





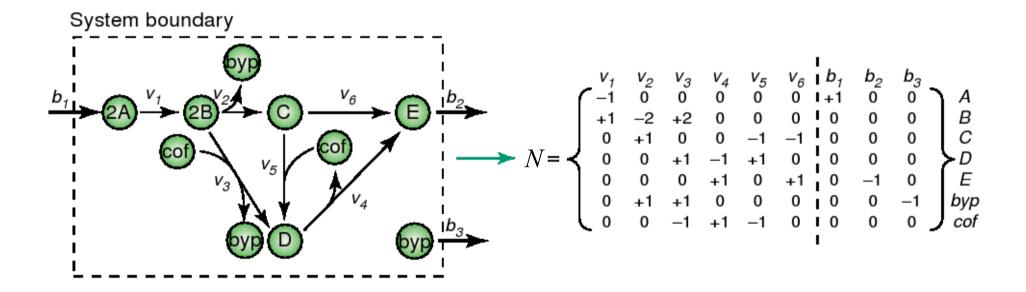
Linear programming applied to genome-scale metabolic network models

Oliver Ebenhöh



Stoichiometric modelling

A metabolic system is defined by internal reactions and exchange fluxes



The temporal change of the concentrations is given by

$$\frac{dX}{dt} = N \cdot v$$

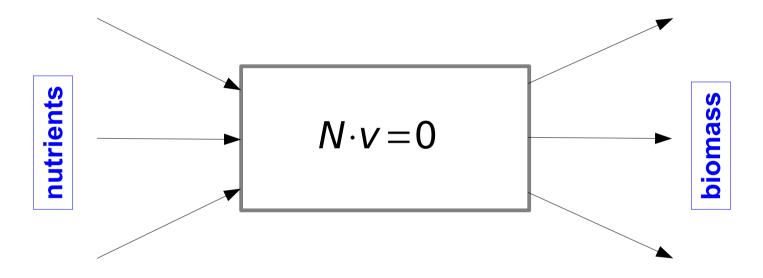
Steady state is characterised by $N \cdot v = 0$

$$N \cdot v = 0$$

$$N \cdot v = 0$$

Problem: This equation has many solutions!

Which one is correct?



- We know what goes in
- We know what goes out
- We do not know what is happening inside!

Constraints allow to reduce the number of possible solutions

Optimisation allows to find special fluxes (and answer 'what if...?')

Linear Programming

Linear Programming (LP) is an optimisation technique

Identify variable values which result in a maximal (or minimal) value of a function which <u>linearly</u> depends on the parameters under given constraints

Introduction

General idea: optimise a linear function under inequality constraints

Variables: x_i , i=1...N

Constraints: $l_i \le x_i \le u_i$

Objective: $\Omega = \sum_{i}^{N} c_{i} \cdot x_{i}$

EXAMPLES

A farmer has 10 acres to plant in wheat and rye. He has to plant at least 7 acres. However, he has only \$1200 to spend and each acre of wheat costs \$200 to plant and each acre of rye costs \$100 to plant. Moreover, the farmer has to get the planting done in 12 hours and it takes an hour to plant an acre of wheat and 2 hours to plant an acre of rye. If the profit is \$500 per acre of wheat and \$300 per acre of rye how many acres of each should be planted to maximize profits?

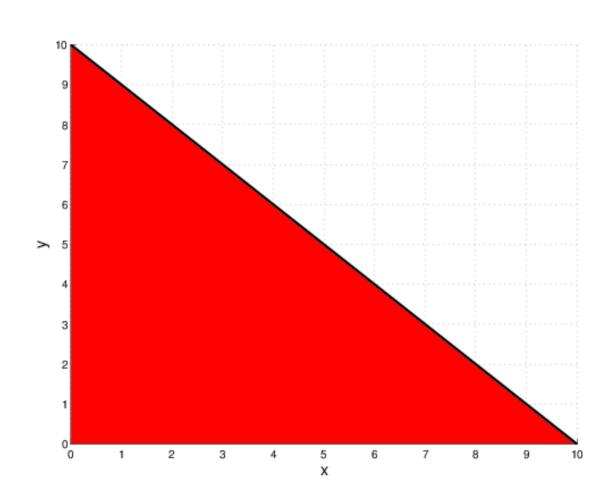
Steps to solve the problem:

- 0. Read the whole problem.
- 1. Define your unknowns.
- 2. Express the objective function and the constraints.
- 3. Graph the constraints.
- 4. Find the corner points to the region of feasible solutions.
- 5. Evaluate the objective function at all the feasible corner points.

see http://www.sonoma.edu/users/w/wilsonst/courses/math_131/lp/

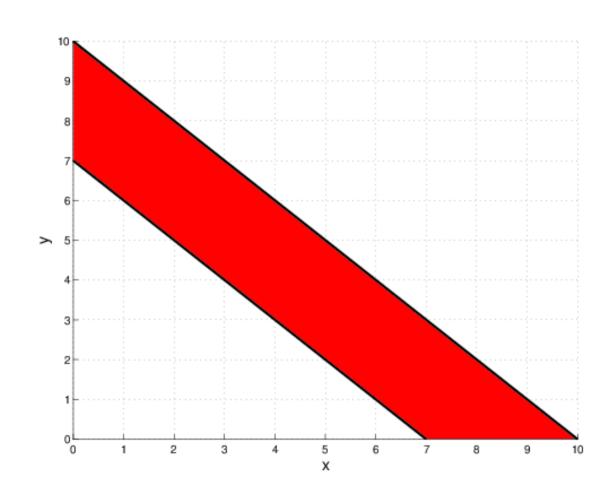
Total land:

$$x+y \leq 10$$



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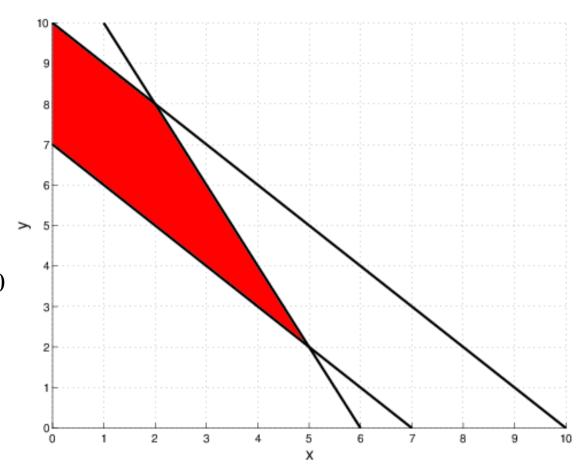
Minimum planting: $x+y \ge 7$



Total land: $x+y \le 10$

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Limited funds: $200 x + 100 y \le 1200$

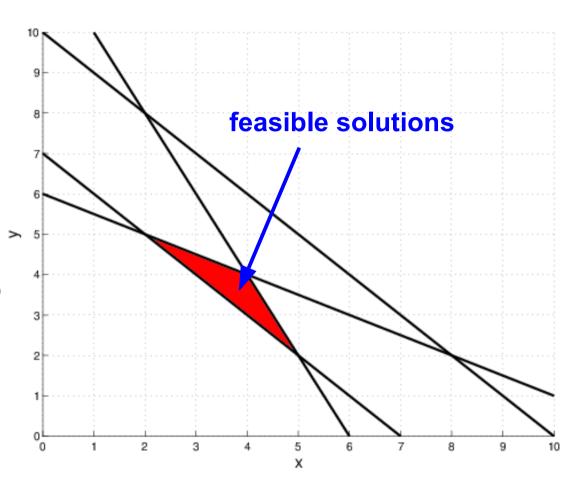


Total land: $x+y \le 10$

Minimum planting: $x+y \ge 7$

Limited funds: $200 x + 100 y \le 1200$

Limited time: $x+2 y \le 12$



Which one of the feasible solutions gives the most profit?

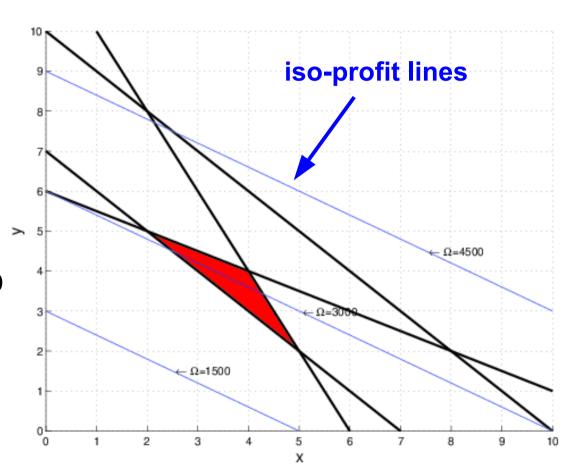
Profit: $\Omega = 500x + 300y$

Total land: $x+y \le 10$

Minimum planting: $x+y \ge 7$

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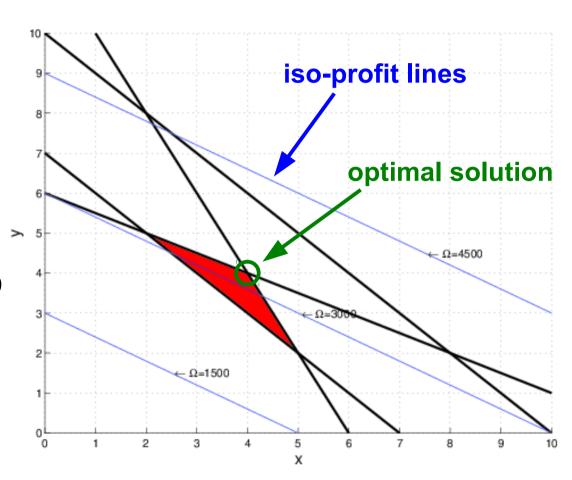
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Maximal profit at: x=4, y=4

In general:

- space of feasible solution is a convex polyhedron
- optimal solution is <u>always</u> at a vertex

A week has 168 hours

- We need time to study (S), party (P) and for everything else (E incl. sleep, eat)
- To survive, we need at least 8h rest per day: *E* ≥ 56
- To maintain sanity, we need to party or rest a bit more: *P*+*E* ≥ 70
- To pass exams, we need to study at least 60h/week: S ≥ 60
- But longer if we don't sleep enough or party too much: 2S+E-3P ≥ 150
 (this means, for every missed hour of sleep, we need to study 30 min longer and for every hour partying, we need to study 1.5h longer because of hangovers)

Objective: Maximise happyness, expressed by $\Omega = 2P + E$ (extra rest makes happy, partying makes twice as happy)

The problem:

maximise

(no negative times) $P \ge 0$

$$\Omega = 2P + E$$

under the constraints that

Eliminate S

 $P \ge 0$

(week)

$$S+P+E=168$$
 $S=168-P-E$

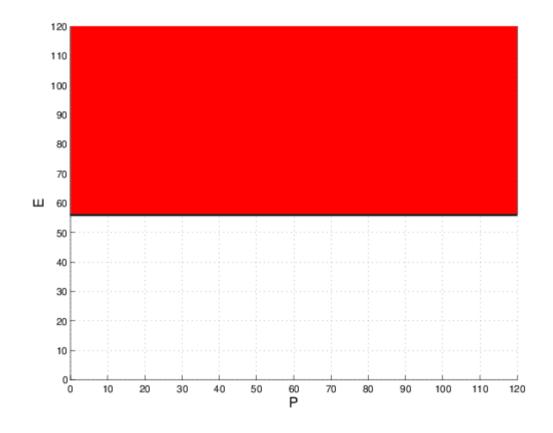
 (survival)
 $E \ge 56$
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 (min. study)
 $S \ge 60$
 $168-P-E \ge 60 \Leftrightarrow P+E \le 108$

 (hangovers)
 $2S+E-3P \ge 150$
 $2(168-P-E)+E-3P \ge 150 \Leftrightarrow E+5P \le 186$

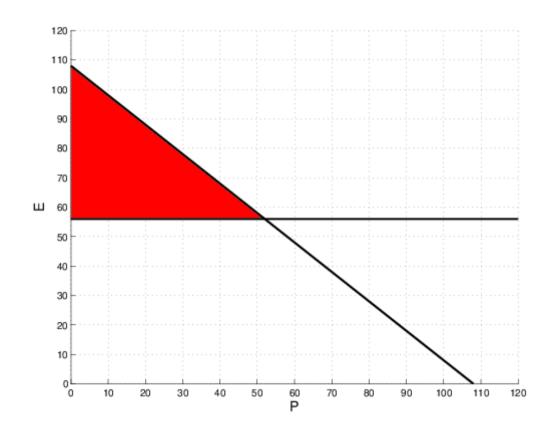
 (sanity)
 $P+E \ge 70$

Survival: $E \ge 56$



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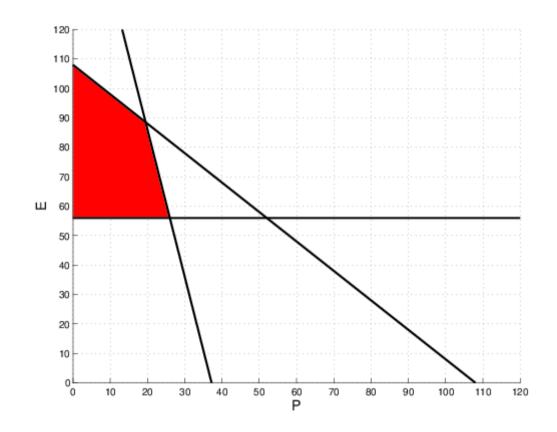
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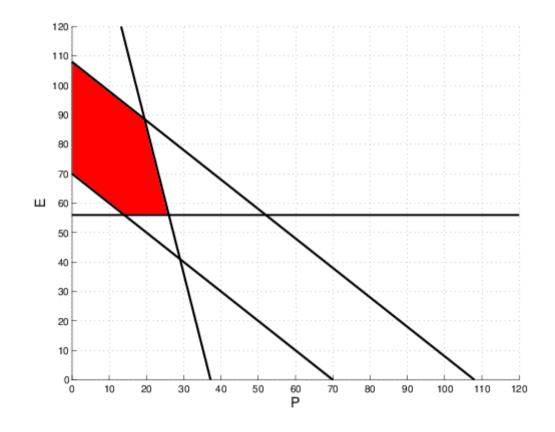


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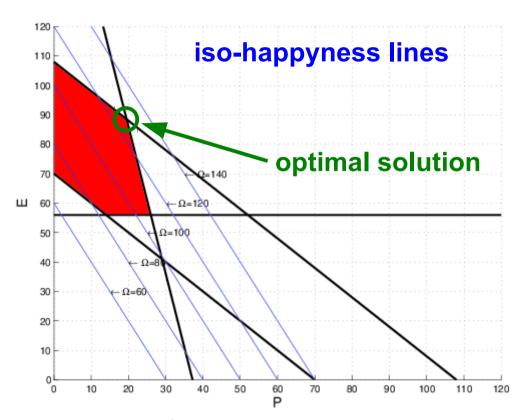


Survival: $E \ge 56$

minimal study: $P + E \le 108$

hangovers: $E + 5P \le 186$

sanity: $P+E \ge 70$



Calculate optimal solution by finding intersection of two lines:

$$186-5P=108-P \Leftrightarrow P=19.5$$

$$E=108-P \Rightarrow E=88.5$$

$$S=168-P-E \Rightarrow S=60$$

Application to metabolic networks

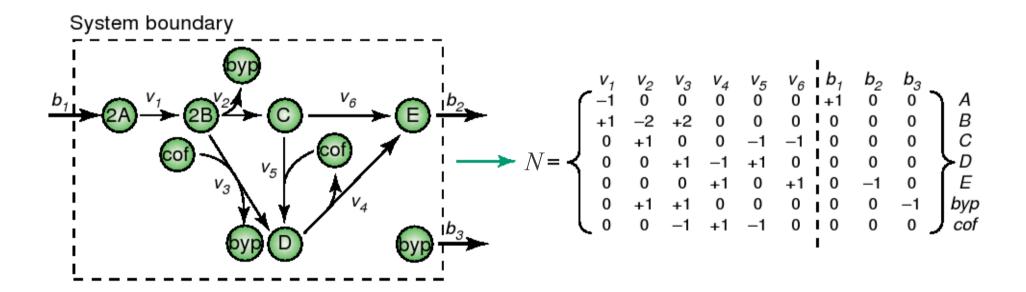
Variables: FLUXES v_i , i=1...R

Constraints: ?

Objective: ?

Application to metabolic networks

A metabolic system is defined by *internal reactions* and *exchange fluxes*



The temporal change of the concentrations is given by

$$\frac{dX}{dt} = N \cdot v$$

Steady state is characterised by

$$N \cdot v = 0$$

Constraint #1

Other constraints

Because of thermodynamic reasons, some reactions can only proceed in one direction

E.g. Glc+ATP
$$\rightarrow$$
G6P+ADP $\Delta G^0 = -24.9 \text{ kJ/mole}$

$$K_{\text{eq}} = \frac{[\text{ADP}]_{\text{eq}} \cdot [\text{G6P}]_{\text{eq}}}{[\text{ATP}]_{\text{eq}} \cdot [\text{Glc}]_{\text{eq}}} = e^{-\Delta G^0/RT} = e^{10.05} = 23000$$

With [ATP]/[ADP]=3 and $[Glc]=1 \,\mathrm{mM}$ the reaction runs in reverse if

$$[G6P] > K_{eq} \cdot [Glc] \cdot \frac{[ATP]}{[ADP]} = 69000 \,\text{mM} = 69 \,\text{M} \quad !!!$$

(pure water has 55.5 M)



directionality implies $v_j \ge 0$ Constraints #2

$$v_i \ge 0$$

Other constraints

Some process have <u>upper bounds</u>

- maximal uptake rates
- known maximal enzyme activities



limitation implies $v_i \le v_j^{\text{max}}$ Constraints #3

$$v_i \leq v_i^{\max}$$

What is constraint based modelling?

Fluxes in metabolic networks are subject to *constraints*

- Thermodynamic (directionality)
- $V_i \ge 0$

Enzyme concentrations

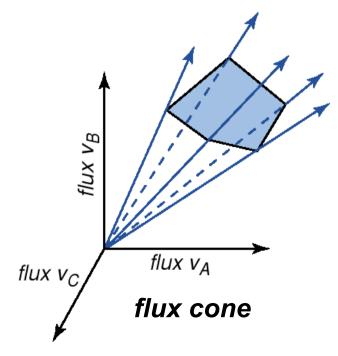
$$V_i \leq V_{i,max}$$

Constraint based models analyse steady state solutions which fulfill the given constraints.

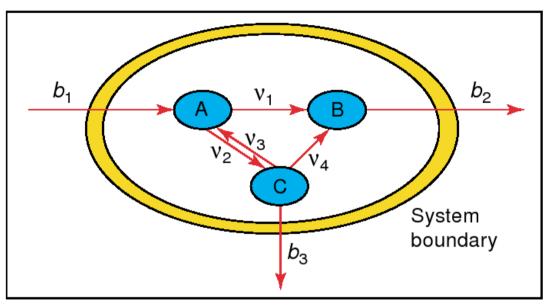
Find a solution vector
$$\mathbf{v} = (\mathbf{v}_1, \dots, \mathbf{v}_r)^T$$
 such that

$$N \cdot v = 0$$

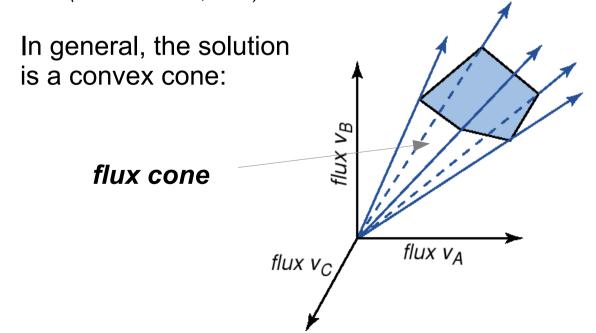
and
$$a_i \leq v_i \leq b_i$$



Example: constraint based model



(Kauffman et. al, 2003)



constraints

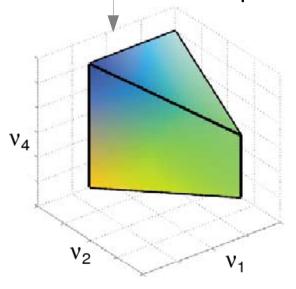
$$v_3 = 0$$
 (thermodynamic)

$$0 \le b_1 \le 1 \Rightarrow 0 \le v_1 + v_2 \le 1$$

$$0 \le b_2 \le 2 \Rightarrow 0 \le v_1 + v_4 \le 2$$

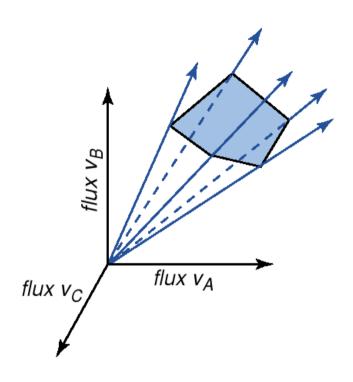
$$0 \le b_3 \Rightarrow 0 \le v_2 - v_4$$

Solution space



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

Which solution?



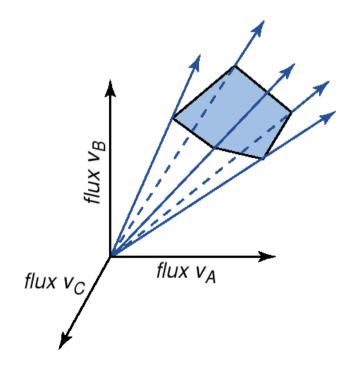
Application to metabolic networks

Variables: FLUXES v_i , i=1...R

Constraints: Stationarity, maximal rates $N \cdot v = 0$, $0 \le v_i \le v_i^{\text{max}}$

Objective: ?

The whole purpose of linear programming is to find <u>one flux distribution</u> from the solution cone which is "optimal"



What is optimal?

No general answer!

Plausible assumptions:

- maximal growth / biomass production
- most 'economic' solution (minimal enzyme usage)

Even if the objective is not 'correct', the computation is useful:

We can investigate the question "what if...?"

A typical LP problem maximising biomass

- assemble r x n stoichiometry matrix N (r reactions, n metabolites)
- identify irreversible reactions $R \subset \{1...r\}$
- define boundary fluxes $B \subset \{1...r\}$
- define "biomass reaction" v_{biomass} : $\sum_{i} \alpha_{i} \cdot S_{i} \rightarrow \text{biomass}$

Example from *E.coli* model (Feist et al. 2007)

(54.613) cpd00001 + (59.98) cpd00002 + (0.001787) cpd00003 + (0.000045) cpd00004 + (0.000335) cpd00005 + (0.000112) cpd00006 + (0.000168) cpd00010 + (0.01128) cpd00013 + (0.000223)+ (0.4991) cpd00035 + (0.2091) cpd00038 + (0.3334) cpd00039 + (0.2342) cpd00041 + (0.000223) cpd00042 + (0.00376) cpd00048 + (0.2874) cpd00051 + (0.1298) cpd00052 + (0.2557) cpd00053 +(0.2097) cpd00054 +(0.000223) cpd00056 +(0.03008) cpd00058 +(0.1493) cpd00060 +(0.1401) cpd00062 +(0.004512) cpd00063 +(0.05523) cpd00065 +(0.18) cpd00066 +(0.134) $\verb|cpd00069+(0.000031)| \verb|cpd00070+(0.000098)| \verb|cpd00078+(0.08899)| \verb|cpd000084+(0.000223)| \verb|cpd000087+(0.004512)| \verb|cpd00099+(0.4378)| \verb|cpd00107+(0.02481)| \verb|cpd000115+(0.03327)| \verb|cpd000118+(0.03327)| | |cpd000118+(0.03327)| | |cpd000118+(0.03327$ + (0.0921) cpd00119 + (0.000223) cpd00125 + (0.2148) cpd00129 + (0.2342) cpd00132 + (0.03008) cpd00149 + (0.1542) cpd00155 + (0.4119) cpd00156 + (0.2465) cpd00161 + (0.000223) cpd00166 + (0.000223) cpd00201 + (0.1692) cpd00205 + (0.000223) cpd00216 + (0.000223) cpd00220 + (0.02561) cpd00241 + (0.007519) cpd00254 + (0.006744) cpd00264 + (0.2823) cpd00322+ (0.000223) cpd00345 + (0.02561) cpd00356 + (0.02481) cpd00357 + (0.000223) cpd00557 + (0.000055) cpd00229 + (0.000223) cpd003453 + (0.006767) cpd10515 + (0.006767) cpd10516 + (0.000223) cpd11313 + (0.003008) cpd11574 + (0.000223) cpd15353 + (0.002944) cpd15428[p] + (0.00229) cpd15429[p] + (0.00118) cpd15431[p] + (0.008151) cpd15432[e] + (0.000223) cpd15499 + (0.001345) cpd15501[p] + (0.006605) cpd15503[p] + (0.005381) cpd15505[p] + (0.005448) cpd15506[p] + (0.00673) cpd15508[p] + (0.0318) cpd15531[p] + (0.02473) cpd15532[p] + (0.01275) cpd15534[p] + (0.004897) cpd15538[p] + (0.003809) cpd15539[p] + (0.001963) cpd15541[p] + (0.000223) cpd15561 => (59.81) cpd00008 + (58.8062) cpd00009 + (0.7498) cpd00012 + (59.81) cpd000012 + (59.81) cpd00008 + (58.8062) cpd00009 + (0.7498) cpd000012 + (59.81) cpd00008 + (58.8062) cpd00009 + (0.7498) cpd000012 + (59.81) cpd00008 + (58.8062) cpd000009 + (0.7498) cpd000012 + (59.81) cpd000008 + (58.8062) cpd000009 + (0.7498) cpd000012 + (59.81) cpd000008 + (58.8062) cpd000009 + (0.7498) cpd000012 + (59.81) cpd000008 + (58.8062) cpd000009 + (0.7498) cpd000012 + (59.81) cpd000008 + (58.8062) cpd000009 + (0.7498) cpd000012 + (59.81) cpd000008 + (58.8062) cpd000009 + (0.7498) cpd000012 + (59.81) cpd000000 + (0.7498) cpd000012 + (59.81) cpd000000 + (0.7498) cpd00000000 + (0.7498) cpd000000000000 + (0.7498) cpd000000000000 + (0.7498) cpd000000cpd00067 + cpd11416

• define upper bounds for uptake rates (boundary fluxes): $v_i \le v_i^{\text{max}}$ for $i \in B$

The LP-problem:

maximise

 $\mathcal{V}_{\text{biomass}}$

Result:

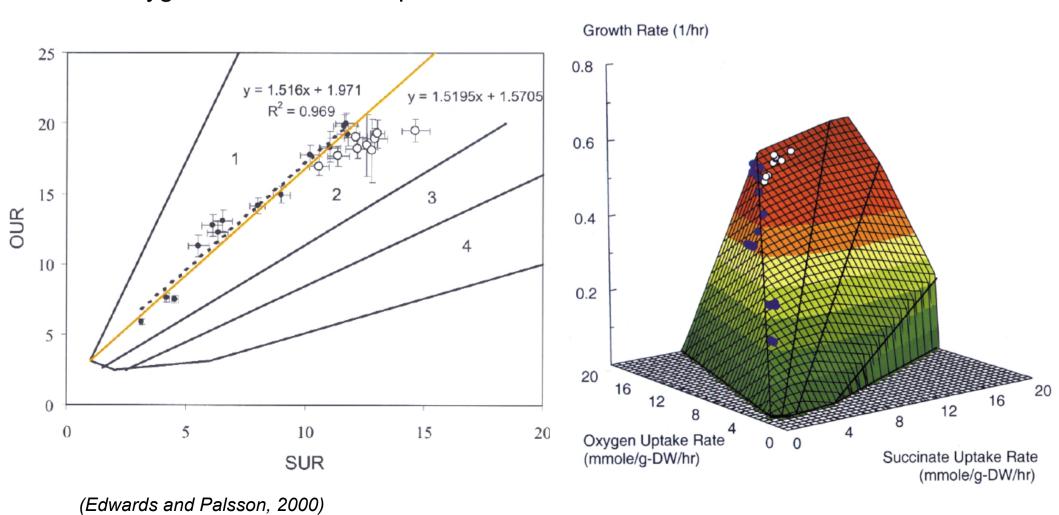
under the constraints $N \cdot v = 0$

 $v_i \leq v_i^{\text{max}} \text{ for } i \in B$ $v_i \ge 0$ for $i \in R$

Flux distribution v

Optimality studies in *E. coli*

- E. coli was grown on succinate
- Optimal growth rates were predicted as extreme fluxes
- Oxygen and succinate uptake rates were measured



A typical LP problem minimising costs

- assemble r x n stoichiometry matrix N (r reactions, n metabolites)
- identify irreversible reactions $R \subset \{1...r\}$
- define boundary fluxes $B \subseteq \{1...r\}$
- define "biomass reaction" v_{biomass} : $\sum_{i} \alpha_{i} \cdot S_{i} \rightarrow \text{biomass}$
- Fix biomass (e.g. from experiments) $v_{\text{biomass}} = v_{\text{biomass}}^{\text{exp}}$

The LP-problem: minimise
$$\sum_{i}^{r} w_{i} \cdot v_{i}$$
 under the constraints $N \cdot v = 0$ $v_{\text{biomass}} = v_{\text{biomass}}^{\text{exp}}$ $v_{i} \ge 0$ for $i \in R$

Variation of constraints to query the model

Objective: study how optimal fluxes change upon perturbation of external conditions Example: impose additional ATP demand (reflecting e.g. external stress conditions)

 \longrightarrow Additional constraint $v_{\text{ATPdemand}} = \gamma$ tunable parameter

(Additional ATP consuming process: $ATP + H_2O \rightarrow ADP + P_i$) RIBULOSE BISPHOSPHATE CARBOXYLASE RXN
PHOSPHORIBULOKINASE RXN ICOŚANOYL-COA-SYNTHASE-RXN-(NADP GLU6PDEHYDROG-RXN 8.0e-05 7.0e-05 5.0e-05 6.0e-05 4.0e-05 5.0e-05 4.0e-05 3.0e-05 3.0e-05 2 Op-05 1.0e-05 1.0e-05 8.0e-04 4.0e-04 8.0e-04 1.2e-03 1.6e-03 1.2e-03 1.6e-03 9.0e-06 RIB5PISOM-RXN 1.4e-04 8.0e-06 7.0e-06 1 0e-04 6.0e-06 5.0e-06 6.0e-05 4.0e-06 3.00-06 1.6e-03 2.3e-04 1.2e-0 NADOxid 2.2e-04 1.0e-04 CisAcon 2.1e-04 8.0e-05 2.0e-04 1.9e-04 4.0e-05 1.8e-04 2.0e-05 1.7e-04 (Poolman et al. 2009) - SucCoA

Simulating availability of nitrogen sources

The LP-problem:

minimise

$$\gamma \cdot v_{\text{NO3_uptake}} + \sum_{i}^{r} v_{i}$$

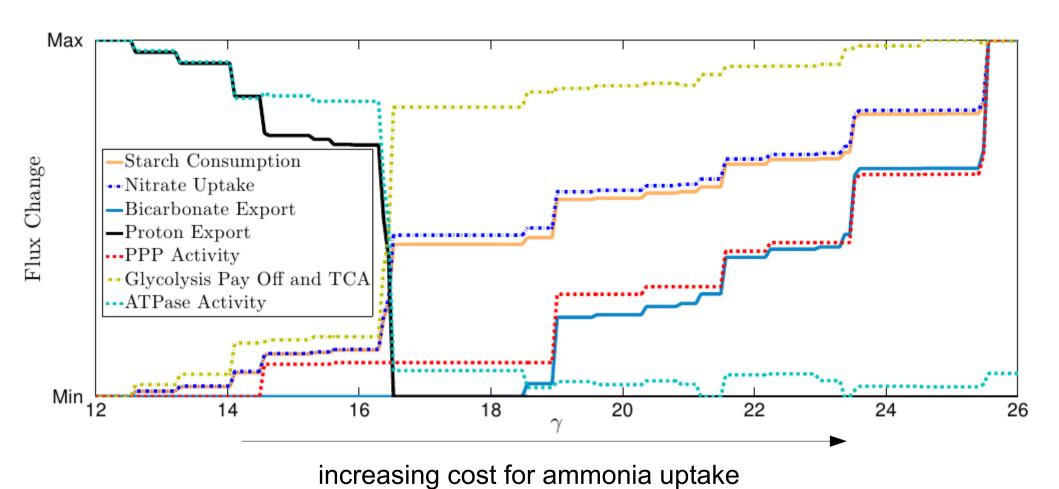
under the constraints

$$N \cdot v = 0$$

$$v_{\text{biomass}} = v_{\text{biomass}}^{\text{exp}}$$

$$v_i \ge 0$$
 for $i \in R$

Simulating availability of nitrogen sources



Results for a network of Medicago truncatula

"What if" questions

Assume, we want to know what is the 'cheapest' metabolic route to produce a certain compound X

Add consuming reaction $v_X : X \rightarrow \emptyset$

Define 'cheap'

- minimal energy requirement (ATP)
- minimal redox requirement (NADPH)

minimise
$$w_1 v_{ATPpro}$$

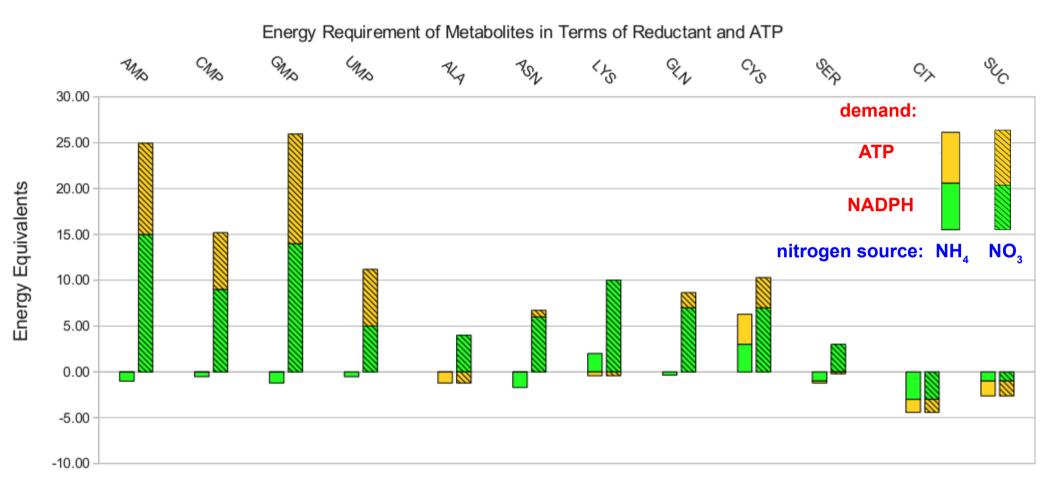
$$w_1 v_{\text{ATPproduction}} + w_2 v_{\text{NADPHproduction}}$$

under the constraints

$$N \cdot v = 0$$

$$v_X = 1$$

$$v_j \ge 0$$
 for $i \in R$



Results for a network of *Medicago truncatula*