Synchronization in complex networks

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(Dated: February 6, 2008)

Synchronization processes in populations of locally interacting elements are in the focus of intense research in physical, biological, chemical, technological and social systems. The many efforts devoted to understand synchronization phenomena in natural systems take now advantage of the recent theory of complex networks. In this review, we report the advances in the comprehension of synchronization phenomena when oscillating elements are constrained to interact in a complex network topology. We also overview the new emergent features coming out from the interplay between the structure and the function of the underlying pattern of connections. Extensive numerical work as well as analytical approaches to the problem are presented. Finally, we review several applications of synchronization in complex networks to different disciplines: biological systems and neuroscience, engineering and computer science, and economy and social sciences.

PACS numbers: 05.45.Xt, 89.75.Fb, 89.75.Hc

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I. INTRODUCTION

Synchronization, as an emerging phenomenon of a population of dynamically interacting units, has fascinated humans from ancestral times. No matter whether the phenomenon is spontaneous or induced, synchronization captivates our minds and becomes one of the most interesting scientific problems. Synchronization processes are ubiquitous in nature and play a very important role in many different contexts as biology, ecology, climatology, sociology, technology, or even in arts ((Osipov et al., 2007; Pikovsky et al., 2001). It is known that synchrony
is rooted in human life from the metabolic processes in our cells to the highest cognitive tasks we perform as a group of individuals. For example, the effect of synchrony has been described in experiments of people communicating, or working together with a background of shared, non-directive conversation, song or rhythm, or of groups of children interacting to an unconscious beat. In all cases the purpose of the common wave length or rhythm is to strengthen the group bond. The lack of such synchrony can index unconscious tension, when goals cannot be identified nor worked towards because the members are "out of sync" (Hall, 1983).

Among the efforts for the scientific description of synchronization phenomena, there are several capital works that represented a breakthrough in our understanding of these phenomena. In 1665, the mathematician and physicist, inventor of the pendulum clock, C. Huygens, discovered an odd "kind of sympathy" in two pendulum clocks suspended side by side of each other. The pendulum clocks swung with exactly the same frequency and 180 degrees out of phase; when the pendula were disturbed, the antiphase state was restored within half an hour and persisted indefinitely. Huygens deduced that the crucial interaction for this effect came from "imperceptible movements" of the common frame supporting the two clocks. From that time on, the phenomenon got into the focus of scientists. Synchronization involves, at least, two elements in interaction, and the behavior of a few interacting oscillators has been intensively studied in the physics and mathematics literature. However, the phenomenon of synchronization of large populations is a different challenge and requires different hypothesis to be solved. We will focus our attention on this last challenge.

In the obituary of Arthur T. Winfree, Strogatz (2003a) summarizes what can be considered the beginning of the modern quest to explain the synchronization of a population of interacting units: "Wiener (1948) posed a problem in his book Cybernetics: How is it that thousands of neurons or fireflies or crickets can suddenly fall into step with one another, all firing or flashing or chirping at the same time, without any leader or signal from the environment? Wiener did not make significant mathematical progress on it, nor did anyone else until Winfree came along". Winfree (1967) studied the nonlinear dynamics of a large population of weakly coupled limit-cycle oscillators with intrinsic frequencies that were distributed about some mean value, according to some prescribed probability distribution. The milestone here was to consider biological oscillators as phase oscillators, neglecting the amplitude. Working within the framework of a mean field model, Winfree discovered that such a population of non-identical oscillators can exhibit a remarkable cooperative phenomenon. When the variance of the frequencies distribution is large, the oscillators run incoherently, each one near its natural frequency. This behavior remains when reducing the variance until a certain threshold is crossed. However, below the threshold the oscillators begin to synchronize spontaneously (see (Winfree, 1980)). Note that the original Winfree model was not solved analytically until recently (Ariaratnam and Strogatz, 2001).

Although Winfree's approach proved to be successful in describing the emergence of spontaneous order in the system, it was based on the premise that every oscillator feels the same pattern of interactions. However, this all-to-all connectivity between elements of a large population is difficult to conceive in real world. When the number of elements is large enough, this pattern is incompatible with physical constraints as for example minimization of energy (or costs), and in general with the rare observation of long range interactions in systems formed by macroscopic elements. The particular local connectivity structure of the elements was missing (in fact, discarded) in these and subsequent approaches.

In 1998, Watts and Strogatz presented a simple model of network structure, originally intended precisely to introduce the connectivity substrate in the problem of synchronization of cricket chirps, which show a high degree of coordination over long distances as though the insects were "invisibly" connected. Remarkably, this work did not end in a new contribution to synchronization theory but as the seed for the modern theory of complex networks (Watts and Strogatz, 1998). Starting with a regular lattice, they showed that adding a small number of random links reduces the distance between nodes drastically, see Fig. 1. This feature, known as small-world (SW) effect, had been first reported in an experiment conducted by S. Milgram (Travers and Milgram, 1969) examining the average path length for social networks of people in the United States. Nowadays, the phenomenon has been detected in many other natural and artificial networks. The inherent complexity of the new model, from now on referred to as the Watts-Strogatz (WS) model, was in its mixed nature in between regular lattices and random graphs. Very soon, it turned out that the nature of many interaction patterns observed in scenarios as diverse as the Internet, the World-Wide Web, scientific collaboration networks, biological networks, was even more "complex" than the WS model. Most of them showed a heavy tailed distribution of connectivities with no characteristic scale. These networks have been since then called scale-free (SF) networks and the most connected nodes are called hubs. This novel structural complexity provoked an explosion of works, mainly from the physicists community, since a completely new set of measures, models, and techniques, was needed to deal with these topological structures.

During one decade we have witnessed the evolution of the field of complex networks, mainly from a static point of view, although some attempts to characterize the dynamical properties of complex networks have also been made. One of these dynamical implications, addressed since the very beginning of the subject, is the emergent phenomena of synchronization of a population of units with an oscillating behavior. The analysis of synchronization processes has benefited from the advance in the understanding of the topology of complex networks, but
it has also contributed to the understanding of general emergent properties of networked systems. The main goal of this review is precisely to revise the research undertaken so far in order to understand how synchronization phenomena are affected by the topological substrate of interactions, in particular when this substrate is a complex network.

The review is organized as follows. We first introduce the basic mathematical descriptors of complex networks that will be used henceforth. Next, we focus on the synchronization of populations of oscillators. Section IV is devoted to the analysis of the conditions for the stability of the fully synchronized state using the Master Stability Function (MSF) formalism. Applications in different fields of science are presented afterwards and some perspectives provided. Finally, the last section rounds off the review by giving our conclusions.

II. COMPLEX NETWORKS IN A NUTSHELF

There exist excellent reviews devoted to the structural characterization and evolution of complex networks (Albert and Barabási, 2002; Boccaletti et al., 2006; Costa et al., 2007; Dorogovtsev and Mendes, 2002; Newman, 2003b; Strogatz, 2001). Here we summarize the main features and standard measures used in complex networks. The goal is to provide the reader a brief overview of the subject as well as to introduce some notation that will be used throughout the review.

The mathematical abstraction of a complex network is a graph $G$ comprising a set of $N$ nodes (or vertices) connected by a set of $M$ links (or edges), being $k_i$ the degree (number of links) of node $i$. This graph is represented by the adjacency matrix $A$, with entries $a_{ij} = 1$ if a directed link from $j$ to $i$ exists, and 0 otherwise. In the more general case of a weighted network, the graph is characterized by a matrix $W$, with entries $w_{ij}$, representing the strength (or weight) of the link from $j$ to $i$. The investigation of the statistical properties of many natural and man-made complex networks revealed that, although representing very different systems, some categorization of them is possible. The most representative of these properties refers to the degree distribution $P(k)$, that indicates the probability of a node to have a degree $k$. This fingerprint of complex networks has been taken for a long time as its most differentiating factor. However, several other measures help to precise the categorization. Examples are the average shortest path length $\ell = \langle d_{ij} \rangle$, where $d_{ij}$ is the length of the shortest path between node $i$ and node $j$, and the clustering coefficient $C$ that accounts for the fraction of actual triangles (three vertices forming a loop) over possible triangles in the graph.

The first classification of complex networks is related to the degree distribution $P(k)$. The differentiation between homogeneous and heterogeneous networks in degree is in general associated to the tail of the distribution. If it decays exponentially fast with the degree we refer to as homogeneous networks, the most representative example being the Erdős-Rényi (ER) random graph (Erdős and Rényi, 1959). On the contrary, when the tail is heavy one can say that the network is heterogeneous. In particular, SF networks are the class of networks whose distribution is a power-law, $P(k) \sim k^{-\gamma}$, the Barabási-Albert (BA) model (Barabási and Albert, 1999) being the paradigmatic model of this type of graph. This network is grown by a mechanism in which all incoming nodes are linked preferentially to the existing nodes. Note that the limiting case of lattices, or regular networks, corresponds to a situation where all nodes have the same degree.

This categorization can be enriched by the behavior of $\ell$. For a lattice of dimension $d$ containing $N$ vertices, obviously, $\ell \sim N^{1/d}$. For a random network, a rough estimate for $\ell$ is also possible. If the average number of nearest neighbors of a vertex is $k$, then about $k^2$ vertices of the network are at a distance $\ell$ from the vertex or closer. Hence, $N \sim k^\ell$ and then $\ell \sim \ln(N) / \ln(k)$, i.e., the average shortest-path length value is small even for very large networks. This smallness is usually referred to as the SW property. Associated to distances, there exist many measures that provide information about "centrality" of nodes. For instance, one can say that a node is central in terms of the relative distance to the rest of the network. One of the most frequently used centrality measures in the physics literature is the betweenness (load in some papers), that accounts for the number of shortest paths between any pair of nodes in the network that go through a given node or link.

The clustering coefficient $C$ is also a discriminating property between different types of networks. It is usually calculated as follows:

$$C = \frac{1}{N} \sum_{i=1}^{N} C_i = \frac{1}{N} \sum_{i=1}^{N} \frac{n_i}{k_i(k_i-1)/2},$$

where $n_i$ is the number of connections between nearest neighbors of node $i$, and $k_i$ is its degree. A large clustering coefficient implies many transitive connections and consequently redundant paths in the network, while a low $C$ implies the opposite.

Finally, it is worth mentioning that many networks have a community structure, meaning that nodes are
linked together in densely connected groups between which connections are sparser. Finding the best partition of a network into communities is a very hard problem. The most successful solutions, in terms of accuracy and computational cost (Danon et al., 2005), are those based on the optimization of a magnitude called modularity, proposed by Newman and Girvan (2004), that precisely allows for the comparison of different partitionings of the network. The modularity of a given partition is, up to a multiplicative constant, the number of links falling within groups minus its expected number in an equivalent network with links placed at random. Given a network partitioned into communities, the mathematical definition of modularity is expressed in terms of the adjacency matrix $a_{ij}$ and the total number of links $M = \frac{1}{2} \sum_i k_i$ as

$$Q = \frac{1}{2M} \sum_{ij} (a_{ij} - \frac{k_i k_j}{2M}) \delta_{c_i, c_j}$$

where $c_i$ is the community to which node $i$ is assigned and the Kronecker delta function $\delta_{c_i, c_j}$ takes the value 1 if nodes $i$ and $j$ are in the same community, and 0 otherwise. The larger the $Q$ the more modular the network is. This property promises to be specially adequate to unveil structure-function relationships in complex networks (Girvan and Newman, 2002; Guimerà et al., 2003, 2005; Newman, 2006).

### III. COUPLED PHASE OSCILLATOR MODELS ON COMPLEX NETWORKS

The need to understand synchronization, mainly in the context of biological neural networks, promoted the first studies of synchronization of coupled oscillators considering a network of interactions between them. In the late 90’s, Strogatz and Mirollo (1988) and later Niebur et al. (1991) studied the collective synchronization of phase non-linear oscillators with random intrinsic frequencies under a variety of coupling schemes in 2D lattices. Beyond the differences with the actual conception of a complex network, the topologies studied in Niebur et al. (1991) can be thought of as a first approach to reveal how the complexity of the connectivity affects synchronization. The authors used a square lattice as a geometrical reference to construct three different connectivity schemes: four nearest neighbors, Gaussian connectivity truncated at $2\sigma$, and finally a random sparse connectivity. These results showed that random long-range connections lead to a more rapid and robust phase locking between oscillators than nearest-neighbor coupling or locally dense connection schemes. This observation is at the root of the recent findings about synchronization in complex networks of oscillators. In the current section we review the results obtained so far on three different kinds of oscillatory ensembles: limit cycle oscillators (Kuramoto), pulse-coupled models, and finally coupled map systems. We reserve for Sect. IV those works that use the MSF formalism. Many other works whose major contribution is the understanding of synchronization phenomena in specific scenarios are discussed in the Applications section.

#### A. Phase oscillators

1. The Kuramoto model

The pioneering work by Winfree (1967) spurred the field of collective synchronization and called for mathematical approaches to tackle the problem. One of these approaches, as already stated, considers a system made up of a huge population of weakly-coupled, nearly identical, interacting limit-cycle oscillators, where each oscillator exerts a phase dependent influence on the others and changes its rhythm according to a sensitivity function (Acebrón et al., 2005; Strogatz, 2000).

Even if these simplifications seem to be very crude, the phenomenology of the problem can be captured. Namely, the population of oscillators exhibits the dynamic analog to an equilibrium phase transition. When the natural frequencies of the oscillators are too diverse compared to the strength of the coupling, they are unable to synchronize and the system behaves incoherently. However, if the coupling is strong enough, all oscillators freeze into synchrony. The transition from one regime to the other takes place at a certain threshold. At this point some elements lock their relative phase and a cluster of synchronized nodes develops. This constitutes the \textit{onset of synchronization}. Beyond this value, the population of oscillators is split into a partially synchronized state made up of oscillators locked in phase and a group of nodes whose natural frequencies are too different as to be part of the coherent cluster. Finally, after further increasing the coupling, more and more elements get entrained around the mean phase of the collective rhythm generated by the whole population and the system settles in the completely synchronized state.

Kuramoto (1975, 1984) worked out a mathematically tractable model to describe this phenomenology. He recognized that the most suitable case for analytical treatment should be the mean field approach. He proposed an all-to-all purely sinusoidal coupling, and then the governing equations for each of the oscillators in the system are:

$$\dot{\theta}_i = \omega_i + \frac{K}{N} \sum_{j=1}^N \sin(\theta_j - \theta_i) \quad (i = 1, ..., N),$$

where the factor $1/N$ is incorporated to ensure a good behavior of the model in the thermodynamic limit, $N \to \infty$, $\omega_i$ stands for the natural frequency of oscillator $i$, and $K$ is the coupling constant. The frequencies $\omega_i$ are distributed according to some function $g(\omega)$, that is usually assumed to be unimodal and symmetric about its mean frequency $\Omega$. Admittedly, due to the rotational symmetry in the model, we can use a rotating frame and redefine
\[ \omega_i \rightarrow \omega_i + \Omega \text{ for all } i \text{ and set } \Omega = 0, \text{ so that the } \omega_i \text{'s denote deviations from the mean frequency.} \]

The collective dynamics of the whole population is measured by the macroscopic complex order parameter,

\[ r(t) e^{i\phi(t)} = \frac{1}{N} \sum_{j=1}^{N} e^{i\theta_j(t)}, \quad (4) \]

where the modulus \( 0 \leq r(t) \leq 1 \) measures the phase coherence of the population and \( \phi(t) \) is the average phase. The values \( r \approx 1 \) and \( r \approx 0 \) (where \( \approx \) stands for fluctuations of size \( O(N^{-1/2}) \)) describe the limits in which all oscillators are either phase locked or move incoherently, respectively. Multiplying both parts of Eq. (4) by \( e^{-i\theta_i} \) and equating imaginary parts gives

\[ r \sin(\phi - \theta_i) = \frac{1}{N} \sum_{j=1}^{N} \sin(\theta_j - \theta_i), \quad (5) \]

yielding

\[ \dot{\theta}_i = \omega_i + Kr \sin(\phi - \theta_i) \quad (i = 1, ..., N). \quad (6) \]

Equation (6) states that each oscillator interacts with all the others only through the mean field quantities \( r \) and \( \phi \). The first quantity provides a positive feedback loop to the system’s collective rhythm: as \( r \) increases because the population becomes more coherent, the coupling between the oscillators is further strengthened and more of them can be recruited to take part in the coherent pack. Moreover, Eq. (6) allows to calculate the critical coupling \( K_c \) and to characterize the order parameter \( \lim_{t \rightarrow \infty} r_t(K) = r(K) \). Looking for steady solutions, one assumes that \( r(t) \) and \( \phi(t) \) are constant. Next, without loss of generality, we can set \( \phi = 0 \), which leads to the equations of motion (Kuramoto, 1975, 1984)

\[ \dot{\theta}_i = \omega_i - Kr \sin(\theta_i) \quad (i = 1, ..., N). \quad (7) \]

The solutions of Eq. (7) reveal two different types of long-term behavior when the coupling is larger than the critical value, \( K_c \). On the one hand, a group of oscillators for which \( |\omega_i| \leq Kr \) are phase-locked at frequency \( \Omega \) in the original frame according to the equation \( \omega_i = Kr \sin(\theta_j) \). On the other hand, the rest of the oscillators for which \( |\omega_i| > Kr \) holds, are drifting around the circle, sometimes accelerating and sometimes rotating at lower frequencies. Demanding some conditions for the stationary distribution of drifting oscillators with frequency \( \omega_i \) and phases \( \theta_i \) (Strogatz, 2000), a self-consistent equation for \( r \) can be derived as

\[ r = Kr \int_{-\pi/2}^{\pi/2} \left( \cos^2 \theta \right) g(\omega) d\theta, \]

where \( \omega = Kr \sin(\theta) \). This equation admits a non-trivial solution,

\[ K_c = \frac{2}{\pi g(0)} \quad (8) \]

beyond which \( r > 0 \). Equation (8) is the Kuramoto mean field expression for the critical coupling at the onset of synchronization. Moreover, near the onset, the order parameter, \( r \), obeys the usual square-root scaling law for mean field models, namely,

\[ r \sim (K - K_c)^{\beta} \quad (9) \]

with \( \beta = 1/2 \). Numerical simulations of the model verified these results. The Kuramoto model (KM, from now on) approach to synchronization was a breakthrough for the understanding of synchronization in large populations of oscillators.

Even in the simplest case of a mean field interaction, there are still unsolved problems that have resisted any analytical attempt. This is the case, e.g., for finite populations of oscillators and some questions regarding global stability results (Acebrón et al., 2005). In what follows, we focus on another aspect of the model’s assumptions, namely that of the connection topology of real systems (Boccaletti et al., 2006; Newman, 2003b), which usually do not show the all-to-all pattern of interconnections underneath the mean field approach.

2. Kuramoto model on complex networks

To deal with the KM on complex topologies, it is necessary to reformulate Eq. (3) to include the connectivity

\[ \dot{\theta}_i = \omega_i + \sum_j \sigma_{ij} a_{ij} \sin(\theta_j - \theta_i) \quad (i = 1, ..., N), \quad (10) \]

where \( \sigma_{ij} \) is the coupling strength between pairs of connected oscillators and \( a_{ij} \) are the elements of the connectivity matrix. The original Kuramoto model is recovered by letting \( a_{ij} = 1, \forall i \neq j \) (all-to-all) and \( \sigma_{ij} = K/N, \forall i, j \).

The first problem when defining the KM in complex networks is how to state the interaction dynamics properly. In contrast with the mean field model, there are several ways to define how the connection topology enters in the governing equations of the dynamics. A good theory for Kuramoto oscillators in complex networks should be phenomenologically relevant and provide formulas amenable to rigorous mathematical treatment. Therefore, such a theory should at least preserve the essential fact of treating the heterogeneity of the network independently of the interaction dynamics, and at the same time, should remain calculable in the thermodynamic limit.

For the original model, Eq. (3), the coupling term on the right hand side of Eq. (10) is an intensive magnitude because the dependence on the size of the system cancels out. This independence on the number of oscillators \( N \) is achieved by choosing \( \sigma_{ij} = K/N \). This prescription turns out to be essential for the analysis of the system in the thermodynamic limit \( N \rightarrow \infty \) in the all-to-all case. However, choosing \( \sigma_{ij} = K/N \) for the governing equations of
the KM in a complex network makes them to become dependent on \( N \). Therefore, in the thermodynamic limit, the coupling term tends to zero except for those nodes with a degree that scales with \( N \). Note that the existence of such nodes is only possible in networks with power-law degree distributions (Boccaletti et al., 2006; Newman, 2003b), but this happens with a very small probability as \( k^{-\gamma} \), with \( \gamma > 2 \). In these cases, mean field solutions independent of \( N \) are recovered, with slight differences in the onset of synchronization of all-to-all and SF networks (Restrepo et al., 2005a).

A second prescription consists in taking \( \sigma_{ij} = K/k_i \) (where \( k_i \) is the degree of node \( i \)) so that \( \sigma_{ij} \) is a weighted interaction factor that also makes the right hand side of Eq. (10) intensive. This form has been used to solve the paradox of heterogeneity (Motter et al., 2005b) that states that the heterogeneity in the degree distribution, which often reduces the average distance between nodes, may suppress synchronization in networks of oscillators coupled symmetrically with uniform coupling strength. This result refers to the stability of the fully synchronized state, but not to the dependence of the order parameter on the coupling strength (where partially synchronized and unsynchronized states exist). Besides, the inclusion of weights in the interaction strongly affects the original KM dynamics in complex networks because it can impose a dynamic homogeneity that masks the real topological heterogeneity of the network.

The prescription \( \sigma_{ij} = K/\text{const} \), which may seem more appropriate, also causes some conceptual problems because the sum in the right hand side of Eq. (10) could eventually diverge in the thermodynamic limit. The constant in the denominator could in principle be any quantity related to the topology, such as the average connectivity of the graph \( \langle k \rangle \), or the maximum degree \( k_{\text{max}} \). Its physical meaning is a re-scaling of the temporal scales involved in the dynamics. However, except for the case of \( \sigma_{ij} = K/k_{\text{max}} \), the other possible settings do not avoid the problems when \( N \rightarrow \infty \). On the other hand, for a proper comparison of the results obtained for different complex topologies (e.g. SF or uniformly random), the global and local measures of coherence should be represented according to their respective time scales. Therefore, given two complex networks \( A \) and \( B \) with \( k_{\text{max}} = k_A \) and \( k_{\text{max}} = k_B \) respectively, it follows that to make meaningful comparisons between observables, the equations of motion Eq. (10) should refer to the same time scales, i.e., \( \sigma_{ij} = K_A/k_A = K_B/k_B = \sigma \). With this formulation in mind, Eq. (10) reduces to

\[
\dot{\theta}_i = \omega_i + \sigma \sum_j a_{ij} \sin(\theta_j - \theta_i) \quad (i = 1, ..., N),
\]

independently of the specific topology of the network. This allows us to study the dynamics of Eq. (11) on different topologies, compare the results, and properly inspect the interplay between topology and dynamics in what concerns synchronization.

As we shall see, there are also several ways to define the order parameter that characterizes the global dynamics of the system, some of which were introduced to allow for analytical treatments at the onset of synchronization. We advance, however, that the same order parameter, Eq. (4), is often used to describe the coherence of the synchronized state.

3. Onset of synchronization in complex networks

Studies on synchronization in complex topologies where each node is considered to be a Kuramoto oscillator, were first reported for WS networks (Hong et al., 2002; Watts, 1999) and BA graphs (Moreno and Pacheco, 2004; Moreno et al., 2004). These works are mainly numerical explorations of the onset of synchronization, their main goal being the characterization of the critical coupling beyond which groups of nodes beating coherently first appear. Hong et al. (2002) considered oscillators with intrinsic frequencies distributed according to a Gaussian distribution with unit variance arranged in a WS network with varying rewiring probability, \( p \), and explored how the order parameter, Eq. (4), changes upon addition of long-range links. Moreover, they assumed a normalized coupling strength \( \sigma_{ij} = K/\langle k \rangle \), where \( k \) is the average degree of the graph. Numerical integration of the equations of motion (10) under variation of \( p \) shows that collective synchronization emerges even for very small values of the rewiring probability.

The results confirm that networks obtained from a regular ring by just rewiring a tiny fraction of links (\( p \geq 0 \)) can be synchronized with a finite \( K \). Moreover, in contrast with the arguments provided by Hong et al. (2002), we notice that their results had been obtained for a fixed average degree and thus the Kuramoto’s critical coupling can not be recovered by simply taking \( p \rightarrow 1 \), which produces a random ER graph with a fixed minimum connectivity. This limit is recovered by letting \( (k) \) increase. Actually, numerical simulations of the same model by Watts (1999) showed that the Kuramoto limit is approached when the average connectivity grows.

Moreno and Pacheco (2004) considered the same problem in BA networks. The natural frequencies and the initial values of \( \theta_i \) were randomly drawn from a uniform distribution in the interval \((-1/2, 1/2)\) and \((-\pi, \pi)\), respectively. The global dynamics of the system, Eq. (11), turns out to be qualitatively the same as for the original KM as shown in Fig. 2, where the dependence of the order parameter Eq. (4) with \( \sigma \) is shown for several system sizes.

The existence of a critical point for the KM on SF networks came as a surprise. Admittedly, this is one of the few cases in which a dynamical process shows a critical behavior when the substrate is described by a power-law connectivity distribution with an exponent \( \gamma \leq 3 \) (Boccaletti et al., 2006; Dorogovtsev et al., 2007; Newman, 2003b). In principle it could be a finite size effect, but it turned out from numerical simulations that this was
not the case. To determine the exact value of $\sigma_c$, one can make use of standard finite-size scaling analysis. At least two complementary strategies have been reported. The first one allows bounding the critical point and is computationally more expensive. Consider a network of size $N$, for which no synchronization is attained below $\sigma_c$, where $r(t)$ decays to a small residual value of size $O(1/\sqrt{N})$. Then, the critical point may be found by examining the $N$-dependence of $r(\sigma, N)$. In the sub-critical regime ($\sigma < \sigma_c$), the stationary value of $r$ falls off as $N^{-1/2}$, while for $\sigma > \sigma_c$, the order parameter reaches a stationary value as $N \rightarrow \infty$ (though still with $O(1/\sqrt{N})$ fluctuations). Therefore, plots of $r$ versus $N$ allow us to locate the critical point $\sigma_c$. Alternatively, a more accurate approach can be adopted. Assume the scaling form for the order parameter (Marro and Dickman, 1999):

$$r = N^{-\alpha} f(N^\nu(\sigma - \sigma_c)),$$  
(12)

where $f(x)$ is a universal scaling function bounded as $x \rightarrow \pm \infty$ and $\alpha$ and $\nu$ are two critical exponents to be determined. Since at $\sigma = \sigma_c$, the value of the function $f$ is independent of $N$, the estimation of $\sigma_c$ can be done by plotting $N^\alpha f$ as a function of $\sigma$ for various sizes $N$ and then finding the value of $\alpha$ that gives a well-defined crossing point, the critical coupling $\sigma_c$. As a by-product, the method also allows us to calculate the two scaling exponents $\alpha$ and $\nu$, the latter can be obtained from the equality

$$\ln[(dr/d\sigma)]_{\sigma_c} = (\nu - \alpha) \ln N + \text{const},$$  
(13)

once $\alpha$ is computed.

Following these scaling procedures, Moreno and Pacheco (2004) and Gómez-Gardeñes et al. (2007a,b) estimated a value for the critical coupling strength $\sigma_c = 0.05(1)$. Moreover, $r \sim (\sigma - \sigma_c)^\beta$ when approaching the critical point from above with $\beta = 0.46(2)$ indicating that the square-root behavior typical of the mean field version of the model ($\beta = 1/2$) seems to hold as well for BA networks.

Before turning our attention to some theoretical attempts to tackle the onset of synchronization, it is worth to briefly summarize other numerical results that have explored how the critical coupling depends on other topological features of the underlying SF graph. Recent results have shed light on the influence of the topology of the local interactions on the route to and the onset of synchronization. In particular, McGraw and Menzinger (2005, 2007) and Gómez-Gardeñes and Moreno (2007) explored the Kuramoto dynamics on networks in which the degree distribution is kept fixed, while the clustering coefficient ($C$) and the average path length ($l$) of the graph change. The results suggest that the onset of synchronization is mainly determined by $C$, namely, networks with a high clustering coefficient promote synchronization at lower values of the coupling strength. On the other hand, when the coupling is increased beyond the critical point, the effect of $l$ dominates over $C$ and the phase diagram is smoothed out (a sort of stretching), delaying the appearance of the fully synchronized state as the average shortest path length increases.

In a series of recent papers (Ichinomiya, 2004, 2005; Lee, 2005; Restrepo et al., 2005a,b, 2006), the onset of synchronization in large networks of coupled oscillators has been analyzed from a theoretical point of view. Despite these efforts no exact analytical results for the KM on general complex networks are available up to date. Moreover, the theory predicts that for uncorrelated SF networks with an exponent $\gamma \leq 3$, the critical coupling vanishes as $N \rightarrow \infty$, in contrast to numerical simulations on BA model networks. It appears that the strong heterogeneity of real networks and the finite average connectivity strongly hampers analytical solutions of the model.

Following Restrepo et al. (2005a), consider the system in Eq. (11), with a symmetric\(^1\) adjacency matrix $a_{ij} = a_{ji}$. Defining a local order parameter $r_i$ as

$$r_i e^{i \phi_i} = \sum_{j=1}^{N} a_{ij} \langle e^{i \theta_j} \rangle_t,$$  
(14)

where $\langle \cdot \cdot \cdot \rangle_t$ stands for a time average, a new global order parameter to measure the macroscopic coherence is readily introduced as

$$r = \frac{\sum_{i=1}^{N} r_i}{\sum_{i=1}^{N} k_i}.$$  
(15)

Now, rewriting Eq. (11) as a function of $r_i$, yields,

$$\dot{\theta}_i = \omega_i - \sigma_i \sin(\theta_i - \phi_i) - \sigma b_i(t).$$  
(16)

---

\(^1\) The reader can find the extension of the forthcoming formalism to directed networks in Restrepo et al. (2005b).
In Eq. (16), \( h_i(t) = \text{Im}\{e^{-i\theta_i} \sum_{j=1}^N a_{ij}(e^{i\theta_j})_t - e^{i\theta_j}) \} \) depends on time and contains time fluctuations. Assuming the terms in the previous sum to be statistically independent, \( h_i(t) \) is expected to be proportional to \( \sqrt{K_i} \) above the transition, where \( r_i \sim k_i \). Therefore, except very close to the critical point, and assuming that the number of connections of each node is large enough \( (k_i \gg 1) \) as to be able to neglect the time fluctuations entering \( h_i \), i.e., \( h_i \ll r_i \), the equation describing the dynamics of node \( i \) can be reduced to

\[
\dot{\theta}_i = \omega_i - \sigma r_i \sin(\theta_i - \phi_i). \tag{17}
\]

Next, we look for stationary solutions of Eq. (17), i.e., \( \sin(\theta_i - \phi_i) = \omega_i / \sigma r_i \). In particular, oscillators whose intrinsic frequency satisfies \( |\omega_i| \leq \sigma r_i \) become locked. Then, as in the Kuramoto mean field model, there are two contributions (though in this case to the local order parameter), one from locked and the other from drifting oscillators such that

\[
r_i = \sum_{j=1}^N a_{ij}(e^{i(\theta_j - \phi_j)})_t = \sum_{|\omega_j| \leq \sigma r_j} a_{ij} e^{i(\theta_j - \phi_j)} + \sum_{|\omega_j| > \sigma r_j} a_{ij} e^{i(\theta_j - \phi_j)}_t. \tag{18}
\]

To move one step further, some assumptions are needed. Consider a graph such that the average degree of nearest neighbors is high (i.e., if the neighbors of node \( i \) are well-connected). Then it is reasonable to assume that these nodes are not affected by the intrinsic frequency of \( i \). This is equivalent to assume solutions \( (r_i, \phi_i) \) that are, in a statistical sense, independent of the natural frequency \( \omega_i \). With this assumption, the second summand in Eq. (18) can be neglected. Taking into account that the distribution \( g(\omega) \) is symmetric and centered at \( \Omega = 0 \), after some algebra one is left with (Restrepo et al., 2005a)

\[
r_i = \sum_{|\omega_j| \leq \sigma r_j} a_{ij} \cos(\phi_j - \phi_i) \sqrt{1 - \left( \frac{\omega_j}{\sigma r_j} \right)^2}. \tag{19}
\]

The critical coupling \( \sigma_c \) is given by the solution of Eq. (19) that yields the smallest \( \sigma \). It can be argued that it is obtained when \( \cos(\phi_j - \phi_i) = 1 \) in Eq. (19), thus

\[
r_i = \sum_{|\omega_j| \leq \sigma r_j} a_{ij} \sqrt{1 - \left( \frac{\omega_j}{\sigma r_j} \right)^2}, \tag{20}
\]

which is the main equation of the time average approximation (recall that time fluctuations have been neglected). Note, however, that to obtain the critical coupling, one has to know the adjacency matrix as well as the particular values of \( \omega_i \) for all \( i \) and then solve Eq. (20) numerically for the \( \{r_i\} \). Finally, the global order parameter defined in Eq. (15) can be computed from \( r_i \).

Even if the underlying graph satisfies the other aforementioned topological constraints, it seems unrealistic to require knowledge of the \( \{\omega_i\} \)'s. A further approach, referred to as the frequency distribution approximation can be adopted. According to the assumption that \( k_i \gg 1 \) for all \( i \), or equivalently, that the number of connections per node is large (a dense graph), one can also consider that the natural frequencies of the neighbors of node \( i \) follows the distribution \( g(\omega) \). Then, Eq. (20) can be rewritten avoiding the dependence on the particular realization of \( \{\omega_i\} \) to yield,

\[
r_i = \sum a_{ij} \int_{-\sigma r_j}^{\sigma r_j} g(\omega) \sqrt{1 - \left( \frac{\omega}{\sigma r_j} \right)^2} d\omega = \sigma \sum a_{ij} r_j \int_{-1}^{1} g(x\sigma r_j) \sqrt{1 - x^2} dx, \tag{21}
\]

with \( x = \omega / (\sigma r_j) \). This equation allows us to readily determine the order parameter \( r \) as a function of the network topology \((a_{ij})\), the frequency distribution \( g(\omega) \) and the control parameter \( \sigma \). On the other hand, Eq. (20) still does not provide explicit expressions for the order parameter and the critical coupling strength. To this end, one introduces a first-order approximation \( g(x\sigma r_j) \approx g(0) \) which is valid for small, but nonzero, values of \( r \). Namely, when \( r_j \to 0^+ \)

\[
r_i^0 = \frac{\sigma}{K_c} \sum_j a_{ij} r_j^0, \tag{22}
\]

where \( K_c = 2/(\pi g(0)) \) is Kuramoto’s critical coupling. Moreover, as the smallest value of \( \sigma \) corresponds to \( \sigma_c \), it follows that the critical coupling is related to both \( K_c \) and the largest eigenvalue \( \lambda_{\text{max}} \) of the adjacency matrix, yielding

\[
\sigma_c = \frac{K_c}{\lambda_{\text{max}}}. \tag{22}
\]

Equation (22) states that in complex networks, synchronization is first attained at a value of the coupling strength that inversely depends on \( g(0) \) and on the largest eigenvalue \( \lambda_{\text{max}} \) of the adjacency matrix. Note that this equation also recovers Kuramoto’s result when \( a_{ij} = 1, \forall i \neq j \), since \( \lambda_{\text{max}} = N - 1 \). It is worth stressing that although this method allows us to calculate \( \sigma_c \) analytically, it fails to explain why for uncorrelated random SF networks with \( \gamma \leq 3 \) and in the thermodynamic limit \( N \to \infty \), the critical value remains finite. This disagreement comes from the fact that in these SF networks, \( \lambda_{\text{max}} \) is proportional to the cutoff of the degree distribution, \( k_{\text{max}} \) which in turn scales with the system size. Putting the two dependencies together, one obtains \( \lambda_{\text{max}} \sim k_{\text{max}}^{-\gamma} \sim N^{-\frac{1}{\gamma-1}} \to \infty \) as \( N \to \infty \), thus predicting

\[\text{This obviously restricts the range of real networks to which the approximation can be applied.}\]
\( \sigma_c = 0 \) in the thermodynamic limit, in contrast to finite size scaling analysis for the critical coupling via numerical solution of the equations of motion. Note, however, that the difference may be due to the use of distinct order parameters. Moreover, even in the case of SF networks with \( \gamma > 3 \), \( \lambda_{\text{max}} \) still diverges when we take the thermodynamic limit, so that \( \sigma_c \rightarrow 0 \) as well. As we shall see soon, this is not the case when other approaches are adopted, at least for \( \gamma > 3 \).

It is possible to go beyond with the latter approximation and to determine the behavior of \( r \) near the critical point. Restrepo et al. (2005a) developed a perturbative approach to higher orders of Eq. (21), which is valid for relatively homogeneous degree distributions (\( \gamma > 5 \)).\(^3\) They showed that for \( (\sigma/\sigma_c) - 1 \sim 0^+ \)

\[
r^2 = \left( \frac{\eta}{\eta_1 K_c} \right) \left( \frac{\sigma}{\sigma_c} - 1 \right) \left( \frac{\sigma}{\sigma_c} \right)^3 ,
\]

where \( \eta_1 = -\pi g''(0) K_c / 16 \) and

\[
\eta = \frac{\langle u \rangle^2 \lambda_{\text{max}}^2}{N \langle k \rangle^2 \langle u^4 \rangle} ,
\]

where \( u \) is the normalized eigenvector of the adjacency matrix corresponding to \( \lambda_{\text{max}} \) and \( \langle u^4 \rangle = \sum_i u_i^4 / N \).

The analytical insights discussed so far can also be reformulated in terms of a mean field approximation (Ichinomiya, 2004, 2005; Lee, 2005; Restrepo et al., 2005a) for complex networks. This approach (valid for large enough \( k \)) considers that every oscillator is influenced by the local field created in its neighborhood, so that \( r_i \) is proportional to the degree of the nodes \( k_i \), i.e., \( r_i \sim k_i \). Assuming this is the case and introducing the order parameter \( r \) through

\[
r = \frac{r_i}{k_i} = \frac{1}{k_i} \left| \sum_{j=1}^{N} a_{ij} e^{i \theta_j} \right| ,
\]

after summing over \( i \) and substituting \( r_i = r k_i \) in Eq. (21) we obtain (Restrepo et al., 2005a)

\[
\sum_{j}^{N} k_j = \sigma \sum_{j}^{N} k_j^2 \int_{-1}^{1} g(x \sigma k_j) \sqrt{1-x^2} dx .
\]

The above relation, Eq. (26), was independently derived by Ichinomiya (2004), who first studied analytically the problem of synchronization in complex networks, though using a different approach. Taking the continuum limit, Eq. (26) becomes

\[
\int kP(k)dk = \sigma \int k^2 P(k)dk \int_{-1}^{1} g(x \sigma k) \sqrt{1-x^2} dx ,
\]

which for \( r \rightarrow 0^+ \) verifies

\[
\int kP(k)dk = \sigma \int k^2 P(k)dk \int_{-1}^{1} g(0) \sqrt{1-x^2} dx
\]

\[
= \frac{\sigma g(0) \pi}{2} \int k^2 P(k)dk ,
\]

which leads to the condition for the onset of synchronization \((r > 0)\) as

\[
\frac{\sigma g(0) \pi}{2} \int k^2 P(k)dk > \int kP(k)dk ,
\]

that is,

\[
\sigma_c = \frac{2}{\pi g(0) \langle k^2 \rangle} K_c \langle k \rangle \langle k^2 \rangle .
\]

The mean field result, Eq. (29), gives as a surprising result that the critical coupling \( \sigma_c \) in complex networks is nothing else but the one corresponding to the all-to-all topology \( K_c \) re-scaled by the ratio between the first two moments of the degree distribution, regardless of the many differences between the patterns of interconnections. Precisely, it states that the critical coupling strongly depends on the topology of the underlying graph. In particular, \( \sigma_c \rightarrow 0 \) when the second moment of the distribution \( \langle k^2 \rangle \) diverges, which is the case for SF networks with \( \gamma \leq 3 \). Note, that in contrast with the result in Eq. (22), for \( \gamma > 3 \), the coupling strength does not vanish in the thermodynamic limit. On the other hand, if the mean degree is kept fixed and the heterogeneity of the graph is increased by decreasing \( \gamma \), the onset of synchronization occurs at smaller values of \( \sigma_c \). Interestingly enough, the dependence gathered in Eq. (29) has the same functional form for the critical points of other dynamical processes such as percolation and epidemic spreading processes (Boccaletti et al., 2006; Dobogókó et al., 2007; Newman, 2003b). While this result is still under numerical scrutiny, it would imply that the critical properties of many dynamical processes on complex networks are essentially determined by the topology of the graph, no matter whether the dynamics is nonlinear or not. The corroboration of this last claim will be of extreme importance in physics, probably changing many preconceived ideas about the nature of dynamical phenomena.

Within the mean field theory, it is also possible to obtain the behavior of the order parameter \( r \) near the transition to synchronization. Lee (2005) also independently derived Eq. (29) starting from the differential equation Eq. (11). Using the weighted order parameter

\[
\bar{r}(t)e^{i \bar{\phi}(t)} = \frac{\sum_{i=1}^{N} \sum_{j=1}^{N} a_{ij} e^{i \theta_j} k_i}{\sum_{i=1}^{N} k_i} ,
\]

and assuming the same magnitude of the effective field of each pair of coupled oscillators one obtains

\[
\dot{\theta}_i = \omega_i - \frac{\sigma}{\langle k \rangle} k_i \bar{r} \sin(\theta_i) ,
\]

\(^3\) The approach holds if the fourth moment of the degree distribution, \( \langle k^4 \rangle = \int P(k)k^4dk \) remains finite when \( N \rightarrow \infty \).
where we have set $\bar{\phi} = 0$. Now, it is considered again that in the stationary state the system divides into two groups of oscillators, which are either locked or rotating in a nonuniform manner. Following the same procedure employed in all the previous derivations, the only contribution to $r$ comes from the former set of oscillators. After some algebra, Lee (2005) showed that the critical coupling $\sigma_c$ is given by Eq. (29) and that near criticality

$$r \sim (\sigma - \sigma_c)^{3},$$

(31)

for $\gamma > 3$, with a critical exponent $\beta = \frac{1}{2}$ if $\gamma \geq 5$, and $\beta = \frac{2}{3}$ when $3 < \gamma \leq 5$. For the most common cases in real networks of $2 < \gamma < 3$, the critical coupling tends to zero in the thermodynamic limit so that $r$ should be nonzero as soon as $\sigma \neq 0$. In this case, one gets $r \sim \sigma^{1/(3-\gamma)}$. Notably, the latter equation is exactly the same found for the absence of critical behavior in the region $2 < \gamma < 3$ for a model of epidemic spreading (Moreno et al., 2002).

One recent theoretical study by Oh et al. (2007) is worth mentioning here. They have extended the mean field approach to the case in which the coupling is asymmetric and depends on the degree. In particular, they studied a system of oscillators arranged in a complex topology whose dynamics is given by

$$\dot{\theta}_i = \omega_i + \frac{\sigma}{k_i} \sum_{j=1}^{N} \sin(\theta_j - \theta_i).$$

(32)

$\eta = 1$ corresponds to the symmetric, non-degree dependent case. Extending the mean field formalism to the cases $\eta \neq 1$, they investigated the nature of the synchronization transition as a function of the magnitude and sign of the parameter $\eta$. By exploring the whole parameter space $(\eta, \sigma)$, they found that for $\eta = 0$ and SF networks with $2 < \gamma < 3$, a finite critical coupling $\sigma_c$ is recovered in sharp contrast to the non-weighted coupling case for which we already know that $\sigma_c = 0$. This result seems phenomenologically meaningful, since setting $\eta = 0$ implies that the coupling in Eq. (10) is $\sigma_{ij} = \sigma/k_i$, which, as discussed before (Motter et al., 2005b), might have the effect of partially destroying the heterogeneity inherent to the underlying graph by normalizing all the contributions $\sum_{j=1}^{N} a_{ij} \sin(\theta_j - \theta_i)$ by $k_i = \sum_{j=1}^{N} a_{ij}$.

4. Path towards synchronization in complex networks

Up to now, we have discussed both numerically and theoretically the onset of synchronization. In the next section, we shall also discuss how the structural properties of the networks influence the stability of the fully synchronized state. But, what happens in the region where we are neither close to the onset of synchronization nor at complete synchronization? How is the latter state attained when different topologies are considered?

As we have seen, the influence of the topology is not only given by the degree distribution, but also by how the oscillators interact locally. To reduce the number of degrees of freedom to a minimum, let us focus on the influence of heterogeneity and study the evolution of synchronization for a family of complex networks which are comparable in their clustering, average distance and correlations so that the only difference is due to the degree distribution. For these networks, the previous theoretical approaches argued that the critical coupling $\sigma_c$ is proportional to $(k)/(k^2)$, so that different topologies should give rise to distinct critical points. In particular, Gómez-Gardeñas et al. (2007a,b) studied numerically the path towards synchronization in ER and SF networks. They also studied several networks whose degree of heterogeneity can be tuned between the two limiting cases (Gómez-Gardeñas and Moreno, 2006). These authors put forward the question: How do SF networks compare with ER ones and what are the roots of the different behaviors observed?

Numerical simulations (Gómez-Gardeñas et al., 2007a,b) confirm qualitatively the theoretical predictions for the onset of synchronization, as summarized in Table I. In fact, the onset of synchronization first occurs for SF networks. As the network substrate becomes more homogeneous, the critical point $\sigma_c$ shifts to larger values and the system seems to be less synchronous. On the other hand, they also showed that the route to complete synchronization, $r = 1$, is sharper for homogeneous networks. No critical exponents for the behavior of $r$ near the transition points have been reported yet for the ER network, so that comparison with the mean field value $\beta = 1/2$ for a SF network with $\gamma = 3$ is not possible.

\begin{table}[h]
\centering
\caption{Topological properties of the networks and critical coupling strengths derived from a finite size scaling analyses, Eq. (12). Different values of $\chi$ corresponds to grown networks whose degree of heterogeneity varies smoothly between the two limiting cases of ER and SF graphs. From Gómez-Gardeñas et al. (2007b).}
\begin{tabular}{cccc}
\hline
$\chi$ & $(k^2)$ & $k_{\text{max}}$ & $\sigma_c$ \\
\hline
0.0 (SF) & 115.5 & 326.3 & 0.051 \\
0.2 & 56.7 & 111.6 & 0.066 \\
0.4 & 44.9 & 47.7 & 0.088 \\
0.6 & 41.1 & 25.6 & 0.103 \\
0.8 & 39.6 & 16.8 & 0.108 \\
1.0 (ER) & 39.0 & 14.8 & 0.122 \\
\hline
\end{tabular}
\end{table}

4 This isolation of individual features of complex networks is essential to understand the interplay between topology and dynamics. As we will discuss along the review, many times this aspect has not been properly controlled raising results that are confusing, contradictory or even incorrect.

5 The numerical value of $\beta$ contradicts the prediction of the mean-field approach (see the discussion after Eq. (31)) The reason of such discrepancy is not clear yet.
the symmetric matrix

\[ \mathbf{L} \]

chronization of pairs of connected oscillators in terms of

\[ r_{\text{link}}(\sigma) \]

to add edges between nodes already belonging to such

\[ \sigma_{\text{SF}} \]

\[ \sigma_{\text{ER}} \]

\[ \langle \tau \rangle \sim k^{-1} \]

\[ T \]

to the effective coupling is increased. On the contrary, in heterogeneous graphs, what is added are links between nodes already belonging to such cluster. From Gómez-Gardeñes et al. (2007a).

numerically, a detailed finite size scaling analysis in SF and ER topologies shows that the critical coupling strength corresponds in SF networks to \( \sigma_{\text{SF}} = 0.051 \), and in random ER networks to \( \sigma_{\text{ER}} = 0.122 \), a fairly significant numerical difference.

The mechanisms behind the differences in the emergence of collective behavior for ER and SF topologies can be explored numerically by defining a local order parameter that captures and quantifies the way in which clusters of locked oscillators emerge. The main difference with respect to \( r \) is that one measures the degree of synchronization of nodes \( (r) \) with respect to the average phase \( \phi \) and the other \( (r_{\text{link}}) \) to the degree of synchronization between every pair of connected nodes. Thus, \( r_{\text{link}} \) gives the fraction of all possible links that are synchronized in the network as

\[ r_{\text{link}} = \frac{1}{2N} \sum_{i,j=1}^{N} a_{ij} \left| \lim_{\Delta t \to \infty} \frac{1}{\Delta t} \int_{t_r}^{t_r + \Delta t} e^{i[\theta_i(t) - \theta_j(t)]} dt \right| , \]

being \( t_r \) the time the system needs to settle into the stationary state, and \( \Delta t \) a large averaging time. Gómez-Gardeñes et al. (2007a,b) measured the degree of synchronization of pairs of connected oscillators in terms of the symmetric matrix

\[ D_{ij} = a_{ij} \left| \lim_{\Delta t \to \infty} \frac{1}{\Delta t} \int_{t_r}^{t_r + \Delta t} e^{i[\theta_i(t) - \theta_j(t)]} dt \right| , \]

which, once filtered using a threshold \( T \) such that the fraction of synchronized pairs equals \( r_{\text{link}} \), allows us to identify the synchronized links and reconstruct the clusters of synchrony for any value of \( \sigma \), as illustrated in Fig.3. From a microscopic analysis, it turns out that for homogeneous topologies, many small clusters of synchronized pairs of oscillators are spread over the graph and merge together to form a giant synchronized cluster when the effective coupling is increased. On the contrary, in heterogeneous graphs, a central core containing the hubs first comes up driving the evolution of synchronization patterns by absorbing small clusters. Moreover, the evolution of \( r_{\text{link}}(\sigma) \) as \( \sigma \) grows explains why the transition is sharper for ER networks: nodes are added first to the giant synchronized cluster and later on the links among these nodes that were missing in the original clusters of synchrony. In SF graphs, oscillators are added to the largest synchronized component together with most of their links, resulting in a much slower growth of \( r_{\text{link}} \). Finally, Gómez-Gardeñes et al. (2007a,b) also computed the probability that a node with degree \( k \) belongs to the largest synchronized cluster and reported that this probability is an increasing function of \( k \) for every \( \sigma \), namely, the more connected a node is, the more likely it takes part in the cluster of synchronized links. It is interesting to mention here that a similar dependence is obtained if one analyzes the stability of the synchronized state under perturbations of nodes of degree \( k \). Moreno and Pacheco (2004) found that the average time \( \langle \tau \rangle \) a node needs to get back into the fully synchronized state is inversely proportional to its degree, i.e., \( \langle \tau \rangle \sim k^{-1} \).

Very recently, Almendral and Díaz-Guilera (2007) also studied the path towards synchronization, looking for the relation between the time needed for complete synchronization and the spectral properties of the Laplacian matrix of the graph,

\[ L_{ij} = k_i \delta_{ij} - a_{ij}. \]

The Laplacian matrix is symmetric with zero row-sum and hence all the eigenvalues are real and non-negative. Considering the case of identical Kuramoto oscillators, whose dynamics has only one attractor, the fully synchronized state, they found that the synchronization time scales with the inverse of the smallest nonzero eigenvalue of the Laplacian matrix. Surprisingly, this relation qualitatively holds for very different networks where synchronization is achieved, indicating that this eigenvalue alone might be a relevant topological property for synchronization phenomena. Donetti et al. (2006) remark the role of this eigenvalue not only for synchronization purposes but also for the flow of random walkers on the network.

5. Kuramoto model on structured or modular networks

In this section, we discuss a context in which synchronization has turned out to be a relevant phenomenon to explore the relation between dynamical and topological properties of complex networks. Many complex networks in nature are modular, i.e., composed of certain subgraphs with differentiated internal and external connectivity that form communities (Boccaletti et al., 2006; Danon et al., 2005; Newman and Girvan, 2004). This is a limiting situation in which the local structure may greatly affect the dynamics, irrespective of whether or not we deal with homogeneous or heterogeneous networks.

Synchronization processes on modular networks have been recently studied as a mechanism for community de-
tinction (Arenas et al., 2006a,b; Lodato et al., 2007; Oh et al., 2005). Arenas et al. (2006a,b), studied the situation in which a set of identical Kuramoto oscillators (i.e., $\omega_i = \omega$, $\forall i$) with random initial conditions evolves after a transient to the synchronized state.\(^6\) Note that in this case full synchronization is always achieved as this state is the only attractor of the dynamics so that the coupling strength sets the time scale to attain full synchronization: the smaller $\sigma$ is, the longer the time scale. Arenas et al. (2006b) guessed that if high densely interconnected motifs synchronize more easily than those with sparse connections (Moreno et al., 2004), then the synchronization of complex networks with community structure should behave differently at different time and spatial scales. In synthetic modular networks, starting from random initial phases, the highly connected units forming local clusters synchronize first and later on, in a sequential process, larger and larger topological structures do the same up to the point in which complete synchronization is achieved and the whole population of oscillators beat at the same pace. Moreover, this process occurs at different time scales and the dynamical route towards the global attractor reveals the topological structures that represent communities, from the microscale at very early states up to the macroscale at the end of the time evolution. The authors studied the time evolution of pairs of oscillators defining the local order parameter

$$\rho_{ij}(t) = \langle \cos[\theta_i(t) - \theta_j(t)] \rangle ,$$

(36)

averaged over different initial conditions, which measures the correlation between pairs of oscillators. To identify the emergence of compact clusters reflecting communities, a binary dynamic connectivity matrix is introduced such that

$$D_i(T)_{ij} = \begin{cases} 1 & \text{if } \rho_{ij}(t) > T, \\ 0 & \text{if } \rho_{ij}(t) < T, \end{cases}$$

(37)

for a given threshold $T$. Changing the threshold $T$ at fixed times reveals the correlations between the dynamics and the underlying structure, namely, for large enough $T$, one is left with a set of disconnected clusters or communities that are the innermost ones, while for smaller values of $T$ inter-community connections show up. In other words, the inner community levels are the first to become synchronized, subsequently the second level groups, and finally the whole system shows global synchronization. Note that since the function $\rho_{ij}(t)$ is continuous and monotonic, we can define a new matrix $D_T(t)$, that takes into account the time evolution for a fixed threshold. The evolution of this matrix unravels the topological structure of the underlying network at different time scales. In the top panels of Fig. 4 we plot the number of connected components corresponding to the binary connectivity matrix with a fixed threshold as a function of time for networks with two hierarchical levels of communities. There we can notice how this procedure shows the existence of two clear time scales corresponding to the two topological scales.

It is also possible to go one step further and show that the evolution of the system to the global attractor is intimately linked to the whole spectrum of the Laplacian matrix (35). The bottom panels of Fig. 4 show the ranked index of the eigenvalues of $L_{ij}$ versus their inverse. As can be seen, both representations (top and bottom) are qualitatively equivalent, revealing the topological structure of the networks. The only difference is that one comes from a dynamical matrix and the other from the spectrum of a matrix that fully characterizes the topology. Thus, synchronization can be used to unveil topological scales when the architecture of the network is unknown.

The relationship between the eigenvalue spectrum of $L_{ij}$ and the dynamical structures of Fig. 4 can be understood from the linearized dynamics of the Kuramoto model close to the global attractor, which reads (Arenas et al., 2006a,b)

$$\dot{\theta}_i = -\sigma \sum_{j=1}^{N} L_{ij} \theta_j \quad i = 1, \ldots, N .$$

(38)

The solution of Eq. (38) in terms of the normal modes

\(^6\) It is worth stressing here that for this purpose the assumption of $\omega_i = \omega$ can be adopted without loss of generality as it makes the analysis easier. Synchronization of non-identical oscillators also reveals the existence of community structures. See Gómez-Gardeñes et al. (2007b).
\[ \psi_i(t) = \sum_{j=1}^{N} B_{ij} \theta_j = \psi_i(0) e^{-\lambda_i t}, \quad (39) \]

where $B_{ij}$ is the matrix of eigenvectors and $\lambda_i$ the eigenvalues of $L_{ij}$. Sorting the set of Eqs. (39) in descending order of the eigenvalues, one realizes that the solutions $\psi_i(0) e^{-\lambda_i t}$ ($i = N, \ldots, 1$) will approach zero in a hierarchical way; exactly in the same qualitative way that groups of synchronized oscillators appear when the synchronization threshold $T$ is varied. If we fix the threshold and look at different times, the gaps in the eigenvalue spectrum of the Laplacian matrix are associated to different time scales between modes representing different topological scales. Summarizing these findings, the modular structure of the network is revealed by the synchronization process, and on the other side, a more interesting fact comes along, the topological structure of networks imposes characteristic times scales on the synchronization dynamics.

The above results refer to situations in which networks have clearly defined communities. The approach we have proposed enables one to deal with different time and topological scales. In the current literature about community detection (Danon et al., 2005; Newman and Girvan, 2004), the main goal is to maximize the modularity, see Eq. (2). In this case the different algorithms try to identify communities. The betweenness is used as a measure of community coparticipation, since links between nodes that are connected to $i$, and $\alpha$ a time-dependent exponent. The authors use this dynamical evolution to identify communities. The betweenness is used as a measure of community coparticipation, since links between nodes that are in the same community have low betweenness (Girvan and Newman, 2002). Starting from a synchronized state, $\alpha$ is decreased from zero and then the corresponding interaction strength on those links is increasingly enhanced. An additional mechanism that adjusts frequencies between neighboring nodes causes the final state to show partial synchronization among nodes that are in the same community.

6. Synchronization by pacemakers

It is worth mentioning the existence of other different approaches to the synchronization of populations of Kuramoto oscillators. So far we have referred to populations where the oscillators are nearly identical in the sense that they can have slightly different frequencies. Whenever there is a subset of units that play a special role, in the sense that they have substantially different frequencies than the rest in the population or they affect some units but are not affected by any of them, one usually refers to them as pacemakers. The effect of pacemakers has

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7 Here stability (relative) of a given structure is understood to be the ratio between the final and initial times a partition remains synchronized. In terms of the number of connected components in Fig. 4 it corresponds to the length of the plateaus.
been studied in regular networks, as for instance in one-dimensional rings, two-dimensional tori and Cayley trees (Radicchi and Meyer-Ortmanns, 2006). So far, the only approach in a complex topology has been performed by Kori and Mikhailov (2004). There, the authors considered a system of identical units (same frequency) and a singular pacemaker. For an ER network they found that for a large coupling the pacemaker entrains the whole system (all units with the same effective frequency, that of the pacemaker), but the phase distribution is hierarchically organized. Units at the same downward distance from the pacemaker form shells of common phases. As the coupling strength is decreased the entrainment breaks down at a value that depends exponentially on the depth of the network. This result also holds for complex networks, as for instance in WS or SF networks, although the analytical explanation is only valid for ER networks.

### B. Pulse-coupled models

In parallel to the studies described so far, some other approaches to synchronization in networks have invoked models where the interaction between units takes the form of a pulse. In particular, much attention has been devoted to models akin to reproduce the dynamics of neurons, e.g., excitable integrate-and-fire oscillators (IFOs). The basics of an IFO system is as follows. The phase dynamics of any oscillator \( i \) is linear in time \( d\phi_i(t)/dt = 1 \) in absence of external perturbations. However, when the oscillator \( i \) reaches the threshold \( \phi_i(t) = 1 \) it sends a signal (or pulse) to the rest of the oscillators to which it is connected, and relaxes to \( \phi_i(t) = 0 \). The pulse can be considered to propagate instantaneously or with a certain time delay \( \tau \), and when it reaches other oscillators induces a phase jump \( \phi_j \rightarrow \phi_j + \Delta(\phi_j) \). The effects of the topology on the synchronization phenomena emerging in a network of IFOs are at least as rich as those presented in Sect. III.A, although far more difficult to be revealed analytically. The main problem here is that the dynamics presents discontinuities in the variable states that are difficult to deal with. Nevertheless, many insights are recovered from direct simulations and clever mappings of the system. From direct simulations the first insights pointed directly to certain scaling relations between the synchronization time and topological parameters of networks. In ER networks, the scaling relation between the time to needed to achieve complete synchronization \( T \), the number of nodes \( N \), and the number of links \( M \), was found to be

\[
T \sim \left( \frac{M}{N^2} \right)^\alpha \sim \left( \frac{M}{N^2} \right)^{\frac{1}{2}},
\]

with \( \alpha = 1.30(5) \) and \( \beta = 1.50(5) \). Comparing this synchronization process with the same system on a regular square lattice, one realizes that the time needed to synchronize a random network is larger, specially in sparse networks (Guardiola et al., 2000). In between of these two extremal topologies, some WS networks with a rewiring probability \( p \) were studied and were found to expand the synchronization time more than the original regular lattice. However, it was first pointed out that an appropriate normalization of the pulses received by each neuron, rescales the time to very short values. This phenomenon of normalization of the total input signal received by each oscillator has been repeatedly used to homogenize the dynamics in heterogeneous substrates.

Excitable IFOs in SW networks were revisited later by Roxin et al. (2004) to study the possibility of self-sustained activity induced by the topology itself. Considering a unidirectional ring of IFOs with density \( p \) of random long-range directed connections, the authors showed that periodical patterns persist at low values of \( p \), while long-transients of disordered activity patterns are observed for high values of \( p \). Responsible for this behavior is a tradeoff between the average path length and the speed of activity propagation. For low \( p \) configurations, the distances in the networks decrease logarithmically with size, while the superposition of activities is almost the same than in the regular configuration, i.e., the same activity occurs but in a ”smaller” network able to self-sustain its excitation. However for large \( p \), the superposition of activity between excited domains plays also an important role, and then both effects make the synchronized self-sustained activity collapse, leading to disordered patterns.

Lago-Fernández et al. (2000) were the first to investigate a network of nonidentical Hodgkin-Huxley (Hodgkin and Huxley, 1952) elements coupled by excitatory synapses in random, regular, and SW topologies. The parameters of the model neurons were kept to stay below the bifurcation point, until the input arrives and forces the system to undergo a saddle-node bifurcation on a limit cycle. The dynamics of the system ends up into a coherent oscillation or in the activation of asynonymous states. In absence of a detailed analysis of the mechanism that generates coherence, the simulations showed several effects of the topology on the dynamics, the most interesting of which is that achieving synchronization in regular networks takes longer compared to SW, where the existence of short-cuts favors faster synchronization. The results obtained in all cases show that the randomness of the topology has strong effects on the dynamics of these models, in particular the average connectivity is a control parameter for the transition between asynchronous and synchronous states. In Fig. 5 we present a phase diagram for the Hodgkin-Huxley model in SW networks with varying average degree \( k \) and rewiring probability \( p \). A detailed analysis of sparse random networks of general IFOs was exposed by Golomb and Hansel (2000). Their analytic results are in agreement with the previous observations. Very recently, Leyva et al. (2006) analyzed a SW network of non-identical Hodgkin-Huxley units in which some of the couplings could be negative; they surprisingly found that a small fraction of such phase-repulsive links can enhance synchronization.
The works reviewed so far in this subsection are based on the assumption that the coupling is fixed, and that the only source of topological complexity is embedded in the connectivity matrix. Denker et al. (2004) showed that for networks of pulse-coupled oscillators with complex connectivity, coupling heterogeneity induces periodic firing patterns, which replace the state of global synchrony. The coupling heterogeneity has a critical value from which the periodic firing patterns become asynchronous aperiodic states. These results are in agreement with the observations described in previous works. We will come back to this issue in the next section, when dealing with the stability of the synchronized state.

In a slightly different scenario, Vragovic et al. (2006) studied numerically a system of pulse-coupled excitable Bonhoeffer-van der Pol-FitzHugh-Nagumo oscillators (Cartwright, 2000) in WS networks, and found a major influence of the average path length of the network on the degree of synchronization, whereas local properties characterized by clustering and loop coefficients seem to play a minor role. In any case, the authors warn that the results are far from being conclusive, since single characteristics of the network are not easily isolated. We will come back to this issue in the next section, when dealing with the stability of the synchronized state.

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C. Coupled maps

Maps represent simple realizations of dynamical systems exhibiting chaotic behavior. At a first sight they can represent discrete versions of continuous oscillators. Coupled populations of such rather simple dynamical systems have been one of the paradigmatic models to explain the emergence and self-organization in complex systems due to the rich variety of global qualitative behavior they give rise to. From a more practical point of view coupled map systems have found a widespread range of applications, ranging from fluid dynamics and turbulence to stock markets or ecological systems (Pikovsky et al., 2001). Since these systems are nowadays known to have complex topologies, populations of maps coupled through a complex pattern of interactions are natural candidates to study the onset of synchronization as an overall characteristic of the population.

Coupled maps have been widely analyzed in regular lattices, trees and also in global connectivity schemes. The first attempt to consider connectivities in between these extreme cases is due to Gade (1996). He proposed a system formed by units, whose individual dynamics are given by the logistic map, that are connected to a fixed number \( k \) of other units randomly chosen (multiple and self-links are permitted). The evolution rule for the units is

\[
x_i(t + 1) = \frac{1}{k} \sum_j a_{ij} f(x_j(t)) . \tag{41}
\]

A linear stability analysis of this system is performed in terms of the eigenvalues of the matrix \( A \). For the logistic map it is shown that for \( k > 4 \) the maps synchronize. The time the system needs to synchronize decreases with the connectivity \( k \) and also with the system size, although in the latter case the time saturates for large values of the system size. When the connectivity pattern is changed to a modified WS model (by adding long-range short-cuts but not rewiring), Gade and Hu (2000) showed that just a nonzero value of the addition probability is enough to guarantee synchronization in the thermodynamic limit.

In another early attempt to include non-regular topologies in chaotic dynamics, Batista et al. (2003) studied a SW network, in which

\[
\theta_i(t + 1) = (1 - \sigma) f(\theta_i(t)) + \frac{\sigma}{4 + \kappa} \sum_{j=1}^{N} a_{ij} f(\theta_j(t)) , \tag{42}
\]

where \( \kappa \) is the number of shortcuts in the network, and \( \sigma \) the coupling constant. Each unit evolves according to a sine-circle map (Ott, 1993)

\[
f(\theta) = \theta + \Omega - \frac{K}{2\pi} \sin(2\pi \theta) \quad (\text{mod 1}) , \tag{43}
\]

which provides a simple example for describing the dynamics of a phase oscillator perturbed by a time-periodic force. Here \( K \) is a constant related to the external force...
amplitude and $0 \leq \Omega < 1$ is the ratio between the natural oscillator frequency and the forcing frequency. It is observed that synchronization, in terms of a parameter related to the winding number dispersion, is induced by long-range coupling in a system that, in the absence of the shortcuts, does not synchronize.

A slightly different approach was conducted by Jost and Joy (2002), who considered a population of units evolving according to

$$x_i(t+1) = (1-\sigma)f(x_i(t)) + \frac{\sigma}{k_i} \sum_{j \in \Gamma_i} f(x_j(t)).$$

They obtain the stability condition of the synchronized state in terms of the eigenvalues of the normalized Laplacian matrix $\left(\delta_{ij} - a_{ij}/k_i\right)$ and the Lyapunov exponent of the map $f(x)$. Furthermore, they also find a sufficient condition for the system to synchronize independently of the initial conditions, namely

$$(1-\sigma\lambda_2) \sup |f'| < 1,$$

where $\lambda_2$ is the smallest non-zero eigenvalue of the normalized Laplacian matrix. They demonstrate their results for regular connectivity patterns as global coupling and one-dimensional rings with a varying number of nearest neighbors, since in these cases the eigenvalues can be computed analytically. Complexity in the connectivity pattern is introduced in different ways. In these cases one needs numerical estimates of the eigenvalues to compare the synchronization condition (45) with the simulation of the model (44). By using a quadratic map $f(x) = 1-ax^2$ (Ott, 1993) and choosing the free parameter $a$ in a range where different regimes are realized, they find that in a random network the system synchronizes for an arbitrary large number of units, whenever the number of neighbors is larger than some threshold determined by the maximal Lyapunov exponent. This implies a remarkable difference to the one-dimensional case where synchronization is not possible when the number of units is large enough. For a WS model their main finding is that a quite high value of the rewiring probability ($p > 0.8$) is needed to achieve complete synchronization. Finally for BA networks the behavior is comparable to the random ER case.

Following a similar line, Lind et al. (2004) study the behavior of a model where the interaction between the units can be strengthened according to the degree. In this case

$$x_{t+1,i} = (1-\sigma)f(x_{t,i}) + \frac{\sigma}{N_i} \sum_{j \in \Gamma_i} k_i^j f(x_{t,j}),$$

where $N_i = \sum_{j \in \Gamma_i} k_i^j$ is the appropriate normalizing constant. Here the function $f(x)$ is also a quadratic map. The authors study first BA networks. When $\alpha = 0$ the model is equivalent to that discussed previously; in this case they find the existence of a first-order transition between the coherent and the noncoherent phases that depends on both the mean connectivity and the coupling $\sigma$. As varying $a$, the parameter of the quadratic map, they find that these two critical values are related by the power law $\sigma_c \propto k^{-\nu}$. The effect of $\alpha$ being larger than zero is only quantitative, since in this case the transition appears at smaller values of the interaction as compared with the usual case. Additionally, the authors studied two types of deterministic SF networks: a pseudofractal SF network introduced by Dorogovtsev et al. (2002) and the Apollonian network introduced by Andrade et al. (2005). In both cases, there is no coherence when $\alpha = 0$ and $a = 2$. This fact leads the authors to conclude that some degree of randomness that shortens the mean distance between units is needed for achieving a synchronized state, since in these networks the SF nature is not related to a short mean distance. Nevertheless, this situation is avoided if the contributions from the hubs are strengthened, by making $\alpha > 0$.

Another set of papers deals with units that are coupled with some transmission delay. Atay et al. (2004), Masoller and Martí (2005), and Martí et al. (2006), investigate the effect of delays in complex topologies. In the first paper, Atay et al. (2004) propose a model in which all units have the same time delay (in discrete units) with respect to the unit considered:

$$x_i(t+1) = f(x_i(t)) + \frac{\sigma}{k_i} \sum_{j \in \Gamma_i} f(x_j(t - \tau_{ij})) - f(x_i(t)).$$

For a uniform delay $\tau_{ij} = \tau \forall i, j$, they show analytically, and numerically, that the delay facilitates synchronization for general topologies. In any case, this fact confirms the results obtained in Jost and Joy (2002) that ER and SF networks are easier to synchronize than regular or SW ones. Furthermore, one of the implications of connection delays is the possibility of the emergence of new collective phenomena. Masoller and Martí (2005) and Martí et al. (2006) considered uniform distributions of (discrete) time delays. Their main result is that in the presence of random delays the synchronization depends mainly on the average number of links per node, whereas for fixed delays there is also a dependence on other topological characteristics.

In a more general framework (Amritkar and Jalan, 2003; Jalan and Amritkar, 2003; Jalan et al., 2005), the following problem is considered

$$x_i(t+1) = (1-\sigma)f(x_i(t)) + \frac{1}{k_i} \sum_{j \in \Gamma_i} g(x_j(t)).$$

For a logistic map $g(x) = \mu x (1-x)$ (although analyses on other maps as the circle map and the tent map have also been performed) the authors show a phase diagram in which the different stationary configurations are obtained as a function of the coupling strength $\sigma$ and the parameter of the map $\mu$. The stationary configurations are classified in the following way: turbulence (all units behave chaotically), partially ordered states (few synchronized clusters with some isolated nodes), ordered
states (two or more synchronized clusters with no isolated nodes), coherent states (nodes form a single synchronized cluster), and variable states (nodes form different states depending strongly on initial conditions). The critical value of the coupling above where phase synchronized clusters are observed depends on the type of network and the coupling function. As a remarkable point, it is found that two different mechanisms of cluster formation (partial synchronization) can be distinguished: self-organized and driven clusters. In the first case, the nodes of a cluster get synchronized because of intrachannel coupling. In the latter case, however, synchronization is due to intercluster coupling; now the nodes of one cluster are driven by those of the others. For a linear coupling function \( g(x) = x \), self-organization of clusters dominates at weak coupling; when increasing the coupling strength, a transition to driven-type clusters, almost independent of the type of network, appears. However, for a nonlinear coupling function the driven type dominates for weak coupling, and only networks with a tree-like structure show some cluster formation for strong coupling.

Finally, it is worth mentioning the very recent work by Levnajic and Tadic (2007), in which the authors consider a SF tree (preferential-attachment growing network with one link per node) of two-dimensional standard maps:

\[
\begin{align*}
x'(t+1) &= x + y + \mu \sin(2\pi x) \pmod{1} \\
y'(t+1) &= y + \mu \sin(2\pi x).
\end{align*}
\]

(49)

The nodes are coupled through the angle coordinate \( (x) \) so that the complete time-step of the node \( i \) is

\[
\begin{align*}
x_i(t+1) &= (1-\varepsilon)x'_i(t) + \sum_{j \in F_i} \epsilon_j x_j(t) - x'_i(t) \\
y_i(t+1) &= (1-\varepsilon)y'_i(t).
\end{align*}
\]

(50)

Here, ‘(’ denotes the next iteration of the (uncoupled) standard map (49) and \( t \) denotes the global discrete time. The update of each node is the sum of a contribution given by the updates of the nodes, the ‘(‘ part, plus a coupling contribution given by the sum of differences, taking into account a delay in the coupling from the neighbors. By keeping \( \mu = 0.9 \) such that the individual dynamics is in the highly chaotic regime, the authors analyze the dependence on the interaction strength \( \sigma \). For small values of the coupling the motion of the individual units is still chaotic, but the trajectories are contained in a bounded region. With further increments of the coupling, the units follow periodic motions which are highly synchronized. In this case, however, synchronization takes place in clusters, each cluster having a common value of the band center around which the periodic motion occurs, and center values appear in a discrete set of possible values. These clusters form patterns of dynamical regularity affecting mainly nodes at distances 2, 3, and 4, as shown in Fig. 6(left). In fact, the histograms of distances between nodes along the tree and between nodes belonging to the same synchronized cluster have different statistical weights only for these values of the distances.

All previously discussed chaotic models are discrete time maps, which are appropriate discrete versions of chaotic oscillators. Nevertheless, we also notice a couple of works dealing with time continuous maps. Li and Chen (2004) analyze a WS network of Rössler oscillators, where the parameters are chosen to ensure that the system generates chaotic dynamics. The basic observation is that the network synchronizes when the coupling strength is increased, as expected. Another interesting result is that the mean phase difference among the chaotic oscillators decreases with the increasing of the probability of adding long-range random short-cuts. Along the same line, Yook and Meyer-Ortmanns (2006) consider a system of Rössler oscillators on BA networks. The only tuning parameter of the BA networks is \( m \), the number of links that a newly added node has. For \( m = 1 \) (SF trees) there is no synchronized state for a large number of oscillators. Increasing \( m \) synchronization is favored. The topological effect of increasing \( m \) is to create loops, but it is shown that this is not the only fact that improves synchronization.

Finally, a different and interesting proposal was made by Belykh et al. (2004a) where a fixed 1-d connectivity pattern is complemented by a set of switching long-range connections. In this case, it is proven that interactions between nodes that are only sporadic and of short duration are very efficient for achieving synchronization.

As a summary, we can say that most of the works deal with particular models of coupled maps (logistic maps, sine-circle maps, quadratic maps, ...). Thus, it is possible, to obtain in some cases not only the conditions of local stability of the completely synchronized state but the conditions for the synchronization independently of the initial conditions. In general, the addition of short-cuts to regular lattices improves synchronization. There are even some cases, for which synchronization is only attainable when a small fraction of randomness is add to the system. On the contrary, in the next section we will discuss the linear stability of the synchronized state for general dynamical systems.
IV. STABILITY OF THE SYNCHRONIZED STATE IN COMPLEX NETWORKS

In the previous section we have reviewed the synchronization of various types of oscillators on complex networks. Another line of research on synchronization in complex networks, developed in parallel to the studies of synchronization in networks of phase oscillators, is the investigation of the stability of the completely synchronized state of populations of identical oscillators. The seminal work by Barahona and Pecora (2002) initiated this research line by analyzing the stability of synchronization in SW networks using the Master Stability Function (MSF). The framework of MSF was developed earlier for the study of synchronization of identical oscillators on regular or other simple network configurations (Fink et al., 2000; Pecora and Carroll, 1998). The extension of the framework to complex topologies is natural and important, because it relates the stability of the fully synchronized state to the spectral properties of the underlying structure. It provides with an objective criterion to characterize the stability of the global synchronization state, from now on called synchronizability of networks independently of the particularities of the oscillators. Relevant insights about the structure-dynamics relationship have been obtained using this technique.

In this section, we review the MSF formalism and the main results obtained so far. Note that the MSF approach assesses the linear stability of the completely synchronized state, which is a necessary, but not a sufficient condition for synchronization.

A. Master Stability Function formalism

To introduce the MSF formalism, we start with an arbitrary connected network of coupled oscillators. The assumption here for the stability analysis of synchronization is that all the oscillators are identical, represented by the state vector $\mathbf{x}$ in an $m$-dimensional space. The equation of motion is described by the general form

$$\dot{\mathbf{x}} = \mathbf{F}(\mathbf{x}).$$

For simplicity, we consider time-continuous systems. However, the formalism applies also to time-discrete maps. We will also assume an identical output function $\mathbf{H}(\mathbf{x})$ for all the oscillators, which generates the signal from the state $\mathbf{x}$ and sends it to other oscillators in the networks. In this representation, $\mathbf{H}$ is a vector function of dimension $m$. For example, for the 3-dimensional system $\mathbf{x} = (x, y, z)$, we can take $\mathbf{H}(\mathbf{x}) = (x, 0, 0)$, which means that the oscillators are coupled only through the component $x$. $\mathbf{H}(\mathbf{x})$ can be any linear or nonlinear mapping of the state vector $\mathbf{x}$. The $N$ oscillators, $i = 1, \ldots, N$, are coupled in a network specified by the adjacency matrix $A = (a_{ij})$. We have

$$\dot{\mathbf{x}}_i = \mathbf{F}(\mathbf{x}_i) + \sigma \sum_{j=1}^{N} a_{ij} w_{ij} [\mathbf{H}(\mathbf{x}_j) - \mathbf{H}(\mathbf{x}_i)]$$

being $w_{ij}$ the connection weights, i.e., the network is, in general, weighted. The coupling matrix $G$ is $G_{ij} = -a_{ij} w_{ij}$ if $i \neq j$ and $G_{ii} = \sum_{j=1}^{N} a_{ij} w_{ij}$. When the coupling strength is uniform for all the connections ($w_{ij} = 1$), the network is unweighted, and the coupling matrix $G$ is just the usual Laplacian matrix $L$. By definition, the coupling matrix $G$ has zero row-sum. Thus there exists a completely synchronized state in this network of identical oscillators, i.e.,

$$\mathbf{x}_1(t) = \mathbf{x}_2(t) = \ldots = \mathbf{x}_N(t) = \mathbf{s}(t),$$

which is a solution of Eq. (53). In this synchronized state, $\mathbf{s}(t)$ also approaches the solution of Eq. (51), i.e., $\mathbf{s} = \mathbf{F}(\mathbf{s})$. This subspace in the state space of Eq. (53), where all the oscillators evolve synchronously on the same solution of the isolated oscillator $\mathbf{F}$, is called the synchronization manifold.

1. Linear Stability and Master Stability Function

When all the oscillators are initially set at the synchronized manifold, they will remain synchronized. Now the crucial question is whether the synchronization manifold is stable in the presence of small perturbations $\delta \mathbf{x}_i$. To assess the stability, we need to know whether the perturbations grow or decay in time. The linear evolution of small $\delta \mathbf{x}_i$ can be obtained by setting $\mathbf{x}_i(t) = \mathbf{s}(t) + \delta \mathbf{x}_i(t)$ in Eq. (53), and expanding the functions $\mathbf{F}$ and $\mathbf{H}$ to first order in a Taylor series, i.e., $\mathbf{F}(\mathbf{x}_i) = \mathbf{F}(\mathbf{s}) + D\mathbf{F}(\mathbf{s})\delta \mathbf{x}_i$ and $\mathbf{H}(\mathbf{x}_i) = \mathbf{H}(\mathbf{s}) + D\mathbf{H}(\mathbf{s})\delta \mathbf{x}_i$. Here $D\mathbf{F}(\mathbf{s})$ and $D\mathbf{H}(\mathbf{s})$ are the Jacobian matrices of $\mathbf{F}$ and $\mathbf{H}$ on $\mathbf{s}$, respectively. This expansion results in the following linear variational equations for $\delta \mathbf{x}_i$,

$$\delta \dot{\mathbf{x}}_i = D\mathbf{F}(\mathbf{s})\delta \mathbf{x}_i - \sigma D\mathbf{H}(\mathbf{s}) \sum_{j=1}^{N} G_{ij} \delta \mathbf{x}_j.$$
row-sum of $G$. So the first eigenmode $\xi_1 = D\mathbf{F}(s)\xi_1$ corresponds to the perturbation parallel to the synchronization manifold. The other $N-1$ eigenmodes are transverse to the synchronization manifold and should be damped out to have a stable synchronization manifold.

In the following we assume that all the eigenvalues are real. This property is analytically proven for symmetric $G$ but it can also be true for non-symmetric cases, as will be discussed later. Furthermore, the non-negative nature of $G$ ensures that the eigenvalues are also non-negative. Then they can be ordered as

$$0 = \lambda_1 \leq \lambda_2 \cdots \leq \lambda_N. \quad (57)$$

In a connected network of $N$ oscillators, $\lambda_2$ is the minimal and $\lambda_N$ is the maximal non-zero eigenvalue.

Importantly, we observe that all the variational equations in Eq. (56) have the same form

$$\dot{\xi} = [D\mathbf{F}(s) - \alpha D\mathbf{H}(s)]\xi. \quad (58)$$

They only differ by the parameter $\alpha_1 = \sigma \lambda_1$. Now if we know the stability of the solution $\xi = 0$ for any reasonable value of $\alpha$, then we can infer the stability for any eigenmode with $\alpha_1 = \sigma \lambda_1$. To assess the stability of this master variational equation (58), we calculate its largest Lyapunov exponent $\lambda_{\text{max}}$ as a function of $\alpha$, the resulting function is the master stability function. The evolution of small $\xi$ is then described on average as $||\xi(t)|| \sim \exp(\lambda_{\text{max}}(\alpha)t)$, and the mode is stable with $||\xi|| \rightarrow 0$ if $\lambda_{\text{max}}(\alpha) < 0$.

So far, we have assumed that the eigenvalues $\lambda_1$ are real, and the problem is to compute the MSF $\lambda_{\text{max}}(\alpha)$ for real values of $\alpha$. In general, the coupling matrix $G$ is asymmetric, and the eigenvalues $\lambda_1$ can be complex numbers. In this case one needs to evaluate the MSF in the complex plane $(\alpha_R, \alpha_I)$. It has been found that for many oscillator types $\lambda_{\text{max}}(\alpha) < 0$ in a finite region of this plane (Fink et al., 2000; Pecora and Carroll, 1998).

The reader may find the framework of master stability function abstract. Here we present it in a physically intuitive manner with the help of a schematic diagram. For this, we restrict ourself to the case where $\alpha$ is real. Let us consider first two coupled oscillators. The first one is autonomous and evolving along the trajectory $s(t)$, and the second one is driven by the first one with a coupling strength $\alpha$, as depicted in Fig. 7(a). The dynamical equations read,

$$\dot{s} = \mathbf{F}(s), \quad (59)$$

$$\dot{x} = \mathbf{F}(x) + \alpha [\mathbf{H}(s) - \mathbf{H}(x)]. \quad (60)$$

Then immediately, we obtain the linear variational equation for the synchronization difference $\xi(t) = x(t) - s(t)$ as in Eq. (58). This means that the MSF describes the stability of the state in which the two oscillators are synchronized.

In a similar spirit, the equations for the $N-1$ transverse eigenmodes in Eq. (56) can be graphically represented as $N-1$ oscillators driven by a common autonomous oscillator $\dot{s} = \mathbf{F}(s)$, and the corresponding coupling strengths are $\sigma \lambda_1$, as depicted in Fig. 7(b). In this representation, the mode decomposition decouples the complex network connections into many pairs of oscillators and the stability thus can be understood from that of the two coupled oscillators, the MSF. The complete synchronization state $s$ is stable when all the $N-1$ oscillators in Fig. 7(b) are synchronized by the common forcing signal $s$. This picture will be useful later on in this section.

2. Measures of synchronizability

In general, the MSFs verify that $\lambda_{\text{max}}(\alpha) < 0$ within a finite interval $\alpha_1 < \alpha < \alpha_2$ (Fig. 8). This means that for two coupled oscillators, as those in Fig. 7(a), the driven oscillator will be synchronized, i.e., $x(t) = s(t)$, if the coupling strength $\alpha \in (\alpha_1, \alpha_2)$. For the network, all the $N-1$ oscillators in Fig. 7(b) should be synchronized by the common forcing $s(t)$, requiring $\sigma \lambda_l \in (\alpha_1, \alpha_2)$ for $2 \leq l \leq N$. Explicitly, the following condition is necessary for the stability of the synchronization state of the network,

$$\alpha_1 < \sigma \lambda_2 \leq \sigma \lambda_3 \leq \cdots \leq \sigma \lambda_N < \alpha_2. \quad (61)$$
This condition can only be fulfilled for some values of $\sigma$ when the eigenratio $R$ satisfies the following relation

$$R \equiv \frac{\lambda N}{\lambda_2} < \frac{\alpha_2}{\alpha_1}. \quad (62)$$

Therefore, we conclude that it is impossible to completely synchronize the network if $R > \alpha_2/\alpha_1$, since there is no $\sigma$ for which the fully synchronized state is linearly stable. On the contrary, if $R < \alpha_2/\alpha_1$ the synchronous state is stable for $\alpha_1/\lambda_2 < \sigma < \alpha_2/\lambda_N$.

Note that the eigenratio $R$ depends only on the network structure, as defined by $G$. If $R$ is small, the condition in Eq. (62) is, in general, easier to satisfy. From this, it follows that the smaller the eigenratio $R$ the more synchronizable the network and vice versa (Barahona and Pecora, 2002), and we can characterize the synchronizability of the networks with $R$, without referring to specific oscillators.

In the case that $\alpha_2$ is finite, the MSF is bounded. This is what happens always for time discrete maps and in occasionally to time-continuous systems. In other cases, the MSF is unbounded, i.e., $\alpha_2 = \infty$, and the synchronized state is stable when the overall coupling strength $\sigma > \alpha_1/\lambda_2$. For example, this is the case for time-continuous systems and $H(x) = x$, i.e., the oscillators are linearly coupled through all the $m$ corresponding components. In these cases, a larger $\lambda_2$ means a smaller coupling strength $\sigma$ for synchronization, and thus corresponds to a better synchronizability of the networks. Both $\lambda_2$ and $R$ are adopted as measures of synchronizability in the literature. Moreover, as previously discussed in Sect. III.A.4, $\lambda_2$ plays a special role in the time needed to achieve complete synchronization (Almendral and Díaz-Guilera, 2007).

The key advantage of the MSF framework is that it provides an objective criterion for the synchronizability of complex networks without the need to refer to specific oscillators. The drawback is that it only informs about the dynamics towards synchronization from small perturbations of the synchronization manifold. The study of synchronizability is then converted into the investigation of the eigenvalues of the coupling matrix $G$ of the networks. Note that although the more appropriate approach seems to be that of following the general framework of graph theory to investigate the spectral properties of networks, many authors have tried to relate one statistical properties of networks with synchronizability, sometimes misinterpreting the results due to an incorrect approach.

Of course, natural questions are: What is the synchronizability of different types of complex networks? Which structural properties are related to or control the synchronizability? In the following section, we will summarize results on the synchronizability of typical network models and then describe the relationship between graph theoretical measures and the synchronizability of complex networks. Further insights will be obtained with the help of graph theory to get the bounds that constrain the eigenvalues of the networks.

3. Synchronizability of typical network models

We will present the main findings related to synchronizability in the most common classes of complex networks found in the literature: Regular, SW, ER, and SF networks.

**Regular networks** For a long time, synchronization of chaotic oscillators has been studied on regular network configurations (Fink et al., 2000; Pecora and Carroll, 1998). A typical example of a regular network is a cycle (or ring) of $N$ nodes each coupled to its $2k$ nearest neighbors with a total of $Nk$ links. The eigenvalues of the coupling matrix are (Barahona and Pecora, 2002; Monasson, 1999)

$$\lambda_l = 2k - 2 \sum_{j=1}^{k} \cos \left( \frac{2\pi(l-1)j}{N} \right). \quad (63)$$

By a series expansion, we obtain

$$\lambda_2 \simeq \frac{2\pi^2k(k+1)(2k+1)}{3N^2}, \lambda_N = (2k+1)(1+2/3\pi). \quad (64)$$

![FIG. 9 (color online) Decay of the eigenratio $R$ in a $N = 100$ lattice as a fraction $f$ of the $N(N-1)/2$ possible edges are added following purely deterministic, semirandom, and purely random schemes. Networks become synchronizable below the dashed line (betw {\beta = \alpha_2/\alpha_1, where \alpha_1 and \alpha_2 are from Fig. 8). The squares (numerical) and the solid line [analytic Eq. (63)] show the eigenratio decay of regular networks through the deterministic addition of short-range connections. The dotted line corresponds to purely random graphs [Eq. (68)], which become almost surely disconnected and unsynchronizable at $f \simeq 0.0843$. The semirandom approach (dots, shown for ranges $k = 1, 2, 4, 6, 10, 14$) is more efficient in producing synchronization when $k < \ln N$. From Barahona and Pecora (2002).]
and the eigenratio $R$ for $k \ll N$ can be approximated as

$$R \approx \frac{(3\pi + 2)N^2}{2\pi^2k(k + 1)}$$

(65)

Regular networks have a poor synchronizability, since $R$ increases rapidly with the system size $N$. In fact, $\lambda_2 \sim 1/N^2$, and it approaches zero quickly when increasing $N$, thus making synchronization effectively impossible for both bounded and unbounded MSFs given chaotic oscillators with $\alpha_1 > 0$. In such systems with local coupling we usually have complicated pattern formation (waves) instead of a stable spatially homogeneous (completely synchronized) state (Matías and Güeméz, 1998). For fixed $N$, adding connections to generate regular networks with higher range $k$ improves the synchronizability, since $\lambda_2$ increases approximately as $\lambda_2 \sim k^3$ and the eigenratio $R$ decreases as $R \sim 1/k^2$ for $k \ll N$.

**SW networks** In the work by Barahona and Pecora (2002), SW networks are obtained by adding NS links at random between the $N(N - 2k - 1)/2$ remaining pairs in the regular network, so that the average number of shortcuts per node is $S$. As shown in Fig. 9, adding a small fraction of such random connections reduces the eigenratio $R$ so that the synchronizability is improved significantly.

The eigenvalues of the SW networks have been obtained (Barahona and Pecora, 2002) through a perturbation analysis of the SW Laplacian $L = L^0 + L^r$. Here $L^0$ is the deterministic Laplacian of the regular networks and $L^r$ the Laplacian formed by the random shortcuts. In the stochastic Laplacian matrix $L^r$ (symmetric, zero row-sum), any of the remaining entries $N(N - 2k - 1)$ of $L^0$ takes the value 1 with probability $p_s = 2S/(N - 2k - 1)$ and the value 0 with probability $(1 - p_s)$. For $p_s \ll 1$, and $N^{1/3} < k \ll N$, the perturbations of the eigenvalues are

$$\varepsilon \lambda_2^{(1)} \approx Np_s - \sqrt{3\pi p_s}/4, \quad \varepsilon \lambda_N^{(1)} \approx Np_s + \sqrt{3\pi p_s}/4.$$

(66)

The extreme eigenvalues are

$$\lambda_2 = \lambda_2^{(0)} + \varepsilon \lambda_2^{(1)}, \quad \lambda_N = \lambda_N^{(0)} + \varepsilon \lambda_N^{(1)},$$

(67)

where $\lambda_2^{(0)}$ and $\lambda_N^{(0)}$ are the eigenvalues of the regular networks ($L^0$) as in Eq. (64). From this analysis, it follows that for a fixed small value of $S$, the minimal non-zero eigenvalue $\lambda_2$ is driven away from $\lambda_2^{(0)} \approx 0$ to $\varepsilon \lambda_2^{(1)} \approx Np_s \approx 2S$ for any SW network with large $N$ and $N^{1/3} < k \ll N$. This means that the synchronizability is mainly determined by the average number of shortcuts per node $S$.

In SW networks, the variance of the degree distribution raises as the regular network is rewired with an increasing probability $p$ (Hong et al., 2004) or when more shortcuts are added, see Fig. 10(b). This process results in an improvement of the synchronizability (the eigenvalue ratio, $R$, is reduced), as illustrated in Fig. 9 and Fig. 10(a). This is because $\lambda_2$ increases proportionally to the number of shortcuts per node, i.e., $\lambda_2 \approx 2S = kp$.

**Random networks** In purely random graphs, in which a fraction $f$ of the $N(N - 1)/2$ possible links is established at random, the eigenratio is a function of $f$ and $N$ (Barahona and Pecora, 2002). It reads

$$R \approx \frac{Nf - \sqrt{2f(1 - f)}\ln N}{Nf - \sqrt{2f(1 - f)}\ln N}.$$  

(68)

Note that the networks are synchronizable only when $f \geq 1/2 \ln N/(N + 2 \ln N)$, where it is almost sure that the networks are connected (Bollobás, 2001). If this condition is verified the synchronizability is improved with increasing $f$, as seen in Fig. 9.

Note that for small $f \lesssim \ln N/N$, the purely random networks are almost surely disconnected and thus non-synchronizable. On the contrary, the regular backbone of nearest connections with $k \geq 1$ can already make the semirandom SW networks connected regardless of $N$ and thus synchronizable as a whole. In this sense, semirandom SW networks turn out to be more superior to the purely random networks in terms of synchronizability, as can be seen from Fig. 9 for small $k$, $k \lesssim \ln N$ ($k = 1 \sim 4$ in networks with $N = 100$). The improvement is even more pronounced for larger $N$.

**SF networks** Here we discuss the synchronizability of SF networks for different values of the degree distribution exponent $\gamma$. To this end, we consider a random model introduced by Newman et al. (2001) to construct SF networks. The algorithm works as follows, first, a degree $k_i$ is assigned to each node $i$ according to the probability distribution $P(k) \sim k^{-\gamma}$ and $k \geq k_{\min}$. The network is generated by randomly connecting the nodes so that node $i$ has exactly the prescribed $k_i$ links to other nodes, prohibiting self- and repeated links.

The dependence of the eigenvalues $\lambda_2$ and $\lambda_N$ and the eigenratio $R$ on $\gamma$ and $N$ are shown in Fig. 11. We observe that $\lambda_2$ has no noticeable dependence on $\gamma$ and $N$. 

![FIG. 10 Synchronizability (eigenratio $R$, (a)) and heterogeneity of degrees (variance $\sigma_k^2$ of the degree distribution (b)) of SW networks as a function of the rewiring probability $p$. Inset of (b): average path length $\ell$ vs $p$. The network has a size $N = 2000$ and the range $k = 3$. From Hong et al. (2004).](image-url)
However, $\lambda_N$ becomes larger as the degree heterogeneity is increased. So the changes of the eigenratio $R$ follow the trend of $\lambda_N$ closely. This result is somehow expected given that the largest eigenvalue $\lambda_N$ is intimately related to the degree of hubs, and this is the essential fingerprint of SF networks with different exponents $\gamma$. On the other hand, $\lambda_2$ increases with $k_{\text{min}}$, and $R$ is larger at smaller $k_{\text{min}}$ for the same $\gamma$ and $N$. The variation of $R$ as a function of $\gamma$ was reported by Nishikawa et al. (2003). The dependence of the synchronizability on $k_{\text{min}}$ and $N$ in this random SF network model turns out to be very similar in the BA growing model, as shown by Pecora and Barahona (2005).

In an early paper (Wang and Chen, 2002) that studied robustness and fragility of synchronization of SF networks, it was reported that $\lambda_2$ is a constant almost unrelated with $k_{\text{min}}$ ($k_{\text{min}} = 3, 5, 7, 9, 11$). This result by Wang and Chen (2002) is inconsistent with the observation in Fig. 11, with the work by Pecora and Barahona (2005), and with graph theoretical analysis by Wu (2003), which we will discuss in more detail later on. Thus the observation of robustness and fragility of synchronizability by Wang and Chen (2002) (changes of $\lambda_2$ due to random or deliberate attack of the nodes) should be taken cautiously, and a detailed reexamination of this issue is mandatory before assessing conclusions.

4. Synchronizability and structural characteristics of networks

The relationship between structural characteristics of networks and synchronizability has been explored intensively in the literature, mainly based on numerical experiments on various network models. The observations, which are summarized in what follows, are quite confusing. The main problem is that many works have made a naive use of complex network models to assess synchronizability. Very often network models do not allow us to isolate one structural characteristic while keeping the other properties fixed. For this reason many results have been misinterpreted. As we will see in the next section, a more objective graph theoretical analysis sheds some light to the whole problem.

Synchronizability dependence on $\ell$ The average shortest path length $\ell$ is a property of the network closely related to the efficiency of information processing. Most real-world complex networks are characterized by a small $\ell \lesssim \ln N$ (Dorogovtsev et al., 2007). Indeed, it has been conjectured and rationalized that in biological neuronal networks, $\ell$ has been minimized by evolution (Kaiser and Hilgetag, 2006; Stephan et al., 2000). Generally speaking, $\ell$ is lower in SF networks than in ER networks due to the presence of hubs (Cohen and Havlin, 2003), and $\ell$ is lower in SW networks than in regular lattices due to the presence of shortcuts.

Watts and Strogatz (1998) suggested that the decrease in the distance in the WS network would lead to more efficient coupling and thus enhanced synchronization of the oscillators. Investigation of phase oscillators (Hong et al., 2002) or circle maps (Batista et al., 2003) on WS networks has shown that when more and more shortcuts are created at larger rewiring probability $p$, the transition to the synchronization regime becomes easier. On the other hand, the synchronizability of identical oscillators follows the same trend of $\ell$, in networks with fixed $N$ and $k$, as $p$ is increased (Fig. 10). A similar type of behavior is observed if shortcuts are added to the regular networks (Fig. 9). However, more detailed analyses of various network models have shown that there is no direct relationship between $\ell$ and the synchronizability of the networks. The reason is that the transition to the small-world regime occurs at a value of the rewiring probability for which there is no significant effect on $\lambda_2$.

In fact, in WS networks, $\ell$ is a function of the network size $N$, the degree of the nodes in the original regular network, $k$, and the randomness parameter $p$ (Newman et al., 2000)

$$\ell(N, k, p) \sim \frac{N}{k} f(pkN),$$

(69)

where $f(u)$ is a universal scaling function, $f(1) = \text{const}$ if $u \ll 1$ and $f(u) = \ln(u)/u$ if $u \gg 1$. From this result, it turns out that $\ell$ begins to decrease with $p$, and consequently the SW behavior emerges, for $p \gtrsim p_{\text{SW}} = 1/Nk$. At $p = p_{\text{SW}}$ the average number of shortcuts per node is $S \sim 1/N$, and then $\lambda_2 \sim 1/N$ as well. This shows that at this point the synchronizability is not enhanced by the rewiring. To achieve such an enhancement, the density of shortcuts has to be independent of $N$, which happens for $p \gtrsim p_{\text{sync}} = 1/k$, that is deep in the SW regime. In other words, in the intermediate region $p_{\text{SW}} < p < p_{\text{sync}}$, $\ell$ decreases while the synchronizability of the system remains roughly the same.

Barahona and Pecora (2002) showed the existence of two thresholds, one for the small-world transition and the other for the enhancement of synchronizability, in SW networks using a more rigorous analysis. They obtained that the SW regime starts at $S_\ell \sim 1/N$ whereas...
the threshold beyond which the synchronizability is improved goes like $S_{\text{sync}} \propto k$. This means that when $k$ is large, $\lambda_2(0)$ of the underlying regular network contributes significantly to the synchronizability, and then it can be enhanced without additional shortcuts. This is manifested in Fig. 12 by the fast decrease of $S_{\text{sync}}$ when $k$ approaches the critical value $k^0_{\text{sync}}$. On the other hand, at low $k$, the approximation in Eq. (66) is not valid. However, the results in Fig. 12 show that $S_{\text{sync}} \in [0.3, 1]$ depends on $k$, but not noticeably on $N$.

In summary, it is not very meaningful to compare the synchronizability of two SW networks (with different $N$, $k$ or $p$) considering only $\ell$. In other words, it is possible to have better synchronizability for networks (e.g., fixed $N$) with larger $\ell$ in some region of $p$ and $k$. For example, along the line $pk = \text{const}$, $R \sim 1/p$ decreases while $\ell \sim p$.

The relationship between $\ell$ and the synchronizability of the system was also scrutinized for SF networks by Nishikawa et al. (2003). As seen in Fig. 13, for random SF networks, $\ell$ decreases when the degree distribution becomes more heterogeneous by lowering $\gamma$. However, the network becomes less synchronizable, since $R$ increases. Note that in these simulations, the mean degree of the network $\langle k \rangle = k_{\text{min}} \frac{\gamma-1}{\gamma-2}$ also changes with $\gamma$, as well as the standard deviation of the degree distribution. As will be clarified later on, in this case, the synchronizability is controlled by the heterogeneity of the degree distribution, being $R \sim k_{\text{max}}/k_{\text{min}}$ if $k_{\text{min}}$ is large enough (Zhou, Mottet and Kurths, 2006). Similar to the SW networks with different range $k$, where synchronizability is not simply controlled by $\ell$, here it is also not surprising to observe that synchronizability can be reduced when $\ell$ becomes smaller (Nishikawa et al., 2003). Besides, it should be possible to observe the opposite situation in which $R$ decreases with $\ell$ for suitable combinations of parameters $N$, $\gamma$ and $k_{\text{min}}$. Therefore, the conclusion is that the synchronizability of complex networks cannot be assessed solely on the average shortest path length $\ell$. 

**Synchronizability dependence on betweenness centrality**

Nishikawa et al. (2003) argued heuristically that this apparently surprising behavior (smaller $\ell$, less synchronizability in SF networks) is due to the fact that a few central oscillators interacting with a large number of other oscillators tend to become overloaded. When too many independent signals with different phases and frequencies are going through a node at the same time, they can cancel out each other, resulting in no effective communication between oscillators. Thus the authors were motivated to examine the influence of the load (betweenness) of the nodes. Nishikawa et al. (2003) showed (see the inset of Fig. 13 (b)) that the synchronizability follows the same trend as the maximum load $b_{\text{max}}$ (normalized by the total load of the network). However, the load of a node in SF networks is closely related to the degree (Barthélémy, 2004; Goh et al., 2001), i.e., nodes with large degrees or links connecting nodes with large degrees have, on average, a large load. The correlation between reduced synchronizability and heterogeneous load has also been observed in a variant of the WS network model by adding $m$ shortcuts to the network from a randomly selected node to one out of the $n_c$ center nodes (Nishikawa et al., 2003). In this case, when $n_c$ is small, the $m$ shortcuts are connected to a few hubs, and the degree becomes more heterogeneous and the maximum load $b_{\text{max}}$ increases, while...
the synchronizability decreases. As before, in these two examples, it is not very clear whether the change of synchronizability is mainly influenced by the degree heterogeneity or by the load itself, because these two properties are closely related.

In the original WS network (Watts and Strogatz, 1998), the maximum load \( b_{\text{max}} \) decreases when the degree distribution becomes more heterogeneous as \( p \) is increased. Based on this observation, it was claimed that more homogeneous load predicts better synchronizability on complex networks (Hong et al., 2004). However, a direct relationship between load and synchronizability can not be clearly established. In fact, Zhao, Zhou, Wang, Yan, Yang and Bai (2006) showed an example where the network displays improved synchronizability while the load becomes more heterogeneous when an original random SF is rewired to obtain nontrivial clustering.

The heuristic argument by Nishikawa et al. (2003) is not clearly justified. If the picture described is correct, i.e., signals can cancel out each other at the central nodes, resulting in no effective communication, then the synchronizability should also be reduced significantly for unbounded MSF where the synchronizability is only controlled by \( \lambda_2 \). However, as seen in Fig. 11, \( \lambda_2 \) in SF networks is mainly determined by the minimal degree \( k_{\text{min}} \), without a noticeable dependence on the degree distribution exponent or load. In fact, several further investigations have shown that the highly connected oscillators synchronize faster among them and form synchronization clusters (Gómez-Gardeñes et al., 2007a; Lee, 2005; Zhou and Kurths, 2006a,b), which is in contrast with the argument provided by Nishikawa et al. (2003). A different viewpoint, based on a mean field approximation (Motter et al., 2005a; Zhou and Kurths, 2006a,b), leads to a deeper understanding of this issue, as will be discussed later.

**Synchronizability dependence on the clustering coefficient** Motter et al. (2005b) pointed out that, in general, the eigenratio \( R \) increases with increasing clustering in a modified version of the BA model (Steyvers and Tenenbaum, 2005). Unlike the original one, in this model, motivated by the evolution of language, a new node is first linked to an existing node according to the preferential attachment rule, and also linked to the neighbors of this target node. Thus this model displays nontrivial clustering while keeping the same SF degree distribution. Besides, the eigenratio \( R \) is larger than that of the BA model.

A recent work demonstrated that for both SW and SF networks, large values of clustering hinders global synchronization of phase oscillators, since the network splits into dynamical clusters that oscillate at different frequencies (Gómez-Gardeñes and Moreno, 2007; McGraw and Menzinger, 2005). Zhao, Zhou, Wang, Yan, Yang and Bai (2006), using the rewiring scheme proposed by Kim (2004), that changes the clustering but keeps the degree sequence, showed that the eigenratio \( R \) also increases with \( C \), for both SW and SF networks. This indicates that synchronizability is reduced when clustering \( C \) increases.

Note that in the above investigations different structural properties change at the same time (Barabási and Pecora, 2002; Hong et al., 2004; Nishikawa et al., 2003) when the parameters characterizing the original networks are modified. Once again, it is difficult to draw conclusions about the relationship between one single statistical descriptor of the network and its synchronizability. Zhao, Zhou, Wang, Yan, Yang and Bai (2006) paid special attention to this problem and showed that in the rewiring scheme of Kim (2004), \( \ell \) is correlated with the clustering coefficient \( C \) (Fig. 14).

**Synchronizability dependence on degree correlations** In many real-world networks, the degree of a node is often correlated with the degree of the neighboring nodes. Correlated networks show **assortative** (disassortative) mixing when high degree nodes are mostly attached to nodes with high (low) degree (Newman, 2002). In practice, the degree-degree correlation of a network can be calculated as the Pearson correlation coefficient between degrees \( (j_i, k_i) \) of the nodes at the ends of the \( l \)th link, i.e.,

\[
\rho_k = \frac{(j_i k_i) - \langle k_i \rangle^2}{\langle k_i^2 \rangle - \langle k_i \rangle^2},
\]

where \( \langle \cdot \rangle \) denotes average over the total number of links in the network. Positive values of \( \rho_k \) indicate assortative mixing, while negative values refer to disassortative networks. A specific correlation \( \rho_k \) can be obtained by rewiring the links while keeping the degree sequences unchanged (Newman, 2003a). However, one can expect that \( \ell \) and \( b_{\text{max}} \) change also when the networks are rewired.

The influence of degree correlations on synchronizability was addressed by Motter et al. (2005b), showing that the eigenratio \( R \) increases when increasing the assortativity of the network. A more detailed and systematic
analysis was later on carried out by di Bernardo et al. (2007). They generated random SF network as in Newman et al. (2001) with a minimal degree \(k_{\text{min}} = 5\) and obtained the desired degree correlation \(r_k\) using the rewiring procedure of Newman (2003a) and Motter et al. (2005b). The dependence of the synchronizability on \(r_k\) is shown in Fig. 15. The main effect on the eigenratio \(R\) is from the fact that \(\lambda_2\) decreases when \(r_k\) grows, while \(\lambda_2\) remains roughly constant (di Bernardo et al., 2007). As it happens for the other parameters, the dependence of \(\lambda_2\) on \(r_k\) can be obtained from graph theoretical analysis, which is the subject we are going to discuss next.

5. Graph theoretical bounds to synchronizability

Previously we have seen that many structural properties can influence the synchronizability of the networks, but none of them can be regarded as the exclusive factor in the observed dependencies. The moral is that all works described in the preceding paragraphs seem not to be on the most appropriate way to elucidate the dependence of synchronizability on the network characteristics. Since the synchronizability depends on the ratio between the two extreme eigenvalues of the Laplacian matrix, a sound analysis must attack the raw problem of the spectral properties of networks from a mathematical point of view, given that the simulation experiments are far from being conclusive.

Graph theoretical analyses of the Laplacian matrix \(L\) mainly focus on the bounds of its extreme eigenvalues. These bounds have been discussed in several works (Atay et al., 2006a,b; Chung et al., 2003; Donetti et al., 2007; Nishikawa et al., 2003; Pecora and Barahona, 2005; Wu, 2003, 2005). Here we summarize the main results and discuss how they help to understand the synchronizability of complex networks. Some detailed proof of the results can be found in the corresponding references and in books on graph theory, e.g. (Chung, 1997; Mohar, 1991a).

First, we follow Wu (2005) and Atay et al. (2006a,b) to discuss the bounds for networks with prescribed degree sequences \(k_{\text{min}} = k_1 \leq k_2 \leq \cdots \leq k_N = k_{\text{max}}\). The bounds are

\[
2\left(1 - \cos\left(\frac{\pi}{N}\right)\right)k_{\text{min}} \leq \lambda_2 \leq \frac{N}{N-1}k_{\text{min}}.
\]

and

\[
\frac{N}{N-1}k_{\text{max}} \leq \lambda_N \leq 2k_{\text{max}}.
\]

It follows that

\[
\frac{k_{\text{max}}}{k_{\text{min}}} \leq R \leq \frac{k_{\text{max}}}{\left(1 - \cos\left(\frac{\pi}{N}\right)\right)k_{\text{min}}}.
\]

Note that the lower bound in Eq. (71) is the same as the minimal nonzero eigenvalue of the regular network with range \(k = k_{\text{min}} = 2\) in Eqs. (63-64). If \(k\) is finite, the bound decreases as \(1/N^2\) which is asymptotically the fastest possible decay. This means that among all networks with the same minimal degree \(k_{\text{min}}\), regular networks have the lowest synchronizability, since \(\lambda_2\) approaches 0 in the fastest way as \(N \to \infty\).

The upper bound in Eq. (71) is approached when the network is random. As shown by Wu (2003, 2005), in a \(k\)-regular random network, where each node is randomly connected to other \(k = k_{\text{min}}\) nodes in the network, \(\lambda_2 = k - O(\sqrt{k})\) as \(N \to \infty\). In fact, \(\lambda_2 = c_k k\), where \(c_k \to 1\) as \(k \to \infty\) both for \(k\)-regular networks and random SF networks (Wu, 2003). In this sense, the observation in (Wang and Chen, 2002) that \(\lambda_2\) is a good approximation. So for large random networks with large enough minimal degree, \(\lambda_2 \approx k_{\text{min}}\) is a good approximation.

Another general lower bound for \(\lambda_2\) is (Mohar, 1991b)

\[
\frac{4}{ND} \leq \lambda_2 \leq \frac{N}{N-1}k_{\text{min}},
\]

so that

\[
\frac{k_{\text{max}}}{k_{\text{min}}} \leq R \leq \frac{NDk_{\text{max}}}{2},
\]

where \(D\) is the diameter of the graph, i.e. the maximum value of the shortest path lengths between any two nodes. Let us discuss the implications of the above bounds for WS networks. In this model, when the probability of having shortcuts is very low, \(pkN \ll 1\), then \(D \sim N/k\) and the lower bound in Eq. (74) approaches zero as \(1/N^2\) (regular network). Beyond the onset of the SW regime \((pkN \sim 1)\), \(D\) decreases and approaches \(D \sim \ln(N)\), and \(\lambda_2\) increases for fixed \(N\). Thus this bound allows us to understand why the network synchronizability is inversely proportional to \(\ell\), as observed numerically by Barahona and Pecora (2002) and Hong et al. (2004) (Figs. 9 and 10). However, \(\lambda_2\) is not immediately bounded away from zero, since it approaches zero faster than \(1/N\) just after the onset of the SW regime. When moving deeper into...
the SW region, \( pkN \sim N \) so that each node has \( S \sim 1 \) shortcuts, and \( \lambda_2 \) is already bounded away from the lower bound \( 4/ND \) and approaching the upper bound \( k_{\text{min}} \). In this regime, \( \lambda_2 \) will not be sensitive to changes in the diameter \( D \). This helps to understand why the synchronization threshold is different from the onset of the SW behavior in Fig. 12.

Thus, in WS networks with high \( p \), \( k_{\text{max}}/k_{\text{min}} \) provides a good lower bound to the eigenratio \( R \). The upper bound, \( NDk_{\text{max}}/2 \), still increases with \( N \), and can be several orders of magnitude larger than the actual value of \( R \) even for small random networks. In fact, \( R \) may not follow the variation of the distant upper bound when the diameter \( D \) or the size \( N \) change. Thus, in complex networks with both local and random connections, a close relationship between the synchronizability and \( \ell \) is not expected.

Nishikawa et al. (2003) obtained similar bounds but as a function of \( \ell \) and \( b_{\text{max}} \) as

\[
\frac{k_{\text{max}}}{k_{\text{min}}} \leq R \leq Nk_{\text{max}}b_{\text{max}}D\ell. \tag{76}
\]

These authors pointed out that experimental values of \( R \) are closer to the lower bound, and far away from the upper bound (Fig. 13). Thus quite probably such upper bound does not provide meaningful understanding of the relationship between synchronizability and \( \ell \), \( D \) or \( b_{\text{max}} \), since the change of the upper bound by these structural measures is quite likely not to be reflected in the actual value of \( R \). The observation by Nishikawa et al. (2003) that the synchronizability is correlated with a more homogeneous load distribution is basically an effect of the correlation between the load and the degree. In fact, when keeping the degree sequence unchanged, it is observed that more heterogeneous load distributions can be correlated with improved synchronizability (Zhao, Zhou, Wang, Yan, Yang, and Bai, 2006).

The above analysis of bounds provides justification about why we can observe different \( R \) for increased or decreased heterogeneity, distances or loads. Moreover, the bounds expressed by these quantities are not tight at all in the particular examples, e.g., in Eq. (76). In general, the graph theoretical analysis states that randomness improves the synchronizability, since \( \lambda_2 \) is well bounded away from 0, while in networks with dominantly local connections, \( \lambda_2 \) approaches to 0 in large networks. In other words, for a prescribed degree sequence, the eigenratio \( R \) changes mainly because of \( \lambda_2 \). A schematic plot of the bounds of \( \lambda_2 \) is shown in Fig. 16.

On the other hand, the local organization of the connections even within a small part of the network, can make the eigenvalue \( \lambda_2 \) deviate significantly from the upper bound \( k_{\text{min}}N/(N-1) \). For an arbitrary network, there is a better upper bound for \( \lambda_2 \) as

\[
\lambda_2 \leq 2i_G, \tag{77}
\]

where \( i_G \) is the isoperimetric number of a graph (Atay et al., 2006a,b; Molnar, 1989), which is defined as

\[
i_G = \min_S \frac{|\partial S|}{|S|}. \tag{78}
\]

Here \( S \subset G \) is a subset of the nodes, with \( G - S \) denoting its complement, and \( |S| \) is the number of nodes within \( S \), with \( 0 < |S| < N/2 \). Besides, \( |\partial S| \) is the number of connections between \( S \) and its complement, namely,

\[
|\partial S| = \sum_{i \in S} \sum_{j \in (G-S)} a_{ij}. \tag{79}
\]

For an arbitrary partition of the network into \( S \) and \( G \), we have (Atay et al., 2006b)

\[
\lambda_2 \leq 2\frac{|\partial S|}{|S|}. \tag{80}
\]

This means that the synchronizability of the network is determined by the sparse connections between the two subnetworks. For instance, if a small set \( S \) made up of, say, 20 nodes, is connected to \( G-S \) with just one link, then \( \lambda_2 \) < 0.1, regardless of how the nodes are connected within \( S \) and within the large complement \( G-S \). It follows that the statistical properties of the network \( G \) are mainly determined by the huge part \( G-S \), while \( \lambda_2 \) is independently constrained by the small subgraph. This result is in complete agreement with the path towards synchronization in modular networks presented in section III.A.5, where the community structure has been demonstrated to impose scales in the synchronization process.

Everything up to now indicates that statistical properties, such as degree distribution, \( \ell \), etc., may not be always correlated with the synchronizability of the network. In fact, it was particularly shown by Wu (2005) and Atay et al. (2006a,b), that networks with very different synchronizability can be constructed for the same prescribed degree sequences, because \( i_G \) can be at any place in a broad range between the lower and upper bounds in Eq. (71), see also Fig. 16.

Another similar bound in graph theory is based on the Cheeger inequality (Cheeger, 1970; di Bernardo et al., 2007),

\[
\lambda_2 \leq h_G, \tag{81}
\]

where \( h_G = \min_S h_G(S) \) and

\[
h_G(S) = \frac{|\partial S|N}{|S|(N-|S|)}. \tag{82}
\]
As for correlated networks, di Bernardo et al. (2007) showed that in random networks with degree correlations $r_k$

$$\frac{\partial h_G(S)}{\partial r_k} < 0,$$  \hspace{1cm} (83)

which explains why the synchronizability is reduced in networks with assortative connections, as shown in Fig. 15. Along the same line, the dependence of the minimum nonzero eigenvalue on the topological properties of the network and its degree-degree correlation coefficient $r$ was also analyzed by Donetti et al. (2007). The authors derived a rigorous upper bound for $\lambda_2$ as,

$$\lambda_2 \leq (1 - r) \frac{\langle k \rangle \langle k^3 \rangle - \langle k^2 \rangle \langle k^2 \rangle}{\langle k \rangle \langle k^2 \rangle - \langle k^2 \rangle \langle k^2 \rangle}.$$  \hspace{1cm} (84)

Another interesting inequality from graph theory is (di Bernardo et al., 2007; Mohar, 1989),

$$\lambda_2 \geq k_{\text{max}} - \sqrt{k_{\text{max}}^2 - i_{G}^2}.$$  \hspace{1cm} (85)

Note that $k_{\text{max}} - \sqrt{k_{\text{max}}^2 - i_{G}^2}$ is a decreasing function of $k_{\text{max}}$ if $i_{G}$ is fixed. Based on this bound, we can expect some apparently counterintuitive effects on the synchronizability of complex networks. Suppose we put more links to the graph $G$, but only add them to the nodes within $G - S$ and $S$, but not to the nodes between $S$ and $G - S$ so that $i_G$ does not change. For simplicity, we can further assume that the nodes within $S$ and $G - S$ are connected with a uniform probability $f$ (random networks), so that $k_{\text{max}}$ and the mean degree $\langle k \rangle$ increase when more and more links are added. In this case, the synchronizability of the subnetworks $S$ and $G - S$ is enhanced at larger $f$, see Fig. 9. However, $\lambda_2$ of the whole network $G$ is reduced according to Eq. (85). Therefore, in the case of two coupled networks, enhancing the synchronizability of the subnetworks may actually reduce the synchronizability of the whole network. Phenomenologically, this is intuitively expected, because the subnetworks tend to form distinct synchronized clusters.

Based on the above arguments (Eqs. (77,81,85)), networks possessing a clear community organization display a small synchronizability, since the density of connection between different communities can be much smaller than the density within the communities (Almendral and Díaz-Guilera, 2007; Huang et al., 2006; Zhou, Zhao, Chen, Yan and Wang, 2006).

Recapitulating, we have seen that for a prescribed degree sequence, it is possible to construct a very large number of networks ranging from fully local connections to fully random networks (Wu, 2005), with many possible structures in between. However, the degree sequence by itself is not sufficient to determine the synchronizability. On the other hand, we have seen that the synchronizability is not directly related to graph measures, such as distance, clustering or maximal betweenness. Admittedly, the weak connections between two subnetworks (characterized by the isoperimetric number $i_G$) determine the behavior of the eigenvalue $\lambda_2$, and hence that of the synchronizability of the whole network.

6. Synchronizability of weighted networks

Up to now, we have considered the influence of the network topology on synchronization, assuming that the connection weights are the same for all the links in the network, i.e., the networks are unweighted. However, this is not the case for many real-world networks. Indeed, many complex networks where synchronization is relevant are actually weighted and display a highly heterogeneous distribution of both degrees and weights (Barrat et al., 2004; Braunstein et al., 2003; Newman, 2001; Yook et al., 2001). Examples include neural networks (Fellerman and Van Essen, 1991; Scannell et al., 1999), airport networks (Barrat et al., 2004) and the structure of the networks characterizing epidemic outbreaks in different cities (Grassly et al., 2005; Grenfell et al., 2001). Furthermore, it has been observed that in many cases, the connection strength is not an independent parameter, but it is correlated to the network topology. The analysis of some real networks (Barrat et al., 2004) yields the following main properties:

(i) the weight $w_{ij}$ of a connection between nodes $j$ and $i$ is strongly correlated with the product of the corresponding degrees as $\langle w_{ij} \rangle \sim \langle k_i \rangle \langle k_j \rangle$;

(ii) the average intensity $S(k)$ of nodes with degree $k$ increases as $S(k) \sim k^\theta$. Here the intensity $S_i$ of a node $i$ is defined as the total input weight of the node:

$$S_i = \sum_{j=1}^{N} a_{ij} w_{ij}.$$  \hspace{1cm} (86)

Note that the inclusion of a distribution of weights in the network affects directly its classification within topological homogeneity or heterogeneity. For example, a regular lattice with a very skewed distribution of weights can eventually represent a SF topology. From a mathematical point of view, the adjacency matrix is in this case simply substituted by the weight matrix. On the contrary, from a physical perspective, it is still interesting to keep separated the topology of interactions from the distribution of weights, and answer questions, whenever possible, discriminating these two topological aspects.

The first works on synchronization in weighted networks considered that the weighted input of a node $i$ from a node $j$ depends on the degree $k_i$ of node $i$ (Motter et al., 2005a,b), with a model of weighted coupling as $w_{ij} = k_i^\theta$, so that the matrix $\tilde{G} = (G_{ij})$ in Eq. (53) reads

$$\tilde{G} = D^\theta (D - A) = D^\theta L.$$  \hspace{1cm} (87)

Here $D_{ij} = \delta_{ij} k_i$ is the diagonal matrix of degrees. $\theta$ is a tunable parameter that keeps the network topology
unchanged, but varies the distribution of the weights of the links.

Within this scheme, the weights between a pair of nodes $i$ and $j$ are in general asymmetric, because $w_{ij} = k_i^\theta$ and $w_{ji} = k_j^\theta$. However, since

$$\det(D^\theta L - \lambda I) = \det(D^{\theta/2}LD^{\theta/2} - \lambda I),$$

is valid for any $\lambda$, the spectrum of eigenvalues of the matrix $G$ is equal to the spectrum of a symmetric matrix defined as $H = D^{\theta/2}LD^{\theta/2}$. As a result, all the eigenvalues of $G$ are real, and the synchronizability can still be characterized in the framework of the MSF.

The synchronizability of various complex networks as a function of the parameter $\theta$ is shown in Fig. 17. Except for $k$-regular networks, in all other cases, including the SW networks, the eigenratio $R$ exhibits a pronounced minimum at $\theta = -1$. Here the SW networks are obtained by adding $M \leq N(N-2k-1)/2$ new links between randomly chosen pairs of nodes on the basic regular array where each node is connected to its $2k$ first neighbors.

Motter et al. (2005a,b) also characterize the synchronizability of the network, related to $\lambda_2$, using the notion of the cost of the network. When Eq. (62) is satisfied, the fully synchronized state is linearly stable for $\sigma > \sigma_{\text{min}} \equiv \alpha_1/\lambda_2$. The cost is defined as the total input strength of the connections of all nodes at the synchronizability threshold: $\sigma_{\text{min}} \sum w_{ij} \alpha_{ij} = \sigma_{\text{min}} \sum_{i=1}^{N} S_i$. A more convenient definition for comparisons is obtained normalizing by the number of nodes, such that

$$C_0 \equiv \frac{\sigma_{\text{min}} \sum_{i=1}^{N} S_i}{N \alpha_1} = \langle S \rangle / \lambda_2,$$

where $\langle S \rangle$ is the average intensity of nodes in the network. Similar to $R$, $C_0$ is also minimal at $\theta = -1$ (Fig. 18).

Interestingly enough, Motter et al. (2005a,b) obtained that in SF networks with fixed minimal degrees $k_{\text{min}}$, the weighted versions ($\theta = 1$) behave differently to the unweighted networks when one looks at the dependence of both the eigenratio $R$ and the cost $C_0$ on the scaling exponent $\gamma$, as shown in Fig. 19.

$\theta = -1$ is a special case. The coupling matrix is now $G = D^{-1}L$, and all the diagonal elements $G_{ii} \equiv 1$. It is usually called the normalized Laplacian of a graph. Based on graph spectral analysis results by Chung et al. (2003) for random networks with arbitrary given degrees, it can be shown that the spectrum of the normalized Laplacian tends to the semicircle law for large networks. In particular, for $k_{\text{min}} \gg \sqrt{\langle k \rangle}$, one has

$$\max\{1 - \lambda_2, \lambda_N - 1\} = [1 + O(1)]^2 \frac{2}{\sqrt{\langle k \rangle}}.$$  \hspace{1cm} (90)

This result is rigorous for ensembles of networks with a given expected degree sequence and sufficiently large minimum degree $k_{\text{min}}$, but the numerical results reported in Fig. 19 support the hypothesis that the approximate relations

$$\lambda_2 \approx 1 - 2/\sqrt{\langle k \rangle}, \quad \lambda_N \approx 1 + 2/\sqrt{\langle k \rangle}$$  \hspace{1cm} (91)
hold under much milder conditions. In particular, the relations (91) are expected to hold for any large network with a sufficient number of random connections, $k_{\text{min}} \gg 1$.

Furthermore, the synchronizability in this case seems to be independent of the degree distribution. It is only controlled by the average degree $\langle k \rangle$, since the synchronizability of the weighted SF networks is almost identical to that of a regular random network where each node has the same degree $k_i = \langle k \rangle$. These results demonstrate that the topological degree of networks is not the only determinant of the synchronizability of the networks; having a heterogeneous distribution for the connection strengths can significantly influence the synchronizability.

7. Universal parameters controlling the synchronizability

What are the leading parameters governing the synchronizability for the more general case in which weighted networks are considered? Zhou, Motter and Kurths (2006) have proved analytically and verified numerically what controls the synchronizability of sufficiently random networks with large enough minimal degree ($k_{\text{min}} \gg 1$). It is the distribution of the intensities $S_i$ defined in Eq. (86). The intensity of a node incorporates both the information about the topology and the weights of the connections in the networks. The main finding is that the synchronizability is sensitively controlled by the heterogeneity of the intensity $S_i$. The eigenratio $R$ and the normalized cost $C_0$ can be expressed as

$$R = A_R \frac{S_{\text{max}}}{S_{\text{min}}}, \quad C_0 = A_C \frac{\langle S \rangle}{S_{\text{min}}},$$

(92)

where $S_{\text{min}}$, $S_{\text{max}}$, and $\langle S \rangle$ are the minimum, maximum and average intensities, respectively. The pre-factors $A_R$ and $A_C$ are expected to approach 1 for large average degree $\langle k \rangle$. Equations (92) are universal in the sense that they apply to many random networks with arbitrary degree and weight distributions provided that the minimal degree is sufficiently large. The main hypothesis behind this result is the assumption that the local mean fields $H_i = \langle 1/k_i \rangle \sum_j a_{ij} H(x_j)$ can be substituted by the global mean field $\bar{H}_i \approx \bar{H} = (1/N) \sum_j H(x_j)$.

In Fig. 20, Eq. (92) is corroborated by numerical results of $R$ and $C_0$ for networks with several degree and intensity distributions of degrees, using the weighted coupling scheme

$$w_{ij} = S_i/k_i.$$  
(93)

This coupling scheme means that the intensity of the nodes, which is not necessarily correlated with the degree, is uniformly distributed into the input links of the nodes. It covers the coupling scheme in Eq. (87) as a special example: $S_i = k_i^{1+\theta}$. Recent analysis by Kim and Kahng (2007) on the spectral density of SF networks with a weighted Laplacian matrix similar to Eq. (87), also confirms that Eq. (92) holds.

Zhou, Motter and Kurths (2006) also presented results for the following coupling scheme

$$w_{ij} = (k_i k_j)^{\theta},$$

(94)

that describes the relationship between the weights and the degrees in some real networks (Barrat et al., 2004; Macdonald et al., 2005). The tunable parameter $\theta$ controls the heterogeneity of the intensity $S_i$ and the correlation between $S_i$ and $k_i$, since $S_i = k_i^\theta \langle k_i^\theta \rangle_i$, where $\langle k_i^\theta \rangle_i = (1/k_i) \sum k_i^\theta_j$ is approximately constant for $k_i \gg 1$ when the degree correlations can be neglected. Variations of $\theta$ have a significant impact on the synchronizability of networks which are heterogeneous in degree (Fig. 21). Note that in heterogeneous in degree networks, the weighted coupling in Eq. (94) may result in a broad distribution of the input weights $w_{ij}$ among the $k_i$ links of the node $i$, especially when $\theta$ is not close to 0. However, as shown in the insets of Fig. 21 for various networks and $\theta$ values, $R$ and $C_0$ collapse again to the universal curves when regarded as functions of $S_{\text{max}}/S_{\text{min}}$ and $\langle S \rangle/S_{\text{min}}$, respectively. The fact that the universal formula holds for a broad range of $\theta$ values shows that the mean field approximation used to obtain Eq. (92) often remains valid under milder conditions.

It is important to stress that these results also hold for unweighted random networks. In this case $S_i = k_i$ for all
the nodes and one gets

\[ k_{\text{min}}(1 - \frac{2}{\sqrt{\langle k \rangle}}) \lesssim \lambda_2 \lesssim k_{\text{min}} \]

\[ k_{\text{max}} \leq \lambda N \leq k_{\text{max}}(1 + \frac{2}{\sqrt{\langle k \rangle}}), \]

for large random networks with minimal degree \( k_{\text{min}} \gg 1 \). These results provide much tighter bounds than those discussed in the previous sections (e.g., cf. Eqs. (73,75, 76)) which depend on the system size \( N \). Interestingly, Eqs. (92) also provide meaningful insights into the problem for other special networks. For example, consider the class of SF networks generated using the BA model. When \( m = 1 \), the network is a tree, and \( R \) is much larger than \( S_{\text{max}}/S_{\text{min}} \). However, it is an increasing function of \( S_{\text{max}}/S_{\text{min}} \), showing that the heterogeneity of the intensity is still an important parameter. But for \( m = 2 ((k) = 4) \) it approaches the universal curve quickly (Fig. 22 (a)). The drastic change of synchronizability from \( m = 1 \) to \( m = 2 \) can be attributed to the appearance of loops (Yook and Meyer-Ortmanns, 2006). In WS networks (Watts and Strogatz, 1998) with \( N = 2^{10} \) and \( \langle k \rangle = 20, R \) collapses to the universal curve even when the networks are dominated by local connections, e.g. for a rewiring probability \( p = 0.3 \) with intensities \( S_{\text{max}}/S_{\text{min}} \gtrsim 10 \), see Fig. 22 (b).

B. Design of synchronizable networks

An interesting subject related to the impact of network structure on synchronization dynamics is the design of synchronizable networks. Here we review several ideas exploring this issue: weighting the couplings leaving the topology unchanged, perturbing part of the network topology, and finally searching for optimal topologies with respect to synchronizability. Note that the following theoretical schemes may not directly apply to real complex networks. It is difficult to conceive real systems where the weights can be tuned at discretion, or where the topological substrate of interactions can be changed accordingly. Nevertheless, the insights given by these works allow for a deeper understanding of the synchronizability of networks.

1. Weighted couplings for enhancing synchronizability

The previous analysis shows that for networks that are heterogeneous in degree, synchronizability can be enhanced by balancing the heterogeneity in the degree distribution with suitable weighted couplings, towards the obtention of a homogeneous distribution of the intensities \( S_i \).

Chavez et al. (2005) presented a different scheme assuming that the weight of a link is related to its betweenness \( b_{ij} \) as

\[ G_{ij} = -\frac{b_{ij}^\alpha}{\sum_{j \in \Gamma} b_{ij}^\alpha}. \]

where \( \alpha \) is a tunable parameter that controls the dependence of the weights \( w_{ij} \) on the loads \( b_{ij} \). The zero-sum requirement of the matrix \( G \) implies that \( G_{ii} = 1 \) for all \( i \). Note that \( \alpha = 0 \) corresponds to the weighted coupling scheme in Eq. (87) at the optimal point \( \theta = -1 \). As seen in Fig. 23, the eigenratio \( R \) depends on \( \alpha \), reaching a minimum at a value \( 0 < \alpha \lesssim 1 \), showing that the synchronizability in SF networks can be slightly enhanced compared to the optimal case of the weighted coupling scheme (Motter et al., 2005a,b; Zhou, Motter and Kurths, 2006). The SF network considered by Chavez et al. (2005) is a generalized BA model with a preferential attachment probability \( \pi_i \sim k_i + B \) (Dorogovtsev et al., 2000), where the parameter \( B \) controls the exponent \( \gamma = 3 + B/m \) of the power law degree distribution, and \( m = k_{\text{min}} \). At large \( \alpha \) values, only the links with the largest loads \( b_{ij} \) are significant, which can lead to effectively disconnected nodes, so that synchronizability is reduced. Chavez et al. (2005) also pointed out that for large minimal degrees, the regimes corresponding to
enhanced synchronizability are reduced so that the minimum approaches to $\alpha = 0$. This demonstrates again that for random networks with large enough minimal degree, Eq. (91) is asymptotically valid regardless of the detailed weighted scheme, as claimed by Zhou, Motter and Kurths (2006).

The explanation of the observed enhanced synchronizability proposed by Chavez et al. (2005) is that the load $b_{ij}$ reflects the global information of the network, while at $\alpha = 0$ only the local information (degree) is employed. Such a heuristic explanation, however, is not supported by several further investigations. In fact, only the local information can also lead to similar enhanced synchronizability. For example, consider that the weights depend on the degrees following Eq. (94), and then normalize to allow fully uniform intensity $S_i = 1$, namely (Motter et al., 2005c)

$$G_{ij} = \frac{(k_i k_j)^{\alpha}}{\sum_{j \in \Gamma_i} (k_i k_j)^{\alpha}} = \frac{k_i^\alpha}{\sum_{j \in \Gamma_i} k_j^\alpha}, \quad \alpha = 0$$

Again, $\alpha = 0$ corresponds to the optimal case (Eq. (87), $\theta = -1$) of the weighted coupling scheme (Motter et al., 2005a,b; Zhou, Motter and Kurths, 2006). Similar to Chavez et al. (2005), the synchronizability can be further enhanced in a range $\alpha > 0$ (Fig. 24). However, the minimum moves closer to $\alpha = 0$ when the networks are larger, indicating that the synchronizability in large random networks with $k_{\text{min}} \gg 1$ can hardly be enhanced further with other weighted coupling schemes different from the optimal case in Eq. (87). This coupling scheme was also considered by Lind et al. (2004) where synchronizability in SF networks of maps is enhanced for $\alpha > 0$. Additionally, we note that the coupling form in Eq. (96) has been recently revisited from the viewpoint of gradient-network (Wang et al., 2007).

Another work (Zhao, Zhou, Wang, Ou and Ren, 2006) introduced an additional parameter $\beta$ into the coupling scheme in Eq. (96), as

$$G_{ij} = \frac{k_i^\alpha}{(\sum_{j \in \Gamma_i} k_j^\alpha)^{\beta}}. \quad \beta \neq 1$$

In this case, the intensity of the node is $S_i = (\sum_{j \in \Gamma_i} k_j^\alpha)^{1-\beta}$.

More weighted coupling methods have been proposed. Hwang et al. (2005) use the information about the age of the nodes in growing networks and introduce asymmetrical coupling between old and young nodes. In particular, they propose

$$G_{ij} = \frac{\theta_{ij}}{(\sum_{j \in \Gamma_i} \theta_{ij})}, \quad \theta_{ij} = (1 - \theta)/(1 + \theta)$$

where $\theta_{ij} = (1 - \theta)/(1 + \theta)$ for the connections from old to young nodes ($i > j$) and $\theta_{ij} = (1 + \theta)/(1 - \theta)$ for the connections from young to old nodes ($i < j$). The limit $\theta = -1$ ($\theta = 1$) gives a unidirectional coupling where the old (young) nodes drive the young (old) ones. It was shown that in SF networks, synchronizability is enhanced when couplings from older to younger nodes are dominant ($\theta < 0$).

In spite of these numerical observations, a clear understanding about why the synchronizability is further
enhanced with the various weighted coupling schemes (Eqs. (95,96,97)) has not been obtained. The same scheme as in (98), but without the normalization \((G_{ij} = -a_{ij}\Theta_{ij})\) was considered by Zou et al. (2006), and again the synchronizability is enhanced for \(\theta < 0\). In this case, the change of synchronizability is mainly due to the heterogeneity of the intensity distribution \(S_i\), which becomes more homogeneous for \(\theta < 0\) because the old nodes are hubs and have most of the connections to young nodes. In this case, the universality in Eq. (92) should apply when the minimal degree is large enough. Finally, it has also been shown that a random distribution of the weights of the connections of regular networks, with only nearest neighbors, can also enhance synchronizability. This fact is related to the effective presence of short cuts in terms of weights (Li et al., 2007).

As it has been already pointed out, the particular scientific course of action taken by proposing different weighting schemes to enhance synchronizability is unfinished, and probably an unfruitful quest given the many possibilities of inventing new weights. A more rigorous analysis of the eigenspectra of general graphs beyond the possibility of inventing new weights. A more rigorous analysis of the eigenspectra of general graphs beyond the framework (Eqs. (95,96,97)) has not been obtained. The same scheme as in (98), but without the normalization \((G_{ij} = -a_{ij}\Theta_{ij})\) was considered by Zou et al. (2006), and again the synchronizability is enhanced for \(\theta < 0\). In this case, the universality in Eq. (92) should apply when the minimal degree is large enough. Finally, it has also been shown that a random distribution of the weights of the connections of regular networks, with only nearest neighbors, can also enhance synchronizability. This fact is related to the effective presence of short cuts in terms of weights (Li et al., 2007).

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The adaptation process makes the intensity more homogeneous, so that it is expected that the synchronizability is enhanced. In Fig. 26 we show the eigenratio \(R\) as a function of network size \(N\). There we compare the original unweighted network with two weighted networks after the adaptation. Suppose that the largest network size synchronizable for the bounded MSF \((R = \alpha_2/\alpha_1)\) is \(N_1\) for unweighted networks and \(N_2\) for weighted networks obtained from adaptation. It then follows that \(N_2/N_1 \sim (\alpha_2/\alpha_1)^{(1+\theta)} \sim (\alpha_2/\alpha_1)^2\) for power law degree distributions regardless of the exponent \(\gamma\) (Zhou and Kurths, 2006a).
2. Topological modification for enhancing synchronizability

Some authors have proposed to enhance synchronizability by perturbing the network topology. Based on the argument that heterogeneity in the betweenness distribution is related to poor synchronizability, Zhao et al. (2005) and Yin et al. (2006) proposed to modify the nodes or links with the highest maximal betweenness. As already noticed, in SF networks, the betweenness of a node and a link is strongly correlated with the degree \( k \), and the product of degrees \( k_i k_j \) of the two nodes at the ends, respectively. The perturbation proposed by Zhao et al. (2005) consists of dividing the node with the highest degree into a group of several fully connected nodes and redistribute the \( k \) links equally over the new nodes. Following this scheme, the synchronizability can be substantially enhanced by modifying a very small portion of the nodes. The enhanced synchronizability follows closely the reduced maximal degree in the networks. It was also shown that the average distance actually increases when the hubs are divided. Yin et al. (2006) propose that the connections with the largest \( k_i k_j \) are broken, and again the synchronizability can be enhanced by cutting a small fraction of links with high betweenness. These ideas can plausibly be implemented in technological networks, where the substitution of hubs by a core of nodes is possible. In this way, the redistribution of load will improve traffic, and as a by-product, the synchronizability.

3. Optimization of synchronizability

A more straightforward approach to the design is that of asking which are the best network architectures to get an optimal synchronizability. Donetti et al. (2005) applied an optimization scheme (e.g. simulated annealing) combined with a network rewiring algorithm to minimize the eigenratio \( R \). In this case, the total number of nodes and links are preserved. The resulting networks, with optimal synchronizability, called entangled networks, are found to be very homogeneous in many topological measures, such as degrees, distance between nodes, betweenness etc. This result is quite relevant because it provides a null model that allows to compare the synchronizability of networks directly with its optimal counterpart.

A similar optimization scheme was applied to study the optimal synchronizability in networks with a preserved SF degree sequence (Liu et al., 2007). In this case, the synchronizability can only be slightly enhanced. An interesting finding is that the optimized networks become disassortative and the clustering and the maximal betweenness is reduced, which is consistent with the observed enhanced synchronizability obtained by changing these features.

Thus, optimization schemes are helpful to identify meaningful topological features that correlate with the synchronizability of complex networks. However, such optimization schemes are computationally demanding when we deal with large networks. The development of analytical tools to attack this optimization problem is currently a major challenge in the subject.

Furthermore, in these two studies, the underlying network topology is bi-directional. In many of the weighted coupling schemes previously mentioned, the connections are effectively directed, since the coupling strength is asymmetrical for the input and output links. Nishikawa and Motter (2006) went to the extreme by imposing an unidirectional information flow so that the networks become optimally synchronizable with \( R = 1 \). For any topology, the maximally synchronizable network can be achieved by imposing that the network: (i) embeds a directed spanning tree, (ii) has no directed loop, and (iii) has normalized input strengths. With these conditions, the original networks are changed into feedforward net-
works without any feedback, and $0 = \lambda_1 < \lambda_2 = \cdots = \lambda_N = \lambda$. Furthermore, the synchronization of the whole network is achieved in a hierarchical way. The conditions (i)-(iii) lead to the following path towards synchrony. There is a node in the directed spanning tree that has no input and acts as a master oscillator driving the network dynamics. If $\alpha = \sigma \lambda$, the oscillators that are just one hierarchical level below the master oscillator will synchronize. Then, the next lower level oscillators will also get synchronized and so on until the whole network reaches complete synchronization. Note that this happens for the entire range of the coupling strength where $\alpha = \sigma \lambda$. Finally, we note that it could take a very long time to achieve complete synchronization when the number of effective layers is large. It would be interesting to see how the topology of the original network is related to the depth of such effective directed trees and how it influences the transient time towards synchronization. Very recently, it was shown (Lu et al., 2007) that an age-based coupling similar to Eq. 98, but with $\Theta_{ij} = e^{-\Theta(i-j)/N}$, will lead to such an effective directed tree with $R = 1$ at large values of $\alpha$ (Lu et al., 2007).

C. Beyond the Master Stability Function formalism

We have discussed how the impact of network topology on synchronizability can be addressed using the MSF and graph theory. Away from the complete synchronization regime, the linear stability does not strictly apply. However, it is still possible to go one step forward to further understand some aspects of the dynamical synchronization patterns.

Zhou and Kurths (2006b) studied effective synchronization patterns in unweighted SF networks of chaotic oscillators (with the coupling function $H(x) = x$) in several situations: away from the complete synchronization regime, when the coupling strength is smaller than the threshold for complete synchronization, when the oscillators have mismatches in parameters, and when there are noise perturbations. They considered a mean field approximation in which each oscillator is influenced by a global mean field $X$, with a coupling strength $\sigma k_i$, namely,

$$x_i = F(x_i) + \sigma k_i (X - x_i), \quad k_i \gg 1.$$  \hfill (101)

The authors compared the synchronization of each oscillator to $X$ by computing $\Delta X_i = |x_i - X|$ and then obtained the average $\Delta X(k)$ over all nodes with the same degree $k$. It was shown that out of the complete synchronized state

$$\Delta X(k) \sim k^{-\gamma},$$  \hfill (102)

where the exponent $\gamma \approx 1$, as seen in Fig. 28. This result shows that in heterogenous networks, the hubs ($k_i > k_{th}$) will synchronize more closely with the mean field and they will form effective synchronization clusters ($|x_i - x_j| < \Delta_{th}$). However, there is not a unique threshold to define such effective clusters (see Fig. 28). This path to synchronization, i.e., the formation of clusters by the hubs, was further described later in Gómez-Gardeñes et al. (2007a), see also Sect. III.A.4.

The formation of such SF or hierarchical clusters could be understood from a linear analysis using the MSF formalism. The linear variational equations of (101) read

$$\dot{\xi}_i = [DF(X) - \sigma k_i I] \xi_i, \quad k_i \gg 1.$$  \hfill (103)

They have the same form as (56), except that $\lambda_i$ is replaced by $k_i$ and $DH$ by the identity matrix $I$. The MSF for the coupling function $H(x) = x$ is $\lambda_{\text{max}}(\alpha) = \lambda^F - \alpha$, where $\lambda^F$ is the largest Lyapunov exponent of the isolated oscillator $F$. Thus the largest Lyapunov exponent $\lambda_{\text{max}}(k_i)$ of the linearized Eq. (103) is a function of $k_i$ and becomes negative for $\sigma k_i > \lambda^F$. For large $k$ values satisfying $\sigma k \gg \lambda^F$, we have $\lambda_{\text{max}}(k) \approx -\sigma k$.

Now suppose that the network is not completely synchronized, but slightly perturbed from the state of complete synchronization, when the coupling strength $\sigma$ is below the complete synchronization threshold, or when there is noise present in the system. For nodes with large degree $k$, so that $\lambda_{\text{max}}(k) \approx -\sigma k$ is large enough in absolute value but negative, the dynamics of the averaged synchronization difference $\Delta X(k)$ over large time scales can be expressed as

$$\frac{d}{dt} \Delta X(k) = \lambda_{\text{max}}(k) \Delta X(k) + c,$$  \hfill (104)

where $c > 0$ is a constant denoting the level of perturbation with respect to the complete synchronization state, and depends either on the noise level $D$ or on the coupling strength $\sigma$. From this we get the asymptotic result $\Delta X(k) = c/|\lambda_{\text{max}}(k)|$, yielding

$$\Delta X(k) \sim k^{-1},$$  \hfill (105)

which explains qualitatively the numerically observed scaling (solid lines in Fig. 28). Interestingly, the same
scaling dependence but for the time needed to get back into the fully synchronized state was obtained by Moreno and Pacheco (2004) for a population of Kuramoto oscillators, see Sect. III.A.4.

To round off this section, let us mention other works about synchronizability in complex networks that make use of linear criteria similar to the MSF (Chen, 2006; Li and Chen, 2003; Wang and Chen, 2002). The analysis of the global stability of the synchronized state, was first carried out for general graphs (Wu and Chua, 1995) and then followed for specific complex networks (Belykh et al., 2006, 2004b; Chen, 2007; Li and Chen, 2006). The global stability requires additional constraints on the dynamical properties of the individual oscillators. For example, consider the form of coupling as in Eq. (52), the requirement imposed by (Belykh et al., 2006, 2004b) is that the following auxiliary system of the synchronization difference \( \dot{X}_{ij} = x_i - x_j \),

\[
\dot{X}_{ij} = \left[ \int_0^1 D \mathbf{F}(\beta x_j + (1 - \beta)x_i) d\beta - \alpha DH \right] X_{ij},
\]

be globally stable at the fixed point \( X_{ij} = 0 \) for \( \alpha > \alpha_c \).

With this requirement, the condition for global synchronization of a network is

\[
\sigma > \sigma^* = \max \left( \frac{\alpha_c b_l}{N} \right),
\]

(106)

where \( b_l \) is the sum of the lengths of all chosen paths that pass through a given link \( l \) in the network. Note that \( b_l \) is related to both the betweenness and the path length of the link. The result, which is for undirected networks, also holds for directed networks where the input and output degrees are equal for every node of the network. Finally, Belykh et al. (2006, 2004b) also derived the condition needed for global synchronization in more general cases where each link in the network may have a different coupling strength that is allowed to vary in time.

V. APPLICATIONS

The focus of the review up to now has been to revise the main contributions, from theoretical and computational points of view, to our understanding of synchronization processes in complex networks. In this section we will overview the applications to specific problems in such different scientific fields as biology and neuroscience, engineering and computer science, and economy and social sciences. There are nowadays several problems where the application of the ideas and techniques developed in relation to synchronization in complex networks are very clear and the results help to understand the interplay between topology and dynamics in very precise scenarios. There are other cases, also included here for completeness, for which most of the applications so far have been developed in simple patterns of interaction, but extension to complex topologies is necessary because it is its natural description.

A. Biological systems and neuroscience

In biology, complex networks are found at different spatio-temporal scales: from the molecular level up to the population level, passing through many intermediate scales of biological systems. In some of these networks, dynamical interactions between units, which are crucial for our current understanding of living systems, can be analyzed in the framework of synchronization phenomena developed so far. Here we review some of these application scenarios where synchronization in networks has been shown to play an essential role. Thus, at the molecular level we can analyze the evolution of genetic networks and at the population level the dynamics of populations of species coupled through diffusion along spatial coordinates and through trophic interactions. Amongst these two extremes we find a clear application in the analysis of circadian rhythms. On a different context, neuroscience offers applications also at two different levels, one for the synchronization of individual spiking neurons and the other for the coupling between cortical areas in the brain.

1. Genetic networks

The finding that a few basic modules are the building blocks of large real regulatory networks has enabled the design and construction of small synthetic regulatory circuits to implement particular tasks. One of the most salient examples of a synthetic gene network is the "repressilator", that has become one of the best studied model systems of this kind. The repressilator is a network of three genes, whose products (proteins) act as repressors of the transcription of each other in a cyclic way. This synthetic network was implemented experimentally in the bacterium *E. coli*, so that it periodically induces the synthesis of a green fluorescent protein as a readout of the repressilator state (Elowitz and Leibler, 2000). It turns out that the temporal fluctuations in the concentration of each of the three components of the repressilator can be reproduced by a system of six ordinary differential equations,

\[
\frac{d[x_i]}{dt} = -[x_i] + \frac{\alpha}{1 + [y_i]^2},
\]

(107)

\[
\frac{d[y_i]}{dt} = -\beta([y_i] - [x_i]),
\]

(108)

where the couples \((i, j)\) assume the values \((1, 3), (2, 1)\) and \((3, 2)\). The variable \([x_i]\) is the mRNA concentration encoded by gene \(x_i\), and \([y_i]\) is the concentration of its translated protein \(y_i\). The parameter \(\alpha\) is the promoter rate, the parameter \(\beta\) is the ratio of the protein decay rate to the mRNA decay rate, and time has been rescaled in units of the mRNA lifetime. This system has a unique steady state which can be stable or unstable depending on the parameter values and constitutes an illustrative example of the experience gained by identifying network
modules and modeling its dynamical behavior in real networks.

Not surprisingly, the repressilator has called much attention from experts on biological synchronization, since it offers good prospects for further insights into the nature of diverse biological rhythms—whose mechanisms remain to be understood—which are generated by thousands of cellular oscillators that operate synchronously. García-Ojalvo et al. (2004) recently proposed a simple modular addition of two proteins to the repressilator original design that could be used to describe the metabolic oscillations observed in a well mixed suspension of yeast cells. In the new setting, one of the new proteins can diffuse through the cell membrane, thus providing a coupling mechanism between cells containing repressilator networks. This inter-cell communication couples the dynamics of the different cell oscillators (with different repressilator periods) and thus allows us to study the transition to synchronization of coupled phase oscillators in a biological system. In particular, in the limit of infinite cell dilution, the system is made up of a population of uncoupled limit-cycle oscillators. This is not anymore the case when the cell density increases, as now extracellular diffusion provides a mechanism of intercell coupling. As a result, the system shows partial frequency locking of the cells. Finally, if the cell density is large enough, complete locking and synchronized oscillations are observed. García-Ojalvo et al. (2004) proposed an order parameter to measure the degree of synchronization of oscillatory behavior. The dependence of this order parameter $R$ defined as the ratio of the standard deviation of the time series of the average signal to the standard deviation of each individual signal $x_i$ averaged over all signals $i$, as a function of the coupling strength, is shown in Fig. 29 for different values of the ratio between the mRNA and protein lifetimes width distribution ($\delta/\beta$). Note that the phase diagram of the coupled repressilators can be explained by the very same mechanism involved in the transition to synchronization for systems of coupled oscillators (e.g., Kuramoto oscillators) studied in Sec. III.A. What is relevant here is that the transition from an unsynchronized to a synchronized regime is caused by an increase in cell density and therefore the experimental observation of a synchronizing transition in biological phase oscillators might be achieved. In fact, Wagemakers et al. (2006) have recently designed a simple electronic circuit analogy of a population of globally coupled repressilators. They show that coupling is more efficient than externally forcing for the achievement of synchronization. In contrast to the existence of a unified rhythm that gives rise to synchronization, Koseska et al. (2007) analyze the mechanisms of intercell communication that can be responsible for multi-rhythmicity in coupled genetic units. We foresee that works on this line of research will incorporate more genetic interactions in the near future, being the complex network substrate and the synchronization dynamics key aspects of the whole problem.

2. Circadian rhythms

A circadian rhythm is a roughly 24-hour cycle in the physiological processes of living systems; usually endogenous, or when it is exogenous it is mainly driven by daylight. Understanding circadian rhythms is crucial for some physiological and psychological disorders. A nice description of experiments carried out in human beings, in which their circadian rhythms are altered, can be found in the books by Strogatz (2003b) and Pikovsky et al. (2001). Circadian rhythms are known to be dependent on the network of interactions between different subsystems. For example, daylight sensed by eyes and processed by the brain develops a chain of interactions that affects even the behavior of certain groups of cells. On a different scenario, the work by Chialvo and Jalife (1987) reports how non-oscillatory cardiac conducting tissues, when driven rhythmically by repetitive stimuli from their surroundings, produce temporal patterns including phase locking, period-doubling bifurcation and irregular activity.

Synchronization phenomena in complex networks of coupled circadian oscillators has been recently investigated experimentally (Fukuda et al., 2007) on plant leaves. The vein system is in this case the complex network substrate of the synchronization process. Plant cells are coupled via the diffusion of materials along two types of connections: one type that directly connects nearest-neighboring cells and the other type that spreads over the whole plant to transport material among all tissues quickly. Analyzing the phase of circadian oscillations, the phase-wave propagations and the phase delay caused by the vein network, the authors describe how global synchronization of circadian oscillators in the leaf can be attained. As we have seen throughout this review, the role of the topology of interactions is again fundamental.
in the development of synchronization. This work is representative of the new type of applications we can find in the very recent literature about synchronization in complex networks. This particular case of circadian rhythms in plants might be extended to other living systems, including humans.

3. Ecology

It is a well known observation in nature that fluctuations in animal and plant populations display complex dynamics. Mainly irregular, but some of them can show a remarkably cyclical behavior and take place over vast geographical areas in a synchronized manner (Keith, 1963). One of the best documented cases of such situation are the population fluctuations in the Canadian lynx, obtained from the records of the fur trade between 1821 and 1939 in Canada. Fluctuations in lynx populations show a 10-year periodic behavior from three different regions in Canada (Blasius et al., 1999; Ranta et al., 1999; Stenseth et al., 1999). On the other hand, there are some evidences that the existence of conservation corridors favoring the dispersal of species and enhancing the synchronization over time increases the danger of global extinctions (Earn et al., 2000).

One of the first explanations for such types of behavior was that of synchronous environmental forcing, this is the so-called Moran effect (Moran, 1953). There are, however, other explanations for this phenomenon (Ripa and Ranta, 2007), but in any case the problem highlights the importance of integrating explicitly spatial and trophic couplings into current metacommunity theories (Leibold et al., 2004; Maser et al., 2007). Some efforts along these lines have already been made by considering very simple trophic interaction in spatially extended systems. For example, Blasius et al. (1999) analyze a three-level system (vegetation, herbivores, and predators), where diffusive migration between neighboring patches is taken into account. They find that small amounts of migration are required to induce broad-scale synchronization. Another interesting study is performed by Vandermeer (2004), who, again in an extremely simple model, find that changing the patterns of interaction between consumers and resources can lead to either in-phase synchrony or antiphase synchrony.

Nowadays we know, however, about the inherent complexity of food-webs (Dunne, 2006). Food webs have been studied as paradigmatic examples of complex networks, because they show many of their non-trivial topological features. Furthermore, the existence of conservation corridors affecting the migration between regions adds another ingredient to the structure of the spatial pattern. It is precisely this complexity in the trophic interactions coupled to the spatial dependence that must to be considered in detail in the future to get a deeper understanding of ecological evolution.

4. Neuronal networks

Synchronization has been shown to be of special relevance in neural systems. The brain is composed of billions of neurons coupled in a hierarchy of complex network connectivity. The first issue concerns neural networks at the cellular level. In the last years, significant progress has been made in the studies about the detailed interconnections of different types of neurons at the level of cellular circuits (Binzegger et al., 2004; Markram et al., 2004; Silberberg et al., 2005). At this level, the neuronal networks of mammalian cortex also possess complex structure, sharing SW and SF features. Here are two basic neuron types: excitatory principal cells and inhibitory interneurons. In contrast to the more homogeneous principal cell population, interneurons are very diverse in terms of morphology and function. There is an apparently inverse relationship between the number of neurons in various interneuron classes and the spatial extent of their axon trees—most of the neurons have only local connections, while a small number of neurons have long-range axons (Buzsáki et al., 2004). These properties of neuronal networks reflects a compromise between computational needs and wiring economy (Chklovskii et al., 2002; Karbowski, 2001).

On the one hand, the establishment and maintenance of neuronal connections require a significant metabolic cost that should be reduced, and consequently the wiring length should be globally minimized. Indeed, the wiring economy is apparent in the distributions of projection length in neural systems, which show that most neuronal projections are short (Braitenberg and Schüz, 1998; Sik et al., 1995). However, there also exists a significant number of long-distance projections (Kaiser and Hilgetag, 2006; Sik et al., 1994).

Large-scale synchronization of oscillatory neural activity has been believed to play a crucial role in the information and cognitive processing (Fries, 2005). At the level of cellular circuits, oscillatory timing can transform unconnected principal cell groups into temporal coalitions, providing maximal flexibility and economic use of their spikes (Konig et al., 1996). Brains have developed mechanisms for keeping time by inhibitory interneuron networks (Buzsáki and Chrobak, 1995). The wiring will be the most economic if the connections were all local. However, in this case physically distant neurons are not connected, and synaptic path length and synaptic delays become exceedingly long for synchronization in large networks. From previous analysis of synchronization of random networks, we know that synchronizability (stability of the synchronized state) is optimal in fully random networks with a uniform connectivity per node, independent of the network size (Zhou, Motter and Kurths, 2006). The same happens if interneuronal oscillators are coupled (Buzsáki et al., 2004). However, fully random connections irrespective of physical distance are not economic if wiring cost is taken into account.

In Buzsáki et al. (2004), it was shown with a model of
interneuronal networks containing local neurons (Gaussian distribution of projection length) and a fraction of long-range neurons (power law distribution of projection length), that the ratio of synchrony to wiring length is optimized in the SW regime with a small fraction of long-range neurons. Thus, most wiring is local and neurons with long-range connectivity and large global impact are rare, as consistent with observations. It was argued that the complex wiring of diverse interneuron classes could represent an economic solution for supporting global synchrony and oscillations at multiple time scales with minimum axon length (Buzsáki et al., 2004). While such mathematical consideration can predict the scaling relationship among the interneuron classes in brain structures of varying sizes, understanding the role of complex neuronal connectivity, most likely mediated by synchronization, is still one of the main challenges in neuroscience. The theory reviewed in this article will surely contribute to their understanding when more systematic information of neuronal connectivity becomes available in the future.

5. Cortical networks

On a larger neurophysiological scale, the activity observed experimentally by electroencephalographs or functional magnetic resonance imaging, is characterized by oscillations occurring over a broad spectrum and by synchronization phenomena over a wide range of spatial and temporal scales. Reliable databases are available now for large-scale systems level connectivity formed by long-range projections among cortical areas in the brains of several animals (Scannell et al., 1999; Sporns et al., 2004). Large-scale brain networks are found to be densely connected, with very complex and heterogeneous connectivity patterns (Hilgetag and Kaiser, 2004; Hilgetag et al., 2006; Sporns and Zwi, 2004). In parallel, the investigation of brain activity has also put significant emphasis on large-scale functional interactions, characterized by coherence and synchronization between the activity of cortical regions (Bassett et al., 2006; Eguíluz et al., 2005; Salvador et al., 2005; Stam, 2004). Both the structural and functional connectivity of the brain display SW and SF features. The relationship between structural and functional connectivity remains an important open problem in neuroscience.

Recent simulations of synchronization dynamics of brain networks have shed light on this challenging problem. In a series of papers (Zemanová et al., 2006; Zhou, Zemanová, Zamora, Hilgetag and Kurths, 2006; Zhou et al., 2007) the dynamics of a realistic cortico-cortical projection network of the cat has been modeled at the level of functional areas (Scannell et al., 1999). At this level, the network (see Fig. 30) displays a hierarchical cluster organization (Hilgetag and Kaiser, 2004). There are four prominent clusters that agree broadly with the four functional subsystems: visual (V), auditory (A), somatomotor (SM), and frontolimbic (FL). They simulated the network dynamics by a 2-level model: each node (cortical area) is represented by a SW subnetwork of neurons (network of networks). It was shown that the model possesses two distinguished regimes, weak and strong synchronization. In the weak synchronization regime, the model displays biologically plausible dynamical clusters. The functional connectivity, obtained by passing the correlation matrix through various thresholds, exhibits various levels of organization. The clusters with the highest levels of synchronization are from respective functional subsystems (Fig. 31 (a,b)) and are related to specialized functions of the subsystems. The specialized clusters are integrated into larger clusters through brain areas having many inter-community connections (Fig. 31 (c,d)). As a whole, the functional connectivity reveals the hierarchical organization of the structural connectivity (Zhou, Zemanová, Zamora, Hilgetag and Kurths, 2006). The dynamics forms four major clusters (Fig. 32), in excellent agreement with the four functional subsystems (Zemanová et al., 2006). Furthermore, brain areas that bridge different dynamical clusters are found to be the areas involved in multisensory associations. In a comparative study (Zhou et al., 2007), it was shown that representing the brain areas with a periodic, low-dimensional neuronal mass oscillator describing alpha waves (da Silva et al., 1974) cannot resolve these four clusters. The detailed network topology becomes rather irrelevant to the dynamical patterns which is not very much changed when the network is randomized. This is the same for the strong synchronization regime in the 2-level model which resembles epileptic-like activity (Zemanová et al., 2006). This can be understood recalling previous analysis based on random networks where is shown that synchronization is mainly determined by the node intensity (Zhou, Motter and Kurths, 2006; Zhou and Kurths, 2006a) (see Section IV 6). Furthermore, the transition from the weak to the strong synchronization regime shares a similar picture with the Kuramoto model in complex networks (Gómez-Gardeñes et al., 2007a,b) (see Section 3 4).

Shortly after these works, a very similar structure-function relationship was observed (Honey et al., 2007). Each area of the macaque neocortex was represented by a neural mass model in the regime of spontaneous activity with complicated temporal patterns. It was shown that the functional connectivity, measured over a very long time, is closely shaped by the underlying structural connectivity as described by Zhou, Zemanová, Zamora, Hilgetag and Kurths (2006), Zemanová et al. (2006), and Zhou et al. (2007). On short time-scales, the functional connectivity changes, forming two anticorrelated clusters, similar to functional networks obtained from brain imaging data (Fox et al., 2005).

These findings support the idea that the brain is an active network, and it can generate activity by itself in the absence of external signals. Classical theories in cognitive neuroscience viewed the brain as a passive, stimul-
driven device and the spontaneous on-going activity of the brain had been regarded as background noise (Engel et al., 2001). It is still customary in data analysis to take the average signals over many trials of electroencephalographs as the event-related activation and associate them with cognitive processes. In the view of active dynamical brain networks, it has been shown that the spontaneous on-going activity imposes significant impact on the selective responses to stimuli (Engel et al., 2001; Fries, 2005). The intricate relationship between large-scale structural and functional networks revealed in these works (Honey et al. (2007); Zemanová et al. (2006); Zhou, Zemanová, Zamora, Hilgetag and Kurths (2006); Zhou et al. (2007)) will contribute to this reorientation of concepts in neuroscience. On the other hand, excessive activation and synchronization of neural networks have been found to associate with dysfunctions and disorders of the brain, such as the epileptic seizure (Stam, 2005) and the Parkinsonian disease (Tass et al., 1998). Understanding synchronization in neuronal networks of various level, especially studying the role played by the complex network topology, is crucial to elucidate how normal brains can maintain desirable levels of synchronization. It will also contribute significantly to biomedical data analysis of pathological brain activities (Lai et al., 2007), for example, the challenging task of detecting precursors that can make prediction much before the clear onset of seizures, and design suitable methods for treatments of neural diseases (Popovych et al., 2006).

B. Computer science and engineering

Complex networks and synchronization dynamics are relevant in many computer science and engineering problems. For example, in computer science, synchronization is desirable for an efficient performance of distributed systems. Sometimes, the goal of the distributed system is to achieve a global common state (consensus). Nowadays these systems are becoming larger and larger and their topologies more and more complex. On the other hand, some engineering problems also face the need of maintaining coordination at the level of large scale complex networks, for example in problems of distribution of information, energy or materials.

FIG. 30 (Color online) Connection matrix $M^4$ of the cortical network of the cat brain. The different symbols represent different connection weights: 1 (◦ sparse), 2 (● intermediate) and 3 (∗ dense). The organization of the system into four topological communities (functional sub-systems, V, A, SM, FL) is indicated by the solid lines. From Zhou, Zemanová, Zamora, Hilgetag and Kurths (2006).

FIG. 31 (Color online) The functional networks (◦) at various thresholds: $R_{th} = 0.070$ (a), $R_{th} = 0.065$ (b), $R_{th} = 0.055$ (c) and $R_{th} = 0.019$ (d). The small dots indicate the anatomical connections. From Zhou, Zemanová, Zamora, Hilgetag and Kurths (2006).

FIG. 32 (Color online) Four major dynamical clusters (◦) in the weak synchronization regime, compared to the underlying anatomical connections (●). From Zemanová et al. (2006).
The simulation of large systems are, nowadays, mainly implemented as parallel distributed simulations where parts of the system are allocated and simulated on different processors. A large class of interacting systems including financial markets, epidemic spreading, traffic, and dynamics of physical systems in general, can be described by a set of local state variables allowing a finite number of possible values. As the system evolves in time, the values of the local state variables change at discrete instants, either synchronously or asynchronously, depending on the dynamics of the system. The instantaneous changes in the local configuration are called discrete events, forming what has been coined as a parallel discrete-event simulation (PDES) (Nicol and Fujimoto, 1994). The main difficulty of PDES is the absence of a global pacemaker when dealing with asynchronous updates. This imposes serious problems because causality and reproducibility of experimental results are desired. In a conservative scheme, processes modeling physical entities are connected via channels that represent physical links in the target system. Since PDES events are not synchronized via a global clock, they must synchronize by communication between nodes.

The essentials of a PDES are: a set of local simulated times of the processors and a synchronization schedule. As the grid-computing networks with millions of processors emerge, fundamental questions of the scalability of the underlying algorithms must be addressed. The properties of a PDES to be scalable are: (i) the virtual time horizon \( \{ \tau_i(t) \}_{i=1}^{N_p} \), a set of time simulated instants in \( N_p \) processors after \( t \) parallel steps, should progress on average with a non-zero rate, and (ii) the width of the time horizon should be bounded when \( N_p \to \infty \).

Korniss et al. (2000) studied the asymptotic scaling properties of a conservative synchronization algorithm in massive PDESs where the discrete events are Poisson arrivals. They found an interesting analogy between the evolution of the simulated time horizon and the growth of a nonequilibrium surface. They showed that the steady-state behavior of the macroscopic landscape of the simulated time horizon, in one dimension, is governed by the Edwards-Wilkinson Hamiltonian (Edwards and Wilkinson, 1982).

The analogy becomes clear by interpreting the virtual times \( \tau_i \) as the height of a surface, and defining the width of the simulated times surface as the mean root square of the virtual times with respect to the mean \( \bar{\tau} \). This width provides a measure for de-synchronization

\[
\langle w^2 \rangle = \langle \frac{1}{N_p} \sum_1^{N_p} [\tau_i(t) - \bar{\tau}(t)]^2 \rangle.
\]

The problem that now is faced is how the width of the synchronization landscape scales with the number of processors. If the scaling diverges, it means that the synchronization is hardly achievable. In one- and two-dimensional regular lattices, the width of the synchronization landscape diverges with the number of processors as \( w \sim N_p^{1/d} \) where \( d \) is the dimension. This effect can be traced back to the lateral correlation length.

\[\text{FIG. 33 (Color online) Virtual time horizon snapshots in the steady state for 10000 sites in one dimension. (a) For a regular network. The lateral correlation length } \xi \text{ and width } w \text{ are shown for illustration in the graph. (b) For the SW network with } p = 0.1, \text{ the heights are effectively decorrelated and both the correlation length and the width are reduced. From Guclu et al. (2006).}
\]
ξ of the surface that also diverges with the number of processors ξ ∼ N (Barabási and Stanley, 1995). An interesting solution to this problem has been proposed by Korniss et al. (2003) and Guclu et al. (2006). It consists of adding a few random links to the regular lattice resulting in a SW structure. This structure has the effect of de-correlating the lateral length, suppressing large fluctuations in the synchronization surface (roughness), and producing finite average values of w in the large system-size limit, see Fig. 33. Moreover, the extreme height diverges only logarithmically in this limit. This latter property ensures synchronization in a practical sense in a SW topology of processors.

2. Data mining

The term data mining refers to the process of automatically searching large volumes of data for patterns that provide some useful information for classification. Miyano and Tsutsui (2007) have proposed a new method of data mining based on spontaneous data clustering using a network of locally coupled limit-cycle phase oscillators. The method is closely related to the determination of community structure via synchronization processes devised by several authors (Arenas et al., 2006b; Boccaletti et al., 2007). The idea is to encode multivariate data vectors (that are the elements of the database) into vectors of natural frequencies for an oscillators’ dynamical model, akin to the KM, expecting that the dynamics of the system will group similar data in clusters of synchronization. These interactions can be directed or undirected, fixed or mobile, constant or weighted, keeping them many of the ingredients we have been discussing in this review. Another interesting fact in this sort of problems is the existence of time delays in the communication process.

Olfati-Saber and Murray (2004) define consensus problems on general graphs. Let us consider a dynamic graph in which the connectivity pattern of the nodes can change in time. At each node, a dynamical agent evolves in time according to the dynamics

\[ \dot{x}_i = f(x_i, u_i), \quad (111) \]

where \( f(x_i, u_j) \) is a function that depends on the state of the unit \( x_i \), and on \( u_i \) that describes the influence from the neighbors. The χ-consensus problem in a dynamical graph is a distributed way to reach an asymptotically stable equilibrium \( x^* \) satisfying \( x^*_i = \chi(x(0)), \forall i \), where \( \chi(x(0)) \) is a prescribed function of the initial values (e.g., the average or the minimum values).

They present two protocols that solve consensus problems in a network of agents:

1. fixed or switching topology and zero communication time-delay:

\[ \dot{x}_i = \sum_{i,j=1}^{n} a_{ij}(t)(x_j(t) - x_i(t)), \quad (112) \]

2. fixed topology and non-zero communication time-delay \( \tau_{ij} > 0 \)

\[ \dot{x}_i = \sum_{i,j=1}^{N} a_{ij}(t)(x_j(t - \tau_{ij}) - x_i(t - \tau_{ij})). \quad (113) \]

We note that the analysis of the asymptotic behavior of such linear system is similar to the stability analysis performed in the framework of the MSF (Sect. IV).

They find very interesting results in terms of network properties. For instance, a network with fixed topology that is a strongly connected digraph (a subgraph connected via a path that follows the direction of the edges of the graph), solves the average consensus problem if and only if all the nodes of the network have the same indegree as the outdegree, i.e. \( k^\text{in} = k^\text{out}, \forall i \), as the balanced networks discussed by Belykh et al. (2006) (see Sect. IV.C). Furthermore, the performance of the network measured in terms of the speed in which the global asymptotic equilibrium state is reached, is proportional

3. Consensus problems

Consensus problems, understood as the ability of an ensemble of dynamic agents to reach a unique and common value in an asymptotically stable stationary state, have a long history in the field of computer science, particularly in automata theory and distributed computation. In many applications, like for instance cooperative control on unmanned air vehicles, formation control or distributed sensor networks, groups of agents need to agree upon certain quantities of interest (Olfati-Saber and Murray, 2004). As a result, it is important to address these problems of agreement within the assumption that agents form a complex pattern of interactions. These interactions can be directed or undirected, fixed or mobile, constant or weighted, keeping then many of the ingredients we have been discussing in this review. Another interesting fact in this sort of problems is the existence of time delays in the communication process.

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They present two protocols that solve consensus problems in a network of agents:

1. fixed or switching topology and zero communication time-delay:

\[ \dot{x}_i = \sum_{i,j=1}^{n} a_{ij}(t)(x_j(t) - x_i(t)), \quad (112) \]

2. fixed topology and non-zero communication time-delay \( \tau_{ij} > 0 \)

\[ \dot{x}_i = \sum_{i,j=1}^{N} a_{ij}(t)(x_j(t - \tau_{ij}) - x_i(t - \tau_{ij})). \quad (113) \]

We note that the analysis of the asymptotic behavior of such linear system is similar to the stability analysis performed in the framework of the MSF (Sect. IV).

They find very interesting results in terms of network properties. For instance, a network with fixed topology that is a strongly connected digraph (a subgraph connected via a path that follows the direction of the edges of the graph), solves the average consensus problem if and only if all the nodes of the network have the same indegree as the outdegree, i.e. \( k^\text{in} = k^\text{out}, \forall i \), as the balanced networks discussed by Belykh et al. (2006) (see Sect. IV.C). Furthermore, the performance of the network measured in terms of the speed in which the global asymptotic equilibrium state is reached, is proportional

3. Consensus problems

Consensus problems, understood as the ability of an ensemble of dynamic agents to reach a unique and common value in an asymptotically stable stationary state, have a long history in the field of computer science, particularly in automata theory and distributed computation. In many applications, like for instance cooperative control on unmanned air vehicles, formation control or distributed sensor networks, groups of agents need to agree upon certain quantities of interest (Olfati-Saber and Murray, 2004). As a result, it is important to address these problems of agreement within the assumption that agents form a complex pattern of interactions. These interactions can be directed or undirected, fixed or mobile, constant or weighted, keeping then many of the ingredients we have been discussing in this review. Another interesting fact in this sort of problems is the existence of time delays in the communication process.

Olfati-Saber and Murray (2004) define consensus problems on general graphs. Let us consider a dynamic graph in which the connectivity pattern of the nodes can change in time. At each node, a dynamical agent evolves in time according to the dynamics

\[ \dot{x}_i = f(x_i, u_i), \quad (111) \]

where \( f(x_i, u_j) \) is a function that depends on the state of the unit \( x_i \), and on \( u_i \) that describes the influence from the neighbors. The χ-consensus problem in a dynamical graph is a distributed way to reach an asymptotically stable equilibrium \( x^* \) satisfying \( x^*_i = \chi(x(0)), \forall i \), where \( \chi(x(0)) \) is a prescribed function of the initial values (e.g., the average or the minimum values).

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to $\lambda_2(\tilde{G})$ where $\tilde{G}$ is the mirror graph induced by $G$, which is defined as an undirected graph with symmetric adjacency matrix $\tilde{a}_{ij} = (a_{ij} + a_{ji})/2$.

For a switching topology, they find that if the dynamics of the network is such that any graph along the time evolution is strongly connected and balanced then the switching system asymptotically converges to an average consensus. Concerning time communication-delays, the important result is that if all links have the same time-delay $\tau > 0$, and the network is fixed, undirected and connected, the system solves the average consensus if $\tau \in (0, \pi/2\lambda_N)$. In this case, in a similar way as discussed in previous applications, there are two tradeoff issues that can be related to problems of network design: one concerns the robustness of the protocol with respect to time-delays, and the other to communication cost.

When applying this framework to a certain class of networks, Olfati-Saber (2005) finds that the speed of convergence, as the inverse of $\lambda_2$ (as was also found in synchronization problems (Almendral and Díaz-Guilera, 2007)), can be increased by orders of magnitude by simply rewiring a regular lattice, while this change has a negligible effect on $\lambda_N$, which measures the robustness to delays of the system. This can be understood by the eigenvalues of the SW network in Eq. (67) as compared to the regular networks in Eq. (64). Some other variations can be found in the recent literature; e.g., Wu et al. (2007) analyze several network models with physical neighborhood connectivity and depending on the precise rules they discuss on the performance and the robustness of the system.

Due to its importance as an application in computer science, consensus problems are interesting by themselves. But understanding its linear dynamics can be also of great importance in the behavior of complex populations of units that evolve according to more complex non-linear dynamics, as it happens in many synchronization problems.

4. Communication networks

Another emerging line of research can be found in the field of synchronizing wireless communication networks. Wireless ad-hoc networks are telecommunication networks that are created by devices equipped with a short-range wireless technology, such as WiFi or Bluetooth. Unlike wired networks, these networks can be created on the fly to perform a task, such as information routing, environmental sensing, etc. (Hekmat, 2006) Furthermore, the topology of these networks can be changed dynamically to achieve a desired functionality. With the rapid growth of the number of portable devices and the increased popularity of wireless communication, it is expected that these types of networks will play a key role as a building block of the next generation Internet. On the other hand, from the perspective of fundamental research, these systems provide a clear-cut example of highly dynamic, self-organizing complex networks.

Synchronization processes in wireless systems naturally arise in two problems. On the one hand, routing and information flow algorithms require synchronization of the clocks of the nodes of the wireless network to establish a universal coordinated time. In a very recent work, Díaz-Guilera et al. (2008) study synchronization of Kuramoto oscillators in random geometric graphs. They consider a wireless system in which the connections vary at a time scale much shorter than the time scale associated to the dynamics, and hence the network is static. Nodes correspond to devices that have a finite transmission range, and are linked to those nodes that are located within the range. This procedure gives rise to a two-dimensional random geometric graph, which are characterized by a high clustering coefficient and a very large average shortest path length, as compared to ER graphs with the same number of nodes and links. The remarkable result is that this type of network is very hard to synchronize, both in terms of the stability of the synchronized state and in terms of the time needed to reach the completely synchronized state. Although they are very homogeneous, the smallest non-zero eigenvalue of the Laplacian matrix is very low. However, just by rewiring a small fraction of the links at random synchronizability is clearly improved, which from a practical point of view is a very interesting issue.

On the other hand, communication channels have a finite bandwidth so that the access times of different users should be desynchronized when their number is large. The latter, on its turn, depends on the density of wireless devices accessing the available resources, which is determined by the topology of the underlying ad-hoc network, and on the dynamical details of the signal transmission process. Given the entangled structural and dynamical complexity of synchronization phenomena in these wireless settings, it is expected that the network formalism and some of the techniques reviewed here will greatly contribute to the development and deploying of more efficient algorithms and protocols for wireless communications in the near future. This is what can be found in the model proposed by Degesys et al. (2007), a biologically inspired algorithm for desynchronization in a single-hop network. They consider a set of $N$ nodes (oscillators) that generate events with a common period. The nodes rearrange their phases, just considering the times in which neighboring (in time) units fire, in such a way that the events become spaced at intervals $T/N$. The final state then corresponds to what is usually called a round-robin schedule. In this way, the use of the bandwidth without collisions between messages is optimized. Inspired by this result, (Díaz-Guilera and Arenas, 2008) considered the units to be Kuramoto oscillators with a common frequency. Introducing some dephase angle in the sinus function and coupling pairs of units along a closed chain, the authors find new stable configurations different from the completely synchronized state. Some of these configurations correspond also to the round-robin
schedule, which turns out to be very robust under the addition or deletion of nodes from the system. In this simple setting we can see how important is the local pattern of connections. There is no doubt that extension of the proposed models to more complex topologies, as for example within some transmission range as considered in the previous paragraph, gives rise to much more complex, and interesting, global behaviors of the system of oscillators.

5. Material and traffic flow

In our globalized and connected world not only information flows, but also material transportation is an important issue that needs to be performed in an efficient way. In many material flow systems coordination of tasks in a parallel way is essential for a proper functioning but difficult to achieve. Typical examples of this are crossroads in road traffic (Nagatani, 2002) and supply chains in production processes (Helbing, 2003). Recent work on supply networks has shown how to treat them as physical transport problems governed by balance equations and equations for the adaptation of production speeds. Although the nonlinear behavior is different, the linearized set of coupled differential equations is formally related to those of mechanical or electrical oscillator networks (Helbing et al., 2004). Whereas traditional optimization techniques can be used to setup single nodes, the inherent topological complexity makes maintenance of coordination at network-wide level to be practically unsolvable by these methods. Furthermore, robustness and flexibility, due to continuous changes in demand and failures, are also required for an efficient transportation.

It is obvious that there is an analogy between material transportation in networks and the flow of chemical substances and nutrients in biological organisms, where synchronization dynamics plays an important role. Lämmer et al. (2006) have proposed a decentralized control model for material flow networks with transportation delays and setup-times, based on phase-synchronization of oscillatory services at the network nodes.

A material transport network is a directed and weighted graph where the flow of material at nodes is conserved. Subsets of links are active at different times, and this makes that the activity of the node is periodic and one can map a continuous phase variable \( \theta(t) \) to a discrete service state \( M : \theta(t) \to s(t) \). While the phase angle \( \theta \) of the oscillator modeling the intersection varies from 0 to \( 2\pi \) at a rate \( \omega \), all states \( s \) are sequentially activated. To achieve a coordination of the switching states on a network-wide level, they propose a suitable synchronization mechanism.

The authors apply this formalism to the control of traffic lights at intersections of road networks. A single traffic light intersection is modeled by an oscillator where the continuous phase maps to a set of states corresponding to green lights (see Fig. 34). The maximum frequency of the oscillator dynamics is calculated in terms of the load at the different lanes and the setup-time. Global coordination of the network is achieved by synchronizing the local phases and frequencies, requiring to reach a phase-locked state where the phase difference between neighboring sites is fixed. They suggest a coupling on two different time scales:

- Adaptation of the phase \( \theta \) a la Kuramoto:

\[
\dot{\theta}_i = \min \left\{ \omega_{i_{\text{max}}} + \frac{1}{T_0} \sum_{j \in \Gamma_i} \sin (\theta_j(t) - \theta_i(t)) \right\}
\]

where \( \omega_i \) is the intrinsic frequency. As long as \( \omega_i < \omega_{i_{\text{max}}} \), \( \theta_i \) tries to adjust to the neighboring phases. The constant \( T_0 \) corresponds to the typical time scale for this adaptation.

- A second decentralized coupling can be used to increase the intrinsic frequencies to approach the possible maximum within a slow time scale:

\[
\dot{\omega}_i = \frac{1}{T_{\omega}} \left( \min_{j \in \Gamma_i} \{ \omega_j(t) \} + \Delta \omega - \omega_i(t) \right).
\]

Here the constant parameter \( \Delta \omega \) provides a linear drift towards higher frequencies.

Under these assumptions two dynamical regimes are possible (see Fig. 35). Starting with a random initial
condition (left), the system quickly settles into a state with growing common frequency and vanishing phase-differences. As soon as the maximum common frequency is found, the system enters the other state with a locked common frequency and phase-differences exponentially converging towards constant values (right).

The extension of the analysis performed on such a simple setting to more realistic complex transportation networks is very promising but challenging (Helbing et al., 2005).

### 6. Power-grids

Finally, we mention an engineering problem that might seem at the first sight unrelated to synchronization phenomena. Power grids are physical networks of electrical power distribution lines of generators and consumers. In the pioneering paper by Watts and Strogatz (1998) it was already reported that the power-grid constitutes one of the examples of a self-organized topology that has grown without a clear central controller. This topology is indeed very sensitive to attacks and failures. From its topological point of view there are several analyses on power-grids in different areas of the world (Chassin and Posse, 2005; Crucitti et al., 2004a) and some models have been proposed to deal with the cascading process of failures (Crucitti et al., 2004b; Sachtjen et al., 2000; Scirè et al., 2005).

The principles of electricity generation and distribution are well known. Synchronization of the system, i.e. every station and every piece of equipment run on the same clock, is crucial for its proper operation. Cascading failures related to de-synchronization can lead to massive power blackouts (Symetricom, 2003).

Then power production, transmission, and consumption represents a dynamical problem and the power grid can be seen as an example of a system of oscillators (Strogatz, 2003b). Along this line, Filatrella et al. (2007) proposed to study synchronization of the various components of the power grid by means of coupled phase oscillators:

\[
\dot{\theta}_i = -\alpha \theta_i + P_i + \sum_{j \in \Gamma(i)} P_{ij}^{\text{max}} \sin(\theta_j - \theta_i),
\]

where \( P_{ij} \) accounts for the type of coupling between the different units (generators, consumers, ...). They analyze under which conditions the system is able to restore to a stable operation after a perturbation in simple networks of machines and generators. We believe that the principles summarized in this review will be very useful for the understanding of the impact of the complex topology of the power grids on their synchronization and robustness.

### C. Social sciences and economy

In the last decades, social sciences and economy have become one of the favorite applications for physicists. In particular, tools and models from statistical physics can be implemented on what some people has called social atoms, (Buchanan, 2007) i.e. unanimated particles are replaced by agents that take decisions, trade stocks or play games. Simple rules lead to interesting collective behaviors and synchronization is one of them, because some of the activities that individual agents do can become correlated in time due to its interaction pattern, which, in turn, is clearly another example of the complex topologies considered along the review.

#### 1. Opinion formation

One of such problems is the study of opinion formation in society. The underlying idea is that individuals (or agents) have opinions that change under the influence of other individuals giving rise to a sort of collective behavior, grouping together a macroscopic part of the whole population with similar social features (Boccaletti et al., 2006). Therefore, the main goal is to figure out whether and when a complete or partial consensus can emerge out of initially different opinions, no matter how long it takes for the consensus to be reached.

In general, the formation of a collective opinion about a certain matter is not equivalent to a transition to some kind of synchrony, but rather to a transition to an absorbing state. However, a recently proposed model (Pluchino et al., 2005) makes explicit use of a modified KM (see Sect. III) and thus in this case the formation of groups of opinions can be thought of a synchronization process. Agreement models deal with \( N \) individuals characterized by an opinion \( x_i \) (either an integer or real number) and a network of contacts that drives the dynamics of opinion formation through deterministic rules (Boccaletti et al., 2006). Pluchino et al. (2005) considered the case in which opinions are neither bounded nor periodic, but that two initially different opinions can also diverge when time goes on. Moreover, they also take into account that two
quite different individuals tend to interact less by assuming that the coupling between these two individuals is a decreasing function of their opinion differences. Finally, Pluchino et al. (2005) considered the situation in which the main source of heterogeneity is not given by the initial positions, but by different rates of changing individuals’ opinions.

Taking all the previous statements into account, Pluchino et al. (2005) proposed the following governing equations for the dynamics of the rate of change of opinions

\[ \dot{x}_i = \omega_i + \frac{\sigma}{N} \sum_{j=1}^{N} \alpha \sin(x_j - x_i)e^{-\alpha|x_j - x_i|}, \quad (117) \]

where \( x_i(t) \in (-\infty, +\infty) \) is the opinion of the \( i \)-th individual at time \( t \) and the \( \omega_i's \) are the natural or intrinsic opinion changing rates. Note that the interactions are assumed to be all-to-all, though the model can be directly generalized to any other topology. Moreover, the \( \omega_i's \) are drawn from a distribution \( g(\omega) \) centered at \( \omega_0 \) with the same properties as in the KM case (Sect. III).

On the other hand, as opposite to the KM, opinions are not periodic anymore, so that a new order parameter is introduced as

\[ R(t) = 1 - \sqrt{\frac{1}{N} \sum_{i=1}^{N} (\dot{x}_i(t) - \bar{X}(t))^2}, \quad (118) \]

with \( \bar{X}(t) \) being the average of \( \dot{x}_i(t) \) over all individuals. From Eq. (118), it follows that \( R = 1 \) if complete synchronization is achieved and \( R < 1 \) when only partial synchronization occurs.

Numerical simulations of the model show that there is a phase transition from incoherence to synchrony at a well defined critical coupling \( \sigma_c \). Pluchino et al. (2005) argued that when \( \sigma < \sigma_c \), the society can be thought of as being formed by isolated, non-interacting cultures or groups of opinions, since mixture or agreement is not achievable. On the contrary, when \( \sigma \gg \sigma_c \), the system fully synchronizes, giving rise in a social context to a polarized or globalized society where social and cultural differences are constrained into a single way of thinking, notwithstanding the different tendencies to changes of the individuals. Finally, the authors reported that bipolarity is only possible if \( \sigma \sim \sigma_c \), although in this case the model shows a rich behavior depending on the way initial opinions are assigned (Pluchino et al., 2005).

2. Finance

When reading the economic news, it is not difficult to identify the existence of economic cycles in which Gross Domestic Products (GDP’s), economic sectors, or stock options raise and fall. Most of the time this does not happen for isolated countries, sectors or options but it occurs in quite a synchronized way, although some delays are noticeable.

Within the framework of the current review, we are focusing on synchronization in complex networks, and this is what we can identify in many economical sectors: there exists a complicated pattern of interactions among companies or countries and the dynamics of each one is quite complex. But, in contrast to many networks with a physical background, here we neither know in detail the node dynamics nor its connectivity pattern. In this situation it is useful to look at the problem from a different angle. By analyzing some macroscopic outcomes, we get some insight into the agents’ interactions.

The application of networks concepts, mainly that of trees, to economical systems dates back to the pioneering work by Mantegna (1999), who found a hierarchical arrangement of stocks through the study of the correlation returns.

Recently, Onnela et al. (2003) have taken a similar approach to analyze the dynamics of markets. They look at the daily closure prices for a total of \( N = 477 \) stocks traded by the New York Stock Exchange over a period of 20 years, from Jan 02, 1980 to Dec 31, 1999. The data is smoothed by looking at time windows of given width. As is usually done in the analysis of financial data, the measured quantity is the logarithmic return of the stocks, defined as

\[ r_i(t) = \ln P_i(t) - \ln P_i(t - 1), \quad (119) \]

where \( P_i(t) \) is the closure price of stock \( i \) at time \( t \). To quantify the degree of synchronization of the data, they use the equal time correlation between assets

\[ \rho_{ij}(t) = \frac{\langle r_i(t)r_j(t) \rangle - \langle r_i(t) \rangle \langle r_j(t) \rangle}{\sqrt{[\langle r_i^2(t) \rangle - \langle r_i(t) \rangle^2][\langle r_j^2(t) \rangle - \langle r_j(t) \rangle^2]}} \quad (120) \]

where \( \langle ... \rangle \) stands for a time average over the consecutive trading days.

From these correlations the asset tree is constructed. The distance between assets is defined as

\[ d_{ij}(t) = \sqrt{2(1 - \rho_{ij}(t))}. \quad (121) \]

The minimum spanning tree is a simply connected graph with \( N - 1 \) edges, such that the sum of all the distances between connected nodes in the graph \( \sum d_{ij}(t) \) is minimum.

This procedure generates a time sequence of asset trees that can be interpreted as a sequence of evolutionary steps of a single dynamics asset tree. For instance, one can identify the leading asset, that, in most instances, corresponds to General Electric. Such a reduction of the whole set of data retains most of the salient features of the stock market. It is a remarkable fact that during crisis periods the markets are very strongly correlated. In terms of the tree its average length is reduced and the tree is very tightly packed. By
reducing the time window, the location of the smallest tree converges to the Black Monday (October 19, 1987).

It is clear that a hidden pattern of interactions between the assets is responsible for such a synchronized behavior and that during crashes the interactions are strengthened. The message here is that the degree of synchronization, quantitatively described in terms of the asset correlation, is an indirect measure of the existence of strongly connected agents in financial markets.

3. World Trade Web

The World Trade Web (WTW) is another example of an empirically known network in economy constructed by collecting data from available databases. There are several studies that have focused on the static complex nature of the links between countries (Serrano and Boguña, 2003) and also the evolution of the statistical properties of the network (Garlaschelli and Loffredo, 2005). But, as expected, the trade volume between countries strongly depends on the internal state of the nodes that form the network. In this case the state of the nodes corresponds to the GDP of the countries. Garlaschelli et al. (2007) have analyzed the interplay between the topology of the WTW and the dynamics of the GDP’s.

In none of the previous studies any reference is made to the precise dynamics of the countries economies. As we have stated before, we find what are usually called economic cycles. Economies rise and fall, although they mainly raise, but without a constant rate, following rather unpredictable evolutions in time. Due to globalization effects, all economies are strongly correlated and they will tend to follow a common trend. In our framework, we can say that following a similar time evolution, economies are synchronized. This cycle synchronization of economies is a topic of current interest in the economic literature; Calderón et al. (2007) perform a comparative analysis between developing and industrial countries, finding that the correlations within the first group are positive but smaller than within the second group. Another example is found in Li et al. (2003) where correlations between countries are measured in a similar way as the financial series reported above. By taking into account that the US is the largest economy, and also the biggest node in the WTW, they look at the degree of synchronization between 22 developed countries and the USA (see Fig. 36).

Another clear effect is that particular economies are tightly connected because of economical agreements or dependence on particular sectors. In the language of communities this stronger relation can be understood as the existence of communities in the overall structure of the world economy.

Along these lines we have observed that there is a tight relationship between synchronization of economic cycles in terms of the GDP and the topological structure of the WTW. The important question raised here is if the structure of the network that can be constructed from the empirical data on the cycles correlations, as is done with the finance data, can be mapped into the WTW also empirically constructed from the world trade transactions.

D. Perspectives

Despite the number of applications revised so far, there is no doubt that synchronization phenomena on complex topologies is still a new emerging field in which many efforts will be invested in the future. We have mentioned some problems where the tools and methodologies revised in the first Sections have not been applied yet. It will perhaps take more time to transfer the acquired theoretical knowledge to the application level. In other cases, the available studies deal with networks of just a few elements. An important question for the near future is how to extend these results to systems of many elements and how the entangled structural and dynamical complexity inherent to such systems contribute to the synchronization of the system’s elements.

On more general grounds, given that complex networks have become a far-reaching interdisciplinary topic and that synchronization processes are present in a wide variety of fields, it would not be surprising that new applications emerge guided by further theoretical developments in the subject. For instance, many real systems may not operate at the fully synchronized state, where most of the theoretical approaches revised are effective. In this sense, an important unsolved challenge is how to characterize partial synchronization and how to exploit and control partially synchronized states in contexts such as wireless communication systems, where the information transfer capacity is regulated by these states.
VI. CONCLUSIONS

Through the current review we have outlined the state of the art towards a theory of synchronization in complex networks. We emphasize the word theory, because, up to now, physicists have made an effort of characterization that certainly deepened our understanding of the complex connectivity of natural and manmade networks, however, we cannot yet state that we have a theory of complex networks. We emphasize the word theory, because, to this specific end, the complex network substrate must be enriched and entangled to the functioning of the system, i.e., to the dynamics run on top of it.

The phenomenon of synchronization is one of the paradigmatic observations in different dynamical systems. It is at the heart of some biological processes, and according to the wide variety of applications presented here, it is a plausible abstraction for many other processes in different contexts. The natural approximation to synchronization, from the simplicity of the Kuramoto model, has been explored, and even in this case an intricate set of questions concerning the uniformity of the equations proposed, or the nature of the critical behavior at the onset of synchronization still have to be definitely settled. The main results, however, have helped to understand the nature of the relationship between the topology of interactions and the synchronization of phase oscillators. We foresee the importance of these results in the basis of a theory of neural dynamics of the brain. Although neural dynamics is far more complex than the phase representation reviewed, main features that describe the path towards synchrony in complex networks have been already stated and can thus be considered as a good starting point. However, regarding the subjects revised here, one easily realizes that the work is far from complete and many questions are in the air, e.g., the evolution of synchronization in evolving topologies, the effect on the phenomenon of several kind of disorder, time delays, the presence of noise, etc.

The other main contribution to the problem comes from the MSF formalism. The elegant structure of the formalism, designed for linear systems or nonlinear systems close to the synchronization state, allows us to make theoretical predictions independent of the specificities of the dynamics. This general framework has been deeply studied recently, and provides one of the few mechanisms that allow to make predictions about the evolution of synchronized systems as a function of its specific topology.

We think that the exploration of new mathematical objects, able to merge the information provided by specific dynamics (as for example the Kuramoto model) along to the whole process towards synchronization, together with the fine description of the dynamics near the synchronization manifold, should be the focus of intense research if we aim to provide a general theory of synchronization processes in complex networks. On the other hand, the myriad of applications that can be cast into mathematical models equivalent to those presented along the review, indicates an explosion of activity in different disciplines that will use the conceptual framework of synchronization processes in networked systems as a fundamental playground for understanding its dynamical behavior.

Acknowledgments

We thank all those that have actively collaborated with us along the years in the subject. In any case, we specially thank J. García-Ojalvo, C. J. Pérez-Vicente, J. Schmidt, M. Thiel, for having carefully read the manuscript and for their enlightening comments. A.A. acknowledges support by the Director, Office of Science, Computational and Technology Research, U.S. Department of Energy under Contract No. DE-AC02-05CH11231. A.A. and A.D.-G. also thank funding from Spanish Government (BFM-2003-08258 and FIS-2006-13321) and Generalitat de Catalunya (2005SGR00889). JK acknowledges support from the EU Network of Excellence BIOSIM (Contract No. LSHB-CT-2004-005137), the EU Network BRACCIA, the DFG SFB 555, and the BMBF GoFORSYS. Y.M. is supported by MEC through the Ramón y Cajal Program, the Spanish DICYT Projects FIS2006-12781-C02-01 and FIS2005-00337, and by the European NEST Pathfinder project GABA (contract 043309).
aneous Order (Hyperion, New York, NY, USA).
Symetricom, 2003, “How time finally caught up with the power grid. white paper”,