

Structural Analysis of Dynamical Networks: **BDC-Decomposition and Influence Matrix**

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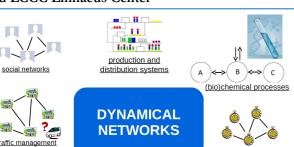
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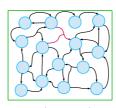
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Local Interactions

Global Behaviour

Graph Representation

• Parameter-free approach: system structure

Structural analysis

Explain behaviours based on the system inherent structure (graph)

• Structurally assess fundamental properties

Structural properties

Satisfied by all the systems of a family specified by a structure, without numerical bounds.

• Applications to biochemical systems

Structural properties in nature

Species: A, B, C

ODE system:

Concentrations: a, b, c

 $\dot{a} = a_0 - g_a(a) - g_{ac}(a,c)$

 $\dot{b} = g_a(a) - g_b(b)$

 $\dot{c} = g_a(a) - g_{ac}(a, c)$

 ${\sf Biological\ systems} \to {\sf extremely\ robust:\ fundamental\ properties}$ preserved despite huge uncertainties and parameter variations.

Example: biochemical reaction network

Reactions: $\emptyset \xrightarrow{a_0} A$, $A \xrightarrow{g_a} B + C$, $A + C \xrightarrow{g_{ac}} \emptyset$, $B \xrightarrow{g_b} \emptyset$

THE BDC-DECOMPOSITION

$$\dot{x}(t) = Sg(x(t)) + g_0,$$
 g monotonic functions

water distribution

Local BDC-decomposition

telecommunication / data

communication networks

The Jacobian can be decomposed as:

$$J(x) = \frac{\partial Sg(x)}{\partial x} = B\Delta(x)C, \qquad \Delta(x) = \operatorname{diag}\left\{\left|\frac{\partial g_k}{\partial x_h}\right|\right\} > 0.$$

The decomposition is unique (up to permutations).

Global BDC-decomposition

Given the equilibrium \bar{x} $(0 = Sg(\bar{x}) + g_0)$, $z \doteq x - \bar{x}$.

$$\dot{z}(t) = Sg(z(t) + \bar{x}) - Sg(\bar{x}) = [BD(z)C] \ z(t),$$

$$Sg(x) - Sg(\bar{x})$$

$$= \left[\int_{-\pi}^{\pi} I(\bar{x} + \sigma(x - \bar{x})) d\sigma \right] (x - \bar{x})$$

$$= B \left[\int_0^1 D(\bar{x} + \sigma(x - \bar{x})) d\sigma \right] C(x - \bar{x})$$

$$J(x) = \frac{\partial Sg(x)}{\partial x} = B\Delta(x)C, \qquad \Delta(x) = \operatorname{diag}\left\{\left|\frac{\partial g_k}{\partial x_h}\right|\right\} > 0.$$

The system can be rewritten as:

diagonal

 $D(z) \succ 0$.

From local to global

For any vector \bar{x}

 $V_X(x) = \inf\{\|w\|_1: Xw = x\}$

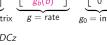
X full row rank

$$\begin{aligned}
sg(x) - sg(x) \\
&= \left[\int_0^1 J(\bar{x} + \sigma(x - \bar{x})) d\sigma \right] (x - \bar{x}) \\
&= \int_0^1 J(\bar{x} + \sigma(x - \bar{x})) d\sigma \\
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&= \int_0^1 J(\bar{x} + \sigma(x - \bar{x}$$

Based on a discrete difference inclusion,

a numerical algorithm computes the unit ball of the polyhedral Lyapunov

function (if any) via set iteration.



$$D = \operatorname{diag}\left\{\frac{\partial g_a}{\partial a}, \frac{\partial g_{ac}}{\partial a}, \frac{\partial g_{ac}}{\partial c}, \frac{\partial g_b}{\partial b}\right\} \succ 0$$

$$B = \begin{bmatrix} -1 & -1 & -1 & 0 \\ 1 & 0 & 0 & -1 \\ 1 & -1 & -1 & 0 \end{bmatrix} \text{ and } C = \begin{bmatrix} 1 & 0 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \\ 0 & 1 & 0 \end{bmatrix}$$



The procedure converges ⇒ structurally stable

BDC-based computation of Polyhedral Lyapunov Functions for structural stability

Monotonic functions g and dissipative reactions $\frac{\partial \dot{x}_i}{\partial x_i} < 0$

exploit the BDC-decomposition! Structurally \Leftrightarrow for any $D_i > 0$

Idea: $D(z(t)) \rightarrow D(t)$

Absorb the nonlinear system in a Linear Differential Inclusion

$$\dot{z}(t) = [BD(t)C] \ z(t), \qquad D(t) \succ 0. \quad (LDI)$$

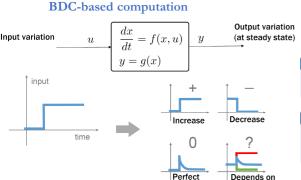
Any trajectory of the original system is also a trajectory of (LDI)

To analyse stability we can assume $0 \le D_i \le 1$.

Polyhedral... why?

The only structural Lyapunov function is polyhedral!

Structural Steady-State Influence:



Structural Influence Matrix

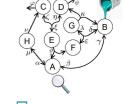
Structural influence of variable j on variable i

Assuming stability, $\Sigma_{ij} \in \{+,-,0,?\}$: sign of the steady-state variation of $x_i(\infty)$ due to a step input acting on x_i .

For systems admitting a BDC decomposition

$$\Sigma_{ij} = H_i(-BDC)^{-1}E_j,$$

H output matrix, E input matrix \rightarrow efficient vertex algorithm





Blanchini, F. and Giordano, G. (2015a). Polyhedral Lyapunov functions for structural stability of biochemical systems in concentration and reaction coordinates. In *Proceedings of the IEEE Conference on Decision and Control*, giulia.giordano@uniud.it 3110-3115. Osaka, Japan.

parameters

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