Une sémantique pour la biologie moléculaire ?

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Scientific inputs to systems biology

- Physics: population dynamics (Monte carlo), free energy (protein folding)...
- Chemistry (obviously)
- Computer Science (essentially algorithmics)
- Can «semantics» (us) bring anything to them?

Total Publication volume?





How many papers should you read if you are a specialist of subject X?





Budget? (USA)



Budget 2012 of NIH : 31 billion \$ (~ total budget of France research!) Budget 2012 of NIST : 750 million \$ (source: nih.giv and nist.gov)



No theorems, only facts...

PTOV1 antagonizes MED25 in RAR transcriptional activation

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No theorems, only facts...

domains (Fig. 1A). MED25 also contains the VWA domain, which is required for mediator binding, and the NR box, which is required for RAR binding. In line with our previous study [13], yeast two-hybrid assays showed that the PTOV domain (amino acid [aa] residues 395–545) of MED25 interacted with the N-terminal region (aa 1–460) of CBP, which is a co-activator with histone acetyltransferase (HAT) activity (Fig. 1B). In similar fashion, the region of PTOV1 that contains the second PTOV domain (aa 253–416) was found to be responsible for CBP binding. To confirm the CBP interactions with MED25 and PTOV1 *in vivo* and *in vitro*, we performed immunoprecipitation (IP) and GST pull-down assays, respectively. For the IP assay, H1299 cells were co-transfected with HA-tagged CBP and Flag-tagged MED25 or Flag-tagged PTOV1. IP with an anti-HA antibody and subsequent Western blotting (WB) with an anti-Flag antibody demonstrated that both MED25 and PTOV1 interacted with CBP (Fig. 1C). The *in vivo* interaction was further verified by IP using an anti-CBP antibody and WB with an anti-PTOV1 or anti-MED25 antibody (Fig. 1D). GST pull-down assays were performed using purified GST-fused CBP (aa 1-460), Histaggod MED25 and Uic taggod DTOV1 Dinding reactions

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What we would like

- Translate facts (in English) into formal rules
- Use machines to connect related facts (negative knowledge, redundancy...)
- Build models from rules and simulate them (stochastic simulation, ODEs generation)

What we can do...

Abstract interpretation

Graph rewriting

Stochasticity/CTMC

Intrinsic Information Carriers in Combinatorial Dynamical Systems

Russ Harmer,^{1,2} Vincent Danos,³ Jérôme Feret,⁴ Jean Krivine,² and Walter Fontana¹

Internal coarse-graining of molecular systems

Jérôme Feret *, Vincent Danos [†], Jean Krivine *, Russ Harmer [‡], and Walter Fontana * *Harvard Medical School, Boston, USA,[†]University of Edinburgh, Edinburgh, United Kingdom, and [‡]CNRS & Paris Diderot, Paris, France Submitted to Proceedings of the National Academy of Sciences of the United States of America

Graphs, Rewriting and Pathway Reconstruction for Rule-Based Models

Vincent Danos³, Jérôme Feret⁴, Walter Fontana⁵, Russell Harmer¹, Jonathan Hayman^{4,2}, Jean Krivine¹, Chris Thompson-Walsh², and Glynn Winskel²

Rule-based modelling of (

Vincent Danos^{1,3,4}, Jérôme Feret², Walter Fo Jean Krivine⁵

Scalable simulation of cellul

Vincent Danos^{1,4 \star}, Jérôme Feret³, Walter

KaSim 3.4

Representation issue

"A model should be a data structure that contains a transparent, formal, and executable representation of the facts it rests upon"

Fontana, POPL'08

Can we do this?



Operational semantics for the biologist!

From 2002 International AIDS Conference in Barcelona, GlaxoSmithKline commissioned movie on the life cycle of HIV

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Cellular machinery



Protein-protein interaction



Protein-DNA interaction



Diffusion



Membrane fusion

Membrane fission Static compartments

Cellular machinery



Protein-protein interaction

Protein-DNA interaction

Diffusion



A language factory...

- **C0: forming molecules** untyped basic reactions
- Cl: naming molecules names as a refinements
- C2: placing molecules compartments
- C3: moving molecules the diffusion problem





C0: Forming molecules

$$D, D' ::= \mathsf{D}^a(x_1, \dots, x_k) \quad \text{for } a \in \mathcal{B}, \ x_i \in \mathcal{S}$$

 $T, S \quad ::= D \mid 0 \mid (T, S) \mid T \setminus v \text{ for } v \in S \cup B$



$$D, D' ::= \mathbf{O}^{a}(x_{1}, \dots, x_{k}) \quad \text{for } a \in \mathcal{B}, \ x_{i} \in \mathcal{S}$$
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$$D, D' ::= \bigcirc^{a} (\bigcirc, \dots, \bigcirc) \quad \text{for } a \in \mathcal{B}, \ x_{i} \in \mathcal{S}$$
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$$D, D' ::= \bigcirc (0), \dots, (0) \quad \text{for } a \in \mathcal{B}, \ x_i \in \mathcal{S}$$
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Terms $D, D' ::= \bigcirc (x_1, \ldots, x_k) \quad \text{for } a \in \mathcal{B}, \ x_i \in \mathcal{S}$ $T, S \quad ::= D \mid 0 \mid (T, S) \mid T \setminus v \text{ for } v \in \mathcal{S} \cup \mathcal{B}$ $(S,T) \equiv (T,S)$ $(T,S), T' \equiv T, (S,T')$ $T, 0 \equiv T$ $T \backslash u \equiv T \quad u \not\in fn(T)$ $(T \setminus u) \setminus v \equiv (T \setminus v) \setminus u$ $T = (T (... / ...)) = ... \neq f_{m} (T)$

Graphical notation

$$fn (D^{a}(x_{1},...,x_{k})) = \bigcup_{i} x_{i} \cup \{a\}$$

$$fn(0) = \emptyset$$

$$fn(T,S) = fn(T) \cup fn(S)$$

$$fn(T \setminus v) = fn(T) - \{v\}$$
free names
bound names
bound names
shared names





 $S = \left(\mathsf{D}^{a}(x), \mathsf{D}^{a}(x)\right) \backslash x$



 $U = (\mathsf{D}^{a}(x, x), \mathsf{D}^{a}(z), \mathsf{D}^{a}()) \setminus a \setminus x$

Dynamics

Contexts:

 $\mathbb{C}[\bullet] ::= \bullet \mid \mathbb{C}[\bullet] \setminus u \mid \mathbb{C}[\bullet], T \mid \mathbb{C}[\bullet] \{u/v\} \quad u, v \in \mathcal{B} \cup \mathcal{S}$

Context free rewriting:

$$\frac{r = \langle T, S \rangle \in \mathcal{R} \quad T' \equiv \mathbb{C}[T] \quad S' \equiv \mathbb{C}[S]}{T' \to_r S'}$$

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A rule

Dynamics

Contexts:

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Context free rewriting:

$$\underbrace{r = \langle T, S \rangle \in \mathcal{R} \quad T' \equiv \mathbb{C}[T]}_{T' \to_r S'} \quad S' \equiv \mathbb{C}[S]$$

A rule A match for T in T'
Dynamics

Contexts:

 $\mathbb{C}[\bullet] ::= \bullet \mid \mathbb{C}[\bullet] \setminus u \mid \mathbb{C}[\bullet], T \mid \mathbb{C}[\bullet] \{u/v\} \quad u, v \in \mathcal{B} \cup \mathcal{S}$

Context free rewriting:



Generators



Fig. 1: Generators for \mathcal{C}_0 .

Generating rules



Generators toolbox

A rule of the model

(using compose and refine)



Names as refinements

 $\begin{array}{lll} D, D' ::= \mathsf{D}_{\mathcal{D}}^{a}(x_{1}, \dots, x_{k}) & a \in \mathcal{B}, \ m \in \mathcal{M}, \ x_{i} \in \mathcal{S} \ \text{(domains)} \\ I, J & ::= \mathsf{Inform} & \mathsf{info} \in \mathcal{I}, \ m \in \mathcal{M} & (\mathsf{info}) \\ T, S & ::= 0 \mid D \mid I \mid (T, S) \mid T \setminus v \ \text{for} \ v \in \mathcal{S} \cup \mathcal{B} \cup \mathcal{M} & (\mathsf{named terms}) \end{array}$

 $\mathsf{D}_m^a(x)$ A domain

 $D_m^a(x)$, Tyrosine_m A Tyrosine domain

 $D_m^a(x), Tyr_m, phos_m$ A pho'ylated Tyrosine domain



(Concretize) $\mathsf{D}_m^a(x_1, \dots, x_k) \to \mathsf{D}_m^a(x_1, \dots, x_k)$, Info_m (Abstract) $\mathsf{D}_m^a(x_1, \dots, x_k)$, $\mathsf{Info}_m \to \mathsf{D}_m^a(x_1, \dots, x_k)$)





C2: Placing molecules

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Terms

$T, S ::= \cdots \mid \mathsf{C}_m(T) \mid X \quad m \in \mathcal{M}, \ X \in \mathcal{V} \text{ (local terms)}$ $P, Q ::= (T \parallel P) \mid P \setminus v \quad v \in \mathcal{M} \cup \mathcal{S} \cup \mathcal{B} \quad \text{(wide terms)}$



















Generators I/2



 $0 \parallel 0 \rightarrow (\mathsf{channel}_m \parallel \mathsf{C}(\mathsf{channel}_m)) \backslash m$





 $(\mathsf{D}_n^{hiv}(x),\mathsf{D}_o^{tcell}(x))\backslash x$ D_m^{hiv} tcell **HIV** membrane T-cell membrane









 D_m^{hiv} $(\mathsf{D}_n^{hiv}(x),\mathsf{D}_o^{tcell}(x))\backslash x$ tcell T-cell membrane **HIV** membrane













Extension relation

(ax.)
$$\frac{|\sigma| = \mathcal{V}(T)}{T \hookrightarrow_{\bullet} \langle \mathbb{C}[\bullet], \sigma \rangle} \qquad \frac{P \hookrightarrow_{\pi} \langle T_{\bullet}, \sigma \rangle \quad m \text{ fresh } (\!\!|\cdot\pi\cdot|\!\!) \not\rightarrow \bot}{P \hookrightarrow_{(\!|\pi\!|)} \langle \mathsf{C}_m(T_{\bullet}), \sigma \rangle} \quad (\text{wrap})$$
$$\frac{P \hookrightarrow_{\pi_0} \langle T_{\bullet}, \sigma \rangle \quad Q \hookrightarrow_{\pi_1} \langle S_{\bullet}, \sigma' \rangle \quad \pi_0 \cdot \pi_1 \not\rightarrow \bot}{P \parallel Q \hookrightarrow_{\pi_0 \pi_1} \langle \mathbb{C}[T_{\bullet}, S_{\bullet}], \sigma; \sigma' \rangle} \quad (\text{comp})$$

Local contexts:

$$\mathbb{C}[\bullet] ::= \bullet \mid \mathbb{C}[\bullet] \setminus u \mid \mathbb{C}[\bullet], T \mid \mathbb{C}[\bullet] \{u/v\}$$

Compatibility relation:

$$\bullet \cdot (\! | \cdot (\! | \rightarrow \bot \qquad) \cdot)\! | \cdot \bullet \rightarrow \bot \qquad) \cdot (\! | \rightarrow \bot$$

 $\bullet \cdot \bullet \to \bot \qquad \bullet \cdot (\!\! (\cdot \bullet \cdot)\!\!) \cdot \bullet \to \bot \qquad \bot \cdot \pi \to \bot \qquad \pi \cdot \bot \to \bot$

Extension relation

(ax.)
$$\frac{|\sigma| = \mathcal{V}(T)}{T \hookrightarrow_{\bullet} \langle \mathbb{C}[\bullet], \sigma \rangle} \qquad \frac{P \hookrightarrow_{\pi} \langle T_{\bullet}, \sigma \rangle \quad m \text{ fresh } (\!\!|\cdot\pi \cdot \!\!|) \not\rightarrow \bot}{P \hookrightarrow_{(\!(\pi)\!)} \langle \mathbb{C}_m(T_{\bullet}), \sigma \rangle} \quad (\text{wrap})$$
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$$\text{.ocal contexts:} \qquad \text{Term abstraction! } \Pi \stackrel{def}{=} \{(\!\!(, \!\!)), \bullet, \bot\}$$

 $\mathbb{C}[\bullet] ::= \bullet \mid \mathbb{C}[\bullet] \setminus u \mid \mathbb{C}[\bullet], T \mid \mathbb{C}[\bullet] \{u/v\}$

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 $\bullet \cdot \bullet \to \bot \qquad \bullet \cdot (\! | \cdot \bullet \cdot)\! | \cdot \bullet \to \bot \qquad \bot \cdot \pi \to \bot \qquad \pi \cdot \bot \to \bot$

Definition 4 (Matches). A wide context $\mathbb{C}^n[\bullet, \ldots, \bullet]$ with exactly n holes and a parameter assignation list σ form a match $\langle \mathbb{C}^n, \sigma \rangle$ for a wide term $P = T_1 \parallel \cdots \parallel T_n$ in S if and only if:



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Results

Theorem 1 (Soundness). Let $\langle \mathbb{C}^n[\bullet, \ldots, \bullet], \sigma \rangle$ be a match for a wide term P in a local term T. For all disjoint local term occurrences $S, S' \in P$, we have $\Delta_{S,S'}(P) = \Delta_{S,S'}(T)$.

Theorem 2 (Completeness). Let $P = T_1 \parallel \cdots \parallel T_n$ be a wide term and $\mathbb{C}^n[\bullet, \ldots, \bullet]$ be a generic context with exactly n holes. Let also $T \equiv \mathbb{C}^n[T_1, \ldots, T_n]\sigma$ for some parameter assignation σ . If for all $i, j \leq n$ one has $\Delta_{T_i, T_j}(P) = \Delta_{T_i, T_j}(T)$, then $P \hookrightarrow_{\pi} \langle \mathbb{C}^n, \sigma \rangle$ is derivable, for some $\pi \in (\Pi \setminus \{\bot\})^*$.



C3: Moving molecules

Terms

$$T, S ::= \dots \qquad (\text{local terms})$$

$$G, H ::= T \mid \mathsf{spec}_{\mathsf{S}}^{\mathsf{B}}(T) \mid (G, H) \text{ (global terms)}$$

$$P, Q ::= G \mid (P \parallel Q) \mid \dots \qquad (\text{wide terms})$$

$$\overline{\mathsf{D}^{a}(x_{1},\ldots,x_{k})} \equiv \mathsf{spec}_{\{x_{1},\ldots,x_{k}\}}^{\{a\}}(\mathsf{D}^{a}(x_{1},\ldots,x_{k}))$$

$$\frac{fn(D) \cap (\mathsf{B} \cup \mathsf{S}) \neq \emptyset \quad \mathsf{B}' = \mathsf{B} \cup (fn(D) \cap \mathcal{B}) \quad \mathsf{S}' = \mathsf{S} \cup (fn(D) \cap \mathcal{S})}{\mathsf{spec}_{\mathsf{S}}^{\mathsf{B}}(T), D \equiv \mathsf{spec}_{\mathsf{S}'}^{\mathsf{B}'}(T, D)}$$

$$\frac{fn(T') \cap (\mathsf{B} \cup \mathsf{S}) \neq \emptyset \quad \mathsf{B}' = \mathsf{B} \cup (fn(T') \cap \mathcal{B}) \quad \mathsf{S}' = \mathsf{S} \cup (fn(T') \cap \mathcal{S})}{\mathsf{spec}_{\mathsf{S}}^{\mathsf{B}}(T), \mathsf{C}(T') \equiv \mathsf{spec}_{\mathsf{S}'}^{\mathsf{B}'}(T, \mathsf{C}(T'))}$$

$$\frac{u \in \mathcal{B} \cup \mathcal{S} \quad \mathsf{B}' \stackrel{def}{=} \mathsf{B} - \{u\} \quad \mathsf{S}' \stackrel{def}{=} \mathsf{S} - \{u\}}{\mathsf{spec}_{\mathsf{S}}^{\mathsf{B}}(T) \backslash u \equiv \mathsf{spec}_{\mathsf{S}'}^{\mathsf{B}'}(T \backslash u)} \quad \frac{T \equiv T'}{\mathsf{spec}_{\mathsf{S}}^{\mathsf{B}}(T) \equiv \mathsf{spec}_{\mathsf{S}}^{\mathsf{B}}(T')}$$



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Generators


Definition 5 (Mixture). Say that a term P is a mixture if: -w(P) = 1, $fn(P) = \emptyset$ and P is parameter free

- Site edges have exactly two sites and do not cross compartments
- Backbone hyper edges cross at most one compartment
- Each species node contains a single connected component

Property 1 (Preservation). Let \mathcal{R} be a set of generated rules and let P be a mixture. If $P \to_r Q$ with $r \in \mathcal{R}$ then Q is a mixture.

Summary

- A language expressive enough to model a large swath of systems biology in an algebraic fashion
- Relies on a minimal set of generators (thanks to projectivity)
- Simulating generators is enough (the rest are refinements)!

«I took biology because I didn't want to do math»

a biologist at HMS