Al and Theorem-Proving for Science:

## Towards AI for mathematical modeling of complex physical and biological systems

with aspects of "Feynman on AI ..." book chapter

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Artificial Intelligence and Theorem Proving (AITP) Conference September 2023

## Some aspects of the "Feynman on AI" chapter

- Back-story: lunchtime student+TA discussions with RPF in the early to mid-1980s, often including EM, MD
- in which RPF evinced an interest in a machine learning approach to AI, with a progressive sequence of tasks
- [EM, Feynman on Artificial Intelligence and Machine Learning.] - my considered version
- Includes tutorial derivations omitted here

#### Symbolic algebraic expressions: Key to new computing paradigms, and a hidden currency of ML

- Gibbs-Bogolubov-Feynman variational method
  - $F \leq \langle H \rangle_0 TS_0$
- Hopfield neural network

• 
$$F \leq E_{\text{MFT}}[v] \equiv -\frac{1}{2} \sum_{i \neq j} T_{ij} v_i v_j - \sum_i h_i v_i + \sum_i \varphi(v_i), \quad v_i \equiv \langle s_i \rangle_0 \in \{0,1\}.$$
  
•  $\varphi(v) = \frac{1}{\beta} \Big( v \log v + (1-v) \log(1-v) \Big) = -TS_{\text{MFT}}[v] \Rightarrow \text{ activation } v_i = \tanh \Big[ \beta \Big( u_i = \sum_j T_{ij} v_j + h_i \Big) \Big].$   
Generalized to multiway 0/1 choice:  $\varphi([v_{i+1}]) = \frac{1}{2} \sum_{i \neq j} v_i \log v_i = -T\tilde{S}[v] \Rightarrow \text{ softmax } v_i = e^{\beta u_{ia}} / \sum_i e^{\beta u_{ib}}.$ 

Generalized to multiway 0/1 choice:  $\varphi([v_{i\star}]) = \frac{1}{\beta} \sum_{a} v_{ia} \log v_{ia} = -T\tilde{S}[v] => \text{ softmax } v_{ia} = e^{\beta u_{ia}} / \sum_{b} e^{\beta u_{ib}}$ .

• Boltzmann Machine  $H_{\text{Ising}}[s] \equiv -\frac{1}{2} \sum_{i \neq j} T_{ij} s_i s_j - \sum_i h_i s_i$ ; learn by minimizing KL divergence to external p(s).

• Quantum computer, per Feynman  $H = \sum_{i=0}^{k-1} q_{i+1}^* q_i A_{i+1}$ .

• Transformer neural network: Inner softmax  $v_{ia} = e^{\beta u_{ia}} / \sum_{b} e^{\beta u_{ib}}$ 

[EM, Feynman on Artificial Intelligence and Machine Learning. Feynman Lectures on Computation: Anniversary Edition; arXiv:2209.00083]

## Symbolic vs. Neural Al

- Feynman lunchtime discussions, early 1980s:
  - G. Sussman (Minsky colleague) vs. (ideas of) J. Hopfield
  - Feynman expressed a NN research program
- Symbolic was ascendant, but experiencing AI winter
- Now reversed:
  - Last 10-15 years: AAAI dominated by ML/neural networks
- Future: Hegelian Synthesis?

## Quoting my own best-effort text ...

1/2

• "What would Feynman have thought of all all these advances? Remembering his commitment to having a personal, independent point of view on everything, and the impossibility of emulating a great mind, there can be little certainty on this point. Nevertheless it seems to me that: ..."



• "3. He would be enthusiastic about transfer learning, and about the particular neural architectures that have led to success in vision, natural language, and limited combinations thereof. These methods have essentially achieved the goals of the machine learning project that he had in mind - insofar as he had expressed them."



• "4. Regarding equations as the hidden currency of learning architecture, he might be of two minds: attracted because he was a mathematical master; repelled because the source of his mastery was the ability to deeply visualize the meaning of each equation. Neural network equations for the most part are just not that conceptually deep. In the old AI dichotomy of "scruffy" vs. "neat" research approaches, ML equations might be neat but not neat enough."

## Quoting my own best-effort text ...

2/2

- "What would Feynman have thought of all all these advances? ...." <....Caveats! ...>
  - "6. He might nevertheless be intrigued by conceptually deeper, less *ad hoc* architectures that connect to physics, such as manifold learning (clearly related to general relativity), and perhaps by attempts to connect neural networks to real neurobiology."
  - "8. Because of his deep commitment to physics, Feynman would be quite interested in - if also a bit skeptical of - the "physics-informed machine learning" agenda which is now being pursued with many different architectures, representations, and methods for the purpose of doing computational physics, chemistry, and biology. If I had to pick one thing about present-day neural networks, machine learning, and AI that Richard Feynman would be most interested in, this would be it."
  - 9. He would *certainly* have creative and potentially powerful new ideas that are not yet on anybody's list or agenda."

### AITP for Science: some logic in favor

- "AI": Reliable AI for Science:
  - Representationalism 1st, ML 2nd. (On top vs. on tap)
  - formal, symbolic DSLs for modeling ...
    - applicable math, sci content, algorithms (model sim, ML, analysis)
- "TP": verify exact and approximate maps between such languages
  - e.g. dynamics to simulation algs: XDEs, Markov jump processes, hybrids e.g. *dynamical grammars*
  - high-level support for scientific knowledge representation

### """ "Tchicoma" Architecture for Mathematical Modeling

To formalize:

Vathematics

+ mappings and hierarchies:



+arXiv:1804.11044]

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## Some scientific simulation paradigms

- Partial differential equations (PDEs)
  - central to PIML so far
- Stochastic models
  - kinetic theory in physics
  - well-motivated in biology

## PDE verification?

- ODEs: Lorenz attractor [Immler 2017]
- Finite element method foundation: Lax-Milgram theorem on unique existence of FEM weak formulation solutions

### Master Equation Semantics

 $A_1(x_1), A_2(x_2), \ \dots, A_n(x_n) \to B_1(y_1), B_2(y_2), \ \dots, \ B_m(y_m) \text{ with } \rho(\{x_i\}, \{y_j\})$ 

- RHS, LHS are multisets
- Founded on **stochastic** processes
- Dynamics from the Master Equation:

- Term creation/annihilation operators: for each param value,

$$\hat{a} = \begin{pmatrix} 0 & 0 & 0 & 0 & \cdots \\ 1 & 0 & 0 & 0 & \\ 0 & 1 & 0 & 0 & \\ \vdots & & \ddots & \ddots \end{pmatrix} = \delta_{n,m+1} \text{ and } a = \begin{pmatrix} 0 & 1 & 0 & 0 & \cdots \\ 0 & 0 & 2 & 0 & \\ 0 & 0 & 0 & 3 & \\ 0 & 0 & 0 & 0 & \ddots \\ \vdots & & & \ddots \end{pmatrix} = m \, \delta_{n+1,m}$$

- Obeying Heisenberg **operator algebra** 

$$[a, \hat{a}] \equiv (a \ \hat{a} - \hat{a} a) = I$$

$$[a(x), \hat{a}(y)] = \delta(x - y) \left[ I + N Q(N \mid n^{(\max)}) \right]$$

- Yet classical, not quantum, probabilities

[Doi 1976; ... Mikhailov '81; ...Mattis & Glasser '98; ... EM+GY Annals of Math. and A. I., 47(3-4), January 2007]



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Example: 
$$\{A + B \xrightarrow{k_f} C, C \xrightarrow{k_r} A + B\}$$

• Operator:

 $H = k_f \left( \hat{a}_3 \; a_1 \; a_2 \; - \; N_1 \; N_2 \; I_3 \right) + k_r \left( \hat{a}_1 \; \hat{a}_2 \; a_3 \; - \; I_1 \; I_2 \; N_3 \right)$ 

- Spectrum:
- EV's with boundary layer approximation:



[Mjolsness and Prasad, J. Chem Phys. 2013]

### Operator algebra for Pure stochastic chemical reactions

• For reaction/rule *r*:

 $\hat{W}_{\{m_i^{(r)}\}\to\{n_i^{(r)}\}} = k^{(r)} \prod_i (\hat{a}_i)^{n_i^{(r)}} (a_i)^{m_i^{(r)}}$ 

$$n_{\alpha} \in \mathbb{N} : \begin{bmatrix} [a_{\alpha}, \hat{a}_{\beta}] = \delta_{\alpha\beta}I \\ a_{\alpha}\hat{a}_{\beta} = \hat{a}_{\beta}a_{\alpha} + \delta_{\alpha\beta}I_{\alpha} \end{bmatrix}$$
  
$$n_{\alpha} \in \{0, 1\} : a_{\alpha}\hat{a}_{\beta} = (1 - \delta_{\alpha\beta})\hat{a}_{\beta}a_{\alpha} + \delta_{\alpha\beta}Z_{\alpha}$$

• For reaction/rules  $r_1$  and  $r_2$ :

where 
$$(n)_l \equiv \begin{cases} n!/(n-l)! & \text{for } l \le n; \\ 0 & \text{otherwise} \end{cases}$$

$$\begin{split} \hat{W}_{\{m_i^{(r_2)}\} \to \{n_i^{(r_2)}\}} \hat{W}_{\{m_i^{(r_1)}\} \to \{n_i^{(r_1)}\}} &= k^{(r_2)} k^{(r_1)} \sum_{\{l_i = 0 \dots \min(m_i^{(r_2)}, n_i^{(r_1)})\}} \left(\prod_i \frac{(m_i^{(r_2)})_l (n_i^{(r_1)})_l}{l_i!}\right) \\ &\times \hat{W}_{\{(m_i^{(r_1)} + m_i^{(r_2)} - l_i)\} \to \{(n_i^{(r_1)} + n_i^{(r_2)} - l_i)\}} \end{split}$$

Why:  $\partial_x^m(x^n f(x)) = \text{binomial sum } (+ \text{Weyl algebra})$ 

#### Generation of valid algorithms

- Compute or sample exp *t H* ; e.g. Euler's formula
- Approximate: Trotter Product formula (interleaving):

$$e^{tH} = \lim_{n \to \infty} \left( 1 + \frac{t}{n} H \right)^{t}$$

$$= \lim_{n \to \infty} \left[ e^{(t/n)H_0} e^{(t/n)H_1} \right]^n$$

• Time-ordered product expansion (TOPE) - convergent Dyson series :

$$e^{tH} = \sum_{k=0}^{\infty} \left[ \int_{0}^{t} dt_{k} \int_{0}^{t_{k}} dt_{k-1} \cdots \int_{0}^{t_{2}} dt_{1} \exp((t-t_{k})H_{0})H_{1} \exp((t_{k}-t_{k-1})H_{0}) \cdots H_{1} \exp(t_{1}H_{0}) \right]$$
[Annals of Math. and A. I., 47(3-4), January 2007]

- can be used recursively !
- restate with time-ordering:

[Physical Biology 10, 2013]

$$\exp(t \left(W_0 + W_1\right)) = \exp(t W_0) \left(\exp\left(\int_0^t \exp(-\tau W_0) W_1 \exp(\tau W_0) d\tau\right)\right)_+$$
$$\equiv \exp(t W_0) \left(\exp\left(\int_0^t W_1 (\tau) d\tau\right)\right)_+$$

• Diagonal/off-diagonal split: obtain Gillespie's Stochastic Simulation Algorithm (SSA)

$$\mathcal{W}(I, t' \mid J, t) \approx \hat{W}_{I,J} \exp(-(t' - t) D_{JJ}) \mathbf{1} (t' \ge t)$$

$$\Pr(. \mid J, k) = \mathcal{W}^k \circ \Pr(. \mid J, 0) = \left[ \hat{W} \exp(-\Delta t D) \mathbf{1} \left( \Delta t \ge 0 \right) \right]^k \circ \Pr(. \mid J, 0)$$

#### TOPE and Feynman diagrams *Time-Ordered Product Expansion*



Figure 1. A time history of the reaction  $A + B \rightleftharpoons C$ . Time flows left to right. Solid blocks represent reaction events, with probability factor  $\times W_1$ . In between reaction events are unimolecular particle propagators  $\exp((t_k - t_{k-1}) W_0)$ , labelled by arrows and particle names (repeated for clarity). This is a non-spatial version of the Lee model in quantum field theory (cf. for example [Bender et al. 2005]).

[Physical Biology 10, 2013]

#### Time-Ordered Product Expansion (TOPE) verification? 1/2

- "Big operator" derivation
- formal power series, changes of variable, induction on dimension
- notation "represents" permutation symmetries

$S(z) = \exp(t \left(H_1 z + H_0\right)) \cdot p \exp(s_k z^k)$
$=\sum_{k=0}^{\infty}\frac{z^{k}}{k!}\left[\partial_{z}^{k}\exp(t\left(H_{1}z+H_{0}\right))\right]_{z=0}\cdot p_{0}$
$= \sum_{k=0}^{\infty} \frac{z^{k}}{k!} \left[ \partial_{z}^{k} \sum_{l=0}^{\infty} \frac{(t (H_{1}z + H_{0}))^{l}}{l!} \right]_{z=0} \cdot p_{0}$
$=\sum_{k=0}^{\infty} \frac{z^{k}}{k!} \left[ \sum_{l=k}^{\infty} \frac{1}{l!} \sum_{\substack{\{0 \leqslant i_{p} \leqslant l-k\} \land \sum_{p=0}^{k} i_{p}=l-k}} k! (tH_{0})^{i_{k}} t \hat{H}(tH_{0})^{i_{k-1}} \cdots t \hat{H}(tH_{0})^{i_{0}} \right] \cdot p_{0}$
$=\sum_{k=0}^{\infty} z^{k} t^{k} \left[ \sum_{l=0}^{\infty} \frac{1}{(l+k)!} \sum_{\substack{\{0 \leqslant i_{p} \leqslant l\} \land \sum_{p=0}^{k} i_{p} = l}} (tH_{0})^{i_{k}} H_{1}(tH_{0})^{i_{k-1}} \cdots H_{1}(tH_{0})^{i_{0}} \right] \cdot p_{0}$
$=\sum_{k=0}^{\infty} z^{k} t^{k} \left[ \sum_{l=0}^{\infty} \sum_{\substack{\{0 \leq i_{p} \leq l\} \land \sum_{p=0}^{k} i_{p}=l}} \frac{\prod_{p=0}^{k} (i_{p})!}{\left(\sum_{p=0}^{k} i_{p}+k\right)!} \frac{(tH_{0})^{i_{k}}}{(i_{k})!} H_{1} \frac{(tH_{0})^{i_{k-1}}}{(i_{k-1})!} \cdots H_{1} \frac{(tH_{0})^{i_{0}}}{(i_{0})} \right] p_{0}$
$=\sum_{k=0}^{\infty} z^{t} t^{t} \left[ \sum_{\{0 \leq i_{p} \leq \infty\}} \frac{\prod\limits_{p=0}^{k} (i_{p})!}{\left(\sum\limits_{p=0}^{k} (i_{p}+1)-1\right)!} \frac{(tH_{0})^{i_{k}}}{(i_{k})!} H_{1} \frac{(tH_{0})^{i_{k-1}}}{(i_{k-1})!} \cdots H_{1} \frac{(tH_{0})^{i_{0}}}{(i_{0})!} \right] \cdot p_{0}$
$=\sum_{k=0}^{\infty} z^{k} t^{k} \left[ \sum_{\{0 \leqslant i_{p} \leqslant \infty\}} \frac{\prod\limits_{p=0}^{k} \Gamma(i_{p}+1)}{\Gamma\left(\sum\limits_{p=0}^{k} (i_{p}+1)\right)} \frac{(tH_{0})^{i_{k}}}{(i_{k})!} H_{1} \frac{(tH_{0})^{i_{k-1}}}{(i_{k-1})!} \cdots H_{1} \frac{(tH_{0})^{i_{0}}}{(i_{0})!} \right] \cdot p_{0}$

Line #

(3)

(4)

Now we use the Multinomial-Dirichlet normalization integral

$$\frac{\prod\limits_{p=0}^{n}\Gamma(i_{p}+1)}{\Gamma\left(\sum\limits_{p=0}^{n}\left(i_{p}+1\right)\right)}=\int_{0}^{1}d\theta_{0}\cdots\int_{0}^{1}d\theta_{k}\delta\left(\sum\limits_{p=0}^{k}\theta_{p}-1\right)\prod\limits_{p=0}^{k}\left(\theta_{p}\right)^{i_{p}}.$$

🖄 Springer

17 [EM+GY Annals of Math. and A. I., 47(3-4), January 2007] Proof Outline: TOPE, line 3 => 4, 1/2

H?

$$\begin{split} & \underset{(I_{2}, i_{2}) \in \mathbb{N}}{\underset{(I_{2}, i_{2}$$

#### Human Justification for 3.3, ~ the Distributive Law for Big Operators

#### Machine justification for 3.3?

• In Coq: [Bertot et al, "Canonical Big Operators", TPHOLs 2008]

```
Lemma bigA_distr_bigA :
    forall (I J : finType) F,
    \big[*%M/1]_(i : I) \big[+%M/0]_(j : J) F i j =
    \big[+%M/0]_(f : {ffun I -> J}) \big[*%M/1]_(i) F i (f i).
```

• In Lean 4: Mathlib.Algebra.BigOperators.Ring (?)

**theorem Finset.prod\_sum** { $\alpha$  : Type u} { $\beta$  : Type v} [inst : CommSemiring  $\beta$ ] { $\delta$  :  $\alpha \rightarrow$  Type u\_1} [inst : DecidableEq  $\alpha$ ] [inst : (a :  $\alpha$ )  $\rightarrow$  DecidableEq ( $\delta$  a)] {s : Finset  $\alpha$ } {t : (a :  $\alpha$ )  $\rightarrow$  Finset ( $\delta$  a)} {f : (a :  $\alpha$ )  $\rightarrow \delta$   $a \rightarrow \beta$ } : (Finset.prod s fun a => Finset.sum (t a) fun b => f a b) = Finset.sum (Finset.pri s t) fun p => Finset.prod (Finset.attach s) fun x => f ( $\uparrow$ x) (p  $\uparrow$ x (\_ :  $\uparrow$ x  $\in$  s))

The product over a sum can be written as a sum over the product of sets, Finset.Pi.

https://leanprover-community.github.io/mathlib4\_docs/Mathlib/Algebra/BigOperators/Ring.html#Finset.prod\_sum

Proof Outline: TOPE, line  $3 \Rightarrow 4$ , 2/2

Expressive for map EJSI is if 
$$j_{0}^{p}$$
 by inductions on p:  
 $p=0$  base case:  $\int_{1}^{2} Such dhet n$   
 $\hat{h}_{0} \in Sp_{0}...,k_{3}^{p}$  for  $j_{1},...,j_{n} = 0$   $\Lambda(J_{2n+1} = 1 \quad \forall \quad \dot{h}_{0} = k)$   
 $p > 1$  induction, up to  $p = k - 1$ :  
 $\int_{1}^{2} \frac{g^{p}(\lambda_{q} + 1)}{g^{2n}} \leq g$  // such ling the of  $H_{0}^{in}(H, ..., H_{0}^{in}(H, ..., H_{0}^{in}$ 

2

#### Time-Ordered Product Expansion (TOPE) verification? 2/2

- "Big operator" derivation
- formal power series, changes of variable, induction on dimension
- notation "represents" permutation symmetries

$$\begin{split} S(z) &= \sum_{k=0}^{\infty} z^{k} t^{k} \left[ \sum_{\{0 \leq i_{p} \leq \infty\}} \int_{0}^{1} d\theta_{0} \cdots \int_{0}^{1} d\theta_{k} \delta\left(\sum_{p=0}^{k} \theta_{p} - 1\right) \left(\prod_{p=0}^{k} (\theta_{p})^{i_{p}}\right) \\ &\quad \times \frac{(tH_{0})^{i_{k}}}{(i_{k})!} H_{1} \frac{(tH_{0})^{i_{k-1}}}{(i_{k-1})!} \cdots H_{1} \frac{(tH_{0})^{i_{0}}}{(i_{0})!} \right] \cdot p_{0} \\ &= \sum_{k=0}^{\infty} z^{k} t^{k} \left[ \sum_{\{0 \leq i_{p} \leq \infty\}} \int_{0}^{1} d\theta_{0} \cdots \int_{0}^{1} d\theta_{k} \delta\left(\sum_{p=0}^{k} \theta_{p} - 1\right) \\ &\quad \times \frac{(\theta_{k} tH_{0})^{i_{k}}}{(i_{k})!} H_{1} \frac{(\theta_{k-1} tH_{0})^{i_{k-1}}}{(i_{k-1})!} \cdots H_{1} \frac{(\theta_{0} tH_{0})^{i_{0}}}{(i_{0})!} \right] \cdot p_{0} \\ &= \sum_{k=0}^{\infty} z^{k} t^{k} \left[ \int_{0}^{1} d\theta_{0} \cdots \int_{0}^{1} d\theta_{k} \delta\left(\sum_{p=0}^{k} \theta_{p} - 1\right) \sum_{\{0 \leq i_{0} \leq \infty\}} \frac{(\theta_{k} tH_{0})^{i_{0}}}{(i_{k})!} H_{1} \\ &\quad \times \sum_{\{0 \leq i_{1} \leq \infty\}} \frac{(\theta_{k-1} tH_{0})^{i_{1}}}{(i_{k-1})!} \cdots H_{1} \sum_{\{0 \leq i_{k} \leq \infty\}} \frac{(\theta_{0} tH_{0})^{i_{k}}}{(i_{0})!} \right] \cdot p_{0} \\ &= \sum_{k=0}^{\infty} z^{k} t^{k} \left[ \int_{0}^{1} d\theta_{0} \cdots \int_{0}^{1} d\theta_{k} \delta\left(\sum_{p=0}^{k} \theta_{p} - 1\right) \exp(\theta_{k} tH_{0}) \\ &\quad \times H_{1} \exp(\theta_{k-1} tH_{0}) \cdots H_{1} \exp(\theta_{0} tH_{0}) \right] \cdot p_{0} \end{split}$$

Stochastic process semantics for dynamical grammars

Accordingly,

Thus

$$S(z) = \sum_{k=0}^{\infty} z^k \left[ \int_0^t d\tau_0 \cdots \int_0^t d\tau_k \delta\left(\sum_{p=0}^k \tau_p - t\right) \exp(\tau_k H_0) \right] \times H_1 \exp(\tau_{k-1} H_0) \cdots H_1 \exp(\tau_0 H_0) \right] \cdot p_0$$
(A59)

In summary (since  $p_0$  was never used in the above calculations),

$$\exp(t (H_1 + H_0)) = \sum_{k=0}^{\infty} \left[ \int_0^t d\tau_0 \cdots \int_0^t d\tau_k \delta\left(\sum_{p=0}^k \tau_p - t\right) \exp(\tau_k H_0) H_1 \exp(\tau_{k-1} H_0) \cdots H_1 \exp(\tau_0 H_0) \right].$$

$$222$$

[EM+GY Annals of Math. and A. I., 47(3-4), January 2007]

#### **UCI** Morphodynamics

## Learning in stochastic reaction networks



#### Stochastic Simulation

• SSA:  $\mathcal{W}(I, t' \mid J, t) \approx \hat{W}_{I,J} \exp(-(t'-t)D_{JJ})\mathbf{1}(t' \ge t)$ 

 $\Pr(. \mid J, k) = \mathcal{W}^k \circ \Pr(. \mid J, 0) = \left[ \hat{W} \exp(-\Delta t D) \mathbf{1} \left( \Delta t \ge 0 \right) \right]^k \circ \Pr(. \mid J, 0)$ 

• Exact R-leap:

$$\left[\prod_{k=L-1>0} \hat{W} \exp\left(-\tau_k D\right)\right]_{I_L, I_0} = \frac{\left(\tilde{D}_{I_0 L-1}\right)^L}{\left(\tilde{D}_{I_0 L-1}\right)^L} \sum_{\left\{s \mid s_r \in \mathbb{N}, \sum_r s_r = L\right\}} \text{Multinomial}(s \mid p, L)$$

$$\times \text{Erlang}\left(\sum_k \tau_k \left| L, D_{I_0 L-1} \right| \text{UniformSimplex}(\tau; L) \text{Accept}(s, L, \tau)$$



#### Rejection Sampling (in general)



Rejection sampling allows one to exploit probability bounds in exact sampling, as follows: given a target distribution P(x) and an algorithm for sampling from a related distribution P'(x) and from the uniform distribution U(u) on [0,1], and if

$$P(x) < M P'(x)$$

for some constant M > 1, then P(x) satisfies

$$P(x) = P'(x) \frac{P(x)}{MP'(x)} + (1 - 1/M)P(x)$$

and therefore also

$$P(x) = \int P'(x') \, dx' \, \int U(u) \, du \left[ \mathbf{1} \left( u < \frac{P(x')}{M \, P'(x')} \right) \cdot \delta(x - x') + \mathbf{1} \left( u \ge \frac{P(x')}{M \, P'(x')} \right) \cdot P(x) \right]$$

which constitutes a mixture distribution, that can be applied recursively as needed to sample from P(x). Pseudocode for sampling P(x) according to Equation 13 is as follows (where "//" introduces a comment):

while not accepted {
 sample P'(x) and U(u); // P'(x) only approximates P(x)
 compute Accept(x) = P(x) / (M P'(x)); // acceptance probability
 if u < Accept(x) then accept x;
} // now P(x) is sampled exactly</pre>

#### Accelerated Rejection Sampling ~ "Squeeze method" [Marsaglia 1961]

But what if P(x) is expensive to compute? Then Accept(x) will also be expensive to compute and rejection sampling may be prohibitively expensive, even for a good approximating P'(x). A solution to this problem is possible if a cheap lower bound for P(x) is available. Suppose there is a function A(x) such that

 $0 \leq \mathcal{A}(x) \leq \operatorname{Accept}(x) \equiv P(x)/(M \, P'(x)) < 1$ 

• *Then* Accept(*x*) is a mixture:

Accept(x) = 
$$\underline{A}(x) \cdot 1 + (1 - \underline{A}(x)) \cdot \underline{Q}(x)$$
, where  
(Accept(x) -  $\underline{A}(x)$ )

$$Q(x) \equiv \left(\frac{Accept(x) - \underline{A}(x)}{1 - \underline{A}(x)}\right),$$

• and we have the generic pseudocode algorithm:

```
while not accepted {

sample P'(x) and U(u); // cheap but approximate

compute \underline{A}(x); // cheap

if u < \underline{A}(x) then accept x;

else {

compute Accept(x) = P(x)/MP'(x); // expensive

compute Q(x) = (Accept(x) - \underline{A}(x))/(1 - \underline{A}(x)); // \underline{A}(x) < 1 \Rightarrow 1 - \underline{A}(x) \neq 0

sample U(u);

if u < Q(x) then accept x;

else reject x;

}
```

### AITP for Science: some logic in favor

- "AI": Reliable AI for Science:
  - Representationalism 1st, ML 2nd. (On top vs. on tap)
  - formal, symbolic DSLs for modeling ...
    - applicable math, sci content, algorithms (model sim, ML, analysis)
- "TP": verify exact and approximate maps between such languages
  - e.g. dynamics to simulation algs: XDEs, Markov jump processes, hybrids e.g. dynamical grammars
  - high-level support for scientific knowledge representation

## Variable-binding via operator integration

- Parameterized grammar rules:  $\{\tau_i(x_i)\} \rightarrow \{\tau'_j(y_j)\}$  with  $\rho_r((x_i), (y_j))$
- Parameterized grammar rule operators:

$$\hat{O}_{r} = \int_{D_{\beta(1)}} \dots \int_{D_{\beta(c)}} \dots \left( \prod_{c} d\mu_{\beta(c)}(X_{c}) \right) \rho_{r} \left( [x_{i}(\{X_{c}\})], [y_{j}(\{X_{c}\})] \right) \\ \times \left[ \prod_{i \in rhs(r)} \hat{a}_{a(i)}(x_{i}(\{X_{c}\})) \right] \left[ \prod_{j \in lhs(r)} a_{b(j)}(y_{j}(\{X_{c}\})) \right]$$
(19)

Thus, syntactic variable-binding has the semantics of multiple integration. This is the same result one would get if each rule with variables were replaced with a (finite, countable, or uncountably infinite) set of rules with all possible values substituted in for all the variables, with firing rates weighted by the relevant measure, and running in parallel.

[Annals of Math. and A. I., 47(3-4), January 2007]

• So, object parameters need measure spaces

#### Particle to Structure Dynamics

• Particle reactions/transitions, with params

 $A_1(x_1), A_2(x_2), \dots, A_n(x_n) \to B_1(y_1), B_2(y_2), \dots, B_m(y_m) \text{ with } \rho(\{x_i\}, \{y_j\})$ 

 $\tilde{O}_r = \rho_r \sum_{\{x'_i, x_j\}} \prod_{i \in \text{rhs}(r)} \hat{a}(\tau_i, x_i) \prod_{j \in \text{lhs}(r)} a(\tau_j, x_j) \Pr(\{x_i\} \mid \{x_j\})$ 

(and can integrate ODE rules too)



### Particle to Structure Dynamics

• Particle reactions/transitions, with params

 $A_1(x_1), A_2(x_2), \dots, A_n(x_n) \to B_1(y_1), B_2(y_2), \dots, B_m(y_m) \text{ with } \rho(\{x_i\}, \{y_j\})$ 

$$\tilde{O}_{r} = \rho_{r} \sum_{\{x'_{i}, x_{j}\}} \prod_{i \in \text{rhs}(r)} \hat{a}(\tau_{i}, x_{i}) \prod_{j \in \text{lhs}(r)} a(\tau_{j}, x_{j}) \Pr(\{x_{i}\} \mid \{x_{j}\})$$

(and can integrate ODE rules too)

$$\begin{bmatrix} a_{ab}, \hat{a}_{ad} \end{bmatrix} = \begin{bmatrix} a_{bb}, (a) \end{bmatrix}$$
  
 $\begin{bmatrix} a_{a}, a \end{bmatrix} = \begin{bmatrix} \hat{a}_{a}, \hat{a} \end{bmatrix} = 0$ 

^2 o

• Labelled graph (structure) transitions

$$\hat{W}_r \propto \int d\lambda d\lambda' \, \rho_r(\lambda,\lambda') \sum_{\langle i_1,\dots,i_k \rangle_{\neq}} \hat{a}_{i_1,\dots,i_k}(G^{r \text{ out}}) a_{i_1,\dots,i_k}(G^{r \text{ in}})$$

(and can integrate ODE rules too)

[EM, MFPS Proc., ENTCS 2010]

$$a_{\alpha} = 0 = a_{\alpha}$$
$$a_{\alpha} \hat{a}_{\beta} = (1 - \delta_{\alpha\beta}) \hat{a}_{\beta} a_{\alpha} + \delta_{\alpha\beta} Z_{\alpha}$$
$$Z_{\alpha} \equiv I_{\alpha} - N_{\alpha}$$
$$N_{\alpha} \equiv \hat{a}_{\alpha} a_{\alpha}$$

2

$$\begin{aligned} \hat{a}_{i_1,\dots,i_k}(G') &= \hat{a}_{i_1,\dots,i_k}(G'_{\text{links}})\hat{a}_{i_1,\dots,i_k}(G'_{\text{nodes}}) \\ &= \left[\prod_{s',t'\in\text{rhs}(r)} \left(\hat{a}_{i_{s'}i_{t'}}\right)^{g'_{s'}t'}\right] \left[\prod_{v'\in\text{rhs}(r)} \hat{a}_{i_{v'}\lambda'_{v'}}\right] \\ a_{i_1,\dots,i_k}(G) &= a_{i_1,\dots,i_k}(G_{\text{links}})a_{i_1,\dots,i_k}(G_{\text{nodes}}) \\ &= \left[\prod_{s,t\in\text{lhs}(r)} \left(a_{i_si_t}\right)^{g_s t}\right] \left[\prod_{v\in\text{lhs}(r)} a_{i_v\lambda_v}\right]. \end{aligned}$$

Ψ

## Algebra of Labelled-Graph Rewrite Rules

$$\hat{W}_{G^{r_2} \text{ in}_{\rightarrow} G^{r_2} \text{ out}} \hat{W}_{G^{r_1} \text{ in}_{\rightarrow} G^{r_1} \text{ out}} \simeq \sum_{\substack{H \subseteq G^{r_1} \text{ out}_{\rightarrow} \cong \tilde{H} \subseteq G^{r_2} \text{ in}_{\rightarrow} \tilde{H} \subseteq G^{r_2} \text{ in}_{\rightarrow} \tilde{H}} \sum_{\substack{h: H \stackrel{1-1}{\rightarrow} \tilde{H}}} \hat{W}_{G^{r_1} \text{ in}_{\rightarrow} (G^{r_2} \text{ in}_{\wedge} \tilde{H})_{h} G^{r_2} \text{ out}_{\rightarrow} (G^{r_1} \text{ out}_{\wedge} H) }$$

$$| \text{ edge-maximal}$$

$$G_{\text{nodes}}^{1;2 \text{ in}} = G_{\text{nodes}}^{r_1 \text{ in}} \dot{\cup} (G_{\text{nodes}}^{r_2 \text{ in}} \setminus \tilde{H}_{\text{nodes}}) \qquad G_{\text{nodes}}^{1;2 \text{ out}} = G_{\text{nodes}}^{r_2 \text{ out}} \dot{\cup} (G_{\text{nodes}}^{r_1 \text{ out}} \setminus H_{\text{nodes}}) G_{\text{links}}^{1;2 \text{ in}} = G_{\text{links}}^{r_1 \text{ in}} \cup h^{-1*} (G_{\text{links}}^{r_2 \text{ in}} \setminus \tilde{H}_{\text{links}}) \qquad G_{\text{links}}^{1;2 \text{ out}} = G_{\text{links}}^{r_2 \text{ out}} \cup h^* (G_{\text{links}}^{r_1 \text{ out}} \setminus H_{\text{links}})$$

$$K_a = G_{\text{nodes}}^{r_a \text{ in}} \cap G_{\text{nodes}}^{r_a \text{ out}}$$
$$K_{1;2} = (K_1 \setminus H_{\text{nodes}} \cup h^{-1}(K_2 \setminus \tilde{H}_{\text{nodes}}) \cup (K_1 \cap h^{-1\star}(K_2))$$

Verifying this result by machine would be both valuable, and a big challenge. It was the longest, trickiest calculation I've ever done by hand. Possibly we need an easier hand-verification first.

Ψ

[EM, http://arxiv.org/abs/1909.04118]

### Point of maximum intermediate expression swell



[EM, http://arxiv.org/abs/1909.04118]

## Algebra of Labelled-Graph Rewrite Rules: ODE case

**Operator for ODE rules:** 

$$\begin{split} W_{\text{ODE }2} &= \hat{W}_{\text{ODE }2} = \int d\mathbf{x} d\mathbf{y} \; \rho_2(\mathbf{y}, \mathbf{x}) \sum_{\langle i_1, \dots, i_k \rangle_{\neq}} \hat{a}_{i_1, \dots, i_k} (G^{(2)}(\mathbf{y})) a_{i_1, \dots, i_k} (G^{(2)}(\mathbf{x})) \; \text{, where} \\ \rho_2(\mathbf{y}, \mathbf{x}) &= -\operatorname{grad}_{\mathbf{y}} \cdot (\mathbf{v}(\mathbf{y}) \delta(\mathbf{y} - \mathbf{x})) \; = -\sum_a \operatorname{grad}_{y_a} (v_a(\mathbf{y}) \prod_b \delta(\mathbf{y} - \mathbf{x})) \; \text{.} \end{split}$$



Ψ

Commutator of ODE & ODE rules:

 $\mathbf{v}_{[2,1]}(\mathbf{x}) = (\mathbf{v}_1 \cdot \mathbf{grad}_{\mathbf{x}})\mathbf{v}_2(\mathbf{x}) - (\mathbf{v}_2 \cdot \mathbf{grad}_{\mathbf{x}})\mathbf{v}_1(\mathbf{x})$ 

$$\begin{aligned} \text{Commutator of GG \& ODE rules:} \\ [W_{\text{ODE }(2)}, \hat{W}_{\text{SPG }(1)}] &= \frac{1}{C_1(N_{\text{max free}})} \sum_{h:H \to \tilde{H}} \int d\mu_{(1)h}(X_{h\perp}) \% \\ &\int dz_{h\parallel} \int dx \int dy \quad \rho_2(\mathbf{y}, \mathbf{x}) \\ &\times \left\{ \rho_1(z_{h\parallel} = \mathbf{x}, X_{h\perp}) \hat{W}_{G^{1;2}} \inf_{(\tilde{H}, \mathbf{x}, z_{h\perp}) \to \tilde{h}} G^{1;2} \operatorname{out}_{(H, \mathbf{y}, z_{h\perp})} \right. \end{aligned}$$

## Synapse morphodynamics





# E.g. actin cytoskeleton in synapse

• Branching rule - one of a dozen or so



Plenum DGG simulations: Matthew Hur Earlier software: Guy Yosiphon and Arthur York
### Dendritic spine head DGG Rules



#### **Actin Network Biomechanics Rules**

- Stochastic Angle Bending
- Pairwise Lennard-Jones Force
- Tri-Nodal Anisotropic Buckling
- Hessian Boltzmann Sampling
- Euler-Bernoulli Deflection

#### **Spine Head Membrane Rules**

 Area, Line Tension, & Helfrich Bending Energy

### **Biomechanical Rules**

#### Implement dynamics for combined energy

$$E_{\text{tot}} = \sum_{ijk}^{N} \underbrace{U_{\text{tri-nodal}}(x_{i}, x_{j}, x_{k})}_{\text{Arisotropic Buckling}} + \underbrace{U_{H}(x_{i}, x_{j}, x_{k})}_{\text{Hessian Thermal Noise}} + \underbrace{U_{LJ-\text{midpoint}}(x_{1}, x_{2}, x_{3})}_{\text{Stochastic Midpoint Minimization}} + \underbrace{P\sum_{ij}^{M} \frac{x^{i}y^{j} - y^{i}x^{j}}{2}}_{\text{Membrane Area}} + \underbrace{\Omega\sum_{ijk}^{M} \frac{||x_{i} - x_{j}||_{2} + ||x_{k} - x_{j}||_{2}}{2}}_{\text{Membrane Length}} + \underbrace{2x\sum_{ijk}^{M} \frac{\partial H^{2}}{\partial x_{j}}(x_{h}, x_{h}, x_{h}, x_{h})}_{\text{Membrane Helfrich Curvature}}$$
  
where  

$$U_{H}(x_{1}, x_{2}, x_{3}) = \frac{1}{2}(x_{2} - x_{1})^{T} \frac{\partial^{2}U_{LJ}(x_{2}, x_{1})}{\partial x_{2}\partial x_{1}}(x_{2} - x_{1}) + \frac{1}{2}(x_{2} - x_{3})^{T} \frac{\partial^{2}U_{LJ}(x_{2}, x_{3})}{\partial x_{2}\partial x_{3}}(x_{2} - x_{3})$$

E.g. (pretty-printed rules) :

Hessian Thermal Noise:

 $(\bigcirc_{1} - \bigcirc_{2} - \bigcirc_{3}) \langle\!\langle (\boldsymbol{x}_{1}, \boldsymbol{\theta}_{1}), (\boldsymbol{x}_{2}, \boldsymbol{\theta}_{2}), (\boldsymbol{x}_{3}, \boldsymbol{\theta}_{3}) \rangle\!\rangle$   $\rightarrow (\bigcirc_{1} - \bigcirc_{2} - \bigcirc_{3}) \langle\!\langle (\boldsymbol{x}_{1}, \boldsymbol{\theta}_{1}), (\boldsymbol{x}_{2} + \Delta_{\mathrm{H}}, \boldsymbol{\theta}_{2}), (\boldsymbol{x}_{3}, \boldsymbol{\theta}_{3}) \rangle\!\rangle$ with  $k_{\mathrm{biomech}} p(U_{\mathrm{H}}(\boldsymbol{x}_{1}, \boldsymbol{x}_{2} + \Delta_{\mathrm{H}}, \boldsymbol{x}_{3}) - U_{\mathrm{H}}(\boldsymbol{x}_{1}, \boldsymbol{x}_{2}, \boldsymbol{x}_{3}))$  $\left\{ \begin{aligned} \Delta_{\mathrm{H}} \sim \mathcal{N}_{2}(0, \Sigma_{H}) \\ p(\Delta U(\boldsymbol{x}_{1}, \boldsymbol{x}_{2}, \boldsymbol{x}_{3})) = \frac{e^{-\frac{\Delta U(\boldsymbol{x}_{1}, \boldsymbol{x}_{2}, \boldsymbol{x}_{3})}{k_{B}T}}{1+e^{-\frac{\Delta U(\boldsymbol{x}_{1}, \boldsymbol{x}_{2}, \boldsymbol{x}_{3})}{k_{B}T}} \end{aligned} \right.$  LJ Anisotropic Buckling:

$$(\bigcirc_1 \longrightarrow \bigcirc_2 \longrightarrow \bigcirc_3) \langle\!\langle (\boldsymbol{x}_1, \boldsymbol{\theta}_1), (\boldsymbol{x}_2, \boldsymbol{\theta}_2), (\boldsymbol{x}_3, \boldsymbol{\theta}_3) \rangle\!\rangle$$
$$\longrightarrow (\bigcirc_1 \longrightarrow \bigcirc_2 \longrightarrow \bigcirc_3) \langle\!\langle (\boldsymbol{x}_1, \boldsymbol{\theta}_1), (\boldsymbol{x}_2 + \Delta_{\mathrm{LJ}}(\boldsymbol{x}_2), \boldsymbol{\theta}_2), (\boldsymbol{x}_3, \boldsymbol{\theta}_3) \rangle\!\rangle$$

with  $k_{\text{biomech}} p(U_{\text{tri-nodal}}(x_1, x_2 + \Delta_{\text{LJ}}(x_2), x_3) - U_{\text{tri-nodal}}(x_1, x_2, x_3))$ 

where 
$$\begin{cases} \Delta_{\rm LJ}(x_1, x_2, x_3) = {\rm Clip}(-\frac{1}{\zeta} \frac{\partial U_{\rm tri-nodal}(x_1, x_2, x_3)}{\partial x_2}) \\ p(\Delta U(x_1, x_2, x_3)) = \frac{e^{-\frac{\Delta U(x_1, x_2, x_3)}{k_B T}}}{1 + e^{-\frac{\Delta U(x_1, x_2, x_3)}{k_B T}}} \end{cases}$$

# Plenum/Mathematica implementation

#### E.g.: Rule for Hessian Thermal Noise Boltzmann Sampling of Random Displacements for a Chain of Three Actins:

{

actin[aIDP, coordsP, aIDPP, aID, angleP, CLP, arpIDP, endP, membraneIDP, nucIDP], actin[aID, coords, aIDP, aIDN, angle, CL, arpID, INT, membraneID, nucID], actin[aIDN, coordsN, aID, aIDNN, angleN, CLN, arpIDN, endN, membraneIDN, nucIDN] } → { (\*Move position of middle actin and update angle\*) actin[aID, addVectors[coords, {dx, dy}], aIDP, aIDN, middleThetaMovement[coordsP, addVectors[coords, {dx, dy}], coordsN], CL , arpID, INT, membraneID, nucID], (\*Update angle of first actin\*) actin[aIDP, coordsP, aIDPP, aID, If[endP = INT, angleP + startActinDeltaTheta[coordsP, coords, addVectors[coords, {dx, dy}]], 0], CLP, arpIDP, endP, membraneIDP, nucIDP], (\*Update angle of last actin\*) actin[aIDN, coordsN, aID, aIDNN, If[endN = INT, angleN + endActinDeltaTheta[coords, coordsN, addVectors[coords, {dx, dy}]], 0], CLN, arpIDN, endN, membraneIDN, nucIDN] }, with[networkFactor + heatBathHessian[coordsP, coords, l0, l0, {dx, dy}, step] × grammarPDF[NormalDistribution[0, transverseDeviationA/biomechanicalRate], dx] × grammarPDF[NormalDistribution[0, transverseDeviationA/biomechanicalRate], dy]],

## **2D Actin Network**

#### Network Remodeling Rules Reach Non-Zero Steady-State



- 0.06 fraction of actin are monomers at steady-state. Addition of membrane would increase this value making it consistent with 0.12 reported in literature
- Average number of actins per branch is 7 in the simulation while it is 8 in the literature

Plenum DGG simulations: Matthew Hur Earlier software: Guy Yosiphon and Arthur York

### 2D Actin Network with Membrane



Plenum DGG simulations: Matthew Hur Earlier software: Guy Yosiphon and Arthur York

#### MT fiber Stochastic Parametrized Graph Grammar



[EM, Bull. Math Biol. 81:8 Aug 2019 +arXiv:1804.11044]

# Large-scale simulation

#### Dynamical graph grammar

...

$$\begin{pmatrix} \bigcirc_1 \longrightarrow \bigcirc_2 \longrightarrow \bigcirc_3 \\ \bullet_4 \end{pmatrix} \langle \langle (x_1, u_1), (x_2, u_2), (x_3, u_3), (x_4, u_4) \rangle \rangle \\ \longrightarrow \begin{pmatrix} \bigcirc_1 \longrightarrow \bullet_2 \longrightarrow \bigcirc_3 \\ \bigcirc_4 \end{pmatrix} \langle \langle (x_1, u_1), (x_2, u_2), (x_3, u_3), (x_4, u_4) \rangle \rangle \\ \text{with } \hat{\rho}_{\text{bundle}}(|u_2 \cdot u_4|/|\cos \theta_{\text{crit}}|) \exp\left(-|x_2 - x_4|^2/2L^2\right)$$

$$(\bigcirc_1 \longrightarrow \bullet_2) \langle\!\langle (\boldsymbol{x}_1, \boldsymbol{u}_1)(\boldsymbol{x}_2, \boldsymbol{u}_2) \rangle\!\rangle$$
$$\longrightarrow (\bigcirc_1 \longrightarrow \bullet_2) \langle\!\langle (\boldsymbol{x}_1, \boldsymbol{u}_1), (\boldsymbol{x}_2 + d\boldsymbol{x}_2, \boldsymbol{u}_2) \rangle\!\rangle$$
solving  $d\boldsymbol{x}_2/dt = \hat{\rho}_{\text{grow}}([Y_{\text{g}}])(1 - L/L_{\text{div}})\boldsymbol{u}_2$ 





To infer:  $L, L_{div}, \theta_{crit}$ , params for  $\rho_{bundle}, \rho_{grow}$ 

[Medwedeff and Mjolsness, Physical Biology, June 2023]

### Parallel DGG algorithm: Simulated MT bundling, catastrophe





Figure 5.15: Randomly Selected 5x5 Zoomed of 5,000 MTs, T = 400

[Medwedeff and Mjolsness, Physical Biology, June 2023]

### Cell Complex Operator Splitting

$$\begin{split} e^{tW} &\approx \left(\prod_{d\downarrow} e^{\frac{t}{n}W_{(d)}}\right)^{n \to \infty} \\ e^{t'W_{(d)}} &= \prod_{c \ c \ d} e^{t'W_{(c,d)}} \quad \text{where} \quad \left[W_{(c,d)}, W_{(c',d)}\right] &\approx 0 \quad \text{and} \quad t' \equiv \frac{t}{n} \\ W_{(c,d)} &= \sum_{r} W_{r,c} \equiv \sum_{r} \sum_{\substack{r \ R \ | \ \varphi(R) = c, \\ R \ \text{instantiates} \ r}} W_{r}(R \mid c, d) \end{split}$$

Requires different c of the same d be well-separated.

[Medwedeff and Mjolsness, Physical Biology, June 2023]

#### Expanded Cell Complex, for separation within dimension



# MT MD model reduction



#### MaxEnt Problem



$$S = \int_{0}^{\infty} dt \ \mathcal{D}_{\mathcal{KL}}(\boldsymbol{p}||\boldsymbol{\tilde{p}})$$
$$w/ \ \mathcal{D}_{\mathcal{KL}}(\boldsymbol{p}||\boldsymbol{\tilde{p}}) = \sum_{\boldsymbol{n=0}}^{\infty} \int d\boldsymbol{x} \ \boldsymbol{p} \ln \frac{\boldsymbol{p}}{\boldsymbol{\tilde{p}}}$$
$$\tilde{p}(n, \boldsymbol{x}, \boldsymbol{\alpha}, t) = \frac{1}{Z} \exp \left[ -\sum_{k=1}^{K} \sum_{\langle j \rangle} \nu_k(\boldsymbol{x}_{\langle j \rangle}, \boldsymbol{\alpha}_{\langle j \rangle}, t) \right],$$

Variational problem

$$\frac{\delta S}{\delta F_k[\{\nu_k(\boldsymbol{x})\}_{k=1}^K]} = 0 \text{ for } k = 1, \dots, K \text{ at all } \boldsymbol{x}$$
(12)

where the variation is with respect to a set of functionals

$$\dot{\nu}_k(\mathbf{x}) = F_k[\{\nu_k\}_{k=1}^K]$$
 (13)

... Higher-order calculus!

$$\frac{\delta S}{\delta F_{k}[\boldsymbol{\nu}(\boldsymbol{x})]} = \sum_{k'=1}^{K} \int d\boldsymbol{x}' \int dt \, \frac{\delta S}{\delta \nu_{k'}(\boldsymbol{x}', t)} \frac{\delta \nu_{k'}(\boldsymbol{x}', t)}{\delta F_{k}[\boldsymbol{\nu}(\boldsymbol{x})]} = 0 \quad (19)$$

$$\underbrace{1}_{\substack{\delta S \\ \delta \nu_{k'}(\boldsymbol{x}', t)}} = \left\langle \sum_{\langle i \rangle_{k'}^{n}} \delta(\boldsymbol{x}' - \boldsymbol{x}_{\langle i \rangle_{k'}^{n}}) \right\rangle_{p} - \left\langle \sum_{\langle i \rangle_{k'}^{n}} \delta(\boldsymbol{x}' - \boldsymbol{x}_{\langle i \rangle_{k'}^{n}}) \right\rangle_{\tilde{p}} \quad (20)$$
e.g.  $k' = 1 : \left\langle \sum_{i=1}^{n} \delta(x_{i} - x') \right\rangle$  for all  $x'$   
 $k' = 2 : \left\langle \sum_{i=1}^{n} \sum_{j > i} \delta(x_{i} - x'_{1}) \delta(x_{j} - x'_{2}) \right\rangle$  for all  $x'_{1}, x'_{2}$ 

Need to choose a parametrization for functional!



#### Computational problem





### Adjoint method BMLA-like learning algorithm

Algorithm 1 Stochastic Gradient Descent for Learning Restricted Boltzmann Machine Dynamics

#### 1: Initialize

- 2: Parameters  $\boldsymbol{u}_k$  controlling the functions  $F_k(\boldsymbol{\theta}; \boldsymbol{u}_k)$  for all  $k = 1, \ldots, K$ .
- Time interval  $[t_0, t_f]$ , a formula for the learning rate  $\lambda$ . 3:
- 4: while not converged do
- Initialize  $\Delta F_{k,i} = 0$  for all  $k = 1, \ldots, K$  and parameters  $i = 1, \ldots, M_k$ . 5:
- for sample in batch do 6:
- $\triangleright$  Generate trajectory in reduced space  $\boldsymbol{\theta}$ : 7:
- Solve the PDE constraint (27) for  $\theta_k(t)$  with a given IC  $\theta_{k,0}$  over  $t_0 \le t \le t_f$ , for all k. Wake phase: 8:
- 9:
- Evaluate moments  $\mu_k(t)$  of the data for all k, t. 10:
- $\triangleright$  Sleep phase: 11:
- 12:
- Evaluate moments  $\tilde{\mu}_k(t)$  of the Boltzmann distribution.  $\Rightarrow$  Solve the adjoint system: Solve the adjoint system (31) for  $\phi_k(t)$  for all k, t.  $\Rightarrow \frac{d}{dt}\phi_k(t) = \tilde{\mu}_k(t) \mu_k(t) \sum_{k=1}^K \frac{\partial F_l(\boldsymbol{\theta}(t); \boldsymbol{u}_l)}{\partial \theta_k(t)} \phi_l(t),$ 13:14:
- ▷ Evaluate the objective function: 15:
- Update  $\Delta F_{k,i}$  as the cumulative moving average of the sensitivity equation (30) over the batch. 16:
- $\frac{dS}{du_{k,i}} \stackrel{\clubsuit}{=} -\int_{t}^{t_{f}} dt \; \frac{\partial F_{k}(\boldsymbol{\theta}(t);\boldsymbol{u}_{k})}{\partial u_{k,i}} \phi_{k}(t)$ ▷ Update to decrease objective function: 17: $u_{k,i} \to u_{k,i} - \lambda \Delta F_{k,i}$  for all k, i. 18:



O. Ernst, T. Bartol, T. Sejnowksi, and E. Mjolsness, J of Chem Phys 149, 034107, July 2018. Also arXiv 1803.01063

### """ "Tchicoma" Architecture for Mathematical Modeling

To formalize:

Vathematics

+ mappings and hierarchies:



+arXiv:1804.11044]

### "Eclectic Algebraic Type Theory" for mathematical type hierarchy



# **Eclectic Types**

**Context:** The previous bio modeling languages shown above have a well-developed semantics of scientific "processes" modeled mathematically, but not for extended science-modeling "objects" beyond parameter-labelled graphs (which however covers a lot). E.g. no manifolds, dynamic function spaces, etc.. Such semantics could also undergird machine-assisted proof of some of the "arrows" in the Tchicoma architecture, e.g. algorithm-generation. The hope was that type theory could help with all this. So far no great luck.

#### Goals:

 Generalize (loosen) the idea of a single "universal" Cartesian Closed Category as the semantics of a type theory of mathematical objects, to allow many base categories in a progressive curriculum, or none, while licensing a rich collection of type constructor notations to the extent possible. So, "eclectic" about foundations and expressive notations.
 Define larger units of organization in a "type hierarchy", to allow for looser coordination in expert development.

Caveat: This is all in the service of automated tools for scientific modeling, so might be (even?) worse for other areas of mathematics.

# **Eclectic Types**

**Definition.** An *eclectic type t* is a triple  $(C_t, \tau_t, L_t)$ , subject to certain constraints, where  $C_t$  is a content category whose category-objects are the mathematical objects of eclectic type t,  $\tau_t$  is a type object in a logical type category A, and  $L_t$  is a language equivalent to the minimal subset of the internal language of A that includes  $\tau_t$  and all its predecessors in a partial ordering < on the objects of A. The constraints on  $C_t$  and  $\tau_t$  are:

(1) finite product types  $\tau_t = \prod_{p} \tau_{\sigma_p}$  in A: [technical condition on  $C_t$ ], and likewise for sums; & (2) function types in A,  $\tau_f = \tau_x \to \tau_t$ , [technical condition on  $C_f$ ].

The partial ordering relation < is related to definability <# within a type hierarchy as follows:

$$< \# \implies <$$
 i.e.  $\forall a, b (a < \# b \implies a < b$ 

This way, required definitional statements can be made in the available sub-language  $L_t$  of each type.

**Definition**. A *subtype*  $(C_s, \tau_s, L_s)$  of an eclectic type  $(C_t, \tau_t, L_t)$  is an eclectic type that has both a [technical condition on  $C_s$  and  $C_t$ ] and a subtyping relationship  $\tau_s <: \tau_t$ .

By the Liskov substitution principle, subtype relationships <: imply further such relationships among product types, sum types, and function types ("contravariantly" among the function argument types). This relationship contributes to generativity of the language fragment connected with an eclectic type in a type hierarchy. Likewise, the <# and hence < partial order relations must be consistent with product, sum, and function type constructors.

**Definition**. Assuming these kinds of consistency relations among <:, <#, and <, then we have a *type hierarchy*.

# **Eclectic Types**

**Definition**. In general, <# and < are related not only by type constructor consistency, but also by refinement:  $<# \implies <$ . However, <: need not be related to < by refinement. When it is, the preferred relation has the opposite sense:

 $(:> \implies <\# \implies <)$  i.e.  $\forall a, b \ (a :> b \implies a <\# b \implies a < b)$ . A subset of eclectic types, within a type hierarchy, that satisfies this relationship is called a *type hierarchy module* or where unambiguous a "*type module*".

**Definition**. A *curriculum* is a type hierarchy together with a set-covering collection of type modules within the type hierarchy, that form a DAG and hence a partial order, under the quotient new-generalization relationship: (<# and <:)/modules .

Spiral development is a special case of a curriculum.

A type module is easily extensible, but only by further definition (old <# new) and specialization (old :> new) which can of course be applied in combinatorial ways. Type hierarchy modules could be fruitful for symbolic AI search algorithms. When in the course of mathematical development new generalization supertypes are required, thus exceeding the limitations of a type module, the options are

(1) to begin a new type module within an existing type hierarchy, preserving existing <# definitional type relationships while potentially also creating a "curriculum" of modules of increasing generalization content, or

(2) to map the old type hierarchy into a new and more encompassing one, thus "restructuring" it.

In this way three levels and timescales of mathematical type-elaboration activity naturally emerge: incremental exploitation within modules, development of new modules, and foundational restructuring.

### AITP for Science: some logic in favor

- "AI": Reliable AI for Science:
  - Representationalism 1st, ML 2nd. (On top vs. on tap)
  - formal, symbolic DSLs for modeling ...
    - applicable math, sci content, algorithms (model sim, ML, analysis)
- "TP": verify exact and approximate maps between such languages
  - e.g. dynamics to simulation algs: XDEs, Markov jump processes,
  - high-level support for scientific knowledge representation

# Eg: Plant gene expression model Declarative, with cell growth & division

$$\begin{split} &\{ \emptyset \rightarrow U, k_1 TIP[t] \}, \{ U \rightarrow \emptyset, k_2 \}, \{ U \longrightarrow U, \text{Diffusion}[D_U] \}, \\ &\{ \emptyset \rightarrow V, k_3 L1[t] \}, \{ V \rightarrow \emptyset, k_4 \}, \{ V \longrightarrow V, \text{Diffusion}[D_V] \}, \\ &\{ \emptyset \rightleftharpoons Z, k_7, k_8 U[t] \}, \{ X \mapsto V, \text{GRN}[v_V, T_{WV}, 1, h_V] \}, \\ &\{ \{ U, V, W \} \mapsto W, \text{GRN}[v_W, \{ T_{UW}, T_{VW}, T_{WW} \}, 1, h_W] \}, \{ W \rightarrow \emptyset, k_6 Z[t] + k_9 L2[t] \} \\ &\{ W \mapsto X, \text{GRN}[v_X, T_{WX}, 1, h_X] \}, \{ X \rightarrow \emptyset, k_5 \}, \{ X \longrightarrow X, \text{Diffusion}[D_X] \}, \\ &\{ \text{cell} \longrightarrow \text{cell}, \text{Grow}[\text{GrowthRate}[\mu, f_\mu], \text{Pressure}[P, f_P], \text{Spring}[k, f_k] \}, \\ &\{ \text{cell} \longrightarrow \text{cell} + \text{cell}, \text{Errera}[\text{cell}, \mu, \sigma \} \} \end{split}$$



[Shapiro et al Frontiers in Plant Science 2013]

### Abstract

The complexity of biological systems (among others) makes demands on the complexity of the mathematical modeling enterprise that could be satisfied with mathematical artificial intelligence of both symbolic and numerical flavors. Technologies that I think will be fruitful in this regard include:

(1) the use of machine learning to bridge spatiotemporal scales, which I will illustrate with the "Dynamic Boltzmann Distribution" method for learning model reduction of stochastic spatial biochemical networks and the "Graph Prolongation Convolutional Network" approach to coursegraining the biophysics of microtubules;

(2) a meta-language for stochastic spatial graph dynamics, "Dynamical Graph Grammars", that can represent structure-changing processes including microtubule dynamics and that has an underlying combinatorial theory related to operator algebras; and

(3) an integrative conceptual architecture of typed symbolic modeling languages and structurepreserving maps between them, including model reduction and implementation maps. Machine learning for model reduction incorporating stat mech knowledge:

# *Dynamic Boltzmann Distributions* for stochastic reaction-diffusion systems



Oliver Ernst, Salk & UCSD

#### $\mathscr{R}$

### Model Reduction

- For ...
  - understandability
  - computational escalation
- Commutation:



- To reduce within the paradigm, we need ...
  - stochastic + deterministic dynamics
  - dynamic particles, fields, and graphs



### Mapping: Model reduction



[Johnson, Bartol, Sejnowski, and Mjolsness. Physical Biology 12:4, July 2015]  $\frac{dp}{dt} = W \cdot p$ 

 $\Psi \mathscr{R} \simeq \mathscr{R} \Psi$ 

- Nonspatial:  $p(R,t) = \exp\left[-\sum_{\alpha} \mu_{\alpha}(t)V_{\alpha}(R)\right]/\hat{Z}(\mu(t))$
- [Johnson, Bartol, Sejnowski, and Mjolsness. Physical Biology 12:4, July 2015]
- -Graph-Constrained Correlation Dynamics
- -warmup case for ...
- Spatial generalization:

$$\tilde{p}(n, \boldsymbol{x}, \boldsymbol{\alpha}, t) = \frac{1}{Z} \exp \left[ -\sum_{k=1}^{K} \sum_{\langle j \rangle} \nu_k(\boldsymbol{x}_{\langle j \rangle}, \boldsymbol{\alpha}_{\langle j \rangle}, t) \right],$$

-Dynamic Boltzmann distributions

Stat mech knowledge context for ML: Master eq. at fine scale Dynamic Boltzmann distributions at coarse scale

#### MaxEnt Problem



$$S = \int_{0}^{\infty} dt \ \mathcal{D}_{\mathcal{KL}}(\boldsymbol{p}||\boldsymbol{\tilde{p}})$$
$$w/ \ \mathcal{D}_{\mathcal{KL}}(\boldsymbol{p}||\boldsymbol{\tilde{p}}) = \sum_{\boldsymbol{n=0}}^{\infty} \int d\boldsymbol{x} \ \boldsymbol{p} \ln \frac{\boldsymbol{p}}{\boldsymbol{\tilde{p}}}$$
$$\tilde{p}(n, \boldsymbol{x}, \boldsymbol{\alpha}, t) = \frac{1}{Z} \exp \left[ -\sum_{k=1}^{K} \sum_{\langle j \rangle} \nu_k(\boldsymbol{x}_{\langle j \rangle}, \boldsymbol{\alpha}_{\langle j \rangle}, t) \right],$$

Variational problem

$$\frac{\delta S}{\delta F_k[\{\nu_k(\boldsymbol{x})\}_{k=1}^K]} = 0 \text{ for } k = 1, \dots, K \text{ at all } \boldsymbol{x}$$
(12)

where the variation is with respect to a set of functionals

$$\dot{\nu}_k(\mathbf{x}) = F_k[\{\nu_k\}_{k=1}^K]$$
 (13)

... Higher-order calculus!

$$\frac{\delta S}{\delta F_{k}[\boldsymbol{\nu}(\boldsymbol{x})]} = \sum_{k'=1}^{K} \int d\boldsymbol{x}' \int dt \, \frac{\delta S}{\delta \nu_{k'}(\boldsymbol{x}', t)} \frac{\delta \nu_{k'}(\boldsymbol{x}', t)}{\delta F_{k}[\boldsymbol{\nu}(\boldsymbol{x})]} = 0 \quad (19)$$

$$\underbrace{1}_{\substack{\delta S \\ \delta \nu_{k'}(\boldsymbol{x}', t)}} = \left\langle \sum_{\langle i \rangle_{k'}^{n}} \delta(\boldsymbol{x}' - \boldsymbol{x}_{\langle i \rangle_{k'}^{n}}) \right\rangle_{p} - \left\langle \sum_{\langle i \rangle_{k'}^{n}} \delta(\boldsymbol{x}' - \boldsymbol{x}_{\langle i \rangle_{k'}^{n}}) \right\rangle_{\tilde{p}} \quad (20)$$
e.g.  $k' = 1 : \left\langle \sum_{i=1}^{n} \delta(x_{i} - x') \right\rangle$  for all  $x'$   
 $k' = 2 : \left\langle \sum_{i=1}^{n} \sum_{j > i} \delta(x_{i} - x'_{1}) \delta(x_{j} - x'_{2}) \right\rangle$  for all  $x'_{1}, x'_{2}$ 

Need to choose a parametrization for functional!



#### Computational problem


## Linearity of process operators

### Proposition

Fix: reaction network  $\{\mathbf{W}_r\}$  and graphical model  $\nu$ Then: linearity of CME in reaction operators:  $\dot{\mathbf{p}} = \sum_r \mathbf{W}^{(r)} \mathbf{p}$ extends to the reduced model:  $\tilde{F}_{1\alpha} = \sum_r \tilde{F}_{1\alpha}^{(r)}$  and  $\tilde{F}_{2\alpha\beta} = \sum_r \tilde{F}_{2\alpha\beta}^{(r)}$ 

• Reason: Chain rule on MaxEnt optimal inverse statement



O. Ernst, T. Bartol, T. Sejnowksi, and E. Mjolsness, J of Chem Phys 149, 034107, July 2018. Also arXiv 1803.01063



## Adjoint method BMLA-like learning algorithm

Algorithm 1 Stochastic Gradient Descent for Learning Restricted Boltzmann Machine Dynamics

#### 1: Initialize

- 2: Parameters  $\boldsymbol{u}_k$  controlling the functions  $F_k(\boldsymbol{\theta}; \boldsymbol{u}_k)$  for all  $k = 1, \ldots, K$ .
- Time interval  $[t_0, t_f]$ , a formula for the learning rate  $\lambda$ . 3:
- 4: while not converged do
- Initialize  $\Delta F_{k,i} = 0$  for all  $k = 1, \ldots, K$  and parameters  $i = 1, \ldots, M_k$ . 5:
- for sample in batch do 6:
- $\triangleright$  Generate trajectory in reduced space  $\boldsymbol{\theta}$ : 7:
- Solve the PDE constraint (27) for  $\theta_k(t)$  with a given IC  $\theta_{k,0}$  over  $t_0 \le t \le t_f$ , for all k. Wake phase: 8:
- 9:
- Evaluate moments  $\mu_k(t)$  of the data for all k, t. 10:
- $\triangleright$  Sleep phase: 11:
- 12:
- Evaluate moments  $\tilde{\mu}_k(t)$  of the Boltzmann distribution.  $\triangleright$  Solve the adjoint system: Solve the adjoint system (31) for  $\phi_k(t)$  for all k, t.  $\Rightarrow \frac{d}{dt}\phi_k(t) = \tilde{\mu}_k(t) \mu_k(t) \sum_{l=1}^K \frac{\partial F_l(\boldsymbol{\theta}(t); \boldsymbol{u}_l)}{\partial \theta_k(t)} \phi_l(t),$ 13:14:
- ▷ Evaluate the objective function: 15:
- Update  $\Delta F_{k,i}$  as the cumulative moving average of the sensitivity equation (30) over the batch. 16:
- $\frac{dS}{du_{k,i}} \stackrel{\clubsuit}{=} -\int_{t}^{t_{f}} dt \; \frac{\partial F_{k}(\boldsymbol{\theta}(t);\boldsymbol{u}_{k})}{\partial u_{k,i}} \phi_{k}(t)$ ▷ Update to decrease objective function: 17: $u_{k,i} \to u_{k,i} - \lambda \Delta F_{k,i}$  for all k, i. 18:

# Benefit of Hidden Units

### Network: fratricide + lattice diffusion



# Benefit of Hidden Units

Network: fratricide + lattice diffusion

• Learned DBD ODE RHS, without and with hidden units



FIG. 2. Top row: Learned time-evolution functions for the fully visible model (38), using the  $Q_3$ ,  $C_1$  finite-element parametrization (34) with cells of size  $0.5 \times 0.5 \times 0.5$  in (b, J, K). Left panel: Training set of initial points (b, J, K) (cyan) sampled evenly in [-1, 1]. Stochastic simulations for each initial point are used as training data (learned trajectories shown in black, endpoints in magenta). Middle three panels: The time evolution functions learned, where the heat map indicates the value of  $F_{\gamma}$  in (38). Right panel: Vertices of the finite-element cells used. Bottom row: Hidden layer model (40) and parametrization (34) with cells of size  $0.5 \times 0.5 \times 0.05$  in (b, W, b'). Initial points are generated by BM learning applied to the points of the visible model. Note that the coefficients corresponding to the other seven degrees of freedom at each vertex are also learned (not shown), i.e., the first derivatives in each parameter.

[Ernst, Bartol, Sejnowski, Mjolsness, Phys Rev E 99 063315, 2019]

# **Rössler Oscillator in 3D**

• Function: • Learned DBD ODE RHS:



[Ernst, Bartol, Sejnowski, Mjolsness, arXiv:1808.08630 v2 April 2019]

### Graph Prolongation Convolutional Networks for biomechanical model reduction

from molecular dynamics



Cory Scott

### **Graph Lineage Definitions**

- *Hierarchical Graph Sequence:* a mapping from  $\mathbb{N}$  into some sequence of graphs which obeys the following:
  - G<sub>0</sub> is the graph with one vertex and one loop on that vertex
  - Edge and vertex cardinality of graphs in the sequence grow at most "exponentially" in some base, b:  $O(b^{l^{1+\epsilon}})$



• *Graded Graph:* G is a graded graph if all of the vertices of G are labeled with nonnegative integers such that if  $(v_1, v_2)$  is an edge, the labels of  $v_1$  and  $v_2$  differ by at most 1.



- *Graph Lineage:* a graded graph where the sequence of  $\Delta L = 0$  subgraphs is a HGS and the subgraphs with  $\Delta L = 1$  are a HGS of bipartite graphs. The above is a graph lineage of path graphs of length  $2^n$ .
- *Hierarchitecture*: A graph lineage, used as a model architecture.

# **Generating Graph Lineages**

• One way to generate a graph lineage (or more generally, graded graphs) is via local graph rewrite rules.



• Rules can be applied locally, or to all cells in a graph simultaneously:



### **Local Firing**

**Global Firing** 

- Graph labels suppressed, but necessary
- More: (i + j) = (i + j



## Multiscale numerics: Alg. Multigrid Methods for Graphs



E.g. in optimization-based neural networks. [EM et al., IEEE Trans. Neural Networks 1991] Cf.  $MG' \simeq GM$  for graph-matching [Gold et al. Neural Computation 8 1996]

Here: use graph Laplacian  $L' \simeq P^T L P$ , or  $PL' \simeq L P$  with orthogonal  $P^T P = I$ 

#### **UCI** Morphodynamics

## Define Graph Process Directed "Distances"

• Definition requires constrained opt of diffusion operator:

 $D(G_1, G_2 | R, \alpha > 0, t) = \inf_{P | C(P)} || P \exp(\alpha^{-1/2} t W_1^{(R)}) - \exp(\alpha^{1/2} t W_2^{(R)}) P ||_F$  $D(G_1, G_2 | R, t) = \inf_{\alpha > 0} D(G_1, G_2 | R, \alpha, t)$ 

• Constraints: orthogonality; sparsity?

 $C(P): \qquad P^T P = I \qquad ; \qquad \max \text{ fanout}(P) \le (n_{P \text{fine}}/n_{P \text{course}})^s$ restriction.prolongation

• Optimize time & time dilation due to graph size:

 $\tilde{D}(G_1, G_2|R) = \sup_{t>0} \inf_{\alpha>0} D(G_1, G_2|R, \alpha, t)$ 

• Can obtain *P* at early times ("rigid" vs "flexible" def of *D*):

$$D_{\text{rigid}}(G_1, G_2|R, t) = \inf_{P|C(P)} ||P^* \exp(\alpha^{*-1/2} t W_1^{(R)}) - \exp(\alpha^{*1/2} t W_2^{(R)}) P^*||_F, \text{ where}$$
  
$$(\alpha^*, P^*) = \operatorname{argmin}_{\alpha > 0, P|C(P)} ||\alpha^{-1/2} P W_1^{(R)} - \alpha^{1/2} W_2^{(R)} P||_F$$

•  $\triangle \leq$  provable with weaker  $\alpha$ :  $\alpha = \left(\frac{n_1}{n_2}\right)^r$ 

# MT MD model reduction

