Q-matrix: An Algebraic Formulation for the Analysis and Visual Characterization of Network Graphs

Roldan Pozo

National Institute of Standards and Technology, Gaithersburg, MD 20899

roldan.pozo@nist.gov

Given an undirected network, we describe a two-dimensional graphical measure based on the connected component distribution of its degree-limited subgraphs. This process yields an unambiguous visual *portrait*, which reveals important network properties. It can be used as a classification tool, as graphs from related application areas have striking similarities. It can also be used as an efficient algorithm to demonstrate graph non-isomorphism for large graphs with identical degree distributions. Finally, it can be used as an analysis tool to help distinguish real-world networks from their synthetic counterparts.

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1. Introduction

Attempting to represent a large-scale network as a small picture or a thumbnail image can prove to be a challenging task. Most application networks (e.g. biological, information, social) tend to have large hubs (heavy-tailed degree distributions) [1] and exhibit small-world properties [2], making their layout difficult to embed in two or three-dimensional spaces [3]. Current state-of-the-art algorithms for graph layout and visualization often render such objects as densely colored disks, or entangled "hairballs," making it difficult to extract meaningful information from their appearance. Furthermore, graph layout algorithms do not yield unique images; a single graph may yield many variations, depending on parameter and algorithmic choices. This situation is certainly understandable – it would be rather optimistic to expect graphs containing millions of vertices and edges crammed into a small snapshot (say, a 300×300 pixel image) to yield much insight.

Instead, we offer a different approach based on a simple idea: rather than draw the graph itself, represent the component size distribution of its degree-limited subgraphs. We define the Q-matrix of an undirected graph G to be the matrix formulation Q, where Q_{ij} is the number of connected components of size j of the degree-limited subgraph of G consisting of vertices with degree i or lower. The matrix Q, which is typically sparse, can be thought of as a generalization of the graph's degree-distribution. It also reveals such characteristics as the number of connected components, the formation and growth of the giant component, and the effect of node-removal (site percolation) [4] on the connectivity of the remaining subgraphs. These attributes are useful, for example, in simulations of network reliability [5] and the spread of infectious diseases [6].

Visualizations of the matrix Q can serve as useful network *portraits*. That is, networks from different application areas yield visually distinct portraits (Fig. 3) while networks from the *same* application area bear a strong resemblance. Furthermore, given a network graph, there is only one Q-matrix representation. Visualizations, such as those in Fig. 3, are just a three-dimensional view obtained by mapping the nonzero values of the Q-matrix to the z-axis, and can be easily rendered within scientific or visualization software environments.

2. Mathematical Formulation

Given an undirected graph G = (V, E), with n vertices, m edges, its degree distribution can be described as a vector $\vec{d}(G) \equiv \langle d_0, d_1, d_2, \dots d_{n-1} \rangle$, where each d_i is the number of vertices in G with degree equal to i. Note that if D denotes the largest degree of any vertex in G, then $d_i = 0$ for all i > D, so \vec{d} is typically truncated to a shorter length. We can then define the **degree-limited subgraph** $G_i = G[V_i]$ as the induced subgraph created from vertices of G which have degree less than or equal to i. That is, $G_i \equiv (V_i, E_i)$ where

$$V_i = \{ v \in V | \mathbf{degree}(v, G) \le i \}$$
 (1)

and

$$E_i =: \{ (u, v) \in E | u, v \in V_i \}$$
 (2)

where degree(v, G) denotes the degree of vertex v in graph G.

Alternatively, the subgraphs G_i can be thought of as what remains when every vertex of degree larger than i, together with every edge touching such vertices are removed from the original graph. Viewed either way, these degree-limited subgraphs are often comprised of disconnected components, even if the original graph is completely connected. By analyzing not only the number of components, but also their size distribution we can render interesting visualizations that are unique for each network (i.e., invariant under graph isomorphisms) and illustrate fundamental properties of the graph's structure.

Define the Q-matrix of a graph G as the two-dimensional component size distribution of its degree-limited subgraphs. Specifically, let $\Pi(A)$ denote the number of connected components in graph A, and let $\Pi_i(A)$ be the number of connected components which have exactly j vertices. Then

$$Q_{i,j} \equiv \Pi_j(G_i). \tag{3}$$

In other words, $Q_{i,j}$ is the number of connected components of size j in G_i . Note that $G_i = G$ for $i \ge D$. If M(G) denotes the number of vertices in the largest component of G, then Q is a matrix with row indices $\left[0,1,\ldots,D\right]$ and column indices $\left[1,2,\ldots,M(G)\right]$. Although $Q_{i,j}$ is defined for any $i \ge 0$ and $j \ge 1$, it is zero beyond these values, so we typically truncate Q to be of size $(D+1)\times M(G)$. If the degree distribution is sparse then there will be repeated degree-limited subgraphs, as $(d_i = 0) \Rightarrow (G_i = G_{i-1})$ for $i \ge 1$. In such cases, the Q-matrix will therefore have duplicate rows.

Consider for example the graph G in Fig. 1, which has 8 vertices and 10 edges. It has a degree distribution of $\vec{d} = \langle 0, 4, 0, 2, 1, 0, 1 \rangle$ and gives rise to four distinct degree-limited subgraphs, G_1, G_3, G_4 , and G_6 . Since the maximum degree of G is 6, and the largest component size is 8, the corresponding Q-matrix of G is given by the 7×8 matrix

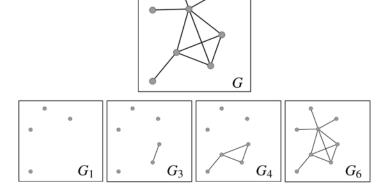


Fig. 1. A small graph G and its 4 distinct degree-limited subgraphs.

The *i*th row gives the component size distribution for G_i . At i=4, for example, we see that there are three components of size 1 (i.e., isolated vertices) and one component of size 4 in G_4 . Thus, $Q_{4,1}=3$ and $Q_{4,4}=1$. (The first element in the upper left-hand corner of Q is $Q_{0,1}$, rather than $Q_{1,1}$.) Furthermore, $G_D=G$, so G_6 contains the original graph, consisting of a single connected component of size 8, thus $Q_{6,8}=1$. We note that Q is sparse and contains redundant rows: $G_2=G_3$ and $G_5=G_4$. For practical considerations, we define a compact representation, the Q^* -matrix

$$Q_{i,j}^* \equiv \begin{cases} \Pi_j(G_i) & \text{if } d_i > 0\\ 0 & \text{otherwise} \end{cases}$$
 (5)

which zeros out these redundant rows of Q:

Here, only the non-zero values are explicitly shown. The matrices Q and Q^* convey the same information. Given one, the other can be easily derived. In practice, Q^* provides an economical storage format which more clearly conveys the information content of Q.

The graph characteristics captured by the Q-matrix may not be fully apparent for this small example. It is simple enough to explain the basic ideas, but too coarse to reveal structural patterns. In the next section we examine large real networks in which the usefulness of this representation becomes apparent.

For **directed graphs**, the Q-matrix can be interpreted as the number of *weakly connected components*. This essentially ignores the direction of edges and allows the same algorithms and analysis to be applied to both directed and undirected graphs. Other extensions to the Q-matrix are described in later sections.

The process of removing or adding specific vertices to a graph, as is done here, is a particular type of site percolation process and arises in several areas of network science, such as modeling the failure of routers in computer networks (information technology) or the spread of infectious diseases in populations (epidemiology). Various mathematical models have been developed to analyze the resilience to targeted attacks [7]. In the Q-matrix formulation, the site percolation process is rather specific (by ordering the removal of nodes by their degree) and occurs in discrete "bulk" steps (i.e., at each percolation step all nodes of a given degree are processed simultaneously). This last stipulation differs from conventional approaches in percolation studies, but this slight twist ensures that the process yields consistent results that do not exhibit statistical fluctuations and reduces the overall size of the Q-matrix.

3. Q-matrix Visualization

For large networks, it is impractical to display the Q-matrix explicitly, as in Eq. (4) or even Eq. (6). Instead, we project the matrix onto the x-y plane and plot the nonzero values on the z-axis, creating a three dimensional scatter plot of component size distributions. In this way the degree, component size, and number of components comprise the x, y, and z-axis, respectively. Furthermore, because the values on these axes span several orders of magnitude, it is convenient to render the plot on a log-log-log scale and use the Q* formulation to provide images which are less cluttered. We refer to this representation as the Q-matrix plot to distinguish it from the array representation in Eq. (6). In the sequel we use the term Q-matrix to refer both to the matrix and its plot; the context should make it clear which we mean.

For example, the Q-matrix in Fig. 2 is that of an undirected email communication network [8] with 36,692 vertices and 183,831 edges, where each vertex is an individual email address and two vertices are connected by an edge if there was at least one message sent from one to the other. The original graph is too large to render in its entirety, but its Q-matrix values consist of individual points (non-zeros) which can be effectively plotted. The top-left (0,1) corner of the Q-matrix is now on the floor in the rear corner, with the degree values running along the left rear wall, and component sizes running along the right rear wall.

The comb-like "lines" appearing in the plot are constant component size contours, for component sizes of k = 1, 2, 3, and so on. They are discrete points, but they are so densely represented as to appear as continuous lines when viewed at these scales.

Moving from the left wall (x-z plane) along the positive y-axis (towards us) the resulting image appears to resemble a hill, with a hook-like trail appendage closest to us, moving towards the lower right of the matrix, where the degree and component size are greatest. Upon closer inspection we can identify three loosely-defined regions in this type of image: the **wall** occurs near the x-z plane and shows how the small component sizes vary for each G_i ; the **hill** middle region shows how small and medium component sizes vary, and the characteristic **hook** on the floor (x-y plane) represents the birth and growth of the largest component. These are not precise mathematical boundaries, but these characteristics do seem prevalent in O-matrix plots, so the nomenclature is useful in describing these renderings.

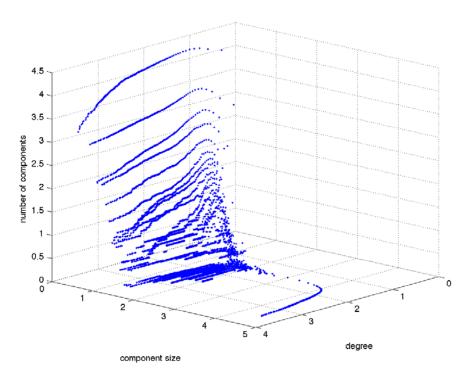


Fig. 2. The Q-matrix for the email-Enron network graph (Table 1) with 36,692 nodes and 183,831 edges.

4. Application Examples

Figure 2 shows the Q-matrix plots of real-work networks found in the Stanford Large Network Collection [9]. Their detailed descriptions are found in Table 1. In some cases these are directed graphs, and as previously noted, the Q-matrix then refers to the distribution of weakly connected components.

First and foremost, the experimental data shows that Q-matrix images of graphs from distinct application areas do, in fact, appear different. In each subfigure of Fig. 2 the wall, the hill and hook all have different shapes and aspect ratios.

Surprisingly, Q-matrices of graphs from the *same* application area appear to have *similar* characteristics, as demonstrated in Figs. 3-8. In such cases, each group shares similar shape and slope of the wall, hill, and hook regions for every network studied in our experiments. This suggests that the Q-matrix may be used as a crude classification tool to help identify "families" of large network graphs. Indeed, it is a canonical visual representation of the original graph, and unlike matrix structure plots, or graph drawing algorithms, there is *only one representation* for each graph, invariant under graph isomorphisms. This makes it useful for labeling large graphs with a compact image, and using this visual representation to categorize graphs into distinct groups. In particular, it is useful for tagging network graphs in databases with thumbnail images that actually yield distinguishable characteristics. In other words, Q-matrix plots provide a compact data set and a thumbnail image that may serve as a network "identification badge", or a "photo ID," capturing important characteristics beyond its size and degree distribution.

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¹ Current graph-drawing techniques have difficulty rendering meaningful visualizations for large graphs with heavy-tail degree distributions.

NETWORK NODES EDGES REFERENCE Collaboration Astro Physics 396,160 18,772 ca-AstroPh networks Condensed Matter 23,133 186,936 ca-CondMat 12,008 237,010 ca-HepPh (Fig. 4) High Energy Physics HE Physics Theory 9,877 51,971 ca-HepTh Web graphs Google 875,713 5,105,039 web-Google Notre Dame 325,729 1,497,134 web-NotreDame (Fig. 5) Stanford 281,903 2,312,497 web-Stanford Berkeley-Stanford 7,600,595 web-BerkStan 685,230 Road networks California 1,965,206 5,533,214 roadNet-CA (Fig. 6) Pennsylvania 1,088,092 3,083,796 roadNet-PA Texas 1,379,917 3,843,320 roadNet-TX Citation High Energy Physics 34,546 421,578 cit-HepPh networks HE Theoretical Physics 27,770 352,807 cit-HepTh cit-Patents (Fig. 7) **US** Patents 3,774,768 16,518,948 amazon0302 Co-purchasing March 2 262,111 1,234,877 March 12 3,200,440 amazon0312 networks 400,727 May 5 3,356,824 amazon0505 (Fig. 8) 410,236 403,394 June 1 3,387,388 amazon0601 **Email networks** Enron 36,692 183, 831 email-Enron European University 420,045 (Fig. 9) 265,214 email-EuAll Online social 5,879 508,837 **Epinons** soc-Epinions (Fig. 10) LiveJournal 4,847,571 68,993,773 soc-LiveJournal Slashdot (11-2008) 77,360 905,468 soc-Slashdot0811

Table 1. Example datasets from the Stanford Large Network Collection used in Q-martrix experiments

5. Extracting Conventional Measures

Slashdot (02-2009)

Embedded within the Q-matrix are basic networks measures, which can be inspected visually, or can be computed exactly with simple matrix/vector operations. For example, the nonzeros in the bottom row ($Q_{D+1,*}$) enumerate the connected components of each size in the original network; the first element, $Q_{0,1}$, tells us how many isolated vertices, if any, are present in G; the height and extension of the left wall capture leaf and low-degree vertex behavior as one increases the degree i for G_i .

82,168

Using $|x|_1$ to denote the vector 1-norm and [0:N] to denote the vector of N+1 nonnegative integers <0,1,2,...,N>, we can derive the following quantities:

• number of components in G_i is the row sum of $Q_{i,*}$

$$\Pi(G_i) = \sum_{j} Q_{i,j}$$

$$= |Q_{(i,*)}|_1$$
(7)

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In particular, $G_D = G$, so $\Pi(G) = |Q_{(D,*)}|_1$.

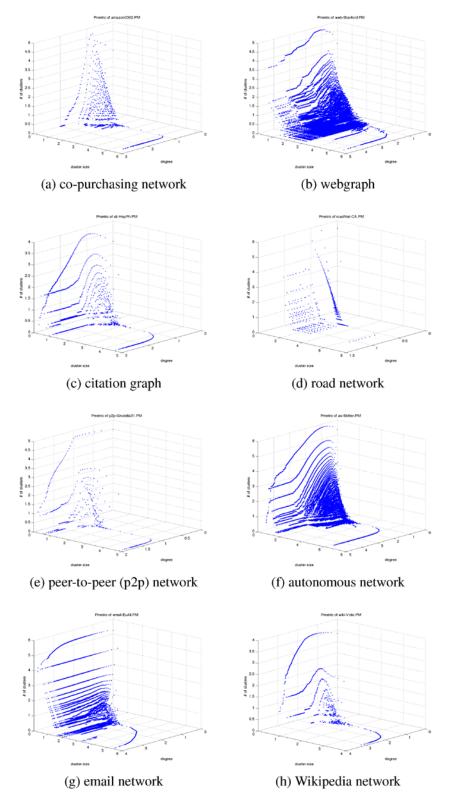


Fig. 3. Q-matrices of networks graphs from different application areas.

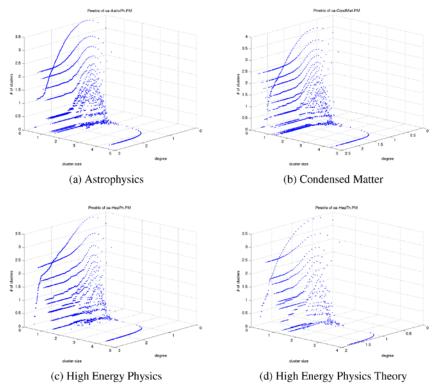


Fig. 4. Q-matrices of co-authorship networks.

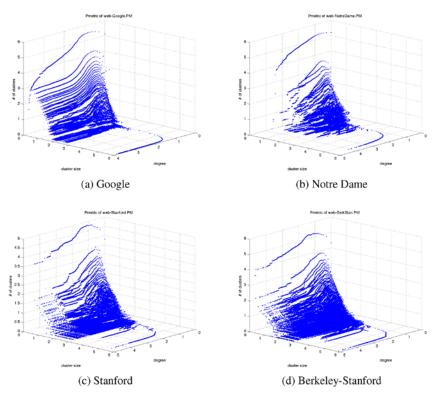


Fig. 5. Q-matrices of Web graphs.

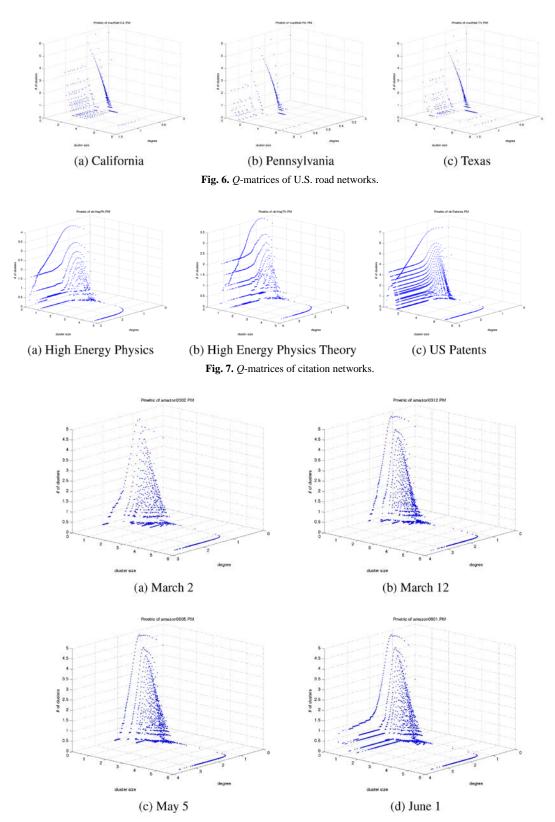


Fig. 8. Q-matrices of Amazon co-purchasing networks (2003).

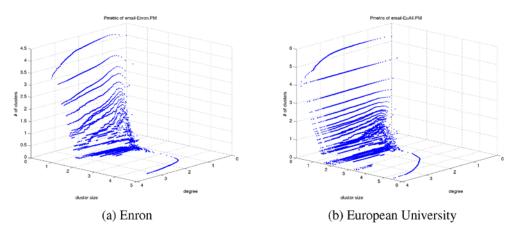


Fig. 9. Q-matrices of email networks.

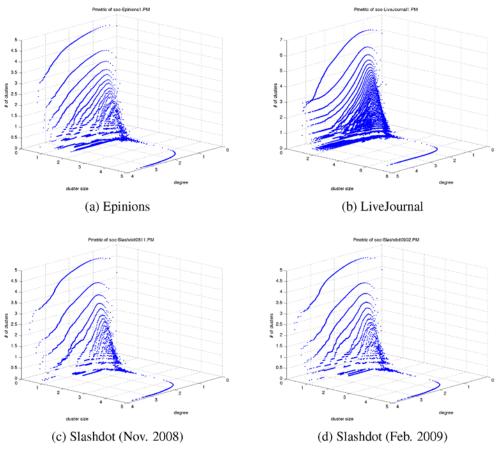


Fig. 10. Q-matrices of online social networks.

• size of largest component in G_i , denoted $M(G_i)$, is the index of the last nonzero in the i-th row:

$$M(G_i) = \max_{j} \{ j | Q_{i,j} > 0 \}.$$
 (8)

• number of vertices in G_i is the number of vertices with degree less than or equal to i, which is the number of components in the i-th row of Q multiplied by their respective sizes:

$$|V_{i}| = \sum_{j=0}^{M(G)} (Q_{(i,j)} \times j)$$

$$= Q_{(i,*)} \cdot [0:M(G)].$$
(9)

In particular, $|V|=|V_D|=Q_{(D,*)}\cdot [0:M(G)]$.

• **degree distribution of G**, $\vec{d} = \langle d_i \rangle$: The number of vertices in G with exactly degree i can be seen as the difference between the number of those with degree i or less and those with degree i-1 or less:

$$d_{i} = |V_{i}| - |V_{i-1}|$$

$$= \left[Q_{(i,*)} - Q_{(i-1,*)}\right] \cdot \left[0 : M(G)\right].$$
(10)

For i=0, we just have $d_0=|V_0|=Q_{(0,*)}\cdot \left[0:M(G)\right]$ as the number of isolated nodes in the original graph.

• number of edges in G for an undirected graph is the sum of the degrees of each vertex divided by two:

$$|E| = \frac{1}{2} \sum_{i} i \times d_{i}$$

$$= \frac{1}{2} [0:D] \cdot \vec{d}$$
(11)

where $\vec{d} = \{d_0, d_1, ..., d_D\}$ is given by Eq. (10).

6. Practical Considerations

The Q-matrix plot works best for large, complex networks with non-trivial degree distributions, where the Q-matrix contains sufficient non-zeros to yield a visually interesting image. For small graphs, like our toy example (Fig. 1) it is difficult to identify the wall, the hill and the hook. In fact, the Q-matrix plot works best precisely where other approaches, such as conventional graph drawing layouts fail, thus creating a useful complement to conventional methods for annotating network graphs.

In practice, the Q-matrix is sparse for large network graphs: there are relatively few distinct degrees (nonzeros in \vec{d}) and it is unlikely to find a component of particular size j in G_i . Hence, although the dimensions of Q are (D+1) by M(G), the actual number of nonzeros is quite small. Figure 11 illustrates the ratio between the number of nonzeros in the Q-matrix (in its compact Q^* form) and the number of edges in the original graph from a sample of 67 applications, ranging in size from several hundred to several million. The results are plotted on a log-log scale, and we see that the number of nonzeros in Q^* is roughly one to three orders smaller than the number of edges in G, with larger networks favoring higher

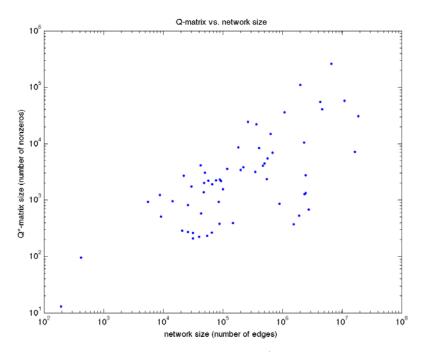


Fig. 11. In practice, the size of the Q-matrix (number of nonzeros in compact Q^* form) is roughly one to three orders of magnitude smaller in size than the original graph (number of edges), with larger graphs favoring higher compression ratios.

compression ratios. Networks with millions of edges are often represented by Q-matrices with just a few thousand numbers.

Computing the Q-matrix is practical for large networks. One can employ an efficient algorithm which builds the Q-matrix incrementally, without calculating G_i explicitly in the intermediate steps. It begins by sorting and partitioning the vertices by their percolation order (degree) and growing equivalence classes corresponding to the intermediate subgraph components. Large network graphs with millions of edges can be processed in a few seconds on a desktop computer. A separate paper describes the implementation and optimization strategies for developing an efficient algorithm in C++[14].

Furthermore, because a small perturbation to the graph structure (e.g., edge swap) could have a cascading effect to the resulting Q-matrix, it provides a fast method for identifying **graph non-isomorphism** of two large networks with identical size and degree distributions. On the other hand, proving the converse remains challenging—different graphs may yield the same Q-matrix. For example, any k-regular graph (i.e., where every vertex has the same degree k) will yield a Q-matrix that has exactly one non-zero: $Q_{k,|V|} = 1$. As a specific example, consider two non-isomorphic 3-regular (cubic) graphs: the Desargue graph, and the Dodecahedral graph (Fig. 12). Both have 20 nodes and 30 edges each, with identical degree distributions [10]. Both yield identical Q-matrices. Thus, the Q-matrix for a graph is not an invertible representation. (This is expected, as computing graph isomorphism remains a computationally challenging problem. [11]) There are indeed specific counter-examples where comparing two Q-matrix plots may yield little insight. Nevertheless, the interesting idea here is that the Q-matrix works best precisely when there is diversity in the degree distribution, and when large hubs are present: two key characteristics that separate real-world networks from structured and uniformly random graphs.

We can also use the Q-matrix to investigate how various graph properties compare to those of random graphs with similar degree sequences. For example, how does the growth of the giant component compare between real-world graphs and randomized versions with the same degree distribution? While formulations exist for calculating expected size at a given degree point [12], it is insightful to see how the variations

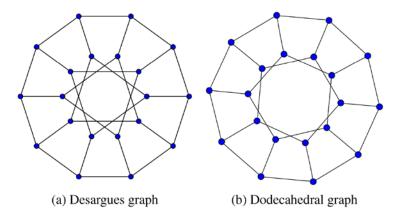


Fig. 12. Two non-isomorphic cubic graphs of same size (20 nodes and 30 edges) and degree distributions, which yield the same Q-matrix, i.e. $G \rightarrow Q(G)$ is not one-to-one. (Images courtesy of David Eppstein, CC0 2.0 license.)

behave over the complete degree spectrum. Figure 13, for example, shows the giant component growing much faster for the randomized graphs, in some regions by almost three orders of magnitude. In this experiment, we computed random variations of the original network by accumulative edge swaps that preserved the degree distribution. (That is, each edge in the graph was randomly swapped with another edge in such a way to preserve the original degree distribution.) We then computed the Q-matrix for these randomized versions, and compared the largest component size growth. The results demonstrate that original and randomized graphs have a completely different signature, and that the Q-matrix can be used as a validation tool to help separate real-world graphs from their synthetic counterparts.

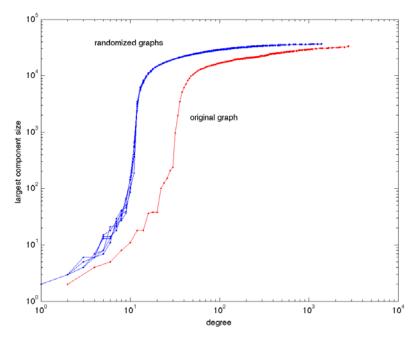


Fig. 13. The growth of the giant component for the email network (Fig. 2) shown in red, compared to random graphs of same size and degree distribution, all computed directly from their corresponding *Q*-matrices. The original graph behaves significantly different, and this technique can be used to identify real networks from their synthetic counterparts.

7. Comparing Graphs

Given two graphs, A and B, and their respective Q-matrices, Q(A) and Q(B), we may define a distance function $\Delta(A,B)$ between these two graphs as the Q-metric:

$$\Delta(A,B) \equiv ||Q(A) - Q(B)||$$

$$\Delta(A,B) = \sum_{i} \sum_{j} |Q(A)_{i,j} - Q(B)_{i,j}|.$$
(12)

In cases where the matrices Q(A) and Q(B) are of different sizes, they are padded with zeros so they are conformant. This formulation is essentially the vector 1-norm, interpreting the elements Q(A) and Q(B) as a long vector. This definition is chosen over the more common Frobenius matrix norm to keep all computation in integer arithmetic, and hence its numerical value exact.

Note that Δ does satisfy the requirement for a **pseudometric space**. Namely, for any graph A, B, C

$$\Delta(A, A) = 0 \tag{13}$$

$$\Delta(A,B) = \Delta(B,A) \tag{14}$$

$$\Delta(A,C) \le \Delta(A,B) + \Delta(B,C). \tag{15}$$

(Since $\Delta(A, B) = 0$ does not imply that A = B, the requirements for conventional metric space are not met.) We can use this distance function as a way to measure how different two graphs are in respective Q-matrix formulation. For example, Fig. 14 shows this metric applied to the email communication network (Fig. 2) and 100 random graphs generated as before with identical degree distribution. Here, a distribution of the $\left(\frac{101}{2}\right) = 5,050$ pairwise Q-metrics are plotted on a logarithmic x-axis. The result is a

bimodal distribution illustrating the difference between random graphs (left mode) and the original graph. That is, the pairwise Δ for each random graph is over 40 times smaller than the Δ between the original graph and its random counterparts. If we normalize this difference by the number of vertices in the graph, the mean difference between random matrices is 0.7964 with a standard deviation of 0.2494, while the mean difference between the original graph and all 100 random graphs is 34.9570, with a standard deviation of 0.2941. Once again, the original graph behaves significantly different than its random counterparts and such a test can help identify real networks from those generated synthetically.

8. Generalizations and Extensions

The Q-matrix has been defined for directed and undirected graphs, but further refinements could be made for the **directed graph** case by distinguishing between weakly-connected and **strongly-connected components**. One possible generalization of the Q-matrix formulation would be to define a version that creates two Q-matrices for directed graphs: one each for in-degree and out-degree distributions, and measure strongly-connected components for each.

Likewise, the Q-matrix formulation could also be extended to **weighted graphs**, where each edge has a weight, ω_e for $e = \{1, 2, ..., |E|\}$, by extending the notion of degree of a vertex to the sum of its edge weights.

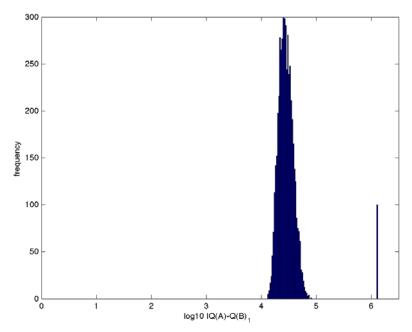


Fig. 14. Comparison between the email network (Fig. 2) and 100 random graphs with identical degree distribution. Here, a distribution of the 5,050 pairwise *Q*-metrics are plotted on a logarithmic *x*-axis, showing that the original graph (right mode) is quite different than its random counterparts (left mode).

Also, we may create alternate versions of the Q-matrix using other node orderings (centralities) in place of degree, e.g. between-ness, eigenvalue, or Pagerank centralities. A similar framework can be developed for **edge centralities**, in which edges, rather than vertices are removed (sometimes referred to as **bond percolation**).

Finally, we note that the Q-matrix of G can *itself* be interpreted as a weighted graph, written in adjacency format. That is, Q(G) is itself a graph. In this case one could apply this formulation twice, Q(Q(G)), to create a Q^2 -matrix, or any number of times to create the Q^n -matrix. Such an approach would produce a family of graph reductions that could collapse a large network graph into a single number. We are just beginning to investigate the implications of these extended interpretations.

9. Conclusion

The Q-matrix is a condensed representation of a network graph, which provides a meaningful visualization and encodes several measures of the graph's underlying topological structure. It is small, relatively easy to compute, and provides a convenient identification of the original network graph. (The Q^* formulation is used in practice, but both are mathematically equivalent.)

We have illustrated Q-matrix identities for computing the degree distribution, giant component growth, and basic parameters of undirected graphs (Eqs. (7)-(11)).

Computing the Q-matrix is computationally efficient [14]. Optimized algorithms allow networks with millions of edges to be processed in a few seconds on a laptop. The transformation $G \to Q(G)$ is a form of lossy compression. Furthermore, in its compact form, Q^* , the resulting structure is made even smaller. In practice, its size is roughly one to three orders of magnitude smaller than the original graph, with larger networks favoring higher compression ratios. For example, the LiveJournal network [9] has nearly 69

million edges, yet its compact Q^* representation requires less than 67 thousand values – a reduction ratio of about 1,000:1.

Experimental data indicates that the visualization provided by the Q-matrix distinguishes between graphs from different applications areas (Fig. 3) and that graphs from the same application area share visual similarities (Figs. 4-9). This includes examples from citation graphs, web graphs, road networks, peer-to-peer networks, autonomous networks, email networks, and Wikipedia networks, ranging from sizes of just a few thousand to nearly 70 million edges [13, 9]. While these experiments are not exhaustive of all network data available, they do suggest that the approach appears promising in practice.

The Q-matrix approach can also reveal differences between an organic (real-world) graph and randomized variations from its corresponding configuration model (ensemble of random graphs with identical degree distribution). We have shown example cases where the giant component grows much slower, by as much as three orders of magnitude, and such a difference can be computed exactly from their corresponding Q-matrices.

Furthermore, the difference between Q-matrices of different graphs may be quantified by the induced Q-metric $\Delta(A,B)$, as given by Eq. (13). This defines an exact, reproducible measure for network graphs which can also be useful in identifying application graphs from their randomized counterparts (Fig. 14).

Finally, we have outlined extensions to this approach for directed and weighted graphs, as well as generalized percolation orderings, like eigenvalue or between centrality. We have also proposed a recursive *O* -matrix formulation approach that can reduce a large network graph to a single number.

The understanding of large-scale networks remains a challenging problem, and hopefully such approaches may shed light on our comprehension of these systems. There is still much work to be done, and we hope that these formulations can help further that understanding.

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About the author: Roldan Pozo is a computer scientist in the Applied and Computational Mathematics Division of the Information Technology Laboratory at NIST. He conducts research in complex networks, high performance computing, and software tools for scientific and numerical problems. The National Institute of Standards and Technology is an agency of the U.S. Department of Commerce.