



# Reaction Network Modeling

Molecular Programming tutorial  
ASE 2015

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A **programming language** for molecular programming in a well-mixed solution.

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Essentially equivalent to

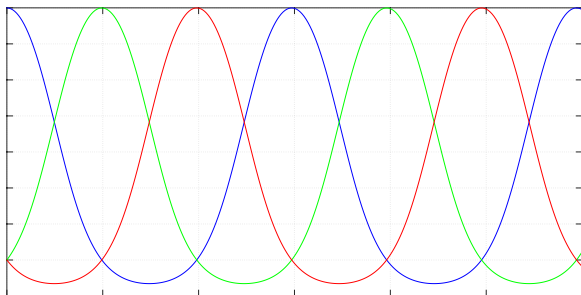
- Population protocols

- Petri nets

- Vector addition systems

# Chemical Reaction Networks

**Example:** A three-phase oscillator ( $\approx$  Lotka-Volterra)



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2. Mathematical (20th century): Use algebra, analysis, and probability to analyze structure and dynamics of CRNs.



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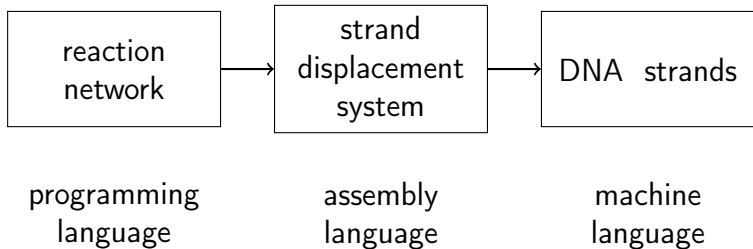
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## 4. Systematic prescriptive (21st century):

Replace (c) with **uniform translation** into DNA strand displacement systems.

Soloveichik, Seelig, and Winfree 2010



Chen, Dalchau, Srinivas, Phillips, Cardelli, Soloveichik, and Seelig 2013.

“Real programmers code in chemistry!”

Soloveichik, 2011



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$$\rho = (\mathbf{r}, \mathbf{p}, k) \in \mathbb{N}^S \times \mathbb{N}^S \times (0, \infty)$$

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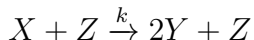
$\mathbf{r} = \mathbf{r}(\rho) =$  reactant vector of  $\rho$

$\mathbf{p} = \mathbf{p}(\rho) =$  product vector of  $\rho$

$k = k(\rho) =$  rate constant of  $\rho$

# The CRN Model

**Example:** Let  $S = \{X, Y, Z\}$ . The intuitive notation



refers to the formal reaction  $(\mathbf{r}, \mathbf{p}, k)$ , where

$$\begin{aligned}\mathbf{r}(X) &= \mathbf{r}(Z) = 1, & \mathbf{r}(Y) &= 0, \\ \mathbf{p}(X) &= 0, & \mathbf{p}(Y) &= 2, & \mathbf{p}(Z) &= 1.\end{aligned}$$

# The CRN Model

The two most widely used semantics (operational meanings) ascribed to a CRN  $N = (S, R)$ .

1. Deterministic mass action

A **state** is a vector  $\mathbf{x} \in [0, \infty)^S$ .

$\mathbf{x}(Y)$  = the **concentration** of  $Y$  in state  $\mathbf{x}$ .

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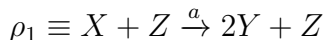
Behavior of  $N$  governed by autonomous polynomial ODEs.

Used to model systems with **many** molecules of each species.

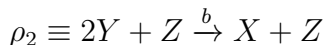
# Law of Mass Action (deterministic)

The **rate** of a reaction depends on its rate constant **and the state**!

*Example.* In state  $\mathbf{x}$ , the reactions



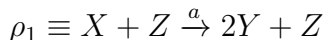
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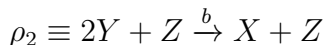
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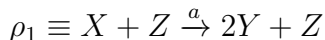
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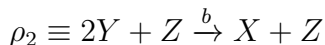
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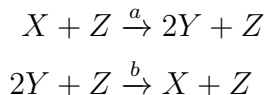
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$$rate_{\mathbf{x}}(\rho_1) = a\mathbf{x}(X)\mathbf{x}(Z)$$

and

$$rate_{\mathbf{x}}(\rho_2) = b\mathbf{x}(Y)^2\mathbf{x}(Z).$$

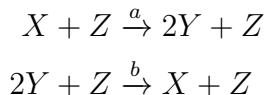
*Example* (continued). The CRN



is governed by the system

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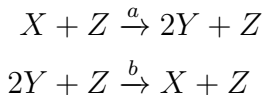
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$$\begin{aligned}\frac{dx}{dt} &= -axz + by^2z \\ \frac{dy}{dt} &= 2axz - 2by^2z \\ \frac{dz}{dt} &= 0 \quad (Z \text{ is a catalyst.})\end{aligned}$$

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**A harder question:** If so, is this universality robust?

What is **robustness** here?



# The CRN Model

Klinge, Lathrop, and Lutz 2015: Uniform translation of a nondeterministic finite automaton  $M$  into a deterministic mass action CRN  $N$ .

- $N$  simulates  $M$

- $\text{size}(N)$  linear in  $\text{size}(M)$

- $N$  robust with respect to

  - input signal

  - initial concentrations

  - output

  - rate constants of reactions

- Proof, not simulation.

# The CRN Model

## 2. Stochastic mass action

A **state** is a vector  $\mathbf{x} \in \mathbb{N}^S$ .

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Used to model systems in which **small** counts of molecules are significant.

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**Theorem** (Kurtz 1972). Roughly stated, deterministic mass action is the limiting case of stochastic mass action as populations and volumes go to infinity.

# The CRN Model

The computational power of stochastic mass action CRNs.

- Angluin, Aspnes, and Eisenstat 2006;

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Stable computation (correctness on **all** paths) can only compute semilinear predicates and functions.

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- The known simulation is **not** robust.



# The CRN Model

**Recall** For  $\lambda \in (0, \infty)$ , the **exponential distribution** with **rate**  $\lambda$  is the probability measure on  $[0, \infty)$  given by the c.d.f.

$$F_\lambda : [0, \infty) \rightarrow [0, 1]$$

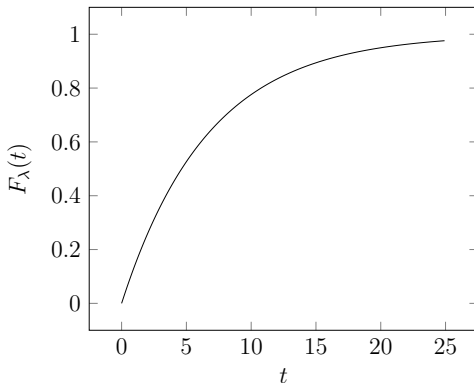
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$$\Delta(Q) = \{\text{probability measures on } Q\}.$$

Semantics of a CTMC  $C = (Q, \lambda, \pi)$ .

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- For each state  $q \in Q$ ,

$$\lambda_q = \sum_{r \in Q} \lambda(q, r)$$

is the **rate out of**  $q$ .

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When  $C$  does move to a new state, this state is  $r$  with probability

$$p(q, r) = \frac{\lambda(q, r)}{\lambda_q}.$$

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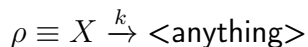
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- $\pi \in \Delta(Q)$  is how you want to initialize the species counts in  $N$ .
- Still need  $\lambda : \mathbb{N}^S \times \mathbb{N}^S \rightarrow [0, \infty)$ .



# Law of Mass Action (stochastic)

Intuition: The rate of a **unary** reaction

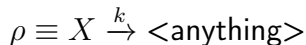


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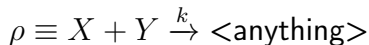
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The rate of a **binary** reaction



in a state  $\mathbf{x} \in \mathbb{N}^S$  and volume  $V$  is

$$\text{rate}_{\mathbf{x}}(\rho) = \begin{cases} k\mathbf{x}(X)\mathbf{x}(Y)/V & \text{if } X \neq Y \\ k\mathbf{x}(X)(\mathbf{x}(X) - 1)/V & \text{if } X = Y \end{cases}$$

The **rate** of a reaction  $\rho = (\mathbf{r}, \mathbf{p}, k)$  in a state  $\mathbf{x} \in \mathbb{N}^S$  is

$$\text{rate}_{\mathbf{x}}(\rho) = \frac{k}{V^{\|\mathbf{r}\|-1}} \prod_{Y \in S} \frac{\mathbf{x}(Y)!}{(\mathbf{x}(Y) - \mathbf{r}(Y))!},$$

where  $V$  is the volume of the solution and

$$\|\mathbf{r}\| = \sum_{Y \in S} \mathbf{r}(Y).$$

Gillespie, 1977

In the CTMC  $C = (\mathbb{N}^S, \lambda, \pi)$  for a CRN  $N = (S, R)$  we define the rate matrix  $\lambda : Q \times Q \rightarrow [0, \infty)$  by

$$\lambda(\mathbf{x}, \mathbf{y}) = \sum \{rate_{\mathbf{x}}(\rho) \mid \rho \in R \text{ takes } \mathbf{x} \text{ to } \mathbf{y}\}.$$

We have now completely specified the most often-used semantics (kinetics) for CRNs, namely,

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Caution: These are sometimes called

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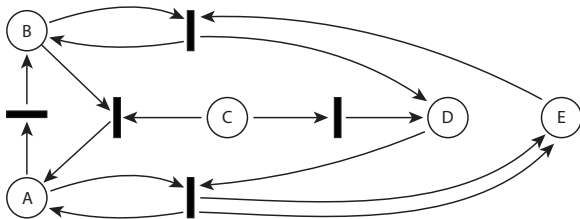
respectively.

# Relating CRNs to Other Models

## 1. Petri nets (Petri 1962).

A **Petri net** is a directed bipartite graph whose vertices are **places** (circles) and **transitions** (bars).

*Example* (CSWB 2009)



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A **state** of a Petri net is a placement of zero or more **tokens**, all of which are identical, on its places.



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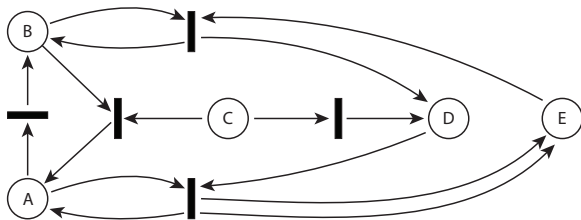
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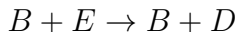
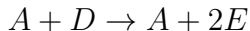
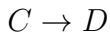
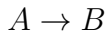
In this case, the transition may **fire**, causing these tokens to be removed and tokens to be placed at the targets of its outgoing edges.

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It is routine to check that the Petri net



has the same behavior as the CRN



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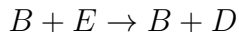
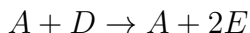
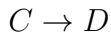
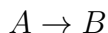
## 2. Vector addition systems (Karp and Miller 1969)

Fix  $n \in \mathbb{Z}^+$  and a finite set  $V$  of vectors  $\mathbf{v} \in \mathbb{Z}^n$ .

Consider **walks** in  $\mathbb{N}^n$  that start at a location  $\mathbf{x}_0 \in \mathbb{N}^n$  and proceed in **steps**, each of which adds a vector in  $V$  to the current location, leading to a new location in  $\mathbb{N}^n$ .

# Relating CRNs to Other Models

*Example* (CSWB 2009) The CRN



is equivalent to the vector addition system

|   | A  | B  | C  | D  | E  | F  | G  |   |
|---|----|----|----|----|----|----|----|---|
| ( | -1 | 1  | 0  | 0  | 0  | 0  | 0  | ) |
| ( | 0  | 0  | -1 | 1  | 0  | 0  | 0  | ) |
| ( | 1  | -1 | -1 | 0  | 0  | 0  | 0  | ) |
| ( | -1 | 0  | 0  | -1 | 0  | 1  | 0  | ) |
| ( | 1  | 0  | 0  | 0  | 2  | -1 | 0  | ) |
| ( | 0  | -1 | 0  | 0  | -1 | 0  | 1  | ) |
| ( | 0  | 1  | 0  | 1  | 0  | 0  | -1 | ) |

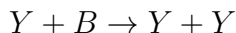
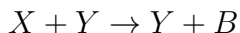
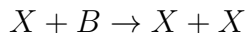
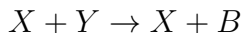
# Relating CRNs to Other Models

## 3. Population protocols ([Angluin, Aspnes, and Eisenstat 2008](#))

These are essentially CRNs in which every reaction has exactly two reactants and two products.

Note that this fixes the total population.

*Example* ([AAE 2008](#)) The population protocol

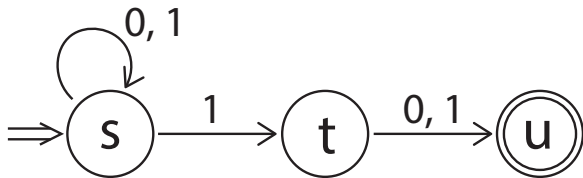


computes **approximate majority**.

# A CRN That Simulates an NFA

with Titus Klinge and Jim Lathrop

Let  $M$  be the NFA



Objective: Design a CRN  $N$  that simulates  $M$  in deterministic mass action.

# A CRN That Simulates an NFA

$M$  gets its input  $w \in \{0, 1\}^*$  sequentially, so  $N$  should get its input sequentially.





# A CRN That Simulates an NFA

We thus have **input species**  $X_0, X_1, X_c, X_r$ .

# A CRN That Simulates an NFA

We thus have **input species**  $X_0, X_1, X_c, X_r$ .

We also have

- **state species**  $Y_s, Y_t, Y_u$ ,

# A CRN That Simulates an NFA

We thus have **input species**  $X_0, X_1, X_c, X_r$ .

We also have

- **state species**  $Y_s, Y_t, Y_u$ ,
- **portal species**  $Z_s, Z_t, Z_u$ , and

# A CRN That Simulates an NFA

We thus have **input species**  $X_0, X_1, X_c, X_r$ .

We also have

- **state species**  $Y_s, Y_t, Y_u$ ,
- **portal species**  $Z_s, Z_t, Z_u$ , and
- **dual species**  $\overline{Y}_s, \overline{Y}_t, \overline{Y}_u, \overline{Z}_s, \overline{Z}_t, \overline{Z}_u$ .

# Reactions

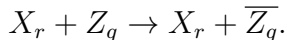
- For each state  $q$ ,

$$X_r + Z_q \rightarrow X_r + \overline{Z}_q.$$

reset

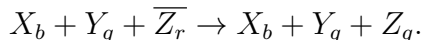
# Reactions

- For each state  $q$ ,



reset

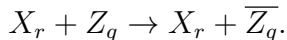
- For each transition  $\textcircled{q} \xrightarrow{b} \textcircled{r}$ ,



compute

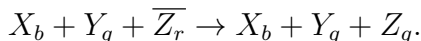
# Reactions

- For each state  $q$ ,



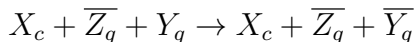
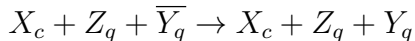
reset

- For each transition  $\textcircled{q} \xrightarrow{b} \textcircled{r}$ ,



compute

- For each state  $q$

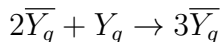
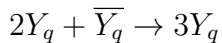


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# Reactions

- For each state  $q$ ,



error correction

# Summary

CRNs are programs.

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CRNs are magnets for good students.

# Summary

CRNs are programs.

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Thank you!