Graphical Models for Stochastic Verification and Synthesis

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Research Goals

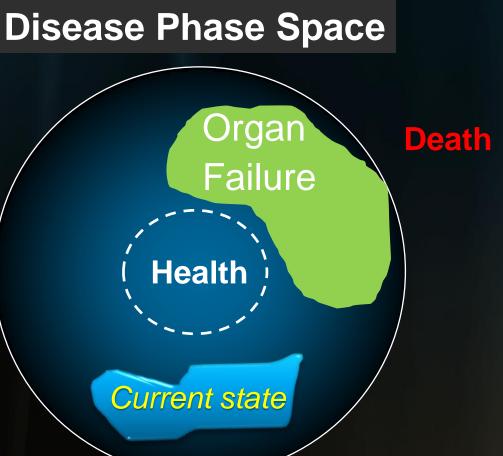
 Develop new methods for reasoning about stochastic processes by adapting and combining methods from Machine Learning and Formal Verification

- Machine Learning addresses uncertainty
- Verification addresses complexity and scalability
- Application Domains:
 - Computational Biology
 - Embedded Systems

Context & Motivation Personalized Medicine Developing patient-specific treatment plans



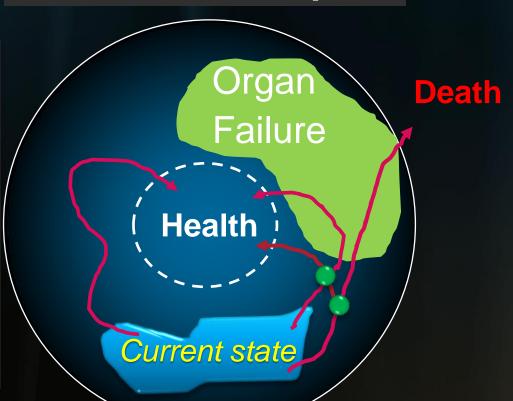
Intensive Care Unit



Context & Motivation Personalized Medicine Developing patient-specific treatment plans

Disease Phase Space

Primary Tasks:
(1) Determine where the patient is now (approximately)
(2) Characterize the patient's trajectory (approximately)
(3) Select interventions based on (1) and (2)

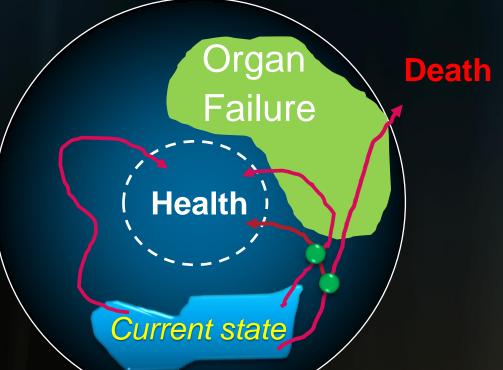


Context & Motivation

Physicians routinely use simple (nondynamic) models in these tasks

We believe that dynamic models will be more useful

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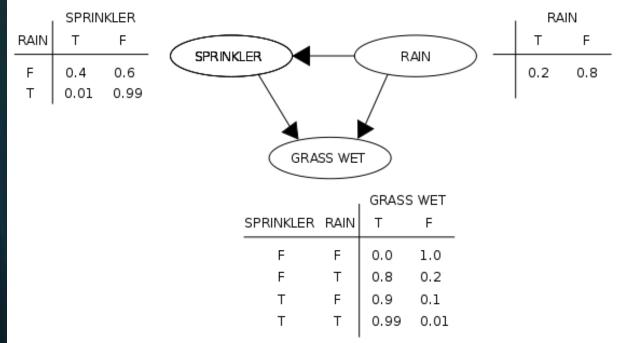
Context and Motivation

Dynamic models of disease processes
 (Stochastic) ODEs/PDEs
 Graphical models

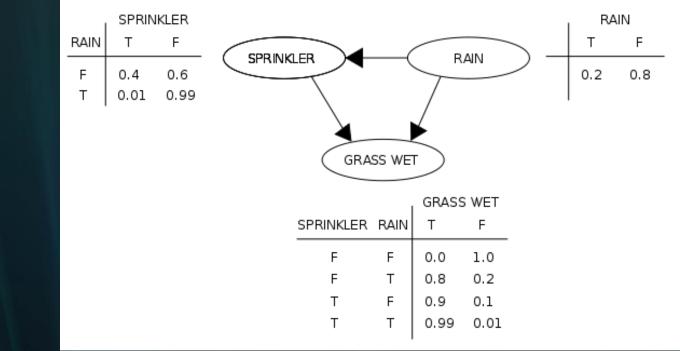
Graphical Models

- Let X = {x₁,..., x_n} be a set of random variables
 Each X_i can be continuous or discrete
- A probabilistic graphical model (PGM) is a factored encoding of P(X)
 - $\mathcal{M} = (\mathsf{G}, \Psi, \Theta)$
 - G = (V,E) is a graph over the random variables
 - V_i corresponds to x_i
 - Edges reveal conditional independencies
 - Ψ is a set of functions over V and E
 - Θ is a vector of parameters

Example



 This models P(Sprinkler, Rain, Grass Wet)
 V = {Sprinkler, Rain, Grass Wet}
 Ψ = { P(rain), P(Sprinkler | Rain), P(Grass Wet | Sprinkler, Rain) }
 Θ = The elements in the 3 tables



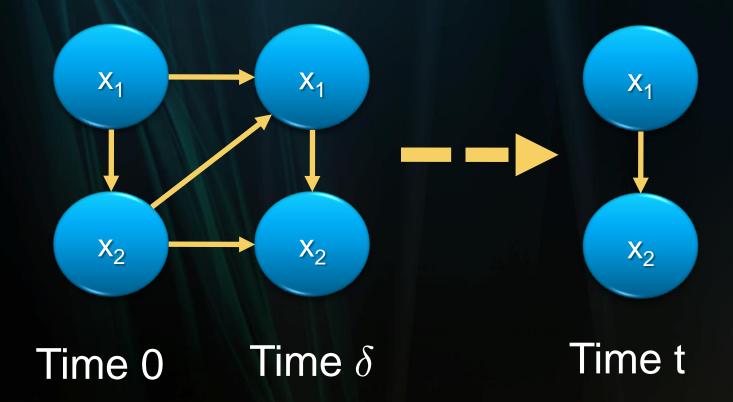
What is the probability that it is raining, given that the grass is wet?

Example

$$\mathbf{P}(R = T \mid G = T) = \frac{\mathbf{P}(G = T, R = T)}{\mathbf{P}(G = T)} = \frac{\sum_{S \in \{T, F\}} \mathbf{P}(G = T, S, R = T)}{\sum_{S, R \in \{T, F\}} \mathbf{P}(G = T, S, R)}$$

 $\frac{(0.99 \times 0.01 \times 0.2 = 0.00198_{TTT}) + (0.8 \times 0.99 \times 0.2 = 0.1584_{TFT})}{0.00198_{TTT} + 0.288_{TTF} + 0.1584_{TFT} + 0_{TFF}} \approx 35.77\%.$

Models distributions over time series P(X_{0:t})
 Time can be continuous or discrete



- Traditional Tasks
 - Inference
 - Computing P($A_{0:t+\delta} | B_{0:t}; \mathcal{M}$)
 - $A \cap B = \emptyset, A \cup B = X$
 - Learning
 - Computing $\operatorname{argmax}_{\theta} \mathsf{P}(\mathcal{D}; \mathcal{M})$
 - \mathcal{D} is a set of observations over $\mathbf{Y}_{0:t} \subseteq \mathbf{X}_{0:t}$
 - Structure Learning
 - Computing $\operatorname{argmax}_{G,\theta} \mathsf{P}(\mathcal{D};\mathcal{M})$
 - I.e., simultaneously learning graph topology and parameters

We introduced the following generalizations:

- Inference over temporal logic formulas*
 - Computing P($\phi_1 | \phi_2; \mathcal{M}$)
- Learning over temporal logic formulas*
 - Computing $\operatorname{argmax}_{\theta} \mathsf{P}(\phi ; \mathcal{M})$
- Structure Learning over temporal logic formulas*
 Computing argmax_{G,θ} P(φ; M)

*Formulas are in bounded LTL

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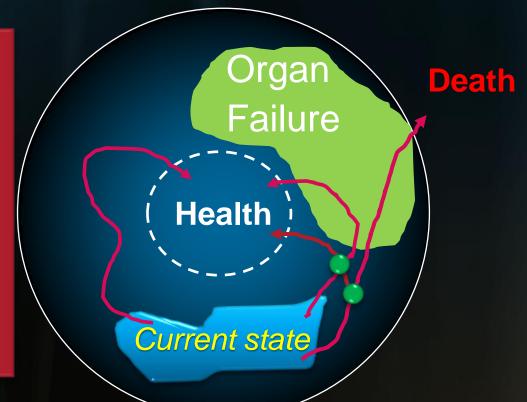
These generalizations provide a more expressive framework for using graphical models

In Context

These generalizations are also relevant to our three tasks

• Eg., $\phi := \neg$ "Organ Failure" U^t "Health"

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New Algorithms (1)

Inference over temporal logic formulas [L09]

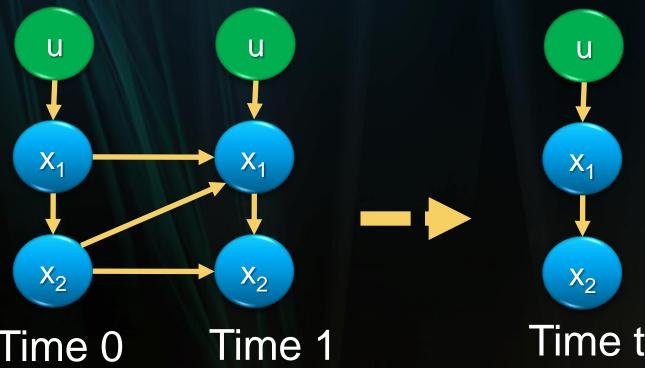
Dynamics	Distribution	Time	Inference
Linear	Gaussian	Discrete	Exact
Non-linear	Gaussian	Discrete	Accurate to 2 nd order
Non-linear	Non-Gaussian	Discrete	Approximate
Non-linear	Multinomial	Discrete	Exact

Also: new sampling algorithms for rare events

New Algorithms (2)

 Control policy synthesis and structure learning for synchronous or asynchronous Boolean networks [LJ08,LJ09]

Relies on symbolic model checking



New Algorithms (3)

Learning over temporal logic formulas [L08]

Distribution	Time	Learning
Multinomial	Discrete	Approximate*
Continuous	Discrete	Approximate

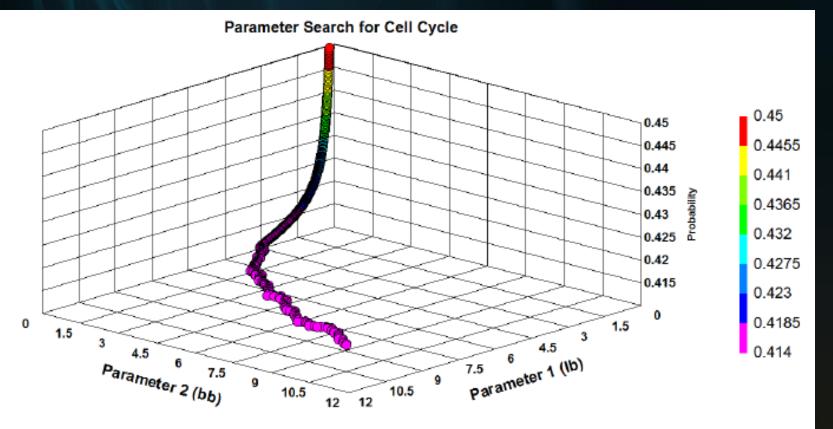
* Uses bit-vector decision procedures Parameter Synthesis [DCL09,JL10]

Distribution	Time	Learning
Continuous	Continuous	Approximate
Multinomial	Discrete	Approximate*

* Uses abstraction-refinement

Example: Parameter Estimation

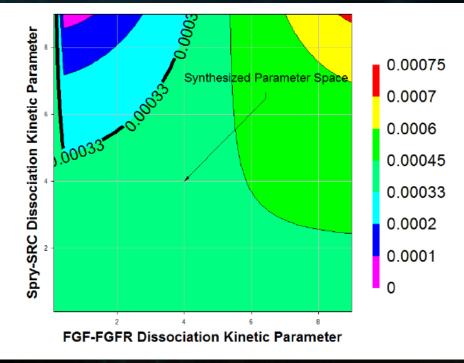
Cell-cycle model φ := F^t "Cyclin bound = 0"



Example: Synthesis

Fibroblast growth receptor pathway model φ := F^t (A>0 & B = 0)

2D synthesis



Note: our method has been used to synthesize up to 6 parameters, simultaneously

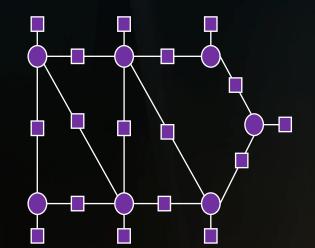
Applications and Extensions

Medical

- Sepsis
- Pancreatitis
- Chronic Myeloid Leukemia

Biological

- Embryogenesis
- Signaling Pathways
- Spatial Models
 - Markov Random Fields



Potential Collaborations

Jim Faeder

- Our algorithm for parameter synthesis for stochastic systems was developed with BioNetGen in mind
- Bud Mishra
 - Applications to GOALIE
- Atrial Fibrillation
 - Markov Random Fields
- Rance Cleaveland
 - Our rare event sampling procedure might be relevant to Reactis[©]

