



Critical brain networks[☆]

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Abstract

Highly correlated brain dynamics produces synchronized states with no behavioral value, while weakly correlated dynamics prevents information flow. We discuss the idea put forward by Per Bak that the working brain stays at an intermediate (critical) regime characterized by power-law correlations.

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1. Introduction

The human brain is a large system, with no more than a hundred specialized modules with different functions. At the smallest grain, the cerebral cortex consists of about 10^{10} neurons that comprise of a highly interconnected network. Each cell receives continuously a few thousands of excitatory inputs from other neurons. One of the simplest things we do not know about the brain is how the cortex, being a mainly excitatory network, prevents the expected explosive propagation of activity and still transmits information across areas. If the average number of neurons activated by one neuron is too high (i.e., supercritical) this results in the massive activation of the entire network, while if it is too low (i.e., subcritical), propagation dies out. The critical regime is the one in which these opposing processes are balanced. It was

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Table 1

Complicated systems are not complex

| | |
|---------------------|--|
| Complicated systems | Many linear pieces + a central supervisor + blueprint = “whole” (example: tv set) |
| Complex systems | Many nonlinear pieces + coupling + injected energy = “emergent properties” (example: society) |

Turing, about 50 years ago [1], the first to speculate in these terms, arguing that brains should be at a barely critical state. We review here the proposal [2] that most behaviorally relevant brain states, are associated with dynamics which is critical in this sense. This perspective places the emphasis on understanding the brain’s large number of dynamical nonlinear degrees of freedom, and in the dynamical attractors that are expected to emerge from the interaction of these elements. Less emphasis is given to some other aspects, including probable “computation” properties of the emergent circuits. It sees the brain as the solution of a very large dynamical system. It says nothing about anything else.

The paper is dedicated to discuss recent experimental findings of critical correlations in brain dynamics. The paper is organized as follows. The next section reminds ourselves where the brain problem is in the general context of dynamical systems. In Section 3 we discuss results from brain imaging experiments showing a broad distribution of functional connectivity, implying that brain networks are scale-free. This is contrasted with the known cortical connectivity. Section 4 contains recent experimental evidence at the *in vitro* level indicating that cultured cortical networks are critical at the neuronal level. The paper concludes with a comment on the ideas of our late friend Per, who foresaw many of these results.

2. Brains in “DynamicsLand”

Brain activity happens in bursts, in which pauses, silence or boredom suddenly and unpredictably are followed by brief activity. From a dynamical viewpoint, brain dynamics is not different from other natural processes. Nature is clearly nonhomogeneous and intermittent, the analysis of any natural object reveals an ever surprising amount of details; there is no single relevant scale at which Nature becomes homogeneous. Complexity is this lack of uniformity associated with the scale-free spatiotemporal feature. The driving force in this field continues to be the effort to understand what generates complexity, and how many different dynamical mechanisms can produce scale-free objects. It is now widely recognized that, under a variety of conditions, nonlinear systems with many degrees of freedom tend to evolve towards complexity and criticality [2,3]. It is the interaction of many nonlinear degrees of freedom which produces emergent dynamics we call complex. The latest “heavenly example” is the sun’s sudden bursts of radiation emanating from quick re-arrangements of the magnetic field network in the corona [4]. This is different from the dynamics arising in complicated systems comprised of the simple addition of interconnected *linear* pieces (Table 1). Examples

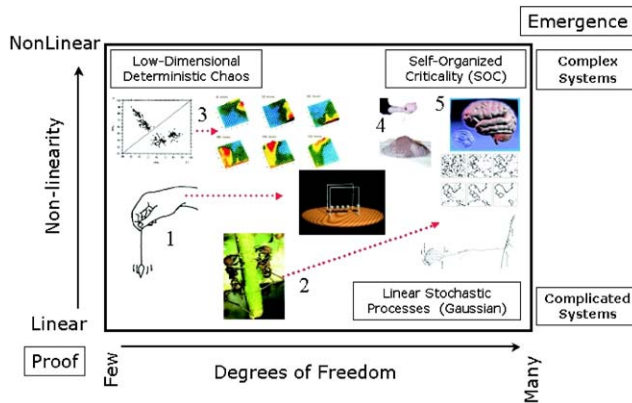


Fig. 1. “DynamicsLand”: A cartoonish representation of the parameter space for various classes of dynamical systems. The simplest ones “live” in the left-bottom corner, where analysis and formal proofs are the techniques expected, but many fundamental problems in biology correspond to areas distant from that land. Relatively simple dynamics gets sophisticated as the nonlinear term acquires relevance (moving upward in the graph) or as the number of degrees of freedom increases (moving to the right). Pictorial examples include: (1) the transition from one to many coupled pendulums, (2) few foraging ants to the entire swarm [5], (3) from the chaotic dynamics of an isolated cardiac cell [6] to the spatiotemporal spiral waves in the heart [7], (4) a sandpile and, of course, (5) the brain.

of complicated systems are a television set, or a car. They do not “emerge”, they are manufactured by following a blueprint given by the designer. Evolution to more complicated designs requires always of the supervisor (or designer) intervention. Complex systems such as species, ecologies, societies or brains do not arise from blueprints, they are very robust emergent consequences of dynamical laws we still do not understand. We can not replay the tape of evolution to investigate whether other ecologies will arise or self-organize, but in the case of brains we witness the high percentage of brains that end up well connected and, with few exceptions, working. This is a marvellous thing, considering that during one year of development a brain is continuously adding (or “connecting”) 10^5 new neurons per minute. What are the basis of the self-organizing mechanism able to achieve such feats? One motivation for biologists to look at the physical laws governing complex systems of all kinds is the hope that universality will give us an edge. The rationale is that a good understanding of these universal laws will provide a breakthrough and shed light on related biological problems. The considerations in the cartoon of Fig. 1 remind us that the brain is at a region of parameter space where complex dynamics can emerge. In that top-right corner, theory is scarce, but some insight and numerical tools can be borrowed from related work in self-organized criticality and complex networks, as discussed in the next section.

3. Functional networks are scale-free

Brain activity is eminently spatio-temporal, as such the monitoring of the complicated cortical patterns have greatly benefited from techniques developed in the context

of functional magnetic resonance imaging (fMRI). However, the numerical analysis of such spatiotemporal patterns is less developed, lacking mathematical tools and approaches specifically tailored to grasp the complexity of brain cortical activity. One possibility is to get insight from recent work showing that disparate systems can be described as complex networks, that is assemblies of nodes and links with nontrivial topological properties [8–10].

The brain creates and reshapes continuously complex functional networks of correlated dynamics responding to the traffic between regions, during behavior or even at rest. We have recently studied these networks, using functional magnetic resonance imaging in humans (see methods in Ref. [15]). The data is analyzed in the context of the current understanding of complex networks (for reviews see Refs. [8–13]). During any given task the networks are constructed in the following way. Magnetic resonance brain activity is measured, at each time step (typically 400 spaced 2.5 s), from $36 \times 64 \times 64$ brain sites (so-called “voxels” of dimension $3 \times 3.475 \times 3.475 \text{ mm}^3$). The activity of voxel x at time t is denoted as $V(x, t)$. To define the links, we denote as functionally connected those brain sites whose temporal evolution are correlated beyond a positive pre-established value r_c , following an approach used previously in Ref. [14]. (Networks can be built as well by defining negative or positive correlations.) Specifically, we calculate the linear correlation coefficient between any pair of voxels, x_1 and x_2 , as

$$r(x_1, x_2) = \frac{\langle V(x_1, t)V(x_2, t) \rangle - \langle V(x_1, t) \rangle \langle V(x_2, t) \rangle}{\sigma(V(x_1))\sigma(V(x_2))}, \quad (1)$$

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

In Fig. 2, we show a typical network extracted with this technique for one subject in a finger tapping task [15]. The top panel shows the network’s nodes (only a portion for illustration) colored according to its degree and the bottom panel the degree distribution of the network. Degree is the mathematical term for each voxel connectivity, being represented here as how many other voxels are temporally correlated with it. We find that the degree distribution has a skewed distribution with a tail approaching a power-law distribution with an exponent around 2. As the threshold r_c is decreased a maximum appears which shifts to the right as r_c is lowered. Other measures reveal that the number of links as a function of distance also decays as a power law. This is so, from one voxel, the smallest scale able to be measured with this technique, to the largest, the size of the brain. When we looked how the connectivity was arranged in the neighborhood of a node we found that highly connected nodes were connected, on the average, with highly connected ones. This feature, only seen before in social networks is inverse to what one expects from a hierarchical organization [16–18]. We looked at two other statistical properties of these networks, path length and clustering. 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. If voxel i has degree k_i , then the maximum number of links between the k_i neighbors is $k_i(k_i - 1)/2$. Thus, if E_i is the number of links connecting the neighbors then the clustering of voxel i , $C_i = 2E_i/k_i(k_i - 1)$. The average clustering of a network is given by $C = 1/N \sum_i C_i$,

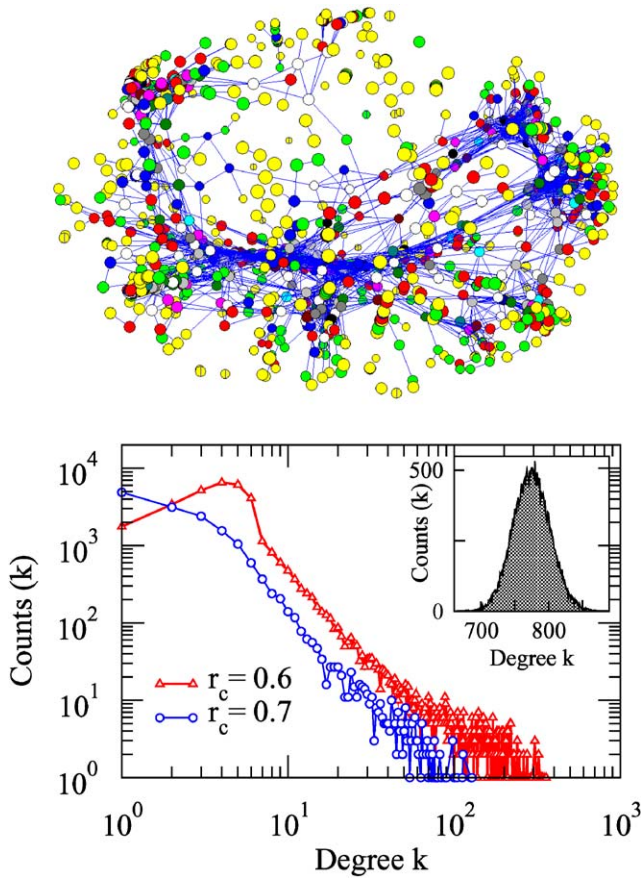


Fig. 2. A typical brain network extracted from functional magnetic resonance imaging. Top panel shows a pictorial representation of the network where the nodes are colored according to its degree: yellow = 1, green = 2, red = 3, blue = 4, etc. The bottom panel shows the degree distribution for two correlation thresholds. The inset depicts the degree distribution for an equivalent randomly connected network. Data re-plotted from Ref. [15].

Table 2
Statistical properties of human fMRI functional networks [15] and macaque cerebral cortex connectivity [21]

| Networks | N | C | L | $\langle k \rangle$ | λ | C_{rand} | L_{rand} |
|--------------|------|------|-----|---------------------|-----------|----------------------|------------|
| fMRI network | 4891 | 0.15 | 6.0 | 4.12 | 2.2 | 8.9×10^{-4} | 6.0 |
| Macaque C.C. | 71 | 0.46 | 2.3 | 10.6 | NA | 0.15 | 2.0 |

where N is the number of voxels. The results are presented in Table 2 (average values for $n = 22$ datasets, $r_c = 0.8$). From left to right are listed N , (number of nodes) C (clustering coefficient), L (shortest path length), the average degree $\langle k \rangle$, and λ the

exponent of the degree distribution. The clustering (C_{rand}) and path length (L_{rand}) values of an equivalent random network are also included for comparison. In all cases, the coefficient C remains four orders of magnitude larger than C_{rand} , the clustering of a random network. This feature, together with the similarity of path length of the original network and their randomized controls (L and L_{rand}), is indicative of a small-world structure.

In this approach we define two voxels as “linked” if they are temporally correlated beyond some value. This, of course, does not mean they are mutually connected (via chemical or neuro-transmitter or anatomical paths). The simplest counter-example is the case of a common input activating both sites. Further analysis will clarify if this is the case for the networks we are studying. In this regard it is relevant to look at work done over the last decade on the cortical connectivity (now “linked” is used in the strong sense of being mutually connected) by Sporns et al. [19–21], Hilgetag et al. [22], Young et al. [23] and Scannel et al. [24]. The analysis of connectivity matrix of these data sets reveals that although the networks are highly clustered and exhibit relatively short path lengths (see Table 2) as in small world networks, they exhibit a rather homogeneous degree distribution. In Fig. 3 the macaque cerebral cortex connectivity matrix [23] is plotted in the top panel. Because of the small statistics (only 71 nodes), its degree distribution is computed as the cumulative density, plotted in the bottom panel of the same figure. It can be seen that the degree distribution is not a power law, instead one sees that no area has less than four links and that the majority are linked with about ten other areas. The scale-free features illustrated in Fig. 2 reflects underlying long range correlations, i.e., brain activity on a given area can be correlated with far away and apparently unrelated regions, something already documented with other technology [25]. In qualitative terms this means that, for instance, a concurrent sound or simply imagery can influence thoughts or pain perception. Using the network approach here described, various dynamical brain behavioral states can be studied in the future.

4. Neuronal avalanches are critical

What are the neuronal mechanisms responsible for the correlations described in the previous section? From a top-down approach, Varela was among the first to be concerned with the brain large scale dynamical properties (reviewed in Ref. [26]). Varela assumed that “For every cognitive act, there is a singular and specific large cell assembly that underlies its emergence and operation” [27]. Efforts are underway to formulate in neuronal terms this working hypothesis, also termed “dynamic core” [28]. Other bottom-up complementary, approaches involve sophisticated recording in cortical structures with multielectrode arrays (see a review in Ref. [29]). A recent study [30] deserves special mention, because it provides quantitative estimations of the dynamical properties and characteristic exponents opening the possibility to model the neural mechanisms. Beggs and Plenz [30] studied the spontaneous neuronal activity of cultured and acute slices of rat cortex using 60-channel multielectrode arrays. They documented that during spontaneous activity the cortex typical activity shows

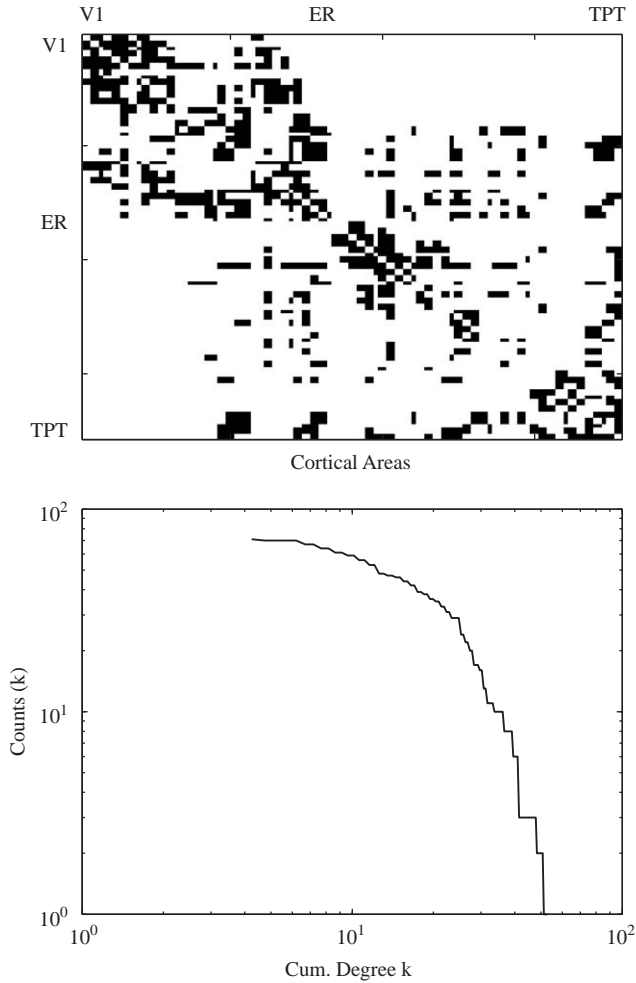


Fig. 3. Top panel: Connectivity matrix of the macaque cortex [23]. A black square denotes a connection between the 71 cortical areas: V1, V2, V3, VP, V3A, V4, VOT, V4T, MT, MSTD, MSTL, FST, PITD, PITV, CITD, CITV, AITD, AITV, STPP, STPA, TF, TH, PO, PIP, LIP, VIP, DP, A7A, FEF, A46, TGV, ER, HIPP, A3A, A3B, A1, A2, A5, R1, S2, A7B, IG, ID, A35, A4, A6, SMA, A3, A23, A24, A9, A32, A25, A10, A45, A12, A11, A13, G, PAAR, PAAL, PAAC, KA, PAL, PROA, REIT, TGD, TS1, TS2, TS3, TPT (top to bottom, left to right). For reference are labelled visual (V1), entorhinal (ER) and temporoparietal (TPT) cortices. Bottom panel: Cumulative degree distribution for the same data in the top panel. In contrast with the data plotted in Fig. 2, here there is no scale-free connectivity. Matrix data from Ref. [21] (available from website <http://www.indiana.edu/~cortex/connectivity>).

intermittent avalanches. After computing the statistics of several days worth of continuous cortical activity with a few millions of events, they showed that the avalanche size distributions, expressed as total number of electrodes activated per run, demonstrates the existence of a power law, with exponent $\alpha \sim 1.5$ as shown in Fig. 4. They

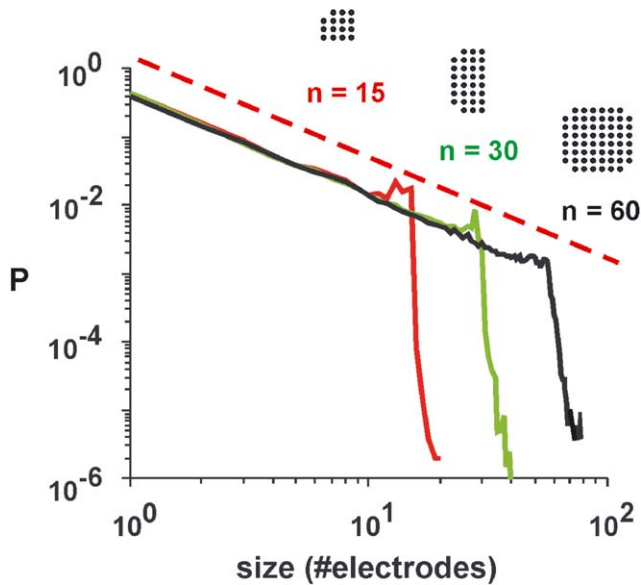


Fig. 4. The size distribution of neuronal avalanches in mature cortical cultured networks follows a power law with an exponent $\sim \frac{3}{2}$ (dashed line). The data, re-plotted from Fig. 4 of Ref. [30], shows the probability of observing an avalanche covering a given number of electrodes for three sets of grid sizes shown in the insets with $n = 15, 30$ or 60 sensing electrodes (equally spaced at $200 \mu\text{m}$). The statistics is taken from data collected from 7 cultures in recordings lasting a total of 70 h and accumulating $58000 (\pm 55000)$ avalanches per hour (mean \pm SD).

also computed an average branching activation ratio close to unity (calculated as the ratio between the current and future number of excited electrodes). The authors showed that this branching ratio optimizes information transmission in feedforward networks models. These aspects demonstrating criticality should be quantitatively accounted for by futures theories of cortical dynamics.

5. Per foresaw it

“Is biology too difficult for biologists? And what can physics, dealing with the simple and lawful, contribute to biology, which deals with the complex and diverse?”. In such challenging terms Per Bak encouraged colleagues to look at biology. Much before mainstream physics embraced biology, Per Bak was already convinced that “The big targets for physics theorists are biological evolution and the brain. These complex many-body problems might have similarities to problems studied in particle- and solid-state physics.” [31]. In an attempt to shake the stagnating state of these fields, by way of models and provocative metaphors, he suggested useful ways to re-think the most important issues in these areas. At the same time, he often insisted, in his unforgiving way, that mainstream ideas about neural models were an “example of physicists

leading a whole field astray”, because it was “hard to imagine a biological foundation for the complicated procedures for updating the synaptic strengths in those models” [31]. It was fun, in a way, to witness him, a *physicist*, reminding *biologists* and everybody else about the *biological* unplausibility of current brain models and the need to reconsider the constraints of self-organization as designing principle [32–35]. In his opinion, it was absolutely obvious that self-organization is the driving mechanism designing Nature around us, regardless if it was a human society, millions of neurons or an ant swarm. It is, in this view, mandatory to understand first the general laws before attempting to proceed with anything else. Per used to joke that, working out the further details will be just as easy (and boring) as the “cleaning after the party”. It is fascinating to see that a decade later, self-organization issues are newsworthy even to engineering [36].

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