

Constraint-based Modeling of Cellular Networks

Exercise 8 – ROOM

08. 12. 2022

ROOM

Step 1: FBA to find wild type flux distribution w

$$\max c^T w$$

$$s. t.$$

$$Nw = 0$$

$$w_{min} \leq w \leq w_{max}$$

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Step 2: Determine the significance thresholds for the flux changes (w^u and w^l).

$\delta = 0.05$ and $\varepsilon = 0.001$

for $1 \leq i \leq r \rightarrow r$ being the number of reactions

$$w_i^u = w_i + \delta |w_i| + \varepsilon$$

$$w_i^l = w_i - \delta |w_i| - \varepsilon$$

Q: What is the dimension of w^u and w^l ?

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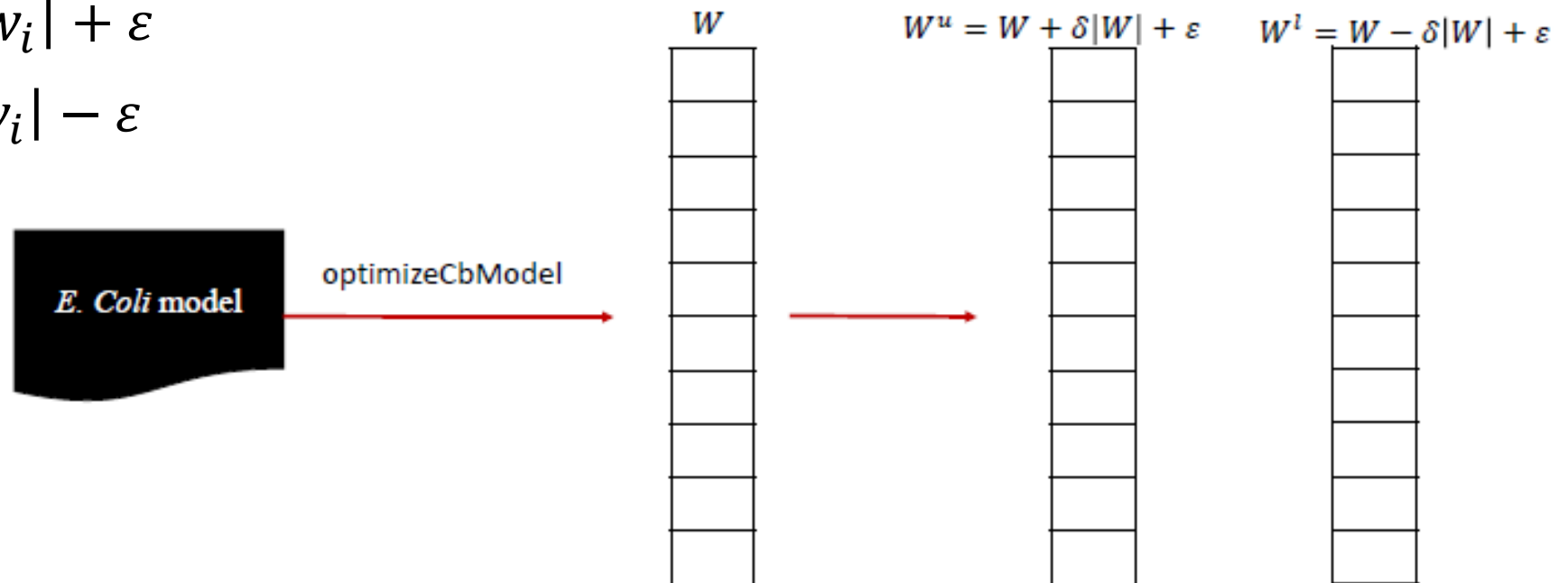
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for $1 \leq i \leq r \rightarrow r$ being the number of reactions

$$w_i^u = w_i + \delta|w_i| + \varepsilon$$

$$w_i^l = w_i - \delta|w_i| - \varepsilon$$



ROOM

Step 3: Put inequality constraints in a matrix form for the optimization problem.

Q: What are the variables, we try to estimate?

for $1 \leq i \leq r$

$$v_i - y_i(v_{\max,i} - w_i^u) \leq w_i^u$$

$$v_i - y_i(v_{\min,i} - w_i^l) \geq w_i^l$$

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Step 3: Put inequality constraints in a matrix form for the optimization problem.

Here the variables for the optimization problem are the vectors of v and y .

for $1 \leq i \leq r$

$$v_i - y_i(v_{max,i} - w_i^u) \leq w_i^u$$

$$v_i - y_i(v_{min,i} - w_i^l) \geq w_i^l$$

variable vector

$$\begin{bmatrix} v_1 \\ \vdots \\ v_r \\ y_1 \\ \vdots \\ y_r \end{bmatrix}$$

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for $1 \leq i \leq r$

$$v_i - y_i(v_{max,i} - w_i^u) \leq w_i^u$$

$$\begin{array}{ccccccc}
 & v_1 & \dots & v_r & & y_1 & \dots & y_r \\
 i = 1 & \left[\begin{array}{ccccccc}
 1 & 0 & 0 & -(v_{max,1} - w_1^u) & 0 & & 0 \\
 \dots & & & & & & \\
 0 & 0 & 1 & 0 & 0 & -(v_{max,r} - w_r^u)
 \end{array} \right] & \cdot & \begin{array}{c} v_1 \\ \dots \\ v_r \\ y_1 \\ \dots \\ y_r \end{array} \leq w_i^u
 \end{array}$$

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Step 3: Put inequality constraints in a matrix form for the optimization problem.

Here the variables for the optimization problem are the vectors of v and y .

for $1 \leq i \leq r$

$$v_i - y_i(v_{max,i} - w_i^u) \leq w_i^u$$

Implement it similarly for

$$v_i - y_i(v_{min,i} - w_i^l) \geq w_i^l$$

$$\begin{array}{ccccccc}
 & v_1 & \dots & v_r & & y_1 & \dots & y_r \\
 i = 1 & \left[\begin{array}{ccccccc}
 1 & 0 & 0 & -(v_{max,1} - w_1^u) & 0 & & 0 \\
 \dots & & & & & & \\
 0 & 0 & 1 & 0 & 0 & -(v_{max,r} - w_r^u)
 \end{array} \right] & \cdot & \begin{array}{c} v_1 \\ \dots \\ v_r \\ y_1 \\ \dots \\ y_r \end{array} \leq w_i^u
 \end{array}$$

ROOM

$$\min_{y,v} \sum_{i=1}^r y_i$$

s.t

$$Sv = 0, \quad v_{\min} \leq v \leq v_{\max}$$

$v_j = 0$, for $\forall j$ blocked by gene knock – out

for $1 \leq i \leq r$

$$v_i - y_i(v_{\max,i} - w_i^u) \leq w_i^u$$

$$v_i - y_i(v_{\min,i} - w_i^l) \geq w_i^l$$

$$y_i \in \{0,1\}$$

ROOM

$$\begin{aligned} \min_{y,v} \quad & \sum_{i=1}^r y_i \\ \text{s.t} \quad & \end{aligned}$$

$$Nv = 0, \quad v_{min} \leq v \leq v_{max}$$

$$v_j = 0, \text{ for } \forall j \text{ blocked by gene knock-out}$$

$$\text{for } 1 \leq i \leq r$$

$$\begin{aligned} v_i - y_i(v_{max,i} - w_i^u) &\leq w_i^u \\ v_i - y_i(v_{min,i} - w_i^l) &\geq w_i^l \\ y_i &\in \{0,1\} \end{aligned}$$

$$\begin{bmatrix} \mathbf{N} & \begin{matrix} 0 & 0 & 0 \\ 0 & 0 & 0 \\ 0 & 0 & 0 \end{matrix} \end{bmatrix} \cdot \begin{matrix} v_1 \\ \dots \\ v_r \\ y_1 \\ \dots \\ y_r \end{matrix} = 0$$

$$\begin{array}{ccccccc}
v_1 & \dots & v_r & & y_1 & & \dots & & y_r \\
i = 1 & \left[\begin{array}{ccccccc} 1 & 0 & 0 & -(v_{max,1} - w_1^u) & 0 & & 0 \end{array} \right. \\
\dots & & & & & & & & \\
i = r & \left[\begin{array}{ccccccc} 0 & 0 & 1 & & 0 & & -(v_{max,r} - w_r^u) \end{array} \right. \\
& & & & & & & & \\
& & & & & & & &
\end{array} \cdot \begin{array}{c} v_1 \\ \dots \\ v_r \\ y_1 \\ \dots \\ y_r \end{array} \leq w_i^u$$

$$\begin{matrix} i = 1 \\ \vdots \\ i = r \end{matrix} \begin{bmatrix} 1 & 0 & 0 & -(v_{min,1} - w_1^l) & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ 0 & 0 & 1 & 0 & 0 & -(v_{min,r} - w_r^l) \end{bmatrix} \cdot \begin{matrix} v_1 \\ \vdots \\ v_r \\ y_1 \\ \vdots \\ y_r \end{matrix} \geq w_i^u$$

ROOM

$$\min_{y,v} \sum_{i=1}^r y_i$$

s.t

$$Nv = 0, \quad v_{\min} \leq v \leq v_{\max}$$

$v_j = 0$, for $\forall j$ blocked by gene knock – out

for $1 \leq i \leq r$

$$\begin{aligned} v_i - y_i(v_{\max,i} - w_i^u) &\leq w_i^u \\ v_i - y_i(v_{\min,i} - w_i^l) &\geq w_i^l \\ y_i &\in \{0, 1\} \end{aligned}$$

MILP problem!

Use intlinprog() to solve

$$\begin{bmatrix} v_1 & \dots & v_r & y_1 & \dots & y_r \\ & & & 0 & 0 & 0 \\ & & & 0 & 0 & 0 \\ & & & 0 & 0 & 0 \end{bmatrix} \cdot \begin{bmatrix} v_1 \\ \dots \\ v_r \\ y_1 \\ \dots \\ y_r \end{bmatrix} = 0$$

$$\begin{matrix} & v_1 & \dots & v_r & & y_1 & \dots & y_r \\ i = 1 & \left[\begin{array}{cccccc} 1 & 0 & 0 & -(v_{\max,1} - w_1^u) & 0 & 0 \end{array} \right] & \begin{matrix} v_1 \\ \dots \\ v_r \\ y_1 \end{matrix} & \leq w_i^u \\ \dots & & & & & & & \\ i = r & \left[\begin{array}{cccccc} 0 & 0 & 1 & 0 & 0 & -(v_{\max,r} - w_r^u) \end{array} \right] & \begin{matrix} \dots \\ y_1 \\ \dots \\ y_r \end{matrix} \end{matrix}$$

$$\begin{matrix} & v_1 & & & & & & \\ i = 1 & \left[\begin{array}{cccccc} 1 & 0 & 0 & -(v_{\min,1} - w_1^l) & 0 & 0 \end{array} \right] & \begin{matrix} \dots \\ v_r \\ y_1 \end{matrix} & \geq w_i^l \\ \dots & & & & & & & \\ i = r & \left[\begin{array}{cccccc} 0 & 0 & 1 & 0 & 0 & -(v_{\min,r} - w_r^l) \end{array} \right] & \begin{matrix} \dots \\ y_1 \\ \dots \\ y_r \end{matrix} \end{matrix}$$

intlinprog Mixed integer linear programming.

$X = \text{intlinprog}(f, \text{intcon}, A, b)$ attempts to solve problems of the form

$$\begin{array}{ll} \min f'x & \text{subject to: } A*x \leq b \\ x & Aeq*x = beq \\ & lb \leq x \leq ub \\ & x(i) \text{ integer, where } i \text{ is in the index} \\ & \text{vector intcon (integer constraints)} \end{array}$$

$X = \text{intlinprog}(f, \text{intcon}, A, b)$ solves the problem with integer variables in the `intcon` vector and linear inequality constraints $A*x \leq b$. `intcon` is a vector of positive integers indicating components of the solution X that must be integers. For example, if you want to constrain $X(2)$ and $X(10)$ to be integers, set `intcon` to `[2,10]`.

$X = \text{intlinprog}(f, \text{intcon}, A, b, Aeq, beq)$ solves the problem above while additionally satisfying the equality constraints $Aeq*x = beq$. (Set `A=[]` and `b=[]` if no inequalities exist.)

$X = \text{intlinprog}(f, \text{intcon}, A, b, Aeq, beq, LB, UB)$ defines a set of lower and upper bounds on the design variables, X , so that the solution is in the range $LB \leq X \leq UB$. Use empty matrices for `LB` and `UB` if no bounds exist. Set `LB(i) = -Inf` if $X(i)$ is unbounded below; set `UB(i) = Inf` if $X(i)$ is unbounded above.

How to find reactions associated with a gene?

Use the GPR rules in the model structure to find related genes

$$\min_{y,v} \sum_{i=1}^r y_i$$

s.t

Use model field rxnGeneMat

$$Nv = 0, \quad v_{\min} \leq v \leq v_{\max}$$

You can also check the Cobra Toolbox function *findRxnsFromGenes*

$v_j = 0$, for $\forall j$ blocked by gene knock – out

for $1 \leq i \leq r$

$$\begin{aligned} v_i - y_i(v_{\max,i} - w_i^u) &\leq w_i^u \\ v_i - y_i(v_{\min,i} - w_i^l) &\geq w_i^l \\ y_i &\in \{0,1\} \end{aligned}$$