

FUNCTIONAL COLLECTIVITY OF COMPLEX BRAIN NETWORKS

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Contents

1. Introduction. The brain is a network.
 2. Extracting brain functional networks from the data
 3. Brain correlation networks are scale free.
 4. The origin of the brain networks scale-free property.
 - 4.1. Average Statistical Properties
 - 4.2. Degree Distribution
 - 4.3. Brain Connectivity versus Brain Collectivity
 5. Implications
 - 5.1. Scale Free Networks Imply Critical Dynamics?
 - 5.2. An Evolutionary Perspective. Why Do We Need A Brain?
 6. Conclusion
- Acknowledgements
Glossary
Bibliography
Biographical Sketches

Summary

Brain function is carried out by an astronomically large network of interconnected

neurons. Thanks to the impressive progress in brain imaging techniques an increasing amount of spatiotemporal brain data is now available. The analysis of this data using statistical physics approaches allows for a reduction of the brain spatiotemporal dynamics into information flow on a complex network. Using this view, recent studies revealed that these brain networks share several principles with natural, social and technological systems. These notes summarize the most relevant advances in our understanding of the brain using this approach.

1. Introduction - The Brain is A Network

Brain function is carried out by a network of approximately 10^{10} interconnected neurons. An increasing amount of spatiotemporal brain data is now available thanks to the impressive progress in brain imaging techniques in particular with functional Magnetic Resonance Imaging (fMRI). To analyze such a large and complex body of information, conceptual approaches grounded in statistical physics have been used recently. This work showed that the dynamics of these patterns can be reduced to complex networks that share several principles with natural, social and technological systems. The aim of these notes is to describe the most relevant advances in our understanding of the brain using this approach, focusing only on those results providing clues about the underlying critical dynamics of the brain. Other excellent surveys providing a broader perspective, including medical applications can be found in Sporns et al (2004), Bassett and Bullmore (2006), Reijneveld et al (2007), Bullmore and Sporns, (2009).

The chapter is organized as follows. After introducing the problem, section 2 describes how brain networks are derived. Section 3 describes the initial brain imaging experiments showing a broad distribution of functional connectivity, implying that brain networks are scale-free. Section 4 discusses how these scale-free properties are conceptually linked to critical dynamics in physical systems. This includes a detailed comparison between the brain results with those extracted from a paradigmatic critical system; the Ising model. Section 5 is dedicated to discuss the implications and the final section summarizes the conclusions.

2. Extracting Brain Functional Networks from the Data

Brain activity evolves continuously over a network of gargantuan size and complexity, therefore and it is crucial before attempting any meaningful analysis to reduce the data to a manageable size. One possible focus is on the *interactions* between brain sites, and uses that information to construct the adjacency matrix defining a network, often called “correlation network”. This is something done previously in many other systems as for instance, to study the interactions between cell proteins (Jeong et al, 2001) or gene expression (Stuart et al, 2003). In this manner, the problem is reduced to the study of a graph composed by nodes (sites) and links (interactions). The cartoon in Figure 1 illustrates these steps after the brain data is collected, as described by Eguiluz and colleagues (Eguiluz et al, 2005). Each snapshot of brain activity in this type of experiments is a fMRI image composite from the so-called BOLD (blood oxygenated level dependent) signal, which is a directly related to the level of neuronal activity in any given brain site. Typically the spatial resolution is of about 36 slices (3 mm thick)

from top to bottom of the brain, divided equally into a 64 by 64 matrix (3.475 mm x 3.475 mm wide) resulting in approximately cubic regions called voxels. The activity is recorded at consecutive intervals (in this case every 2.5 sec.) to produce typically 400 samples. As was stated above, the interest is to use the interactions between brain sizes to define a network. For simplicity, this is done computing the correlation between sites within a time window. Denoting the activity in voxel x at time t as $V(x,t)$ then the linear correlation coefficient between all pair of voxels, x_i and x_j is computed as:

$$r(x_i, x_j) = \frac{\langle V(x_i, t) V(x_j, t) \rangle - \langle V(x_i, t) \rangle \langle V(x_j, t) \rangle}{\sigma(V(x_i)) \sigma(V(x_j))} \quad (1)$$

where $\sigma^2(V(x)) = \langle V(x, t)^2 \rangle - \langle V(x, t) \rangle^2$ and the quantities inside brackets represent temporal averages.

In principle, as it is discussed later, networks can be built by defining negative or positive correlations, although the initial results were obtained using positive correlations. Thus, pairs of voxels whose correlation r exceeds a threshold ρ are considered functionally linked. Of course, if one chooses a threshold too small, then the majority of the voxels will appear to be connected to one another. Likewise if the threshold is too high, then voxels will appear isolated. There is a wide range of threshold values for which networks remain clearly defined indicating that the main conclusions are robust with respect to the selection of parameters.

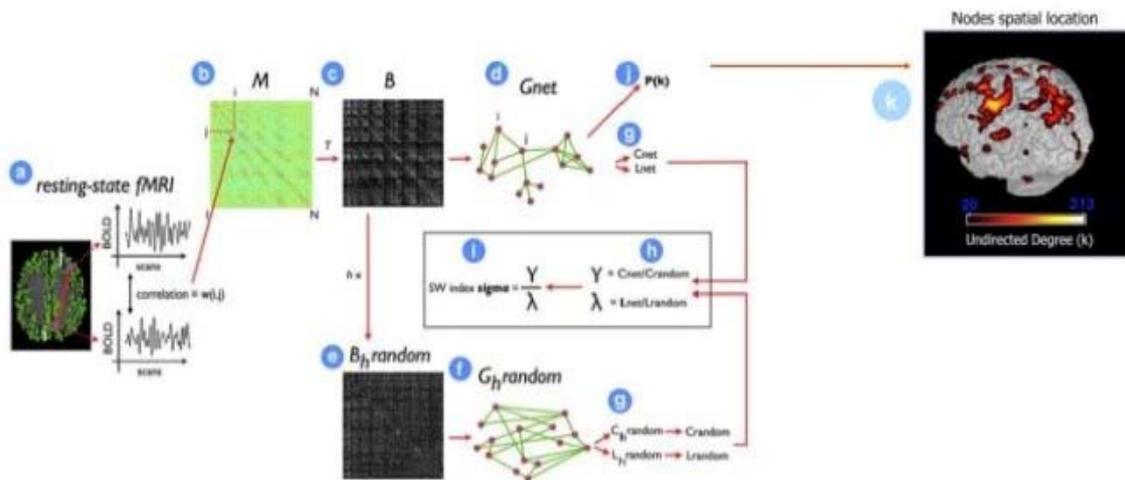


Figure 1. Steps to derive the networks from the BOLD fMRI signals. The correlation between fMRI signals (a) is computed to obtain a correlation matrix (b). The elements in the matrix passing a given threshold (c) define the network (d). Randomizing the nodes location, (e) a random network is obtained (f) and used to compare several statistics (g-j) with the original network. The node spatial location (k) can be also visualized.

3. Brain Correlation Networks Are Scale Free

An example of a brain network extracted using this method is shown in Figure 2. For clarity, only one eighth of the nodes are included in this illustration. The pattern of inhomogeneous connectivity is immediately evident by the fact that a relatively large proportion of nodes have a single link and a smaller proportion of nodes exhibit numerous connections. A quantitative estimate of this inverse relationship is shown in the bottom panel of Figure 2 demonstrating a skewed distribution of the number of links with a tail approaching a distribution $p(k) \sim k^{-\gamma}$ (with γ around 2). This power law is more evident for networks constructed with higher thresholds; i.e., less correlated conditions. For networks constructed with lower thresholds, a maximum appears which shifts to the right as ρ decreases.

The small inset in panel B of Figure 2 shows the distribution of links of a surrogate network constructed by randomly shuffling the original time sequence of each voxel's magnetic resonance signal. This is done to reject the possibility that the network arises from correlations non related to brain dynamics. The degree distribution of these surrogate networks does not fit a power law, but rather display a Gaussian distribution in which the mean and width depend on the ρ threshold used. These results reported by Eguiluz and colleagues demonstrate that brain dynamics, at this relatively large scale, evolves over a scale free network. These initial findings are now confirmed by other authors (Van der Heuvel et al, 2008) in a variety of experimental settings indicating that a few brain regions are very well connected (the right of the distribution), many more regions are only interacting with a few others, and a broad intermediate situation can be found to exist between these two extremes.

Other calculations revealed that the average number of links as a function of -physical-distance between brain sites also decays as a power law, implying that there is no characteristic length for the interactions between any two brain sites (Eguiluz et al 2005). Two other statistical properties of these networks, path length and clustering were computed as well (Boccaletti et al 2006). The path length (L) between two voxels is the minimum number of links necessary to connect both voxels. Clustering (C) is the fraction of connections between the topological neighbors of a voxel with respect to the maximum possible.

Measurements of L and C were also made in a randomized version of the brain network. The main finding was that L remained relatively constant in both cases while C in the random case resulted much smaller, implying that brain networks are "small world" nets, a property with several implications in terms of cortical connectivity. Further calculations revealed that the distribution of links amongst neighbors are positively correlated, in other words, highly connected nodes connect with highly connected ones and vice versa.

This was a surprising finding since this property, known as assortativity, is mostly found in social networks but not seen previously in biological systems, where the rule is that negative correlations regulate connectivity. Summarizing, the initial work of Eguiluz et al. showed that functional brain networks exhibit highly inhomogeneous scale-free functional connectivity with small world and assortativity properties.

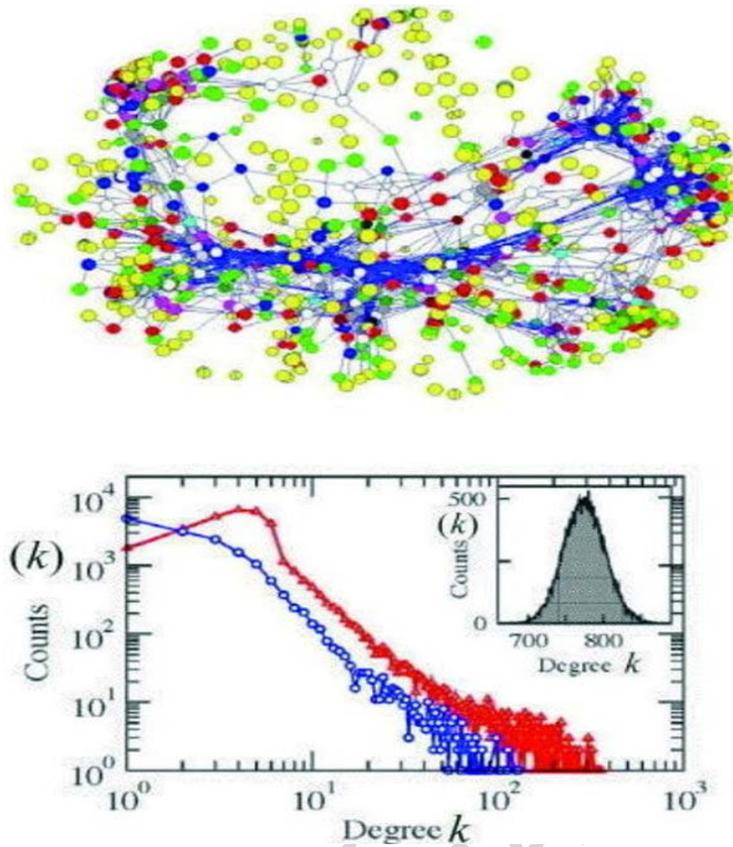


Figure 2. A typical brain network extracted from functional magnetic resonance imaging. Top panel shows a pictorial representation of the network where the nodes are colored according to its degree: yellow=1, green=2, red=3, blue=4, etc. The bottom panel shows the degree distributions for networks constructed with two correlation thresholds: $\rho = 0.7$ (blue circles) and $\rho = 0.6$ (red triangles). The inset depicts the degree distribution for an equivalent randomly connected network.

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Hector H. Berra was born in Rosario, Argentina in 1955. He received his medical degree in 1980 at the National University of Rosario, in Argentina and subsequently accredited as Board Certified Cardiologist by the Medical College of Santa Fe, Argentina in 1985. He received his Biomedical Sciences Ph.D.

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Pablo Balenzuela was born in Buenos Aires, Argentina in 1969. He got his M. Sc. degree in 1998 and his PhD degree in 2002 both in physics at the University of Buenos Aires, Argentina working on “Criticality and non-linearities in fragmentation”. After doing postdoctoral research (supported by the Fundacion Antorchas and the Generalitat of Catalunya) in the Universitat Politecnica de Catalunya, Spain between 2003 and 2006 he returned to Argentina where he was appointed Researcher of the National Council for Scientific and Technological Research (CONICET, Argentina) and Assistant Professor at the Physics Department of the University of Buenos Aires. His main interests are non-linear dynamical systems, criticality in out of equilibrium systems, and the modeling of biological and social systems from a complex system perspective.

Daniel Fraiman was born in Buenos Aires, Argentina in 1974. He received his M.Sc. degree in 2000 and his Ph. D. degree in 2006 both in physics from the University of Buenos Aires, Argentina. Between 2000 and 2002 he was member of the Clinical Research Committee of the National Hospital “Prof. Alejandro Posadas”. His work covers a wide range of topics, including the mathematical modeling of calcium signaling, single channel modeling, the study of complex networks, and brain dynamics. Some of his work is published in Cell Calcium, Physical Review E, and Biophysical Journal. Since 2008 he is Assistant Researcher of the National Council for Scientific and Technological Research (CONICET, Argentina) and Assistant Professor at the University of San Andres, Buenos Aires, Argentina. Prof. Fraiman is member of the US Biophysical Society and the Latin-American Society of Probability and Statistics.

Dante R. Chialvo was born in Rafaela, Argentina in 1956. He received his medical diploma in 1982 from the National University of Rosario, in Argentina where in 1985 he was appointed Professor of the Department of Physiology. From 1987 to 1992 he was Associate Professor in the State University of New York (Syracuse, NY) in the Department of Pharmacology and latter in the Computational Neuroscience Program. Between 1992 and 1995 he was associated with the Santa Fe Institute for the Sciences of Complexity, in Santa Fe, New México. Since 2000 he has been affiliated with Northwestern University (Chicago) as Research Associate Professor and Professor of Physiology, where currently is Adjunct Professor. Throughout these years, he has been Visiting Professor at numerous universities including Wuerzburg University (Germany), University of Copenhagen (Denmark), The Rockefeller University (U.S.A.), University of the Balearic Islands and University Complutense of Madrid, (Spain), University of Rosario and University of Cordoba (Argentina), Universidad Mayor de San Andres, La Paz, (Bolivia) among others. He has published more than 70 scientific papers, all dedicated to understand natural phenomena from the point of view of nonlinear dynamics of complex systems. Prof. Chialvo’s work covers a wide range of topics, including the mathematical modeling of cardiac arrhythmias, the study of molecular motors as stochastic ratchets, neural coding, and self-organization and collective phenomena in ant swarms, brain and communities, among others. In 2005 he was the recipient of a Fulbright US Scholar Award (2005), in 2006 the Distinguished Visiting Professor of the University Complutense, (Psychology Department), Madrid, Spain and elected Fellow of the American Physical Society in 2007.