



# Synchronization in Complex Networks

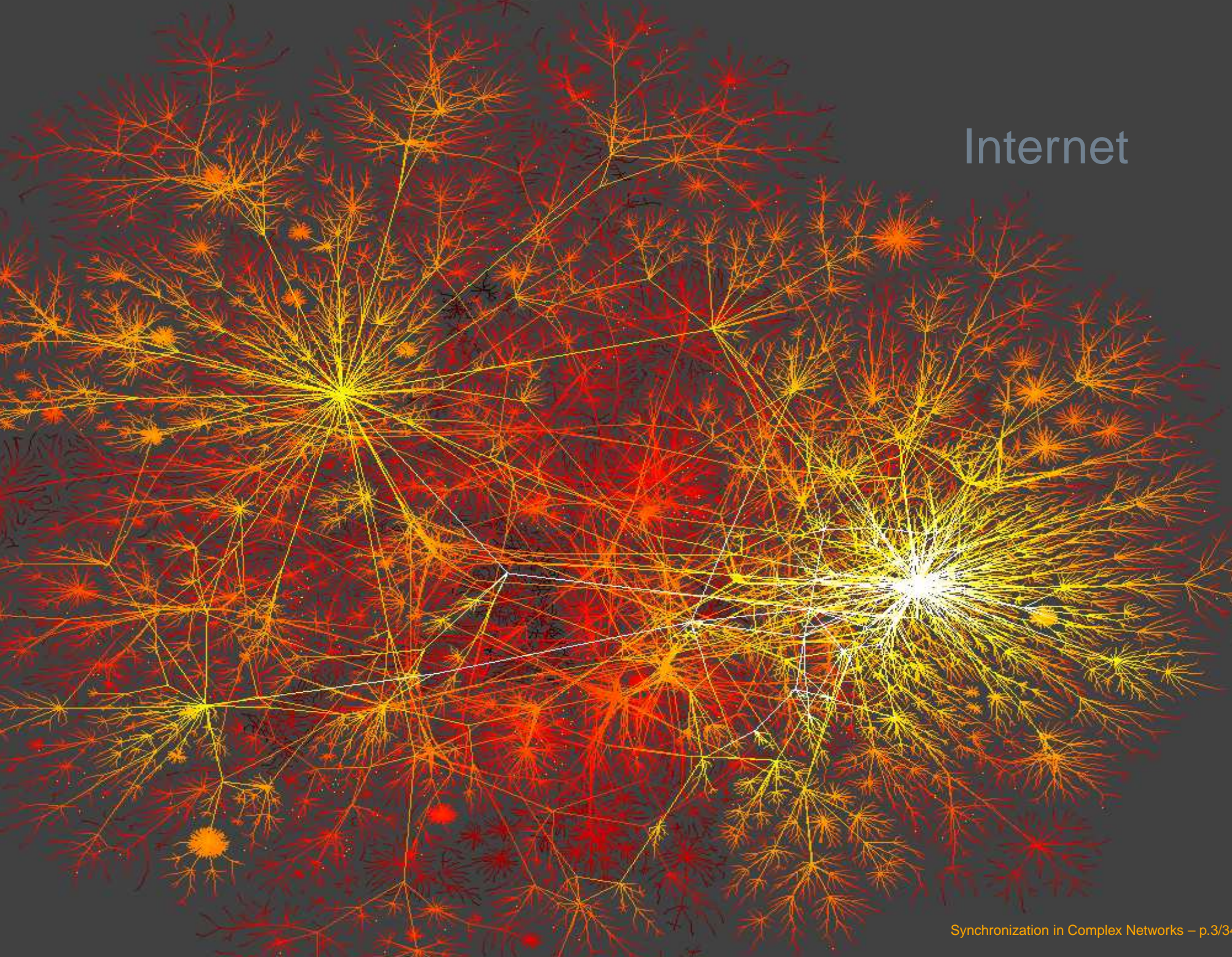
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State University

# Main collaborators

- Frank Hoppensteadt, then ASU now NYU
- Adilson Motter, then ASU now Northwestern U
- Takashi Nishikawa, then ASU now Clarkson U
- Liang Huang, ASU
- Robert A. Gatenby, U. Arizona

# Internet



# Network Synchronization

- The celebrated classical Kuramoto model (1984) - globally coupled network of phase oscillators.
- Synchronization in **complex** networks - highly relevant to biological functions, large-scale parallel processing, coordination in technological systems, etc.

# Relevance to Systems Biology

- Organizing biological information using the network idea has been fundamental to utilizing various systems-level approaches to understanding biological functions.
- **Synchronizability** of oscillator networks, such as the networks of cortical neurons, is thought to play an important role for the functionality of the network.
- Complex topology is common in biological networks.

# Synchronization in Complex Networks

- X.-F. Wang and G. Chen, “Synchronization in small-world dynamical networks,” *Int. J. Bifurcation Chaos* 12, 187 (2002).
- M. Barahona and L. M. Pecora, “Synchronization in small-world systems,” *Phys. Rev. Lett.* 89, 054101 (2002).
- T. Nishikawa, A. Motter, Y.-C. Lai, and F. Hoppensteadt, “Heterogeneity in oscillator networks: Are smaller worlds easier to synchronize?” *Phys. Rev. Lett.* 91, 014101 (2003).

# Oscillator-Network Approach

$$\dot{x}_i = F(x_i) - \varepsilon \sum_{j=1}^N G_{ij} H(x_j)$$

$G$  - coupling matrix, e.g.,

$$G_{ij} = \begin{cases} -1/k_i, & \text{if nodes } i \text{ and } j \text{ interact with each other} \\ 1, & \text{if } i = j \\ 0, & \text{otherwise,} \end{cases}$$

- $\sum_{j=1}^N G_{ij} = 0$  for all  $i$
- How to choose  $G$  to achieve stronger network synchronizability has been an active topic of research.

# Stability of Synchronous Solution

- For **any** solution  $s(t) \in \mathbb{R}^n$  of  $\dot{x} = F(x)$  (periodic, chaotic, or else),  $x_i(t) = s(t)$  for all  $i$  is a **synchronized solution** for the whole oscillator network.
- Stability of the solution is determined by linearizing and diagonalizing the equations, after which it takes the form

$$\dot{y}_i = [DF(s(t)) - \varepsilon \lambda_i DH(s(t))] y_i, \quad i=1, \dots, N$$

where  $\lambda_1 = 0 \leq \lambda_2 \leq \dots \leq \lambda_N$  are the eigenvalues of  $L$ .

- Fujisaka and Yamada, 1983, 1986.



# Transverse Stabilities

- $\lambda_1 = 0$  corresponds to the **synchronization manifold**:

$$\dot{y}_i = [DF(s(t))]y_i,$$

where  $DF$  is the Jacobian matrix of node dynamics.

- $0 \leq \lambda_2 \leq \dots \leq \lambda_N$  correspond to  $(N - 1)$ -dimensional transverse subspace. Let  $K \equiv \varepsilon \lambda_i$  be the normalized coupling parameter. **Transverse stabilities** are determined by

$$\dot{y}_i = [DF(s(t)) - KDH(s(t))]y_i, \quad i=2,\dots,N$$

# Master Stability Function

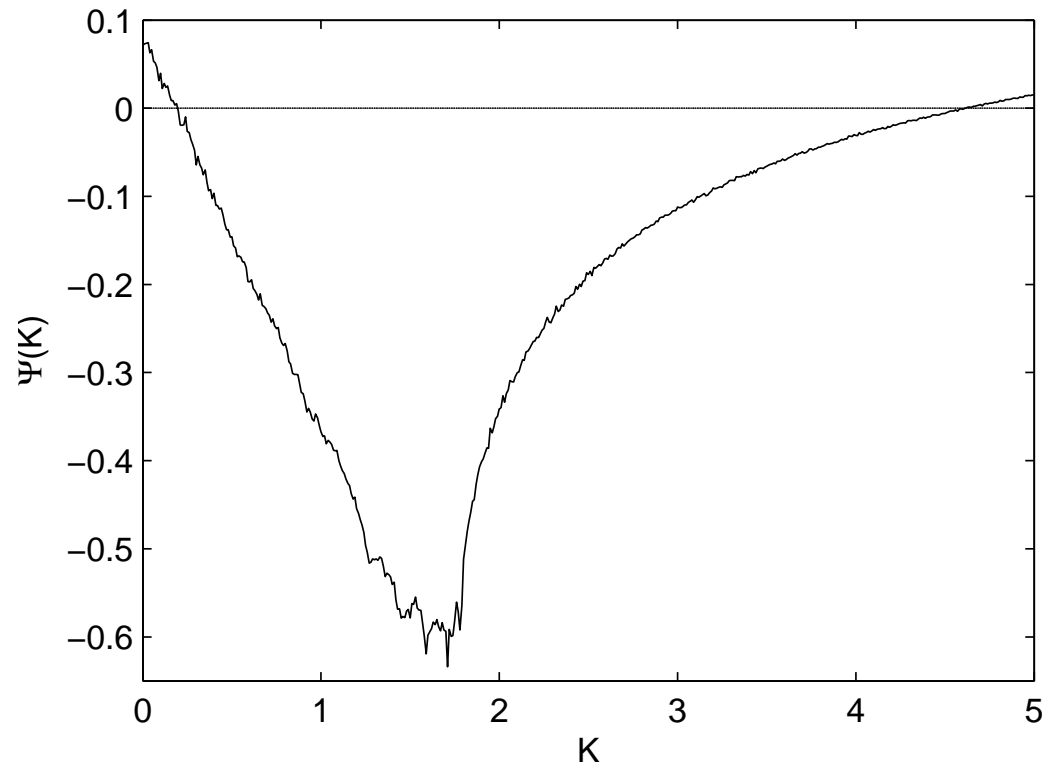
- The largest Lyapunov exponent  $\Psi(K)$  (= the Floquet exponent, if periodic) for the system

$$\dot{y} = [DF(s(t)) - KDH(s(t))]y$$

is called the **master stability function**. Pecora and Carroll (1998).

- Physically realizable synchronization requires  $\Psi(K) < 0$ .
- If  $\Psi(K) < 0$  for  $K_1 < K < K_2$ , since  $K \equiv \varepsilon \lambda_i$  ( $i = 2, \dots, N$ ), network synchronization occurs if  $\varepsilon \lambda_2 > K_1$  and  $\varepsilon \lambda_N < K_2$ .

# Example of Master-Stability Function



- Rössler oscillator:

$$\mathbf{F}(\mathbf{x}) = [-(y + z), x + 0.2y, 0.2 + z(x - 9)]^T.$$

- $K_1 \approx 0.2$  and  $K_2 \approx 4.6$ . Say  $\varepsilon = 0.5$ . Network synchronization requires  $\lambda_2 > 0.4$  and  $\lambda_N < 9.2$ .

# Generic MSF?

- Question raised by Lou Pecora at the September 07 Aberdeen Chaos Conference.
- Recent work at ASU: analysis and computation of MSFs for known chaotic oscillators.
- Systems studied so far: Rössler oscillator, Lorenz oscillator, Chua's circuit, Chen's oscillator, Duffing's oscillator, and van der Pol oscillator.
- For each system, examined all 9 common and experimentally implementable coupling configurations.

# Generic MSF?

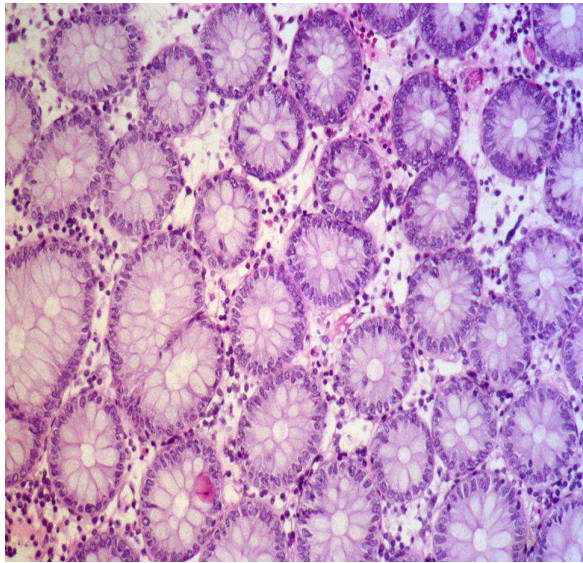
- **Results:** there always exists a coupling configuration for which the MSF is negative in a finite parameter interval, regardless of system details.
- L. Huang and Y.-C. Lai, “Generic master-stability function in chaotic systems,” to be published.
- L. Huang, Y.-C. Lai, and R. Gatenby, “Dynamics-based scalability in complex networks,” to be published.

# Synchronization in Scale-Free Networks

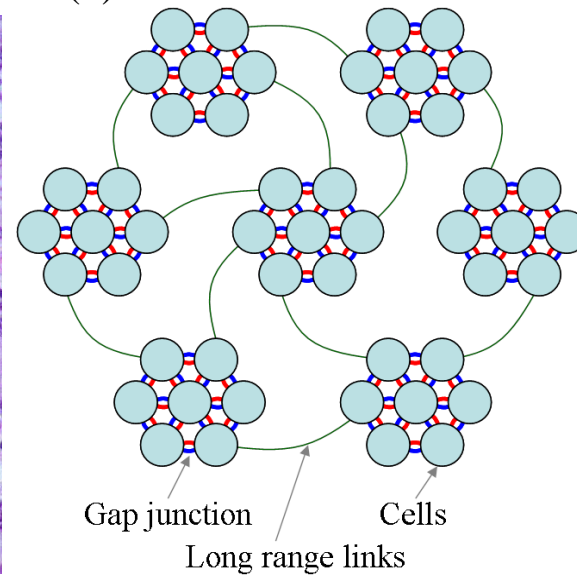
- Smaller average network distance generally does **not** imply enhanced synchronizability.
- Heterogeneity in the degree distribution is an obstacle for synchronization. Thus, **some degree of homogeneity is expected** in networks for which synchronizability is essential (e.g., neuronal networks) even if network distance is not optimal.
- Heterogeneity of load distribution appears to be responsible for the counter-intuitive effect.
- T. Nishikawa, A. Motter, Y.-C. Lai, and F. Hoppensteadt, Phys. Rev. Lett. **91**, 014101 (2003).  
**[202 - Google Scholar]**

# Cell Organization in Normal Tissue

(a)



(b)



- Microscopic image of the colon (crypts);
- Local communication subnetworks based on gap-junction conductance coupling;
- Long-range diffusive interactions based on specific cell-membrane receptors.

# Complex Clustered Networks in Biology

- **Metabolic networks**

E. Ravasz, A. L. Somera, D. A. Mongru, Z. Oltvai, and A.-L. Barabási, “Hierarchical organization of modularity in metabolic networks,” *Science* 297, 1551 (2002).

- **Protein-protein interaction networks**

V. Spirin and L. A. Mirny, “Protein complexes and functional modules in molecular networks,” *Proc. Natl. Acad. Sci. USA* 100, 12123-12128 (2003).



# Clustered Networked Systems in Engineering

Earth Simulator

Centralized single-stage crossbar



Blue Gene/L

Distributed switching 3D torus topology



ASCI Purple

Multilevel fat-tree topology



Interconnection network

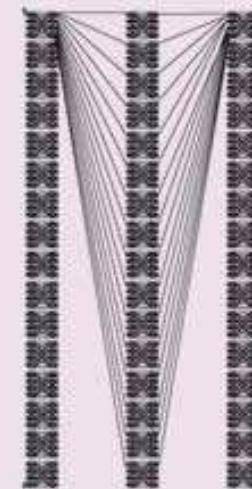
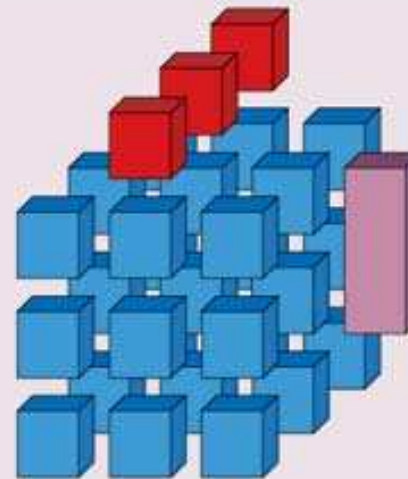
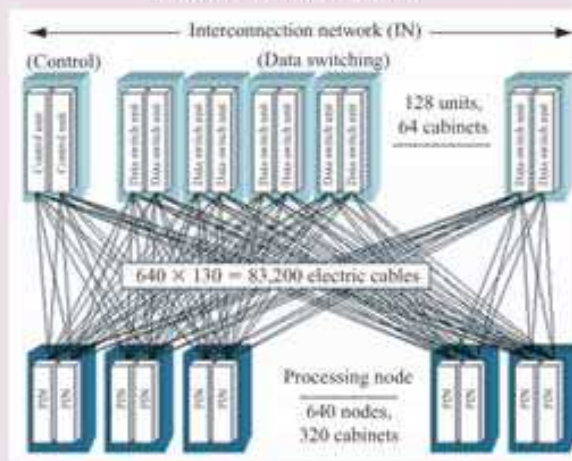
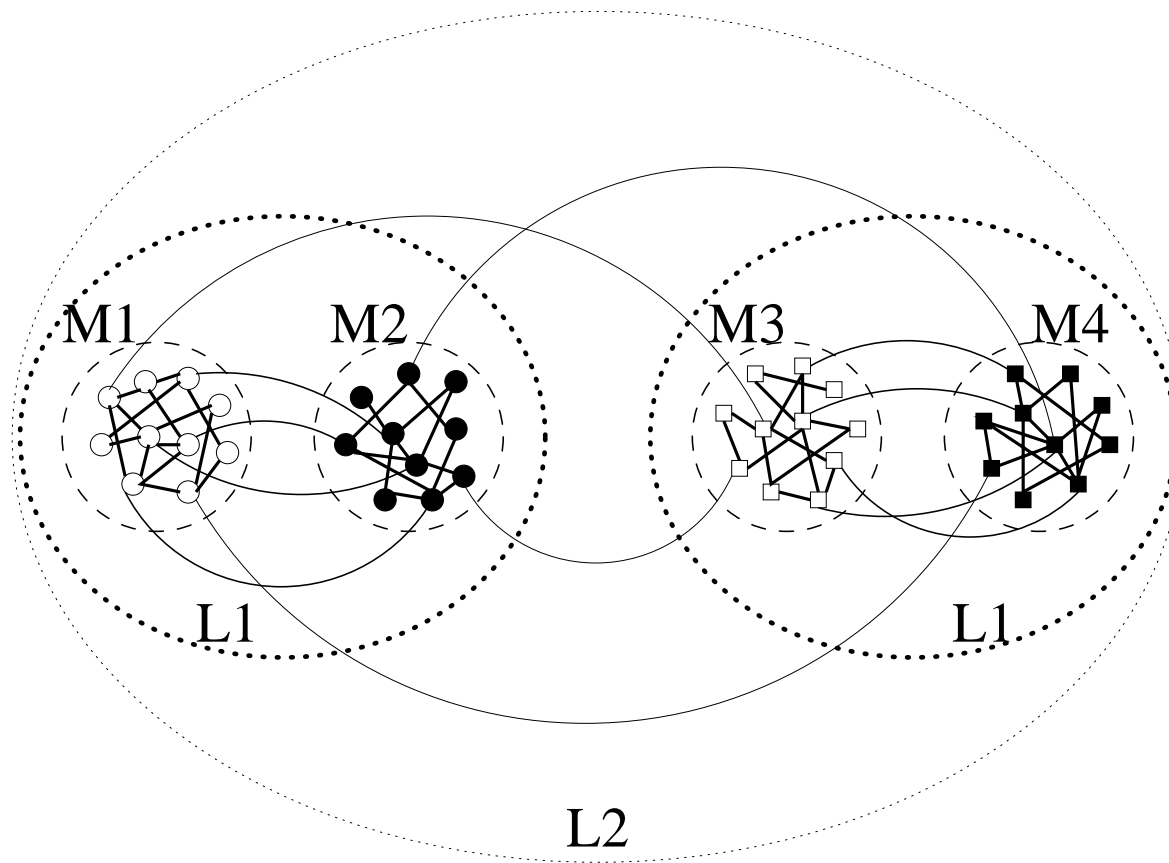


Figure 10

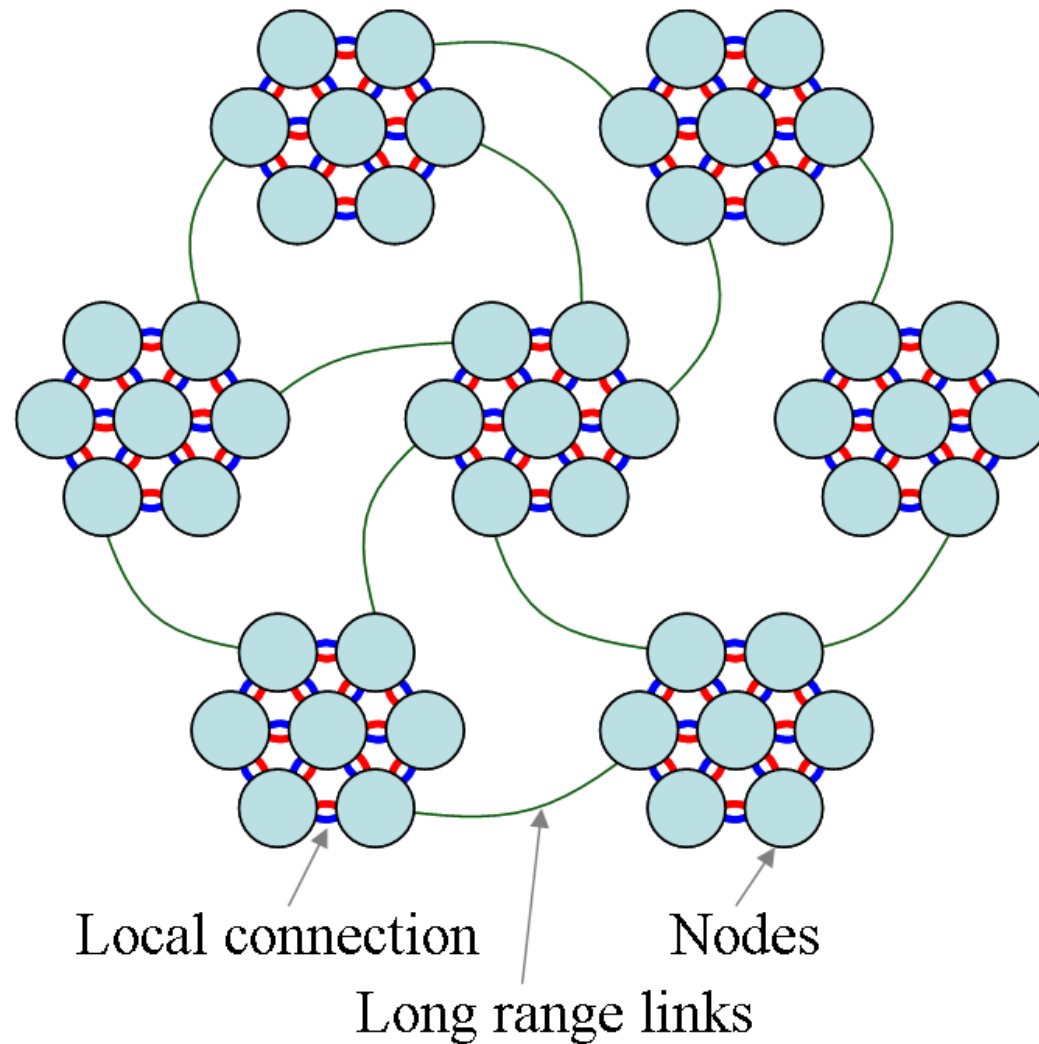
Clusters and massively parallel machines: Earth Simulator, Blue Gene/L, and ASCI Purple.

# Clustered Topology in Social Networks



- W. W. Zachary, J. Anthropol. Res. 33, 452 (1977).

# Complex clustered Networks



# Optimizing Synchronization

- Two key parameters of interest:
  1.  $p_l$  - probability of inter-cluster links;
  2.  $p_s$  - probability of intra-cluster links.
- In the two-dimensional parameter plane defined by  $0 \leq (p_s, p_l) \leq 1$ , where can the network be best synchronized?
- Intuition: as  $p_l$  or  $p_s$  is increased, there are more links in the network, so it should be easier to achieve synchronization.

# Optimizing Synchronization: Main Result

- **Optimal synchronizability occurs for  $p_l \approx p_s$ .**  
For small  $p_l$ , increasing  $p_s$  can lead to destruction of synchronization.
- L. Huang, K. Park, Y.-C. Lai, L. Yang, and K.-Q. Yang, “Abnormal synchronization in complex clustered networks,” PRL 97, 164101 (2006).
- X.-G. Wang, L. Huang, Y.-C. Lai, and C. H. Lai, “Optimization of synchronization in gradient clustered networks,” PRE 6, 056113(1-5) (2007).
- L. Huang, Y.-C. Lai, and R. A. Gatenby, “Optimization of synchronization in complex clustered networks,” Chaos 18, 013101(1-10) (2008).

# Computational Criterion for Synchronization

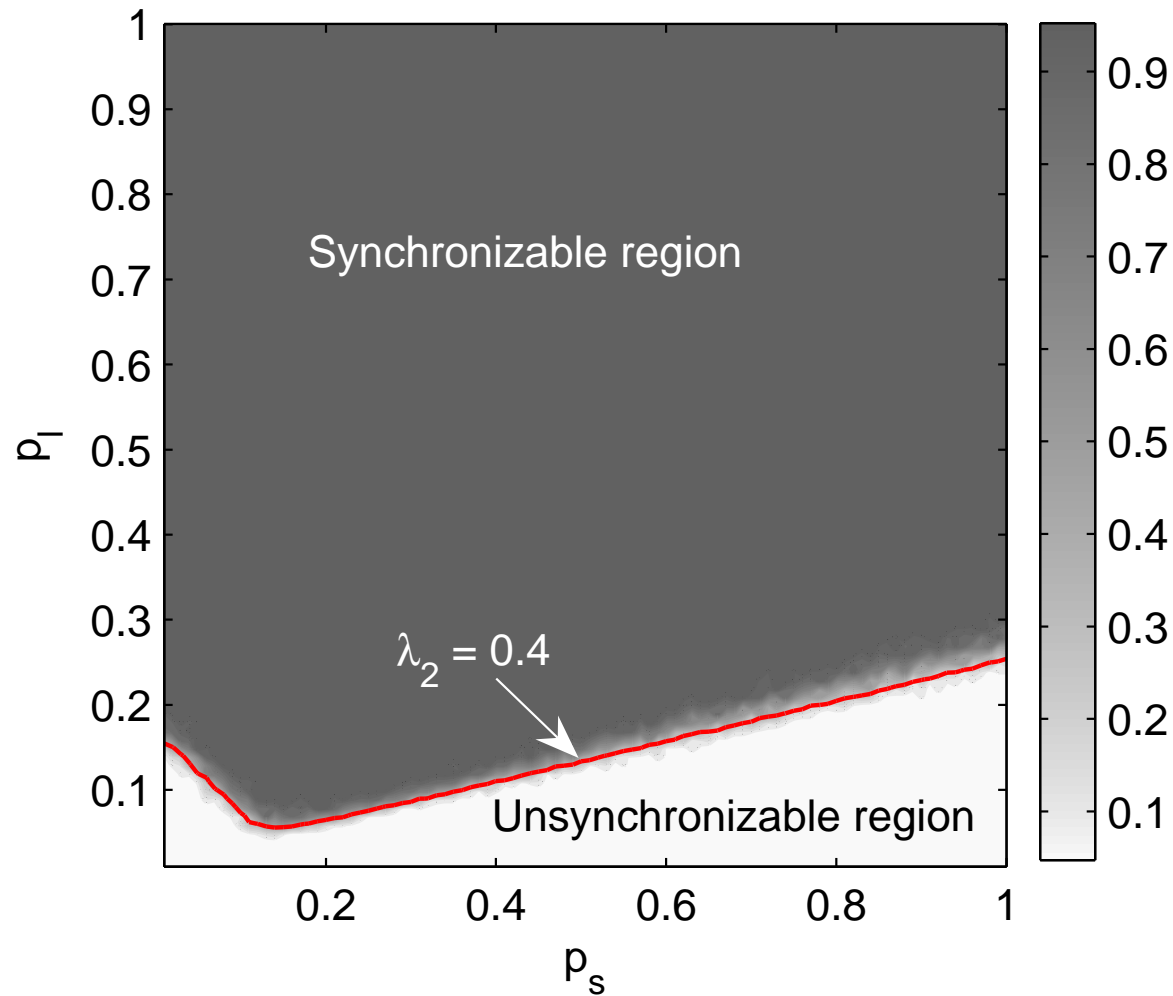
- Network synchronization is meaningful in a probabilistic sense.
- Calculate “synchronization width” defined by

$$W(t) = \langle |x(t) - \langle x(t) \rangle| \rangle,$$

where  $\langle \cdot \rangle$  is average over nodes in the network.

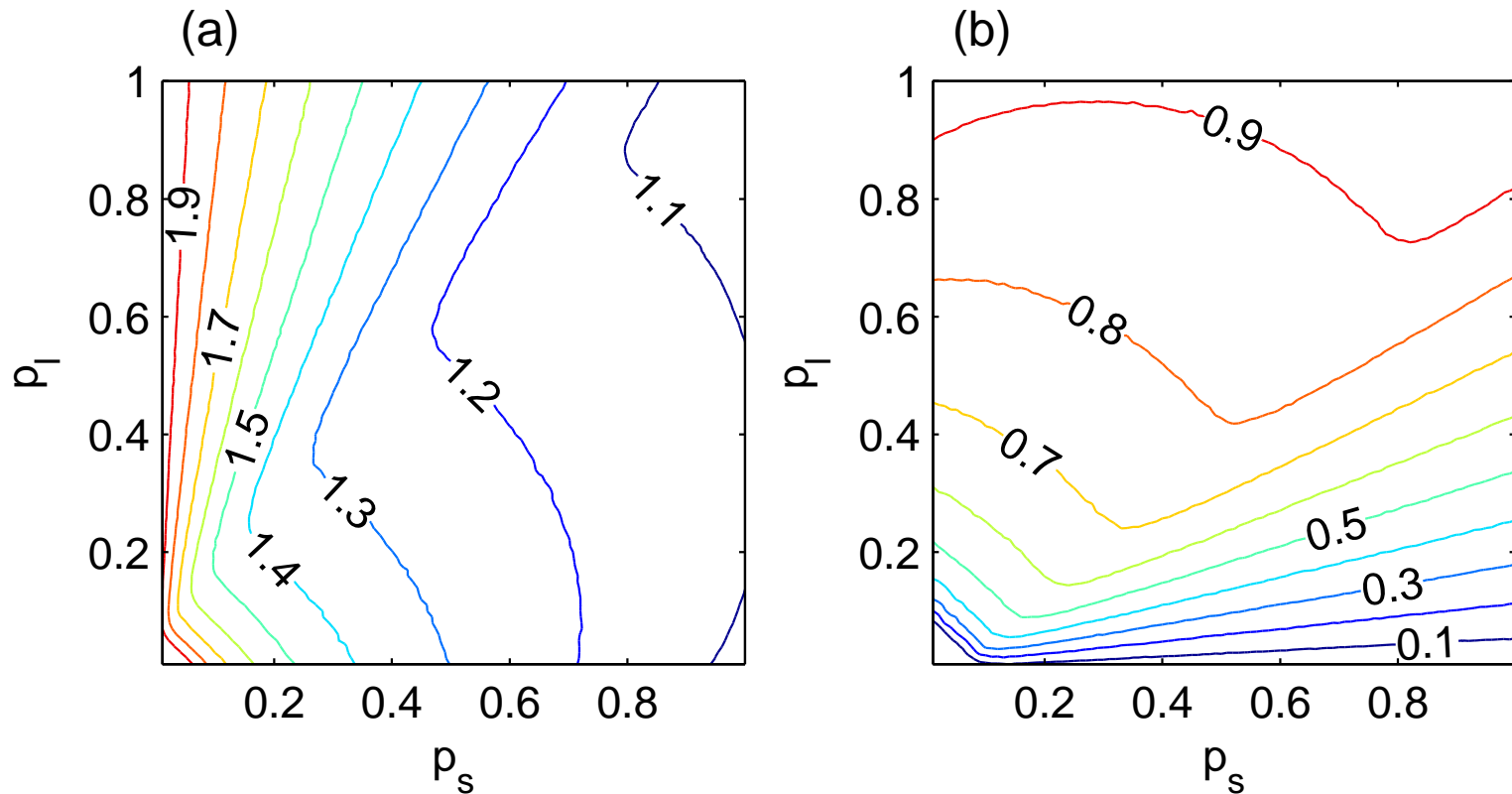
- Define  $P_{syn}$  - probability for  $W(t) < \delta$  (small number, say 0.01) for a large time period.
- $P_{syn} \approx 0$  - network is unsynchronizable;
- $P_{syn} > 0$  - network is synchronizable;
- $P_{syn} \approx 1$  - network is strongly synchronizable.

# Clustered Rössler Network



- Classical Rössler oscillator on each node;  $N = 100$ ,  $M = 2$ , and  $\varepsilon = 0.2$ .

# Contour Plots of $\lambda_N$ and $\lambda_2$



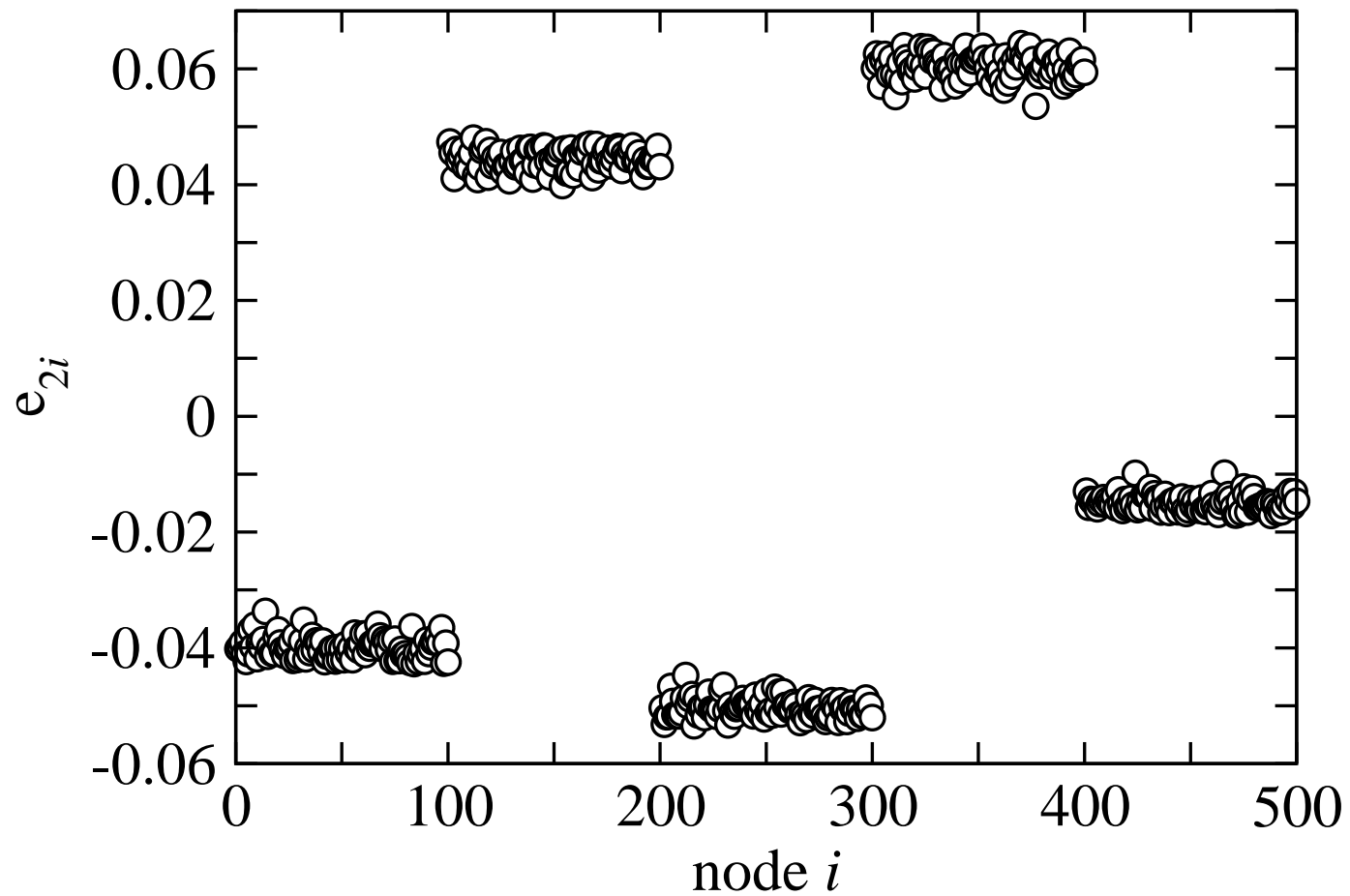
- Reminder: synchronizable if  $\lambda_2 > 0.4$  and  $\lambda_N < 9.2$ .
- Key:  $\lambda_2$ . For fixed  $p_l$  (small),  $\lambda_2(p_s)$  is a decreasing function for  $p_s$  large.



# Theory (1)

- For a clustered network, the components of the eigenvector  $\mathbf{e}_2$  have approximately the same value within any cluster, while they can be quite different for different clusters.
- So write
$$\mathbf{e}_2 \approx [\tilde{e}_1, \dots, \tilde{e}_1, \tilde{e}_2, \dots, \tilde{e}_2, \dots, \tilde{e}_M, \dots, \tilde{e}_M]^T.$$
- For each  $I$ ,  $1 \leq I \leq M$ , there are  $n$   $\tilde{e}_I$ 's in  $\mathbf{e}_2$ .
- By definition,  $\mathbf{G} \cdot \mathbf{e}_2 = \lambda_2 \mathbf{e}_2$  and  $\mathbf{e}_2 \cdot \mathbf{e}_2 = 1$ .

# Components of eigenvector $e_2$



## Theory (2)

- $\lambda_2 = \mathbf{e}_2^T \cdot \mathbf{G} \cdot \mathbf{e}_2 = \sum_{i,j=1}^N e_{2i} G_{ij} e_{2j}$ , where  $e_{2i}$  is the  $i$ th component of  $\mathbf{e}_2$ .
- We have

$$\lambda_2 = \sum_{i=1}^N e_{2i} \{ G_{i1} \tilde{e}_1 + G_{i2} \tilde{e}_1 + \cdots + G_{in} \tilde{e}_1 + G_{in+1} \tilde{e}_2 + \cdots + G_{iN} \tilde{e}_M \}.$$

## Theory (3)

- If  $i$  and  $j$  belong to the same cluster,  $G_{ij}$  equals  $-1/k_i$  with probability  $p_s$  and 0 with probability  $1 - p_s$ .
- If  $i$  and  $j$  belong to different clusters,  $G_{ij}$  equals  $-1/k_i$  with probability  $p_l$  and 0 with probability  $1 - p_l$ . Thus

$$\lambda_2 = \sum_{i=1}^N e_{2i} \left\{ -n \frac{p_l}{k_i} \tilde{e}_1 - n \frac{p_l}{k_i} \tilde{e}_2 + \dots \right. \\ \left. + \tilde{e}_I - n \frac{p_s}{k_i} \tilde{e}_I - \dots - n \frac{p_l}{k_i} \tilde{e}_M \right\},$$

where  $\tilde{e}_I$  is the value corresponding to the cluster that contains node  $i$ .

# Theory (4)

- For random subnetworks, the degree distribution is centered about  $k = np_s + (N - n)p_l \rightarrow k_i \approx k$ .

$$\begin{aligned}\lambda_2 &\approx \sum_{I=1}^M n\tilde{e}_I \left\{ N\frac{p_l}{k}\tilde{e}_I - n\frac{p_l}{k} \sum_{J=1}^M \tilde{e}_J \right\} \\ &= N\frac{p_l}{k} \sum_{I=1}^M n\tilde{e}_I^2 - \left( n \sum_{J=1}^M \tilde{e}_J \right)^2 \frac{p_l}{k}.\end{aligned}$$

- Since  $\sum_{I=1}^M n\tilde{e}_I^2 \approx \sum_{i=1}^N e_{2i}^2 = 1$ , and  $n \sum_{J=1}^M \tilde{e}_J = \sum_{i=1}^N e_{2i}$ , we have

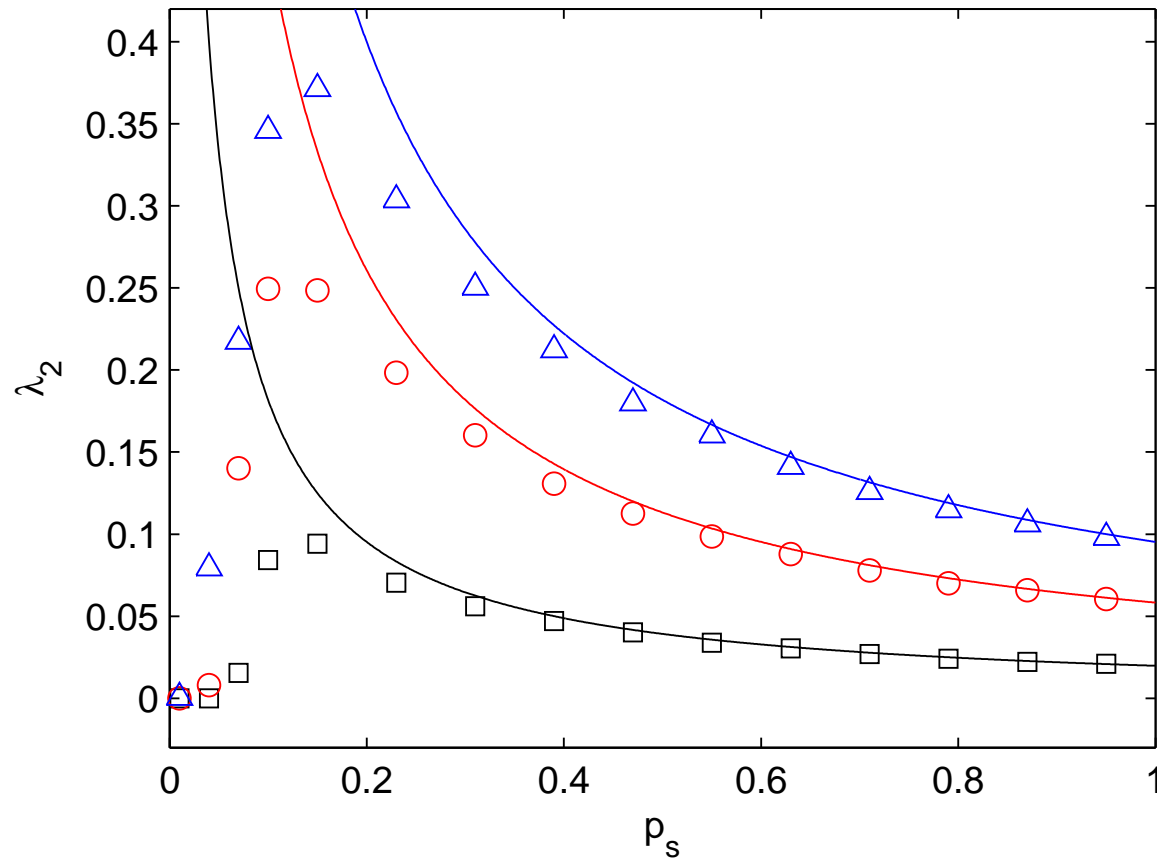
$$\lambda_2 = \frac{Np_l}{np_s + (N - n)p_l} - \left( \sum_{i=1}^N e_{2i} \right)^2 \frac{p_l}{k}.$$

# Theory (5)

- $\lambda_2 = Np_l / [np_s + (N - n)p_l] - (\sum_{i=1}^N e_{2i})^2$
- The normalized eigenvector  $e_1$  of  $\lambda_1$  is associated with the synchronized state, thus its components have constant values:  $e_1 = [1/\sqrt{N}, \dots, 1/\sqrt{N}]^T$ .
- If  $G$  is symmetric, then eigenvectors for different eigenvalues are orthogonal:  $e_i \cdot e_j = \delta_{ij}$ , indicating  $\sum_{l=1}^N e_{2l} = 0$ .
- If  $G$  is slightly asymmetric, we expect  $\sum_{i=1}^N e_{2i}$  to be small - safe to neglect the second term in  $\lambda_2$ . We have

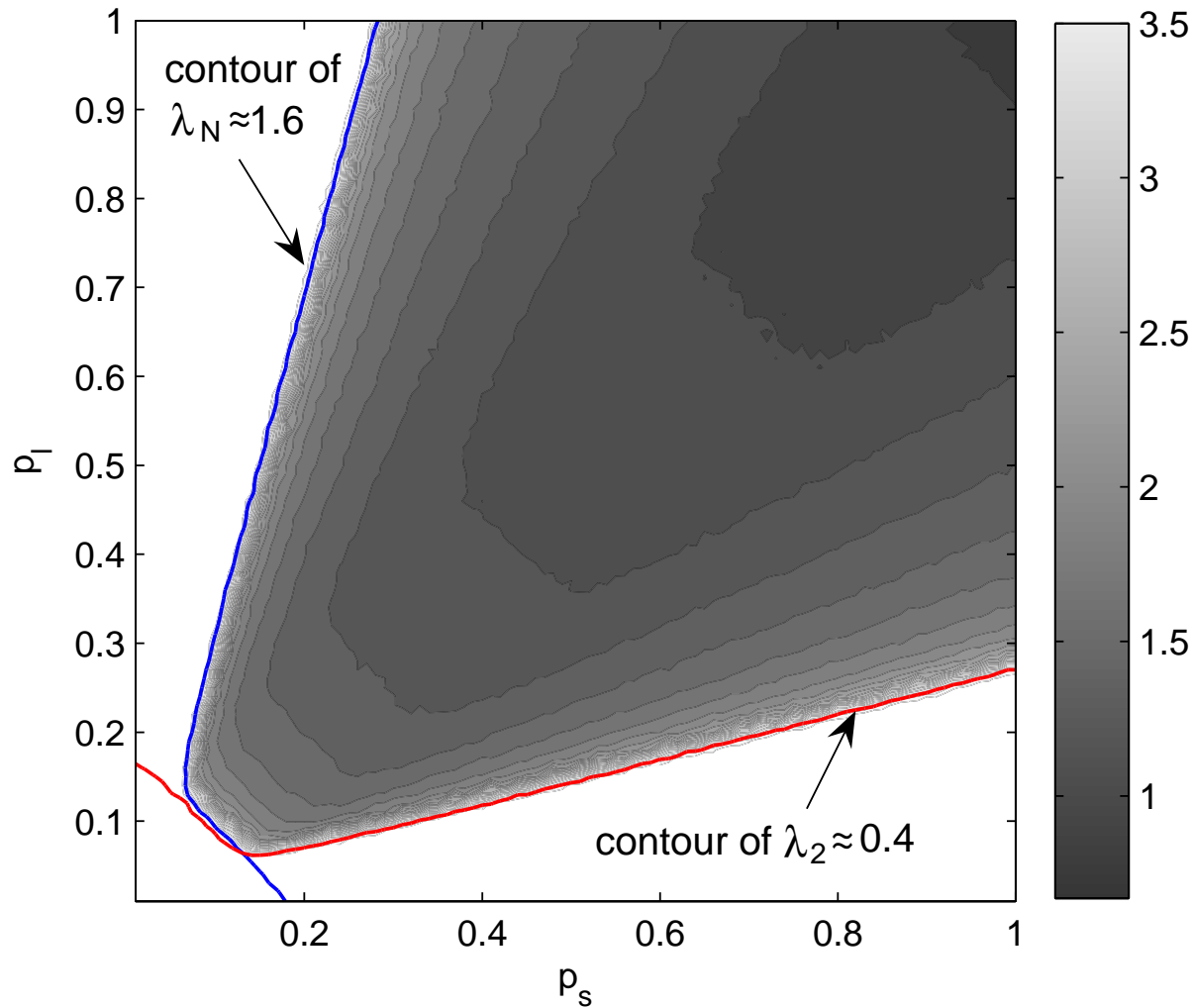
$$\lambda_2 \approx \frac{Np_l}{np_s + (N - n)p_l}$$

# Numerical Verification



- $p_l = 0.01$  (squares),  $p_l = 0.03$  (circles), and  $p_l = 0.05$  (triangles);  $N = 100$ ,  $n = 50$ , and 100 network realizations.

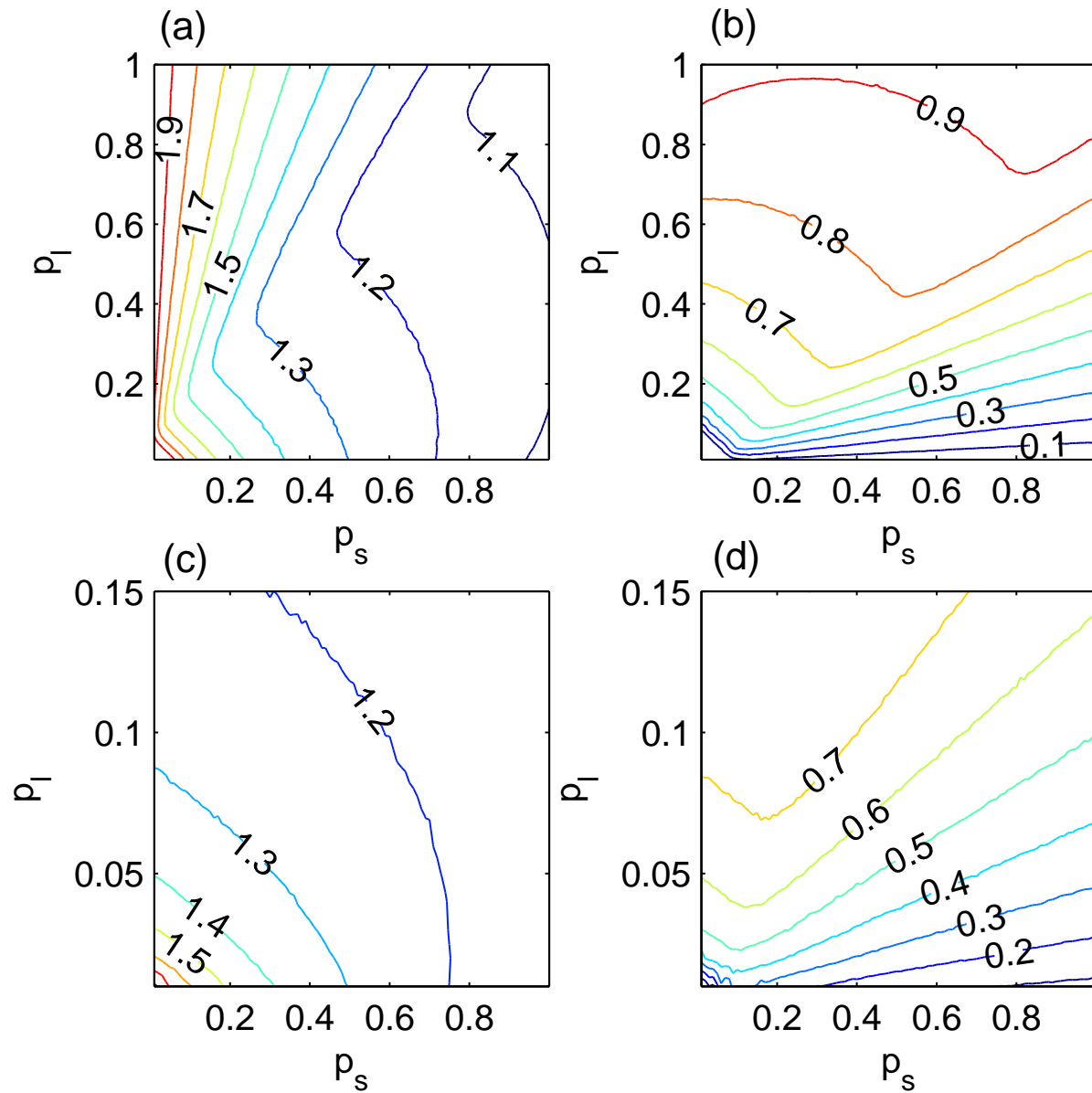
# Clustered Network of Chaotic Logistic Maps



- $N = 100$ ,  $M = 2$ ; chaotic logistic map on each node.



# Contour plots of $\lambda_N$ and $\lambda_2$



# Conclusion

- **Optimization:** network is most synchronizable when the numbers of the inter-cluster and intra-cluster links are approximately equal.

## Potential use:

- Insights into synchronization-dependent biological functions;
- Synchronous timing in large computer networks;
- Better coordination in transportation networks;
- ...