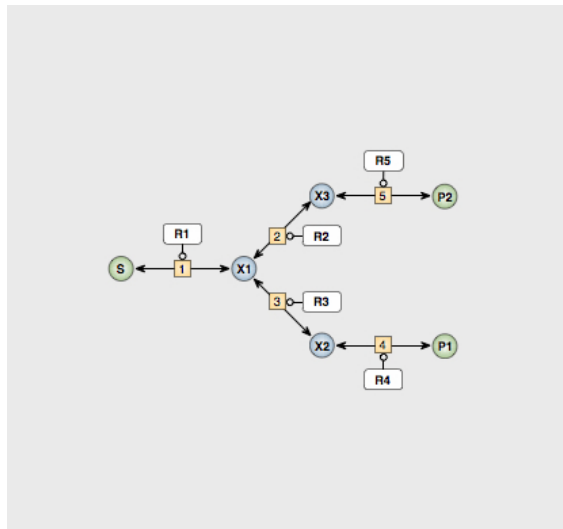


# Systems Biology Tutorial 4: Structural analysis of reaction networks

1. Consider the linear branched pathway:



- (a) Construct the stoichiometric matrix  $N$ .

$$N = \begin{bmatrix} 1 & -1 & -1 & 0 & 0 \\ 0 & 0 & 1 & -1 & 0 \\ 0 & 1 & 0 & 0 & -1 \end{bmatrix}$$

where

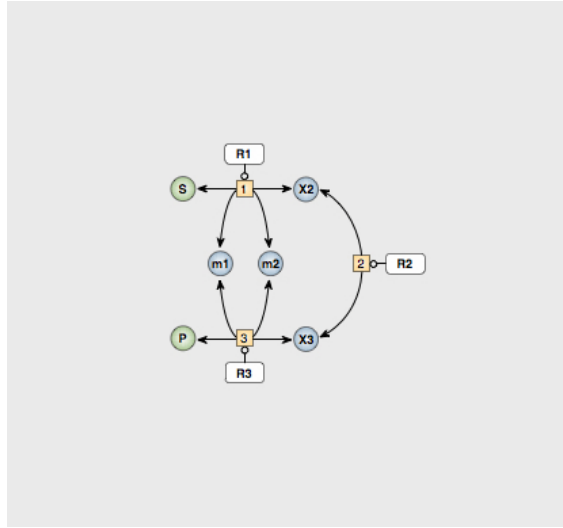
$$\begin{bmatrix} \dot{x}_1 \\ \dot{x}_2 \\ \dot{x}_3 \end{bmatrix} = N \cdot \begin{bmatrix} v_1 \\ v_2 \\ v_3 \\ v_4 \\ v_5 \end{bmatrix}$$

- (b) Are there any dependent metabolites?  
*No*
- (c) Derive the steady-state flux relations by hand from  $Nv = 0$ . How many independent fluxes are there?  
*2 independent fluxes. If you choose  $J_1$  and  $J_2$  the relations are:*

$$\begin{aligned} J_3 &= J_1 - J_2 \\ J_4 &= J_3 = J_1 - J_2 \\ J_5 &= J_2 \end{aligned}$$

- (d) Check your answers by running the branch5 model on JWS Online and generating the  $N$ ,  $L$  and  $K$  matrices.

2. Consider the linear pathway with a moiety:



- (a) Construct the stiochiometric matrix.

$$N = \begin{bmatrix} -1 & 0 & 1 \\ 1 & -1 & 0 \\ 1 & 0 & -1 \\ 0 & 1 & -1 \end{bmatrix}$$

where

$$\begin{bmatrix} \dot{m}_1 \\ \dot{x}_2 \\ \dot{m}_2 \\ \dot{x}_3 \end{bmatrix} = N \cdot \begin{bmatrix} v_1 \\ v_2 \\ v_3 \end{bmatrix}$$

- (b) Are there any dependent metabolites?  
*Yes.  $m_1 + m_2 = \text{constant1}$  and  $x_3 + x_2 + m_1 = \text{constant2}$ .*
- (c) Derive the steady-state flux relations by hand from  $Nv = 0$ . How many independent fluxes are there?  
*1 independent flux.  $J_1 = J_2 = J_3$ .*
- (d) Check your answers by running the lin3moi model on JWS Online and generating the N, L and K matrices.

The diagram illustrates the metabolic map of the glycolysis pathway, showing the flow of metabolites and the activity of various enzymes across different cellular compartments: cytosol, glycosome, and mitochondrion.

**Key Enzymes and Reactions:**

- GlcE** (green circle) enters the cytosol.
- GlcTr** (1) (yellow box) converts GlcE to Glc.
- HK** (2) (yellow box) converts Glc to Glc6P.
- PGI** (3) (yellow box) converts Glc6P to Fru6P.
- PFK** (4) (yellow box) converts Fru6P to Fru16BP.
- ALD** (5) (yellow box) converts Fru16BP to GAP.
- TPI** (6) (yellow box) converts GAP to DHAP.
- GAP** (8) (yellow box) converts DHAP to BPG13.
- BPG13** (9) (yellow box) converts BPG13 to Pyr.
- PK** (10) (yellow box) converts Pyr to PyrE.
- ATPase** (11) (yellow box) converts ATP to ADP.
- GlyK** (13) (yellow box) converts Gly to Gly3P.
- GPO** (14) (yellow box) converts H<sub>2</sub>O to O<sub>2</sub>.
- GDH** (12) (yellow box) converts NADH to NAD.

**Regulatory Elements:**

- Prb** (green box) is a regulatory element that inhibits the activity of HK (2) and PFK (4).
- Nb** (green box) is a regulatory element that inhibits the activity of PK (10).

**Metabolites and Cofactors:**

- ATP** (blue circle) and **ADP** (blue circle) are involved in energy metabolism.
- AMP** (blue circle) is a product of ATP hydrolysis.
- NAD** (blue circle) and **NADH** (blue circle) are involved in redox reactions.
- H<sub>2</sub>O** (blue circle) and **O<sub>2</sub>** (blue circle) are involved in water metabolism.

**Cellular Compartments:**

- cytosol**: The main compartment for glycolysis.
- glycosome**: A specialized organelle for glycolysis.
- mitochondrion**: The organelle responsible for energy production.

- 3

- (e) Would glycolysis reach a steady state in the absence of the glycerol branch (anaerobically)?  
*No, Redox balance will not be maintained i.e. the organism will run out of NADH*

The diagram illustrates the central carbon metabolism of *E. coli*, showing the conversion of Glucose (Glc) into various products. The pathways are interconnected, involving numerous enzymes and cofactors. Key features include:

- Glucose Entry:** Glc enters the pathway and is converted to Pyruvate (Pyr) via several steps involving enzymes like R1, R2, R3, and R4, and cofactors like NAD, NADH, ADP, and ATP.
- Pyruvate (Pyr) Hub:** Pyr is a central intermediate that can be converted to Lactate (Lac) via LDH, to Acetyl-CoA (AcCoA) via PDHc, or to Acetyl-CoA (AcCoA) via AS.
- Acetyl-CoA (AcCoA) Hub:** AcCoA is a central intermediate that can be converted to Acetate (Acet) via ATR, to Ethanol (EtOH) via ADH, to Butyrate (But) via BDH, or to Acetyl-CoA (AcCoA) via ACALDH.
- Acetate (Acet) Output:** Acetate is the final product of the central carbon metabolism, released from the cell via the Cellwall.
- Other Products:** Ethanol (EtOH) and Butyrate (But) are also products of the central carbon metabolism.
- Enzymes and Cofactors:** The diagram shows various enzymes (R1, R2, R3, R4, LDH, PDHc, AS, ATR, BDH, ADH, ACALDH, ACK, PTA) and cofactors (NAD, NADH, ADP, ATP, CoA, H<sub>2</sub>O, O<sub>2</sub>, H<sub>2</sub>, CO<sub>2</sub>) involved in the reactions.

- 5