

Computational Systems Biology: Biology X

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L#1:(September-13-2010)
Cancer and Signals

Outline

1 Administrivia

2 Theme

*Cancer's a Funny Thing:
I wish I had the voice of Homer
To sing of rectal carcinoma,
This kills a lot more chaps, in fact,
Than were bumped off when Troy was sacked....*

–JBS Haldane, *The Enchantress of Florence*, 2008.

Outline

1 Administrivia

2 Theme

Administrivia

- **Instructor:** Bud Mishra
- Room 1002, 715 Broadway
- email: mishra@nyu.edu
- phone: 212-998-3464
- Office Hours: Mondays 1:30 pm - 2:30 pm (and by appt)

Administrivia

- BIOLOGY X
- **Course Details:** G22.3033-002
|| Bioinformatics
- **Time and Place:** Monday, 5:00 PM - 6:50 PM EST
|| Room 1221, 719 Broadway
- **Number of Credits:** 3 credits
- **Course Work:** Software Project, Analyzing Genetics Data, Review Articles
- **Languages of Choice:** R (May be Python, Matlab, Mathematica — But no Perl please)

Some Ideas

- Signaling is a well-studied phenomenon both in evolutionary game theory and in cell biology.
- In game theory, signaling frameworks have been used to study the evolution of such fundamental phenomena as conventions and cooperation, while in biology, signal transduction has been extensively studied as a basic ingredient to multicellularity, enabling cells to communicate and coordinate.
- However, approaches that span both fields are scarce.

- In this course, we explore the idea of viewing multicellular organisms as signaling systems in the game-theoretic sense, attempting to unify these two perspectives on signaling.
- A multicellular organism corresponds to a population of cells in a cooperative state, with a working signaling system in place.
- We will discuss how the evolution of such a system may be modeled. Then, we will in particular be interested in the breakdown of cooperation, leading to an interpretation of cancer as a disease of multicellularity.
- The course will be as self-contained as possible and include introductions to evolutionary game theory and signaling systems, signal transduction in cell biology, and the biology of cancer.

Text Books

- **Required Textbook (1):** *Signals: Evolution, Learning, and Information* || Brian Skyrms || Oxford University Press || 2010.
- **Required Textbook (2):** *Biology of Cancer* || Robert A. Weinberg || Garland Science, 2006.
 - **Recommended textbook (1):** *Game Theory Evolving* || Herbert Gintis || Princeton University Press || 2000
 - **Recommended textbook (2):** *Computational Biology of Cancer* || Dominik Wodarz and Natalia L. Komarova || World Scientific Publishing Company || 2005
 - **Recommended textbook (3):** *Information Theory, Inference & Learning Algorithms* || David J. C. MacKay || Cambridge University Press || 2002

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Cancer and Signals

- **Main Thesis**
- Cancer is a complex disease.
 - 1 *It is associated with the natural somatic **evolution**.*
 - 2 *It has its origin in **multicellularity**.*
 - 3 *It makes use of every **mechanism in biology**.*

Areas we wish to touch on...

- *Bioinformatics to study Mechanisms in Biology*: Genomics, Transcriptomics (c- and ncRNA), Proteomics, Metabolomics and Signaling
- *Structure of Multicellularity*: Signaling and Signaling Games; Multicellularity, Information Theory and Rate Distortion Theory
- *Evolutionary Structure of Somatic Modifications*: Repeated Games, Signaling Games and Neologism

Let us think about these inter-connected questions from a single global perspective...

A Tentative Syllabus

I would like to focus this course on three basic questions...

- 1 **Biology** of **cancer**
- 2 **Signaling** game models of multicellular **biology**
- 3 **Information** theoretic utility functions in **signaling** games

Possible Sets of Lectures

- **Lecture 1:** Causes of Cancer
- **Lecture 2:** A Very Brief Introduction to Game Theory
- **Lecture 3:** Introduction to Biology
- **Lecture 4:** The Nature of Cancer
- **Lecture 5:** Growth Factors, Receptors and Cancer
- **Lecture 6:** Cytoplasmic Signaling Circuitry
- **Lecture 7:** Game Theory: Signaling Games

Possible Sets of Lectures (Contd.)

- **Lecture 8:** Repeated Games and Evolutionary Dynamics
- **Lecture 9:** Probability and Information Theory
- **Lecture 10:** Rate-Distortion Theory and Game Utility
- **Lecture 11:** Cancer Models: Genomics
- **Lecture 12:** Cancer Models: Signaling
- **Lecture 13:** Cancer Models: Genetics
- **Lecture 14:** Cancer Models: Therapeutics
- **Lecture 15:** The Future Challenges

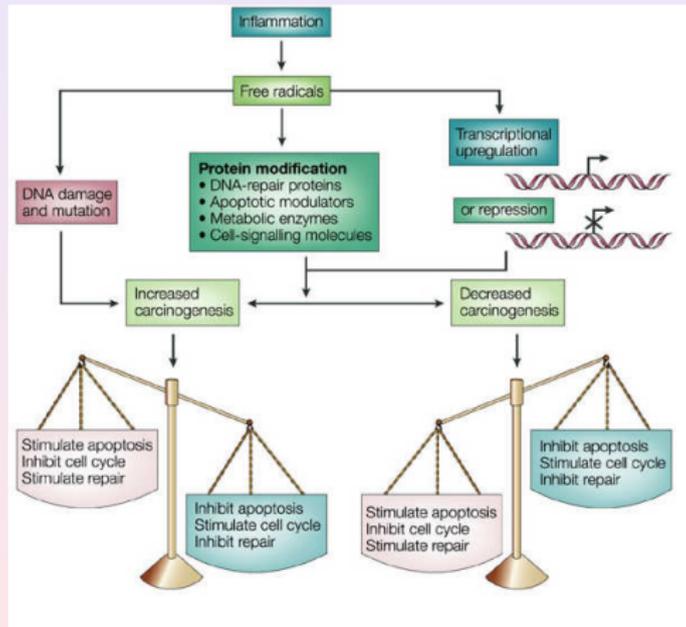
Cancer

- **What “causes” cancer?**

- 1 Cancer as a disease of the genome;
- 2 Cancer as a somatic evolutionary process;
- 3 Cancer as a price of symbiosis (mitochondrial);
- 4 Cancer as a response to multi-cellularity;
- 5 Cancer as a price of repair/regeneration (stem cells);
- 6 Cancer as a consequence of energy consumption (Anaerobic Glycolysis); *Warburg Effect*.
- 7 Cancer as a response to external stress; and
- 8 Cancer as a response to the micro-environment (hyper- and hypo-methylation).

Processes involved in Cancer

- Coordination of processes in cancer progression
 - 1 Inflammation
 - 2 Autophagy and mitophagy
 - 3 Apoptosis
 - 4 Hypoxia
 - 5 Anaerobic glycolysis
 - 6 Fibrosis
 - 7 Signaling



Information Bottleneck

- The computational principles to create a Kripke model (with states and state transitions)
- It uses Information Bottleneck Theory (a special case of RDT)

Data

- D_j = The random variable; Samples are the rows in submatrix of D_j

States

- **States** Cluster D_i ; effectively identifying the state variable X_i such that the mutual information between D_i and X_i is minimized

$$\min I(D_i; X_i) \quad \text{subj DISTORTION}$$

Transitions

- **State-Transitions** Preserve “ontologies”... Relate the conditional distribution $P(O|X_i)$ with

$$\begin{aligned} & P(O|X_{i+1}) && (1 \leq i < k) \\ \& \quad & P(O|X_{i-1}) && (1 < i \leq k) \end{aligned}$$

Measure distortions using Kullback-Liebler Distances.

Algorithm

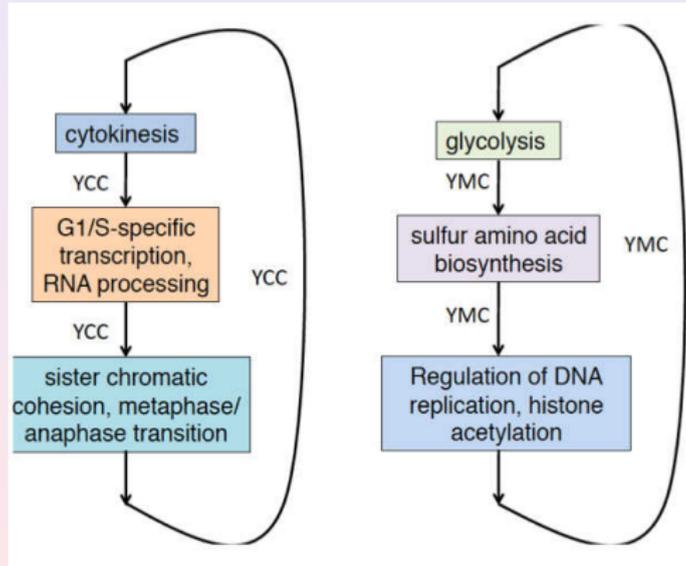
- Cluster the data over time as well as gene sets...
- Cluster and cluster-edges optimize the mutual information terms

$$I(D_i; X_i)$$

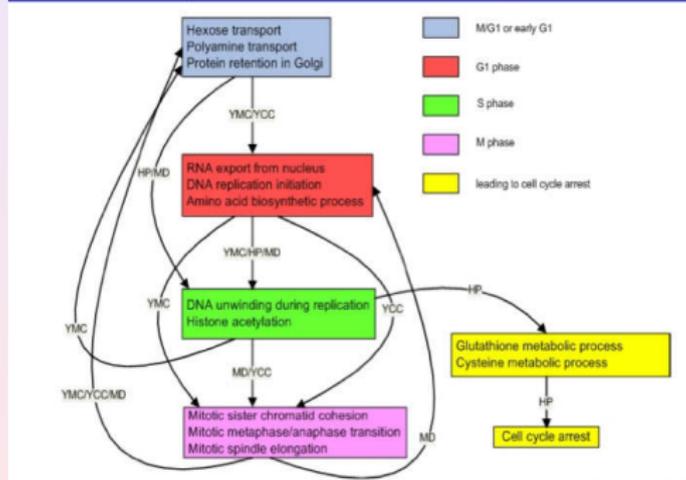
$$- \beta_+ D_{KL}(O|X_i || O|X_{i+1})$$

$$- \beta_- D_{KL}(O|X_i || O|X_{i-1})$$

Measure distortions using Kullback-Liebler Distances.



Combined Kripke Model related to cell cycle



CAN THESE ALGORITHMS BE USED TO UNDERSTAND ALL THE PROCESSES INVOLVED IN CANCER????

Questions???

**Heated Discussions on the Suggested Topics...
Resulting in a New and Better Syllabus...
That EVERYONE Loves!**

Projects

- Agent-Based Models of Cancer
- Mathematical Models (ODE and PDE) of Cancer
- Copy-Number Variation Analysis of Cancer Genomics
- Systems Biology of Cancer: Temporal, transcriptomic data
- PPI Networks and Cancer
- Progression in Pancreatic Cancer, Therapeutics (Abraxane + Gem)
- Cancer Genetics Model
- Biological Lab Projects (Single Cell analysis of tumors, Tumor heterogeneity, Tumorigenesis, Dedifferentiation)

[End of Lecture #1]