

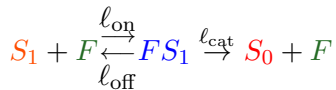
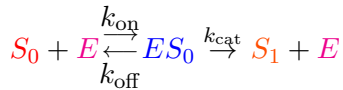
MSRI-MPI LEIPZIG SUMMER GRADUATE
SCHOOL 2023
MESSI SYSTEMS

Alicia Dickenstein

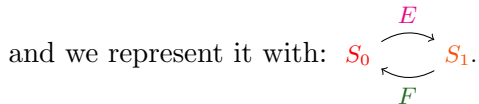
Departamento de Matemática, FCEN,
Universidad de Buenos Aires,
and Instituto de Matemática Luis A. Santaló, UBA-CONICET

Leipzig, June 19, 2023

PHOSHO-DEPHOSPHORYLATION: “FUTILE” CYCLE

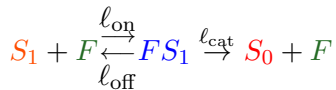
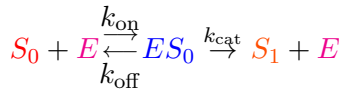


E and F enzymes, S_0 and S_1 substrates, S_0E and S_1F intermediates

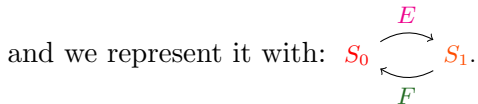


There are 6 species, 6 complexes (nodes) and 6 reactions (edges)

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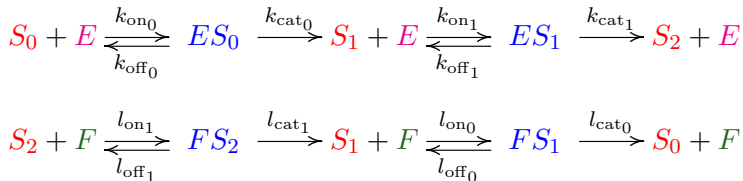


E and F enzymes, S_0 and S_1 substrates, S_0E and S_1F intermediates



There are 6 species, 6 complexes (nodes) and 6 reactions (edges)

TWO SEQUENTIAL PHOSPHORYLATIONS



WE NUMBER THE SPECIES AND THEIR CONCENTRATIONS

x_1, x_2, x_3 = concentrations of S_0, S_1, S_2

y_1, y_2, y_3, y_4 = concentrations of the intermediate species

x_4 = concentration of the kinase E

x_5 = concentration of the phosphatase F .

THE DIFFERENTIAL EQUATIONS AND THE CONSERVATION LAWS

$$\frac{dx_1}{dt} = -k_{on_0}x_1x_4 + k_{off_0}y_1 + l_{cat_0}y_4$$

$$\begin{aligned} \frac{dx_2}{dt} = & -k_{on_1}x_2x_4 + k_{cat_0}y_1 + k_{off_1}y_2 \\ & -l_{on_0}x_2x_5 + l_{cat_1}y_3 + l_{off_0}y_4 \end{aligned}$$

$$\frac{dx_3}{dt} = k_{cat_1}y_2 - l_{on_1}x_3x_5 + l_{off_1}y_3$$

$$\frac{dy_1}{dt} = k_{on_0}x_1x_4 - (k_{off_0} + k_{cat_0})y_1$$

$$\frac{dy_2}{dt} = k_{on_1}x_2x_4 - (k_{off_1} + k_{cat_1})y_2$$

$$\frac{dx_4}{dt} = -k_{on_0}x_1x_4 - k_{on_1}x_2x_4 + (k_{off_0} + k_{cat_0})$$

$$+ (k_{off_1} + k_{cat_1})y_2$$

$$\frac{dx_5}{dt} = -l_{on_0}x_2x_5 - l_{on_1}x_3x_5 + (l_{off_1} + l_{cat_1})y_3$$

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$$\frac{dy_3}{dt} = l_{on_1}x_3x_5 - (l_{off_1} + l_{cat_1})y_3$$

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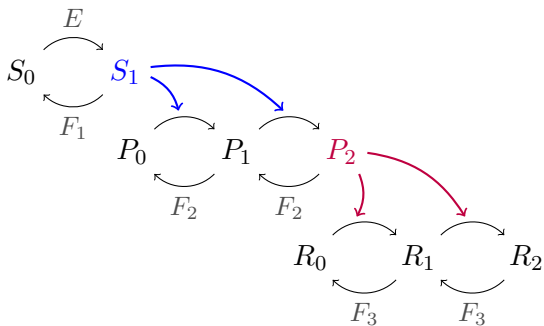
$$x_1 + x_2 + x_3 + y_1 + y_2 + y_3 + y_4 = S_{tot}$$

$$x_4 + y_1 + y_2 = E_{tot}$$

$$x_5 + y_3 + y_4 = F_{tot}$$

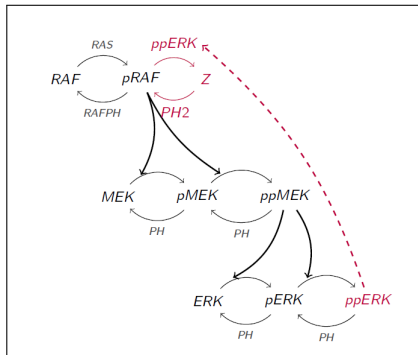
OTHER IMPORTANT EXAMPLES OF NETWORKS

Phosphorylation cascades



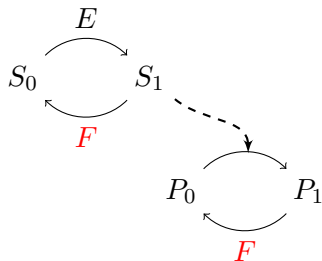
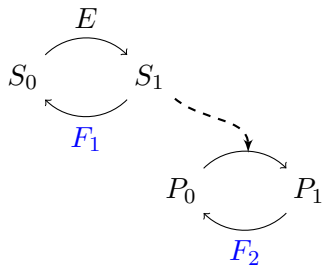
OTHER IMPORTANT EXAMPLES OF NETWORKS

Phosphorylation cascades with retroactivity



OTHER IMPORTANT EXAMPLES OF NETWORKS

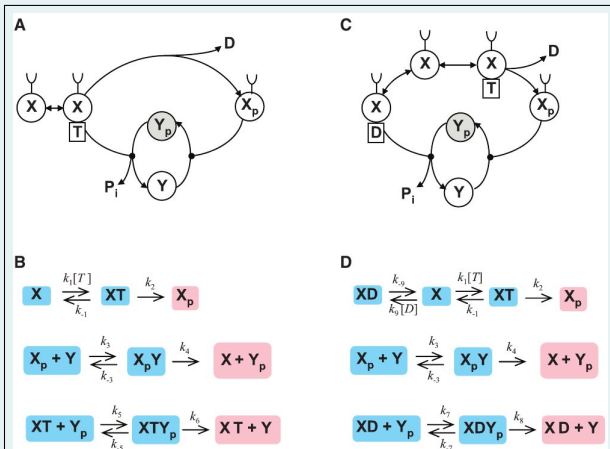
Different phosphatases vs same phosphatase in a cascade



OTHER IMPORTANT EXAMPLES OF NETWORKS

Bifunctional enzyme in *E. coli*, Shinar-Feinberg, Science '10

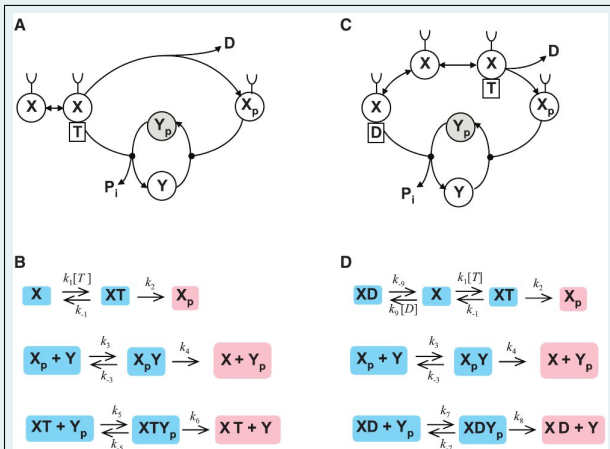
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 $X = \text{EnvZ}, \quad Y = \text{OmpR}$ 

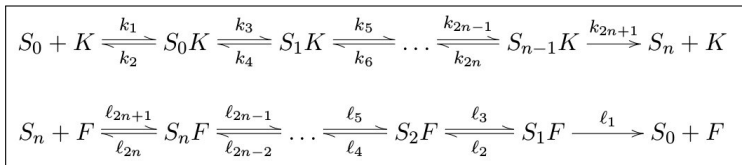
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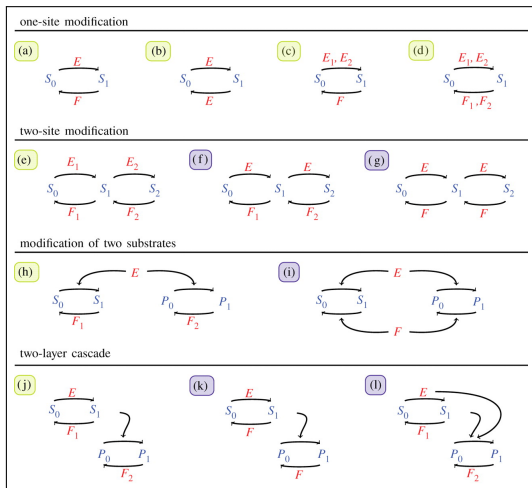
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EXAMPLE: PROGRESSIVE PHOSPHORILATIONS

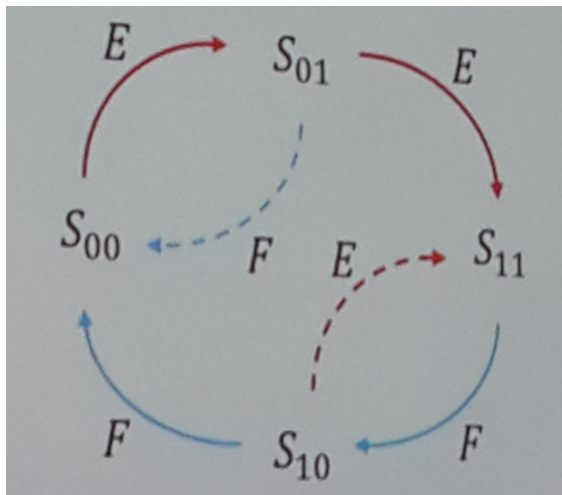


C. CONRADI AND A. SHIU. *A global convergence result for progressive multisite phosphorylation systems*, 2015.

SMALL MOTIFS ([ALON'07, FELIU-WIUF'12])



SHVARTSMAN'S ENZYMATICAL NETWORK



A COMMON STRUCTURE [P.MILLÁN-D.'18: THE STRUCTURE OF MESSI BIOLOGICAL SYSTEMS]

MESSI SYSTEMS

We identified with Mercedes Pérez Millán a **common structure** in many popular biological networks that describe Modifications of type Enzyme-Substrate or Swap with Intermediates, which allows us to prove general results valid in all these networks. MESSI systems include **all** the previous ones.

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A **MESSI network** is a chemical reaction network satisfying the following properties. When endowed with mass-action kinetics, we have a **MESSI system**.

- There exists a **partition** of the set \mathcal{S} of species:

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Examples: All the examples we mentioned, all the linear ones ... plus many other common biochemical models.

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- In a reaction $X_i + X_\ell \rightarrow X_j + X_\ell$, we say that X_ℓ acts as an enzyme.
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A MESSI system has **one (independent) linear conservation relation** associated to each of the subsets $S^{(\alpha)}$, $1 \leq \alpha \leq M$, in the partition of the species set corresponding to **non-intermediate species**:

$$\sum_{\mathbf{X}_i \in S^{(\alpha)}} x_i + \sum_{\mathbf{X}_k \in \text{Int}_\alpha} x_k = \text{constant},$$

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OBSERVATION:

- Theorem 1 implies that all MESSI systems are **conservative** (and thus the solutions are defined for any positive time).
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FROM G TO G_1 (WITHOUT INTERMEDIATES)

Going from G to G_1 we delete intermediates and we put an edge between two core complexes $y_i \rightarrow y_j$ if $y_1 \rightarrow_o y_j$ in G :

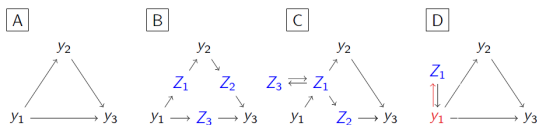


FIGURE: $\mathcal{S}^{(0)} \subseteq \{Z_1, Z_2, Z_3\}$, $\mathcal{S}^{(1)} = \{y_1, y_2, y_3\}$

In all cases $G = A, B, C, D$ (with rate constants κ), the associated digraph G_1 is **A**.

Wiuf and Feliu proved that with rate constants $\tau(\kappa)$ and QSSA style substitutions, G_1 has mass-action kinetics form.

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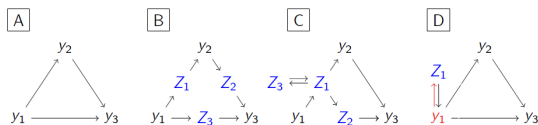


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RELATION BETWEEN THE STEADY STATES OF (G, κ) AND $G_1(\tau, \kappa)$

PROJECTING THE STEADY STATES

The rational map sending the rate constants κ in G to the rate constants $\tau(\kappa)$ in G_1 verifies that the steady states of the mass-action chemical reaction systems defined by G with rate constants κ and G_1 with rate constants $\tau(\kappa)$, are in one-to-one correspondence via the **projection** $\pi(u, x) = x$, where u is the vector of concentrations of the intermediate species and x is the vector of concentrations of the core species.

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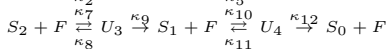
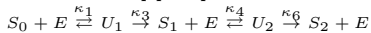
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$G_1 \rightarrow G_2$ (HIDE ENZYMES AND SWAPS IN LABELS)

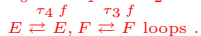
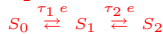
G : Double seq. phospho.



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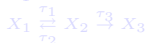
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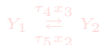
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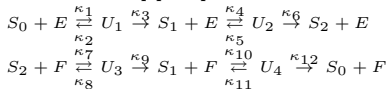
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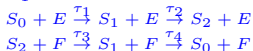
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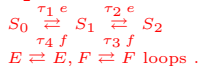
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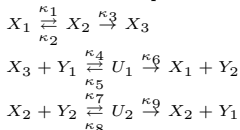
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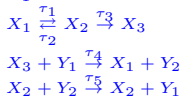
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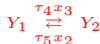
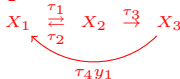
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MG_2, G_2 AND G_2°

- $G_1 \rightarrow MG_2$ produces in general a **multigraph** with **loops**.
- $MG_2 \rightarrow G_2$ replaces multiple arrows between the same two nodes with a **single** edge, and the new label is the **sum of the labels** in all the edges in MG_2 between these two nodes.
- $G_2 \rightarrow G_2^\circ$ **deletes** the loops.

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AN IMPORTANT PROPERTY

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Persistence means that any trajectory starting from a point with **positive coordinates** stays at a **positive distance** from any point in the boundary.

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PRECLUDING RELEVANT BOUNDARY STEADY STATES

If we have a *minimal* partition, we define a new graph G_E , whose **vertices** are the sets $S^{(\alpha)}$ for $\alpha \geq 1$, and there is an **edge** from $S^{(\alpha)}$ to $S^{(\beta)}$ if there is a species in $S^{(\alpha)}$ on a **label** of an edge in G_2 between species of $S^{(\beta)}$.

PERSISTENCE: THEOREM 2

If there is **no directed cycle** in G_E , then G has **no** boundary steady states in any positive stoichiometric compatibility class. So, the system is conservative and there are **no relevant** boundary steady states. Thus, the network is **persistent**.

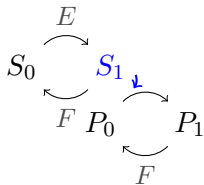
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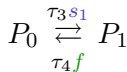
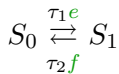
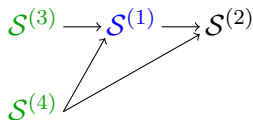
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EXAMPLES



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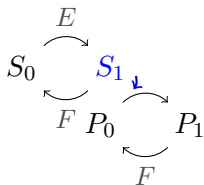
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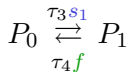
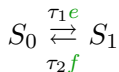
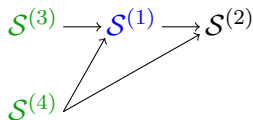
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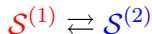


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DECIDING MULTISTATIONARITY: REMINDER

Assume the steady state variety V is cut out by binomials with exponents in a matrix B and coefficients in $\mathbb{Q}(\kappa)$, or equivalently, it is parametrized by monomials with exponents in the dual matrix A . Let W denote a matrix whose rows define the dual of the subspace $S = \langle y_j - y_i, y_i \rightarrow y_j \rangle$. If $\text{rank}(W) = \text{rank}(A) = d$, the following statements are equivalent:

- 1 There is at most a **single** positive solution in $V \cap x(0) + S$ for any $x(0)$ in the positive orthant (**monostationarity**), for any choice of rate constants κ .
- 2 For all subsets $J \subseteq [s]$ of cardinality d , the product $\det(W) \det(A_J)$ either is zero or has the **same sign** as all other nonzero products, and at least one such product is **nonzero**.

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DEFINITION

A *structurally toric*, or *s-toric MESSI system*, is a MESSI system whose digraph G satisfies the following conditions:

- (\mathcal{C}') : For every intermediate complex U_k there exist a *unique* core complex $y_{(k)}$ such that $y_{(k)} \rightarrow_{\circ} U_k$ in G .
- (\mathcal{C}'') : The associated multidigraph MG_2 does not have parallel edges and the digraph G_2 is weakly reversible.
- (\mathcal{C}''') : ...

INTERMEDIATES SATISFY BINOMIAL EQUATIONS AT S.S.

Given a MESSI network G that satisfies condition (\mathcal{C}') , there are (**explicit**) rational functions $\mu_k \in \mathbb{Q}(\kappa)$, $1 \leq k \leq p$, such that for any steady state $x \in \mathbb{R}_{>0}^n$ of the associated MESSI network G_1 , the steady state $\pi^{-1}(x) = (u(x), x)$ of G is given by a **monomial** map:

$$u_k(\mathbf{x}) = \mu_k \mathbf{x}^{y_{(k)}}, \quad k = 1, \dots, p. \quad (2)$$

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COMBINATORIALLY CONSTRUCTED

Any s-toric MESSI system is toric. Moreover, we can choose **explicit binomials** with coefficients in $\mathbb{Q}(\kappa)$ which describe the positive steady states.

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Let G be an s-toric MESSI network. Assume that the partition is **minimal** with m subsets of core species and the associated digraph G_E has **no** directed cycles. Then,

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Let G be the underlying digraph of a MESSI system satisfying condition C' .

Assume that the associated digraph G_E has no directed cycles, the underlying undirected graph of the associated graph G_2^o is a forest (an acyclic graph), and MG_2 has no parallel edges.

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RATIONAL PARAMETRIZATIONS

- A wide class of MESSI systems admit a rational parametrization.
- It is shown by [Feliu-Wiuf'13] that the values of the intermediate species at steady state can be rationally written in terms of the core species, in an algorithmic way.
- The following result extends Theorem 4 in [Thomson-Gunawardena'09].

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Let G be the underlying digraph of a MESSI system. Assume that the associated digraph G_2 is weakly reversible and the associated digraph G_E has no directed cycles. Then, $V_{>0}(I_f)$ admits a *rational parametrization*, which can be *algorithmically computed*.

More explicitly, it is possible to define *levels* for the subsets $\mathcal{S}^{(\alpha)}$, $\alpha \geq 1$, according to *indegree*. Then, given any choice of one index i_α in each $\mathcal{S}^{(\alpha)}$, the concentration of any core species x_i in $\mathcal{S}^{(\alpha)}$ can be rationally expressed in an effective way in terms of x_{i_α} and the variables x_{i_β} for which the *indegree* of $\mathcal{S}^{(\beta)}$ is strictly *smaller* than the indegree of $\mathcal{S}^{(\alpha)}$.

Moreover, if the partition is *minimal* with M subsets of core species, then $\dim(V_{>0}(I_f)) = M$ and $M = \text{rank}(W)$.

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