k_{max}} dk' P(k') \right] \left[N \int_k^{k_{max}} dk' P(k') - 1 \right]}.$$
 (3)

From Eq. (3), it is clear that $\phi(k)$ is also a measure of correlations in the network, although

it represents a different projection of P(k,k') as compared to other degree-degree correlation measures. At the same time, it is possible to see that the rich-club coefficient express a property that is not trivially related to the usual indicators of assortative behavior, such as the Pearson's correlation coefficient 11 or the average nearest neighbor degree 10 . Notice that these assortativity measures quantify two-point correlations and so account for quasi-local properties of the nodes in the network, whereas the rich club phenomenon is computed as a global feature within a restricted subset. The double integral is indeed a convolution of the correlation function that allows the presence of different combinations of the assortative and rich-club features in the same network.

Only in the case of random uncorrelated networks ^{3,4,29}, the joint degree distribution P(k, k') factorizes and takes the simple form $P_{unc}(k, k') = kk'P(k)P(k')/\langle k \rangle^2$. By inserting this expression into Eq. (3), we obtain $\phi(k)$ for uncorrelated networks as

$$\phi_{unc}(k) = \frac{1}{N\langle k \rangle} \left[\frac{\int_{k}^{k_{max}} dk' k' P(k')}{\int_{k}^{k_{max}} dk' P(k')} \right]^{2} \sim \frac{k^{2}}{\langle k \rangle N} , \qquad (4)$$

where we have applied L'Hôpital's rule to derive the behavior for large size networks and large degrees.

Rich club coefficient for weighted networks. If the rich-club is defined as the set of nodes having a strength larger than a given value s, a possible definition of the *weighted rich-club coefficient* can be expressed as

$$\phi^{w}(s) = \frac{2W_{>s}}{\sum_{i|s_{i}>s} s_{i}},\tag{5}$$

where $W_{>s}$ represents the sum of the weights on the links connecting two nodes in the club and the normalization is given by the sum of the strengths of the rich nodes.

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Correspondence and requests for materials should be addressed to A.V.

Table legend

Table 1: Basic topological properties of the analyzed datasets. We considered four real world networks: (1) the *Protein Interaction Network* 12,13 of the yeast *Saccharomyces Cerevisiae* collected with different experimental techniques and documented at the Database of Interacting Proteins (http://dip.doe-mbi.ucla.edu/); (2) the *Scientific Collaboration Network* 14,15 extracted from the electronic database e-Print Archive in the area of condensed matter physics (http://xxx.lanl.gov/archive/cond-mat/), from 1995 to 1998, in which nodes represent scientists and a connection exists if they coauthored at least one paper in the archive; (3) the network of *Worldwide Air Transportation* 16,17 representing the International Air Transport Association (http://www.iata.org/) database of airport pairs connected by direct flights for the year 2002; (4) the *Internet* network at the Autonomous System 4 level 4 , 10,18,19,20 from data collected by the *Oregon Route Views* project (http://www.routeviews.org/) in May 2001, in which nodes represent Internet service providers and edges connections among those. The sizes of the networks in number of nodes and edges are shown, along with the average degree $\langle k \rangle$ and the maximum degree value k_{max} . We also give the value for the corresponding structural cut-off, k_s , in the uncorrelated case 21 .

Figure Legends

Figure 1: Schematic picture of the rich-club phenomenon and rich-club spectrum $\phi(k)$ for real networks. At the top, a conceptual example of disassortative network displaying the presence of the rich-club phenomenon is shown. Disassortative mixing is given by the tendency of hubs to be on average more likely connected to low degree nodes. However, the four rich nodes represented in the schematic picture show a clear rich-club behavior by forming a fully connected clique within the club. At the bottom, results for the four real-world networks and the three models analyzed are shown. The computer generated networks - ER, MR, and BA have size $N=10^5$ and average degree $\langle k \rangle=6$. ER refers to the *Erdös-Rényi graph*, MR is

constructed from the Molloy-Reed algorithm with a given degree distribution $P(k) \sim k^{-3}$, and the BA model is generated by growing a network with preferential attachment that produces a scale-free graph with power-law degree sequence with exponent $\gamma=3$. Results are averaged over $n=10^2$ different realizations for each model. All networks share a monotonic increasing behavior of $\phi(k)$, independent of the nature of the degree distribution characterizing the network and of the possible presence of underlying structural organization principles. Also random networks, either having a Poissonian degree distribution (such as ER) or a heavy-tailed P(k) (such as MR and BA), show a rich club spectrum increasing with increasing values of the degree. This common trend is indeed due to an intrinsic feature of every network structure, for which hubs have simply a larger probability of being more interconnected than low degree nodes.

Figure 2: Assessment for the presence of the rich-club phenomenon in the networks under study. $\phi(k)$ is compared to the null hypothesis provided by the maximally random network with $\phi_{ran}(k)$. The ratio $\rho_{ran} = \phi/\phi_{ran}$ is plotted as a function of the degree k and compared to the baseline value equal to 1. If $\rho(k) > 1$ (< 1) the network displays the presence (absence) of the rich-club phenomenon with respect to the random case. The Protein Interaction Network, the Internet map at the Autonomous System level and the Scientific Collaboration Network show clear behaviors as explained in the main text. The Worldwide Air Transportation network displays a mild rich-club ordering with $\rho_{ran}(k) > 1$. The ER and MR network models show a ratio $\rho_{ran}(k) = 1 \,\forall k$, as expected, whereas the BA model exhibits a mixing behavior with values above 1 for very high degrees.

Figure 3: Graph representations of the rich-clubs. Progressively smaller clubs of $N_{>k}$ rich nodes in the Protein Interaction Network -top- and in the Scientific Collaboration Network -bottom- are shown together with the $E_{>k}$ connections among them. Here $N_{>k}=35$, $E_{>k}=37$ (top left) and $N_{>k}=29$, $E_{>k}=21$ (top right) for the Protein Interactions; $N_{k>}=36$, $E_{>k}=62$ (bottom left) and $N_{k>}=30$, $E_{>k}=54$ (bottom right) for the collaboration network. The two graph representations for each network show progressively smaller clubs made of $N_{>k}$ rich nodes for increasing values of the degree k. The links connecting the rich nodes to the

rest of the network are not represented for sake of simplicity. The Protein Interaction Network shows a club whose hubs are relatively independent being loosely connected among each other, leaving the remaining links to coordinate specific functional modules. A different picture is observed in the Scientific Collaborations case, where most of the hubs form cliques and tightly interconnected subgraphs, thus revealing the tendency of scientists to form densely interconnected collaborative groups. The graphs have been produced with the Pajek software (http://vlado.fmf.uni-lj.si/pub/networks/pajek/).

	Protein Interactions	Scientific Collaborations	Air Transportation	Internet
# nodes	4713	15179	3880	11174
# links	14846	43011	18810	23409
$\langle k \rangle$	6.3	5.7	9.7	4.2
k_{max}	282	97	318	2389
$k_s = \sqrt{\langle k \rangle N}$	172	294	194	216

Table 1.

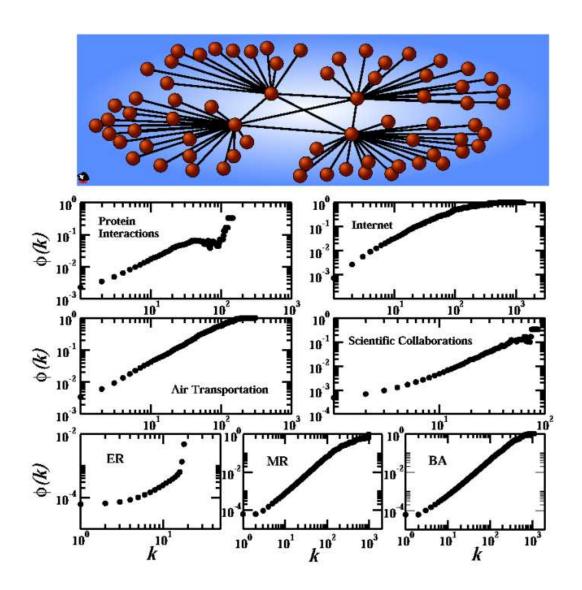


Figure 1.

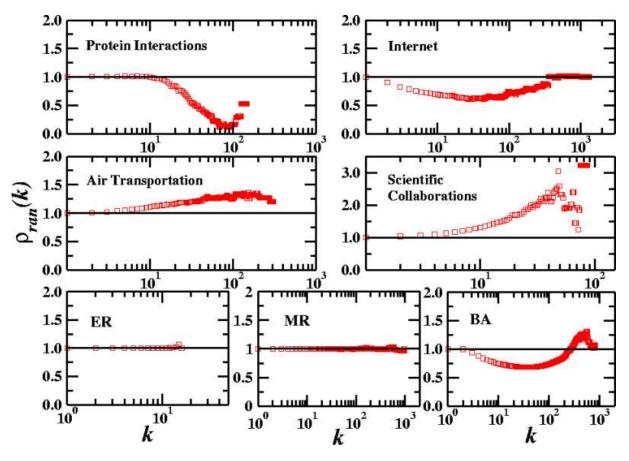


Figure 2.

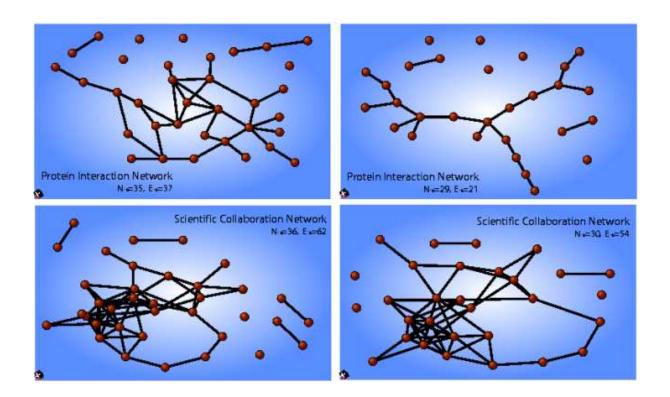


Figure 3.