## Computation with chemistry

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ECS 232: Theory of Molecular Computation, UC Davis



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$$R \rightarrow P_1 + P_2$$
 product(s)

monomers 
$$M_1 + M_2 \rightarrow D$$
 dimer

catalyst  $C+X \rightarrow C+Y$ 

Traditionally a descriptive modeling language... Let's instead use it as a prescriptive programming language







### Computation with chemical reaction networks

- Key ideas setting chemical computation apart from others:
  - <u>cannot</u> control order in which molecules collide
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- Key ideas setting chemical computation apart from others:
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  - <u>can</u> control how they react when they collide
- Related model of distributed computing called *population protocols* 
  - originally motivated by mobile wireless sensor networks, e.g., attached to a birds in a flock

[Computation in networks of passively mobile finite-state sensors, Angluin, Aspnes, Diamadi, Fischer, Peralta. <u>PODC</u> 2004]



### Example: Chemical caucusing

opposite opinions cancel

 $X+Y \rightarrow U+U$ 

#### distributed algorithm for *"approximate majority"*: initial majority (*X* or *Y*) quickly overtakes whole population (with high probability)

### Example: Chemical caucusing

opposite opinions cancel

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both opinions influence the unopinionated  $\begin{array}{c} X + U \rightarrow X + X \\ Y + U \rightarrow Y + Y \end{array}$ 

distributed algorithm for *"approximate majority"*: initial majority (*X* or *Y*) quickly overtakes whole population (with high probability)



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### **Does** chemistry compute?



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[Cardelli, Csikász-Nagy. The cell cycle switch computes approximate majority. *Nature Scientific Reports* 2012] [Cardelli, Morphisms of reaction networks that couple structure to function, *BMC Systems Biology* 2014]



versus





versus



speed?



slow

versus





fast



versus



fast

slow



<u>k</u>d :







spied?

fast

≈ 10-100 nm

comportent size?

≈ 10-100 nm



#### Why compute with chemistry? versus cells fast slow smart drug released only in comportent size? certain cellular ≈ 10-100 nm ≈ 10-100 nm conditions compatible with not easily yes "wet environments"? **DNA** storage bioreactors in-place computation chemical controller to replacing expensive optimize yield of read/write lab steps metabolically produced biofuels/drugs/etc.

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 $X_1 + X_2 \rightarrow X_3$ 

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8

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$$X_1 + X_2 \to X_3$$



### DNA strand displacement implementing $A+B \rightarrow C$



# Experimental implementations of synthetic chemical reaction networks with DNA





#### actual chemicals

#### found in biology


## Theoretical Computer Science Approach





What computation is possible and what is not? (*Computability theory*)

## Theoretical Computer Science Approach





What computation is possible and what is not? (*Computability theory*)



What computations necessarily take a long time and what can be done quickly? (*Computational complexity theory*)

• finite set of d species  $\Lambda = \{ A, B, C, D, \dots \}$ 

• finite set of <u>reactions</u>: e.g.  $A+B \xrightarrow{k_1} A+C$  $C \xrightarrow{k_2} A+A$  $C+2B \xrightarrow{k_3} C$ 

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• finite set of <u>reactions</u>: *e.g.*  $A+B \xrightarrow{k_1} A+C$ 

 $k_1, k_2, k_3$  are called <u>rate constants</u>; if not specified, assume = 1.  $\begin{array}{c} C \xrightarrow{k_2} A + A \\ C + 2B \xrightarrow{k_3} C \end{array}$ 

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• finite set of <u>reactions</u>: e.g.  $k_1, k_2, k_3$  are called <u>rate constants</u>; if not specified, assume = 1. •  $A+B \xrightarrow{k_1} A+C$   $C \xrightarrow{k_2} A+A$  $C+2B \xrightarrow{k_3} C$ 

• <u>configuration</u>  $\mathbf{x} \in \mathbb{N}^d$ : molecular counts of each species

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- <u>configuration</u>  $\mathbf{x} \in \mathbb{N}^d$ : molecular counts of each species
- reaction is <u>applicable</u> to **x** if **x** has enough of each reactant.

- $\alpha: \qquad A+B \rightarrow A+C \qquad A \quad B \quad C$
- β: C → A+A **x** = (2, 2, 0) α applicable but not β



Example reaction sequence (a.k.a. *execution*)

α:A+B → A+CABCβ:C → A+Ax = (2, 2, 0)α applicable but not β







α:	$A+B \rightarrow A+C$	A B	С	
β:	$C \rightarrow A + A$	x = (2, 2,	0)	$\alpha$ applicable but not $\beta$
		α ↓		
		(2, 1,	1)	$\alpha$ , β both applicable
		β ↓		
	B	(4, 1,	0)	
	A			







Example reaction sequence (a.k.a. *execution*)

R  $A+B \rightarrow A+C$ α:  $C \rightarrow A + A$  $\alpha$  applicable but not  $\beta$ x = (2, 2, 0)β: α ]]  $\alpha,\beta$  both applicable (2, 1, 1)  $\searrow \alpha$  (another possibility) β↓ (2, 0, 2) (4, 1, 0)α∥ (4, 0, 1)

Formally, an execution is a sequence of *configurations*  $\mathbf{x}_1$ ,  $\mathbf{x}_{2}$ , ... such that each  $\mathbf{x}_{i} \Longrightarrow \mathbf{x}_{i+1}$  by a single reaction. If initial configuration  $\mathbf{x}_1$  is understood, the sequence of reactions is sometimes called the execution.



start with *n* copies of molecule X

 $X \stackrel{1}{\longleftrightarrow} Y$ 

start with *n* copies of molecule X

#Y = n/2 expected at equilibrium 100 -#Х #Y 80 E[#Y] 60 counts 40 20 0 0 2 3 1 4 time

$$X \xleftarrow{1}{1} Y \qquad \begin{array}{c} \text{Count of } Y \\ \text{never stabilizes} \end{array}$$

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$$\begin{array}{c} X \xrightarrow{1} Y \\ X \xrightarrow{1} \end{array}$$

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 $\begin{array}{c} X \xrightarrow{1} & Y \\ X \xrightarrow{1} & \end{array}$ 

#### **#***Y* <u>stabilizes</u>, with expected value *n*/2



 $X \xleftarrow{1}{1} Y$ Count of Y
never stabilizes

start with *n* copies of molecule X

#Y = n/2 expected at equilibrium



 $\begin{array}{c} X \xrightarrow{1} \\ Y \xrightarrow{1} \end{array}$ 

Count of Y stabilizes, but not to a deterministic value based on initial count of X

#### **#***Y* <u>stabilizes</u>, with expected value *n*/2



 $X \xleftarrow{1}{\underbrace{1}{\underbrace{1}{2}}} Y$  Count of Y never stabilizes

start with *n* copies of molecule X*n*/3

 $#Y = \frac{n/2}{n/2}$  expected at equilibrium

Worse yet, both depend crucially on rate constants.

 $\begin{array}{c} X \xrightarrow{1} Y \\ X \xrightarrow{\pm 2} \end{array}$ 

Count of Y stabilizes, but <u>not</u> to a deterministic value based on initial count of X

*n*/3 #Y stabilizes, with expected value <del>*n*/2</del>

 $X \xleftarrow{1}{\underbrace{1}{\underbrace{1}{2}}} Y$  Count of Y never stability

never stabilizes

start with *n* copies of molecule X

n/3#Y = n/2 expected at equilibrium



Worse yet, both depend crucially on rate constants.

 $\begin{array}{c} X \xrightarrow{1} Y \\ X \xrightarrow{\pm 2} \end{array}$ 

Count of Y stabilizes, but <u>not</u> to a deterministic value based on initial count of X

n/3
#Y stabilizes, with expected value n/2



# Examples of **stable** (*rateindependent*) CRN computation

division by 2: f(a) = a/2goal: end up with a/2 copies of Y



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??
division by 2: f(a) = a/2
goal: end up with a/2 copies of Y

division by 2: f(a) = [a/2]
goal: end up with a/2 copies of Y



division by 2: f(a) = [a/2]
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multiplication by 2: f(a) = 2a





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 $A \rightarrow 2Y$ 



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Y

multiplication by 3: f(a) = 3a



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division by 3:  $f(a) = \lfloor a/3 \rfloor$ 



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f(a) = 3a using ( $\leq 2$ )-product reactions



f(a) = 3a using ( $\leq 2$ )-product reactions

 $\begin{array}{c} A \rightarrow \mathbf{Y} + \mathbf{Y'} \\ \mathbf{Y'} \rightarrow \mathbf{2Y} \end{array}$ 



f(a) = 3a using ( $\leq 2$ )-product reactions

$$\frac{A \rightarrow Y + Y'}{Y' \rightarrow 2Y}$$



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YYYYYYYY

 $f(a) = \lfloor a/3 \rfloor$  using bimolecular (( $\leq 2$ )-reactant) reactions, starting in config { 1  $L_0$ ,  $a \land A$  } (a.k.a., *leader-driven*)



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YYYYYYYY

$$L_0 + A \rightarrow L_1$$

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$$L_2 + A \rightarrow L_0 + Y$$



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 $L_0 + A \rightarrow L_1$   $L_1 + A \rightarrow L_2$   $L_2 + A \rightarrow L_0 + Y$ 

L<sub>1</sub> Y A

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L<sub>1</sub> Y

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$$L_0 + A \rightarrow L_1$$

$$L_1 + A \rightarrow L_2$$

$$L_2 + A \rightarrow L_0 + Y$$

ends with 1 copy of *L<sub>i</sub>* for *i* = ???



 $f(a) = \lfloor a/3 \rfloor$  using bimolecular ( $\leq 2$ -reactant) reactions, starting in config {a A} (a.k.a., *leaderless*)
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$$A + A \rightarrow A_2$$
$$A_2 + A \rightarrow Y$$
$$A_2 + A_2 \rightarrow A + Y$$

Calling  $A = A_1$ , in general to divide by constant c:  $A_i + A_j \rightarrow A_k$  if i+j < c, where k = i + j  $A_i + A_j \rightarrow A_k + Y$  if i+j > c, where k = i + j - c $A_i + A_j \rightarrow Y$  if i+j = c

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Calling  $A = A_1$ , in general to divide by constant c:  $A_i + A_j \rightarrow A_k$  if i+j < c, where k = i + j  $A_i + A_j \rightarrow A_k + Y$  if i+j > c, where k = i + j - c  $A_i + A_j \rightarrow Y$  if i+j = ci.e., A's start with 1 "ball" and pass balls to each other; whenever someone gets  $\geq c$  balls, throw away c balls and produce a Y

addition: *f*(*a*,*b*) = *a*+*b* 



addition: *f*(*a*,*b*) = *a*+*b* 

 $\begin{array}{c} A \to Y \\ B \to Y \end{array}$ 



addition: f(a, b) = a+b

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addition: f(a, b) = a+b

 $\begin{array}{c} A \to Y \\ B \to Y \end{array}$ 

subtraction: f(a,b) = a-b



addition: f(a, b) = a+bsubtraction: f(a, b) = a - b $A \rightarrow Y$  $A \rightarrow Y$  $B \rightarrow Y$  $B+Y \rightarrow \emptyset$ A A A A (A) $\left( Y \right)$ (A)[Y]Y B  $\left( B\right)$ 

addition: f(a, b) = a+bsubtraction: f(a, b) = a - b $A \rightarrow Y$  $A \rightarrow Y$  $B \rightarrow Y$  $B+Y \rightarrow \emptyset$ Y [Y]Y В B

addition: f(a,b) = a+b  $A \rightarrow Y$   $B \rightarrow Y$  YY

subtraction: f(a, b) = a - b

 $A \to \mathbf{Y}$  $\mathbf{B} + \mathbf{Y} \to \mathbf{\emptyset}$ 

subtraction: f(a,b) = a-b addition: f(a, b) = a+b $A \rightarrow Y$  $A \rightarrow Y$  $B \rightarrow Y$  $B+Y \rightarrow \emptyset$ Y  $\left( Y \right)$ Y Y Y Y Y

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B

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 $\begin{array}{c} A \to Y \\ B \to Y \end{array}$ 

subtraction: f(a, b) = a - b

 $A \to \mathbf{Y}$  $\mathbf{B} + \mathbf{Y} \to \mathbf{\emptyset}$ 





addition: f(a, b) = a+b

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 $\begin{array}{c} A \to Y \\ B \to Y \end{array}$ 

Y

 $\left( Y \right)$ 

(Y)

 $A \to \mathbf{Y}$  $\mathbf{B} + \mathbf{Y} \to \mathbf{\emptyset}$ 

В

addition: f(a,b) = a+b

??? **subtraction:** *f*(*a*,*b*) =

 $A \rightarrow Y$ 

 $\begin{array}{c} A \to Y \\ B \to Y \end{array}$ 

Y

[Y]

(Y)

 $B+Y \rightarrow \emptyset$ 

B

addition: f(a, b) = a+b $A \rightarrow Y$  $A \rightarrow Y$  $B \rightarrow Y$  $B+Y \rightarrow \emptyset$ (Y)Y Y

subtraction:  $f(a,b) = \longrightarrow (max(0, a-b))$ 

B

**composition:** *f*(*a*,*b*) = 3*a*–*b* 

composition: f(a,b) = 3a-b  $A \rightarrow 3Y$  $B+Y \rightarrow \emptyset$ 

composition: f(a,b) = 326??  $A \rightarrow 3Y$  $B+Y \rightarrow \emptyset$ 

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composition:  $f(a,b) = 3 \rightarrow 6$ ???  $A \rightarrow 3Y$  $2B+Y \rightarrow \emptyset$  maximum:  $f(a,b) = \max(a,b) = a+b-\min(a,b)$   $A \rightarrow Y + A_2$  $B \rightarrow Y + B_2$  addition

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only linear functions computable?

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## Examples of predicate computation

**Detection:**  $\varphi(a,b) = yes \Leftrightarrow b > 0$ 



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**Majority:**  $\varphi(a, b) = yes \Leftrightarrow a \ge b$ 

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 $A+B \rightarrow A_f+B_f$  (both become "followers" but <u>preserve difference</u> between A's and B's)

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$$A_o + a_e \rightarrow A_o + a_o$$
 leader overwrites  
 $A_e + a_o \rightarrow A_e + a_e$  bit of follower

# Formal definition of CRN computation

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(assuming finite set of reachable configurations) equivalent to: The system <u>will</u> reach a correct stable configuration with probability 1.
# Probability-1 correctness can be characterized with only reachability

### Probability-1 correctness can be characterized with only reachability To understand this slide, only need the following fact: if a reaction

is applicable, then there is a positive probability it occurs.

**Definition**: Let **i** be a configuration and *Y* be a set of configurations. Write  $Pr[\mathbf{i} \Rightarrow Y]$  to denote the probability of the random event that, starting in configuration **i**, the CRN eventually reaches some configuration  $\mathbf{o} \in Y$ .

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**Theorem**: Let **i** be a configuration where Reach(**i**) is finite, and let Y be a set of configurations. Then  $(\Pr[\mathbf{i} \Longrightarrow Y] = 1) \Leftrightarrow (\forall \mathbf{x} \in \operatorname{Reach}(\mathbf{i})) (\exists \mathbf{o} \in \operatorname{Reach}(\mathbf{x})) \mathbf{o} \in Y.$ 

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This theorem lets us use (often simpler) reachability arguments and avoid discussing probability, while still ensuring probability-1 correctness.

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- 1. (⇒): Assume ( $\exists x \in \text{Reach}(i)$ ) ( $\forall o \in \text{Reach}(x)$ )  $o \notin Y$ .
- 2. Since  $Pr[i \implies x] > 0$ , which prevents ever reaching Y,  $Pr[i \implies Y] < 1$ . (Note this didn't assume Reach(i) is finite.)
- 3. ( $\Leftarrow$ ): Assume ( $\forall x \in \text{Reach}(i)$ ) ( $\exists o \in \text{Reach}(x)$ )  $o \in Y$ .
- 4. For each  $\mathbf{x} \in \text{Reach}(\mathbf{i})$ , let  $E_{\mathbf{x}} = (\mathbf{x},...,\mathbf{o})$  be any finite execution leading from **x** to some  $\mathbf{o} \in Y$ .
- 5. Let  $k = \max_{x \in \text{Reach}(i)} |E_x|$  be the maximum length of any of these finite executions reaching **o**.
- 6. Let  $p_{\mathbf{x}} = \Pr[E_{\mathbf{x}} \text{ occurs from } \mathbf{x}] > 0$ .
- 7. Let  $\varepsilon = \min_{\mathbf{x} \in \text{Reach}(\mathbf{i})} p_{\mathbf{x}}$ . Since Reach(**i**) is finite,  $\varepsilon > 0$ .
- 8. Then for each  $\mathbf{x} \in \text{Reach}(\mathbf{i})$ ,  $\Pr[E_{\mathbf{x}} \text{ does not occur from } \mathbf{x}]$ after the next k steps]  $\leq 1 - \varepsilon < 1$ .
- 9. So, breaking the infinite execution into segments of length k, the probability  $E_{x}$  is <u>never</u> followed within k steps after any visit to an  $\mathbf{x} \in \text{Reach}(\mathbf{i})$  is at most  $\prod_{i=1}^{\infty} (1-\varepsilon) = 0.$  QED

**False statement**: If  $Pr[i \Rightarrow Y] = 1$ , then every sufficiently long execution starting at **i** reaches to some **c**  $\in$  *Y*.

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- Lesson: it is too strict to require <u>all</u> sufficiently long executions to reach Y.

**Definition**: An infinite execution  $\mathbf{x}_0, \mathbf{x}_1, \dots$  is <u>fair</u> if

Goal of definition of <u>fair</u> is to make this theorem true:

**Theorem**: Let **i** be a configuration, and let *Y* be a finite set of configurations. Then (every fair execution starting at **i** reaches some  $\mathbf{o} \in Y$ )  $\Leftrightarrow (\forall \mathbf{x} \in \text{Reach}(\mathbf{i})) (\exists \mathbf{o} \in \text{Reach}(\mathbf{x})) \mathbf{o} \in Y.$  **Definition**: An infinite execution  $\mathbf{x}_0, \mathbf{x}_1, \dots$  is <u>fair</u> if

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Proof:

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- 1.  $(\Rightarrow)$ : Suppose every fair execution from **i** reaches *Y*.
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- 8. Since  $\mathbf{x}_0, \mathbf{x}_1, \dots$  is fair and **o** is infinitely often reachable, there is k such that  $\mathbf{x}_k = \mathbf{o} \in Y$ , i.e., the fair execution reaches Y. **QED**

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  - **Recall**: this is equivalent to saying that **i** reaches to a correct, stable **o** with probability 1, <u>and</u> equivalent to saying that every fair execution from **i** reaches to a correct, stable **o**.

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  - We say the CRN stably decides the set  $\varphi^{-1}(Y)$  = set of inputs mapping to output Y

A class of CRNs with a simpler definition/proofs for computation

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**Note**: A configuration can be *stable without being terminal*. Example?

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37

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- 3. CRN produces #A+#B count of Y by rxns 1 and 2, and consumes min(#A,#B) Y's by rxn 4, so computes #A+#B-min(#A,#B) = max(#A,#B). QED

# In feed-forward CRNs, if there is a terminal configuration, any long enough execution reaches it

**Lemma** (*restated*): Suppose that in a feedforward CRN,  $\mathbf{i} \Rightarrow \mathbf{c}$  by execution *P*, and  $\mathbf{i} \Rightarrow \mathbf{d}$ by execution *Q*. If any reaction occurs less in *P* than *Q*, then **c** is not terminal.

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- 4. Finally, to rule out that we might have some <u>shorter</u> terminal execution, any execution Q with |Q| < |P| must have some reaction r occurring more in P than Q, so by the Lemma, Q cannot reach a terminal configuration. QED</li>

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3. $A_2 + B_2 \rightarrow K$	$(A_2, B_2 \text{ aren't reactants elsewhere})$
4. $K+Y \rightarrow Ø$	(K, Y aren't reactants elsewhere)

We often convince ourselves a CRN works by examining just one execution that stabilizes to the correct output, and thinking, "The other executions probably/hopefully end up with the same output." This reasoning becomes sound with non-competitive CRNs.

#### Proof:

- 1.  $Q' = \text{longest prefix } (\mathbf{i}, \mathbf{x}_1, ..., \mathbf{x}_p) \text{ of } Q \text{ such that } \#(r, P) \ge \#(r, Q) \text{ for all reactions } r.$ 
  - i.e.,  $\mathbf{x}_{p+1}$  is the first time in Q that some reaction exceeds its count in P.
- 2. Let *r* be the reaction such that  $\mathbf{x}_p \Longrightarrow \mathbf{x}_{p+1}$  via *r*.
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  - 1. Some other reactions *t* might <u>produce</u> *A*.
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**Definition**: A CRN is <u>non-competitive</u> if, for every species R, if R is net consumed in some reaction (e.g.,  $R \rightarrow A$  or  $2R \rightarrow R$ ), then R is not a reactant in any other reaction. (R can be a <u>non-consumed catalyst</u> in any number of reactions, e.g.,  $R \rightarrow 2R$  or  $R+X \rightarrow R+Y$ , but then <u>no</u> reaction can net consume it)

**Lemma**: Suppose in a non-competitive CRN that  $\mathbf{i} \Rightarrow \mathbf{c}$  by execution *P*, and  $\mathbf{i} \Rightarrow \mathbf{d}$  by execution *Q*. If any reaction occurs less in *P* than *Q*, then **c** is not terminal.

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Example of a non-feedforward CRN that stably computes a function?

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$$f(x) = x/2$$
  
1. X \Rightarrow Y+A  
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## Time complexity of CRNs

### What is **probable**:

### Stochastic kinetic model of chemical reaction networks

Solution volume v

reaction typerate / propensity $A \xrightarrow{k} \dots$  $k \cdot #A$  $A+B \xrightarrow{k} \dots$  $k \cdot #A \cdot #B / v$ 

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expected time until next reaction is 1 / (sum of all reaction rates)

### **population protocol** = list of *transitions* such as

 $x,y \rightarrow x,x$   $a,b \rightarrow c,d$   $a,a \rightarrow a,a$  (null transition)

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## population protocols $\subsetneq$ chemical reactions, but "most" ideas that apply to one model also apply to the other

[Angluin, Aspnes, Diamadi, Fischer, Peralta, *Computation in networks of passively mobile finite-state sensors*, <u>PODC</u> 2004]: Winner of 2020 Dijkstra Prize in Distributed Computing: <u>https://www.podc.org/dijkstra/2020-dijkstra-prize/</u>

### Time complexity in population protocols

- pair of agents picked uniformly at random to interact (possibly null interaction)
- *parallel time* = number of interactions / *n*

i.e., each agent has O(1) interactions per "unit time"
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n = total molecular count

reasonable requirement on volume: v = O(n)

*i.e.*, <u>require bounded concentration</u> (finite density constraint)

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  - In general, with r reactants, propensity is number of ways to pick reactants, times k, divided by  $v^{r-1}$

В

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both are <i>me</i>	moryless:	$\forall s, r > 0$
Pr[1>S+r]		
no rxn after <i>r</i>	no rxn in	no rxn in
additional secs.	first <i>s</i> secs.	first <i>r</i> secs.

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  - If we treat interactions symmetrically, (*i.e., a,b* → *c,d* is an interaction if and only if b,a → d,c is an interaction), then we have twice the probability, i.e., expected time becomes (n-1) / #a·#b ~ n / (#a·#b), essentially the same as the CRN model

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  - one possible convention to avoid symmetric interactions is simply define time = 2.#interactions/n

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Can use Chernoff bounds to show it is very likely that they end up taking very close to the same amount of time for any event.

a #b)	
an	

no rxn in

first r secs.

both are *memoryless*:  $\forall s,r > 0$ 

 $\Pr[\mathbf{T} > s + r \mid \mathbf{T} > s] = \Pr[\mathbf{T} > r]$ 

no rxn after r

additional secs.

no rxn in

first s secs.

*n* molecules volume *v* = *O*(*n*)



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$$A+B \rightarrow Y+B$$

propensity:  $#A \cdot #B / v = O(1/n)$ 

expected time to produce *Y*:

*n* molecules volume v = O(n)

produce *Y*:





*n* molecules volume v = O(n)

produce *Y*:



49

# Time complexity analysis (basic motifs)

"direct communication"

 $|A+B \rightarrow Y+W| \quad \#A=\#B=1, \ \#X=n-2$
"direct communication"

$$A+B \rightarrow Y+W \quad \#A=\#B=1, \ \#X=n-2$$

**population protocol time complexity**: time until non-null interaction is geometric random variable with success probability p = 1 / (n choose 2) = 2 / (n(n-1))

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time until reaction is exponential random variable with rate  $\lambda = #A \cdot #B / n = 1 / n$ E[time] =  $1/\lambda = n$  "epidemic", "gossip", "rumor spreading"

$$B+X \rightarrow B+B$$

"direct communication"

"epidemic", "gossip", "rumor spreading"

$$A + B \rightarrow Y + W \quad \#A = \#B = 1, \ \#X = n - 2$$

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time until reaction is exponential random variable with rate  $\lambda = #A \cdot #B / n = 1 / n$ E[time] =  $1/\lambda = n$  population protocol time complexity:

"direct communication"

$$A+B \rightarrow Y+W$$
 # $A=$ # $B=1$ , # $X=n-2$ 

population protocol time complexity:

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when #B = k, we have #X = n-k $\Pr[B+X \rightarrow B+B$  is pertipheraction | #B-k | = k(n-k) / (n choose - k)

Pr[ $B+X \rightarrow B+B$  is next interaction | #B=k] = k(n-k) / (n choose 2)

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**population protocol time complexity**: when #B = k, we have #X = n-kPr[ $B+X \rightarrow B+B$  is next interaction | #B=k] = k(n-k) / (n choose 2)= 2k(n-k) / ((n(n-1)))

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, we have  $\#X = n-k$   
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 $= 2k(n-k) / ((n(n-1)))$   
expected time until one X converted to  $B = 1/(n \cdot \text{probability})$   
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expected time until all X converted to  $B =$ 

$$\frac{n-1}{2} \sum_{k=1}^{n-1} \frac{1}{k(n-k)}$$

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$$\frac{n-1}{2}\sum_{k=1}^{n-1}\frac{1}{k(n-k)} = \frac{n-1}{2}\sum_{k=1}^{n-1}\frac{1}{n}\left(\frac{1}{k} + \frac{1}{n-k}\right)$$

"direct communication"

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$$\frac{n-1}{2} \sum_{k=1}^{n-1} \frac{1}{k(n-k)} = \frac{n-1}{2} \sum_{k=1}^{n-1} \frac{1}{n} \left(\frac{1}{k} + \frac{1}{n-k}\right)$$

$$\approx \frac{1}{2} \left(\sum_{k=1}^{n} \frac{1}{k} + \sum_{k=n}^{1} \frac{1}{k}\right) = \sum_{k=1}^{n} \frac{1}{k}$$

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#### population protocol time complexity:

time until non-null interaction is geometric random variable with success probability p = 1 / (n choose 2) = 2 / (n(n-1))E[# interactions] = 1/p = (n(n-1)) / 2E[time] = E[# interactions]/n = (n-1) / 2 = O(n)

#### **CRN time complexity**:

time until reaction is exponential random variable with rate  $\lambda = #A \cdot #B / n = 1 / n$ E[time] =  $1/\lambda = n$  "epidemic", "gossip", "rumor spreading"

#**B**=0

"no communication"

? here means "every species" (including A)

$$A+? \rightarrow B+? \qquad \#A=n,$$

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population protocol time complexity:

When #*A*=*k*, time until non-null interaction is geometric random variable with success probability p = k(n-1) / (n choose 2) = k / (2n)E[# interactions] = 1/p = n / kE[time until non-null interaction] = E[# interactions] / n = 1 / kE[time to convert all *A*] =  $\frac{1}{2} \sum_{k=1}^{n} \frac{1}{k} \approx (1/2) \ln n$ 

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 $A+? \rightarrow B+?$ 

#**A**=n, #**B**=0

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$$A \rightarrow B$$

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### **CRN time complexity**: When #4-k time until n

When #A=k, time until next reaction is exponential random variable with rate  $\lambda = k$ 

"no communication"

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 $A+? \rightarrow B+?$ 

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#### **CRN time complexity**:

When #*A*=*k*, time until next reaction is exponential random variable with rate  $\lambda = k$ E[time until next reaction] =  $1/\lambda = 1/k$ E[time for all *n* reactions] =  $\sum_{k=1}^{n} \frac{1}{k} \approx \ln n$ 

"pairing off"

$$A+B \rightarrow C$$

#A=n, #B=n, total volume = O(total count) = n

"pairing off"

 $A+B \rightarrow C$ 

#A=n, #B=n, total volume = O(total count) = n

**CRN time complexity**: When #A=#B=k, next reaction has rate  $\lambda = k^2/n$ 

"pairing off"

#A=n, #B=n, total volume = O(total count) = n

CRN time complexity:

 $A+B \rightarrow C$ 

When #A=#B=k, next reaction has rate  $\lambda = k^2/n$ E[time until next reaction] =  $1/\lambda = n/k^2$ 

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"pairing off"

 $A+B \rightarrow C$ 

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**CRN time complexity**: When #*A*=#*B*=*k*, next reaction has rate  $\lambda = k^2/n$ E[time until next reaction] =  $1/\lambda = n/k^2$ E[time for all *n* reactions] =  $\sum_{k=1}^{n} \frac{n}{k^2}$   $< n \sum_{k=1}^{\infty} \frac{1}{k^2}$  $= n \cdot \pi^2/6 = \Theta(n)$ 

"pairing off"

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"pairing off" (symmetric version)



similar analysis

"coupon collecting"

$$L + A \rightarrow L + B$$

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 $= \Theta(n \log n)$ 

multiplication by 2: f(a) = 2a $A \rightarrow 2Y$ 

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O(log n) "unimolecular decay"

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Suppose *a* > *b*.

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subtraction: f(a,b) = a-b  $A \rightarrow Y$  $B+Y \rightarrow \emptyset$ 

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E[time] = E[time for first to finish] + E[time for second to finish] E[time for first to finish] =  $O(\log n)$  (unimolecular decay) E[time for second to finish] = O(n) in worst case: similar to minimum, worst case when a=b, but  $O(\log n)$  time if  $|a-b| = \Omega(n)$ .

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 $\mathsf{E}[\mathsf{time}] = O(\log n) + O(n) = O(n)$ 

maximum:  $f(a,b) = \max(a,b)$ 1.  $A \rightarrow Y + A_2$ 2.  $B \rightarrow Y + B_2$ 3.  $A_2 + B_2 \rightarrow K$ 4.  $K + Y \rightarrow \emptyset$ 

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- Assume reaction 3 waits for reactions 1 and 2 before starting, and reaction 4 waits for reaction 3.
- $E[\text{time for 1 and 2}] = O(\log n)$
- E[time for 3] = O(n)
- E[time for 4] = O(n)
- So  $E[time] = O(\log n) + O(n) + O(n) = O(n)$

# Possibilities of stable computation

What <u>can</u> be stably computed?

#### Summary: Possibilities and limits of stable computation

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#### Predicates

•  $\varphi$  is stably computable if and only if  $\varphi$  is *semilinear*.

[Angluin, Aspnes, Diamadi, Fischer, Peralta, *Computation in networks of passively mobile finite-state sensors, PODC* 2004] [Angluin, Aspnes, Eisenstat, *Stably computable predicates are semilinear, PODC* 2006]

#### Summary: Possibilities and limits of stable computation

#### **Predicates**

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- semilinear = Boolean combination of <u>threshold</u> and <u>mod</u> predicates: take weighted sum s = w<sub>1</sub>·a<sub>1</sub> + ... w<sub>d</sub>·a<sub>d</sub> of inputs and ask if s > t?
  (threshold) s = c mod m?
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  a>b? a=b? a is odd? a>0? a>1?
  NOT a=b<sup>2</sup>? a is a power of 2? a is prime?

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#### **Functions**

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a+b a-b 2a a/2 min(a,b) a+1 a-1 f(a) = 2a-b/3 if a+b is odd, else f(a) = a/4+5b

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f(a) =	2 <i>a</i> —b/3	if <i>a+b</i>	is odd,	else $f(a) = a_i$	/4+5 <i>b</i>	

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## All semilinear predicates/functions are known to be computable in O(n) time.

**Definition**: A set  $X \subseteq \mathbb{N}^d$  is <u>linear</u> if there are vectors  $\mathbf{b}$ ,  $\mathbf{u}_1$ , ...,  $\mathbf{u}_p \in \mathbb{N}^d$  such that  $X = \{ \mathbf{b} + n_1 \cdot \mathbf{u}_1 + ... + n_p \cdot \mathbf{u}_p \mid n_1, ..., n_p \in \mathbb{N} \}$ 

multi-dimensional generalization of *eventually periodic* 

Example in dimension *d*=2: **b** = (2,1)  $u_1 = (4,1)$ 5 3 **u**<sub>2</sub> = (2,2) 2 3 2

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#### Semilinear sets



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examples:

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**Definition 3**:  $X \subseteq \mathbb{N}^d$  is <u>semilinear</u> if it is definable in the first-order theory of Presburger arithmetic. (*original definition*, *hardest to understand; we won't use it.*)

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Other places semilinear sets show up in computer science:

- Sets decidable by *reversal-bounded counter machines*.
- In 2D, they are conjectured to be the sets weakly selfassembled by temperature τ=1 tile systems.

**Theorem 1**: A set  $X \subseteq \mathbb{N}^d$  is stably decided by some CRN if and only if it is semilinear.

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Full proof is too complex to do in this course. But we'll show:

- 1. All semilinear sets <u>can</u> be stably decided.
- 2. The non-semilinear "squaring" set  $X = \{ (a,y) \in \mathbb{N}^2 \mid a^2 = y \}$  cannot be stably decided.

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**Theorem 2**: A function  $f: \mathbb{N}^d \to \mathbb{N}$  is stably computed by some CRN if and only if it is semilinear.



# Possibilities of stable computation

All semilinear functions/predicates <u>can</u> be stably computed by CRNs

## Stably decidable sets are closed under Boolean operations

**Theorem**: If sets  $X_1, X_2 \subseteq \mathbb{N}^d$  are stably decided by some CRN, then so are  $X_1 \cup X_2, X_1 \cap X_2$ , and  $\overline{X_1}$ .

## Stably decidable sets are closed under Boolean operations

For this proof, we assume that the voting species can be a <u>strict</u> subset of all species.

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**Proof**:

1. To stably decide  $\overline{X_1}$ , swap the yes and no voters.
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- 1. To stably decide  $\overline{X_1}$ , swap the yes and no voters.
- 2. For  $\cup$  and  $\cap$ , let  $C_1$  and  $C_2$  stably decide  $X_1$  and  $X_2$ .

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- 3. Add the reaction  $A \rightarrow A_1 + A_2$  for each input species A, and let  $A_i$  be the input species for  $C_i$ .

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- 5. To "record" the votes of  $C_1$  and  $C_2$ :

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1. 
$$L_{Y} + N \rightarrow L_{N}$$
  
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**Corollary** (*since stably decidable sets are closed under Boolean combinations*): Every semilinear set is stably decided by some CRN.

#### Also true for <u>leaderless</u> CRNs.

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[Computation in networks of passively mobile finite-state sensors, Angluin, Aspnes, Diamadi, Fischer, Peralta. PODC 2004]

**Lemma**: If  $f: \mathbb{N}^d \to \mathbb{N}$  is a semilinear function, then it is <u>piecewise affine</u>: a finite union of partial affine functions  $g_i: \mathbb{N}^d \to \mathbb{N}$ .

Each  $g_i$  is <u>affine</u> (*linear with constant offsets*): there are  $w_1 \dots w_d \in \mathbb{Q}$  and  $b, c_1, \dots, c_d \in \mathbb{N}$  such that each  $g_i(x_1, \dots, x_d) = w_1 \cdot (x_1 - c_1) + \dots + w_d \cdot (x_d - c_d) + b$ .

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start with: (input) X output: Y

 $X{+}X \to Y$ 



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## Semilinear function examples



linear:

f(a,b,c) = 2a + (4/3)b - (5/6)c

 $\frac{\text{linear}}{f(a,b,c)} = 2a + (4/3)b - (5/6)c$   $A \rightarrow 2Y$ 

 $\frac{\text{linear}}{f(a,b,c)} = 2a + (4/3)b - (5/6)c$   $A \rightarrow 2Y$   $3B \rightarrow 4Y$ 

 $\frac{\text{linear}:}{f(a,b,c)} = 2a + (4/3)b - (5/6)c$   $A \rightarrow 2Y$   $3B \rightarrow 4Y$   $6C + 5Y \rightarrow \emptyset$ 

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**General form**:  $w_1 \dots w_d \in \mathbb{Q}$  and  $b, c_1, \dots, c_d \in \mathbb{N}$  $g_i(x_1, \dots, x_d) = w_1 \cdot (x_1 - c_1) + \dots + w_d \cdot (x_d - c_d) + b.$ 

add constant offset: start with 1 *L*, *a A*'s, *b B*'s f(a,b) = 2a + 3b + 4

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add constant offset: start with 1 *L*, *a A*'s, *b B*'s f(a,b) = 2a + 3b + 4  $L \rightarrow 4Y$   $A \rightarrow 2Y$  $B \rightarrow 3Y$
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point of species  $\hat{Y}_i$ ?

**Answer 2**: Consuming  $Y_i$  can disrupt computation of  $g_i$ . Can be solved using *dual-rail encoding*. (not shown)

# Limits of stable computation

Non-semilinear functions/predicates cannot be stably computed by CRNs

## Impossibility of stably deciding nonsemilinear sets

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To start, we use the above theorem to prove the following:

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- 8. Add reactions to test for equality between  $\#Y_P$  and  $\#Y_C$ . (*not shown, but easy*)

### Impossibility of stably deciding a non-semilinear set

goal:

**Theorem**: The "squaring set"  $S = \{ (x,y) \in \mathbb{N}^2 \mid x^2 = y \}$  is <u>not</u> stably decidable by any CRN.

### Additivity, nondecreasing sequences, minimal elements

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**Definition**: Given  $A \subseteq \mathbb{N}^d$ , we say  $\mathbf{y} \in A$  is <u>minimal</u> if, for all  $\mathbf{x} \in A$ ,  $\mathbf{x} \leq \mathbf{y}$  implies  $\mathbf{x} = \mathbf{y}$ , i.e., nothing in A is strictly smaller than  $\mathbf{y}$ . Let min(A) = minimal elements of A.



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- 5. ...
- 6. Since there are only a finite number of **y** in  $\mathbb{N}^d$  such that  $\mathbf{y} < \mathbf{x}$ , this process must terminate with a minimal vector  $\mathbf{m} \in \min(A)$ . **QED**

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- 4. For condition (2), suppose that min(A) is infinite; put them in any order to make an infinite sequence.

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- 2. <u>Base case d = 1</u>: Let  $X = x_0, x_1, \dots$  be an infinite sequence of nonnegative integers.
  - 1. case 1: some  $x \in \mathbb{N}$  appears infinitely often. Let the subsequence be (x, x, ...), e.g. 1,1,5,3,4,3,4,3,4,3,4,3,4,3,4,3,4,3,4,...
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    - 1. First element of subsequence is  $y_0 = x_0$ .
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#### 3. <u>Inductive case d > 1:</u>

- 1. Inductively pick infinite subsequence X' such that the length (d-1) prefix vectors are nondecreasing.
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- 4. For condition (2), suppose that min(A) is infinite; put them in any order to make an infinite sequence.
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**Dickson's Lemma**: (1) Every infinite sequence  $(\mathbf{x}_0, \mathbf{x}_1, ...)$  of vectors in  $\mathbb{N}^d$  has an infinite nondecreasing subsequence, and (2) every set  $A \subseteq \mathbb{N}^d$  has a finite number of minimal elements.

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**Corollary**: The <u>stable configurations are closed downwards</u>: for all stable **c** and all  $\mathbf{b} \leq \mathbf{c}$ , **b** is also stable.

### Upper cones

**Definition**: For all  $\mathbf{c} \in \mathbb{N}^d$ , let  $\nabla(\mathbf{c}) = \{ \mathbf{d} \in \mathbb{N}^d \mid \mathbf{c} \leq \mathbf{d} \}$  denote the <u>upper cone</u> of  $\mathbf{c}$ .



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- 6. Thus  $\mathbf{x} \in \nabla(\mathbf{m}) \subseteq C$ , so  $U \subseteq C$ . **QED**



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**Proof**: By picture.  $\tau = 6$ ,  $\mathbf{c}(S) = 6$ ,  $\mathbf{d}(S) = 8$ . If **c** is not already in a cone  $\nabla(\mathbf{m})$  defining the unstable configurations *U*, we cannot enter any cone by adding more *S*.



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Proof:

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=  $2(z-x)^2$ , which contradicts  $x \neq z$ . **QED** 



# Limits of *efficient* stable computation

## Predicates

Boolean combination of detection predicates

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## Functions

 $\mathbb{N}$ -linear functions (coefficients are nonnegative integers)

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 $\varphi(a,b,c) = a > 0 \text{ OR } (b > 0 \text{ AND } c = 0)$ 

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## **Functions**

N-linear functions (coefficients are nonnegative integers)

e.g., f(a,b) = 2a + 3b $a \rightarrow y+y$  $b \rightarrow y+y+y$ 

## Both computable in O(log n) time

[Angluin, Aspnes, Eisenstat, Fast computation by population protocols with a leader, *DISC* 2006] [Chen, Doty, Soloveichik, Deterministic function computation with chemical reaction networks, *DNA* 2012]

## Known time lower bounds: leader election/majority

## **Leader election**

Leader election (computing the constant function f(a)=1) requires  $\Omega(n)$  time

## **Majority (and other "explicit" predicates)**

Majority (and many other "explicit" predicates such as equality) require  $\Omega(n / \text{polylog } n)$  time, even with up to  $\frac{1}{2} \log \log n$  states.\*

If the protocol satisfies a technical condition called "output dominance", then even with up to log *n* states,  $\Omega(n^{0.999})$  time is required.\*\*

[Doty, Soloveichik, Stable leader election in population protocols requires linear time, DISC 2015]

\*[Alistarh, Aspnes, Eisenstat, Gelashvili, Rivest, SODA 2017]

\*\*[Alistarh, Aspnes, Gelashvili, SODA 2018]: "output dominance"
= changing positive counts of states in a stable configuration
leaves it able to reach a stable configuration with the same output

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  - Both definitions allow exceptions "near a face of  $\mathbb{N}^{k''}$
- <u>Formal theorem</u>: Every predicate that is not eventually constant, and every function that is not eventually  $\mathbb{N}$ -linear, requires at least time  $\Omega(n)$  to compute.
  - They're all computable in at most O(n) time, so this settles their time complexity.

	Predicates	Functions
computable in <mark>O(log n)</mark> time	<u>detection</u> (constant unless changing between 0 and positive) <i>a</i> >0 <b>AND</b> ( <i>b</i> >0 <b>OR</b> <i>c</i> =0)	$\frac{N-linear}{3a+b+2c}$
not computable in less than $\Omega(n)$ time	non-eventually constant a>b? a=b? a is odd?	non-eventually $\mathbb{N}$ -linear $a/2$ $a-b$ $a+1$ $a-1$ 1min( $a,b$ )max( $a,b$ )max( $a$ , min( $b$ + 3, 2 $c$ )) - $c$ - 1
unknown (best known protocol is <i>O</i> ( <i>n</i> ) time)	<u>eventually constant but not</u> <u>constant on <i>all</i> positive values</u> <i>a</i> >1?	eventually N-linear but not N-linear $f(a) = \begin{cases} a \text{ if } a > 1, \\ 0 \text{ otherwise} \end{cases} \begin{cases} f(a) 5 \\ 4 \\ 3 \\ 2 \\ 1 \\ 0 \end{cases}$

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# Other modeling choices?

Modeling choices in formalizing "Computing with chemistry"

- **integer** counts ("stochastic") or **real** concentrations ("mass-action")?
- what is the object being "computed"?
  - yes/no decision problem? "number of A's > number of B's?"
  - numerical function? "m
- *"make Y become double the amount of X"*
- first part of slides
- guaranteed to get correct answer? or allow small probability of error?
  - if Pr[error] = 0, system works *no matter the reaction rates*
- to represent an input  $n_1, ..., n_k$ , what is the initial configuration?
  - only input species present
  - auxiliary species can be present?
- when is the computation finished? when...
  - the output stops changing? (convergence)
  - the output becomes unable to change? (stabilization)
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summarized in next few slides

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But *fundamental computability* doesn't change: exactly the semilinear predicates/functions can be computed (same as without a leader).

[Angluin, Aspnes, Diamadi, Fischer, Peralta, *PODC* 2004] [Angluin, Aspnes, Eisenstat, *PODC* 2006] [Chen, Doty, Soloveichik, *DNA* 2012] [Doty, Hajiaghayi, *DNA* 2013]
#### <u>Convergence vs stabilization</u> and <u>leader vs anarchy</u>









**Theorem**: Without a leader, all non-eventually constant predicates and non-eventually- $\mathbb{N}$ -linear functions require at least  $\Omega(n)$  stabilization time. [Belleville, Doty, Soloveichik, *ICALP* 2017]



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[resolved negatively by Kosowski, Uznański, Population Protocols are Fast, PODC Brief Announcement 2018]

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[Soloveichik, Cook, Bruck, Winfree, Natural Computing 2008]

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**Conjecture**: *Even without a leader,* any computable function can be efficiently computed with high probability.

# What if we use real-valued concentrations... **and** allow reaction rates to influence outcome??

# **Theorem**: A function is computable by a real-valued chemical reaction network using mass-action kinetics if and only if it is computable by <u>any algorithm whatsoever</u>.

[Fages, Le Guludec, Bournez, Pouly. Strong Turing completeness of continuous chemical reaction networks and compilation of mixed analog-digital programs. <u>Computational Methods in Systems Biology – CMSB</u> 2017]

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mass-action kinetics:

$$[X] = -k_1[X] + k_2[Y][Z]$$

$$X \xrightarrow{k_1} Y + Y \qquad [Y] = 2k_1[X] - k_2[Y][Z]$$

$$Y + Z \xrightarrow{k_2} X \qquad [Z] = -k_2[Y][Z]$$

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$$[X] = -k_1[X] + k_2[Y][Z]$$

$$\begin{array}{l} X \stackrel{k_1}{\to} Y + Y \\ Y + Z \stackrel{k_2}{\to} X \end{array} \qquad \begin{bmatrix} \bullet \\ Y \end{bmatrix} = 2k_1[X] - k_2[Y][Z] \\ \hline \\ Z \end{bmatrix} = -k_2[Y][Z] \end{array}$$

# ... with only a polynomial-time slowdown.

[Bournez, Graça, Pouly. *Polynomial time corresponds to solutions of polynomial ordinary differential equations of polynomial length*. Journal of the ACM 2017]

# Fast approximate division by 2

initial configuration:
{ n X, εn A, εn B }

 $\begin{array}{c} X + A \rightarrow B + Y \\ X + B \rightarrow A \end{array}$ 

<u>guaranteed</u> to get  $Y = n/2 \pm \varepsilon n$ E[time] = O(log n) /  $\varepsilon$ 

[Belleville, Doty, Soloveichik, Hardness of computing and approximating predicates and functions with leaderless population protocols, <u>ICALP</u> 2017]

# Fast approximate division by 2

 $n = 100 \quad \varepsilon = 0.1$ 



[Belleville, Doty, Soloveichik, Hardness of computing and approximating predicates and functions with leaderless population protocols, <u>ICALP</u> 2017]

# CRN computation with a small chance of error

r s t






















6) inc s





























HALT



Finite state machine with a fixed number of counters c<sub>1</sub>, c<sub>2</sub>, ..., c<sub>k</sub>, each holding a nonnegative integer.

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- may also have accept/reject semantics, or interpret the final value of some counter as the output

input a

- **1. if** a=0 **goto** 6
- 2. dec a
- 3. inc b
- **4. inc** b
- 5. goto 1
- 6. end

input a f(a) = 2a
1. if a=0 goto 6
2. dec a
3. inc b
4. inc b
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1. if a=0 goto 6
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6. end
1. while a>0:
2. <instruction>

3. <instruction>

```
...
i. ...
```

is a shorthand for

```
1. if a=0 goto i
```

- 2. <instruction>
- 3. <instruction>

... i-1. goto 1 i. ...

f(a) = 2a
<b>oto</b> 6

i	n	р	u	t	а
-		-		-	

1.	while	a>0:
2.	dec	а
3.	dec	а
4.	inc	b

<b>1. while</b> a>0:
<pre>2. <instruction></instruction></pre>
<pre>3. <instruction></instruction></pre>
i
is a shorthand for
1. if a=0 goto i
2. <instruction></instruction>
3. <instruction></instruction>
•••
i-1. goto 1
i

input a $f(a) = 2a$		
1.	if a=0	) <b>goto</b> 6
2.	dec	а
3.	inc	b
4.	inc	b
5.	goto 1	
6.	end	

input a		$f(a) = \lfloor a/2 \rfloor$
1.	while	a>0:
2.	dec	а
3.	dec	а
4.	inc	b

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<pre>2. <instruction></instruction></pre>
3. <instruction></instruction>
•••
i
is a shorthand for
1. if a=0 goto i
2. <instruction></instruction>
3. <instruction></instruction>
i-1. goto 1
i

input a	<i>f</i> ( <i>a</i> ) = 2 <i>a</i>
<b>1. if</b> a=	=0 <b>goto</b> 6
2. dec	a a
3. ind	: b
4. ind	: b
5. goto	1
6. end	

in	put a	$f(a) = \lfloor a/2 \rfloor$
1.	while	a>0:
2.	dec	a
3.	dec	a
4.	inc	b

#### input a

- 1. if a=0 goto 7
- 2. dec a
- 3. if a=0 goto 6
- **4. dec** a
- 5. goto 1
- 6. accept
- 7. reject

#### 1. while a>0: 2. <instruction> 3. <instruction> ... i. ... is a shorthand for 1. if a=0 goto i 2. <instruction> 3. <instruction> ... i-1. goto 1 i. ...

input a	f(a) = 2a
<b>1. if</b> a=6	<b>goto</b> 6
2. dec	a
3. inc	b
4. inc	b
5. goto 1	L
6. end	

inp	out a	$f(a) = \lfloor a/2$
1.	while	a>0:
2.	dec	а
3.	dec	а
4.	inc	b

input a  $\varphi(a) = a is odd$ 

- **1. if** a=0 **goto** 7
- **2. dec** a
- 3. if a=0 goto 6
- **4. dec** a
- 5. goto 1

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7. reject

```
1. while a>0:
2. <instruction>
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...
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1. if a=0 goto i
2. <instruction>
3. <instruction>
...
i-1. goto 1
i. ...
```

input a		f(a) = 2a
1.	if a=0	) <b>goto</b> 6
2.	dec	а
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input a	$f(a) = \lfloor a/2 \rfloor$
1. while	a>0:
2. dec	а
3. dec	а
4. inc	b

#### inputs a,b

1.	while a:	»0:
2.	<b>dec</b> a	
3.	while	b>0:
4.	dec	b
5.	inc	С
6.	inc	d
7.	while	c>0:
8.	dec	С
9.	inc	b

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- **1. if** a=0 goto 7
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7. reject

in	out a	f(a) = 2a
1.	if a=0	) <b>goto</b> 6
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5.	goto 1	
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**1. while** a>0: <instruction> 2. 3. <instruction> ••• i. ... is a shorthand for 1. if a=0 goto i 2. <instruction> 3. <instruction> ••• i-1. goto 1 i. ...

input a	$f(a) = \lfloor a/2 \rfloor$
1. while	a>0:
2. dec	а
3. dec	а
4. inc	b

input a $\varphi(a) = a$ is odd
<b>1. if</b> a=0 <b>goto</b> 7
2. dec a
<b>3. if</b> a=0 <b>goto</b> 6
<b>4. dec</b> a
5. goto 1
6. accept
7. reject

innut a

inputs a,b	f(a,b) = ab
1. while a	>0:
2. dec a	
3. while	b>0:
4. dec	b
5. inc	С
6. inc	d
7. while	c>0:
8. dec	С
9. inc	b

input	ta	f(a) = 2a
1. if	a=0 g	<b>goto</b> 6
2.	<b>dec</b> a	
3.	<b>inc</b> b	
4.	<b>inc</b> b	
5. go	<b>to</b> 1	
6. en	d	

1. while a>0: 2. <instruction> 3. <instruction> ... i. ... is a shorthand for 1. if a=0 goto i 2. <instruction> 3. <instruction> ... i-1. goto 1 i. ...

in	out a	$f(a) = \lfloor a/2 \rfloor$
1.	while	a>0:
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4.	inc	b

inp	uts a,b	f(a,b) = ab
1. v	while ax	»0:
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3. if a=0 goto 6
4. dec a
5. goto 1
6. accept
7. reject

#### input a

1.	inc b	
2.	<pre>while a&gt;</pre>	•0:
3.	dec a	
4.	while	b>0:
5.	dec	b
6.	inc	С
7.	inc	С
8.	while	c>0:
9.	dec	С
10	. inc	b

in	out a	<i>f</i> ( <i>a</i> ) = 2 <i>a</i>
1.	<b>if</b> a=0	<b>goto</b> 6
2.	dec	а
3.	inc	b
4.	inc	b
5.	goto 1	
6.	end	

1. while a>0: 2. <instruction> 3. <instruction> ... i. ... is a shorthand for 1. if a=0 goto i 2. <instruction> 3. <instruction> ... i-1. goto 1 i. ...

in	put a	$f(a) = \lfloor a/2 \rfloor$
1.	while	a>0:
2.	dec	a
3.	dec	а
4.	inc	b

in	puts a,b	f(a,b) = ab
1.	while a:	>0:
2.	<b>dec</b> a	
3.	while	b>0:
4.	dec	b
5.	inc	с
6.	inc	d
7.	while	c>0:
8.	dec	с
9.	inc	b

input a	$\varphi(a)$ = "a is odd"	
1. if a=0	goto 7	
2. dec a 3. if a=0	goto 6	
<b>4. dec</b> a		
5. goto 1 6. accept		
7. reject		

input a		$f(a) = 2^a$
<b>1.</b> i	nc b	
2. w	hile a:	>0:
3.	dec a	
4.	while	b>0:
5.	dec	b
6.	inc	С
7.	inc	С
8.	while	c>0:
9.	dec	С
10.	inc	b

## 3-counter machines are Turing universal
Assume Turing machine

- has a single blank on rightmost cell
- if rightmost blank overwritten, it grows a *new* blank cell to right



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Need a third "work" counter *c* to help do the following operations on counters *a* and *b*:

Turing machine operation	Counter machine implementation
read bit under tape head	
change bit under tape head	
move tape head right	
move tape head left	
test if tape head is on blank and if so, change it to 1	

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move tape head left	
test if tape head is on blank and if so, change it to 1	

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test if tape head is on blank and if so, change it to 1	if <b>b</b> =1 then set <b>a</b> = 2 <b>a</b> + 1

Interpret tape on each side of tape head as binary number; append new leading 1 to make this mapping 1-1, in case the binary string has no leading 1 already, since  $00111_2$ ,  $0111_2$ , and  $111_2$  are all considered the number 7.

1-counter machines are not Turing-universal... why?

[Minsky 1967, Computation: Finite and Infinite Machines]

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- To start with a = n and b = c = 0, start with  $x = 2^n \cdot 3^0 \cdot 5^0 = 2^n$ .
- If *f*: N → N is any computable function, this machine can start with *x*=2<sup>n</sup> and halt with *x*=2<sup>*f*(n)</sup>.

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- <u>Caveat about encoding</u>: there is no 2-counter machine that starts with x=n and halts with  $x=2^n$ . [Schroeppel 1972, A Two Counter Machine Cannot Calculate  $2^N$ ]

"<u>Theorem</u>: Any counter machine can be simulated by a 2-counter machine, provided an obscure coding is accepted for the input and output."

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• 2-counter machines can do universal computation on *encoded* inputs (*n* encoded as 2<sup>*n*</sup>), but they *cannot compute the encoding/decoding* themselves.

[Minsky 1967, Computation: Finite and Infinite Machines]

- To represent counter values (a,b,c) in a single counter x, let  $x = 2^{a} \cdot 3^{b} \cdot 5^{c}$  and y = 0.
  - To increment *b*, set *x* = 3*x*. (using *y* as a work counter)
  - To decrement a, set  $x = \lfloor x/2 \rfloor$ .
  - To test if c = 0, test if  $x \equiv 0 \mod 5$ .
- To start with a = n and b = c = 0, start with  $x = 2^n \cdot 3^0 \cdot 5^0 = 2^n$ .
- If *f*: N → N is any computable function, this machine can start with *x*=2<sup>n</sup> and halt with *x*=2<sup>*f*(n)</sup>.
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- 2-counter machines can do universal computation on *encoded* inputs (*n* encoded as 2<sup>n</sup>), but they *cannot compute the encoding/decoding* themselves.
- However, the fact that 2-counter machines can simulate arbitrary 3-counter machines implies that the Halting Problem for 2-counter machines is undecidable.

#### 2-counter machines: Finite automata robots on the plane



Finite automaton occupying a point  $(x,y) \in \mathbb{N}^2$ .

It cannot write anything, or see anything.

It can sense if it is touching the southern wall, or western wall (or both).

It can move north, south, east, or west based on its current state and 2 "wall bits", and of course change state:

$$\delta: S \times \{ \text{wall, no wall} \}^2 \rightarrow S \times \{ \uparrow, \psi, \leftarrow, \rightarrow \}$$

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There is an automaton A so that this problem is undecidable: given  $(x,y) \in \mathbb{N}^2$ , if started at (x,y), will A ever visit the lower-left corner?

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Error occurs when R is present, but reaction  $L_2 \rightarrow L_1$  occurs instead of  $L_2 + R \rightarrow L_3$ . Semantic effect on register machine: when r > 0, it may jump from line 2 to 1 without decrementing. There's a positive probability of error; how to reduce it? Need to slow down  $L_2 \rightarrow L_1$ .

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Problem with adjusting rate constant to slow down reactions for achieving Turing-universal computation

#### Could make rate constant k very small

- If correct reaction  $r_c: L_2 + R \rightarrow L_3$  has rate constant 1, how small should k be to achieve  $\Pr[r_i \text{ occurs instead of } r_c] = \Pr[\text{error}] = \epsilon$ ?
- rate of  $r_c = \lambda_c = \#L_2 \cdot \#R/v = \#R/v \ge 1/v$
- rate of  $r_i = \lambda_i = k \cdot \#L_2 = k$
- $Pr[error] = \lambda_i / (\lambda_i + \lambda_c) \le k / (k + 1/v)$
- For  $Pr[error] = \varepsilon$ , set  $k = \varepsilon / (v v\varepsilon) \approx \varepsilon / v$

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  - To handle Problem 4, see [Soloveichik, Cook, Winfree, Bruck, *Computation with Finite Stochastic Chemical Reaction Networks*, <u>NaCo</u> 2008]

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E[time for  $L_2 + R \rightarrow L_3] \le n$ 

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Problem 1: Now all rate constants = 1.

 $r_i: C_k + L_2 → C_1 + L_1 + B$   $r_c: L_2 + R → L_3 + B$ So Pr[ $r_i$  ever occurs when it shouldn't] ≤  $\sum_{n=1}^{\infty} 1/n^2 = \pi^2/6$ .

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Still not a great probability bound, but we can scale that to any constant error probability  $\varepsilon$  by setting starting value of *B*:

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Two competing reactions,  $r_i$  incorrect, and  $r_c$  correct:  $r_i: C_k + L_2 \rightarrow C_1 + L_1$   $r_c: L_2 + R \rightarrow L_3$ If both possible, worst case is #R=1, whereas #C\_k = 0 or 1.  $\Pr[r_i] = \Pr[\#C_k = 1] \le 1/n^k$ , where n = #B. Setting k = 2,  $\Pr[r_i] \le 1/n^2$ . Solution: increase *B* after every decrement and jump:  $r_i: C_k + L_2 \rightarrow C_1 + L_1 + B$   $r_c: L_2 + R \rightarrow L_3 + B$ So  $\Pr[r_i \text{ ever occurs when it shouldn't}] \leq \sum_{n=1}^{\infty} 1/n^2 = \pi^2/6$ . Still not a great probability bound, but we can scale that to

any constant error probability  $\varepsilon$  by setting starting value of *B*: For  $\varepsilon = 1/100$ , set initial #B = 102, since  $\sum_{n=102}^{\infty} 1/n^2 < 0.01$ .

Recall three problems we claimed we would solve:

- 1. Adjusting rate constants means designing new chemicals.
- 2. Pr[error in <u>any</u> time step] increases for longer computations.
- 3. Reducing error slows down the computation "significantly".

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<u>Problem 3</u>: Also solved! i.e., halving error probability no longer doubles computation time (*derivation not shown*)