

Analysis of Complex Systems

Lecture 1: Introduction

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www.dynamic-connectome.org

Preliminaries - 1

Lecturers



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Practicals



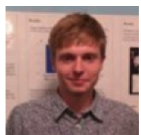
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Preliminaries - 2

- Coursework (specification on NESS):
 - Network analysis 50% due Fri 2 Mar 5pm
 - Dynamical systems 50% due Fri 9 Mar 5pm
- Module contents:
 - Introductory lecture (today)
 - Network analysis (1st week and next Monday)
 - Dynamical systems (2nd week)
 - Dynamical systems (3rd week)

Preliminaries - 3

- Practical uses the widely used Matlab environment (installed on your cluster)
- To try at home:
 - Free licence from university IT service
 - Matlab student bundle
(10 included toolboxes £55)
 - Matlab student version (£29)
 - free version Octave (not 100% compatible)

Team (current and alumni)

Luis Peraza
PostDoc



Frances Hutchings
PhD student



Nishant Sinha
PhD student



Sol Lim
PostDoc



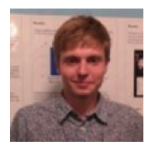
Roman Bauer
Now **faculty**



Chris Hayward
PhD student



Joe Necus
PhD student



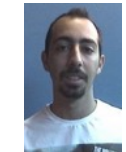
Peter Taylor
Now **faculty**



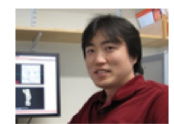
Yujiang Wang
Now **faculty**



Chris Papasavvas
PostDoc



Jinseop Kim
Now **faculty**



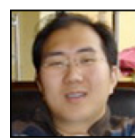
Michael Mackay
PhD student



Chris Thornton
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Collaborators

Justin Dauwels

Sydney S. Cash

M. Brandon Westover

Funders



<http://www.dynamic-connectome.org>



[@ConnectomeLab](https://twitter.com/ConnectomeLab)

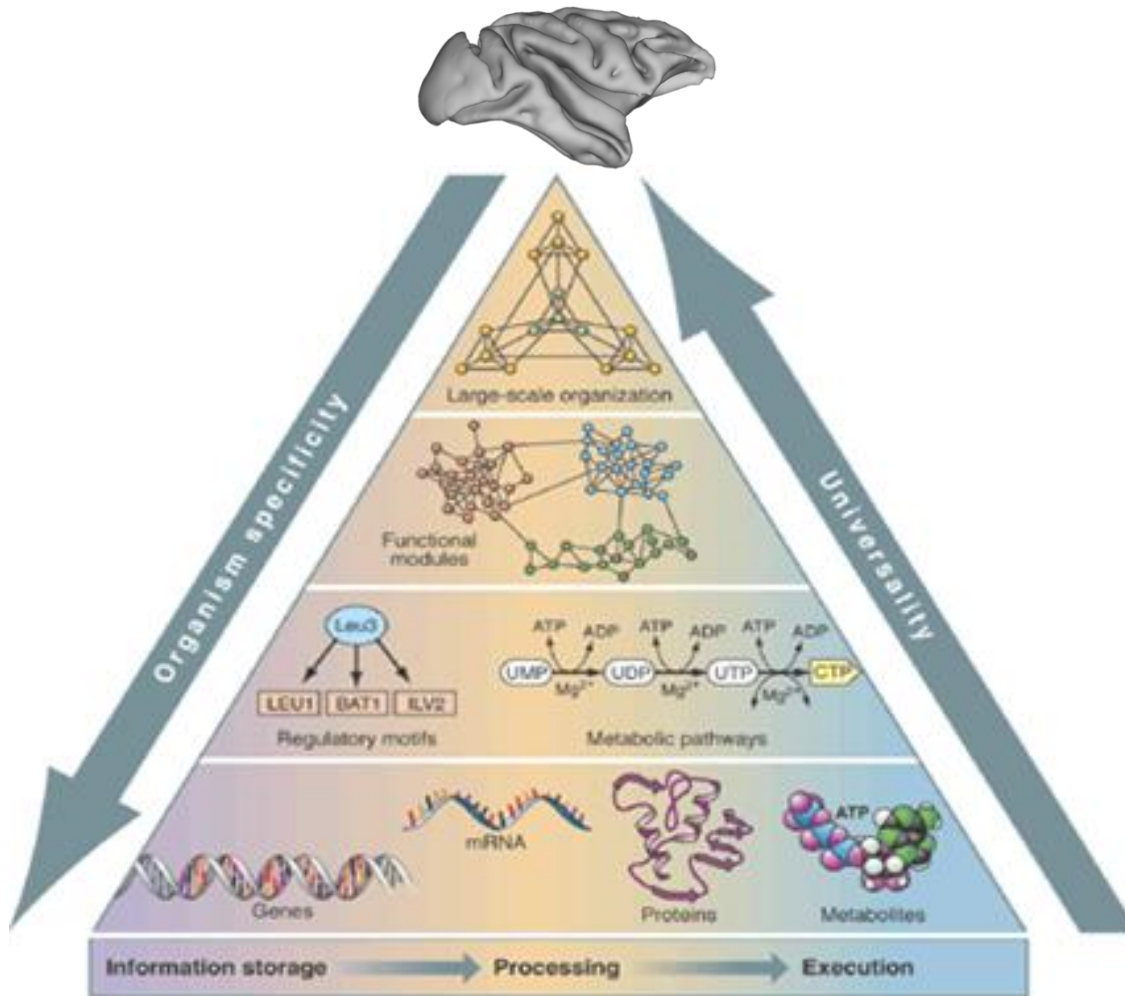
Outline

- Complex Biological systems and selected problems
- Complex Systems approach
- Network analysis
- Dynamical systems

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Complex Biological Systems

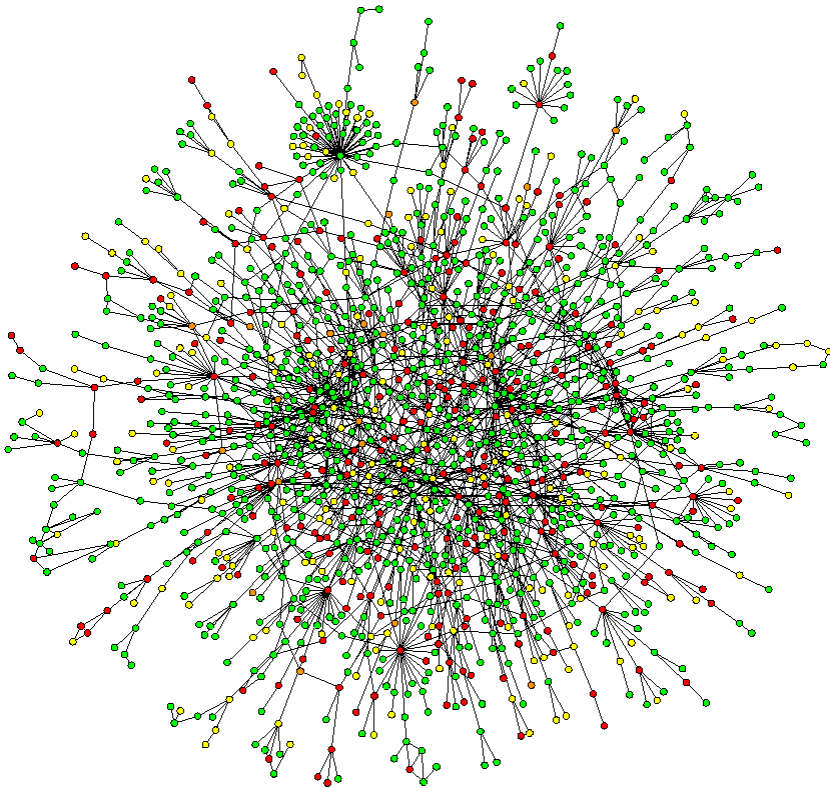


Life's Complexity Pyramid [Z. N. Oltvai and A.-L. Barabási, Science 298, 763 (2002)]

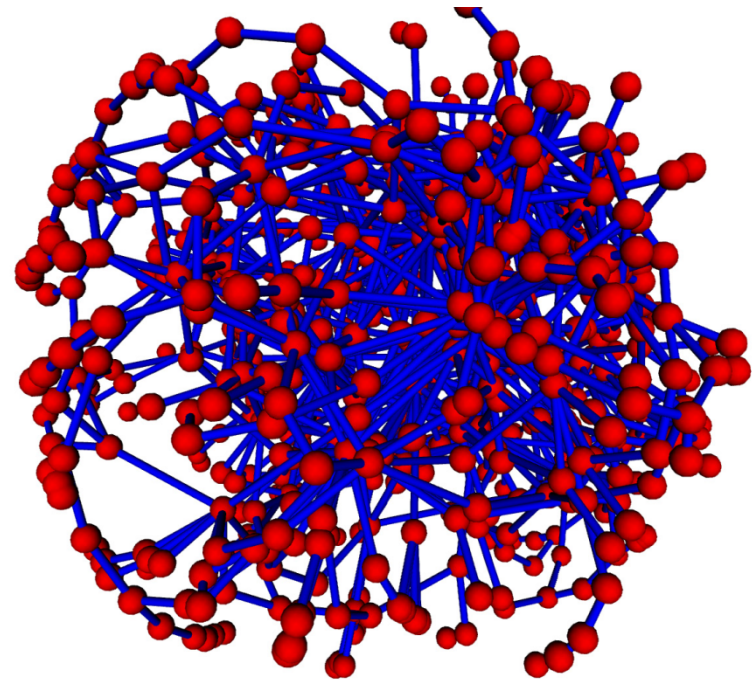
Cellular systems

- Interactions between proteins, lipids, metabolites

Yeast (*Saccharomyces cerevisiae*) protein-protein interaction network



(Jeong et al., Science, 2000)



(Hilgetag & Kaiser, Bioforum, 2005)

Graph Analytics



Government/Security

- Patterns of Activity Analytics
- CyberThreat Discovery
- Tax Fraud Discovery
- Crime Prediction



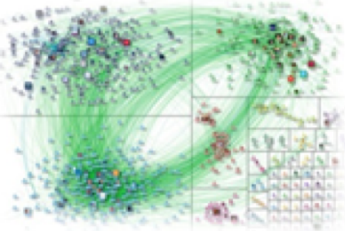
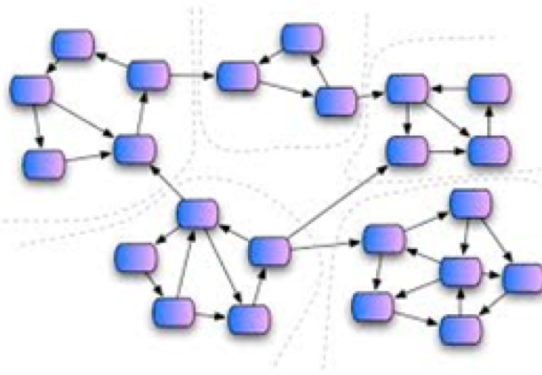
Healthcare

- Personalized Treatment
- Fraud Detection
- Efficacy of Care
- Adverse Event Clustering
- Disease Prediction



Energy/Resources

- Location Discovery
- Field Production Analysis
- Contingency Analysis
- Climate Modeling



Telecom/Media

- Influencer Discovery
- Churn Analytics
- Behavior Analytics

Life Sciences

- Drug Discovery
- Drug Repurposing
- Clinical Trial Mining



Financial Services

- Market Sensing
- News/Trading Analytics
- Counterparty/Risk
- Insider Threat
- AML/Compliance

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- Complex Biological systems and selected problems
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Systems

- System units and interactions between them
- E.g., gene regulation system: the units are genes, the interactions are the regulatory interactions
- E.g., protein interaction systems: the units are proteins, the interactions are the protein interactions

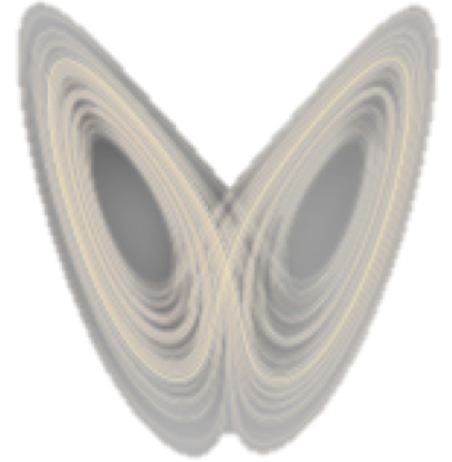
System evolution

- System state: set of units and set of current interactions between units
- Rules: a set of active interactions is followed by another set of interactions, new units may become active depending on the system's state, some units may stop to be active

System analysis

- Description of the system *state*:
current units and interactions
- Analysis of the system's *evolution*:
rules of state change, description of the
state changes for some limited or
unlimited future
- What is a *complex system*?

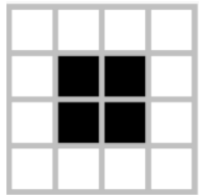
Complex systems



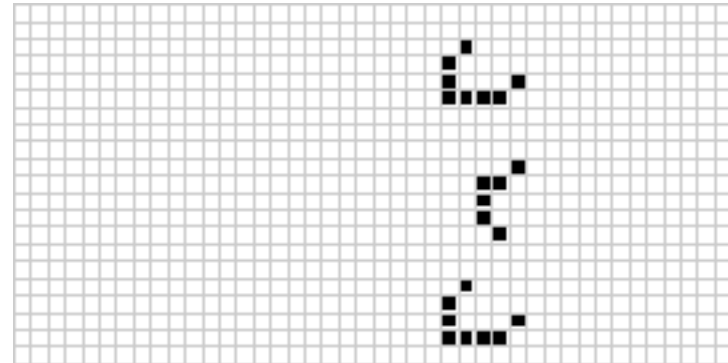
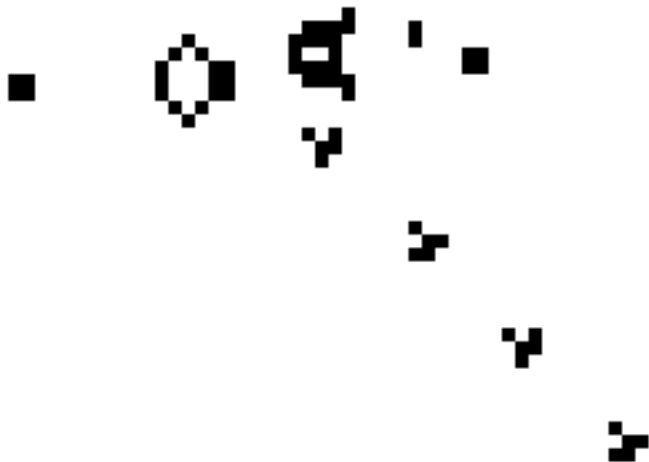
- A complex system is a system composed of interconnected parts that as a whole exhibit one or more properties not obvious from the properties of the individual parts (**emergence**)
- Not obvious: hard to predict or unpredictable (**dependence on initial conditions**)
- E.g., the evolution of a cell, stock market, weather forecast (Edward Lorenz)

Emergence

Conway's Game of Life (the origin of *cellular automata*)

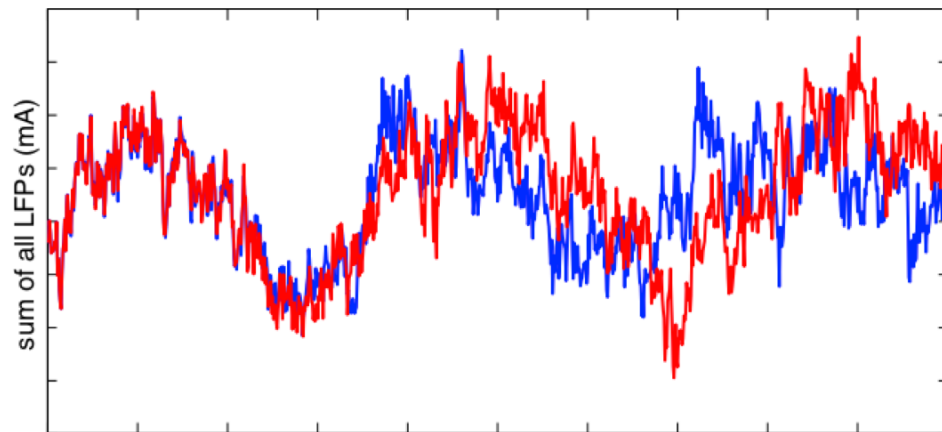


1. Death: if the count is less than 2 or greater than 3, the current cell is switched off.
2. Survival: if (a) the count is exactly 2, or (b) the count is exactly 3 and the current cell is on, the current cell is left unchanged.
3. Birth: if the current cell is off and the count is exactly 3, the current cell is switched on.



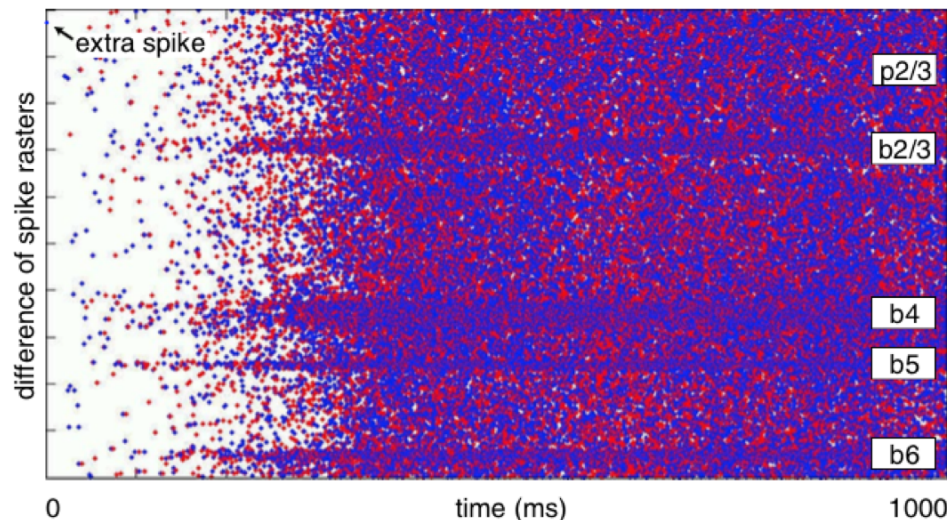
Martin Gardner (1970). Mathematical Games – The fantastic combinations of John Conway's new solitaire game "life". Scientific American 223. pp. 120–123

Dependence on initial conditions



Simulation of 100,000 neurons:
Effect of one additional activation
at the beginning of the simulation

-> divergent system dynamics after
half a second

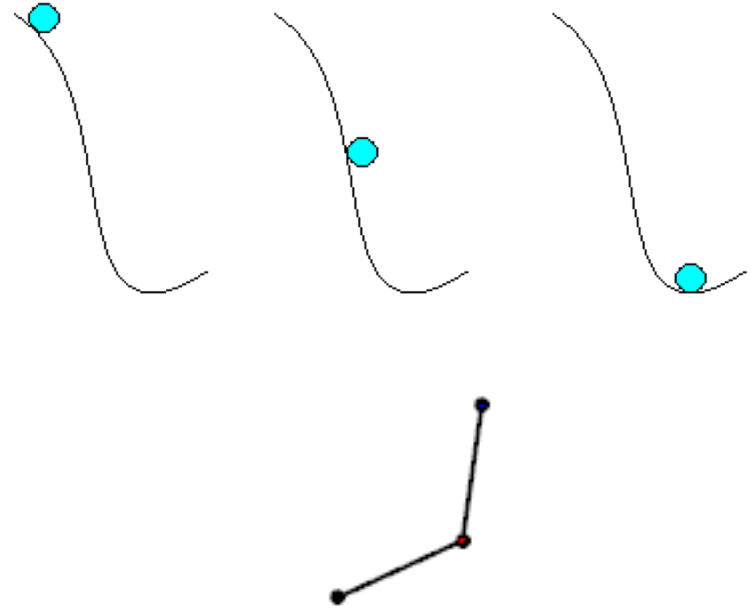


**A small change in initial conditions
can lead to a massive change in the
long-term behaviour of the system**

Izhikevich & Edelman, PNAS, 2008

No analytic solutions for long-term behaviour

- The description of the evolution of the system cannot be summarized in a simple way
- Simple system: ball rolling down on a slope into a valley
- Complex system: double pendulum



Gene regulation systems

- Couple / several / many participant genes
- Possibly highly variable balance of product protein quantities
- → complex system

Protein-protein interaction systems

- Many units and interactions
- Hard to predict and describe the evolution of this system
- Hard to predict the effects of simple changes (e.g., blocking the activity of a couple of proteins)
- → complex system

Analysis methods

- Simplification and abstraction
- Capture key features of the system
- Analysing a model system
- Ignoring non-key features and factors
(and hopefully not missing key features!)

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- **Network analysis**
- Dynamical systems

“Network Science”

Rapidly expanding field:

Watts & Strogatz, *Nature* (June 1998)

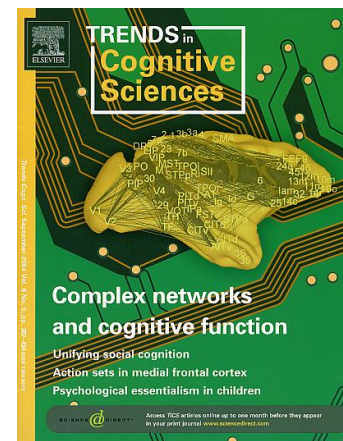
Barabasi & Albert, *Science* (October 1999)

Modelling of SARS spreading over the airline network
(Hufnagel, *PNAS*, 2004)

Identity and Search in Social Networks
(Watts et al., *Science*, 2002)

The Large-Scale Organization of Metabolic Networks.
(Jeong et al., *Nature*, 2000)

Organisation and dynamics of neural systems
(Sporns, Chialvo, Kaiser, Hilgetag, *Trends in Cognitive Sciences*, 2004)



Why use network analysis?

- Many diseases are due to failures of multiple genes
→ single gene knockouts will not be sufficient
- Combinatorial explosion of possible gene knockout combinations (example for 1k genes):

single gene knockout: $1000 = 10^3$

double gene knockout: $1000 * 999 \approx 10^6$

triple gene knockout: $1000 * 999 * 998 \approx 10^9$

...

- Network analysis can help to predict combinations which are more likely to lead to performance changes
→ reduced costs for drug discovery (hopefully!)

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- Complex Biological systems and selected problems
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- **Dynamical systems**

Dynamical systems

- E.g., gene regulatory networks
- Formulation as a set of interdependent differential equations
- $dX / dt = a * X + b * Y$
- $dY / dt = c * X + d * Y$
- X and Y are quantities of proteins

Why use dynamical systems theory?

- Theoretically feasible:
many parameters for computer simulations can now be measured (e.g. gene expression or protein interaction data)
- Computationally feasible:
Increase in processing power
(quad-core microchips today; 100-core in 2010)
- Economically desirable:
In silico experiments are cheaper than wet lab experiments
→ **simulation-based science** (e.g. e-cell project)

Summary - 1

- The state of systems (Cells, genome, organisms) is given by the set of units or nodes (e.g. proteins) and interactions or edges (e.g. protein-protein interaction) between units
- The state of the system evolves over time (e.g. gene expression or metabolite concentration changes) and can lead to observable changes in behaviour, e.g. anaerobic metabolism → aerobic metabolism (not all state changes lead to large-scale changes!)

Summary - 2

- Problem: describe the evolution of protein quantities knowing gene regulatory networks
- Problem: find key functional proteins to damage or restore cell functionality
- Complex systems
- Analysis of complex systems
 - Graph structure analysis
 - Dynamical systems analysis

Q&A - 1

1. What could be nodes or edges for the interaction between cells?
2. Is it true that cell behaviour can be seen as direct interactions between nucleic acids ?
3. Is it true that proteins are encoded by genes ?
4. Is it true that the evolution of a gene regulatory network is relatively simple and can be described easily ?

Q&A - 2

5. Is it true that a way to find antibiotic targets is to analyse the protein interaction network of the bacteria ?
6. Is it true that the dynamical systems analysis can be applied to the analysis of protein interaction networks ?
7. Is it true that graph models use nodes and edges to describe a system ?