# Dynamical Modeling Methods for Systems Biology

# **Eric Sobie**

Icahn School of Medicine at Mount Sinai





# **Outline**

Overall course goals

Topics that will be covered

Structure, grading, assessment

## **Goals**

Teach contemporary methods used in systems biology for dynamical modeling

Teach methods for mathematical analysis of biological systems and simulation output

Demonstrate how dynamical mathematical models can provide insight that cannot be gained from experiments only

# Different categories of mathematical models

Statistical, top down, models versus dynamical, bottom up, models

## **Top down**

- 1) Begin with data set (often very large scale)
- 2) Use statistical methods to find patterns in the data.
- 3) Generate predictions based on the structure within the data.

Network analysis
Gene set enrichment
Clustering
Principal components
Partial least-squares regression

## **Bottom up**

- 1) Begin with hypothesis of biological mechanism.
- 2) Write down equations describing how components interact.
- 3) Run simulations to generate predictions.

Ordinary differential equations
Dynamical systems
Parameter estimation
Partial differential equations
Stochastic models

See Coursera course taught by Dr. Avi Ma'ayan

our focus will be on these

# Statistical versus dynamical models

This is discussed in more detail in several review articles

#### TEACHING RESOURCE

#### COMPUTATIONAL BIOLOGY

## Systems Biology—Biomedical Modeling

Eric A. Sobie,\* Young-Seon Lee, Sherry L. Jenkins, Ravi lyengar

Because of the complexity inherent in biological systems, many researchers frequently rely on a combination of global analysis and computational approaches to gain insight into both (i) how interacting components can produce complex system behaviors, and (ii) how changes in conditions may alter these behaviors. Because the biological details of a particular system are generally not taught along with the quantitative approaches that enable hypothesis generation and analysis of the system, we developed a course at Mount Sinai School of Medicine that introduces first-year graduate students to these computational principles and approaches. We anticipate that such approaches will apply throughout the biomedical sciences and that courses such as the one described here will become a core requirement of many graduate programs in the biological and biomedical sciences.

The Need for a Systems **Biology Course** 

Computational modeling in systems biology

Sobie et al. (2011) Science Signaling 4:tr2.

been part course dev

Medicine ate studer

applicable relevant b

plicit assu

become in

systems b

sciences n

Computa

Systems

Systems b

tational te

varying siz

# Structure of a dynamical modeling study

#### Mechanism

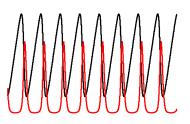
## **Equations**

$$\frac{d[G]}{dt} = V_{in} - k_1[G][ATP]$$

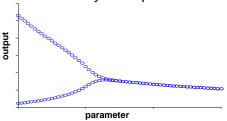
$$\frac{d[ATP]}{dt} = 2k_1[G][ATP] - \frac{k_p[ATP]}{[ATP] + K_m}$$

#### Program to simulate equations

#### Simulations



### Analyze output



# **Course Logistics**

## **Format**

7 week total course, approximately 25 lectures, 20 minutes each

3-4 lectures posted per week

Self-assessment questions during each lecture

5 homework assignments after lecture blocks

# Skills that will be taught

Using MATLAB for data analysis and visualization

Developing models consisting of systems of ordinary differential equations (ODEs)

Implementing ODE models in MATLAB and running simulations with these models

Analyzing ODE models using the tools of dynamical systems analysis

# **Biological applications**

We will see, models are useful for understanding:

Glucose oscillations in yeast

Kinase signaling pathways in mammalian cells

Regulation of the cell cycle

**Electrical signaling in neurons** 

**Goal:** provide you with the tools necessary to apply these types of models to your own questions of interest

# **Assessment and Self-Assessment**

## **Self-assessment**

1-2 questions posted at the end of each lecture explanations provided

#### **Assessment**

homework assignments to perform simulations with dynamical models

General format: start with MATLAB code written for one purpose, modify it to do something else

Assessment questions designed to verify that code works properly and that biological interpretation is correct

# **Homework assignments**

Designed to reinforce concepts discussed in lectures Assignments should require you to demonstrate:

**Technical competence (programming)** 

**Quantitative skills** 

**Biological insight into the problem**