

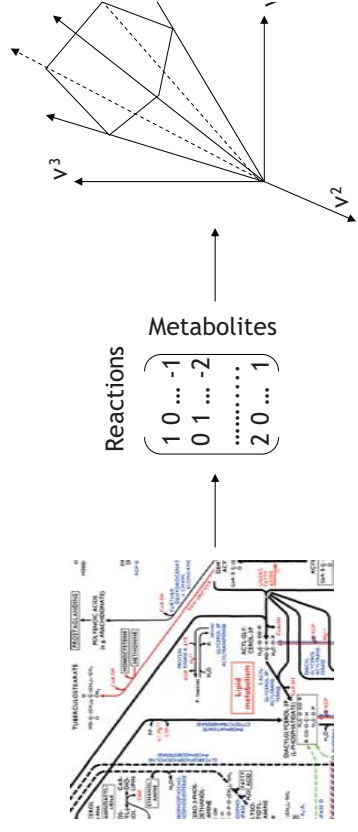


Steady-state flux cone

Set of all possible steady-state flux distributions

$$C = \{v \in \mathbb{R}^n \mid Sv = 0, v_i \geq 0, i \in Irr\}$$

→ polyhedral cone



3. Flux balance analysis (FBA)

- Assume cellular behavior is determined by a certain biological objective.
- Determine a corresponding "best" flux distribution.
- Use mathematical optimization to predict phenotype.

Simplest case: **Linear programming (LP)**

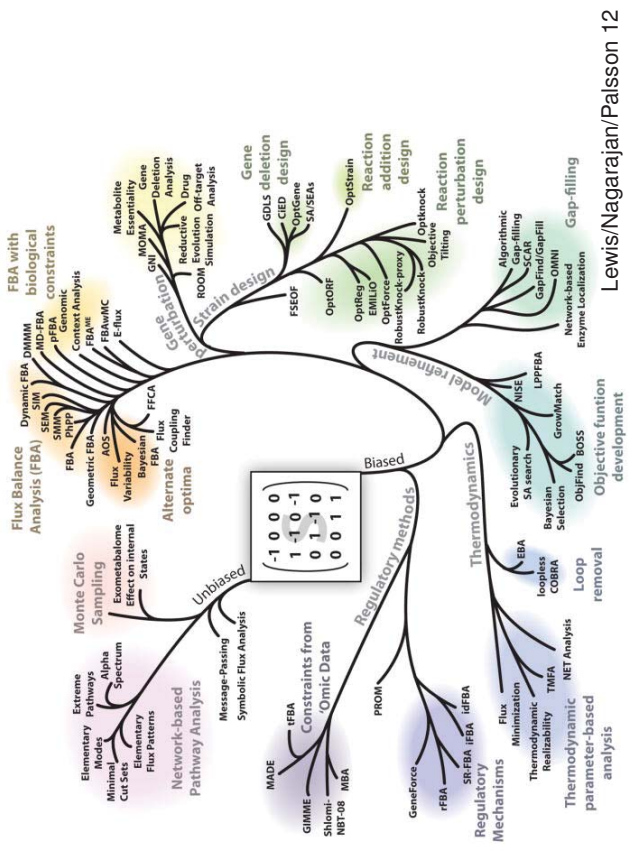
$$\max\{c^T x \mid Ax \leq b, x \in \mathbb{R}^n\}$$

Flux balance problem (FBA)

$$\max\{c^T v \mid Sv = 0, l \leq v \leq u\} \quad (\text{FBA})$$



Constraint-based analysis methods



Example

a Genome-scale metabolic reconstruction

b Mathematically represent metabolic reactions and constraints

c Mass balance defines a system of linear equations

d Define objective function ($Z = c_1 v_1 + c_2 v_2 \dots$)

e Calculate fluxes that maximize Z

$A \leftrightarrow B + C$ Reaction 1
 $B + 2C \rightarrow D$ Reaction 2
 ...
 Reaction n

Metabolites: 1, 2, ..., m
 Reactions: 1, 2, ..., n

Stoichiometric matrix, S

Fluxes, v

To predict growth, $Z = v_{\text{biomass}}$

Solution space defined by constraints

Point of optimal v

- E. coli* metabolism
 - Genome-scale reconstruction (iJO1366)
 - 1336 metabolites, 2251 reactions
 - Objective function: biomass
 - Glucose and oxygen uptake reactions
 - Aerobic and anaerobic growth
 - Software: e.g. COBRA Toolbox 2.0
- Orth/Thiele/Palsson 10



4. Flux variability analysis (FVA)

- Optimal solutions to FBA problems need not be unique.
 - Enumerating all optimal solutions is computationally expensive.
 - Alternative: **Flux variability analysis (FVA)**

$$z_{opt} = \max\{z = c^T v \mid Sv = 0, l \leq v \leq u\} \quad (\text{FBA})$$
- For all $j = 1, \dots, n$:
- $$\max\{\pm v_j \mid Sv = 0, l \leq v \leq u, c^T v = z_{opt}\} \quad (\text{FVA})$$



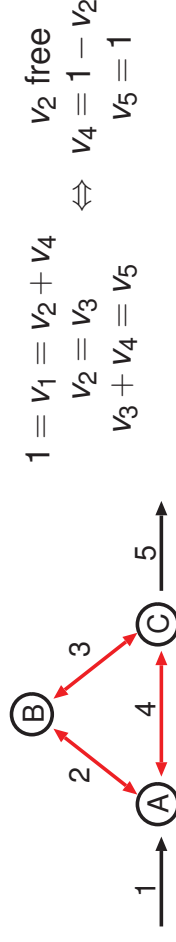
5. Flux coupling analysis (FCA)

Burgard et al. 04

- $C = \{v \mid Sv = 0, v_k \geq 0, k \in Irr\}$ flux cone
- A reaction i is **blocked** if $v_i = 0$, for all $v \in C$.
- Let i and j be two unblocked reactions.
 - i is **directionally coupled** to j , $i \xrightarrow{0} j$, if for all $v \in C$, $v_i = 0$ implies $v_j = 0$.
 - i and j are **partially coupled**, $i \xleftrightarrow{0} j$, if for all $v \in C$, $v_i = 0$ is equivalent to $v_j = 0$.
 - i and j are **fully coupled**, $i \rightsquigarrow j$, if there exists $\lambda \in \mathbb{R} \setminus \{0\}$ such that for all $v \in C$, $v_j = \lambda v_i$.
- $i \rightsquigarrow j$ implies $i \xleftrightarrow{0} j$, which is equivalent to $i \xrightarrow{0} j$ and $j \xrightarrow{0} i$.



Thermodynamic constraints



- Flux through internal cycles is unbounded \rightsquigarrow thermodynamically infeasible
- Include additional (non-linear) thermodynamic constraints (Beard/Qian 02)
- Fast thermodynamic FVA (tFVA)** (Müller/Bockmayr 12)
- Application:** Modules in the optimal flux space (Müller/Bockmayr 13)



Example

