Dynamical Systems Biology

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Executive Summary

- Systems Biology can be interpreted differently depending on where you come from, where you are going to and who judges your research
- Dynamical systems are important for Biology
- Those dynamical systems are **not** necessarily those that you learned about in school (in case you did)
- Some inspiration for dynamic biological models should come from Informatics and Engineering, not only from Physics and Chemistry
- In particular, methodologies for exploring the behavior of under-determined (open) dynamical systems, inspired by formal verification (my own research)

Organization

Some Provocative Views on Systems Biology

- Dynamical Systems and Biology
- The Dynamical Systems of Informatics
- Verification for Dummies
- Exploring the Dynamics of Continuous Systems

Conclusions

Towards Systems Biology

Towards a Paper without the word Towards in the Title

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Abstract. With the advent of post-genomic buzzword-driven big science a need was felt to moderate the level of hype and optimistic false promises in scientific papers and research proposals. In this paper we do not provide a comprehensive and complete solution to the problem mentioned in the title but nevertheless make a promising step towards the fulfillment of the goal.

- The word towards indicates that we are not yet there
- But where is there ?
- Different people will interpret the term systems biology (especially when loaded with money) in their favor
- Arguments over the meaning of words are often the most fierce (and the most stupid in some sense)

Systems Biology: a Cynical View

- Systems Biology: the current gold rush for many mathematical and technical disciplines looking for nutrition (funding, self-esteem) in the scientific food chain
- Biophysics, Biomatics, Bioinformatics, Biostatistics...
- The story goes like this:
- I do X
- I do it for my pleasure, because I studied it, and anyway, this is the only thing I will do in my current incarnation...
- ...fortunately X is very useful for Biology
- ► When you have a hammer, everything looks like a nail
- Personally this is how I came to the domain
 (X = automata, verification and hybrid systems)
- Fortunately, my hammer is universal

Systems Biology: an Arrogant View

- Biologists are essentially very concrete beings, spending most of their time in the kitchen doing manual work
- They were not selected (initially) based on ability to manipulate imaginary concepts or creativity and rigor in the abstract world of ideas but rather..
- ...based on their rigor and efficiency at the bench
- Now when they need to make a real science out of their details they need noble white collar brahmins, namely..
- ... physicists, mathematicians, computer scientists, as spiritual guides
- Like monotheists converting the pagans, these merchants of abstract methodologies try to impress the poor savage with their logics and miracles

Systems Biology: a Humble View

- Biologists are working with the most fascinating, complex and mysterious real-life phenomena
- Living systems are more complex than the hydrogen atom or the electromagnetic field (and are not effectively reducible to them)
- Living systems are more sophisticated than your dumb terminal or smart phone or mobile robot or car
- Living systems are more mysterious and primordial than the prime numbers, the algebra of Boole or the free monoid
- If some of our dry tricks can help them, even a bit, in their grand march toward..
- ..understanding something about Life Itself or helping doctors kill less patients
- We should be very happy and proud for doing, for once, something meaningful

Systems Biology: a (relatively) Sober View

- The dynamics of a scientific discipline may have different periods with various trends and fashions
- This dynamics is not always optimized towards truth
- Many aspects (politics, social dynamics, commercial interests, cognitive inertia, media distortion) play an important role
- Probably most of what is published today in top journals will go to the garbage can of history
- Few centuries ago, the science of this guy (chemistry, medicine, metaphysics) was debated extensively in prime time



Systems Biology: a Sober (but subjective) View

- Today there is an over emphasis on doing something with data provided by new experimental machinery (omics)
- The main question about "knowing" all these details is whether this knowledge:
 - Is sufficient for understanding and learning something about underlying mechanisms ? (certainly not)
 - Is necessary for that ? (very hopefully not)
 - Is helpful or counter-productive ?
- Systems Biology is about seeking some clearer (conceptual and mathematical) models of dynamical systems at various levels of abstraction
- These models, if thoughtfully constructed, and carefully and systematically analyzed/simulated may help reducing the gap between cellular biochemistry and physiology

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Dynamical Systems are Important

- Not news for biologists with a mathematical background
- ► J.J. Tyson, Bringing cartoons to life, Nature 445, 823, 2007:
- "Open any issue of *Nature* and you will find a diagram illustrating the molecular interactions purported to underlie some behavior of a living cell.
- The accompanying text explains how the link between molecules and behavior is thought to be made.
- For the simplest connections, such stories may be convincing, but as the mechanisms become more complex, intuitive explanations become more error prone and harder to believe."

In other Words

What is the relation (if any) between



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Systems and Behaviors



- Left: a model of a dynamical system which "explains" the mechanism in question
- Right: some experimentally observed behavior supposed to have some relation to the behaviors generated by model
- What is this relation exactly?
- Current practice leaves a lot to be desired, at least from a theoreticians' point of view

An Illustrative Joke

- An engineer, a physicist and a mathematician are traveling in a train in Scottland. Suddenly they see a black sheep
- Hmmm, says the engineer, I didn't know that sheeps in Scottland are black
- No my friend, corrects him the physicist, some sheeps in Scottland are black
- To be more precise, says the mathematician, there is a sheep in Scottland having at least one black side
- A discipline is roughly characterized by the number of logical quantifiers ∃ ∀ (and their alternations) its members feel comfortable with
- By the way what would a **biologist** say?
- In the Scottish sheep the agouti isoform is first expressed at E10.5 in neural crest-derived ventral cells of the second branchial arch

Dynamical Systems, a Good Idea

- The quote from Tyson goes on like this:
- "A better way to build bridges from molecular biology to cell physiology is to recognize that a network of interacting genes and proteins is ..
- ... a dynamic system evolving in space and time according to fundamental laws of reaction, diffusion and transport
- These laws govern how a regulatory network, confronted by any set of stimuli, determines the appropriate response of a cell
- This information processing system can be described in precise mathematical terms,
- .. and the resulting equations can be analyzed and simulated to provide reliable, testable accounts of the molecular control of cell behavior"
- No news for engineers..

Models in Engineering

- To build complex systems other than by trial and error you need models
- Regardless of the language or tool used to build a model, at the end there is some kind of dynamical system
- A mathematical entity that generates behaviors which are progressions of states and events in time
- Sometimes you can reason about such systems analytically
- But typically you simulate the model on the computer and generate behaviors
- If the model is related to reality you will learn something from the simulation about the actual behavior of the system
- Major difference: in engineering, the components are often well-understood and we need the simulation only because the outcome of their interaction is hard to predict

My Point: Systems Biology \approx Dynamical Systems, but..

- To make progress in Systems Biology we should upgrade descriptive "models" by dynamic models with stronger predictive power and refutability
- Classical models of dynamical systems and classical analysis techniques tailored for them are **not** sufficient for **effective modeling** and **analysis** of biological phenomena
- Models, insights and computer-based analysis tools developed within Informatics (Computer Science) can help
- The whole systems thinking in CS is more evolved and sophisticated in some aspects than in Physics and Mathematics
- This is true of other information-oriented engineering disciplines such as the design of circuits or control systems

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Conclusions

What "Is" Informatics ?

- Informatics is the study of discrete-event dynamical systems (automata, transition systems)
- A natural point of view for those working on modeling and verification of "reactive systems"
- Less natural for data-intensive software developers and users
- This fact is sometimes **obscured** by fancy formalisms:
- Petri nets, process algebras, rewriting systems, temporal logics, Turing machines, programs
- All honorable topics with intrinsic beauty, sometimes even applications and deep insights
- But in an inter-disciplinary context they should be distilled to their essence to make sense to potential users.. rather than intimidate them
- In fact, the need to impress one's own community is a serious impediment in inter-disciplinary research

Dynamical Systems in General

- The following abstract features of dynamical systems are common to both continuous and discrete systems:
- State variables whose set of valuations determine the state space
- A time domain along which these values evolve
- A dynamic law: how state variables evolve over time, possibly under the influence of external factors
- System behaviors are progressions of states in time produced according to the dynamic law
- Knowing an initial state x[0] the model can predict, to some extent, the value of x[t]

Types of Dynamical Systems

- Dynamic system models differ from each other according to their concrete details:
- State variables: numbers or more abstract domains that do not have a quantitative meaning
- ► Time domain: metric (dense or discrete) or logical
- The form of the dynamical law, constrained, of course, by the state variables and time domain
- The type of available analysis (analytic, simulation)
- Other features (open/closed, type of non-determinism, spatial extension)

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Classical Dynamical Systems

- State variables: real numbers (location, velocity, energy, voltage, concentration)
- Time domain: the real time axis $\mathbb R$ or a discretization of it
- Dynamic law: differential equations

$$\dot{x} = f(x, u)$$

or their discrete-time approximations

$$x[t+1] = f(x[t], u[t])$$

- Behaviors: trajectories in the continuous state space
- Typically presented in the form of a collection of waveforms or time-series, mappings from time to the state-space
- What you would construct using tools like Matlab Simulink, Modelica, SPICE simulators, etc.

Discrete-Event Dynamical Systems (Automata)

- An abstract discrete state space
- State variables need not have a numerical meaning
- A logical time domain defined by the events (order but not metric)
- Dynamics defined by transition rules: input event a takes the system from state s to state s'
- Behaviors are sequences of states and/or events
- Composition of large systems from small ones using different modes of interaction: synchronous/asynchronous, state-based/event-based
- What you will build using tools like Raphsody or Stateflow (or even C programs or digital hardware simulators)

Preview: Timed and Hybrid Systems

- Mixing discrete and continuous dynamics
- Hybrid automata: automata with a different continuous dynamics in each state
- Transitions = mode switchings (valves, thermostats, gears, genes, walking)
- Timed systems: an intermediate level of abstraction
- Timed Behaviors = discrete events embedded in metric time, Boolean signals, Gantt charts
- Used implicitly by everybody doing real-time, scheduling, embedded, planning in professional and real life
- Formally: timed automata (automata with clock variables)

Automata: Modeling and Analysis

- Automata model processes viewed as sequences of steps: software, hardware, ATMs, user interfaces administrative procedures, cooking recipes, smart phones...
- Unlike continuous systems there are no simple analytical tools to predict their long-term behavior
- We can **simulate** and sometimes do **formal verification**:
- Check whether all behaviors of a system, exposed to some uncontrolled inputs, exhibit some qualitative behavior:
- Never reach some part of the state space; Always follow some sequential pattern of behavior...
- These temporal properties include transients and are much richer than classical steady states or limit cycles
- There are tools for the verification of huge systems by sophisticated graph algorithms and powerful SAT solvers

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Conclusions

Illustration: The Coffee Machine

- Consider a machine that takes money and distributes drinks
- The system is built from two subsystems: one takes care of payment and one handles choice and preparation of drinks
- They communicate by sending messages



Remark: Signalling

- Modern systems separate information-processing from the physical interface
- An inserted coin, a pushed button or a full cup are physical events translated by sensors into uniform low-energy signals
- These signals are treated as information, without thinking too much about their material realization
- Unless you are an engineer specialized in such mechanisms



Automaton Models

- The two systems are modeled as automata
- transitions are triggered by external events and by events coming from the **other** subsystem



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The Global Model

- The behavior of the whole system is captured by a composition (product) M₁ || M₂ of the components
- States are elements of the Cartesian product of the respective sets of states, indicating the state of each component
- Some transitions are independent and some are synchronized, taken by the two components simultaneously
- Behaviors of the systems are paths in this transition graph



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Normal Behaviors



 Customer inserts a coin, then sees the bus arriving, cancels and gets the coin back

0A coin-in 1B cancel coin-out 0A

 Customer inserts a coin, requests coffee, gets it and the systems returns to initial state

0A coin-in 1B req-coffee st-coffee 1C drink-ready 0A

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An Abnormal Behavior



Suppose the customer presses the cancel button after the coffee starts being prepared..

0A coin-in 1B req-coffee st-coffee 1C cancel coin-out 0C drink-ready 0A

Not so attractive for the owner of the machine

Fixing the Bug

- ▶ When *M*₂ starts preparing coffee it emits a **lock** signal
- ► When M₁ received this message it enters a new state where cancel is ignored



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The Moral of the Story I

- Many complex systems can be modeled as a composition of interacting automata
- Behaviors of the system correspond to paths in the global transition graph of the system
- The size of this graph is exponential in the number of components (state explosion, curse of dimensionality)
- So if you have an interaction diagram which covers the wall, its state-space can cover the universe
- These paths are labeled by input events representing influences of the external environment
- Each input sequence may induce a different behavior, a different scenario

The Moral of the Story II

- We want to ensure that the system responds correctly to all conceivable inputs
- That it is robust and behaves properly in many contexts, not only where users never push the cancel button inappropriately
- We can choose an individual input sequence and simulate the behavior it induces, but we cannot do it exhaustively
- Verification is a collection of automatic and semi-automatic methods to analyze all the paths in the graph

This type of analysis we export to the assessment of biological models and hypotheses

Organization

- Some Provocative Views on Systems Biology
- Dynamical Systems and Biology
- The Dynamical Systems of Informatics
- Verification for Dummies
- Exploring the Dynamics of Continuous Systems

Conclusions
Under-Determined Continuous Dynamical Systems

We study open dynamical systems of the form

$$x[t+1] = f(x[t], p, u[t])$$

- Such systems are incomplete, under-determined in the following sense:
- The initial state x[0] is not precisely known, only that it is in some set X₀
- The system has a vector of parameters p whose value is not precisely known, only that it is in some parameter-space P
- The exact form of the dynamic disturbance u[t] is not known, only that it is constrained to be in some U for every t
- ► In order to produce a simulation trace x[0], x[1], x[2], · · · you need to fix values for those

Static/Punctual Under-Determination

- Let us ignore dynamic inputs and focus on the first two types of under-determination that we call **punctual**
- In both cases, in order to simulate your model and produce a trace x[0], x[1], x[2], · · · you need to fix one point/vector:
- x[0] in the state space
- p in the parameter space
- Technically their treatment is similar and I will use parameters as motivation and initial states for graphical illustration

Models, Reality and Parameters

- Whenever models are supposed to represent something non-trivial they are just approximations
- This is evident for anybody working in modeling concrete physical systems
- It is less evident for those working on the functionality of digital hardware or software
- In these domains you have powerful deterministic abstractions (logical gates, program instructions) that work
- A common way to pack our ignorance in a compact way is to introduce parameters ranging in some parameter space

Examples:

- Voltage level modeling and simulation of circuits:
- A lot of variability in transistor characteristics depending on production batch, place in the chip, temperature, etc.
- Timing performance analysis of a new application (task graph) on a new multi-core architecture:
- Precise execution times of tasks are not known before the application is written and the architecture is built
- Biochemical reactions in cells following the mass action law:
- Many parameters related to the affinity between molecules cannot be deduced from first principles
- They are measured via isolated experiments under different conditions and only wide bounds on their values can be known

So What is the Problem?

- So you have a model which is under-determined, or equivalently an infinite number of models
- For simulation you need to determine, to make a choice to pick a point p in the parameter space
- The simulation shows you something about one possible behavior of the system, or a behavior of one possible model
- But another choice of parameter values could have produced a completely different behavior

Ho do you live with that?

Possible Attitudes

- The answer depends on many factors
- One is the **responsibility** of the modeler/simulator
- What are the consequences of not taking under-determination seriously
- Is there a **penalty** for jumping into conclusions based on one or few simulations?
- Another factor is the mathematical and real natures of the system you are dealing with

And as usual, it may depend on culture, background and tradition in the industrial or academic community

Non Responsibility: a Caricature

- Suppose you are a scientist not engineer, say biologist
- You conduct experiments and observe traces
- You propose a model and tune the parameters until you obtain a trace similar to the one observed experimentally
- These are nominal values of the parameters
- Then you can **publish** a paper about your model
- Except for picky reviewers there are no real consequences for neglecting under-determination
- The situation is different if some engineering is involved (pharmacokinetics, synthetic biology)
- Or if you want others to compose their models with yours

Justified Nominal Value

- You can get away with using a nominal value if your system is very smooth and well-behaving
- Points in the neighborhood of p generate similar traces
- There are also mathematical techniques (bifurcation diagrams, etc.) that can tell you sometimes what happens when you vary parameters
- This smoothness is easily broken by mode switching
- Another justification for ignoring parameter variability:
- When the system is anyway adaptive to deviations from nominal behavior (control, feedback)

Taking Under-Determination More Seriously: Sampling

- One can sample the parameter space with or without probabilistic assumptions
- Make a grid in the parameter space (exponential in the number of parameters)
- Or pick parameter values at random according to some distribution
- In the sequel I illustrate a technique (due to A. Donze) for adaptive search in the parameter space
- Local sensitivity information from the numerical simulator tells you where to refine the coverage
- Arbitrary dimensionality of the state space, but no miracles against the dimensionality of the parameter space

Sensitivity-based Exploration I

- We want to prove all trajectories from X₀ do not reach a bad set of states
- Take $x_0 \in X_0$ and build a ball B_0 around it that covers X_0



- Simulate from x_0 and generate a sequence of balls B_0, B_1, \ldots
- B_i contains all points reachable from B_0 in *i* steps

Sensitivity-based Exploration II

After k steps, three things may happen:



- 1. No ball intersects bad set and the system is safe (due to over-approximation)
- 2. The concrete trajectory intersects the bad set and the system is unsafe
- ► 3. Ball B_k intersects the bad set but we do not know if it is a real or spurious behavior

Sensitivity-based Exploration III

In the latter case we refine the coverage and repeat the process for two smaller balls



- Can prove correctness using a finite number of simulations, focusing on the interesting values
- Can approximate the **boundary** between parameter values that yield some qualitative behaviors and values that do not

The Breach Toolboox

- Parameter-space exploration for arbitrary continuous dynamical systems relative to quantitative temporal properties expressed in STL (signal temporal logic)
- Applied to embedded control systems, analog circuits, biochemical reactions (haematopoiesis, angiogenesis, apoptosis) and anasthesia.



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Dynamic Under-Determination

- The system is modeled as open, exposed to external disturbances
- Dynamics of the form

$$x[i+1] = f(x[i], v[i])$$

- The natural way to represent the influence of other unmodeled subsystems and external environment
- Under-determination is dynamic: to produce a trace you need to give the value of v at every time step, a signal/sequence v[1],...,v[k]
- ► A priory a much larger space to sample from: dimension *mk* compared to *m* for static
- One can use a nominal value: constant, step, sinusoid, random noise, etc.

Taking Under-Determination Seriously: Guided Sampling

- A method due to T. Dang:
- Use ideas from robotic motion planning (RRT) to generate inputs that yield a good **coverage** of the reachable state space
- Applied to analog circuits



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Taking Under-Determination More Seriously: Verification

- Paranoid worst-case formal verification attitude:
- ► If we say something about the system it should be provably true for all choices of p, x[0] and v[1],...,v[k]
- Instead of doing a simple simulation you do set-based simulation, computing tubes of trajectories covering everything
- Breadth-first rather than depth-first exploration



- Advantages: works also for hybrid (switched) systems
- Limitations: manipulates geometric objects in high dimension

State of the Art

- ► Linear and piecewise-linear dynamics ~ 200 variables using algorithms of **C. Le Guernic and A. Girard**
- ► Nonlinear dynamics with 10 20 variables an ongoing research activity
- Implemented into the SpaceEx tool developed under the direction of G. Frehse
- Available on http://spaceex.imag.fr with model editor, visualization and more

Waiting for more beta testers

The State-Space Explorer (SpaceEx)







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Example: Lac Operon (T. Dang)

$$\begin{aligned} \dot{R}_{a} &= \tau - \mu * R_{a} - k_{2}R_{a}O_{f} + k_{-2}(\chi - O_{f}) - k_{3}R_{a}I_{i}^{2} + k_{8}R_{i}G^{2} \\ \dot{O}_{f} &= -k_{2}r_{a}O_{f} + k_{-2}(\chi - O_{f}) \\ \dot{E} &= \nu k_{4}O_{f} - k_{7}E \\ \dot{M} &= \nu k_{4}O_{f} - k_{6}M \\ \dot{I}_{i} &= -2k_{3}R_{a}I_{i}^{2} + 2k_{-3}F_{1} + k_{5}I_{r}M - k_{-5}I_{i}M - k_{9}I_{i}E \\ \dot{G} &= -2k_{8}R_{i}G^{2} + 2k_{-8}R_{a} + k_{9}I_{i}E \end{aligned}$$





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Conclusions

Back to the Big Picture

- Biology needs (among other things) more dynamic models to form verifiable predictions
- These models can benefit from the accumulated understanding of dynamical system within informatics and cannot rely only on 19th century mathematics
- The views of dynamical system developed within informatics are, sometimes, more adapted to the complexity and heterogeneity of Biological phenomena
- Biological modeling should be founded on various types of dynamical models: continuous, discrete, hybrid and timed
- These models should be strongly supported by computerized analysis tools offering a range of capabilities from simulation to verification and synthesis

Back to the Big Picture

- Systems Biology should combine insights from:
- Engineering disciplines: modeling and analysis of very complex man-made systems (chips, control systems, software, networks, cars, airplanes, chemical plants)
- Physics, Chemistry: experience in mathematical modeling of natural systems with measurement constraints

 Mathematics and Informatics as a unifying theoretical framework

Thank You