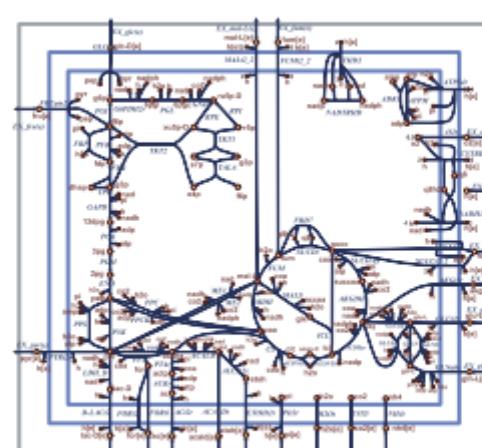


(i)

Genome-scale
metabolic reconstruction



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

(ii) Mathematically represent metabolic reactions and constraints

$$\begin{array}{c}
 \text{Reactions} \\
 \begin{matrix} & 1 & 2 & \dots & n & \text{Biomass} \\
 \text{Metabolites} & A & -1 & & & \\
 & B & 1 & -1 & & \\
 & C & 1 & -2 & & \\
 & D & & 1 & & \\
 & \dots & & & & \\
 & m & & & & \end{matrix} \\
 \text{Stoichiometric matrix, } \mathbf{S} \\
 \end{array}$$

\ast

v_1
 v_2
 \dots
 v_n
 v_{biomass}
 v_{glucose}
 v_{oxygen}

= 0

Fluxes, \mathbf{v}

(iii) Mass balance defines a system of linear equations

$$\begin{aligned}
 -v_1 + \dots &= 0 \\
 v_1 - v_2 + \dots &= 0 \\
 v_1 - 2v_2 + \dots &= 0 \\
 v_2 + \dots &= 0 \\
 \text{etc.}
 \end{aligned}$$

(iv) Define objective function ($Z = c_1^* v_1 + c_2^* v_2 + \dots$)

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

(v)

Calculate fluxes that maximize Z

