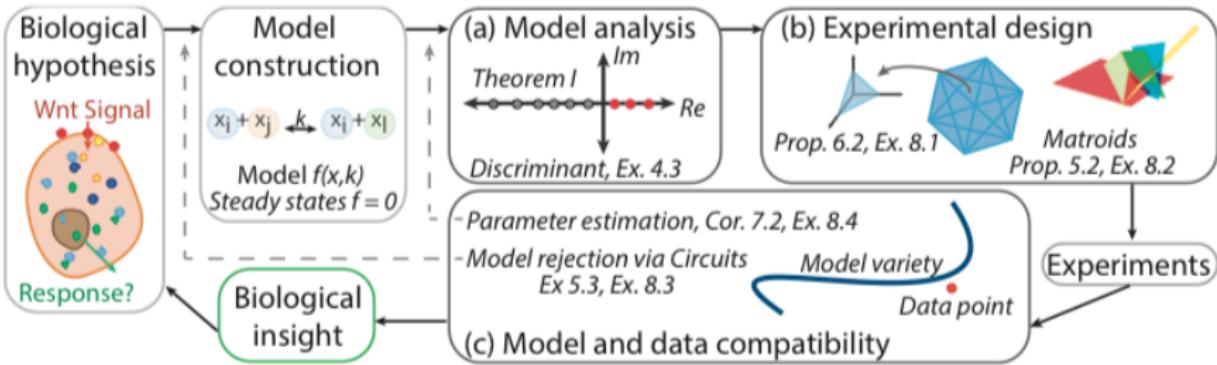


# IDENTIFIABILITY OF LINEAR COMPARTMENT MODELS

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ICERM  
15 November 2018



From *Algebraic Systems Biology: A Case Study for the Wnt Pathway*

(Elizabeth Gross, Heather Harrington, Zvi Rosen, Bernd Sturmfels 2016).

# OUTLINE

- ▶ Introduction: Linear compartment models
- ▶ Identifiability (via differential algebra)
- ▶ The singular locus

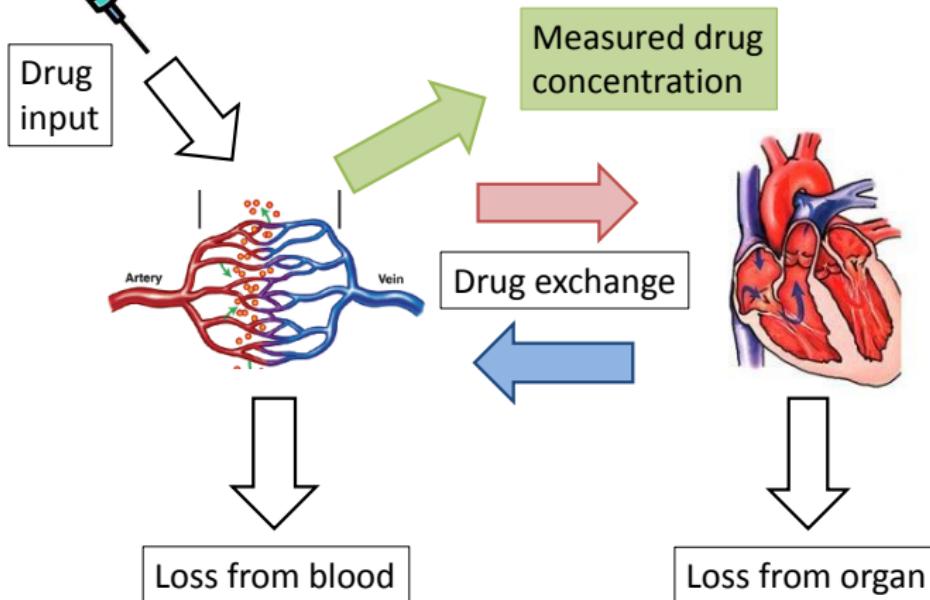
Joint work with

*Elizabeth Gross, Heather Harrington, and Nicolette Meshkat*

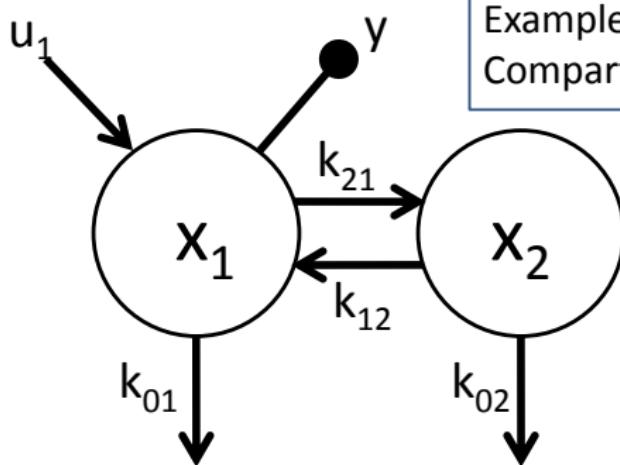
arXiv:1709.10013 and arXiv:1810.05575

# INTRODUCTION

# Motivation: biological models



# COMPARTMENT MODEL



$$\begin{aligned}\dot{x}_1 &= -(k_{01} + k_{21})x_1 + k_{12}x_2 + u_1 \\ \dot{x}_2 &= k_{21}x_1 - (k_{02} + k_{12})x_2 \\ y &= x_1\end{aligned}$$

*Structural identifiability:* Recover parameters  $k_{ij}$  from perfect input-output data  $u_1(t)$  and  $y(t)$ ? (Bellman & Astrom 1970)

# IDENTIFIABILITY VIA DIFFERENTIAL ALGEBRA<sup>1</sup>:

*Which models are identifiable?*

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<sup>1</sup>Ljung and Glad 1994

# INPUT-OUTPUT EQUATIONS

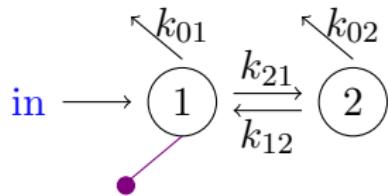
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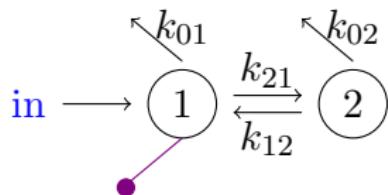
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- ▶ Example, continued:



$$y_1^{(2)} + (k_{01} + k_{02} + k_{12} + k_{21}) y'_1 + (k_{01}k_{12} + k_{01}k_{02} + k_{02}k_{21}) y_1 = (k_{02} + k_{12}) u_1$$

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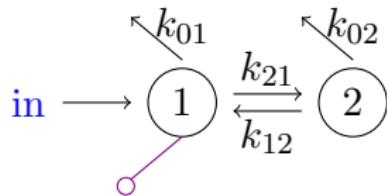


$$y_1^{(2)} + (k_{01} + k_{02} + k_{12} + k_{21}) y'_1 + (k_{01}k_{12} + k_{01}k_{02} + k_{02}k_{21}) y_1 = (k_{02} + k_{12}) u_1$$

- ▶ *Input-output equations* come from the elimination ideal:  
⟨ differential eqns., output eqns.  $y_i = x_j$  , their  $m$  derivatives ⟩

$$\cap \mathbb{C}(k_{ij})[u_i^{(k)}, y_i^{(k)}]$$

# INPUT-OUTPUT EQUATIONS, CONTINUED

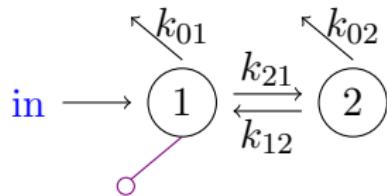


$$A = \begin{pmatrix} -k_{01} - k_{21} & k_{12} \\ k_{21} & -k_{02} - k_{12} \end{pmatrix} \quad x'(t) = Ax(t) + u(t)$$

- ▶ **Proposition** (Meshkat, Sullivant, Eisenberg 2015):  
For a linear compartment model with input and output in compartment-1 only, the **input-output equation** is:

$$\det(\partial I - A)y_1 = \det((\partial I - A)_{11}) u_1 .$$

## INPUT-OUTPUT EQUATIONS, CONTINUED



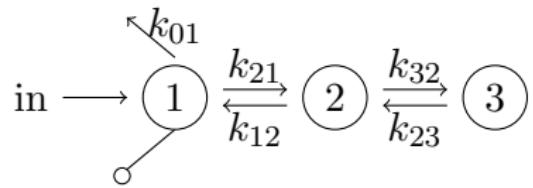
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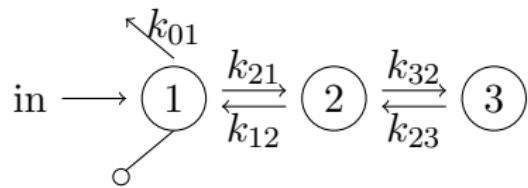
$$\det(\partial I - A)y_1 = \det((\partial I - A)_{11}) u_1 .$$

- ▶ Proof uses Cramer's Rule and Laplace expansion

## INPUT-OUTPUT EQUATIONS, CONTINUED



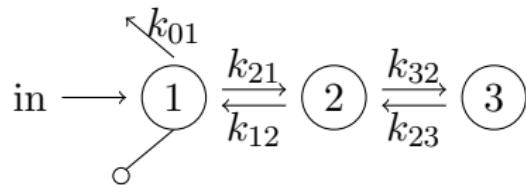
## INPUT-OUTPUT EQUATIONS, CONTINUED



$$\det(\partial I - \textcolor{red}{A}) \textcolor{violet}{y}_1 = \det ((\partial I - \textcolor{red}{A})_{11}) \textcolor{blue}{u}_1$$

$$\begin{aligned} & \det \begin{pmatrix} d/dt + k_{01} + k_{21} & -k_{12} & 0 \\ -k_{21} & d/dt + k_{12} + k_{32} & -k_{23} \\ 0 & -k_{32} & d/dt + k_{23} \end{pmatrix} y_1 \\ &= \det \begin{pmatrix} d/dt + k_{12} + k_{32} & -k_{23} \\ -k_{32} & d/dt + k_{23} \end{pmatrix} u_1 \end{aligned}$$

## INPUT-OUTPUT EQUATIONS, CONTINUED



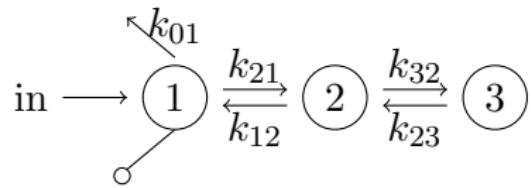
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... expands to the *input-output equation*:

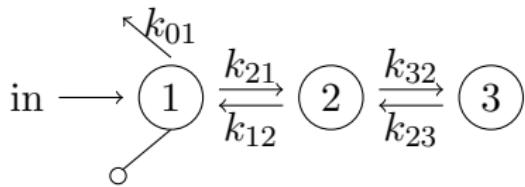
$$\begin{aligned} & y_1^{(3)} + (\textcolor{teal}{k}_{01} + \textcolor{teal}{k}_{12} + \textcolor{teal}{k}_{21} + \textcolor{teal}{k}_{23} + \textcolor{teal}{k}_{32}) y_1^{(2)} \\ &+ (\textcolor{teal}{k}_{01} \textcolor{teal}{k}_{12} + \textcolor{teal}{k}_{01} \textcolor{teal}{k}_{23} + \textcolor{teal}{k}_{01} \textcolor{teal}{k}_{32} + \textcolor{teal}{k}_{12} \textcolor{teal}{k}_{23} + \textcolor{teal}{k}_{21} \textcolor{teal}{k}_{23} + \textcolor{teal}{k}_{21} \textcolor{teal}{k}_{32}) y_1' + (\textcolor{teal}{k}_{01} \textcolor{teal}{k}_{12} \textcolor{teal}{k}_{23}) y_1 \\ &= \textcolor{teal}{u}_1^{(2)} + (\textcolor{teal}{k}_{12} + \textcolor{teal}{k}_{23} + \textcolor{teal}{k}_{32}) \textcolor{teal}{u}_1' + (\textcolor{teal}{k}_{12} \textcolor{teal}{k}_{23}) \textcolor{teal}{u}_1 . \end{aligned}$$

# COEFFICIENTS OF INPUT-OUTPUT EQUATIONS



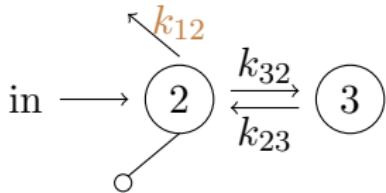
$$\begin{aligned} & y_1^{(3)} + (k_{01} + k_{12} + k_{21} + k_{23} + k_{32}) y_1^{(2)} \\ & + (k_{01}k_{12} + k_{01}k_{23} + k_{01}k_{32} + k_{12}k_{23} + k_{21}k_{23} + k_{21}k_{32}) y'_1 + (k_{01}k_{12}k_{23}) y_1 \\ & = u_1^{(2)} + (k_{12} + k_{23} + k_{32}) u'_1 + (k_{12}k_{23}) u_1 . \end{aligned}$$

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- ▶ coefficient of  $y_1^{(i)}$  corresponds to forests with  $(3 - i)$  edges and  $\leq 1$  outgoing edge per compartment
- ▶ coefficient of  $u_1^{(i)}$  corresponds to  $(n - i - 1)$ -edge forests:



- ▶ **Thm 1:** The coefficients correspond to forests in model.

# IDENTIFIABILITY

$$\begin{aligned} & y_1^{(3)} + (\textcolor{teal}{k_{01}} + \textcolor{teal}{k_{12}} + k_{21} + k_{23} + \textcolor{teal}{k_{32}}) y_1^{(2)} \\ & + (\textcolor{teal}{k_{01}k_{12}} + \textcolor{teal}{k_{01}k_{23}} + k_{01}k_{32} + \textcolor{teal}{k_{12}k_{23}} + k_{21}k_{23} + \textcolor{teal}{k_{21}k_{32}}) y'_1 + (\textcolor{teal}{k_{01}k_{12}k_{23}}) y_1 \\ & = u_1^{(2)} + (\textcolor{teal}{k_{12}} + k_{23} + \textcolor{teal}{k_{32}}) u'_1 + (\textcolor{teal}{k_{12}k_{23}}) u_1 . \end{aligned}$$

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$$\mathbb{R}^5 \rightarrow \mathbb{R}^5$$

$$(k_{01}, k_{12}, k_{21}, k_{23}, k_{32}) \mapsto (\textcolor{teal}{k_{01}} + \textcolor{teal}{k_{12}} + \textcolor{teal}{k_{21}} + \textcolor{teal}{k_{23}} + \textcolor{teal}{k_{32}}, \dots)$$

- ▶ Solve directly, or use ...
- ▶ **Proposition** (Meshkat, Sullivant, Eisenberg 2015):  
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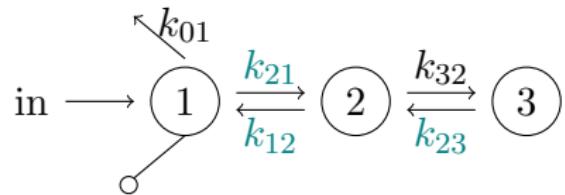
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- ▶ Solve directly, or use ...
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Identifiable  $\Leftrightarrow$  Jacobian matrix of coefficient map has (full) rank = number of parameters *generically*

## THE SINGULAR LOCUS

# DEFINITION

- ▶ Focus on the *non-identifiable* parameters:  
the **singular locus** is where the Jacobian matrix of coefficient map is rank-deficient.
- ▶ Example, continued:



The equation of the singular locus is:

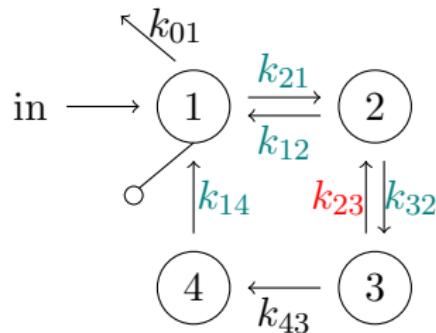
$$\det \text{Jac} = k_{12}^2 k_{21} k_{23} = 0 .$$

## IDENTIFIABLE SUBMODELS

- ▶ *Motivation:* drug targets
- ▶ **Thm 2:** Let  $\mathcal{M}$  be an identifiable linear compartment model, with singular-locus equation  $f$ . Let  $\widetilde{\mathcal{M}}$  be obtained from  $\mathcal{M}$  by deleting edges  $\mathcal{I}$ .  
If  $f \notin \langle k_{ji} \mid (i, j) \in \mathcal{I} \rangle$ , then  $\widetilde{\mathcal{M}}$  is identifiable.

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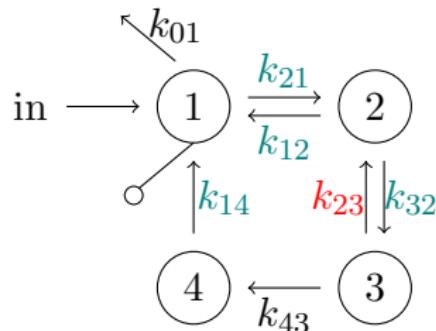
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$$f = k_{12}k_{14}k_{21}^2k_{32}(k_{12}k_{14} - k_{14}^2 - \dots)(k_{12}k_{23} + k_{12}k_{43} + k_{32}k_{43}) .$$

# IDENTIFIABLE SUBMODELS

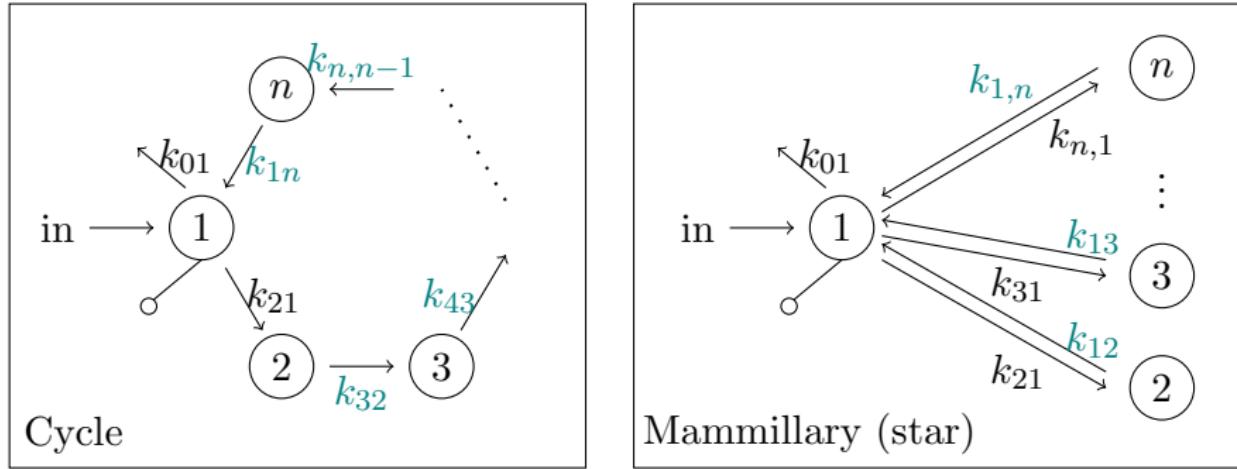
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- ▶ *Converse is false:* deleting  $k_{12}$  and  $k_{23}$  is identifiable!

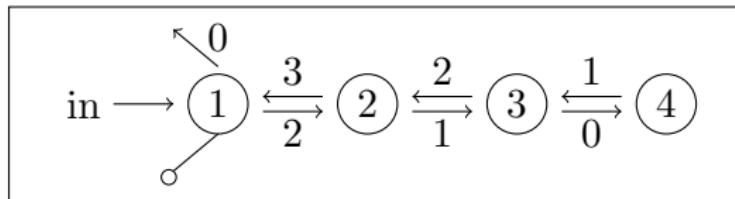
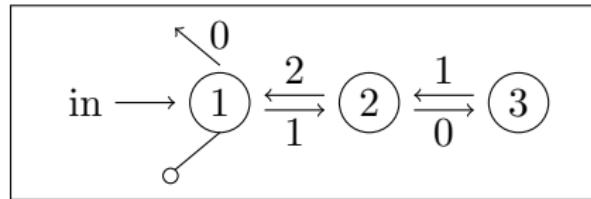
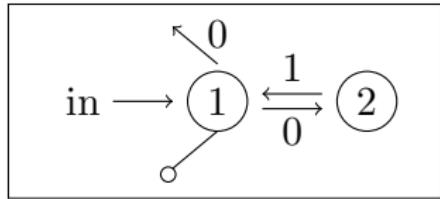
# CYCLE AND MAMMILLARY MODELS



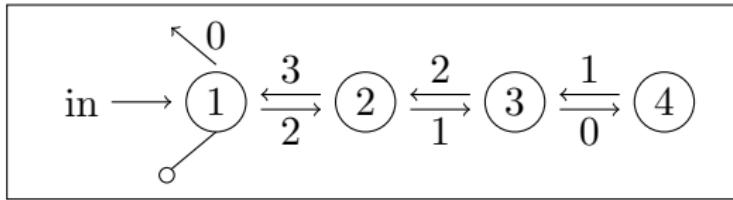
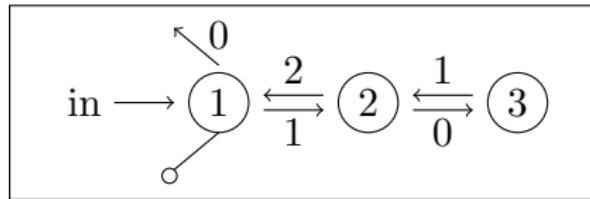
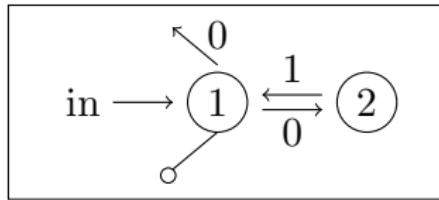
## ► Thm 3:

- The singular-locus equation for the Cycle model is  $k_{32}k_{43}\dots k_{n,n-1}k_{1,n} \prod_{2 \leq i < j \leq n} (k_{i+1,i} - k_{j+1,j})$ .
- The singular-locus equation for the Mammillary model is  $k_{12}k_{13}\dots k_{1,n} \prod_{2 \leq i < j \leq n} (k_{1i} - k_{1j})^2$ .

# CATENARY (PATH) MODELS

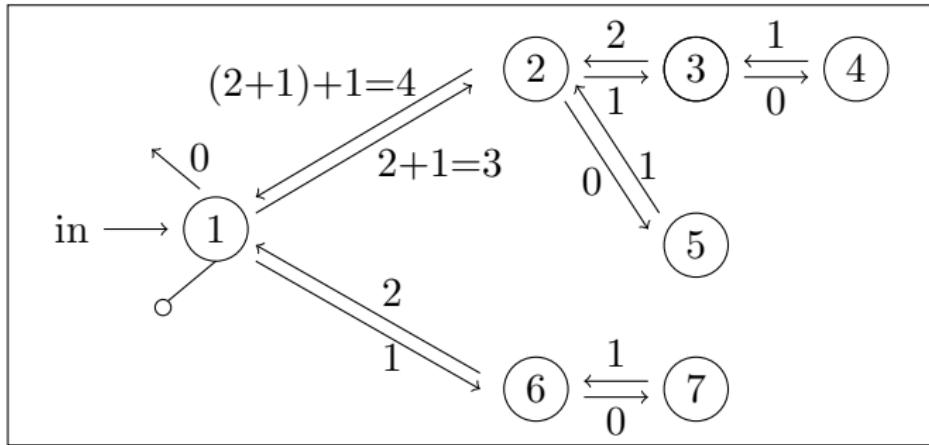
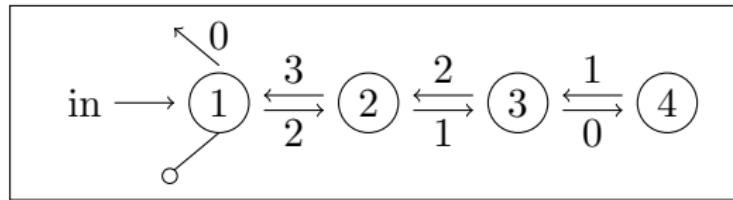


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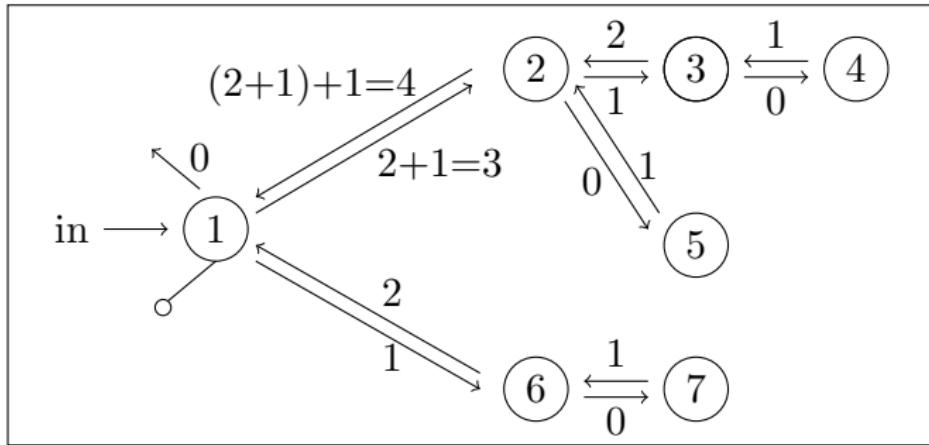
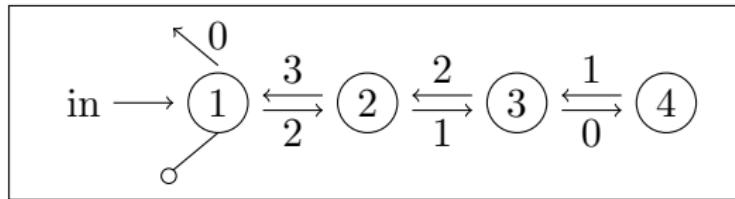


**Conjecture:** For **catenary models**, the exponents in the singular-locus equation generalize the pattern above.

# TREE CONJECTURE



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**Conj.:** (Hoch, Sweeney, Tung) For **tree models**, the exponents in the singular-locus equation generalize the pattern above.

## IDENTIFIABLE SUBMODELS (AGAIN)

- ▶ **Thm 4:** Let  $\widetilde{\mathcal{M}}$  be obtained by:
  - ▶ adding a **leak** to a strongly connected model  $\mathcal{M}$  with *no* leaks, or
  - ▶ deleting the **leak** from a strongly connected model  $\mathcal{M}$  with input, output, and leak in *one* compartment.

Then, if  $\mathcal{M}$  is identifiable, then so is  $\widetilde{\mathcal{M}}$ .

---

<sup>2</sup>Can delete edges *without* making the singular-locus equation = 0.

## IDENTIFIABLE SUBMODELS (AGAIN)

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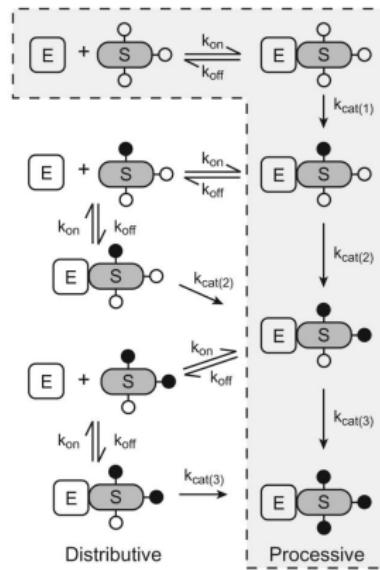
| Operation     | Preserves identifiability?              |
|---------------|---|
| Add input     | Yes                                     |
| Add output    | Yes                                     |
| Add leak      | Not always (and see above)              |
| Add edge      | Not always                              |
| Delete input  | Not always                              |
| Delete output | Not always                              |
| Delete leak   | Open (and see above)                    |
| Delete edge   | Not always (recall Thm 2 <sup>2</sup> ) |

---

<sup>2</sup>Can delete edges *without* making the singular-locus equation = 0.

# FUTURE WORK

## Nonlinear models



From **Processive phosphorylation: mechanism and biological importance**,  
Patwardhan and Miller, *Cell Signal.* 2007.

# SUMMARY

The **singular locus** is an interesting mathematical object that can help us answer the question, *which linear compartment models are identifiable?*

THANK YOU.

# IDENTIFIABILITY DEGREE

- ▶ the **identifiability degree** of a model is the number of parameter vectors that match (generic) input-output data

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- ▶ the **identifiability degree** of a model is the number of parameter vectors that match (generic) input-output data
- ▶ **Proposition** (Cobelli, Lepschy, Romanin Jacur 1979)

| Model             | Identifiability degree |
|-------------------|------------------------|
| Catenary (path)   | 1                      |
| Mammillary (star) | $(n - 1)!$             |

- ▶ **Thm 5**

| Model | Identifiability degree |
|-------|------------------------|
| Cycle | $(n - 1)!$             |