Proving Global Stability of Processive Phosphorylation Systems

Using Graph Reductions

Mitchell Eithun July 18, 2016

Ripon College



From last time:

• A chemical reaction network is a graph with species, reactions and complexes

$$A + B \xrightarrow{k_1} 3A + C \xrightarrow{k_2} D$$

- Mass-action kinematics converts a network into a dynamical system
- Monotone theory is used to prove global stability





We use the setup from [1].

- The **SR-graph** is a directed graph $G_{SR} = (V_{SR}, E_{SR}, L_{SR})$
- The set of vertices *V_{SR}* is the union of all species and reactions
- Rules governing edges and labels:
 - 1. If a species S is a reactant in any reaction or a product in a reversible reaction, then $\overrightarrow{SR}, \overrightarrow{RS} \in E_{SR}$.
 - 2. If a species S is a reactant in an irreversible reaction, then $\stackrel{\rightarrow}{RS} \in E_{SR}$.
 - Let SR ∈ E. If S is a reactant, L(S, R) := + and if S is a product, L(S, R) := −.



- The **R-graph** is an undirected graph $G = (V_R, E_R, L_R)$
- The set of vertices V_R is the set of reactions
- Rules governing edges:
 - 1. If there is a length-2 path connecting R_i and R_j in the SR-graph, then $R_iR_j \in E$
 - 2. Edges are labeled with the opposite of the product of the labels on the length-2 path



Definition

An SR-graph is *R*-strongly connected if there exists a directed path between every pair of reaction vertices.

Example (1-site phosphorylation)

SR-graph



Definition

We say an R-graph has the *positive loop property* if every simple loop has an even number of negative edges.



Theorem (Global Stability)

Let G be a reaction network satisfying several assumptions. Suppose

- 1. the R-graph of G has the positive loop property,
- 2. the directed SR-graph of G is strongly connected,
- 3. concentrations do not not vanish asymptotically (bounded-persistence), and
- 4. ker $\Gamma \cap int K \neq \emptyset$.

Then G has a unique steady state and it is a global attractor.

Graph Reductions

To remove an **intermediate** *Y*, the following two conditions must be met:

- (l1) Y consists of exactly one species and does not appear in any other complex
- (l2) there exist unique complexes y and y' such that
 - either $y \rightarrow Y$ or $y \rightleftharpoons Y$ is a reaction
 - either $Y \rightarrow y'$ or $Y \leftrightarrows y'$ is a reaction
 - shared species in y and y' (represented by e) have the same weights

•
$$y - e \rightarrow y' - e$$
, $y' - e \rightarrow y - e$, $y - e \rightleftharpoons y' - e$ and $y' - e \rightleftharpoons y - e$ are *not* reactions

From the network $G = (S, C, \mathcal{R})$, we construct the reduced network $G^* = (S^*, C^*, \mathcal{R}^*)$.

We define $\mathcal{R}^* := \mathcal{R}^*_c \cup \mathcal{R}^*_{Y}$, where \mathcal{R}^*_Y is the subset of reactions in \mathcal{R} that do not have Y as a product or reactant and

$$R_{Y}^{*} := \begin{cases} \{y - e \rightleftharpoons y' - e\}, \text{ if } y \rightleftharpoons Y, Y \rightleftharpoons y' \in \mathcal{R} \\ \{y - e \rightarrow y' - e\}, \text{ if } y \rightarrow Y \in \mathcal{R}, \text{ or } Y \rightarrow y' \in \mathcal{R} \end{cases}$$
(1)

$$S_0 + E \stackrel{R_1}{\Longrightarrow} S_0 E \stackrel{R_2}{\longrightarrow} S_1 + E$$

$$S_1 + F \xrightarrow{R_3} S_1F \xrightarrow{R_4} S_0 + F$$

$$R_1^*: S_0 \longrightarrow S_1$$

$$R_2^*: S_1 \longrightarrow S_0$$



Theorem (Invariance Under Removing Intermediates)

Let G be a reaction network some basic network assumptions. Suppose G* is a reaction network obtained from G by successive removal of intermediates. Then

- 1. the directed SR-graph of G* is R-strongly connected if, and only if, the directed SR-graph of G is strongly connected,
- 2. the R-graph of G^{*} has the positive loop property if, and only if, the R-graph of G has the positive loop property,
- 3. *G*^{*} is bounded persistent if and only if *G* is bounded persistent
- 4.

5. ker $\Gamma^* \cap int \ K^* \neq \emptyset \iff ker \ \Gamma \cap int \ K \neq \emptyset$.

Processive Phosphorylation

n-site Processive Phosphorylation

$$S_{0} + K = \frac{k_{1}}{k_{2}} S_{0}K = \frac{k_{3}}{k_{4}} S_{1}K = \frac{k_{5}}{k_{6}} \dots = \frac{k_{2n-1}}{k_{2n}} S_{n-1}K = \frac{k_{2n+1}}{k^{*}} S_{n} + K$$
$$S_{n} + F = \frac{\ell_{2n+1}}{\ell_{2n}} S_{n}F = \frac{\ell_{2n-1}}{\ell_{2n-2}} \dots = \frac{\ell_{5}}{\ell_{4}} S_{2}F = \frac{\ell_{3}}{\ell_{2}} S_{1}F = \frac{\ell_{1}}{\ell^{*}} S_{0} + F$$

This model captures systems that are reversible, irreversible and have product inhibition.

Last time we showed that the n-site model is globally stable.

$$P_{1} + E_{1} = - C_{11} = - C_{12} = - C_{12} = - C_{1n_{1}} = - P_{2} + E_{1}$$

$$P_{2} + E_{2} = - C_{21} = - C_{22} = - C_{22} = - C_{2n_{2}} = - P_{3} + E_{2}$$

$$\vdots \qquad \vdots \qquad \vdots \qquad \vdots$$

$$P_{m} + E_{m} = - C_{m_{1}} = - C_{m_{2}} = - C_{m_{2}} = - C_{m_{m_{m_{1}}}} = - P_{1} + E_{m_{1}}$$

Theorem (ME)

The dynamical system of the generalized model arising from mass-action kinematics has a unique positive steady state and it is a global attractor.

$$P_{1} + E_{1} = - C_{11} = - C_{12} = - C_{12} = - C_{1n_{1}} = - P_{2} + E_{1}$$

$$P_{2} + E_{2} = - C_{21} = - C_{22} = - C_{22} = - C_{2n_{2}} = - P_{3} + E_{2}$$

$$\vdots \qquad \vdots \qquad \vdots \qquad \vdots$$

$$P_{m} + E_{m} = - C_{m_{1}} = - C_{m_{2}} = - C_{m_{2}} = - C_{m_{m_{m_{1}}}} = - P_{1} + E_{m_{1}}$$









$$P_1 = P_2$$

$$P_2 = P_3$$

$$\vdots \qquad \vdots$$

$$P_m = P_1$$

$$P_1 = - P_3$$

$$P_3 = - P_4$$

$$\vdots \qquad \vdots$$

$$P_m = - P_1$$

$$P_1 = \stackrel{\sim}{=} P_m$$
$$P_m = \stackrel{\sim}{=} P_1$$

$$R_1^*: P_1 = P_m$$

$$R_2^*$$
: $P_m = P_1$

Processive Phosphorylation





Thank You!

References

[1] M. M. de Freitas, C. Wiuf, and E. Feliu. Intermediates and generic convergence to equilibria. 2016.