Coarse-Grained Brownian Dynamics Simulation of Rule-Based Models

Michael Klann¹, Loïc Paulevé¹, Tatjana Petrov¹, Heinz Koeppl^{1,2}

¹ ETH zürich ² IBM Research

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Motivation

Aim: impact of complexes (rings) stability for signalling cascades, etc.

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Rule-based modelling

- Cope with the combinatorics of potential complexes.
- Compact and non-redundant specification of reactions.

Spatial simulation

- Decompose diffusion (Brownian motion) and reaction.
- Finer expression of reaction probabilities.
- Impact of non-homogeneity.

Coarse grain

- Approximated stochastic simulation.
- Simple parameterization.

Contribution: combine the three ingredients, in a comprehensive manner.

Overview

Use of ruled-based paradigm

- Never expand the rules: directly look for and apply rule embeddings.
- Enforce *atomic* rules: only one single modification at a time, arity ≤ 2 .

Reaction probabilities refinement

- Distinguish bindings occurring within a complex vs between complexes.
- Emerging behaviour: rings become very stable.

Approximations

- Spherical approximation of complexes; ignore rotation, etc.
- Discrete-time simulation (approximation of Poisson distribution).

Proof of concept

- Initiation of the yeast pheromone response signalling cascade.
- Impact of the binding rates rules for the stability of the Ste5 scafold.

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Rule-based modelling Syntax



- Agents: e.g., A, B.
- Sites: e.g., A.x, A.b, A.c, B.a, B.c, B.y
- A site can have values: e.g., p, u.
- A site can be bound to another site of a different agent instance.

Rule-based modelling Rule application



Rule-based modelling Rule application



Rule-based modelling Rule application



Coarse-graining complexes



Spherical approximation

- No orientation, no rotation (considered as occurring on a much faster scale).
- Radius of agent a: ra
- Radius of complex C_i : rad $(C_i) = \left(\sum_{v \in C_i} r_{type(v)}^3\right)^{1/3}$

Brownian diffusion

Each complex instance is tracked individually (particle-based simulation).

Discrete time continuous space random walk for diffusion

$$\mathbf{x}_{\mathbf{i}}(t + \Delta t) = \mathbf{x}_{\mathbf{i}}(t) + \boldsymbol{\xi}\sqrt{2D_{i}\Delta t}$$

- $x_i(t)$: position of complex *i* at time *t*.
- **ξ**: standard normal random variable.
- D_i: diffusion coefficient of complex *i* (derived using Stokes-Einstein equation)

$$D_i = \frac{D_0 r_0}{\operatorname{rad}(C_i)}$$

Rules probability Modifcation / unbinding



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Rules probability

Binding within/between complexes

One rule, two different instanciantions.

Binding between two complexes



$$\Rightarrow \operatorname{prob}(\{C_1, C_2\}, \operatorname{rule}, \Delta t) \approx \frac{3}{4\pi (\operatorname{rad}(C_1) + \operatorname{rad}(C_2))^3} k\Delta t$$

Only when collision!

Binding within a complex



$$\Rightarrow \operatorname{prob}(\{C_1\}, \operatorname{rule}, \Delta t) \approx \frac{3}{4\pi (r_A + r_B)^3} k\Delta t$$

Always active!

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Stochastic abstract machine

Machine term

$$\begin{aligned} M &::= (t, C, X, \mathcal{R}) & \text{Time } t, \text{ complex maps } C \text{ and } X, \text{ rules } \mathcal{R} \\ C &::= \{i_1 \mapsto \mathcal{G}_1, \cdots, i_N \mapsto \mathcal{G}_N\} & \text{Map from a complex } i \text{ to its site-graph } \mathcal{G} \\ X &::= \{i_1 \mapsto \mathbf{x}_1, \cdots, i_N \mapsto \mathbf{x}_N\} & \text{Map from a complex } i \text{ to its position } \mathbf{x} \in \mathbb{R}^3 \end{aligned}$$

 i_1, \cdots, i_N are the complex identifiers, assumed all distinct.

Execution

$$\begin{array}{l} X' &= \text{diffuse}(C, X, \Delta t) \\ (C', X'') &= \text{react}(C, X', \mathcal{R}, \Delta t) \\ \hline (t, C, X, \mathcal{R}) \rightarrow (t + \Delta t, C', X'', \mathcal{R}) \end{array}$$

diffuse: Brownian diffusion

react: apply unary reactions and binary reactions restricted to collisions. Probability scaled by Δt .

Movie



Application: Ste5 scafold



Initiation of the Ste5/MAPK signalling cascade















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Application: Ste5 scafold Scafold stability



Coarse-Grained Brownian Dynamics Simulation of Rule-Based Models

Application: Ste5 scafold Impact on concentration gradients





Conclusion

Stochastic simulation framework combining three ingredients:

- Rule-based modelling: scale with the species combinatorics.
- Particle-based simulation: decompose diffusion and reaction.
- Approximate simulation: discrete time step.

Coarse-grained Brownian dynamics

- Spherical approximation of complexes.
- Spatial dimension: take into account concentration gradients
- Proof-of-concept: influence of weak bonds on scafold stability.

Further work

- More modulation of rates (paper)
- Constrained diffusion (membrane-bound complexes, etc.) (appendix)
- Geometric constraints (related work)

Thank you for your attention.

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