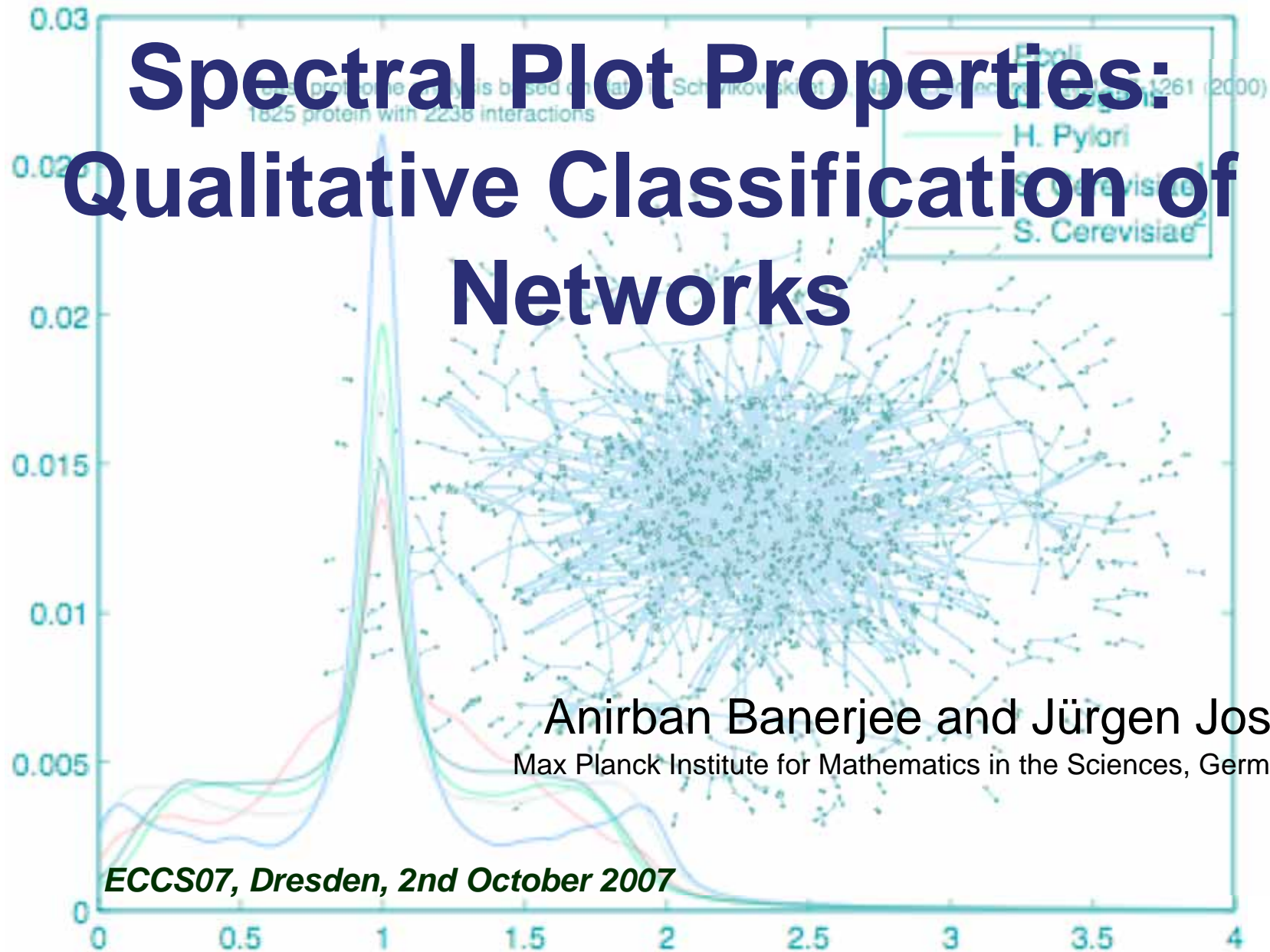


# Spectral Plot Properties: Qualitative Classification of Networks



# Overview of the talk

- Introduction
- Graph Laplacian operator
- Eigenfunction and graph structure
- Visualization techniques
- Qualitative classification of networks
- Conclusion and remarks

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- **Introduction**

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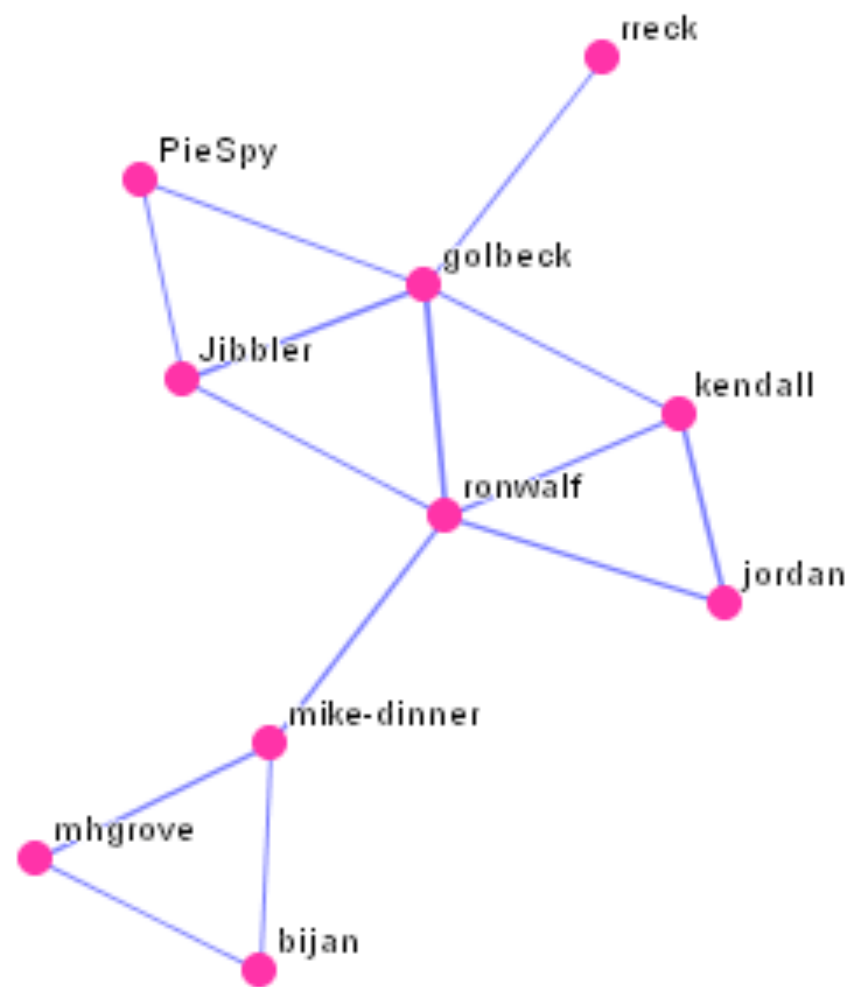
# Network construction

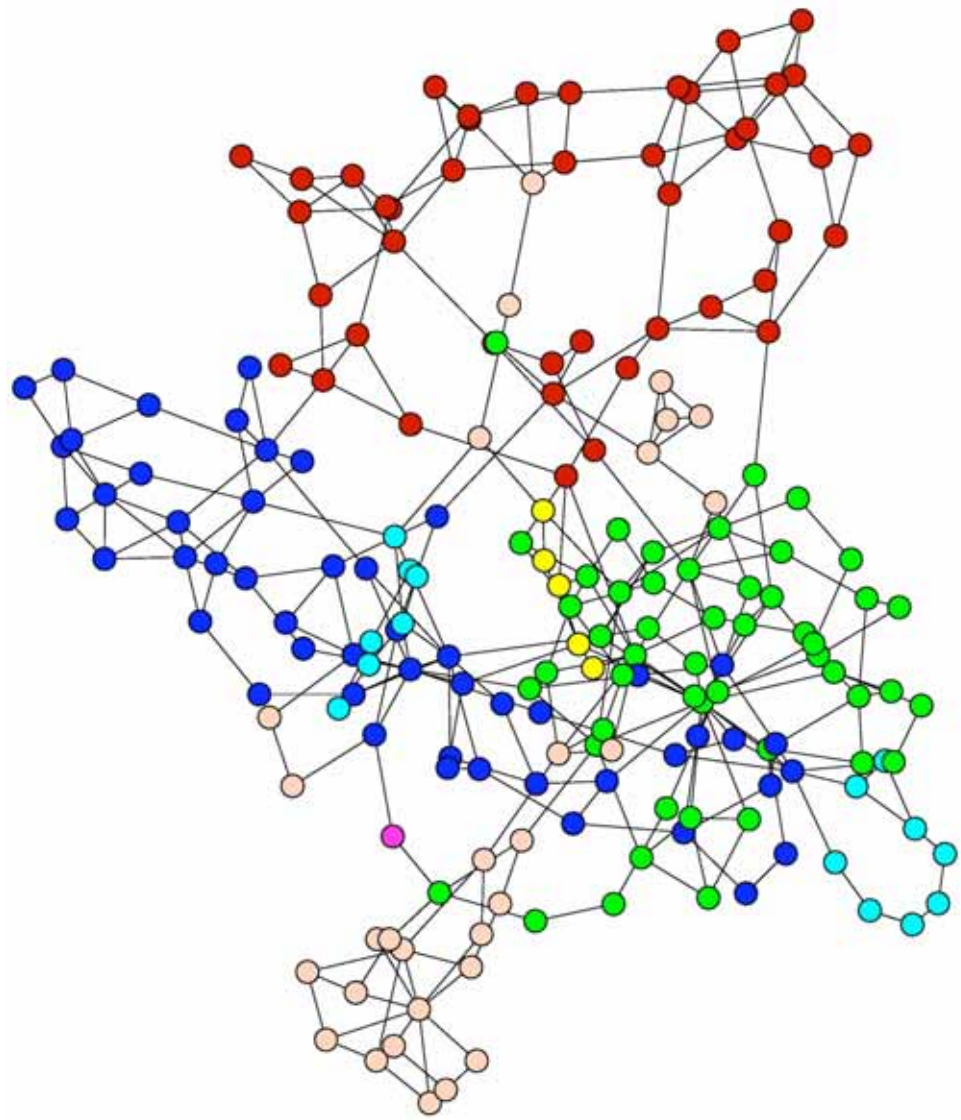
- Component  $\Rightarrow$  Vertex
- Interaction or relation  $\Rightarrow$  Edge

Degree of a vertex = number of its connected neighbors

# Few examples of real networks

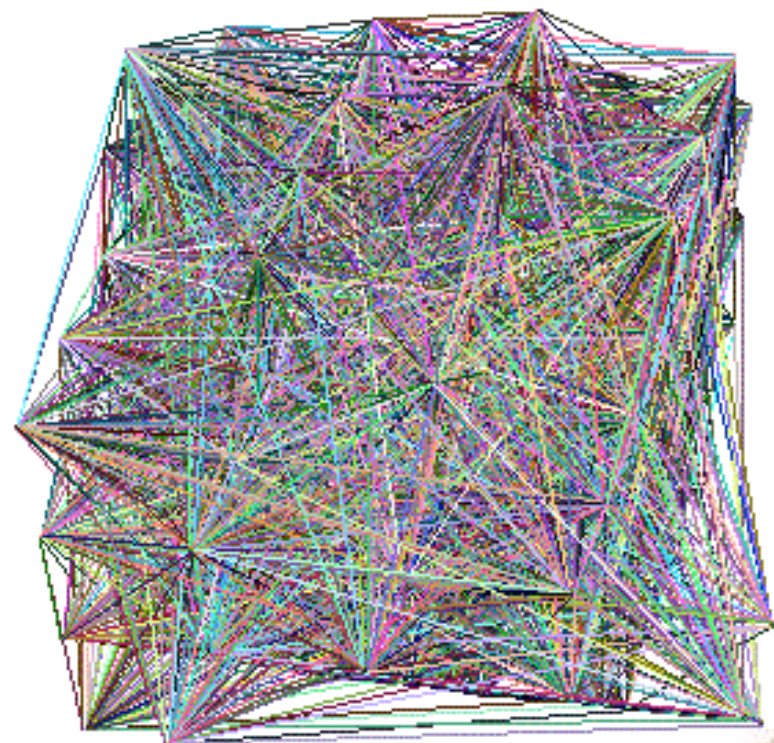
1. Protein-protein interaction network
  - Protein  $\rightarrow$  Vertex
  - Direct physical interactions (binding)  $\rightarrow$  Edges
2. Power grid network
  - Generator, transformers, substations  $\rightarrow$  Vertices
  - High-voltage transmission lines  $\rightarrow$  Edges
3. Neuronal network
  - Neuron  $\rightarrow$  Vertex
  - Synaptic connections  $\rightarrow$  Edges
4. Scientific collaboration network
  - Scientists  $\rightarrow$  Vertices
  - Having joint publication  $\rightarrow$  Edge
5. Food-web network
  - Species  $\rightarrow$  Vertices
  - Predator-prey relation  $\rightarrow$  Edge

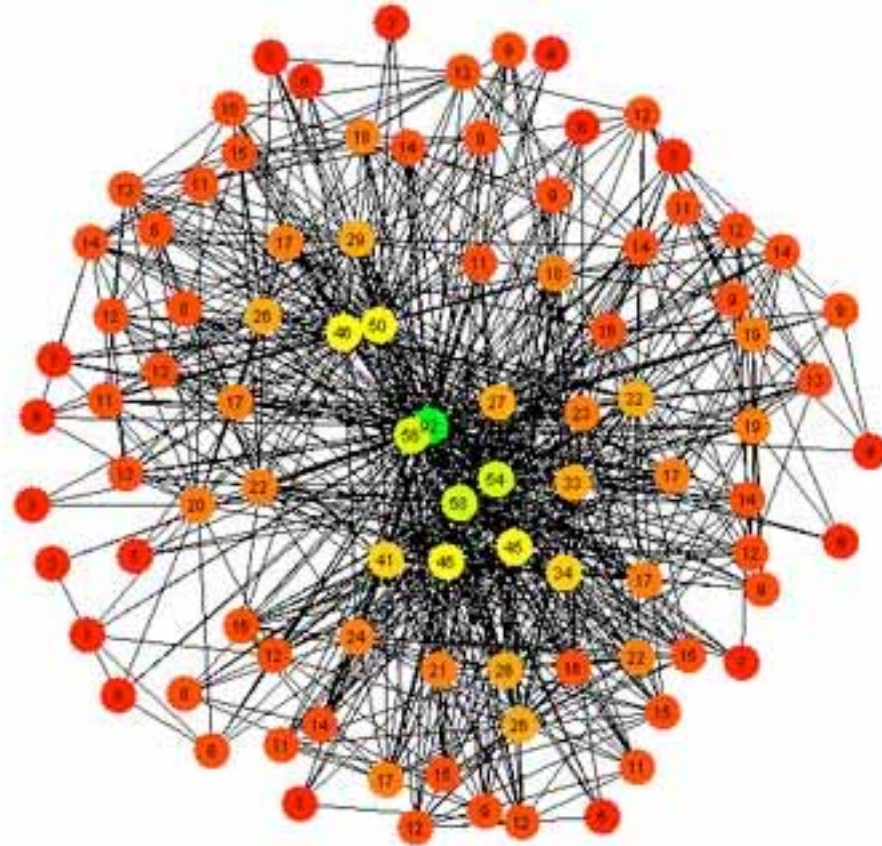


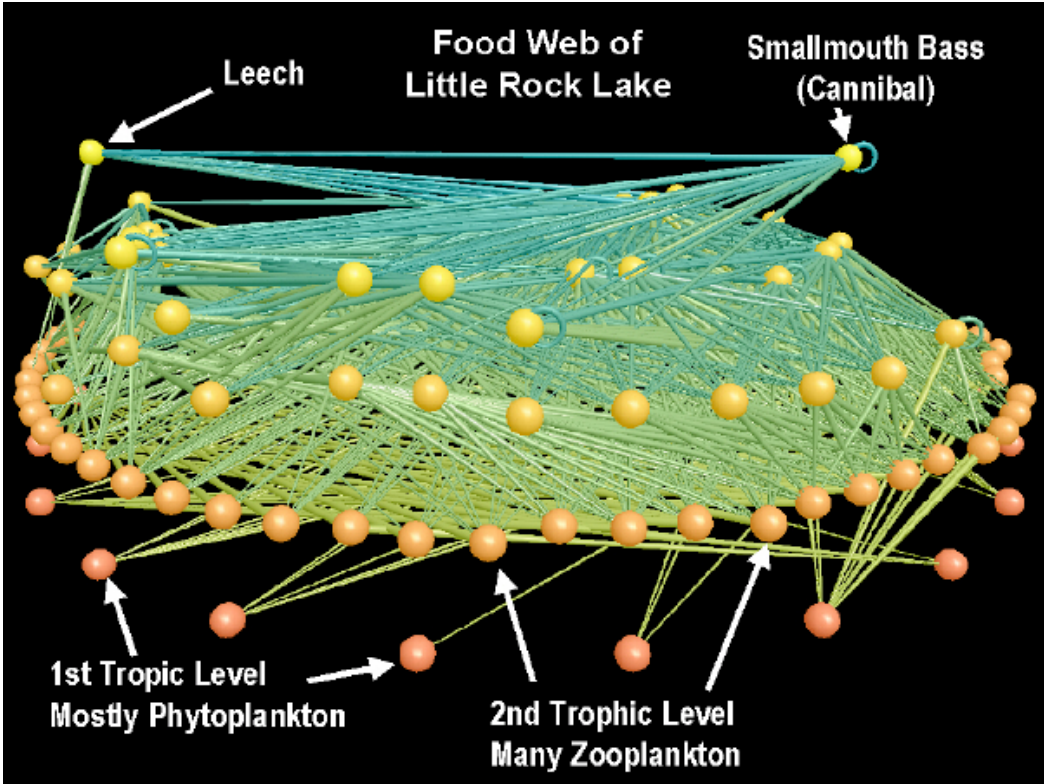


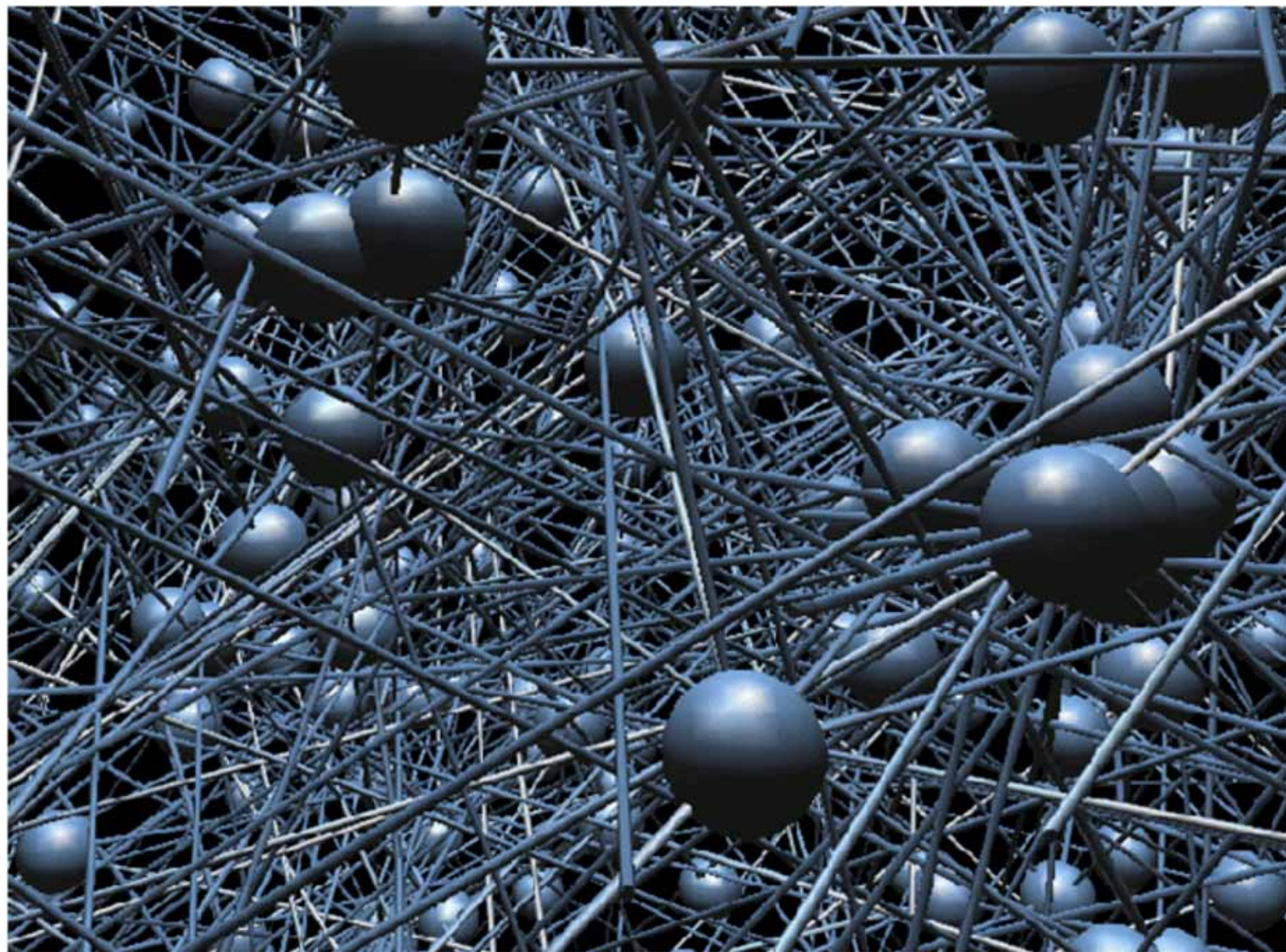


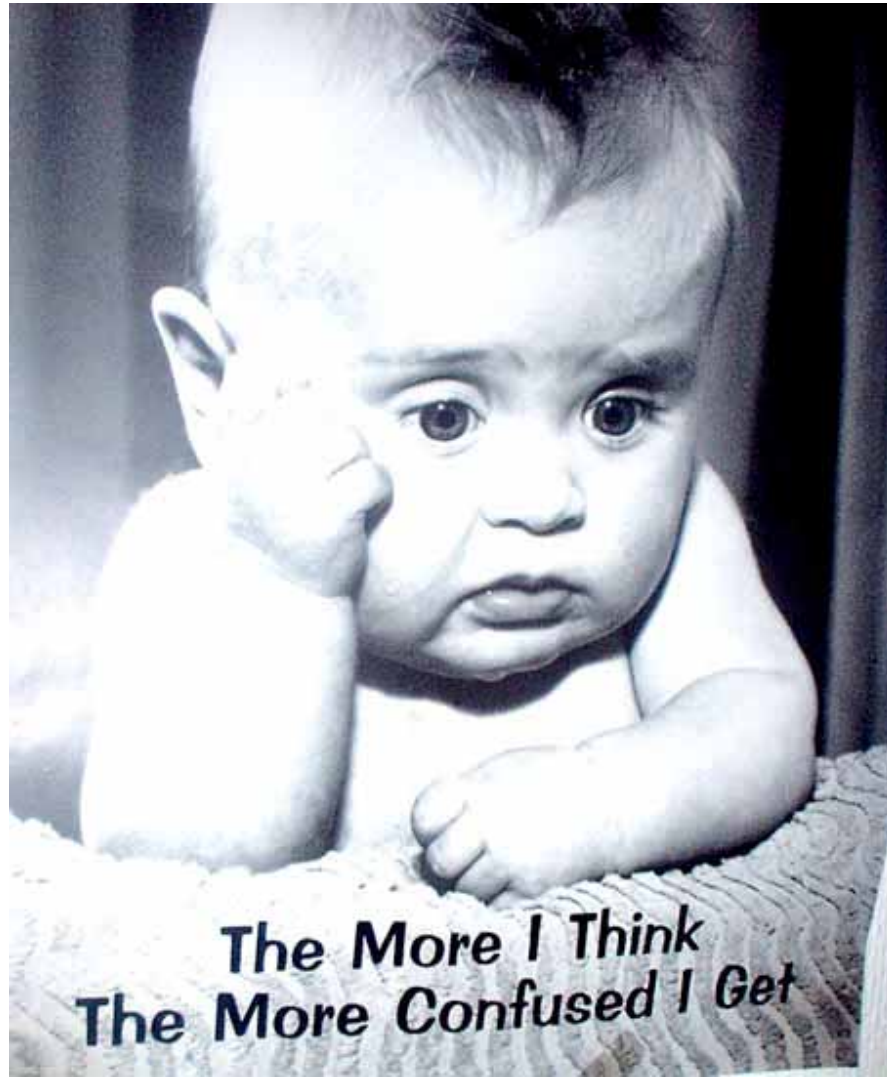












**The More I Think  
The More Confused I Get**

# Parameters for analyzing network structure

1. Degree distribution
  2. Average path length
  3. Diameter
  4. Clustering coefficient
  5. Betweenness centrality
- etc.

These invariants can not capture all qualitative aspects of a graph.

e.g. graphs with same degree distribution can have completely different structural (and dynamical) properties.

And two graphs with very different structure can have same clustering coefficient.

# Power-law every where

Network	Size	$\langle k \rangle$	$\kappa$	$\gamma_{out}$	$\gamma_{in}$	$\ell_{real}$	$\ell_{rand}$	$\ell_{pow}$	Reference	Nr.
WWW	325, 729	4.51	900	2.45	2.1	11.2	8.32	4.77	Albert, Jeong, Barabási 1999	1
WWW	$4 \times 10^7$	7		2.38	2.1				Kumar <i>et al.</i> 1999	2
WWW	$2 \times 10^8$	7.5	4,000	2.72	2.1	16	8.85	7.61	Broder <i>et al.</i> 2000	3
WWW, site	260,000				1.94				Huberman, Adamic 2000	4
Internet, domains*	3, 015 - 4, 389	3.42 - 3.76	30 - 40	2.1 - 2.2	2.1 - 2.2	4	6.3	5.2	Faloutsos 1999	5
Internet, routers*	3, 888	2.57	30	2.48	2.48	12.15	8.75	7.67	Faloutsos 1999	6
Internet, routers*	150,000	2.66	60	2.4	2.4	11	12.8	7.47	Govindan 2000	7
Movie actors*	212, 250	28.78	900	2.3	2.3	4.54	3.65	4.01	Barabási, Albert 1999	8
Coauthors, SPIRES*	56, 627	173	1, 100	1.2	1.2	4	2.12	1.95	Newman 2001b,c	9
Coauthors, neuro.*	209, 293	11.54	400	2.1	2.1	6	5.01	3.86	Barabási <i>et al.</i> 2001	10
Coauthors, math*	70, 975	3.9	120	2.5	2.5	9.5	8.2	6.53	Barabási <i>et al.</i> 2001	11
Sexual contacts*	2810			3.4	3.4				Liljeros <i>et al.</i> 2001	12
Metabolic, E. coli	778	7.4	110	2.2	2.2	3.2	3.32	2.89	Jeong <i>et al.</i> 2000	13
Protein, S. cerev.*	1870	2.39		2.4	2.4				Mason <i>et al.</i> 2000	14
Ythan estuary*	134	8.7	35	1.05	1.05	2.43	2.26	1.71	Montoya, Solé 2000	14
Silwood park*	154	4.75	27	1.13	1.13	3.4	3.23	2	Montoya, Solé 2000	16
Citation	783, 339	8.57			3				Redner 1998	17
Phone-call	$53 \times 10^6$	3.16		2.1	2.1				Aiello <i>et al.</i> 2000	18
Words, cooccurences*	460, 902	70.13		2.7	2.7				Cancho, Solé 2001	19
Words, synonyms*	22, 311	13.48		2.8	2.8				Yook <i>et al.</i> 2001	20



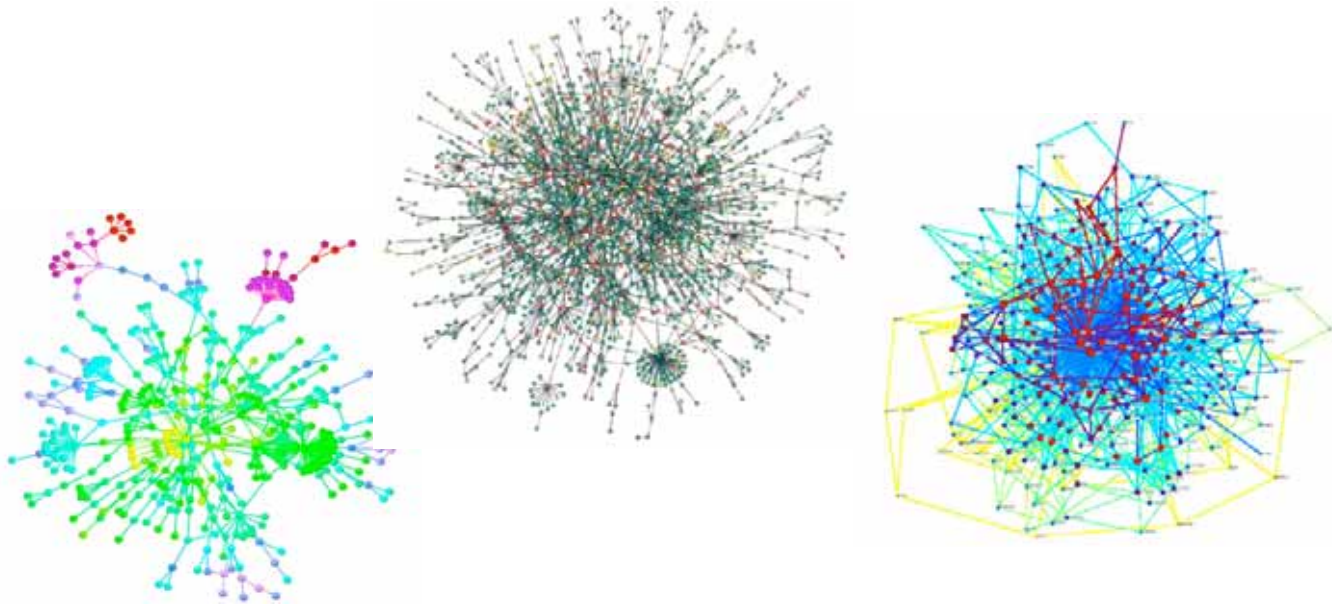
R. Albert and A Barabási, *Rev. of Modern Phys.* 74, 47-97, 2002



How many invariants do we need to consider to investigate the systematic structural difference and similarities between the graphs from different classes?



For a given a particular structure, which of each features or qualities are universal, that is, shared by other structures?



What is unique and special for the structure from a particular class?

Are those invariants qualitatively good enough to identify the domain of a given an empirical graph?

So instead of focusing on particular and specific aspects and quantities in details, of a given a large and complex structure, we can try to obtain, at least at some rough level, a simultaneous representation of all its qualitative features.

Therefore, we are advocating here a tentative classification scheme for empirical networks based on underlying global qualitative properties detected through the *graph Laplacian spectrum*

that,

on one hand, are complete qualitative characterization of a graph and on other hand, can be easily graphically represented and therefore visually analyzed and compared.

# Overview of the talk

- Introduction
- **Graph Laplacian operator**
- Eigenfunction and graph structure
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$\Gamma$  : A finite undirected graph with  $N$  vertices.

Vertices:  $i, j \in \Gamma$  are connected  $\Rightarrow i \sim j$

$n_i \Rightarrow$  degree of vertex  $i$

For any function  $v : \Gamma \rightarrow \mathbb{R}$

$$\Delta v(i) := v(i) - \frac{1}{n_i} \sum_{j, j \sim i} v(j)$$

A.Banerjee, J.Jost, *On the spectrum of the normalized graph Laplacian*, submitted.

$$\mathcal{L}u(i) := u(i) - \frac{1}{n_i} \sum_{j, j \sim i} \frac{1}{\sqrt{n_i n_j}} u(j)$$

F.Chung, *Spectral graph theory*, 1997

$$a_{ij} = \begin{cases} 1, & \text{if } i = j \text{ and } n_i \neq 0 \\ -\frac{1}{\sqrt{n_i n_j}}, & \text{if } ij \text{ is an edge} \\ 0, & \text{otherwise.} \end{cases}$$

~~$$Lu(i) := n_i u(i) - \sum_{j, j \sim i} u(j)$$~~

~~A.Bollobás, *Modern graph theory*, 1998~~

~~$$a_{ij} = \begin{cases} n_i, & \text{if } i = j \\ -1, & \text{if } ij \text{ is an edge} \\ 0, & \text{otherwise} \end{cases}$$~~

# Properties of this operator

- This operator is symmetric about the product

$$(u, v) := \sum_{i \in \Gamma} n_i u(i) v(i) \Rightarrow \text{eigenvalues are real}$$

- $(\Delta u, u) \geq 0 \Rightarrow$  eigenvalues are non-negative

- $\Delta u = 0$  for  $u = \text{constant} \Rightarrow \lambda_{min} = 0$



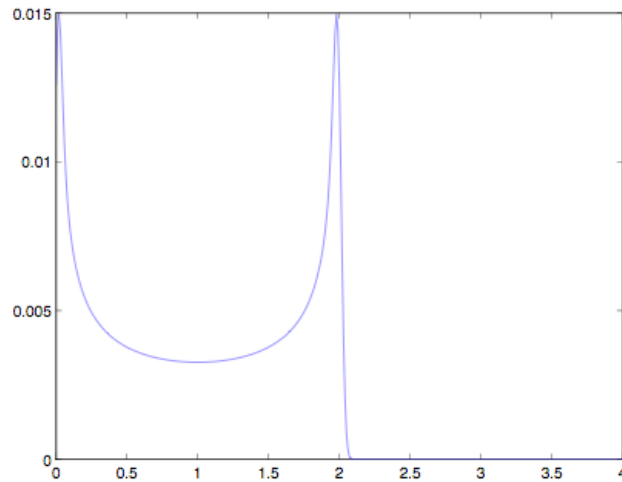
# Eigenvalues of this operator

$$0 = \lambda_0 \leq \lambda_1 \leq \cdots \leq \lambda_{N-1} \leq 2$$

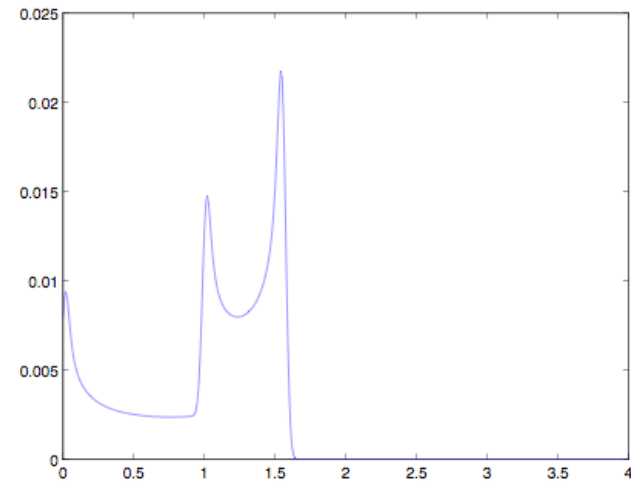
- Multiplicity of 0  $\Leftrightarrow$  # components in the graph
- Eigenvalues  $\lambda_1$  and  $\lambda_{N-1}$
- $\lambda_{N-1} = 2 \Leftrightarrow$  graphs is bipartite
  - Spectrum is symmetric about 1
- Complete graph with  $N$  vertices  $\Leftrightarrow$   
$$\lambda_1 = \lambda_2 = \cdots = \lambda_{N-1} = \frac{N}{N-1}$$

Few example of spectral plots

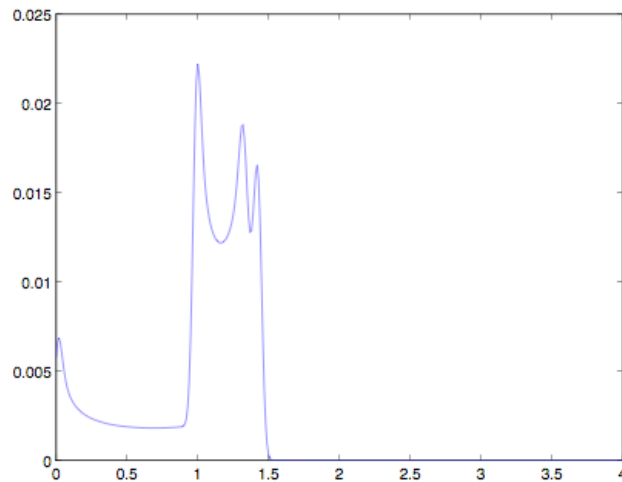
# 1D regular ring lattices



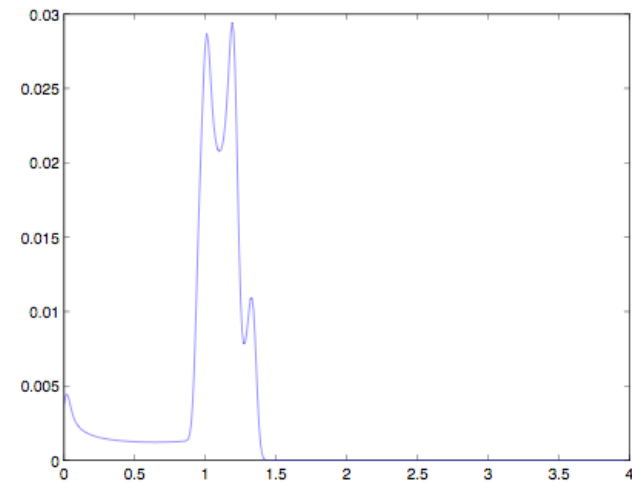
Deg. of each node = 2



Deg. of each node = 4

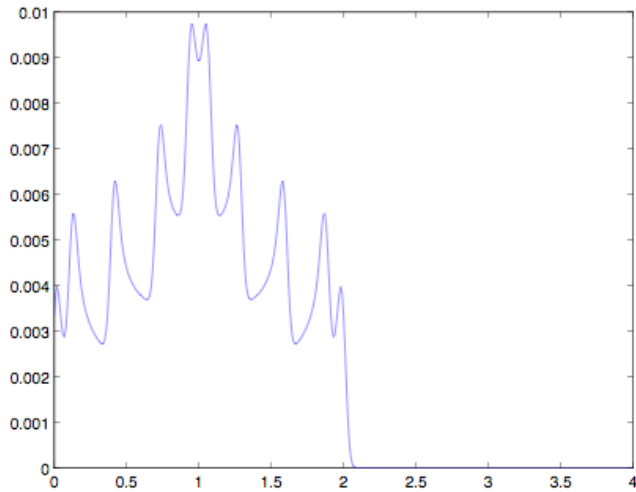


Deg. of each node = 6

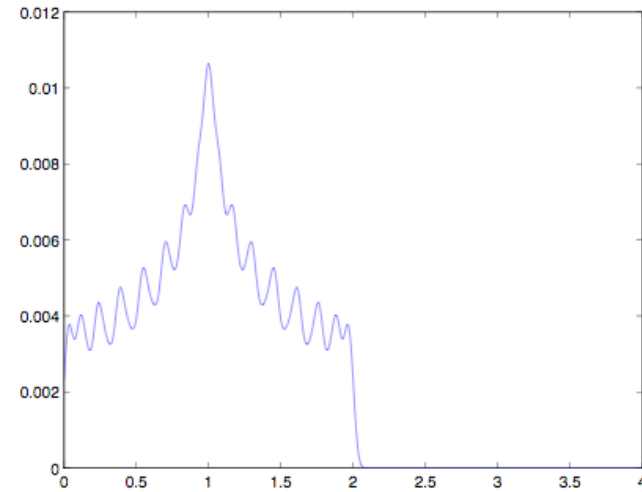


Deg. of each node = 10

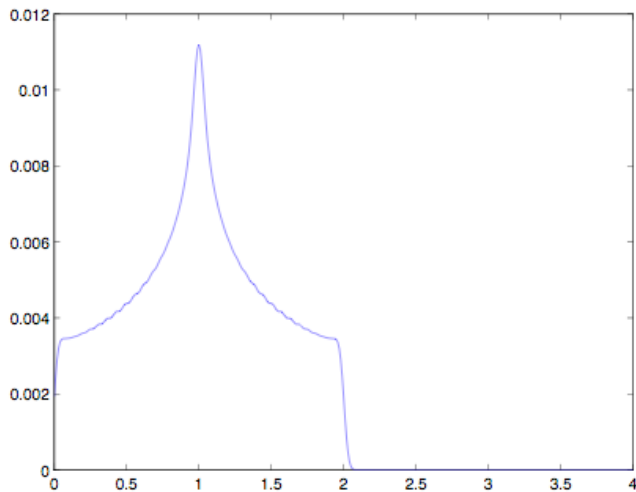
# 2D square grids



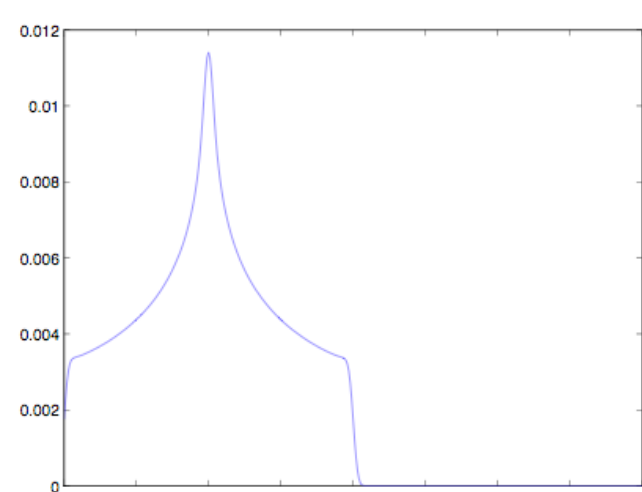
Dimension = 5 X 2000



Dimension = 10 X 1000

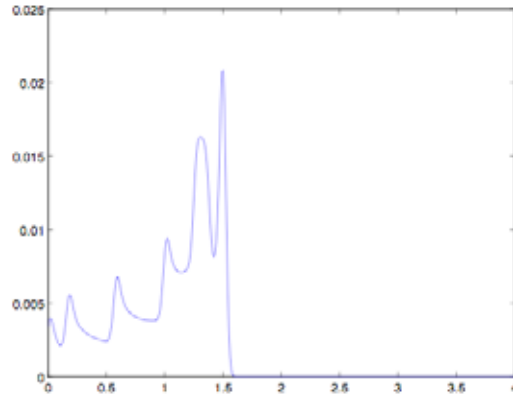


Dimension = 25 X 400

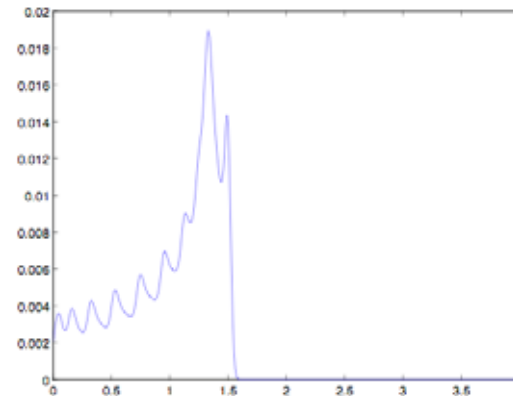


Dimension = 100 X 100

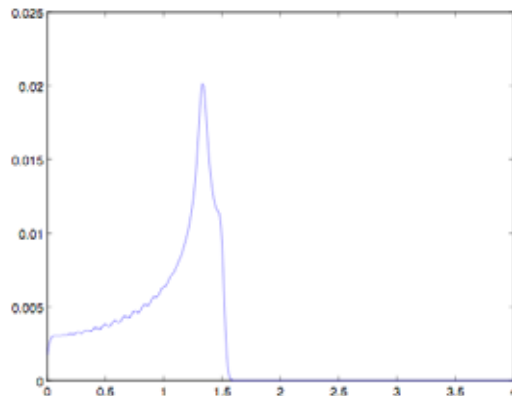
# 2D square grid with one diagonal



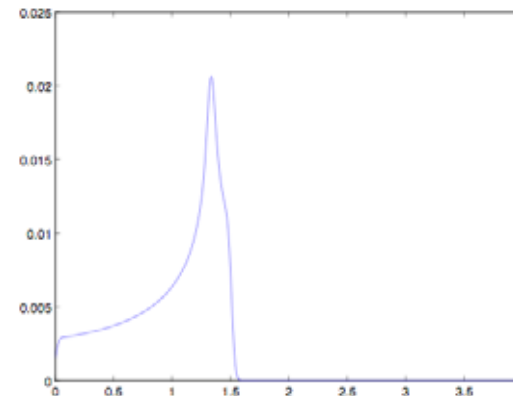
Dimension = 5 X 2000



Dimension = 10 X 1000

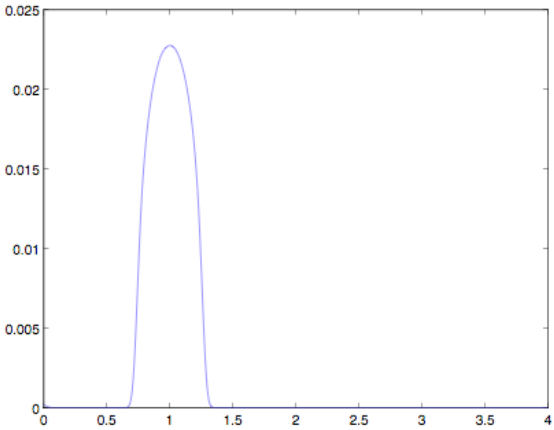


Dimension = 25 X 400

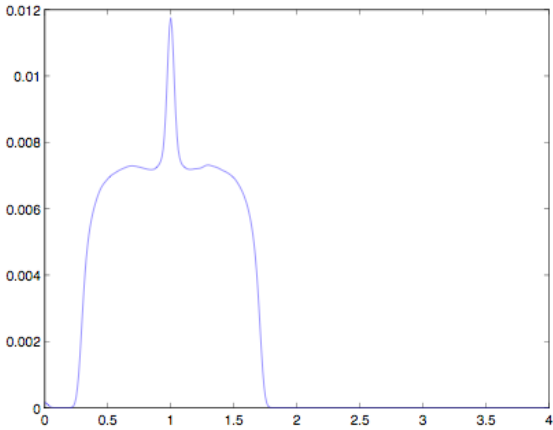


Dimension = 100 X 100

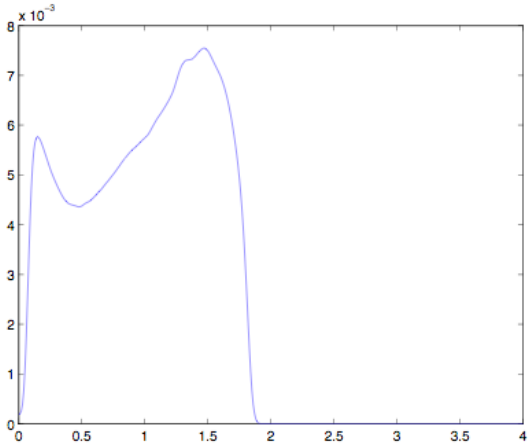
# Three generic models



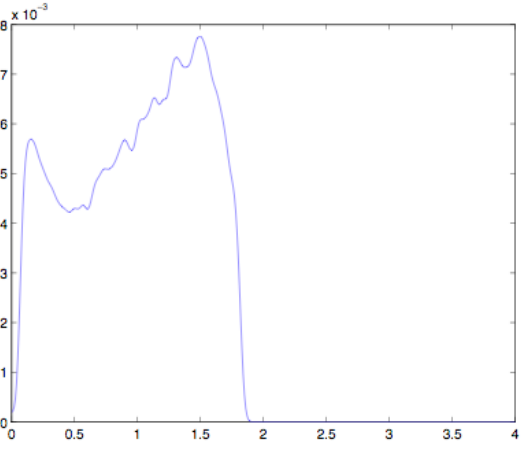
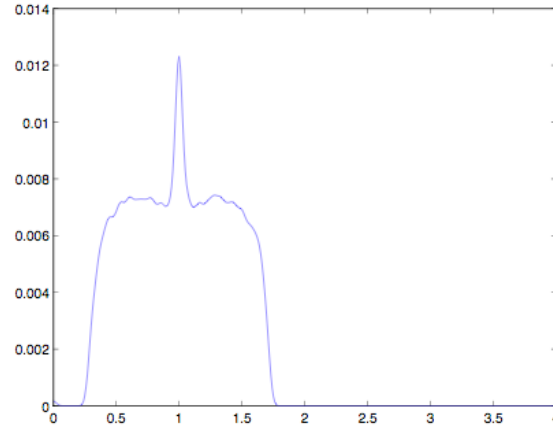
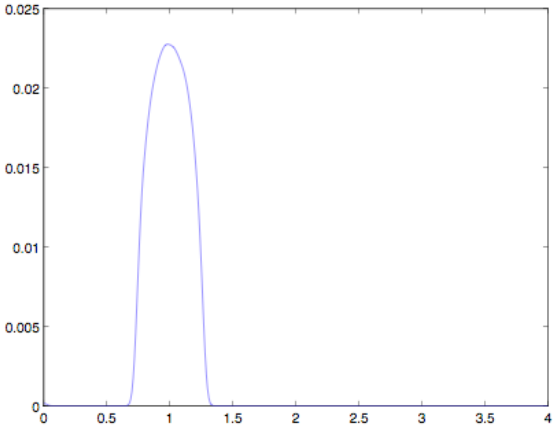
Random network



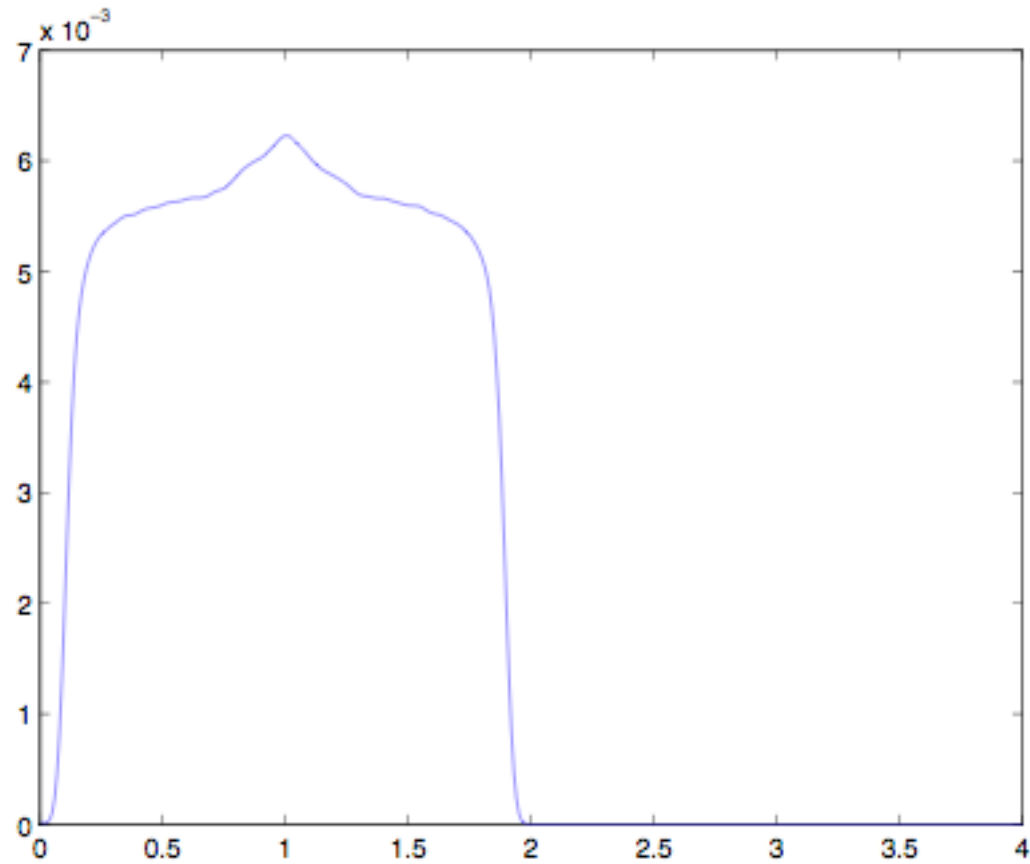
Small-world network



Scale-free network

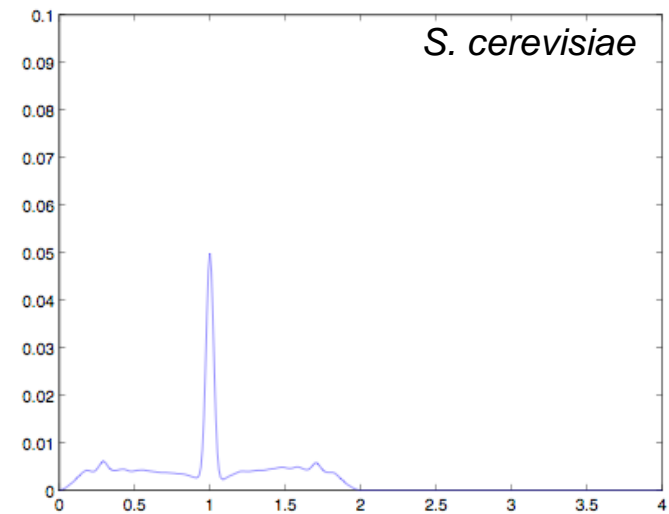
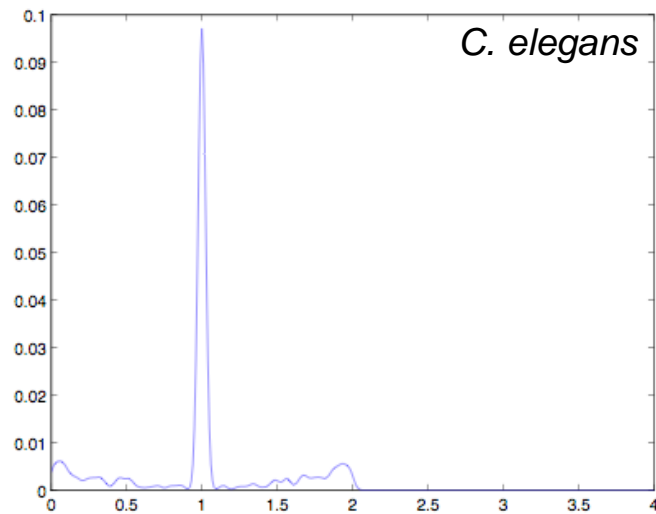
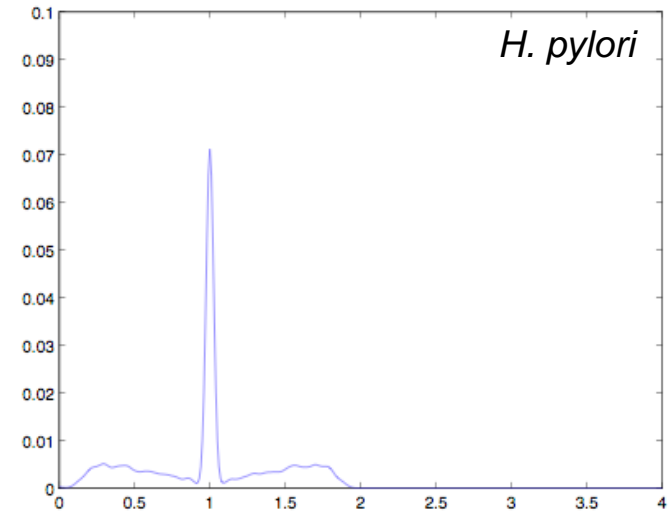
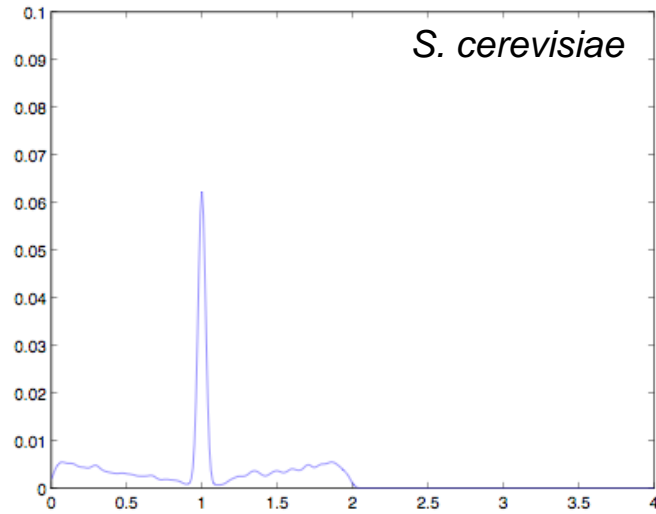


# Small world network (created from 2D grid)



Small-world network, created by rewiring 2-D square grid

# Protein-protein interaction networks





# Overview of the talk

- Introduction
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- **Eigenfunction and graph structure**
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Let think of a graph, representing real data as a structure that has evolved from some simpler precursors.

Constructions with different graph operation related to the evolution of a network describe certain processes of graph formation that leave characteristic traces in the spectrum.

A solution  $u_k$  of the eigenvalue equation

$$\Delta u_k - \lambda_k u_k = 0$$

- can be localize
- can be global

# Vertex doubling

Doubling a single vertex  $j_0 \in \Gamma$

$i_0$  is double of  $j_0$

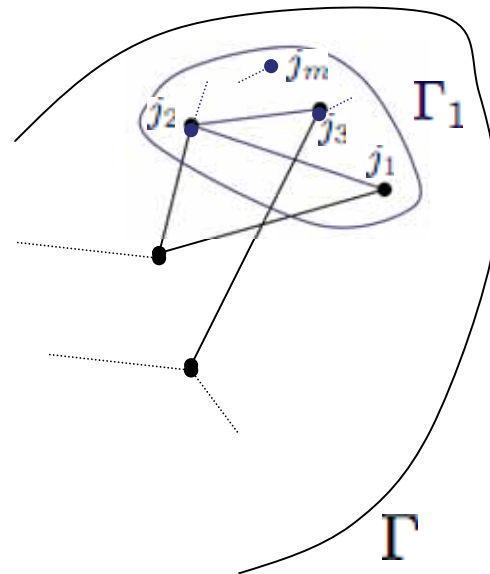
-- generates eigenvalue 1 with an eigenfunction

$$u_1(i) = \begin{cases} 1, & \text{for } i = j_0 \\ -1, & \text{for } i = i_0 \\ 0, & \text{otherwise} \end{cases}$$

High peak at 1: evolve by sequence of vertex duplication

# Motif duplication

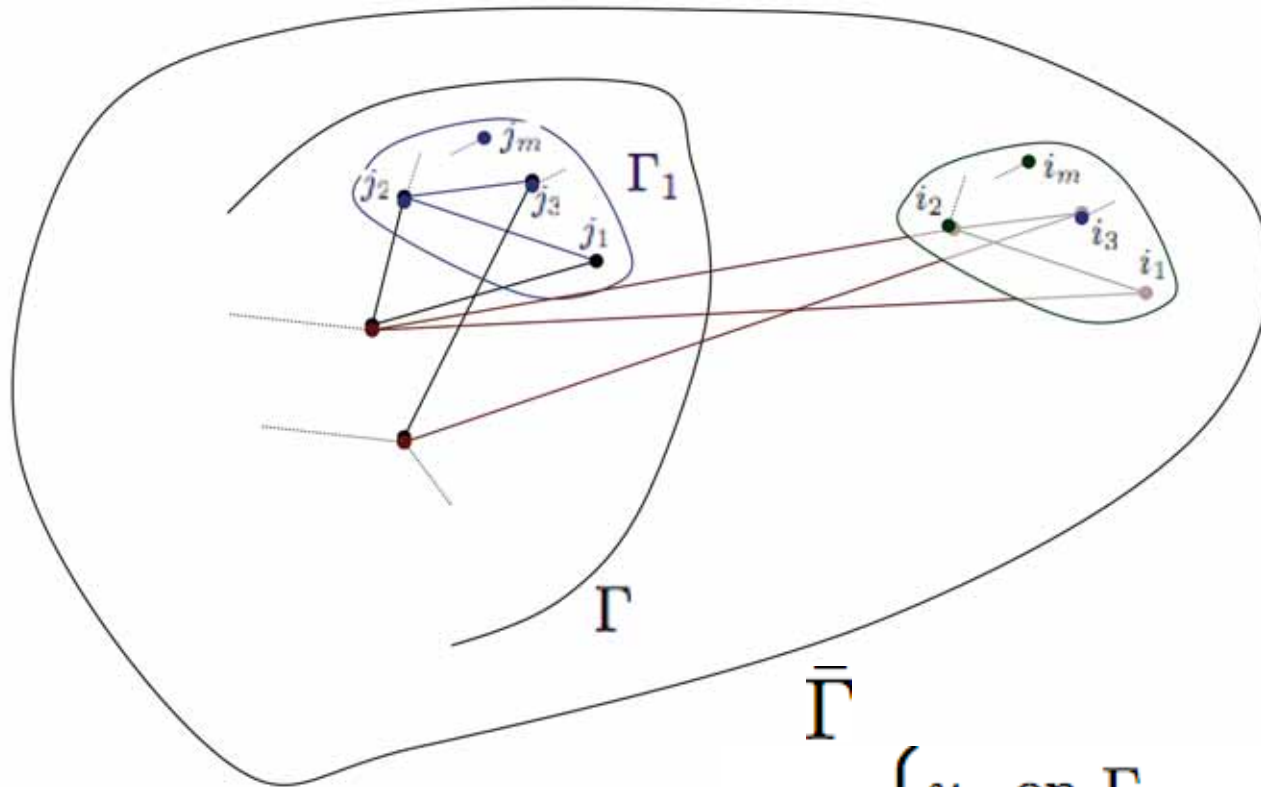
*Motif*: small sub-graph (where as the graph is supposed to be large) containing all edges of the graph between vertices of that subgraph



$$\frac{1}{n_i} \sum_{j \in \Gamma_1, j \sim i} u(j) = (1 - \lambda)u(i) \text{ for all } i \in \Gamma_1 \text{ and some } \lambda$$

# Motif duplication

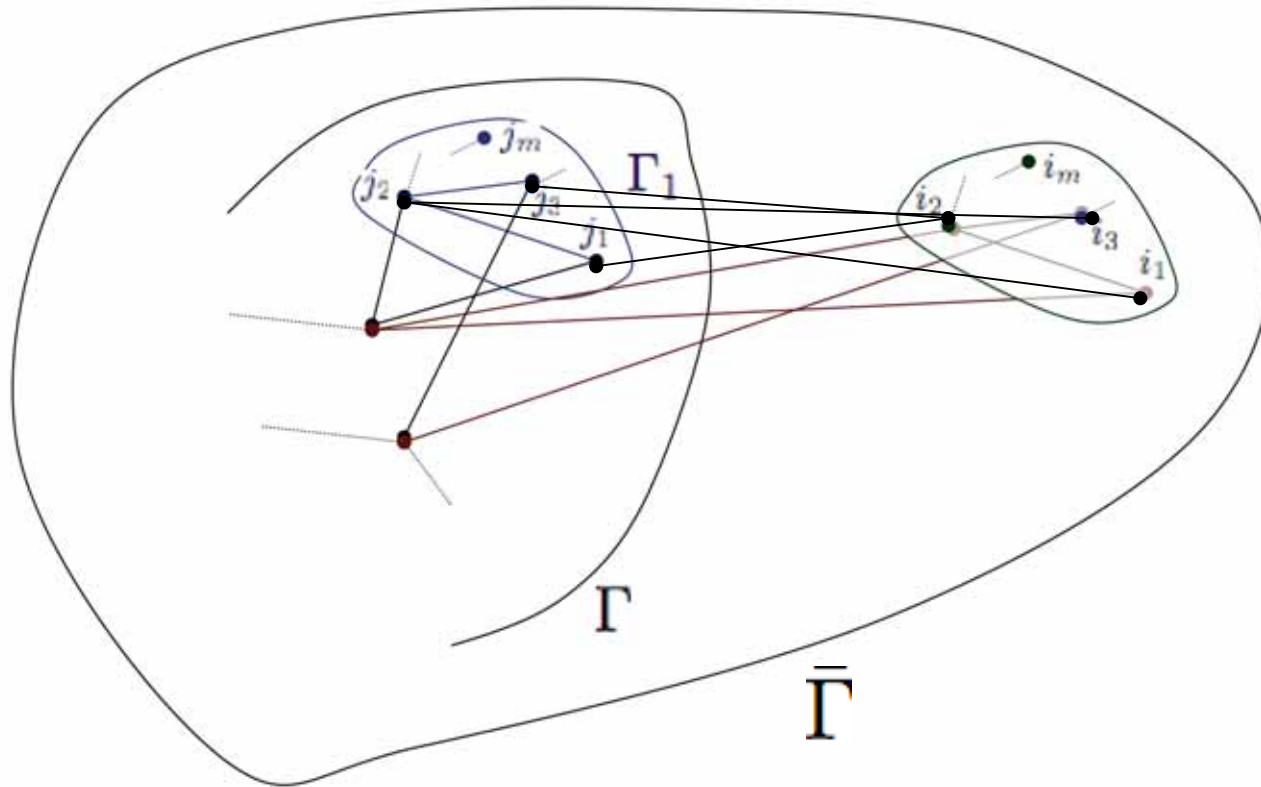
*Motif*: small sub-graph (where as the graph is supposed to be large) containing all edges of the graph between vertices of that subgraph



$$u_{\bar{\Gamma}}^{\lambda} = \begin{cases} u, & \text{on } \Gamma_1 \\ -u. & \text{on the double of } \Gamma_1 \end{cases}$$

# Another motif duplication

*Motif*: small sub-graph (where as the graph is supposed to be large) containing all edges of the graph between vertices of that subgraph



Increment of multiplicity of the eigenvalue 1 by  $m$

# Edge doubling

Doubling an edge that connects vertices  $j_1, j_2$

-- produce eigenvalues

$$\lambda_{\pm} = 1 \pm \frac{1}{\sqrt{n_{j_1} n_{j_2}}}$$

Symmetric about 1 and close to 1 when  $n_{j_1}, n_{j_2}$  are sufficiently large.

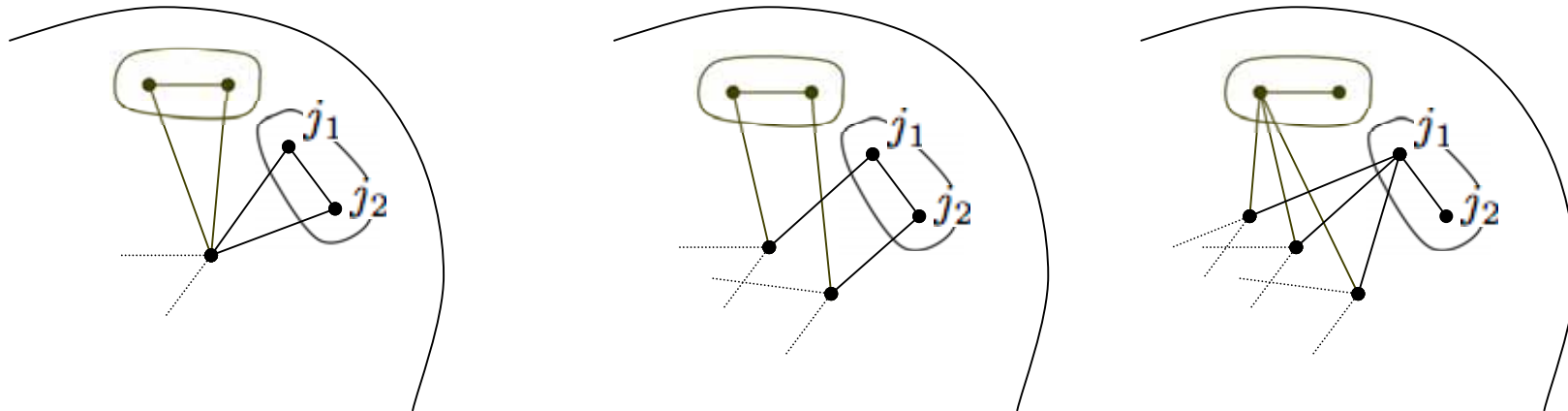
High peak at 1, but not too sharp



# Edge doubling

Doubling an edge that connects vertices  $j_1, j_2$   
with  $n_{j_1} n_{j_2} = 4$

-- produce eigenvalues  $3/2$  and  $1/2$



# Entire graph doubling

Double the entire graph  $\Gamma$  with vertices  $p_1, \dots, p_N$   
 $\Gamma'$  be copy of  $\Gamma$  with vertices  $q_1, \dots, q_N$  and with same connection pattern.

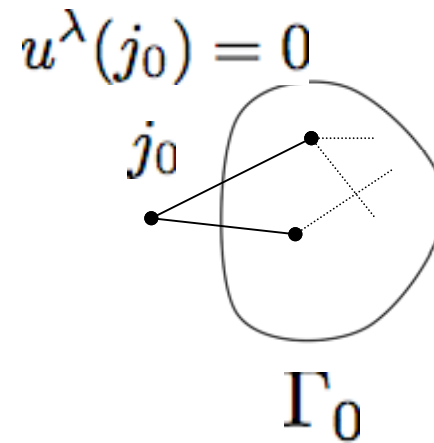
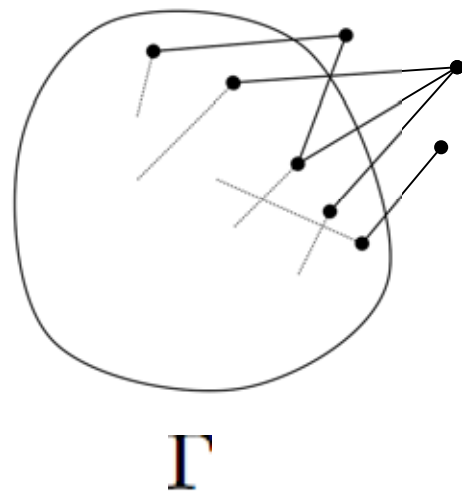
Connect each  $q_\alpha$  to all neighbors of  $p_\alpha$

New graph has the same eigenvalues as  $\Gamma$ , plus the eigenvalue 1 with the multiplicity  $N$ .

protein-protein interaction network do have high multiplicity, not of the order of half of the system size -- subsequent mutations after the genome duplication.

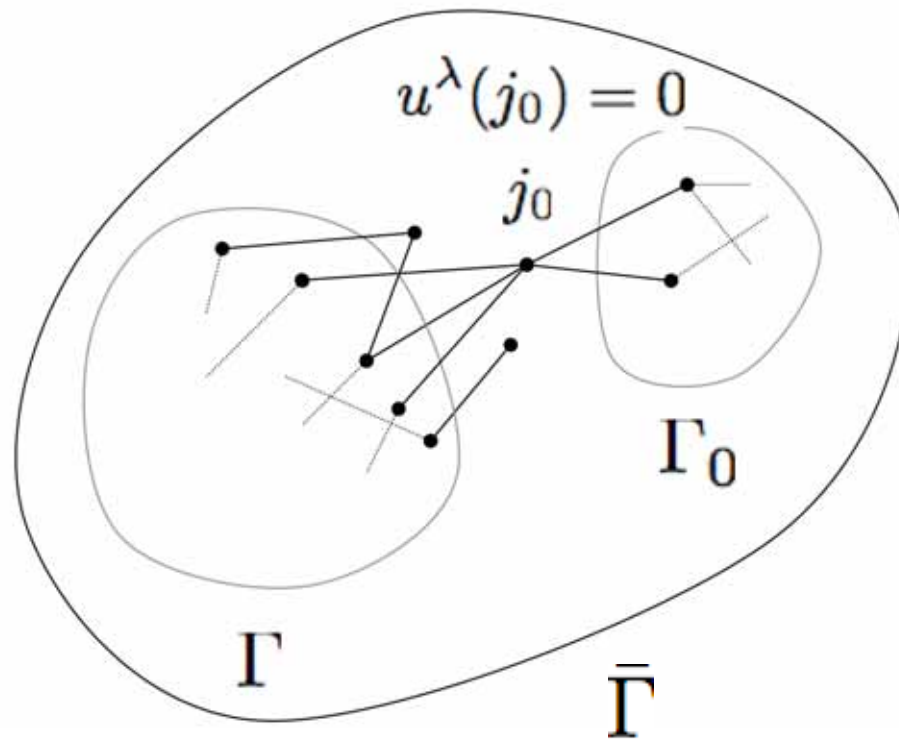
# Motif joining

*Motif*: small sub-graph (where as the graph is supposed to be large) containing all edges of the graph between vertices of that subgraph



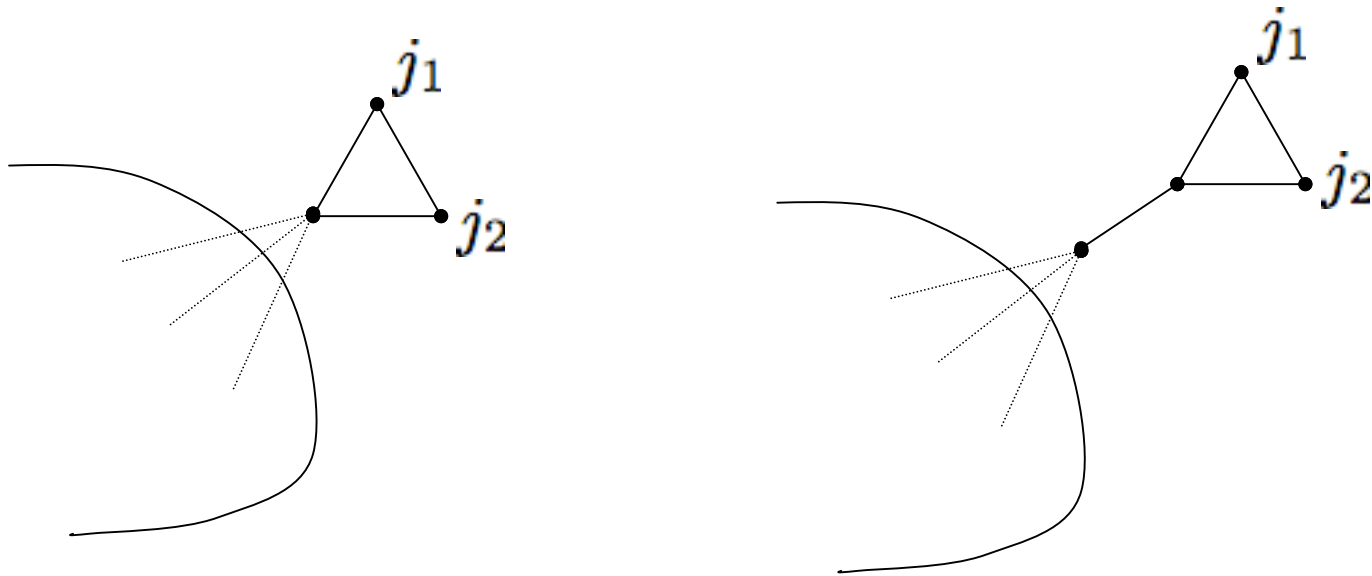
# Motif joining

*Motif*: small sub-graph (where as the graph is supposed to be large) containing all edges of the graph between vertices of that subgraph



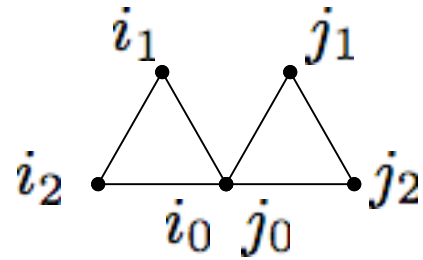
$$u_{\bar{\Gamma}}^\lambda = \begin{cases} u^\lambda, & \text{on } \Gamma_0 \\ 0, & \text{elsewhere} \end{cases}$$

# Triangle joining



$$u_{3/2}(i) = \begin{cases} 1, & \text{for } i = j_1 \\ -1, & \text{for } i = j_2 \\ 0, & \text{otherwise} \end{cases}$$

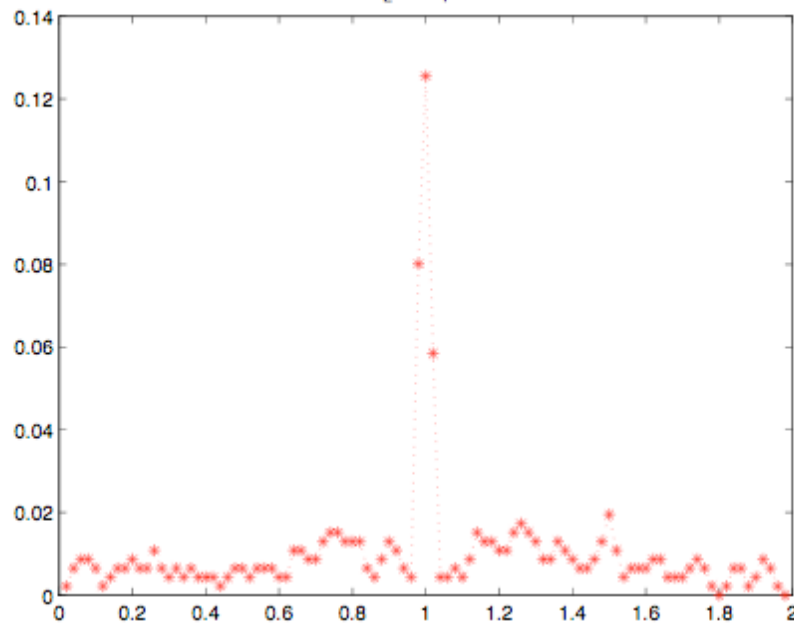
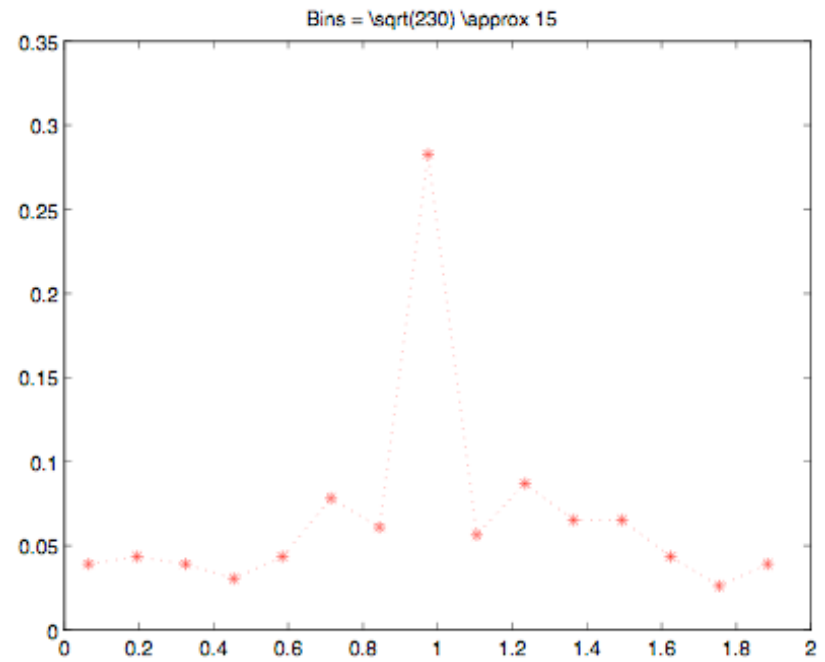
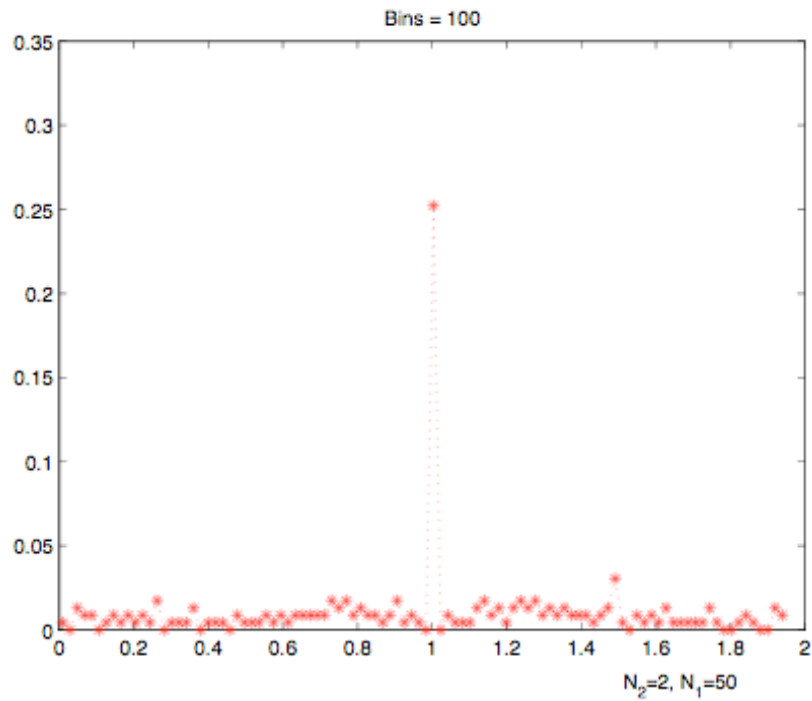
# Triangle joining



Generate not only eigenvalue  $3/2$ , but also  $1/2$

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Protein-protein interaction network *E.coli*. Size = 230

(Overlapping bins such as,  $[0, \frac{N_2}{N_1}]$ ,  $[\frac{1}{N_1}, \frac{N_2+1}{N_1}]$ ,  $\dots$ ,  $[\frac{2N_1-N_2}{N_1}, \frac{2N_1}{N_1}]$ )

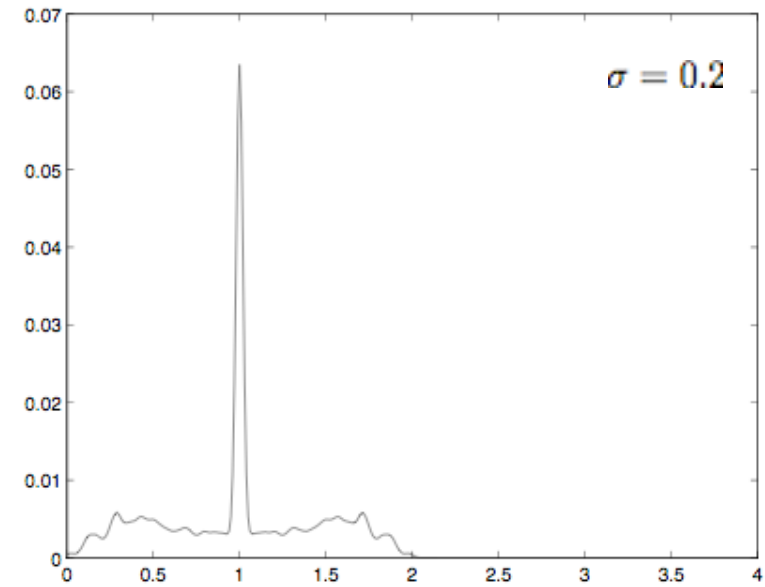
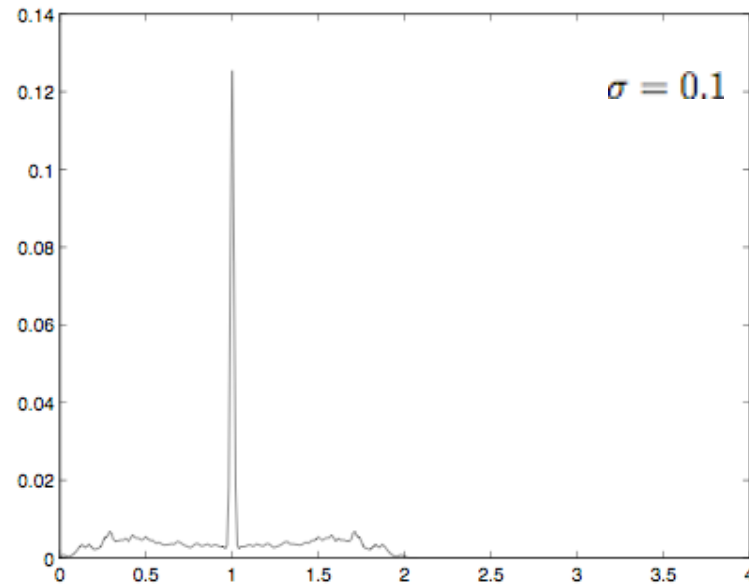


## Convolve with a kernel

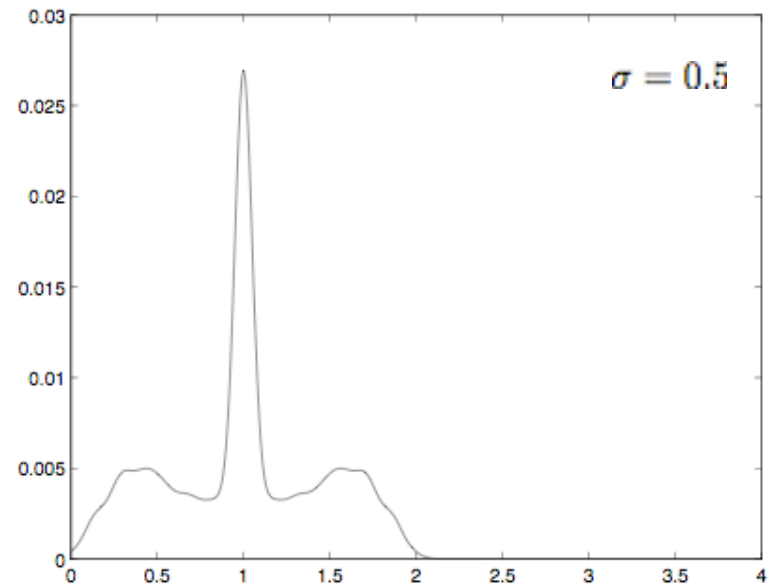
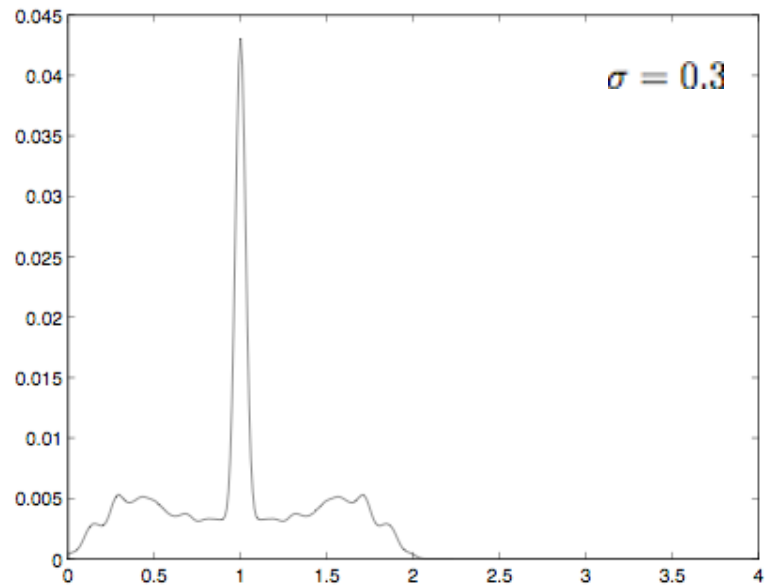
- Gaussian:  $\frac{1}{\sqrt{2\pi\sigma}} \exp\left(-\frac{(x-m_x)^2}{2\sigma^2}\right)$
- Cauchy-Lorentz:  $\frac{1}{\pi} \frac{\gamma}{(x-m)^2 + \gamma^2}$

$$f(x) = \int g(x, \lambda) \sum_k \delta(\lambda, \lambda_k) d\lambda = \sum_k g(x, \lambda_k).$$

# Plots with different kernel values



*C. elegans*

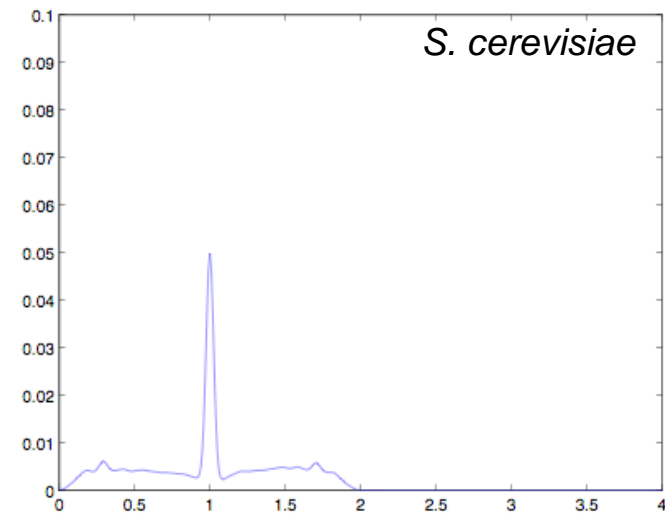
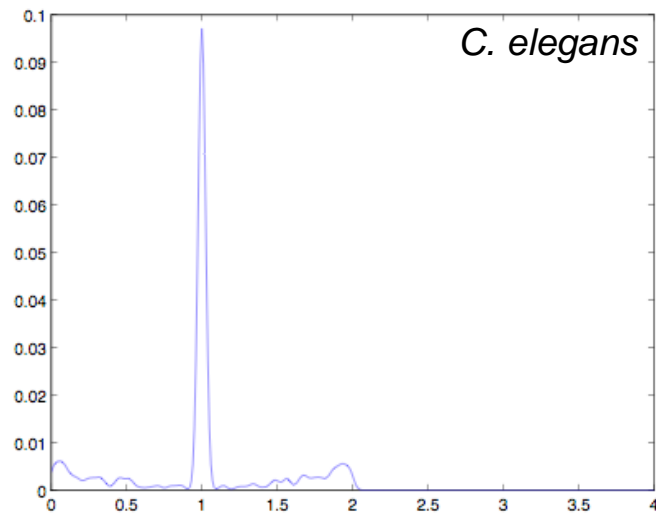
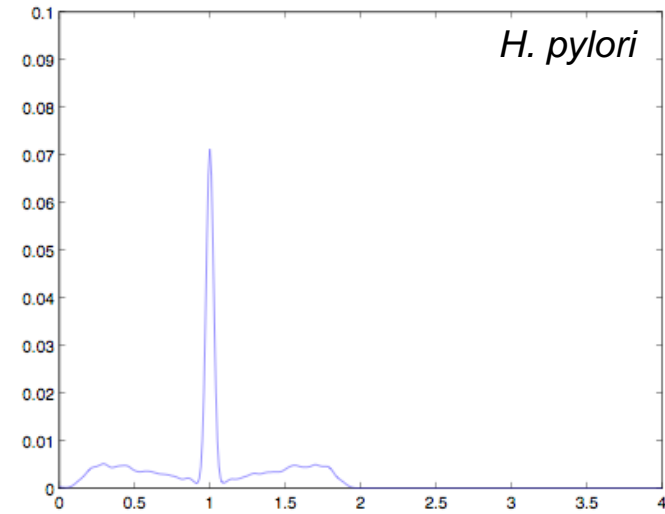
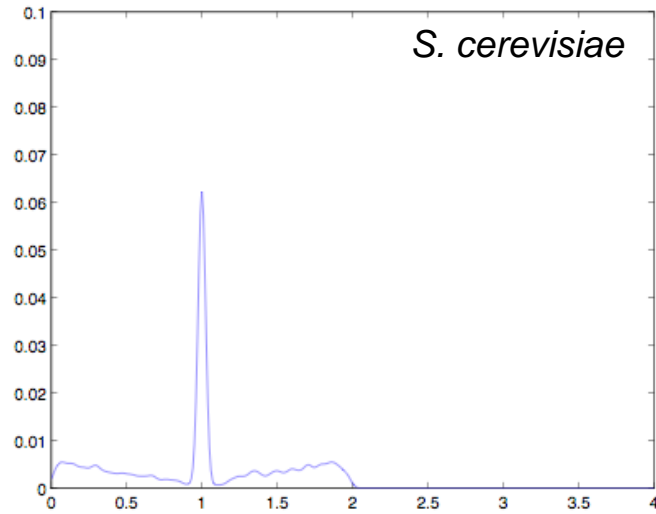


# Overview of the talk

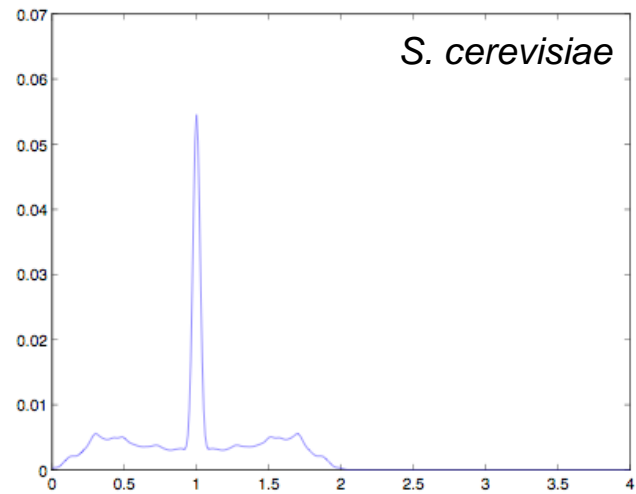
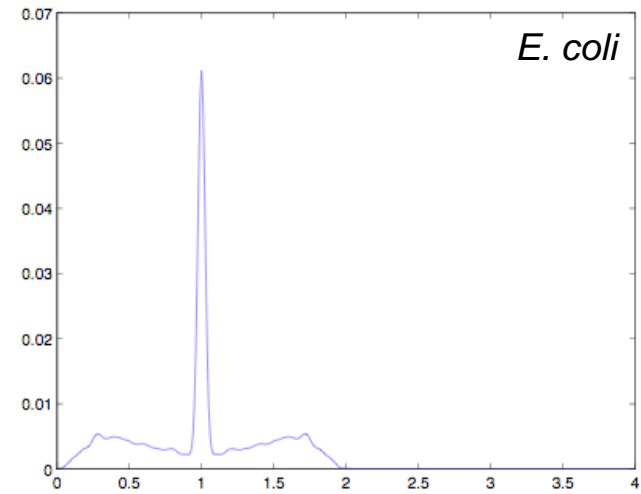
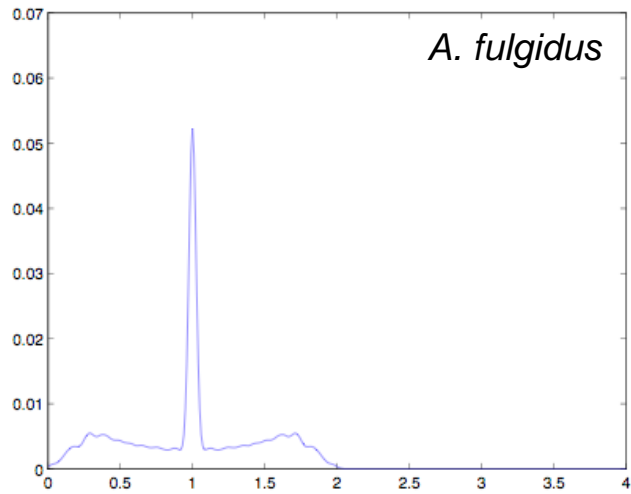
- Introduction
- Graph Laplacian operator
- Eigenfunction and graph structure
- Visualization techniques
- **Qualitative classification of networks**
- Conclusion and remarks

Type I

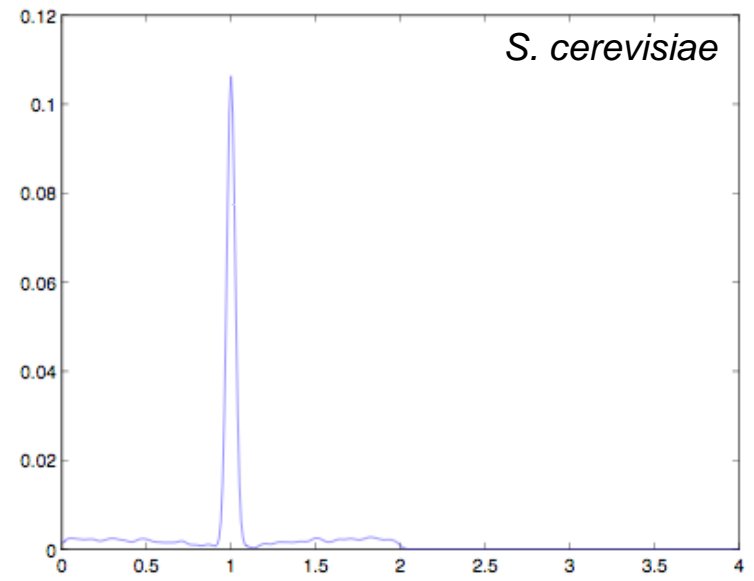
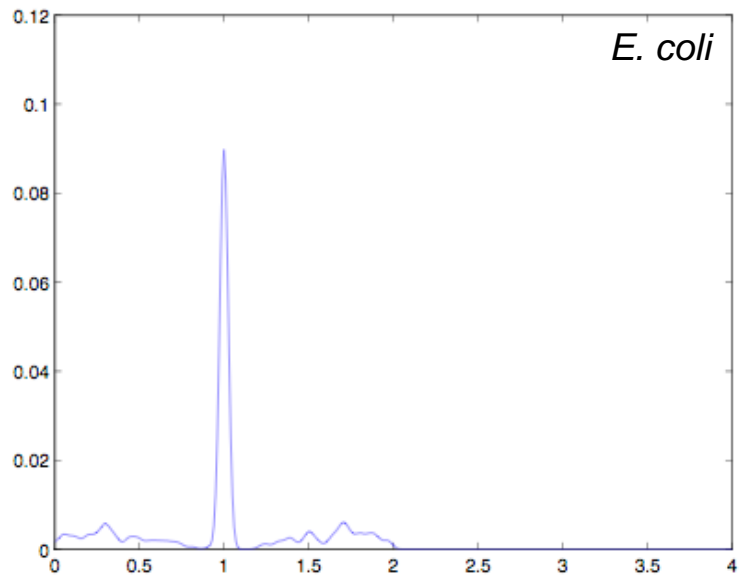
# Protein-protein interaction networks



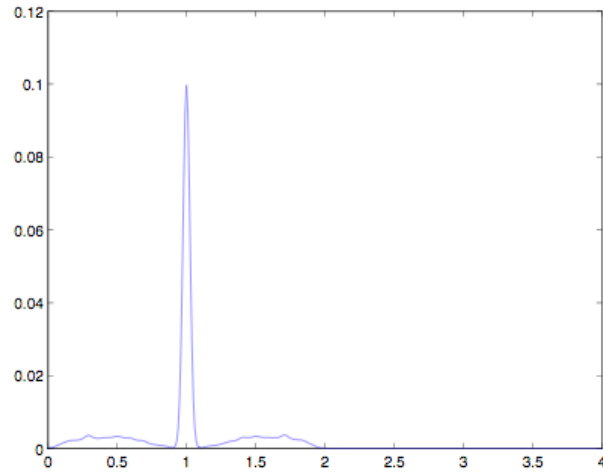
# Metabolic networks



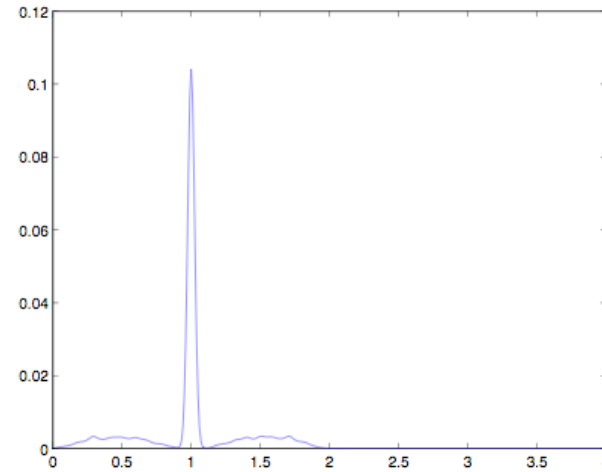
# Transcription networks



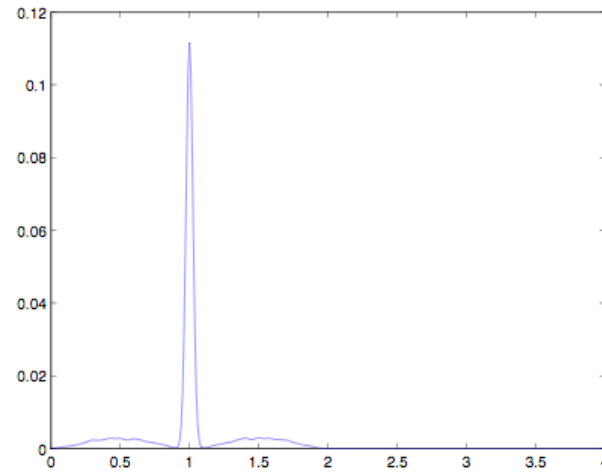
# Autonomous Systems topology of the Internet



AS graph of 1997/11/08



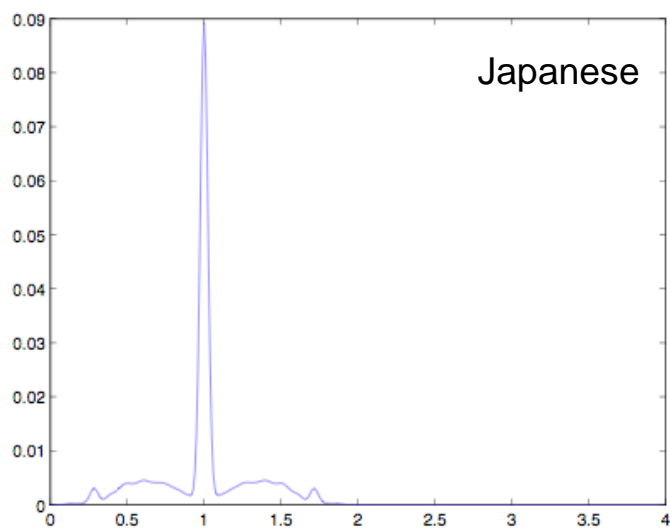
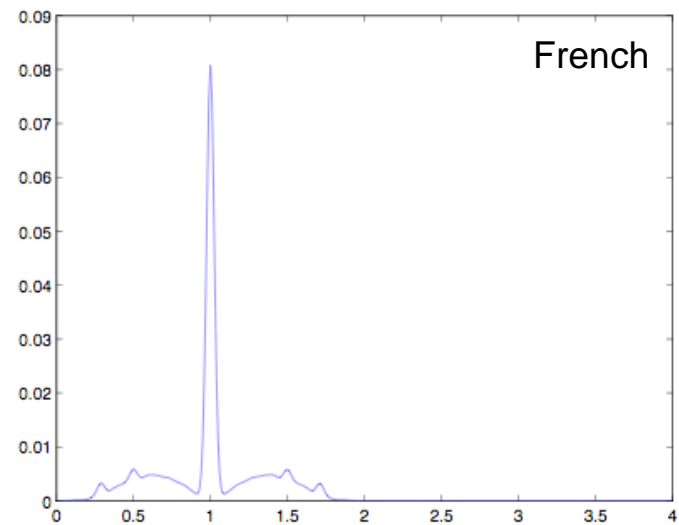
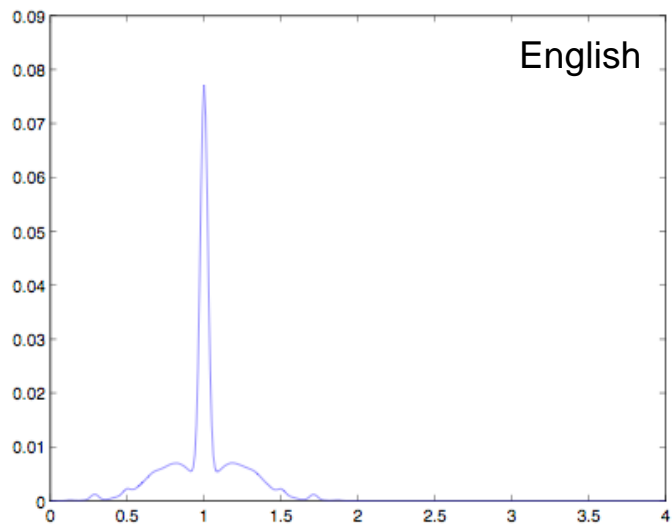
AS graph of 1999/07/02



AS graph of 2001/03/16

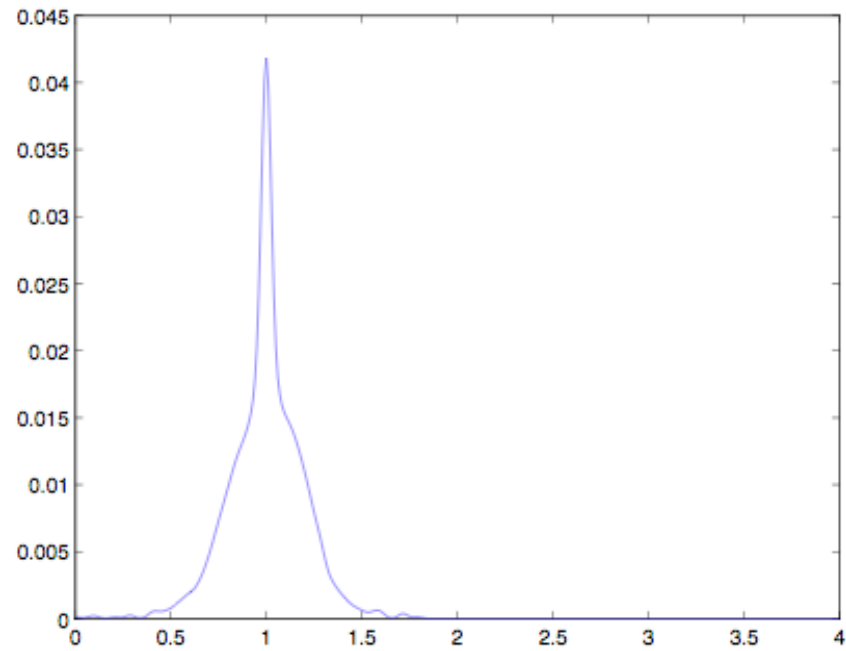


# Word-adjacency networks

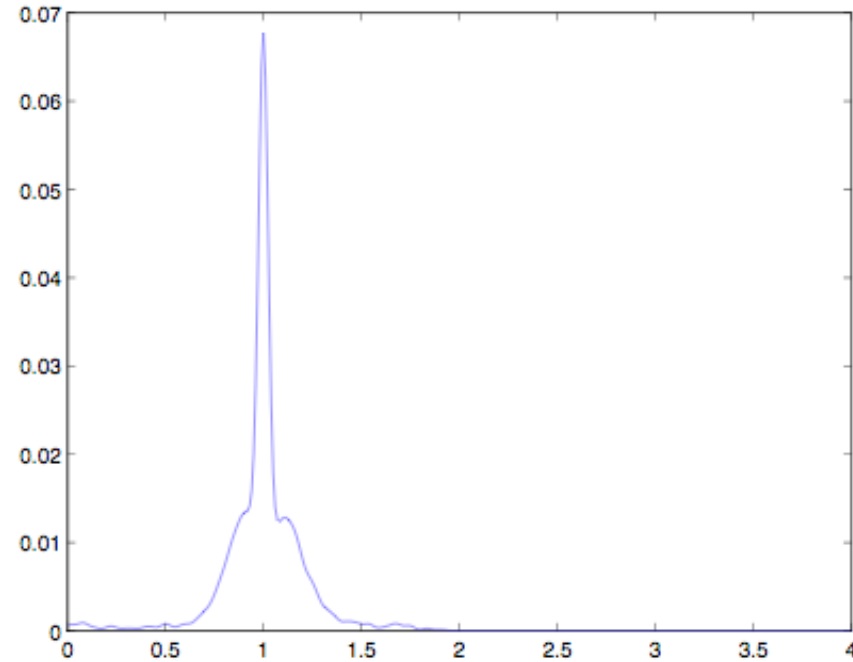


Type II

# Network of hyperlinks between weblogs of US politics

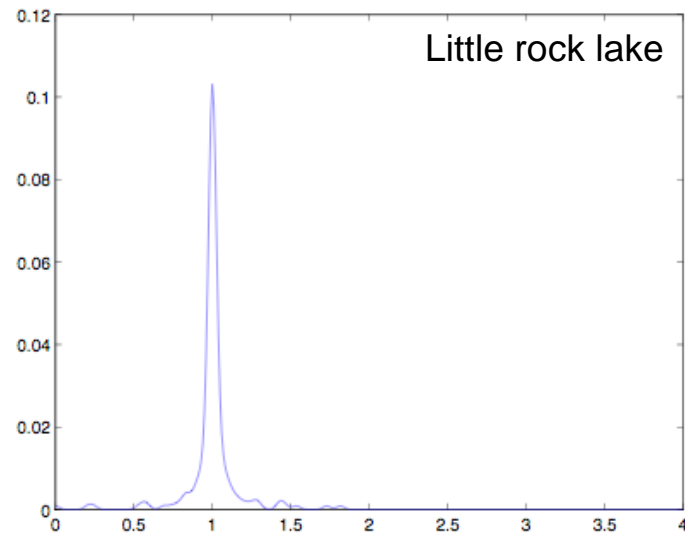
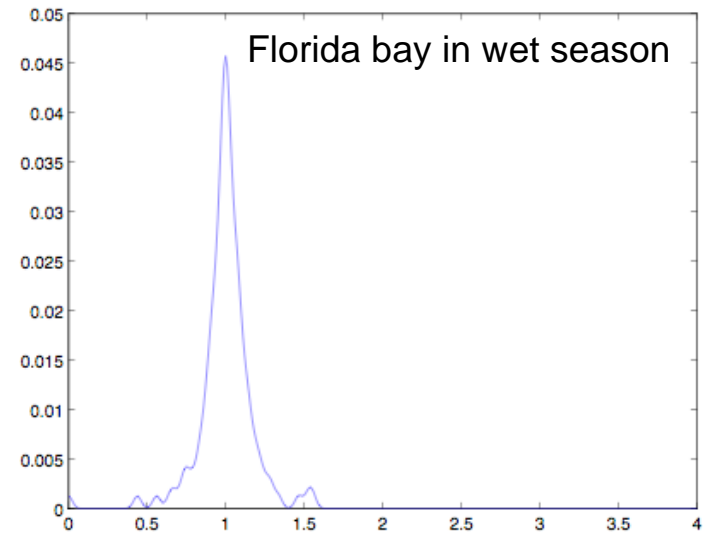
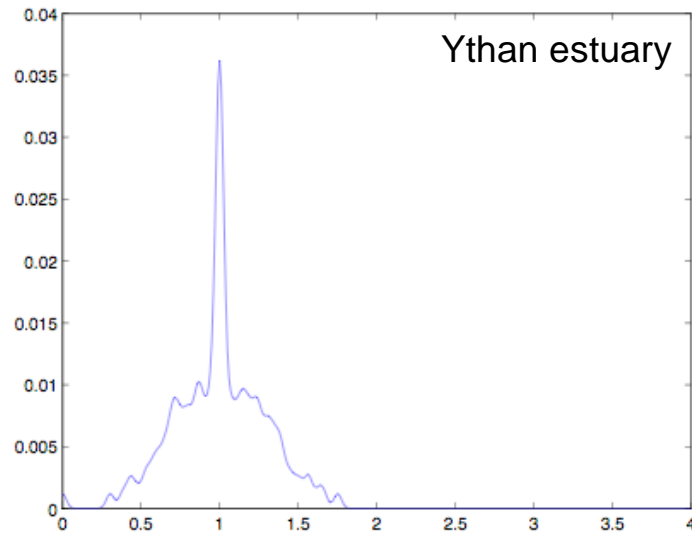


# Network of conformation space

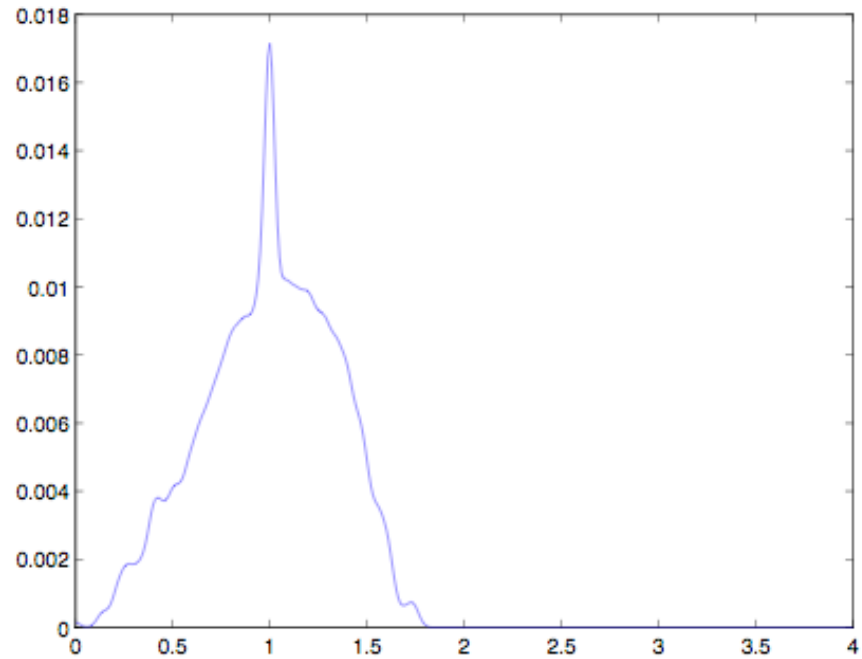


Only conformations that are visited at least 20 times during the simulation are considered in the building of the network.

# Food-web networks

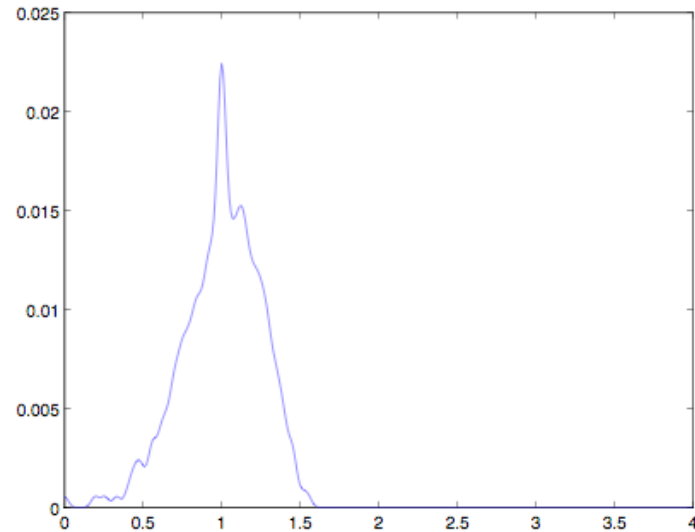


# E-mail interchanges network

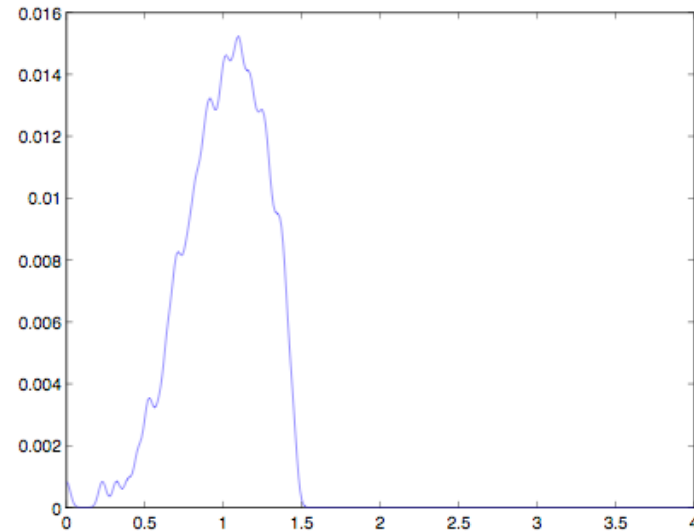


E-mail interchanges between members of the Univeristy Rovira i Virgili (Tarragona)

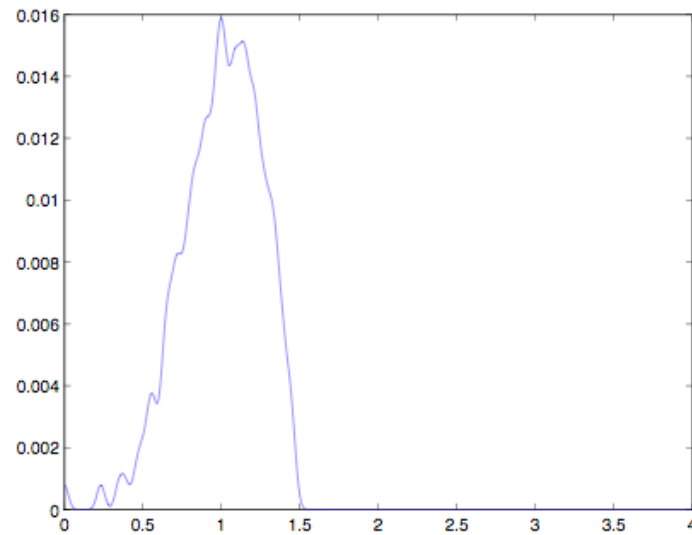
# Neural networks



*C. elegans*



*C. elegans* (animal JSH, L4 male) in RVG region

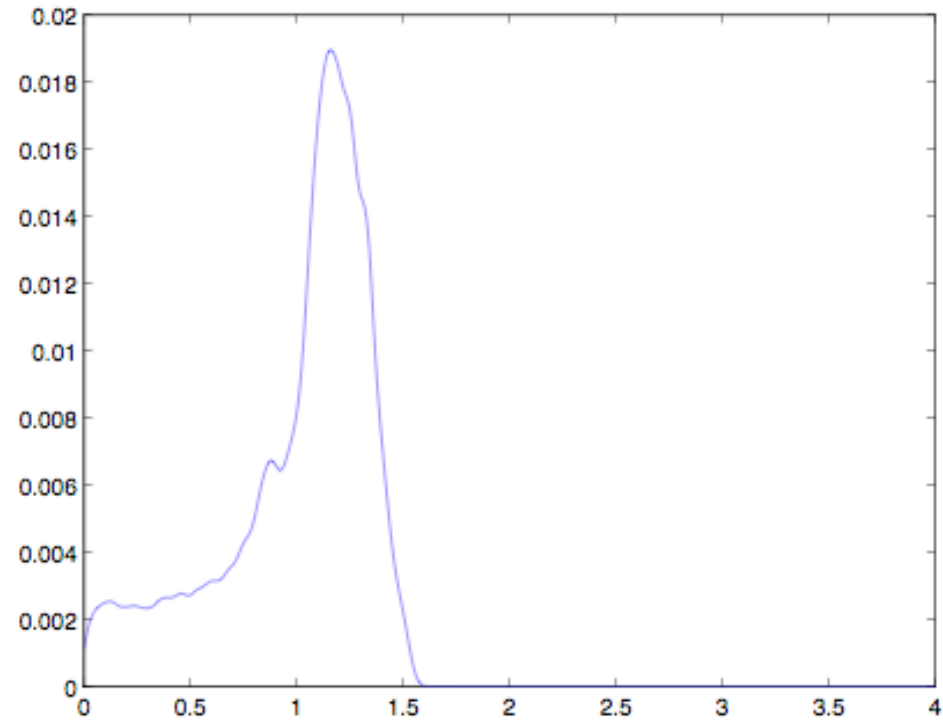


*C. elegans* (animal N2U, adult hermaphrodite) in RVG region

Type III

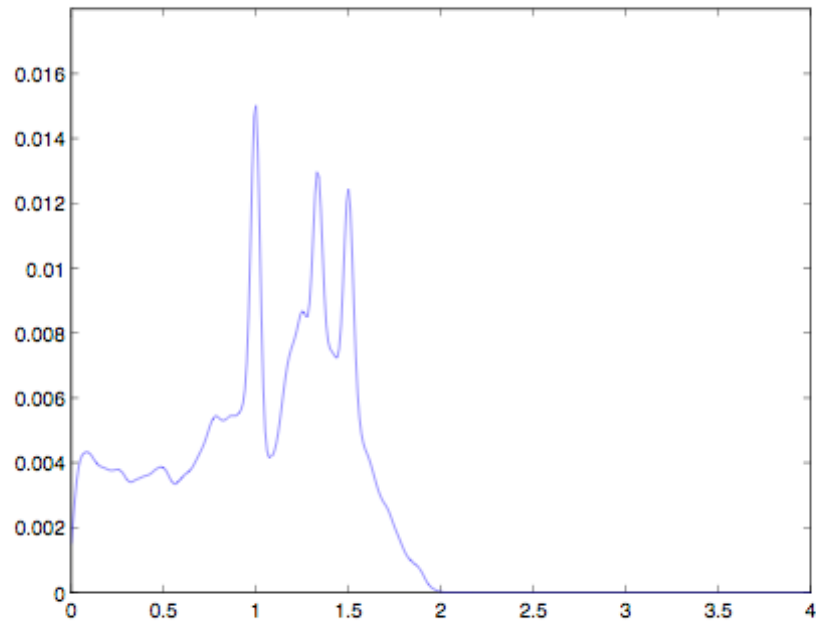


# Power-grid network

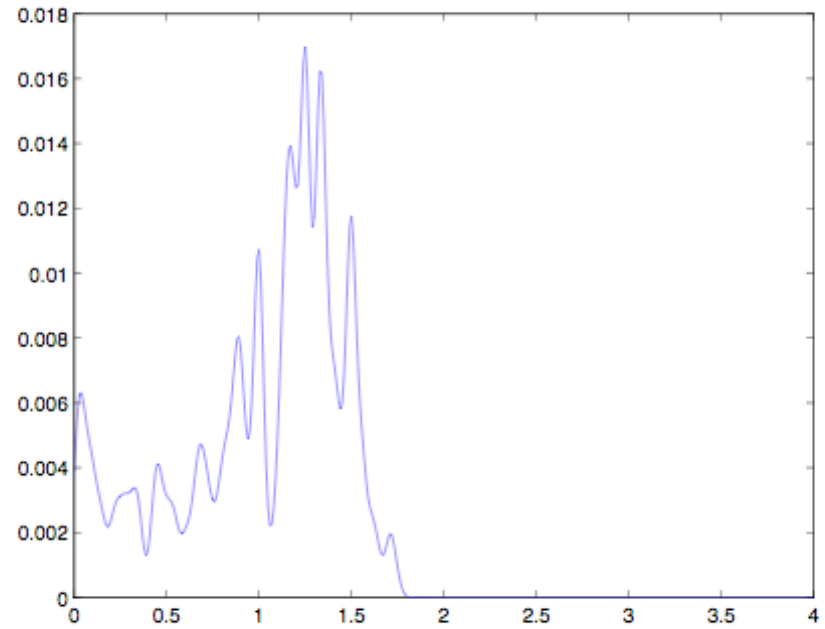


Topology of the Western States Power Grid of the United States

# Networks of co-authorships

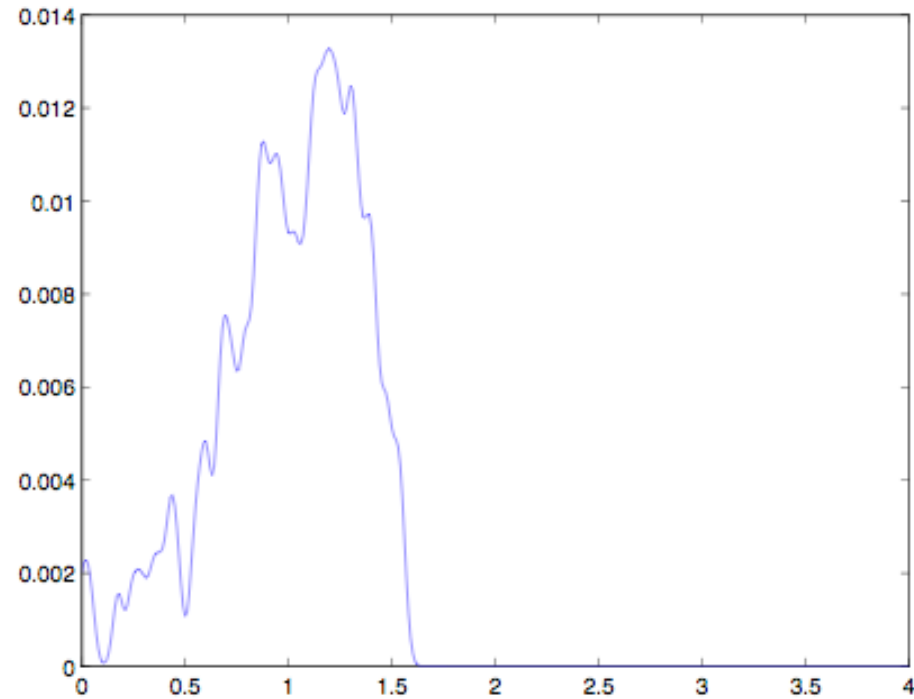


Scientists in high-energy theory



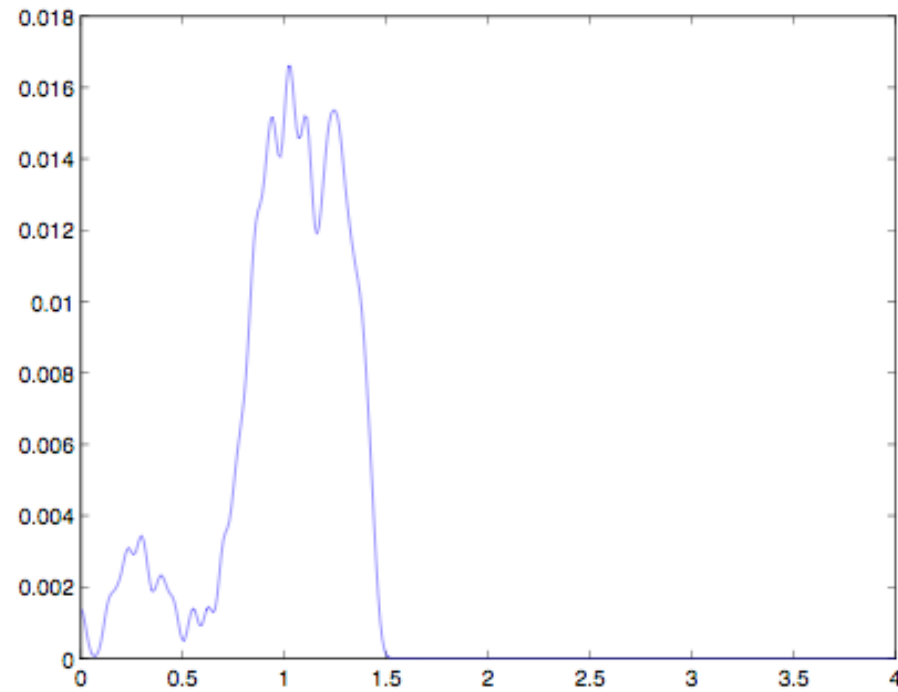
Scientists in network theory and experiment

# Network of co-purchasing of books



Books about recent US politics sold by the online bookseller Amazon.com.

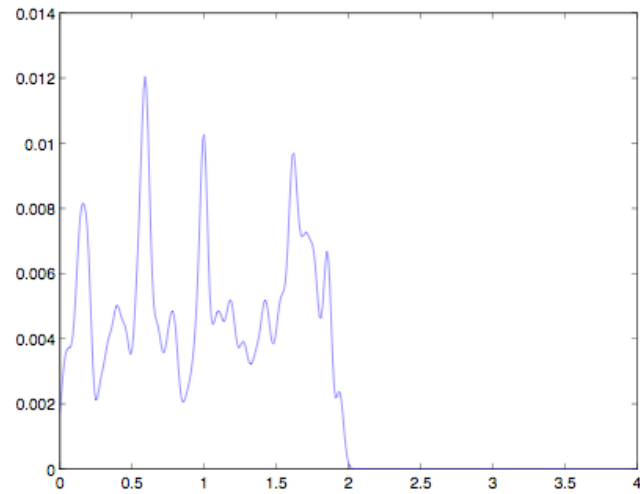
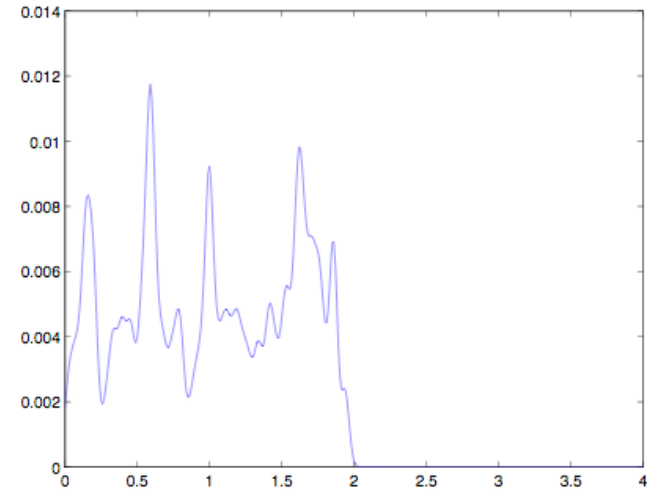
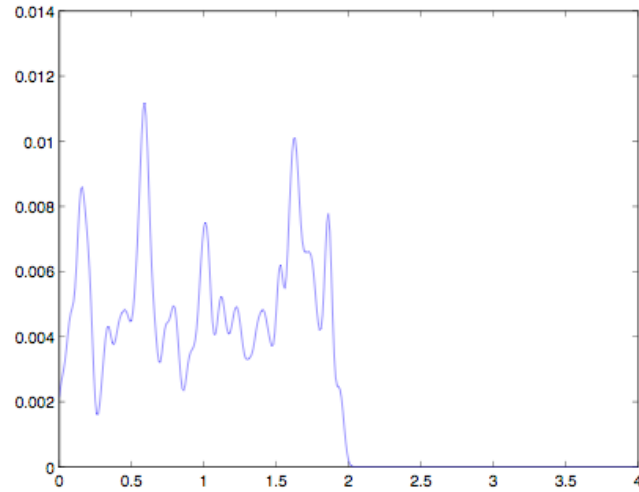
# Network of US football games



American football games between Division IA colleges during regular season Fall 2000

Type IV

# Networks in electronic circuits



# Overview of the talk

- Introduction
- Graph Laplacian operator
- Eigenfunction and graph structure
- Visualization techniques
- Qualitative classification of networks
- **Conclusion and remarks**

Thank you



# Spectral Plot Properties: Qualitative Classification of Networks

