Hierarchical Networks

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- Introduction to networks
- How to model complex networks?
- Clustering and hierarchy
- Hierarchical organization of cellular metabolism
- The *E. Coli* genetic network

Introduction to Networks

Networks in society

- Co-authorship
- Scientific citations
- Friendships
- Sexual contacts
- Movie actor network
- Business relationships







Days of Thunder (1990) Far and Away (1992) Eyes Wide Shut (1999)



Communication networks

- Internet
- > World Wide Web

• And more:

- Power grid
- > Airline routes
- Neuron networks
- > Words linked by synonyms



Networks in the cell



GENOME

protein-gene interactions

PROTEOME

protein-protein interactions

METABOLISM Biochemical reactions

How to model complex networks?

• The Erdős-Rényi model (1960)



The scale-free model

- Real networks grow continuously.
- > Highly connected nodes are more likely to receive new links.

> GROWTH

At each time-step a new node is added with *m* links.

> PREFERENTIAL ATTACHMENT

The probability that an existing node receives a link from the new one is proportional to it's connectivity.

Mean field solution

$$k_i(t) = m \left(\frac{t}{t_i}\right)^{0.5}$$

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$$k_i(t) = \frac{2m^2}{k^3}$$









Clustering and hierarchy

Clustering coefficient

Defined as the probability that a node's neighbors are linked.

> Scale-free model $C \sim N^{-0.75}$

> Real networks have high average clustering coefficients!



• Why?

> Modular organization



- Protein complexes
- WWW communities
- Scientific groups
- Research areas
- Circles of friends

Modular and scale-

free?
♦ Inspired by the *E. Coli* metabolic network
▶ Hierarchical modularity

Hierarchical scale-free model



Hierarchy captured in real networks



Hierarchical organization of cellular metabolism

 $A + B \longrightarrow C + D$

B

D

- Graph representation of metabolism
- **Clustering coefficient is size** independent





• Modules in the *E. Coli* metabolic network



Reduced graph representation



> Clustering





Clusters and their biological function



The *E. Coli* genetic network

Transcription and it's regulation

- Promoter for Promoter regions and operons regulatory gene - Structural loci Regulatory -Transcriptional regulation gene Activation Promoter for lac operon Pi) i) Plac **Operator** Structural Structural Structural SRC 1 gene for gene for gene for u300/CBP β-galacto-RNA Poll β-galactoside β-galactoside sidase transacetylase permease Repression lac Operon transcription RNA pol normal GENE scenario repressor RNA pol. transcription blocked
- Available data on E. Coli
 - ✤ Gene content and activation-repression interactions for 400 operons
 - mRNA concentration of all E. Coli genes in 50+17 experimental conditions

Operon activation states:

***** Observation:

- mRNA levels of genes in an operon can be very different
- Relative levels are very similar across some experiments



clustering based on profile similarity leads to 2-3 clusters



clusters have different average expression levels!

• Next step:

- Only a few "states" of activation?
- Proposed test:



* Modeling

Publications

- "Deterministic scale-free networks", A.-L. Barabási, E. Ravasz and T. Vicsek, *Physica A* 299, 559 (2001).
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 L. Somera, D. A. Mongru, Z. N. Oltvai and A.L. Barabási, *Science* 297,1551 (2002).
- "Hierarchical organization in complex networks", E. Ravasz and A.-L. Barabási, Phys. Rev. E 67, 026112 (2003).
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- "Scale-free and hierarchical structures in complex networks", A.-L. Barabási, Z. Desző, E. Ravasz, S. H. Yook, and Z. Oltvai, Sitges Proceedings on Complex Networks, (2004).
- --- *"Experimental Determination and System-Level Analysis of Essential Genes in E. coli MG1655",* S.Y. Gerdes et al, submitted.

Some networks are not hierarchical



Common features

- ✤ geographical localization
- ✤ economic pressure to minimize link lengths