Finding and breaking Lie symmetries: implications for structural identifiability and observability of dynamic models

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BIRS Workshop "Model Theory of Differential Equations, Algebraic Geometry, and their Applications to Modeling" 1–5 June, 2020

Motivation: identifiability and observability in dynamic modelling

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Observability Structural Identifiability as Observability (SIO) Importance for modelling

Lie Symmetries

Lie Symmetries and SIO Finding Lie symmetries

Examples

Discussion and open questions

Lie symmetries:

- Bluman, G.; Anco, S. Symmetry and integration methods for differential equations; Vol. 154, Springer Science & Business Media, 2008.
- Arrigo, D.J. Symmetry analysis of differential equations: an introduction; John Wiley & Sons, 2015.

SIO:

 Villaverde, A.F. "Observability and Structural Identifiability of Nonlinear Biological Systems". *Complexity* Vol. 2019, Article ID 8497093, https://doi.org/10.1155/2019/8497093.

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Observability and Structural Identifiability: the concepts

We consider the following type of dynamic models of ODEs:

$$M_{NL} := \begin{cases} \dot{x}(t) &= f(x(t), \theta, u(t), w(t)) ,\\ y(t) &= g(x(t), \theta, u(t), w(t)) ,\\ x(t_0) &= x^0(\theta) \end{cases}$$

with states $x(t) \in \mathbb{R}^m$, parameters $\theta \in \mathbb{R}^q$, outputs $y(t) \in \mathbb{R}^n$, known inputs $u(t) \in \mathbb{R}^{m_u}$, unknown inputs $w(t) \in \mathbb{R}^{m_w}$, f and g vectors of analytical functions.

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Structural Identifiability

A model is structurally identifiable if it is theoretically possible to infer its parameters, θ , by observing its outputs, y(t)

Structural Identifiability and Observability (SIO)

Structural Local Identifiability as Observability Extend the state vector as:

$$\widetilde{x}(t) = \begin{bmatrix} x(t) \\ \theta \end{bmatrix}, \dot{\widetilde{x}}(t) = \begin{bmatrix} f(\widetilde{x}(t), u(t)) \\ 0 \end{bmatrix} \Rightarrow M_{NL} := \begin{cases} \dot{\widetilde{x}} = \widetilde{f}(\widetilde{x}, u) \\ y = g(\widetilde{x}, u) \end{cases}$$

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Structurally locally Identifiable or Observable (SIO)

A variable (state or parameter) \tilde{x}_i is structurally locally identifiable or observable (SIO) if there is a neighbourhood $V(\tilde{x}_i^*)$ s.t.

$$\hat{\widetilde{x}_i} \in V(\widetilde{x}_i^*)$$
 and $y(\hat{\widetilde{x}_i}) = y(\widetilde{x}_i^*) \Rightarrow \hat{\widetilde{x}_i} = \widetilde{x}_i^*$

Otherwise it is Structurally Unidentifiable or Unobservable (SU).

Why it matters: SU models provide wrong insights







Model of the glucose-insulin system If $y(t) = [\beta, G] \Rightarrow [p, s_i]$ are SU and I is unobservable. c, α, γ , and the product $p \cdot s_i$ are SLI.

LIE SYMMETRIES

Assessing SIO with Lie Symmetries

- ► Existence of Lie symmetries ⇒ existence of similarity transformations¹ ⇒ existence of transformations of x̃ that leave y unchanged: non-observability (SU).
- Similarity transformations are one-parameter Lie group morphisms that map solutions of a differential equation onto themselves.
- Algorithm for finding Lie symmetries using Ansatz polynomials² + some modifications³.

¹Yates, J.W.; Evans, N.D.; Chappell, M.J. Structural identifiability analysis via symmetries of differential equations. *Automatica* 2009, 45, 25852591.

²Merkt, B., Timmer, J., and Kaschek, D. "Higher-order Lie symmetries in identifiability and predictability analysis of dynamic models". *Phys Rev E* 92.1, 2015.

³Massonis, G., and Villaverde, A.F. "Finding and Breaking Lie Symmetries: Implications for Structural Identifiability and Observability in Biological Modelling". Symmetry 12(3):469, 2020.

One-parameter Lie group of transformations:

$$x^* = X(x;\varepsilon)$$
,

We say that:

•
$$\eta(x) = \frac{\partial X(x;\varepsilon)}{\partial \varepsilon}|_{\varepsilon=0}$$
 is an infinitesimal

• X is the infinitesimal generator, $X = X(x) = \sum_{i=1}^{n} \eta_i(x) \frac{\partial}{\partial x_i}$

• $x + \varepsilon \eta(x)$ is the infinitesimal transformation of the Lie group of transformations.

First, augment the state vector x:

$$x := \begin{cases} \dot{x}_i(t) = f_i(x(t), u(t)), & i = 1, ..., m \\ x_i(t) = \theta, & i = m + 1, ..., m + q \\ x_i(t) = w_i(t), & i = m + q + 1, ..., n^* = m + q + m_w. \end{cases}$$

Then, consider different types of polynomial *Ansatz* for the infinitesimals (univariate, partially variate, and multivariate).

Univariate:

$$\eta_i(\mathbf{x}) = \sum_{d=0}^{d_{max}} r_{i,d} x_i^d, \ i = 1, ..., n^*$$

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Methodology

Creation of infinitesimal generators

Partially variate:

$$\begin{split} \eta_i(\mathbf{x}) &= \sum_{\substack{d_i, d_{m+1}, \dots, d_{m+q} = 0}}^{|d| = d_{max}} r_{i,d} x_i^{d_i} x_{m+1}^{d_{m+1}} \cdots x_{m+q}^{d_{m+q}}, \quad i = 1, \dots, m \;, \\ \eta_i(\mathbf{x}) &= \sum_{\substack{d_{m+1}, \dots, d_{m+q} = 0}}^{|d| = d_{max}} r_{i,d} x_{m+1}^{d_{m+1}} \cdots x_{m+q}^{d_{m+q}}, \quad i = m+1, \dots, m+q \;, \\ \eta_i(\mathbf{x}) &= \sum_{\substack{d_i, d_{m+1}, \dots, d_{m+q} = 0}}^{|d| = d_{max}} r_{i,d} x_i^{d_i} x_{m+1}^{d_{m+1}} \cdots x_{m+q}^{d_{m+q}}, \quad i = m+q+1, \dots, n^* \;. \end{split}$$

Multivariate:

$$\eta_{i}(\mathbf{x}) = \sum_{d_{1},...,d_{m+q}=0}^{|d|=d_{max}} r_{i,d} x_{1}^{d_{1}} \cdots x_{m+q}^{d_{m+q}}, \quad i = 1, ..., m ,$$

$$\eta_{i}(\mathbf{x}) = \sum_{d_{m+1},...,d_{m+q}=0}^{|d|=d_{max}} r_{i,d} x_{m+1}^{d_{m+1}} \cdots x_{m+q}^{d_{m+q}}, \quad i = m+1, ..., m+q ,$$

$$\eta_{i}(\mathbf{x}) = \sum_{d_{1},...,d_{n^{*}}=0}^{|d|=d_{max}} r_{i,d} x_{1}^{d_{1}} \cdots x_{n^{*}}^{d_{n^{*}}}, \quad i = 1, ..., n^{*} .$$

Methodology

Criterion for admittance of a Lie group of transformations

Theorem

The system
$$M_{NL} := \begin{cases} \dot{x}(t) = f(x(t), \theta, u(t)), \\ y(t) = g(x(t), \theta, u(t)) \end{cases}$$
 admits a one-parameter Lie group of transformations defined by $X \iff z$

$$\mathbf{X}' \cdot (\dot{x}_k - f_k(x)) = 0, \ k = 1, ..., m$$

$$\mathbf{X} \cdot (y_l - g_l(x)) = 0, \ l = 1, ..., n$$

where X' is the derivative of infinitesimal generators:

$$X' = \sum_{i=1}^{n^*} \eta_i(x) \frac{\partial}{\partial x_i} + \sum_{i=1}^{n^*} \eta_i'(x) \frac{\partial}{\partial \dot{x_i}} , \quad \text{where} \quad \eta_i'(x) = \sum_{j=1}^{n^*} \dot{x}_j \frac{\partial \eta_i}{\partial x_j}$$

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Methodology Criterion for admittance of a Lie group of transformations

The previous theorem leads to:

$$\begin{split} \sum_{j=1}^{n^*} \dot{x}_j \frac{\partial \eta_k}{\partial x_j}(\mathbf{x}) &- \sum_{i=1}^{n^*} \eta_i(\mathbf{x}) \frac{\partial f_k}{\partial x_i}(\mathbf{x}) = 0, \quad k = 1, ..., m ,\\ \sum_{i=1}^{n^*} \eta_i(\mathbf{x}) \frac{\partial g_l}{\partial x_i}(\mathbf{x}) = 0, \quad l = 1, ..., n . \end{split}$$

The above system of PDEs can be converted to a system of ODEs if we assume **rational** functions...

$$\dot{x}_k = f_k(\mathbf{x}) = rac{P^k(\mathbf{x})}{Q^k(\mathbf{x})}, \ k = 1, ..., m$$

 $y_l = g_l(\mathbf{x}) = rac{R^l(\mathbf{x})}{S^l(\mathbf{x})}, \ l = 1, ..., n$.

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Methodology Computing polynomials

... leading to:

Univariate + Partially variate:

$$P^{k}Q^{k}\frac{\partial \eta_{k}}{\partial x_{k}} - \sum_{i=1}^{n^{*}} \eta_{i}[P^{k}_{x_{i}}Q^{k} - P^{k}Q^{k}_{x_{i}}] = 0, \quad k = 1, ..., m ,$$
$$\sum_{i=1}^{n^{*}} \eta_{i}[R^{l}_{x_{i}}S^{l} - R^{l}Q^{l}_{x_{i}}] = 0, \quad l = 1, ..., n .$$

Multivariate:

$$\sum_{j=1}^{m} P^{j} Q^{k} \left(\prod_{b \neq j} Q^{b} \right) \frac{\partial \eta_{k}}{\partial x_{j}} - \sum_{i=1}^{n^{*}} \eta_{i} \left(\prod_{b \neq k} Q^{b} \right) \left[P_{x_{i}}^{k} Q^{k} - P^{k} Q_{x_{i}}^{k} \right] = 0,$$
$$\sum_{i=1}^{n^{*}} \eta_{i} \left[R_{x_{i}}^{l} S^{l} - R^{l} Q_{x_{i}}^{l} \right] = 0.$$

If the model contains specific initial conditions, they should be included in the equations.

$$\mathbf{X} \cdot (x_k - \mathbf{p}_{ini})|_{x=\mathbf{p}_{ini}} = 0, \quad k = 1, ..., m$$
 (1)

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Thus, following the same procedure as before:

$$\sum_{i=1}^{n^*} \eta_i(\mathbf{p}_{ini}) - \sum_{i=1}^{n^*} \eta_i \frac{V_{x_i}^k W^k - V^k W_{x_i}^k}{(W^k)^2}\Big|_{x=\mathbf{p}_{ini}} = 0, \quad k = 1, ..., m .$$
 (2)

Methodology Obtaining transformations

1. Consider the vector $\mathbf{r} = (r_{1,0}, r_{1,1}, ..., r_{n^*, d_{max}}),$

$$\sum_{i_1,\ldots,i_n} c_{i_1,\ldots,i_n}(\mathbf{r}) x_1^{i_1}\cdots x_n^{i_n} = 0 \Longrightarrow \mathbf{C} \cdot \mathbf{r} = 0$$

(Coefficients c_{i_1,\ldots,i_n} are linear in **r**).

2. To find symmetries, solve the linear system by computing the kernel of $\mathbf{C} = \begin{pmatrix} \cdots & \cdots & \cdots \\ \vdots & \ddots & \vdots \\ \ddots & \cdots & \cdots \end{pmatrix}$. 3. Take the vectors \mathbf{r} : (\vdots) , (\vdots) , ... and replace them in η_i to obtain the infinitesimal generators $\mathbf{X} = \sum_{i=1}^n \eta_i(\mathbf{x}) \frac{\partial}{\partial x_i}$

- Build the expression of x* with the infinitesimal generators X
- When the infinitesimal transformation is given by powers of one variable → "elementary" transformation. Examples:

$$\begin{split} x_i^* &= x_i + \varepsilon, \ \mathbf{X} = \frac{\partial}{\partial x_i} \ (\text{translation}) \ ,\\ x_i^* &= \exp(\varepsilon) x_i, \ \mathbf{X} = x_i \frac{\partial}{\partial x_i} \ (\text{scaling}) \ ,\\ x_i^* &= \frac{x_i}{1 - \varepsilon x_i}, \ \mathbf{X} = x_i^2 \frac{\partial}{\partial x_i} \ (\text{Mobius}) \ ,\\ x_i^* &= \frac{x_i}{[1 - (p - 1)\varepsilon x_i^{p - 1}]^{\frac{1}{p - 1}}}, \ \mathbf{X} = x_i^p \frac{\partial}{\partial x_i} \ (\text{higher order}) \ . \end{split}$$

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The most common ones are translation and scaling.

1. Choose the type of polynomial *Ansatz* (uni-, partial, multi-) and the maximum degree.

- 2. Create infinitesimal polynomials, η_i
- 3. Build the expressions for states, outputs, (& ICs)
- 4. Cast as $\mathbf{C} \cdot \mathbf{r} = 0$ and find \mathbf{r} by kernel(\mathbf{C})
- 5. Replace **r** in η_i to obtain transformations **X**

Implementations

- MinimalOutputSets (Mathematica)⁴
- SADE (Maple) ⁵
- symmetryDetection (Python)⁶
- LieSymmetries (Matlab) ⁷
 - Maximizes number of elementary transformations.
 - Computes non-elementary transformations.
 - Choose the states for which initial conditions are considered.

⁴Anguelova, M.; Karlsson, J.; Jirstrand, M. "Minimal output sets for identifiability". *Mathe Biosci*, 239:139153, 2012.

⁵Rocha Filho, T.M.; Figueiredo, A. "[SADE] a Maple package for the symmetry analysis of differential equations". *Comput Phys Commun*, 182:467476, 2011.

⁶Merkt, B., Timmer, J., and Kaschek, D. "Higher-order Lie symmetries in identifiability and predictability analysis of dynamic models". *Phys Rev E* 92.1, 2015.

⁷Massonis, G., and Villaverde, A.F. "Finding and Breaking Lie Symmetries: Implications for Structural Identifiability and Observability in Biological Modelling". Symmetry 12(3):469, 2020.

EXAMPLES

Simple chemical reaction

(1) Model diagram:

(3) Two infinitesimal generators:

$$\mathbf{X} = \mathbf{A} \frac{\partial}{\partial \mathbf{A}} - k \frac{\partial}{\partial k} - s_1 \frac{\partial}{\partial s_1} - s_2 \frac{\partial}{\partial s_2} \,.$$
$$\mathbf{X} = \mathbf{A}^2 \frac{\partial}{\partial \mathbf{A}} + \frac{\partial}{\partial s_2} \,.$$

(2) Model equations:

$$\dot{A} = -2kA^2 ,$$
$$A^{obs} = s_1 \frac{A}{1 + s_2 A} .$$

(4) New variables (all transformations are elementary):

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$$A^* = e^{\varepsilon}A, k^* = e^{-\varepsilon}k ,$$

$$s_1^* = e^{-\varepsilon}s_1, s_2^* = e^{-\varepsilon}s_2 .$$

$$A^* = \frac{A}{1 - \varepsilon A}, s_2^* = s_2 + \varepsilon .$$

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Simple chemical reaction MATLAB output

```
>> Lie_Symmetry
Ansatz --> OK
Derivatives Ansatz --> OK
Numerator and denominator --> OK
Derivatives numerator and denominator --> OK
States Polynomial --> OK
Observation Polynomial --> OK
System --> OK
Kernel --> OK
```

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Elapsed time is 5.325316 seconds.

Pharmacokinetic model

(1) Model diagram:

(2) Model equations:



$$\begin{aligned} \dot{x_1} &= u - (k_1 + k_2) x_1 , \\ \dot{x_2} &= k_1 x_1 - (k_3 + k_6 + k_7) x_2 + k_5 x_4 , \\ \dot{x_3} &= k_2 x_1 + k_3 x_2 - k_4 x_3 , \\ \dot{x_4} &= k_6 x_2 - k_5 x_4 , \\ x_2^{obs} &= s_2 x_2 , \\ x_3^{obs} &= s_3 x_3 . \end{aligned}$$

(3) Infinitesimal generator:

$$\mathbf{X} = k_1 \left(\frac{\partial}{\partial k_1} - \frac{\partial}{\partial k_2} \right) - \frac{k_3 (k_1 + k_2)}{k_2} \left(\frac{\partial}{\partial k_3} - \frac{\partial}{\partial k_7} \right) - s_2 \frac{\partial}{\partial s_2} + \frac{k_1 s_3}{k_2} \frac{\partial}{\partial s_3} + x_2 \frac{\partial}{\partial x_2} - \frac{k_1 s_3}{k_2} \frac{\partial}{\partial x_3} + x_4 \frac{\partial}{\partial x_4} .$$

Pharmacokinetic model

(4) New variables (I):

$$\begin{split} x_2^* &= x_2 e^{\varepsilon}, \ x_4^* = x_4 e^{\varepsilon}, \ k_1^* = k_1 e^{\varepsilon}, \ s_2^* = s_2 e^{-\varepsilon} \\ x_3^* &= x_3 - \frac{\varepsilon k_1 x_3}{k_2} - \frac{\varepsilon^2 k_1 x_3}{2k_2} - \frac{\varepsilon^3 k_1 x_3}{6k_2} - \frac{\varepsilon^4 k_1 x_3}{24k_2} , \\ k_2^* &= k_2 - \varepsilon k_1 - \frac{\varepsilon^2 k_1}{2} - \frac{\varepsilon^3 k_1}{6} - \frac{\varepsilon^4 k_1}{24} , \\ k_3^* &= k_3 - \frac{k_3 (k_1 + k_2) \varepsilon}{k_2} + \frac{\varepsilon^2 k_3 (k_1 + k_2)}{2k_2} - \frac{\varepsilon^3 k_3 (k_1 + k_2)}{6k_2} + \frac{\varepsilon^4 k_3 (k_1 + k_2)}{24k_2} \\ k_7^* &= k_7 + \frac{k_3 (k_1 + k_2) \varepsilon}{k_2} - \frac{\varepsilon^2 k_3 (k_1 + k_2)}{2k_2} + \frac{\varepsilon^3 k_3 (k_1 + k_2)}{6k_2} - \frac{\varepsilon^4 k_3 (k_1 + k_2)}{24k_2} \\ s_3^* &= s_3 + \frac{\varepsilon k_1 s_3}{k_2} + \frac{\varepsilon^2 k_1 s_3 (2k_1 + k_2)}{2k_2^2} + \frac{\varepsilon^3 k_1 s_3 (6k_1^2 + 6k_1 k_2 + k_2^2)}{6k_2^3} + \\ &+ \frac{\varepsilon^4 k_1 s_3 (24k_1^3 + 36k_1^2 k_2^2 + 14k_1 k_2^2 + k_2^3)}{24k_2^4} . \end{split}$$

Pharmacokinetic model

(4) New variables (II):

$$\begin{split} x_2^* &= x_2 e^{\varepsilon}, \ x_4^* = x_4 e^{\varepsilon}, \ k_1^* = k_1 e^{\varepsilon}, \ s_2^* = s_2 e^{-\varepsilon} \ , \\ k_2^* &= k_1 + k_2 - k_1 e^{\varepsilon} \ , \\ k_3^* &= \frac{k_3 e^{-\varepsilon} (k_1 + k_2 - k_1 e^{\varepsilon})}{k_2} \ , \\ k_7^* &= k_7 + \frac{k_3 (k_1 + k_2)}{k_2} - \frac{k_3 e^{-\varepsilon} (k_1 + k_2)}{k_2} \ , \\ x_3^* &= \frac{x_3 (k_1 + k_2 - k_1 e^{\varepsilon})}{k_2} \ , \\ s_3^* &= \frac{k_2 s_3}{(k_1 + k_2 - k_1 e^{\varepsilon})} \ . \end{split}$$

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JAK-STAT signaling pathway



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Infinitesimal generators:

$$\mathbf{X} = t_{13} \frac{\partial}{\partial t_{13}} - t_{17} \frac{\partial}{\partial t_{17}} + t_{22} \frac{\partial}{\partial t_{22}} ,$$

$$\mathbf{X} = -x_{10} \frac{\partial}{\partial x_{10}} - t_{11} \frac{\partial}{\partial t_{11}} - t_{15} \frac{\partial}{\partial t_{15}} + t_{21} \frac{\partial}{\partial t_{21}} .$$
(3)

New variables:

$$t_{13}^* = t_{13}e^{\varepsilon}, \quad t_{17}^* = t_{17}e^{-\varepsilon}, \quad t_{22}^* = t_{22}e^{\varepsilon},$$
 (4)

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$$x_{10}^* = x_{10}e^{-\varepsilon}, \quad t_{11}^* = t_{11}e^{-\varepsilon}, \quad t_{15}^* = t_{15}e^{-\varepsilon}, \quad t_{21}^* = t_{21}e^{\varepsilon}$$
. (5)

DISCUSSION

Conclusions

- Symmetries inform about lack of SIO and about its source.
- Their study can replace or complement other SIO tests.
- We have illustrated the use of a symbolic computation tool that finds Lie symmetries and the corresponding transformations automatically.
- Open-source implementation in MATLAB. Integrated in the STRIKE-GOLDD toolbox.
- Other tools in Mathematica, Python, Maple.
- Based on previous results (Merkt et al) + a few additions, incl. automatically calculating symmetry-breaking transformations.
- Symmetry-breaking transformations fix observability... but the mechanistic meaning is generally lost (so are they any good?).

Bonus: other uses of symmetry in biological modelling

The study of symmetries can inform about observability. But there are other possible uses, see e.g. (& recent, open special issues in MDPI Symmetry journal):

- morphological (a)symmetries in development
- homeostasis processes
- ▶ ...

PLOS COMPUTATIONAL BIOLOGY

RESEARCH ARTICLE

Conservation laws by virtue of scale symmetries in neural systems

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They provided the funding

 Spanish Ministry of Science, Innovation and Universities ref. DPI2017-82896-C2-2-R (SYNBIOCONTROL)

And thank you for your attention