# Mathematical Foundations of Constraint-Based Modelling

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### Topics to cover

- Kinetic models of metabolism and the quasi-steady state assumption
- Constraints on metabolic flux distributions and feasibility space
- Objective functions, constrained optimization, and LP problems

# Metabolism is a network of reactions

• In biology, metabolites are inter-converted by chemical reactions



• Metabolites can be intra- or extracellular, and compartmentalized

# Modelling the rate of a reaction

- The Law of Mass Action says that the rate of a reaction is proportional to the probability of collisions (and therefore concentrations) of the metabolites participating in the reaction
- For a reversible reaction of the form

$$S_1 + S_2 \rightarrow 2S_3$$

this means the rate broken down in terms of **forward** and **reverse** directions is

$$v = v_{+} - v_{-} = k_{+}S_{1}S_{2} - k_{-}S_{3}^{2}$$

## Modelling the rate of a reaction contd.

 The rate of reaction can be used to express the derivatives of metabolite concentrations as a system of ordinary differential equations (ODEs)

$$\frac{dS_1}{dt} = \frac{dS_2}{dt} = -v = -(k_+ S_1 S_2 - k_- S_3^2)$$
$$\frac{dS_3}{dt} = 2v$$

# Stochiometric coefficients

• From the previous slide we have

$$\frac{dS_1}{dt} = \frac{dS_2}{dt} = -\nu, \qquad \frac{dS_3}{dt} = 2\nu$$

- We say that the stochiometric coefficient of  $S_1$  and  $S_2$  is -1 and that of  $S_3$  is  $\mathbf{2}$
- In general, stochiometric coefficients denote the proportion of each metabolite involved in a reaction

### Representing a metabolic reaction network

• A **metabolic network** with *n* reactions can represented as



• This system of ODEs can be written in vector form as

$$\frac{d\mathbf{S}}{dt} = \mathbf{N} \cdot \mathbf{v}$$

### Some terminology

• In the system of ODEs

$$\frac{d\mathbf{S}}{dt} = \mathbf{N} \cdot \mathbf{v}$$

- a) S is called the vector of **metabolite concentrations** (of length *m*)
- **b)** N is called the **stochiometric matrix** (of dimension  $m \times n$ )
- c) v is called the **flux vector** (of length n)

### Quasi-steady state assumption

• Intracellular metabolite concentrations are modelled with units per gram of biomass, X , which accumulates with growth rate  $\mu$ :

$$\frac{d\mathbf{S}}{dt} = \mathbf{N} \cdot \mathbf{v} - \mu \mathbf{S}$$

• The quasi-steady state assumption (QSSA) assumes that metabolic transients are fast compared to cell growth effects so that

$$0 = \mathbf{N} \cdot \mathbf{v}$$

### Stochiometric constraints

• From the QSSA, the flux vector **v** must satisfy the **stochiometic constraints** (**v** is in the **null space** of **N**)

$$\mathbf{N}\cdot\mathbf{v}=0$$

- Solving for v instead of S (remember, v = v(S) is a function of S) means knowledge of kinetic rate parameters in not necessary
- This defines m algebraic equations for n variables. In general however,  $n \gg m$  and therefore the system is **underdetermined**

# The flux cone

- We will assume all reactions are irreversible so that the components of  ${\bf v}$  are non-negative, i.e.  ${\bf v} \ge 0$
- This is always possible to achieve by splitting each reversible reaction into two: a forward and reverse reaction, both irreversible
- The flux vector **v** therefore must satisfy the constraints that define the **flux cone**:

$$\mathbf{N} \cdot \mathbf{v} = 0, \qquad \mathbf{v} \ge 0$$

# The flux cone contd.

• The flux cone is really a cone:



• Each stochiometric constraint is a **linear equation of the fluxes** and defines a **hyperplane** in *n*-dimensional space. The flux cone is the intersection of these with the positive region of *n*-dimensional space

## Bounds on fluxes

- The flux cone is unbounded in that infinite flux values are allowed
- In reality, we know that physical arguments prevent certain reactions proceeding at rates greater than a given upper bound, i.e.

#### $\mathbf{UB} \ge \mathbf{v} \ge \mathbf{0}$

• This further constrains **v** to a **feasibility space** 

### Feasibility space

• The full set of constraints on **v** are

$$\mathbf{N} \cdot \mathbf{v} = 0, \qquad \mathbf{v} \ge 0, \qquad \mathbf{UB} \ge \mathbf{v} \ge 0$$

• These define a (bounded or unbounded) convex polytope



# The objective function

- We have avoided parameterization at a cost of underdetermination. The feasible space defines an infinite set of flux distributions, but we want just one...
- Introducing an **objective function** based on some biological assumptions can help us to find an **optimal flux distribution**
- This turns the problem of finding v into a constrained optimization problem

# The objective function contd.

• An objective function will reduce the set of flux distributions



• Although the optimal value is unique, an optimal flux distribution might not be

# Examples of objective functions

- By far the most common objective is maximization of growth rate μ.
  In bio-engineering, sometimes the experimentalist wants to maximize the rate of production of a certain metabolite (e.g. ethanol)
- In these cases, the objective function is a linear combination of the components of the flux vector and **objective coefficients**

$$f(\mathbf{v}) = \sum_{i=1}^{n} c_i v_i = \mathbf{c} \cdot \mathbf{v}$$

# Examples of objective functions contd.

- Alternative objective functions sometimes considered are not linear (e.g., ratios of linear functions for yields, quadratic objectives in MOMA and for experimental data)
- All objectives usually considered are convex (we know how to solve convex optimization problems!)

### Constrained optimization problem

- The constrained optimization problem for finding an optimal flux distribution  $\boldsymbol{v}^*$  is

Maximize:  $f(\mathbf{v})$ 

Such that:  $\mathbf{N} \cdot \mathbf{v} = 0, \ \mathbf{v} \ge 0, \ \mathbf{UB} \ge \mathbf{v} \ge 0$ 

•  $\mathbf{v}^*$  is a flux distribution that solves this problem

# Linear programming

When f(v) is linear the problem is a Linear Programming (LP) problem and several methods exist for finding a solution



# Linear programming contd.

• LP problems are nice because if an (finite) optimal solution exists it can always found at a vertex



• The **simplex algorithm** for solving LP problems walks around vertices of the convex polytope looking for a solution!

# Flux balance analysis

• Flux balance analysis (FBA) applies the simplex method to find an optimal flux distribution maximizing a linear objective function



• Stochiometric constraints come from the metabolic reaction network, objective function and flux bounds from biological assumptions

### **Constraint-Based Reconstruction and Analysis**

- FBA is just one technique falling under the umbrella of Constraint-Based Reconstruction and Analysis (COBRA)
- Other COBRA techniques are based on further biological assumptions and/or integration with experimental data
- Tools available through the open source COBRA code base:



https://opencobra.github.io/

Open-source, community-developed code base for COnstraint-Based Reconstruction and Analysis.