

Inductive Logic Programming Applied to Systems Biology

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internship at NII, University of Grenoble and ENSIMAG, 20min speech

September 29, 2009

Introduction

- National Institute of Informatics, Tokyo
- Supervising: Katsumi Inoue, Taisuke Sato, Pierre Bessière, Augustin Lux
- Head (first 2 weeks)
- Tail (last 4 months)

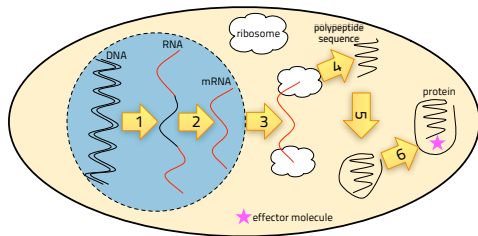
- 1 Systems Biology and Inductive Logic Programming
 - Systems Biology
 - Inductive Logic Programming
 - Previous Works
- 2 Limitations
 - Limits of The Previous Models
 - Many Levels
- 3 A New Logic Modeling
 - Precision - Generality Trade-off
 - Kinetic Modeling
- 4 Implementation
 - Discretization
 - Logic Modeling
 - Results (ranked)

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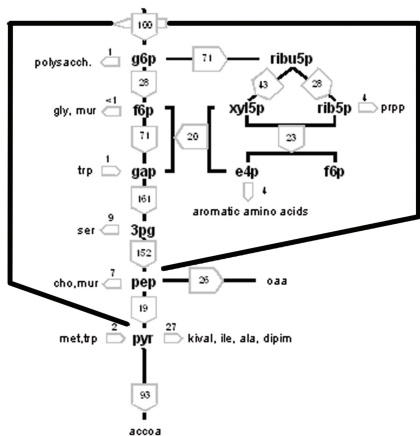
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The Cell



- 1 Transcription
- 2 Post-transcription
- 3 Migration of the mRNA
- 4 Translation
- 5 Post-transl./Folding
- 6 Binding of an effector

Metabolic Pathways



A metabolic pathway: Glucolysis and Pentose Phosphate of *Escherichia Coli*

- Graphs of interconnected reactions
- Glucose enters $ATP \Rightarrow ADP$ G6P
- Chain of reactions to take energy and store it in ATP/NADH (2 per molecule of Glucose)
- Acetyl CoA is at the origin of the Krebs cycle (part of cellular respiration)

Abduction & Induction

ILP strength lies in the fact that learnt rules/clauses are directly useable in a logical program.

Induction & Abduction

From \mathcal{B} Background Knowledge \wedge \mathcal{E} examples

◇ Find \mathcal{H} Hypotheses satisfying $\mathcal{B} \wedge \mathcal{H} \models \mathcal{E}$ and $\mathcal{B} \cup \mathcal{H} \neq \perp$

Abduction: ground (or \exists quant.) formulae, direct causes of observations that are called explanations.

Induction: universally (\forall) quantified formulae (small \mathcal{B}), more general hypotheses.



R. J. Mooney: Integrating abduction and induction in machine learning. IJCAI97 Workshop on Abduction and Induction in AI, 37–42 (1997).



Flach P. A., Kakas A. C.: Abduction and induction: Essays on their relation and integration. Kluwer (2000).

Inverse Entailment (Consequence Finding)

ILP is interested in the formulas derived from $\mathcal{B} \wedge \neg\mathcal{E}$ that are not derived from \mathcal{B} alone.

Inverse Entailment

The previous definition is equivalent to $\mathcal{B} \wedge \neg\mathcal{E} \models \neg\mathcal{H}$ and $\mathcal{B} \not\models \neg\mathcal{H}$.

We can then use a consequence finding procedure (resolution, tableaux) to find $\neg\mathcal{H}$. In Inoue's lab, it is now done with SOLAR.



Inoue, K.: Linear resolution for consequence finding. *Artificial Intelligence* 56:301-353 (1992).



Inoue K.: Induction as consequence finding. *Machine Learning*, 55:109-135 (2004).



Nabeshima H., Iwanuma K., and Inoue K.: SOLAR: A Consequence Finding System for Advanced Reasoning. *TABLEAUX 2003, LNAI, Vol. 2796*, pp. 257-263, Springer (2003).

Previous Works

Why? Inhibitionary effects of toxins on chemical reactions.

How? Metabolic flux analysis through induction with rules that explain the concentration changes (up or down) between 2 experiments, with and w/o toxin.



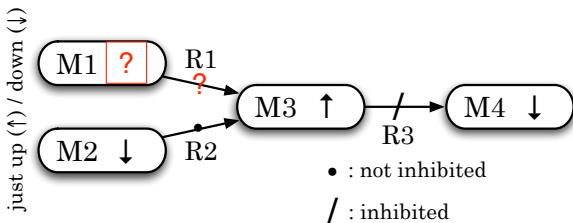
Doncescu, A., Inoue K., Yamamoto Y.: Knowledge Based Discovery in Systems Biology Using CF-Induction. LNCS N.4570, pages 395-404 (2007).



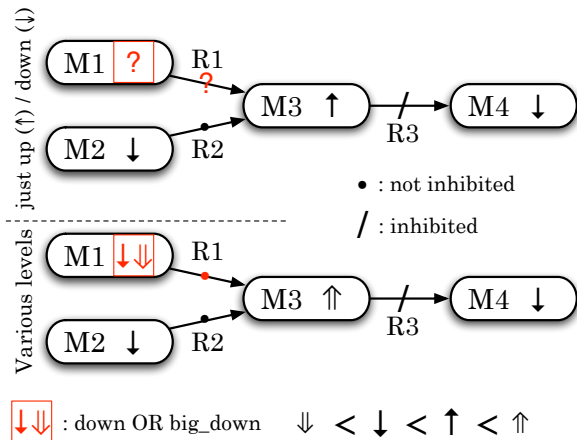
King R.D., Whelan K.E., Jones F.M., Reiser P.G.K., Bryant C.H., Muggleton S.H., Kell D.B. & Oliver S.G.: Functional genomic hypothesis generation and experimentation by a robot scientist. Nature 427, 247-252 (2004).

Limits of The Previous Models

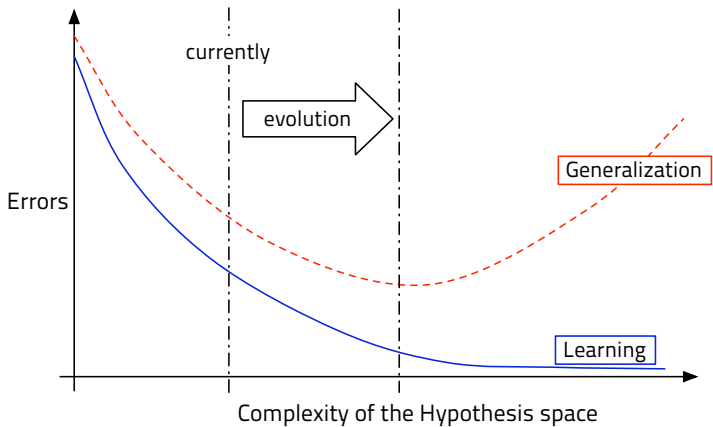
- No models for dynamic transitions
- Not enough information to be precise enough:



Dealing With More Knowledge



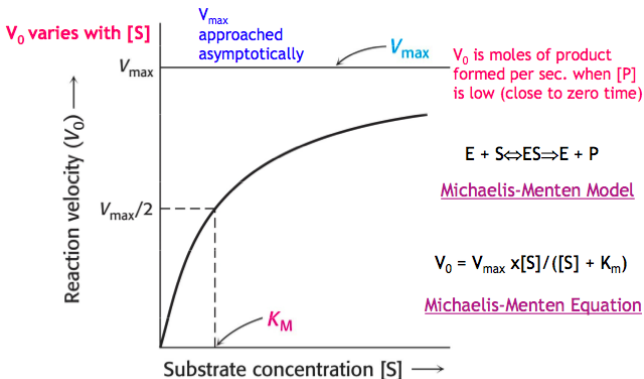
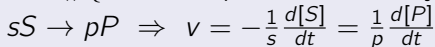
Be More Precise, Avoid Overfitting



Chemical Kinetic and Michaelis-Menten Equation

Speed of a one-way reaction

$v = \#\{\text{reactions per second catalyzed per mole of the enzyme}\}$



Simplification of Michaelis-Menten Equation



Michaelis - Menten equation :
$$\frac{d[P]}{dt} = V_m \frac{[S]}{[S] + K_m} \quad (1)$$

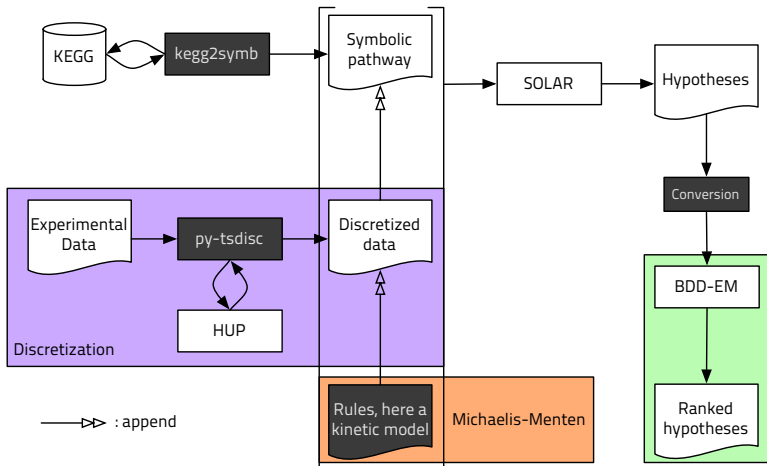
$$\frac{d[P]}{dt} \xrightarrow{\text{disc.time}} \frac{[P]_{T+\text{timestep}} - [P]_T}{(T + \text{timestep}) - T} \quad (2)$$

$$(1) \text{ and } (2) \implies V_m \frac{[S]_T}{[S]_T + K_m} \approx \frac{[P]_{T+\text{timestep}} - [P]_T}{(T + \text{timestep}) - T}$$

We chose to work with a constant timestep :

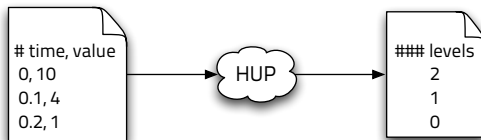
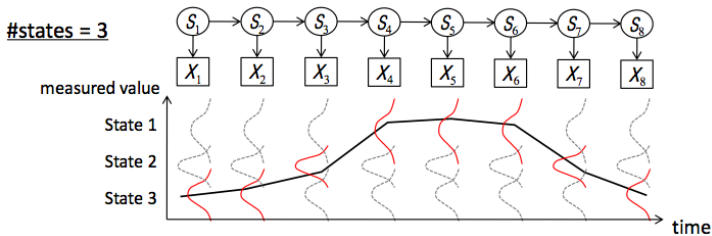
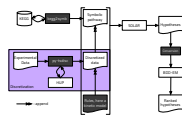
$$\implies [P]_{T+1} = V_m \frac{[S]_T}{[S]_T + K_m} + [P]_T \quad (3)$$

My Work



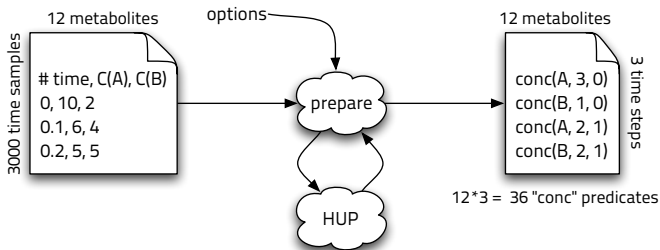
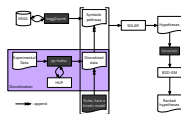
HUP: HMM Utility Program

A work from Yoshitaka Kameya.
 Clustering method that uses Continuous
 Hidden Markov Model + Bayesian Score:



“prepare.py” - Wrapping HUP

Adapts the entry file (concentration, speed, acceleration) and feed it to HUP with the right options, calculates subsets for time discretization and then write the solar-like file.

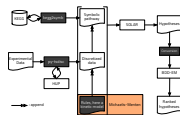


For the presented results on *E. Coli*, I remarked that the values of K_m and concentration were very sparse \Rightarrow we took the log.

Logic Kinetic Modeling

If we make the approximations for extreme values in:

$$[P]_{T+1} = V_m \frac{[S]_T}{[S]_T + K_m} + [P]_T$$



With only 3 levels, as we have in our discretization of *E.Coli* experiments, we will get the following simple rules:

$$[S] \ll K_m$$

$\text{reaction}(S, P, K_m) \wedge \text{concentration}(S, 0, 0) \wedge \text{concentration}(K_m, 2, 0) \wedge \text{concentration}(P, L, 0) \rightarrow \text{concentration}(P, L, 1)$

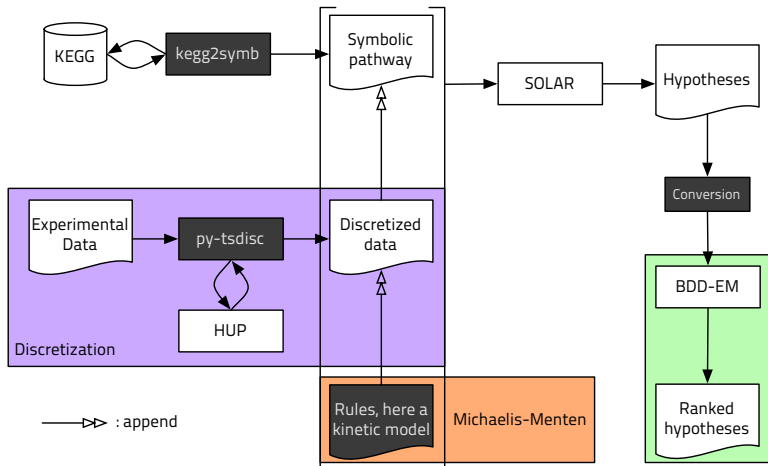
$$[S] \simeq K_m$$

$\text{reaction}(S, P, K_m) \wedge \text{concentration}(S, 1, 0) \wedge \text{concentration}(K_m, 1, 0) \wedge \text{concentration}(P, L, 0) \rightarrow \text{concentration}(P, L, 1)$

$$[S] \gg K_m$$

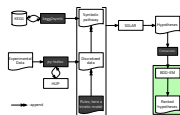
$\text{reaction}(S, P, K_m) \wedge \text{concentration}(S, 2, 0) \wedge \text{concentration}(K_m, 0, 0) \rightarrow \text{concentration}(P, 2, 1)$

The Whole Process

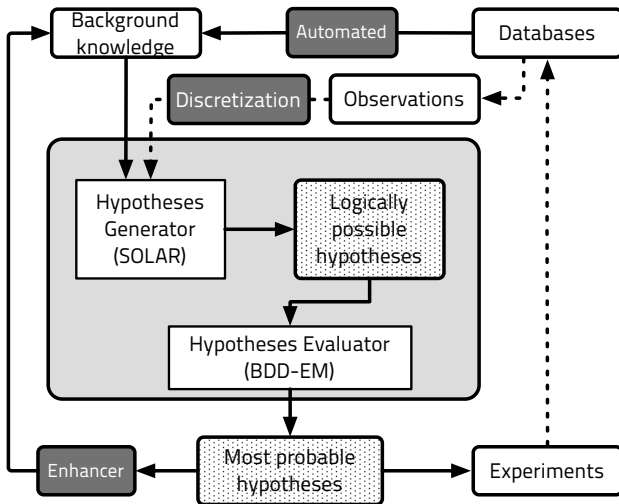


Ranking the hypotheses with BDD-EM

Hyp. no.	Probability	Abducted concentrations levels at T=0
H130	≈ 1.0	pg3: 2, adp: 0
H392	$4.879.E^{-1}$	sed7p: 0, e4p: 2, f6p: 0, pg3: 2, adp: 0
H216	$7.567.E^{-2}$	pg3: 2, adp: 0, pep: 0, atp: 2, pyr: 2
H196	$6.930.E^{-2}$	fdp: 0, dhap: 2, gap: 0, pg3: 2, adp: 0
H356	$5.621.E^{-2}$	pg3: 2, adp: 0, g6p: 1, nadph: 1
H94	$3.692.E^{-2}$	sed7p: 0, e4p: 2, f6p: 0, pg3: 2, adp: 0, pep: 0, atp: 2, pyr: 2
H251	$3.497.E^{-2}$	glucose: 2, adp: 0, pg3: 2
H286	$3.382.E^{-2}$	sed7p: 0, e4p: 2, f6p: 0, fdp: 0, dhap: 2, gap: 0, pg3: 2, adp: 0
H405	$2.796.E^{-2}$	pg3: 2, adp: 0, pep: 2, atp: 0
H167	$2.743.E^{-2}$	sed7p: 0, e4p: 2, f6p: 0, pg3: 2, adp: 0, g6p: 1, nadph: 1
.	.	.
H378	$1.974.E^{-8}$	glucose: 2, adp: 0, sed7p: 0, e4p: 2, f6p: 0, fdp: 0, dhap: 2, gap: 0, pg3: 2, pep: 0, atp: 2, pyr: 2, g6p: 0, nadph: 2, pg6: 1



Conclusion: My Ideas



Conclusion: What's left?

To be able to deal with *any* pathways for which we would have concentrations and michaelis-menten constants. 2 problems:

- 1 more than 3 levels
 - A proof of concept has been made with a compute predicate working in the Java part of SOLAR
 - Kinetic rules can be automatically generated
- 2 reactions with more than “1 substrate - 1 product”
 - kegg2symb already deals with all kind of reactions
 - Allosteric kinetics

1 accepted poster @ ILP'09 (Leuven), 6 pages

1 submission (rejected) @ DS'09 (Porto), 14 pages

Thanks

Many thanks go to `shuffle(thanks)`:
(>'-'> (Katsumi Inoue, Andrei Doncescu, Taisuke Sato, Pierre Bessière, Yoshitaka Kameya, Yoshitaka Yamamoto, Takehide Soh, Nicolas Dumazet, Elsa Prieto, (all_my_friends))).

Thank you for your attention.

Arigato gozaimasu.

Any questions?

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