

Total communicability as a centrality measure

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We examine node centrality measures based on the notion of *total communicability*, defined in terms of the row sums of matrix functions of the adjacency matrix of the network. Our main focus is on the matrix exponential and the resolvent, which have natural interpretations in terms of walks on the underlying graph. While such measures have been used before for ranking nodes in a network, we show that they can be computed very rapidly even in the case of large networks. Furthermore, we propose the (normalized) total sum of node communicabilities as a useful measure of network connectivity. Extensive numerical studies are conducted in order to compare this centrality measure with the closely related ones of subgraph centrality [E. Estrada and J. A. Rodríguez-Velázquez, *Phys. Rev. E*, **71** (2005), 056103] and Katz centrality [L. Katz, *Psychometrika*, 18 (1953), pp. 39–43]. Both synthetic and real-world networks are used in the computations.

Keywords: centrality; communicability; adjacency matrix; matrix functions; network analysis.

1. Introduction

Over the past several years, the analysis of networks has become increasingly important in a number of disciplines [1–8]. Network analysis is used in many situations: from determining network structure and communities, to describing the interactions between various elements of the network, to investigating the dynamics of phenomena taking place on the network (e.g., information flow).

One of the fundamental questions in network analysis is to determine the ‘most important’ elements in a given network. Measures of node importance are usually referred to as *node centrality*, and many centrality measures have been proposed, starting with the simplest of all, the node degree. This crude metric has the drawback of being too ‘local’, as it does not take into effect the connectivity of the immediate neighbours of the node under consideration. A number of more sophisticated centrality measures have been introduced that take into account the global connectivity properties of the network. These include various types of eigenvector centrality for both directed and undirected networks, betweenness centrality, and others which are discussed below. Overviews of various centrality measures can be found in [3,9–14]. The centrality scores can be used to provide *rankings* of the nodes in the network. There are many different ranking methods in use (most of which depend on centrality measures), and many algorithms have been developed to compute these rankings. Information about the many different ranking schemes can be found, e.g., in [3,6,12,15–19].

One now standard method of measuring node importance is *subgraph centrality* [20], which is based on the diagonal entries of a matrix function applied to the adjacency matrix A of the network in question.

Here, the matrix exponential e^A is frequently used. While this approach has been successfully used in a number of problems [3,5,21], obtaining estimates of the diagonals of e^A for a large network with adjacency matrix A can be quite expensive. Indeed, computing individual entries of matrix functions $f(A)$ is generally costly for large A even with the best available algorithms [5,22].

In recent years, efficient algorithms have been developed for computing the *action* of a matrix function on a vector, that is, for computing the vector $f(A)\mathbf{v}$ for a given matrix A (usually large and sparse), vector \mathbf{v} and function f . A particularly important case is that of the matrix exponential, since this provides a solution method for initial value problems for first-order systems of linear ordinary differential equations. These algorithms, based on variants of the Lanczos, Arnoldi or other Krylov subspace method, access the matrix A only in the form of (sparse) matrix-vector products and have $O(n)$ storage cost for a sparse $n \times n$ matrix A [23, Chapter 13]. When $\mathbf{v} = \mathbf{1}$, the vector with all its entries equal to 1, the i th entry of the resulting vector $f(A)\mathbf{1}$ contains the i th row sum of $f(A)$:

$$[f(A)\mathbf{1}]_i = \sum_{j=1}^n [f(A)]_{ij}, \quad 1 \leq i \leq n.$$

This quantity, which has a graph-theoretic interpretation in terms of subgraph centrality and communicability [5,24], can be computed much faster than subgraph centrality using current computational techniques. Of course, the same is true if vector $\mathbf{1}$ is replaced by some other vector—typically, an ‘external importance vector’ which can be used to take into account intrinsic, not network-related contributions to the centrality of each node [13, pp. 174–175].

Such centrality measures have long been in use in network analysis. Note that for the case of the ‘identity function’ $f(A) = A$, and symmetric A (undirected networks), we recover degree centrality. The off-diagonal row sums of e^A have been used in social network analysis to measure the resilience of an individual in the face of hostile attacks from within the network [3, Chapter 6]. More recently, row and column sums of e^A have been applied to the identification of hubs and authorities in directed networks [15]. For resolvent-type functions, such as $f(A) = (I - \alpha A)^{-1}$ (with I the $n \times n$ identity matrix), for suitable values of $\alpha > 0$, we recover the well-known *Katz centrality* and its variants, also known as α -centrality; see, e.g. [10,16,25–27]. None of these previous studies, however, considered algorithmic aspects such as computational cost, storage and so forth.

This paper considers the implications of using the row sums of e^A or similar matrix functions as a measure of node centrality, focusing for the sake of brevity on undirected networks. The interpretation of this measure in terms of total communicability of a node is given, and compared with the one for subgraph centrality in Section 3. In Section 4, the concept of *total network communicability* is introduced and discussed. Section 5 contains experimental comparisons of subgraph centrality and total communicability using various synthetic and real-world networks. Sections 6 and 7 discuss computational aspects and the use of row sum centrality with other standard matrix functions, respectively. We offer some conclusive remarks in Section 8.

2. Background and definitions

The analysis of networks requires the use of notions from graph theory, linear algebra, numerical analysis and computer science. Here, we list some basic definitions and ideas from graph theory. A more complete overview can be found in [28].

A graph $G = (V, E)$ is a set of nodes (vertices) V with $|V| = n$ and edges $E = \{(i, j) | i, j \in V\}$. A graph is *undirected* if the edges are unordered pairs of vertices and *directed* if the pairs are ordered (edges have

a direction). The *degree* of a vertex in an undirected graph is the number of edges which are adjacent to the node. In a directed graph, nodes have both an *in-degree*, the number of edges pointing into the node, and an *out-degree*, the number of edges starting at the node and pointing away. A *simple* graph is a graph with no *loops* (edges from node i to itself), no multiple edges and unweighted edges. In this paper, all networks correspond to simple, undirected graphs unless otherwise specified.

A *walk* of length k on a graph G is a sequence of vertices v_1, v_2, \dots, v_{k+1} such that $(v_i, v_{i+1}) \in E$ for all $1 \leq i \leq k$. A *path* is a walk with no repeated vertices. A *closed walk* is a walk that starts and ends at the same vertex. A *cycle* is a closed walk with no repeated vertices. If any vertex in the graph is reachable from any other vertex, the graph is said to be *connected*.

Every graph can be viewed as a matrix through the use of its *adjacency matrix*. The adjacency matrix of a network with graph G is given by

$$A = (a_{ij}); \quad a_{ij} = \begin{cases} 1 & \text{if } (i, j) \text{ is an edge in } G, \\ 0 & \text{else.} \end{cases}$$

The requirement of unweighted edges causes A to be binary and that of no loops in the graph forces A to have zeros along its diagonal. If the network is undirected, A will be symmetric but if the network is directed, A will generally be unsymmetric. In the case of an undirected network, the eigenvalues of A will be real. We label the eigenvalues of A in non-increasing order: $\lambda_1 \geq \lambda_2 \geq \dots \geq \lambda_n$. Note that the Perron–Frobenius theorem implies that $\lambda_1 > \lambda_2$ if the graph is connected (equivalently, if A is irreducible).

3. Diagonal entries vs. row sums

In [20], the authors introduce the concept of subgraph centrality as a centrality measurement for nodes in a network. This provides a ranking based on the diagonal entries of a matrix function applied to the adjacency matrix. Although there are various choices of function to use, the most common is the matrix exponential. The *subgraph centrality* of node i is given by $[e^A]_{ii}$ where A is the adjacency matrix of the network. The *subgraph communicability* between nodes i and j is given by $[e^A]_{ij}$ (note that in the case of an undirected network, A is symmetric and $[e^A]_{ij} = [e^A]_{ji}$). A node with a (relatively) large subgraph centrality is considered to be more important in the network and is given a higher ranking than nodes with lower subgraph centrality. A (relatively) large subgraph communicability between a pair of nodes i and j indicates that information flows more easily between those two nodes than between pairs of nodes with lower communicability. In other words, a low subgraph communicability indicates that the two nodes cannot easily exchange information. Network communicability can also be interpreted in terms of the correlations between different components of physical systems; see, e.g. [5].

The reasoning behind using the diagonal entries of e^A as a measure of the centrality of a node in the network can be seen by considering the power series expansion of e^A [23]:

$$e^A = I + A + \frac{A^2}{2!} + \frac{A^3}{3!} + \dots + \frac{A^k}{k!} + \dots = \sum_{k=0}^{\infty} \frac{A^k}{k!}. \quad (3.1)$$

It is well known in graph theory (and fairly easy to prove) that if A is the adjacency matrix of a network with unweighted edges, then $[A^k]_{ij}$ counts the number of walks of length k between nodes i and j . Thus, the subgraph centrality of node i , which is equal to $[e^A]_{ii}$, counts the number of closed

walks centered at node i weighting a walk of length k by a penalty factor of $1/k!$. In this way, shorter walks are deemed more important than longer walks. Although some of these walks can be described as ‘illogical’ (for example, the walk $v_i \rightarrow v_j \rightarrow v_i \rightarrow v_j \rightarrow v_i$ is a closed walk of length 4 centred at node i), the subgraph centrality of node i still gives us a measure of how close node i is to everything else in the network.

By contrast, the row sum of e^A for node i is given by $\sum_{j=1}^n [e^A]_{ij}$, which counts all walks between node i and all the nodes in the network (node i included), weighting walks of length k by a penalty factor of $1/k!$. Thus, the i th row sum of e^A can be interpreted as the *total subgraph communicability* of node i , and can be interpreted as a measure of the importance of the i th node in the network, since a node with high communicability with a large number of other nodes in the network is likely to be an important node, and certainly a more important node than one characterized by low total communicability.

An immediate question is how this centrality measure compares with the subgraph centrality of node i in the network. In general, the rankings produced by the total communicability measure will not be the same as those produced by the subgraph centrality measure. The difference between the two rankings is

$$\sum_{j=1}^n [e^A]_{ij} - [e^A]_{ii} = \sum_{j \neq i} [e^A]_{ij} = \sum_{j \neq i} \sum_{k=1}^n e^{\lambda_k} v_{ki} v_{kj}, \quad (3.2)$$

where v_{ik} is the i th element of the normalized eigenvector \mathbf{v}_k of A associated with the eigenvalue λ_k . Note that e^A is always positive-definite and that its diagonal entries are often large compared with the off-diagonals. If the diagonal entries of e^A vary over a wide range while its off-diagonal sums remain confined within a more narrow range, the rankings produced by the two methods will not differ by much. However, this depends both on the spectrum of A and the entries of the eigenvectors.

While it appears to be difficult, in general, to establish a relation between the rankings produced by the subgraph centrality and total communicability, for certain types of simple graphs it is easy to show that the two methods will give identical rankings. These include *complete graphs* and *cycles* (where each node has the exact same ranking under both systems), *paths* and *star graphs*. A star graph on n nodes has one central node that is connected to each of the $n - 1$ remaining nodes and no other edges. Under both ranking systems, the central node is ranked highest and the remaining nodes all have the same scores. This can be shown either using graph theory or by examining the eigenvalues and eigenvectors of the star graph (more information about the spectra of star graphs can be found in [29]).

One case where the two measures could be expected to give similar rankings is that of networks with a large *spectral gap*, which for the purposes of this paper is the difference $\lambda_1 - \lambda_2$ between the first (largest) and second eigenvalue. We have:

$$[e^A]_{ii} = e^{\lambda_1} v_{1i}^2 + \sum_{k=2}^n e^{\lambda_k} v_{ki}^2$$

and

$$[e^A \mathbf{1}]_i = e^{\lambda_1} (\mathbf{v}_1^T \mathbf{1}) v_{1i} + \sum_{k=2}^n e^{\lambda_k} (\mathbf{v}_i^T \mathbf{1}) v_{ki}.$$

Dividing both expressions by the constant e^{λ_1} (which does not affect the rankings) and observing¹ that $\mathbf{v}_1^T \mathbf{1} = \|\mathbf{v}_1\|_1$ shows that for $\lambda_1 \gg \lambda_2$ the two rankings are largely determined by the quantities v_{1i}^2 and $\|\mathbf{v}_1\|_1 v_{1i}$, respectively, and therefore by the entries v_{1i} of the dominant eigenvector of A . Thus, if the difference $\lambda_1 - \lambda_2$ is sufficiently large, the two centrality measures reduce to eigenvector centrality [10] and therefore can be expected to result in very similar rankings, especially for the top nodes. Numerical experiments (not shown here) performed on Erdős–Rényi graphs with large spectral gaps have confirmed this fact.

However, it is difficult to quantify *a priori* how large the spectral gap needs to be for all these rankings to be identical (or even approximately the same). In the section on computational experiments we will see that there can be significant differences between the rankings obtained using subgraph centrality and those using total communicability centrality, even for networks with a relatively large spectral gap.

4. Total network communicability

The total communicabilities of individual nodes give a measure of how well each node communicates with the other nodes of the network. In order to measure how effectively communication takes place across the network as a whole, we consider the sum of all the total communicabilities. For a network with adjacency matrix A , this is given by

$$C(A) = \sum_{i=1}^n [e^A \mathbf{1}]_i = \sum_{i=1}^n \sum_{k=1}^n e^{\lambda_k} (\mathbf{v}_i^T \mathbf{1}) v_{ki} = \mathbf{1}^T e^A \mathbf{1}, \quad (4.1)$$

where, as in Section 3, λ_k is the k th eigenvalue of A and v_{ik} is the i th element of the normalized eigenvector \mathbf{v}_k associated with λ_k . Here, we propose to use the *total network communicability*, $C(A)$, as a global measure of the ease of sending information across a network. We emphasize that while $C(A)$ is defined as the sum of all the entries of e^A , it is not necessary to know any of the individual entries of e^A to compute $C(A)$; indeed, very efficient methods exist to compute quadratic forms of the type $\mathbf{v}^T f(A) \mathbf{v}$ for a given function $f(x)$, matrix A and vector \mathbf{v} , see [22,30,31].

It is instructive to compare the total communicability of a network with the *Estrada index*, an important graph invariant defined as the sum of all the subgraph centralities:

$$EE(A) = \sum_{i=1}^n [e^A]_{ii} = \sum_{i=1}^n e^{\lambda_i} = \text{Tr}(e^A).$$

The following proposition provides simple lower and upper bounds for $C(A)$ in terms of $EE(A)$ and other spectral quantities associated with the underlying network.

PROPOSITION 1 Let A be the adjacency matrix of a simple network on n vertices. Then,

$$EE(A) \leq C(A) \leq n e^{\|A\|_2},$$

where $\|A\|_2$ denotes the spectral norm of A . In particular, for an undirected network we have

$$EE(A) \leq C(A) \leq n e^{\lambda_1}.$$

¹ By the Perron–Frobenius Theorem, the dominant eigenvector can be chosen to have nonnegative entries, and positive entries when the graph G is connected.

Proof. The lower bound is trivial, as

$$EE(A) = \sum_{i=1}^n [e^A]_{ii} \leq \sum_{i=1}^n \sum_{j=1}^n [e^A]_{ij} = \sum_{i=1}^n [e^A \mathbf{1}]_i = C(A).$$

The upper bound follows from noticing that $C(A) = \mathbf{1}^T e^A \mathbf{1} = (e^A \mathbf{1})^T \mathbf{1} = \langle e^A \mathbf{1}, \mathbf{1} \rangle$ and applying the Cauchy–Schwarz inequality to the quadratic form $\langle e^A \mathbf{1}, \mathbf{1} \rangle$:

$$|\langle e^A \mathbf{1}, \mathbf{1} \rangle| \leq \|e^A \mathbf{1}\|_2 \|\mathbf{1}\|_2 \leq \|e^A\|_2 \|\mathbf{1}\|_2 \|\mathbf{1}\|_2 \leq n e^{\|A\|_2}.$$

For an undirected network A is symmetric and $\lambda_1 = \|A\|_2$. □

Note that the lower bound is attained in the case of the ‘empty’ graph with adjacency matrix $A = 0$, while the upper bound is attained on the complete graph, whose adjacency matrix is $A = \mathbf{1}\mathbf{1}^T - I$.

The bounds from Proposition 1 also hold for $e^{\beta A}$, $\beta > 0$. For any connected graph with adjacency matrix A , the bounds get tighter as $\beta \rightarrow 0+$, since both the lower and upper bounds tend to 1. The parameter β can be interpreted as an *inverse temperature* and is a reflection of external disturbances on the network (see, e.g., [5] for details); taking $\beta \rightarrow 0+$ is equivalent to ‘raising the temperature’ of the environment surrounding the network.

When appropriately normalized, $C(A)$ can be used to compare the ease of information exchange on different networks. This could be useful, for instance, in the design of communication networks. In the following sections, we compute the total communicability for various types of networks. The question arises of what would constitute a reasonable normalization factor. There are several possibilities. Normalizing $C(A)$ by the number n of nodes corresponds to the average total communicability of the network per node. Similarly, normalizing $C(A)$ by the number m of edges would correspond to the average total communicability of the network per edge. We also note that the minimum value of $C(A)$ is n , corresponding to the empty graph on n nodes ($V = \emptyset$), while the maximum value is $n^2 e^{n-1} - n$, corresponding to the complete graph on n nodes. The expression

$$\hat{C}(A) := \frac{C(A) - n}{n^2 e^{n-1} - 2n}$$

takes its values in the interval $[0, 1]$, with $\hat{C}(A) = 0$ for ‘empty’ graphs (no communication can take place on such graphs) and $\hat{C}(A) = 1$ on complete graphs (for which the ease of communication between nodes is clearly maximum). Unfortunately, the denominator in this expression grows so fast that for most sparse graphs evaluating $\hat{C}(A)$ results in underflow.

In the experiments below we chose to normalize $C(A)$ by n , the number of nodes, and by m , the number of edges; for the network used in our tests we found that comparing networks based on $C(A)/n$ or on $C(A)/m$ yields exactly the same rankings, therefore we only include results for the former measure.

5. Computational studies

In this section we carry out extensive centrality computations for a variety of networks, with the aim of comparing subgraph centrality with total communicability centrality. In particular, we are interested in determining if, or for what type of networks, the two centrality measures provide similar rankings. Moreover, for those networks where the two measures result in rankings that differ significantly, we would like to obtain some insights on why this is the case. Of course it would be desirable to know

when one measure should be preferred over the other, but this is a difficult problem since it is not easy to come up with objective criteria for comparing ranking methods (see the discussion in [18, Chapter 16]). We will compare the two methods in terms of computational cost in Section 6.

To measure similarities between the rankings obtained with the two methods, we use (Pearson) correlation coefficients and the intersection distance method (see [32] as well as [33,34]) on both the full set V of nodes and on partial lists of nodes. The correlation coefficients are computed using lists of nodes in rank order. The intersection distances are computed using the lists of subgraph centrality and total communicability values. Given two ranked lists x and y , the intersection distance between the two lists is computed in the following way: let x_k and y_k be the top k ranked items in x and y , respectively. Then the top k intersection distance (or *intersection similarity*) is given by

$$\text{isim}_k(x, y) := \frac{1}{k} \sum_{i=1}^k \frac{|x_i \Delta y_i|}{2i}$$

where Δ is the symmetric difference operator between the two sets. If the lists are identical, then $\text{isim}_k(x, y) = 0$ for all k . If the two sequences are disjoint, then $\text{isim}_k = 1$. We denote by cc the correlation coefficient between the two vector rankings, and by cc_p the correlation coefficient between the top $p\%$ of nodes under the two ranking systems. We denote by $\text{isim}_{p\%}$ the intersection distance between the top $p\%$ of nodes.

Unless otherwise specified, all experiments were performed using Matlab version 7.9.0 (R2009b) on a MacBook Pro running OS X Version 10.6.8, a 2.4 GHz Intel Core i5 processor and 4 GB of RAM. In this section, we use the Matlab built-in function `expm` for computing the matrix exponential.

5.1 Test matrices

The synthetic examples used in the tests were produced using the CONTEST toolbox in Matlab [35,36]. The graphs tested were of two types: preferential attachment (Barabási–Albert) model and small-world (Watts–Strogatz) model. In CONTEST, these graphs and the corresponding adjacency matrices can be built using the functions `pref` and `smallw`, respectively.

The preferential attachment model was designed to produce networks with scale-free degree distributions as well as the small-world property [37]. In CONTEST, preferential attachment networks are constructed using the command `pref(n, d)`, where n is the number of nodes in the network and $d \geq 1$ is the number of edges each new node is given when it is first introduced to the network. The network is created by adding nodes one by one (each new node with d edges). The edges of the new node connect to nodes already in the network with a probability proportional to the degree of the already existing nodes. This results in a scale-free degree distribution. Note that with this construction, the minimum degree of the network is d . When $d > 1$, it means that the network has no dangling nodes (nodes of degree 1), whereas in many real-life networks one often observes a high number of dangling nodes. In the CONTEST toolbox, the default value is $d = 2$.

In our experiments, we tested various values of d on a network of size $n = 1000$: 20 networks were tested for all values $1 \leq d \leq 10$, as well as a few larger values. In Table 1, the averages of the correlation coefficients between the subgraph centrality rankings and the total subgraph communicability rankings can be found for various values of d . The intersection distance values can be found in Table 2. The intersection distance values were calculated both for the full set of rankings and for the top 10% of ranked nodes.

TABLE 1 *Comparison, using the correlation coefficient, of rankings based on the diagonal entries and row sums of e^A for 1000-node scale-free networks of various parameters built using the `pref` function in the CONTEST Matlab toolbox. The values reported are the averages over 20 matrices with the same parameters. The parameter d is the initial degree of nodes in the network (and consequently the minimum degree of the network)*

d	1	2	3	4	5	6	7	≥ 8
cc	0.224	0.343	0.517	0.905	0.993	0.999	0.999	1

TABLE 2 *Intersection distance comparisons of rankings based on the diagonal entries and row sums of e^A for 1000-node scale-free networks of various parameters built using the `pref` function in the CONTEST Matlab toolbox. The values reported are the averages over 20 matrices with the same parameters. The parameter d is the initial degree of nodes in the network (and consequently the minimum degree of the network)*

d	isim	isim _{10%}
1	0.174	0.199
2	0.036	0.031
3	0.003	0.005
4	2.04e−4	2.79e−4
5	1.30e−5	1.71e−5
6	9.83e−7	0
7	4.93e−7	0
≥ 8	0	0

The results show that correlation between the two metrics increases and the intersection distance value decreases quickly with the value of the parameter d . The intersection distance values for the top 10% of nodes are very close to those for the complete set of nodes. For sufficiently dense networks, the two measures provide essentially identical rankings, producing correlation coefficients close to 1 and intersection distances close to 0.

A second class of synthetic test matrices used in our experiments corresponds to small-world networks (Watts–Strogatz model). The small-world model was developed as a way to impose a high clustering coefficient onto classical random graphs [38]. The name comes from the fact that, like classical random graphs, the Watts–Strogatz model produces networks with the small world (that is, small graph diameter) property. To build these matrices, the input is `smallw(n,d,p)` where n is the number of nodes in the network, which are arranged in a ring and connected to their d nearest neighbours on the ring. Then each node is considered independently and, with probability p , a link is added between the node and one of the other nodes in the network, chosen uniformly at random. At the end of this process, all loops and repeated edges are removed. For this set of experiments, the size of the network was fixed at $n = 1000$ and the probability of an extra link was left at the default value of $p = 0.1$ while d was varied.

TABLE 3 *Comparison, using the correlation coefficient, of rankings based on the diagonal entries and row sums of e^A for 1000-node small-world networks of various parameters made using the `smallw` function in the CONTEST Matlab toolbox. The values reported are the average over 20 matrices with the same parameters*

(a)		(b)	
d	cc	d	cc
1	0.177	20	0.156
2	0.089	30	0.222
3	0.037	40	0.240
4	0.033	50	0.310
5	0.031	60	0.426
6	0.048	70	0.431
7	0.039	80	0.747
8	0.046	90	0.926
9	0.031	100	0.997
10	0.054	≥ 110	1

TABLE 4 *Intersection distance comparison of rankings based on the diagonal entries and row sums of e^A for 1000-node small-world networks of various parameters made using the `smallw` function in the CONTEST Matlab toolbox. The values reported are the averages over 20 matrices with the same parameters*

(a)			(b)		
d	isim	isim _{10%}	d	isim	isim _{10%}
1	0.015	0.071	20	0.311	0.713
2	0.056	0.160	30	0.239	0.535
3	0.089	0.252	40	0.133	0.351
4	0.117	0.350	50	0.111	0.214
5	0.151	0.479	60	0.039	0.120
6	0.178	0.621	70	0.014	0.041
7	0.218	0.709	80	0.002	0.007
8	0.243	0.731	90	1.71e-4	4.05e-4
9	0.262	0.705	100	5.88e-6	1.09e-5
10	0.284	0.725	≥ 110	0	0

The values of d tested were: all values $1 \leq d \leq 10$, along with all multiples of 10 up to 200. In each case, 20 networks were created with each value of d . The average correlation coefficients between the subgraph centrality rankings and the total communicability rankings are given in Table 3. As before, the correlation coefficients were computed between the complete sets of rankings. The intersection distances, reported in Table 4, were computed on both the complete sets of rankings and the top 10% of ranked nodes.

It is evident from these results that for this class of small-world networks, the similarity between the two ranking measures is much weaker than for the preferential attachment model, at least as long as the networks remain fairly sparse. The intersection distances are also relatively large, further indicating that

TABLE 5 Comparison of the total network communicability $C(A)$ of a ring lattice and small-world rings with increasing probability of a shortcut. The computed values were averaged over 20 instances

Graph	number of edges	$C(A)$	normalized $C(A)$
5000 node ring lattice	5000	3.69e04	7.4
<code>smallw(5000,1,.1)</code>	5492	4.83e04	9.7
<code>smallw(5000,1,.2)</code>	6222	6.22e04	12.4
<code>smallw(5000,1,.3)</code>	6495	7.92e04	15.8
<code>smallw(5000,1,.4)</code>	6990	9.90e04	19.8
<code>smallw(5000,1,.5)</code>	7496	1.24e05	24.8
<code>smallw(5000,1,.6)</code>	7999	1.53e05	30.6

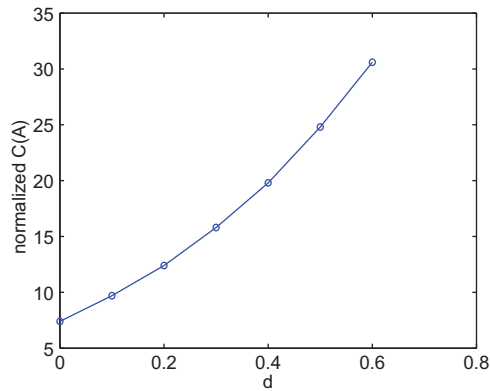


FIG. 1. Plot of the total network communicability $C(A)$ for small-world graphs with increasing probability d of a shortcut. The computed values were averaged over 20 instances.

the two measures are much more weakly related than in the case of the preferential attachment model. For some values of d , the intersection distance between the top 10% of nodes is > 0.7 , indicating that there is little consistency among the rankings of the top 10% of nodes under the two measures. As the networks become increasingly dense, however, the correlation between the two measures becomes stronger and the intersection distance eventually decreases.

5.2 Total communicability in small-world networks

For networks with low connectivity (or high locality), the total network communicability can be expected to be low compared with networks with higher connectivity. For instance, on a 5000 node ring lattice, the total network communicability is $C(A) = 3.69e04$ and the normalized $C(A)$ is 7.4. However, when even a few shortcuts are added across the lattice using the Watts–Strogatz small-world model, this value jumps considerably. If the probability of a shortcut is $p = 0.1$, the normalized total network communicability (averaged over 20 networks created using input `smallw(5000,1,p)`) is 9.7. If the probability of a shortcut is increased to $p = 0.2$, the normalized total network communicability increases to 12.4. These and additional results can be found in Table 5 and Fig. 1.

5.3 Discussion of test results using synthetic data

The results reported so far can be explained as follows. In a (regular) ring-shaped network, no node is more central than the other nodes and no reasonable centrality measure would be able to assign a (strict) ranking of the nodes. In a small-world network obtained by perturbing a regular ring-shaped network, all the nodes have *approximately* the same importance, with the nodes with extra links ('shortcuts') being slightly more important than the others. When d is small, these shortcuts matter more, but the subgraph centrality scores and the total communicability scores do not have a large range. Because of this, the change in the scores due to moving from the subgraph centrality measure to the total communicability measure can have a high impact on node rankings. This leads to a low correlation and a relatively large intersection distance between the two rankings. When d gets very large, the shortcuts matter less and cause less perturbations between the two sets of rankings. By contrast, in a scale-free preferential attachment network both the subgraph centrality scores and total communicability scores are spread out over a large range, even for small d , and adding the corresponding off-diagonal row sums to the diagonal entries does not change the rankings as much.

5.4 Real data

Next, we study correlations between the two ranking methods using various networks corresponding to real data. The networks in this section come from a variety of sources. The Zachary Karate Club network is a classic example in network analysis [39]. The Intravenous Drug User and Yeast PPI networks were provided to us by Prof. Ernesto Estrada. The Yeast PPI network has 440 ones on the diagonal due to the self-interactions of certain proteins. The remainder of the networks can be found in the University of Florida Sparse Matrix Collection [40] under different 'groups'. The Erdős networks are from the Pajek group. They represent various subnetworks of the Erdős collaboration network. The ca-GrQc and ca-HepTh from the SNAP group are collaboration networks for the arXiv General Relativity and High Energy Physics Theory subsections, respectively. The as-735 network, also from the SNAP group, contains the communication network of a group of autonomous systems (AS) measured over 735 days between November 8, 1997 and January 2, 2000. Communication occurs when routers from two ASs exchange information. The Minnesota network from the Gleich group represents the Minnesota road network. The order n and the number of nonzeros nmz of the corresponding adjacency matrices are given in Table 6. These networks exhibit a wide variety of structural properties and together constitute a rather heterogeneous sample of real-world networks. All networks except the Yeast PPI network are simple and all are undirected.

Table 6 reports the correlation coefficients between the two sets of rankings for all the nodes, the top 10% of the nodes and the top 1% of the nodes (limited to the cases where the two methods rank the same nodes in the top 10% and top 1%), as well as the value of the two largest eigenvalues λ_1 and λ_2 of the adjacency matrix. A '-' in the table signifies that different lists of top nodes were produced under the two rankings, hence correlation coefficients could not be computed in such cases. Table 7 reports the intersection distances between the two sets of rankings for all, for the top 10% and for the top 1% of the nodes. Table 8 reports the normalized Estrada index and normalized total network connectivity for each of the networks. For the Zachary Karate Club, which has only 34 nodes, $\alpha_1 = 1$ and $\text{isim}_{1\%} = 0$ indicate that the top two ranked nodes under the two rankings are the same. The top node is node 34, which corresponds to the president of the karate club, and the second is node 1, which corresponds to the instructor. These were the two most influential members of the club and fought with each other to the point that eventually the club split into two factions aligned around each of them [39].

TABLE 6 *Comparison of rankings based on the diagonal and row sum of e^A for various real-world networks*

Graph	n	nnz	λ_1	λ_2	cc	cc_{10}	cc_1
Zachary Karate Club	34	156	6.726	4.977	0.420	—	1
Drug User	616	4024	18.010	14.234	0.083	0.976	1
Yeast PPI	2224	13218	19.486	16.134	0.108	—	1
Pajek/Erdos971	472	2628	16.710	10.199	0.523	1	1
Pajek/Erdos972	5488	14170	14.448	11.886	0.122	—	—
Pajek/Erdos982	5822	14750	14.819	12.005	0.128	—	—
Pajek/Erdos992	6100	15030	15.131	12.092	0.143	—	—
SNAP/ca-GrQc	5242	28980	45.617	38.122	0.021	—	0.995
SNAP/ca-HepTh	9877	51971	31.035	23.004	0.007	—	—
SNAP/as-735	7716	26467	46.893	27.823	0.904	0.771	1
Gleich/Minnesota	2642	6606	3.2324	3.2319	0.087	—	—

TABLE 7 *Intersection distance comparison of rankings based on the diagonal and row sum of e^A for various real-world networks*

Graph	isim	isim _{10%}	isim _{1%}
Zachary Karate Club	0.044	0.111	0
Drug User	0.102	0.002	0
Yeast PPI	0.025	0.056	0
Pajek/Erdos971	0.004	0	0
Pajek/Erdos972	0.081	0.075	0.047
Pajek/Erdos982	0.079	0.065	0.044
Pajek/Erdos992	0.077	0.055	0.034
SNAP/ca-GrQc	0.043	0.091	5.49e−4
SNAP/ca-HepTh	0.142	0.319	0.134
SNAP/as-735	1.81e−4	0.001	0
Gleich/Minnesota	0.096	0.341	0.709

The results indicate that there is a good deal of variation between the correlation coefficients for these networks. The correlation coefficient between the rankings of all the nodes ranges from a low of 0.007 for the SNAP/ca-HepTh network to a high of 0.904 for the SNAP/as-735 network. Even for networks that come from similar datasets, the correlation coefficients can be very different. For example, the networks in the Pajek group are all subsets of the Erdős collaboration network, but correlations between the two sets of rankings range between 0.122 for the Erdos972 network and 0.583 for the Erdos971 network.

For most of the networks, the correlation coefficient (when defined) increases when only the top 1% of nodes are considered (cc_1), sometimes greatly. Five of the networks (Zachary Karate Club, Drug User, Yeast PPI, Pajek/Erdos971, and SNAP/as-735) produce the exact same rankings on the top 1% of nodes. Another network (SNAP/ca-GrQc) has a correlation coefficient greater than 0.9 on the top 1% of nodes.

The intersection distance values behave in a similar way, although there is not as much variation in the values. Among all the nodes, the smallest intersection distance is 1.81e−4 for the as-735 network

TABLE 8 Comparison of the normalized Estrada index $EE(A)/n$, the normalized total network connectivity $C(A)/n$, and $e^{\|A\|_2}$ ($= e^{\lambda_1}$) for various real-world networks

Graph	Normalized $EE(A)$	Normalized $C(A)$	$e^{\ A\ _2}$
Zachary Karate Club	30.62	608.79	833.81
Drug User	1.12e05	1.15e07	6.63e07
Yeast PPI	1.37e05	3.97e07	2.90e08
Pajek/Erdos971	3.84e04	4.20e06	1.81e07
Pajek/Erdos972	408.23	1.53e05	1.88e06
Pajek/Erdos982	538.58	2.07e05	2.73e06
Pajek/Erdos992	678.87	2.50e05	3.73e06
SNAP/ca-GrQc	1.24e16	8.80e17	6.47e19
SNAP/ca-HepTh	3.05e09	1.06e11	3.01e13
SNAP/as-735	3.00e16	3.64e19	2.32e20
Gleich/Minnesota	2.86	14.13	35.34

and the largest is 0.142 for the ca-HepTh network. These networks also had the largest and smallest correlation coefficients, respectively, for the full set of nodes. For 5 of the 11 networks examined, the intersection distance value decreases when only the top 10% of nodes are considered and for all cases except for the Minnesota road network, it decreases when only the top 1% of nodes are considered.

It is interesting to note that the similarity between the two ranking methods is very different on the ca-GrQc and the ca-HepTh networks. The two networks are both arXiv collaboration networks from subsections of physics so, intuitively, one would assume that they behaved similarly. However, the two rankings are very different on the ca-HepTh network and are highly correlated on the ca-GrQc network. The ca-GrQc network has a spectral gap of ~ 7.5 while the spectral gap of ca-HepTh is ~ 8 , only slightly larger. The relative spectral gaps are also comparable. Thus, it is clear that the spectral gap alone cannot be used to differentiate between the two ranking methods. It appears that while the two networks are both physics collaboration networks, there are significant structural differences between the two groups that cause the two ranking systems to behave very differently. Some insights can be gleaned by looking at the degree distributions of the two networks. Although the ca-HepTh network is almost twice as large as the ca-GrQc, the maximum degree on the network is only 65 while the maximum degree on the ca-GrQc network is 81. See Fig. 2 for the degree distributions of the two networks. Additionally, the total communicability scores achieved by nodes in the ca-GrQc network range from 2.7 to 8.5e19 (the subgraph centrality scores range from 1.5 to 1.6e18). In contrast, even with many more nodes, the total communicability scores of the ca-HepTh network have a smaller range, from 2.7 to 3.2e13 (the subgraph centrality scores range from 1.5 to 9.7e11). It appears that the wider range of scores in the ca-GrQc network helps one to prevent rankings from being changed when the scores are perturbed by the addition of off-diagonal communicabilities. This can be observed when looking at the intersection distances between the two sets of rankings on the networks, which are plotted in Fig. 3. Overall, the intersection distances are much lower for the ca-GrQc network than for the ca-HepTh network. Additionally, for $k \leq 34$, $\text{isim}_k(\text{ca-GrQc}) = 0$, indicating that the first 34 nodes are ranked exactly the same. In contrast, $\text{isim}_k(\text{ca-HepTh}) = 0$ only for $k \leq 5$, after which there is a large jump in the intersection distances.

Similar behaviour can be observed on the various instances of the Erdős collaboration network. Erdos971, which is very small, shows a high correlation between the two rankings; indeed, the rankings of the top 10% of nodes are exactly the same. On the other instances of the collaboration network, however, the rankings are somewhat different, as can be seen from the relatively low values of the

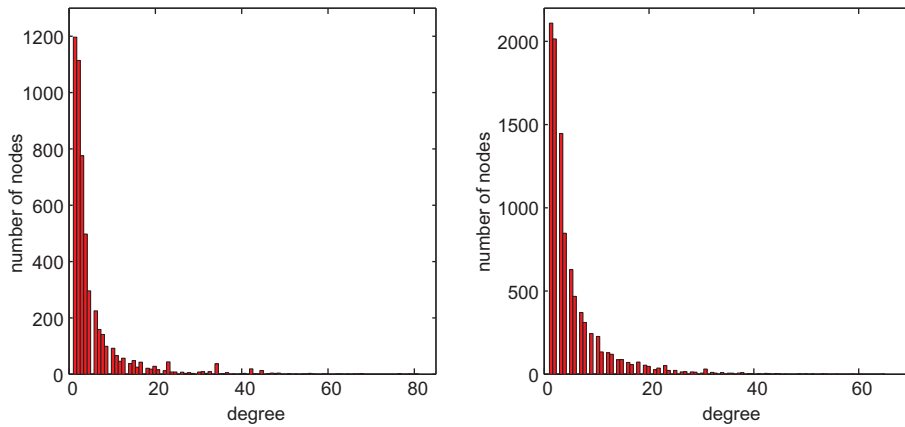


FIG. 2. The degree distributions of the ca-GrQc (left) and the ca-HepTh (right) collaboration networks.

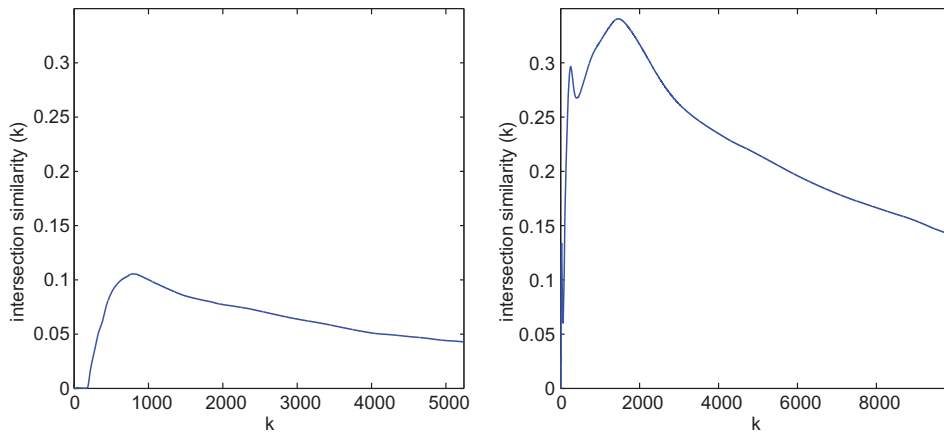


FIG. 3. The intersection distance values (isim_k) of the ca-GrQc (left) and the ca-HepTh (right) collaboration networks.

correlation coefficients. The intersection distance values, while not very high, are somewhat higher than for most other networks. The maximum subgraph centrality and total communicability scores of the Erdos972 network are the smallest of any of the Erdős collaboration subgraphs. The maximum subgraph centrality score is $1.18\text{e}05$ and the maximum total centrality score is $9.20\text{e}06$. By comparison, on the (much smaller) Erdos971 network, the maximum subgraph centrality score is $1.11\text{e}06$. On the Erdos982 network, the maximum subgraph centrality score is $1.71\text{e}05$ and on the Erdos992 network it is $2.47\text{e}05$. Although the top five nodes of the Erdos972 network are exactly the same under the two ranking schemes, the relatively narrow range of possible scores means that the addition of off-diagonal values to the diagonal ones perturbs the rankings of the other nodes so much as to result in a relatively high value of the intersection distance among the top 1% of nodes.

As before, the spectral gap for these networks does not give much insight into the behaviour of the two ranking schemes, unless it is really large; the largest spectral gap for this set of test problems occur

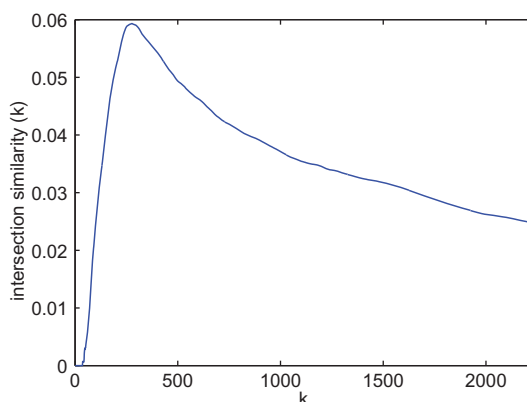


FIG. 4. The intersection distance values (isim_k) of the Yeast PPI network.

for SNAP/as-735, and indeed here we observe a strong correlation and a small intersection distance between the two metrics. Conversely, for the (planar, fairly regular) Gleich/Minnesota network, the spectral gap is smallest and not surprisingly the correlation is very weak and the intersection distance for the top 1% of the nodes, $\text{isim}_{1\%}$, is very high at 0.709.

When examining the (normalized) total network connectivities of the various networks (see Table 8), it can be seen that the ease of information sharing across the networks varies widely. Some networks, such as the collaboration networks ca-HepTh and ca-GrQc, have a high normalized $C(A)$ ($8.80\text{e}17$ and $1.06\text{e}11$, respectively). The value is even higher for the SNAP/as-735 router network ($C(A)/n = 3.64\text{e}19$). The Minnesota road network, on the other hand, has a normalized $C(A)$ of only 14.13, indicating that the network is relatively poorly connected, as one would expect in a graph characterized by wide diameter, small bandwidth and high locality.

5.5 Identification of essential proteins in PPI network of yeast

One important application of node centrality measures is to rank nodes in protein–protein interaction (PPI) networks in an attempt to determine which proteins are essential, in the sense that their removal would result in the death of the cell. The goal of such rankings is for as many of the top-ranked nodes as possible to correspond to essential proteins. In [41], various centrality measures were tested on their ability to identify essential proteins in the Yeast PPI network. It was shown that, among the centrality measures tested, subgraph centrality identified the highest percentage of essential proteins ranked in the top 30 nodes, identifying 18 essential proteins (in [41], subgraph centrality was said to identify 19 essential proteins, but this was later corrected [42]). When total communicability is used instead, the top 30 nodes are the same, so the same percentage of essential proteins are identified. The intersection distances between the two sets of rankings are displayed in Fig. 4. Here, it can be seen that the intersection distances are small for approximately the top 50 nodes, then they begin to rise. The two rankings are least similar for nodes ranked 200–500, then their similarity increases again. As already noted, total communicability rankings can be calculated much more quickly than subgraph centrality rankings (see also Section 6). Although there are currently methodologies which do better in protein ranking (see [43] for example), our findings suggest that total communicability does provide valuable information about the relative importance of nodes in the network.

5.6 Further discussion of test results using real networks

The results just described indicate that in general the two centrality measures can produce significantly different rankings, even when one restricts the attention to the top 1% of nodes, and even for networks belonging to the same ‘family’. As in the case of synthetic networks, a wider range of values in the two sets of centralities leads to stronger correlations between the corresponding rankings than in the case of a narrow range.

Two extreme cases are represented by the SNAP/as-735 and Gleich/Minnesota data sets. The first one exhibits a large value of the spectral gap, and thus (as expected) a strong correlation between the two rankings; the second one has tiny spectral gap and results in very weakly correlated rankings. For networks that fall somewhere in between these two extremes, the observed correlation coefficients can vary significantly. The subgraph centrality scores measure how ‘well-connected’ a node is in the network as a whole while the communicability score between nodes i and j measures how well information travels between node i and node j . Thus, the total communicability of node i is a measure of how well information travels between node i and any node in the network (node i itself included). Although these two measures are closely related, they are not quite the same. This observation suggests that the two centrality measures reflect somewhat different structural properties of the networks. Thus, they should be applied in concert rather than in alternative of one another, unless computational considerations dictate otherwise.

6. Computational aspects

There are various methods available to compute (or approximate) the matrix exponential. One of the most used schemes (which is implemented in Matlab as the `expm` function) is based on Padé approximations combined with scaling and squaring [23,44]. For a generic $n \times n$ matrix, this requires $O(n^3)$ arithmetic operations and $O(n^2)$ storage. The prefactor multiplying n^3 in the arithmetic complexity can vary widely depending on the sparsity and structural properties of A .

Once the matrix exponential is computed, both the subgraph centrality and the total communicability rankings are readily obtained. However, to compute the subgraph centrality rankings, we do not need the complete matrix exponential, we only need the diagonal entries of e^A . Methods for efficiently estimating individual entries of matrix functions have been developed by Golub, Meurant, and others [30,31] and these methods have previously been applied to network analysis [15,22]. They are based on Gaussian quadrature and the Lanczos algorithm, and they have been implemented in the Matlab toolbox `mmq` [45]. The cost per node of estimating the subgraph centrality is typically $O(n)$, giving a total cost of approximately $O(n^2)$ for estimating the subgraph centrality for every node and computing the subgraph centrality rankings. However, the coefficient of the $O(n)$ estimate can be quite large. Additionally, the `mmq` toolbox-based implementation for calculating subgraph centrality that we use here has not been optimized, unlike the built-in Matlab function `expm`. We mention in passing that methods for quickly determining the top k nodes and only calculating the exact rankings on this subset have also been developed [15,46].

The individual entries of the matrix exponential are not necessary for computing the total communicability rankings; only the row sums of e^A are necessary. An efficient algorithm for evaluating $f(A)\mathbf{v}$ using a restarted Krylov method has recently been presented in [47,48]. In this approach, the basic operation is represented by matrix-vector products with A . This method has been implemented in the Matlab toolbox `funm_kryl` by Stefan Güttel [49]. We apply this algorithm with $f(A) = e^A$ and $\mathbf{v} = \mathbf{1}$. Clearly, the same algorithm can be used to rapidly compute $C(A) = \mathbf{1}^T e^A \mathbf{1}$. For many network of

TABLE 9 Timings (in seconds) to compute centrality rankings based on the diagonal and row sum of e^A for various test problems using different methods

Graph	expm	mmq	funm_kryl
Zachary Karate Club	0.062	0.138	0.120
Drug User	0.746	2.416	0.363
Yeast PPI	47.794	9.341	0.402
Pajek/Erdos971	0.542	2.447	0.317
Pajek/Erdos972	579.214	35.674	0.410
Pajek/Erdos982	612.920	39.242	0.393
Pajek/Erdos992	656.270	53.019	0.325
SNAP/ca-GrQc	281.814	23.603	0.465
SNAP/ca-HepTh	2710.802	58.377	0.435
SNAP/as-735	2041.439	75.619	0.498
Gleich/Minnesota	1.956	10.955	0.329

practical interest, the cost is typically $O(n)$, although the prefactor can vary considerably for different types of networks.

Table 9 lists the timings for calculating the matrix exponential directly using `expm`, estimating the subgraph centralities using the `mmq` toolbox (with five iterations of the Lanczos algorithm per node), and estimating the total communicabilities using the `funm_kryl` toolbox to estimate $e^A \mathbf{1}$ (using a very stringent stopping tolerance of $1e-16$). These computations have been performed using Matlab Version 7.9.0 (R2009b) on a 2.4 GHz Intel Core i5 processor with 4 GB of RAM. In general, the timings with `expm` increase for increasing number of nodes, but structural properties of the underlying graph, like the network diameter, can have a very significant impact on the computing times. For example, the yeast PPI network and the Minnesota road network have approximately the same number n of nodes (2224 and 2642, respectively), yet computing the matrix exponential for the yeast network takes almost 25 times longer than for the Minnesota road network. This appears to be due to the fact that the yeast network has a much smaller diameter than the Minnesota network, therefore the powers A^k of the adjacency matrix fill up much more quickly. Since the algorithm implemented in `expm` involves solving linear systems with polynomials in A as coefficient matrices, the execution time for sparse matrices with small diameter tends to be much higher than for matrices exhibiting a high degree of locality.

For the majority of the networks tested, using the `mmq` toolbox to estimate subgraph centrality was faster than using `expm`, frequently by far. The exceptions (Zachary Karate Club, Drug User, Erdos971, and Minnesota) were the networks with a small number of nodes and/or a high diameter.

The computation of the total communicabilities using `funm_kryl` was by far the fastest method for all networks tested, with the only exception of the tiny Zachary Karate Club network. In principle, this is a clear advantage of total communicability over subgraph centrality. However, as we saw, the two methods often result in rather different rankings, therefore we cannot simply replace subgraph centrality with total communicability.

6.1 A large-scale example

In addition to the test results discussed above, we performed tests with the digraph of Wikipedia (as of June 6, 2011), where nodes correspond to entries and directed links to hyperlinks from one entry to

another. In this case, the entries of $e^A \mathbf{1}$ provide a ranking of the hubs in the networks, see [15]. This graph contains 4,189,503 nodes and 67,197,636 links, and it is prohibitively large for centrality measures based on estimating the diagonals of the matrix exponential. For this reason, we limit ourselves to computations using the `funm_kryl` toolbox to estimate the row sum vector $e^A \mathbf{1}$. The restart parameter was set to 10 and we allowed a maximum of 50 restarts. The run time to obtain the rankings on a parallel system comprising 24 Intel(R) Xeon(R) E5-2630 2.30 GHz CPU(s) was 216.7 seconds. This shows that centrality calculations using total communicability are quite feasible even for large networks.

7. Resolvent-based centrality measures

There are matrix functions other than the matrix exponential that may be used to calculate subgraph centrality and subgraph communicability scores. The most common of these is the matrix resolvent

$$(I - \alpha A)^{-1} = I + \alpha A + \alpha^2 A^2 + \cdots + \alpha^k A^k + \cdots = \sum_{k=0}^{\infty} \alpha^k A^k, \quad (7.1)$$

where $0 < \alpha < 1/\rho(A)$, with $\rho(A)$ the spectral radius of A . This was first used by Katz [16] in the 1950s and has been used in various forms since then [5,6,10–12,21,50]. The bounds on α ensure that $I - \alpha A$ is invertible and that the geometric series converges to the inverse. Additionally, the inverse is nonnegative; indeed, $I - \alpha A$ is a nonsingular M -matrix. Note that if A is the adjacency matrix of an undirected network, $\rho(A) = \lambda_{\max}(A) = \|A\|_2$. Since the spectral radius of a nonnegative matrix always satisfies $\rho(A) \geq \min_i \sum_{j=1}^n a_{ij}$, it follows that for a connected undirected graph α must be less than 1.

Like the matrix exponential, $[(I - \alpha A)^{-1}]_{ii}$ counts the number of closed walks centered at node i and $\sum_{j=1}^n [(I - \alpha A)^{-1}]_{ij}$ counts all walks between node i and all other nodes in the network. In this case, however, a walk of length k is penalized by a factor of α^k . One drawback of the use of the matrix resolvent in determining centrality rankings is the need to choose the value of α ; also, different values of α can lead to different rankings. For the purposes of the experiments below, we select $\alpha = 0.85/\lambda_{\max}(A)$ (similar to the choice of parameter in PageRank [6]).

Resolvent-based total network communicability can also be evaluated. As when using the matrix exponential (cf. Section 4), the resolvent-based total network communicability is an upper bound for the resolvent-based Estrada index. In the following, $C_r(A) = \sum_{i=1}^n \sum_{j=1}^n [(I - \alpha A)^{-1}]_{ij}$ denotes the resolvent-based total communicability of a network. The following proposition can be easily proved along the same lines as Proposition 1.

PROPOSITION 2 Let A be the adjacency matrix of a simple, undirected network on n vertices. Then, for any $0 < \alpha < 1/\|A\|_2$,

$$EE_r(A) := \text{Tr}[(I - \alpha A)^{-1}] \leq C_r(A) \leq \frac{n}{1 - \alpha\|A\|_2}.$$

For an undirected network, $\lambda_{\max}(A) = \lambda_1$ can replace $\|A\|_2$ in the upper bound above.

The resolvent-based subgraph centrality and total communicability rankings were compared on the same two sets of synthetic networks used for the tests in Section 5.1.

Table 10 lists the average correlation coefficient between the subgraph centrality and total communicability rankings for the nodes in networks constructed using the preferential attachment model (function `pref` in `CONTEST`) and Table 11 lists the intersection distances for all the nodes and for the top 10% of the nodes. For small values of d ($1 \leq d \leq 3$), the correlation coefficients between the two sets of rankings using the matrix resolvent are close to those using the matrix exponential. However,

TABLE 10 Comparison using correlation coefficients of rankings based on the diagonal entries and row sums of $(I - \alpha A)^{-1}$ for 1000-node scale-free networks of various parameters built using the `pref` function in the CONTEST Matlab toolbox. For each instance, the results are measured for $\alpha = 0.85/\lambda_{\max}(A)$. The values reported are the averages over 20 matrices with the same parameters

(a)		(b)		(c)	
d	cc	d	cc	d	cc
1	0.292	10	0.691	110	0.964
2	0.370	20	0.840	120	0.965
3	0.442	30	0.890	130	0.968
4	0.486	40	0.917	140	0.970
5	0.536	50	0.933	150	0.971
6	0.583	60	0.942	160	0.973
7	0.607	70	0.949	170	0.973
8	0.638	80	0.954	180	0.975
9	0.667	90	0.958	190	0.976
		100	0.962	200	0.976

TABLE 11 Intersection distance comparison of rankings based on the diagonal entries and row sums of $(I - \alpha A)^{-1}$ for 1000-node scale-free networks of various parameters built using the `pref` function in the CONTEST Matlab toolbox. For each instance, the results are measured for $\alpha = 0.85/\lambda_{\max}(A)$. The values reported are the averages over 20 matrices with the same parameters

(a)			(b)			(c)		
d	isim	isim _{10%}	d	isim	isim _{10%}	d	isim	isim _{10%}
1	0.186	0.491	10	0.105	0.051	110	0.006	0.001
2	0.205	0.364	20	0.055	0.020	120	0.005	7.12e-4
3	0.192	0.235	30	0.035	0.012	130	0.005	6.98e-4
4	0.179	0.173	40	0.025	0.007	140	0.004	5.74e-4
5	0.163	0.126	50	0.019	0.005	150	0.004	5.62e-4
6	0.150	0.102	60	0.015	0.004	160	0.003	3.69e-4
7	0.137	0.082	70	0.012	0.003	170	0.003	4.25e-4
8	0.124	0.068	80	0.010	0.002	180	0.003	3.11e-4
9	0.115	0.059	90	0.009	0.002	190	0.003	3.16e-4
			100	0.007	0.001	200	0.002	4.00e-4

when using the matrix exponential the average correlation coefficient was found to be greater than 0.9 for all $d \geq 4$, and exactly 1 for all $d \geq 8$. Using the matrix resolvent the correlation coefficient grows as d increases, but somewhat more slowly than for the matrix exponential. The intersection distances are also larger for all values of d when the matrix resolvent is used, although they also decrease as d increases. Moreover, we did not find a single instance where the two methods produced *exactly* the same rankings.

TABLE 12 *Comparison using correlation coefficients of rankings based on the diagonal entries and row sums of $(I - \alpha A)^{-1}$ for 1000-node small-world networks of various parameters built using the `smallw` function with $p = 0.1$ in the CONTEST Matlab toolbox. For each instance, the results are measured for $\alpha = 0.85/\lambda_{\max}(A)$. The values reported are the averages over 20 matrices with the same parameters*

(a)		(b)		(c)	
d	cc	d	cc	d	cc
1	0.065	10	0.063	110	0.294
2	0.023	20	0.078	120	0.246
3	0.052	30	0.080	130	0.275
4	0.052	40	0.135	140	0.311
5	0.052	50	0.144	150	0.312
6	0.051	60	0.141	160	0.321
7	0.062	70	0.144	170	0.301
8	0.037	80	0.133	180	0.293
9	0.050	90	0.248	190	0.354
		100	0.190	200	0.300

TABLE 13 *Intersection distance comparison of rankings based on the diagonal entries and row sums of $(I - \alpha A)^{-1}$ for 1000-node small-world networks of various parameters built using the `smallw` function with $p = 0.1$ in the CONTEST Matlab toolbox. For each instance, the results are measured for $\alpha = 0.85/\lambda_{\max}(A)$. The values reported are the averages over 20 matrices with the same parameters*

(a)			(b)			(c)		
d	isim	isim _{10%}	d	isim	isim _{10%}	d	isim	isim _{10%}
1	0.040	0.149	10	0.156	0.435	110	0.147	0.541
2	0.070	0.189	20	0.207	0.508	120	0.148	0.553
3	0.085	0.241	30	0.198	0.517	130	0.160	0.554
4	0.091	0.269	40	0.204	0.571	140	0.142	0.560
5	0.098	0.301	50	0.207	0.621	150	0.123	0.542
6	0.104	0.318	60	0.191	0.588	160	0.121	0.539
7	0.126	0.361	70	0.181	0.582	170	0.124	0.517
8	0.135	0.414	80	0.189	0.607	180	0.125	0.512
9	0.149	0.413	90	0.156	0.597	190	0.114	0.504
			100	0.179	0.585	200	0.123	0.504

For the small-world networks, all values $1 \leq d \leq 10$ as well as all multiples of 10 with $20 \leq 10 \leq 200$ were tested. For each d , 20 networks were tested. The averages of the correlation coefficients between the subgraph centrality and total communicability rankings can be found in Table 12 and the average intersection distances for both all the nodes and the top 10% of the nodes can be found in Table 13. As was the case for the matrix exponential, the two methods (diagonal entries and row sums) using the matrix resolvent exhibit much weaker correlations for this class of networks than for the preferential attachment networks; indeed, the correlations tend to be even smaller for the resolvent than

TABLE 14 *Comparison using correlation coefficients of rankings based on the diagonal entries and row sums of $(I - \alpha A)^{-1}$ with $\alpha = 0.85/\lambda_{\max}(A)$ for various real-world networks*

Graph	cc	cc_{10}	cc_1
Zachary Karate Club	0.589	1	1
Drug User	0.189	—	—
Yeast PPI	0.177	—	—
Pajek/Erdos971	0.233	—	1
Pajek/Erdos972	0.215	—	—
Pajek/Erdos982	0.207	—	—
Pajek/Erdos992	0.197	—	—
SNAP/ca-GrQc	0.070	—	—
SNAP/ca-HepTh	0.072	—	—
SNAP/as-735	0.204	—	—
Gleich/Minnesota	0.019	—	—

for the exponential. For $d = 1$, the average correlation is 0.065 and the average intersection distance was 0.040 using the resolvent, compared to a correlation of 0.177 and an intersection distance of 0.015 using the exponential. For the values of d tested, the highest average correlation coefficient was 0.354, for $d = 190$. When looking at the intersection distances for other values of d , the picture is somewhat different. Comparing Table 13 with Table 4, we see that for small d the intersection distance between the two ranking schemes tends to be somewhat higher with the matrix exponential than with the resolvent. However, as d increases the intersection distance eventually drops with the matrix exponential, but not with the resolvent. This is true both when looking at the ranking of all the nodes and when looking at only the top 10%.

Next, we consider tests with real-world networks. As shown in Table 14, the correlation coefficients between the two ranking systems for the whole set of nodes were higher (in a majority of cases) using the matrix resolvent than they were using the matrix exponential. (Again, a ‘—’ signifies that correlation coefficients could not be computed due to the fact that the two ranking schemes produced different lists of nodes.) Only the Erdos971, as-735 and the Minnesota networks had a higher correlation coefficient between the two ranking systems under the exponential than under the matrix resolvent. This can be understood when looking at the normalized Estrada indexes and total network communicabilities in Table 15. The smaller the factor α , the more it minimizes the contribution of the network data from A to the scores produced by the diagonal entries or row sums of $(I - \alpha A)^{-1}$. This can also be seen by noticing that as $\alpha \rightarrow 0$, $(I - \alpha A)^{-1}$ approaches the identity. In these experiments, $\alpha = 0.85/\lambda_{\max}(A)$. However, this also means that for the networks tested with a large maximum eigenvalue (such as ca-GrQc, ca-HepTh, and as-735) α is quite small, causing the resulting subgraph centrality scores to be small and, consequently, close together. In the case of a network with a small maximum eigenvalue (such as the Minnesota network), the effect of α is not as pronounced. The compression of the score values means that a perturbation of the scores (such as occurs when switching from subgraph centrality scores to total communicability scores) has a large effect on the node rankings, especially for the higher ranked nodes.

When only the top 1% of nodes were considered, the exponential subgraph centrality and exponential total communicability rankings were much closer together than their resolvent counterparts, where often the top 1% of nodes were not even the same. This seems to indicate that when using the resolvent,

TABLE 15 *Comparison of the normalized resolvent-based Estrada index $EE_r(A)/n$ and total network connectivity $C_r(A)/n$ for various real-world networks. Here, $f(A) = (I - \alpha A)^{-1}$ with $\alpha = 0.85/\lambda_{\max}(A)$*

Graph	normalized $EE_r(A)$	normalized $C_r(A)$
Zachary Karate Club	1.21	5.13
Drug User	1.03	2.36
Yeast PPI	1.03	2.17
Pajek/Erdos971	1.03	2.44
Pajek/Erdos972	1.01	1.70
Pajek/Erdos982	1.01	1.66
Pajek/Erdos992	1.01	1.60
SNAP/ca-GrQc	1.00	1.21
SNAP/ca-HepTh	1.01	1.24
SNAP/as-735	1.00	1.86
Gleich/Minnesota	1.27	3.44

TABLE 16 *Intersection distance comparison of rankings based on the diagonal entries and row sums of $(I - \alpha A)^{-1}$ with $\alpha = 0.85/\lambda_{\max}(A)$ for various real-world networks*

Graph	isim	isim _{10%}	isim _{1%}
Zachary Karate Club	0.061	0	0
Drug User	0.125	0.145	0.069
Yeast PPI	0.204	0.363	0.187
Pajek/Erdos971	0.080	0.050	0
Pajek/Erdos972	0.110	0.273	0.263
Pajek/Erdos982	0.109	0.269	0.264
Pajek/Erdos992	0.109	0.271	0.247
SNAP/ca-GrQc	0.047	0.122	0.033
SNAP/ca-HepTh	0.058	0.159	0.236
SNAP/as-735	0.247	0.513	0.271
Gleich/Minnesota	0.102	0.301	0.557

the subgraph centrality and total communicability tend to rank the less important nodes more similarly than they do under the matrix exponential. Under the matrix exponential, the two rankings seem to agree more closely on the important nodes than they do when using the resolvent. This can also be seen when looking at the intersection distance, which gives more weight to differences in the top ranked nodes than in the lower ranked nodes (Table 16). For all networks except ca-HepTh, the intersection distance between the two rankings is smaller when using the exponential than when using the resolvent. When looking at the top 1% of nodes, the intersection distances are also smaller (often much smaller) in the case of the exponential, for all except three of the networks. The exceptions are the Minnesota road network (which has a large intersection distance on the top 1% of nodes for both the exponential and the resolvent) and the Zachary Karate Club and Erdos971 networks (which have $\text{isim}_{1\%} = 0$ for both cases).

Another observation that can be made is that the resolvent-based total network communicability $C_r(A)$ is unable to discriminate between highly connected networks and poorly connected ones, in stark

contrast with the exponential-based one. For instance, in the case of the Minnesota road network α is relatively large (since λ_1 is small for this graph), hence the off-diagonal contributions to $C_r(A)$ are more significant than for other networks where λ_1 is large (thus forcing a small value of α , leading to a resolvent very close to the identity matrix). Thus, only the exponential-based total network communicability should be used when comparing different networks in terms of ease of communication.

When the identification of essential proteins in the Yeast PPI network is considered using resolvent-based total communicability, the results are comparable to those using the exponential. The resolvent-based total communicability rankings with $\alpha = 0.85/\lambda_{\max}(A)$ identified 17 essential proteins in the top 30 (when compared with 18 identified by exponential subgraph centrality and total communicability). The resolvent-based subgraph centrality, however, identified 19 essential proteins in the top 30, slightly outperforming the other methods.

Concerning the computational complexity, when dealing with large networks the use of the conjugate gradient method (possibly with some type of preconditioning) to solve the linear system $(I - \alpha A)\mathbf{x} = \mathbf{1}$ is orders of magnitude faster than trying to estimate the diagonal entries of $(I - \alpha A)^{-1}$. For certain networks, Chebyshev semi-iteration can be even faster [51]. Thus, as was the case for the matrix exponential, rankings based on total communicability (row sums) are a lot cheaper than the rankings based on subgraph centrality (diagonals). Once again, however, the two ranking methods in general produce different rankings, so one should not choose between the two based solely on computational cost.

8. Conclusions

We have examined the use of total communicability as a method for ranking the importance of nodes in a network. Like the subgraph centrality ranking, the total communicability ranking using the matrix exponential counts the number of walks starting at a given node, weighing walks of length k by a penalization factor of $1/k!$. However, instead of only counting closed walks, it counts all walks between the given node and every node in the network. If the matrix resolvent is used, the weight on the walks becomes α^k for a chosen parameter α in a certain range. There are various classes of graphs on which it can be shown that the two exponential-based rankings are always identical or in very good agreement; for instance, certain types of simple regular graphs and Erdős–Rényi random graphs with large spectral gap. However, as is well known, these classes are not realistic models of real-world complex networks.

The two sets of rankings (total communicability and subgraph centrality) have been used to rank the nodes of networks corresponding to both real and synthetic data sets. The synthetic data sets were constructed using the preferential attachment (Barabási–Albert) and the small-world (Watts–Strogatz) models, corresponding to the functions `pref` and `smallw` of the CONTEST toolbox for Matlab. Good agreement between the two ranking methods was observed on the networks obtained with the preferential attachment method, especially as the density of the graphs increased. More pronounced differences between the rankings produced with the two methods were observed in the case of small-world networks. Overall, the two importance rankings matched more closely when the matrix exponential was used than when under the matrix resolvent.

We also presented the results of experiments with real-world networks including social networks, citation networks, PPI networks and infrastructure (transportation) networks. Here, we found that overall, the two (complete) sets of rankings were closer to each other when the matrix resolvent was used instead of the matrix exponential. However, when only the top 1% of nodes was examined, the rankings matched more closely when the matrix exponential was used. This suggests that, for the networks

tested, the resolvent-based rankings match more closely on ‘unimportant’ (low-ranked) nodes and the exponential-based rankings exhibit more agreement on the ‘important’ (top-ranked) nodes.

In general, there is no simple way to compare two ranking schemes and determine that one is ‘better’ than the other. However, the total communicability rankings take into account more of the network topology than the subgraph centrality rankings (all walks starting at node i versus all closed walks starting at node i). This added information often (but not always) changes the ranking of the nodes to a certain degree, although there are many cases where there is still a strong similarity between the two sets of rankings. The main benefit of using total communicability to rank the nodes is that the ranking can be estimated extremely quickly using Krylov subspace methods. Indeed, as the Wikipedia graph calculation described in section 6.1 shows, for very large networks only the total communicability (row sum) method is computationally feasible, the subgraph centrality ranking being prohibitively expensive to compute. Even if total communicability cannot always be recommended as a cheaper alternative to subgraph centrality, it provides valuable information about the network and can be used along with other ranking schemes.

Finally, we have introduced the total communicability of a network as a global measure of connectivity and of the ease of information flow on a given network. This measure can be computed quickly even for very large networks, and could be of interest in the design of communication networks.

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