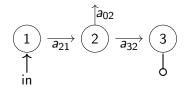
Indistinguishability of Linear Compartmental Models

Cash Bortner* and Nicolette Meshkat†

*California State University, Stanislaus cbortner@csustan.edu

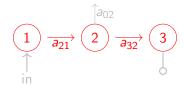
†Santa Clara University

Algebraic Systems Biology SIAM Conference on Applied Algebraic Geometry, 2025 University of Wisconsin, Madison, WI 7/10/2025



$$\mathcal{M} = (G, In, Out, Leak)$$

= $(\overrightarrow{P}_3, \{1\}, \{3\}, \{2\}).$



Directed Graph: $G = \overrightarrow{P}_3$

$$\mathcal{M} = (G, In, Out, Leak)$$

= $(\overrightarrow{P}_3, \{1\}, \{3\}, \{2\}).$



Input Compartment: $In = \{1\}$

$$\mathcal{M} = (G, In, Out, Leak)$$

= $(\overrightarrow{P}_3, \{1\}, \{3\}, \{2\}).$



Output Compartment: $Out = \{3\}$

$$\mathcal{M} = (G, In, Out, Leak)$$

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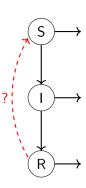
Leak Compartment: $Leak = \{2\}$

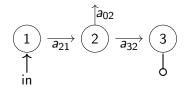
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Compartmental Models in the Wild

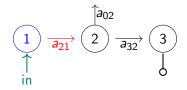
- SIR Model for spread of a virus in Epidemiology (non-linear)
- SIV Model for vaccine efficiency in Epidemiology
- Modeling Pharmacokinetics for absorption, distribution, metabolism, and excretion in the blood
- Modeling different biological systems





$$\mathcal{M} = (G, In, Out, Leak)$$

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Linear Compartmental Model

$$\mathcal{M} = (\textit{G}, \textit{In}, \textit{Out}, \textit{Leak})$$
$$= (\overrightarrow{\textit{P}}_3, \{1\}, \{3\}, \{2\}).$$

ODEs in terms of concentrations $x_i(t)$, input $u_1(t)$, and output $y_3(t)$:

$$\dot{x_1}(t) = -a_{21}x_1(t) + u_1(t)$$

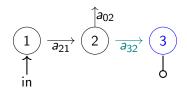


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$$\dot{x_1}(t) = -a_{21}x_1(t) + u_1(t)
\dot{x_2}(t) = a_{21}x_1(t) - (a_{02} + a_{32})x_2(t)$$



Linear Compartmental Model

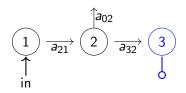
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Linear Compartmental Model

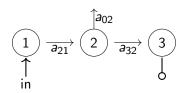
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 $+u_1(t)$
 $\dot{x_2}(t) = a_{21}x_1(t) - (a_{02} + a_{32})x_2(t)$
 $\dot{x_3}(t) = a_{32}x_2(t)$

with

$$y_3(t) = x_3(t)$$
.



Linear Compartmental Model

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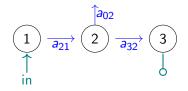
ODEs in terms of concentrations $x_i(t)$, input $u_1(t)$, and output $y_3(t)$:

$$\begin{pmatrix} \dot{x}_1(t) \\ \dot{x}_2(t) \\ \dot{x}_3(t) \end{pmatrix} = \underbrace{\begin{pmatrix} -a_{21} & 0 & 0 \\ a_{21} & -a_{02} - a_{32} & 0 \\ 0 & a_{32} & 0 \end{pmatrix}}_{a_{32}} \begin{pmatrix} x_1(t) \\ x_2(t) \\ x_3(t) \end{pmatrix} + \begin{pmatrix} u_1(t) \\ 0 \\ 0 \end{pmatrix}$$

compartmental matrix A

with

$$y_3(t)=x_3(t).$$

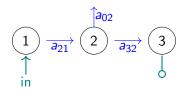


Linear Compartmental Model

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Via an application of Cramer's Rule:

$$\det(\partial I - A)y_3 = \det(\partial I - A)^{1,3} u_1$$



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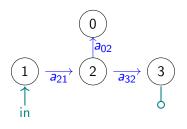
Via an application of Cramer's Rule:

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$$y_3^{(3)} + (a_{21} + a_{02} + a_{32})\ddot{y_3} + (a_{02}a_{21} + a_{21}a_{32})\dot{y_3} = (a_{21}a_{32})u_1.$$

an ODE in only the measurable variables and the parameters:

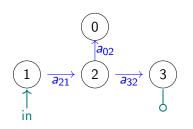
Input/Output Equation



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Linear Compartmental Model

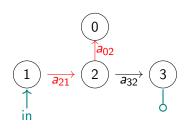
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Theorem (\$, Gross, Meshkat, Shiu, Sullivant [2023])

The coefficients of the input-output equation of a LCM (G, In, Out, Leak) can be generated by *incoming forests* on graphs related to G.

- incoming: no vertex has more than one outgoing edge
- forest: no cycles



Linear Compartmental Model

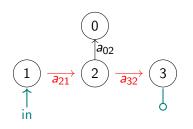
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Motivating Question

How do we know if the model structure we chose which seems to represent the data well is **unique**?

Indistinguishability

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Definition

Two LCMs are *indistinguishable* if for any choice of parameters in the first model, there is a choice of parameters in the second model that will yield the same *dynamics* in both models.

Indistinguishability

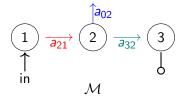
Motivating Question

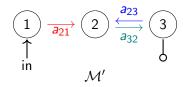
How do we know if the model structure we chose which seems to represent the data well is **unique**?

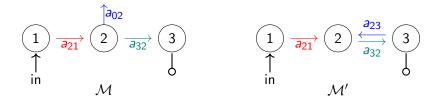
Definition

Two LCMs are *indistinguishable* if for any choice of parameters in the first model, there is a choice of parameters in the second model that will yield the same *dynamics* in both models.

What does "same dynamics" mean?



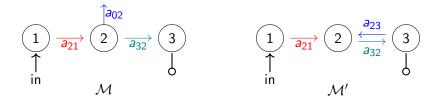




$$\begin{split} \mathcal{M}: y_3^{(3)} + \big(a_{21} + a_{02} + a_{32}\big)\ddot{y_3} + \big(a_{02}a_{21} + a_{21}a_{32}\big)\dot{y_3} &= \big(a_{21}a_{32}\big)u_1. \\ \mathcal{M}': y_3^{(3)} + \big(a_{21} + a_{23} + a_{32}\big)\ddot{y_3} + \big(a_{23}a_{21} + a_{21}a_{32}\big)\dot{y_3} &= \big(a_{21}a_{32}\big)u_1. \end{split}$$

Remark

From the perspective of the input-output equations, we can not *distinguish* between these two very structurally different models.

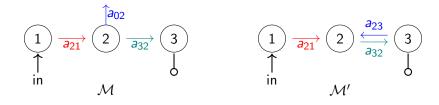


$$\mathcal{M}: y_3^{(3)} + (a_{21} + a_{02} + a_{32})\ddot{y_3} + (a_{02}a_{21} + a_{21}a_{32})\dot{y_3} = (a_{21}a_{32})u_1.$$

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Working Definition

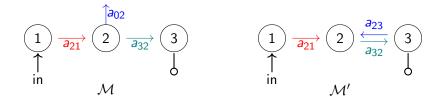
Two models are *permutation indistinguishable* if they have the same input-output equations up to renaming the parameters.



$$\mathcal{M}: y_3^{(3)} + (a_{21} + a_{02} + a_{32})\ddot{y_3} + (a_{02}a_{21} + a_{21}a_{32})\dot{y_3} = (a_{21}a_{32})u_1.$$

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renaming:
$$\begin{pmatrix} a_{21} \\ a_{02} \\ a_{32} \end{pmatrix} \mapsto \begin{pmatrix} a_{21} \\ a_{23} \\ a_{32} \end{pmatrix}$$

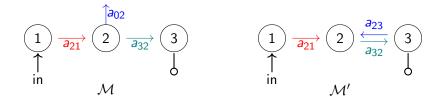


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Remark

Permuation indistinguishability is an equivalence relation!



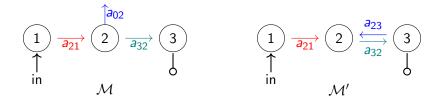
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Question: What is the equivalence class of a model?



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Remark

Permuation indistinguishability is an equivalence relation!

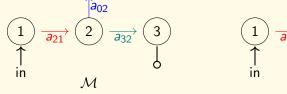
Question: What is the equivalence class of a model? (Size 1?)

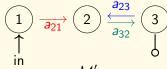
Theorem (Godfrey & Chapman [1990])

Two indistinguishable models must preserve the following:

- 1. The length of the shortest path from the input to the output
- 2. The number of compartments with a path to any output compartment
- 3. The number of compartments that can be reached from an input
- 4. The number of traps*

Example



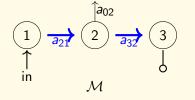


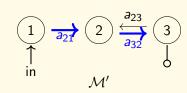
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Example (1. Dist(1,3) = 2 in both!)



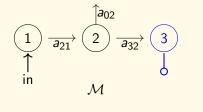


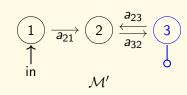
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Example (2. **Two** compartments with a path to the output!)



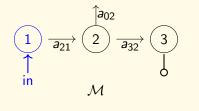


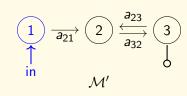
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Example (3. **Two** compartments with a path from the input!)



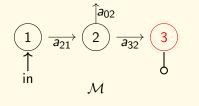


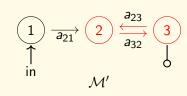
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Example (4. Each model has **one** trap!)





Theorem (Godfrey & Chapman [1990])

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Goal:

Find *sufficient* conditions for permutation indistinguishability of two models based on their graph structures.

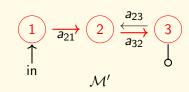
Skeletal Path Models

Definition

A *skeletal path model* is an LCM whose graph contains the directed path $\overrightarrow{P_n}$, i.e $1 \rightarrow 2 \rightarrow \ldots \rightarrow n$, with $In = \{1\}$ and $Out = \{n\}$.

Example





Skeletal Path Moves: Walking the Leak

Question

What *moves* can you perform on a basic skeletal path model resulting in an indistinguishable model?



Skeletal Path Moves: Walking the Leak

Question

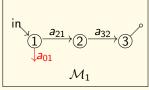
What *moves* can you perform on a basic skeletal path model resulting in an indistinguishable model?

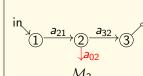
Theorem (\$ & Meshkat [2024]; \$, Gilliana, Patel, & Tamras [2025*])

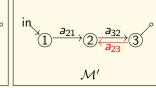
The following skeletal path models are indistinguishable:

- $\mathcal{M}_i = (\overrightarrow{P_n}, \{1\}, \{n\}, \{i\})$ for any $i \in \{1, 2, 3, \dots, n-1\}$
- $\mathcal{M}' = (\overrightarrow{P_n} \cup \{n \rightarrow n-1\}, \{1\}, \{n\}, \emptyset).$

Example







Proof idea

Theorem (\$ & Meshkat [2024]; \$, Gilliana, Patel, & Tamras [2025*])

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Proof idea 1: Left-hand side of the input/output equation of \mathcal{M}_i given by:

$$\det(\partial I - A_i)y_n = \det\begin{pmatrix} \partial + a_{21} & 0 & \cdots & \cdots & 0 & 0 \\ -a_{21} & \ddots & \ddots & \ddots & \ddots & \vdots & \vdots \\ 0 & \ddots & \partial + a_{0i} + a_{i(i-1)} & \ddots & \vdots & \vdots & \vdots \\ \vdots & \ddots & -a_{i(i-1)} & \ddots & 0 & 0 \\ 0 & \cdots & \ddots & \ddots & \partial + a_{n(n-1)} & 0 \\ 0 & \cdots & \cdots & 0 & -a_{n(n-1)} & \partial \end{pmatrix} y_n$$

Proof idea

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Proof idea 2:

• Under a renaming of the parameters, the incoming forests of each \mathcal{M}_i are exactly the same as the incoming forests of each \mathcal{M}_j (and \mathcal{M}').

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Proof idea 2:

- Under a renaming of the parameters, the incoming forests of each \mathcal{M}_i are exactly the same as the incoming forests of each \mathcal{M}_j (and \mathcal{M}').
- Thus, each of the coefficients of the respective input/output equations are indistinguishable.

Detour Models

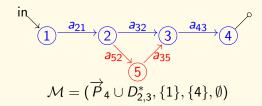
Definition

A detour model is given by

$$\mathcal{M} = (\overrightarrow{P_n} \cup \overrightarrow{D_{i,j}}, \{1\}, \{n\}, Leak)$$

where D is some connected directed graph, and $D_{i,j}^*$ includes one edge from node i and to node j in the skeletal path.

Example



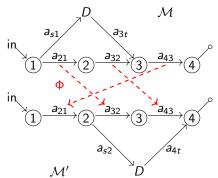
Detour Models

Theorem (\$ & Meshkat [2024])

The following two detour models are indistinguishable:

- $\mathcal{M} = (\overrightarrow{P_n} \cup \overrightarrow{D_{i,j}}, \{1\}, \{n\}, Leak)$
- $\mathcal{M}' = (\overrightarrow{P_n} \cup \overrightarrow{D_{i+1,j+1}}, \{1\}, \{n\}, Leak)$

Proof idea:



 Break the A matrices into blocks, and show equivalent determinants under Φ

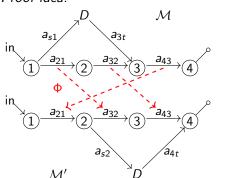
Detour Models

Theorem (\$ & Meshkat [2024])

The following two detour models are indistinguishable:

- $\mathcal{M} = (\overrightarrow{P_n} \cup \overrightarrow{D_{i,j}}, \{1\}, \{n\}, Leak)$
- $\mathcal{M}' = (\overrightarrow{P_n} \cup \overrightarrow{D_{i+1,i+1}}, \{1\}, \{n\}, Leak)$

Proof idea:



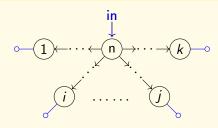
- Break the A matrices into blocks, and show equivalent determinants under Φ
- Or, the incoming forests under the renaming are the same, so the coefficients are the same!

Source and Sink Models

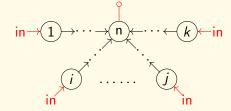
Corollary (\$ & Meshkat [2024]

We can extend results from detour models to source and sink models.

Example



Basic Source Model



Basic Sink Model

Future Work

- These are very specific families of LCMs, but this is a proof of concept moving forward!
- More implementation of graph theory in showing sufficient conditions for other families of models
 - Cycle Models (undergraduate research project)
 - Tree Models
- Look into more general indistinguishability from a graph perspective
- Help biologists determine if the model they are using is the only model which yield the same dynamics

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This Presentation!

