# Constraint-based modeling of cellular networks WS 2023/2024

Master of Bioinformatics University of Potsdam

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#### **Last lecture**

Flux variability analysis feasible or operational ranges Flux variability and classification of reactions blocked reactions Flux variability to generate alternative optima Flux sampling Flux sampling and reaction classification shape probability distribution correlation between sampled fluxes

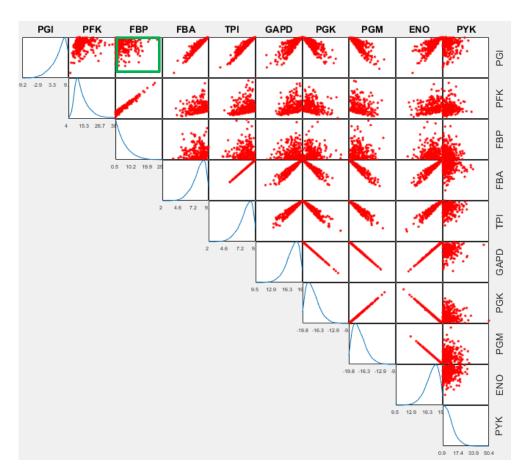
#### Flux sampling and classification of reactions

Pairs of reactions can be classified based on the correlation between their respective fluxes

#### Scatter plot

#### **Correlation of 1**

Correlation of -1
Moderate correlation
No significant or
small correlation



#### Flux sampling and classification of reactions

Flux sampling is costly
three algorithms for sampling
(could be affected by bias, depending on implementation)

#### **Questions**

1. Is there a more efficient way to determine reactions whose fluxes show correlation of 1?

do we have to sample or one LP is sufficient?

- 2. Can we specify how the structure of the metabolic network leads to dependencies between reaction fluxes?
- 3. What are the biological implications of such dependencies?

### Flux coupling analysis (FCA) questions

#### **Questions**

1. Is there a more efficient way to determine reactions whose fluxes show correlation of 1?

do we have to sample or one LP is sufficient?

- 2. Can we specify how the structure of the metabolic network leads to dependencies between reaction fluxes?
- 3. What are the biological implications of such dependencies?

These questions can be answered by

Flux Coupling Analysis (FCA)

Main goal

Classify reaction pairs based on their dependencies.

The dependencies between reactions are called **reaction couplings**.

We will show that this classification can be achieved by a formulation which resembles Flux Variability Analysis

Main goal
Classify reaction pairs based on their couplings.

Three types of reaction couplings

Directional coupling vi!= 0 => vj!=0

Partial coupling vi!= 0 => vj!=0, with vi/vj!= const

Full coupling vi/vj = vij, s.t. for every v, Nv=0

Reaction pairs which are not directionally, partially, or fully coupled will be called **uncoupled**.

### **Directional coupling**

#### **Definition**

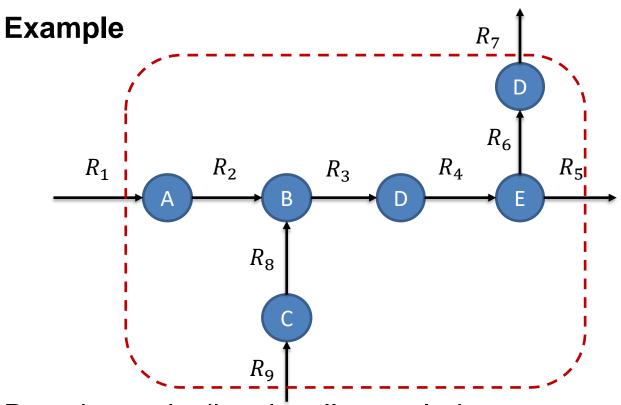
A reaction  $R_i$  is directionally coupled to reaction  $R_j$  if for every steady-state flux distribution non-zero steady-state flux for  $R_i$  implies a non-zero steady-state flux for  $R_j$  but necessarily the reverse.

#### Mathematical

A reaction  $R_i$  is directionally coupled to reaction  $R_j$  if for every v with Nv = 0,  $v_i \neq 0$  implies  $v_j \neq 0$ , but not the reverse.

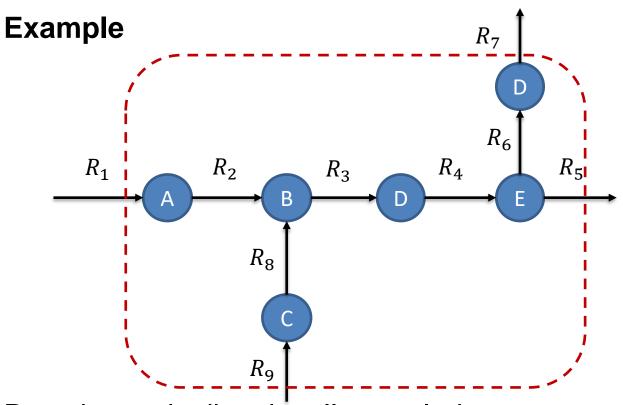
**Notation**  $R_i \rightarrow R_j$  or  $i \rightarrow j$ 

### **Directional coupling**



Reaction  $R_2$  is directionally coupled to  $R_3$ Denoted by  $R_2 \rightarrow R_3$ 

#### **Directional coupling**



Reaction  $R_2$  is directionally coupled to  $R_3$ Denoted by  $R_2 \rightarrow R_3$ 

$$\frac{dc_B}{dt} = v_2 + v_8 - v_3$$

$$v_2 + v_8 - v_3 = 0$$

If  $v_2 \neq 0$ , then  $v_3 \neq 0$  (observe the signs)

### Partial coupling

#### **Definition**

A reaction  $R_i$  is partially coupled to reaction  $R_j$  if non-zero flux for  $R_i$  implies a non-zero, though variable, flux for  $R_j$  and vice versa.

#### Mathematical

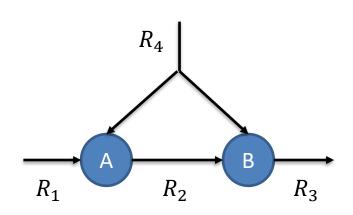
A reaction  $R_i$  is partially coupled to reaction  $R_j$  if for every v with Nv = 0,  $v_i \neq 0$  implies  $v_j \neq 0$ , with  $v_i/v_j \neq const$  and vice versa.

**Notation**  $R_i \leftrightarrow R_j$  or  $i \leftrightarrow j$ 

#### **Example**

It turns out that arriving at an example for partially coupled reactions is a non-trivial task

### Consider the following:



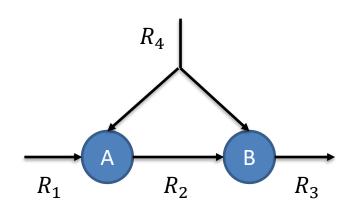
$$\frac{dc_A}{dt} = v_1 + v_4 - v_2$$

$$\frac{dc_B}{dt} = v_2 + v_4 - v_3$$

#### **Example**

It turns out that arriving at an example for partially coupled reactions is a non-trivial task

#### Consider the following:



$$v_1 + v_4 - v_2 = 0$$
 (1)

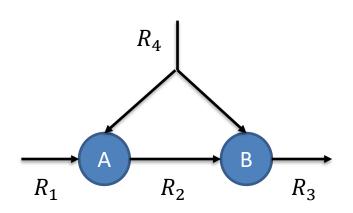
$$v_2 + v_4 - v_3 = 0$$
 (2)

If 
$$v_2 \neq 0$$
 then  $v_3 \neq 0$  (from 2)

#### **Example**

It turns out that arriving at an example for partially coupled reactions is a non-trivial task

#### Consider the following:



$$v_1 + v_4 - v_2 = 0$$
 (1)

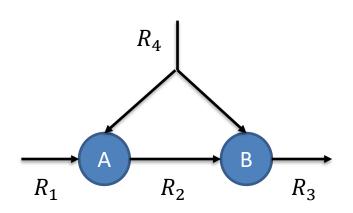
$$v_2 + v_4 - v_3 = 0$$
 (2)

If  $v_3 \neq 0$  then either  $v_2 \neq 0$  or  $v_4 \neq$ 0 (from 2) If  $v_2 \neq 0$ , done! If  $v_4 \neq 0$ , then  $v_2 \neq 0$ (from 1)

#### **Example**

It turns out that arriving at an example for partially coupled reactions is a non-trivial task

#### Consider the following:



 $v_3 \neq 0$  if and only if  $v_2 \neq 0$ 

Note that partial coupling arises as a result of the network as a whole

#### **Emergent property**

#### **Emergent property**

Property of which holds as a result of the network as a whole, rather than some characteristics of a subnetwork or a component itself

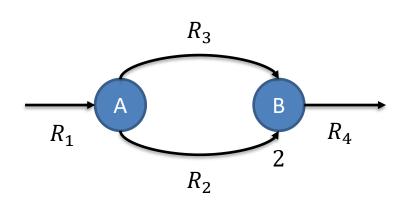
Partial coupling is an emergent property

Is partial coupling requiring multisubstrate reactions (like  $R_4$  in the previous example)?

#### Partial coupling and multisubstrate reactions

Does partial coupling require multisubstrate reactions (like  $R_4$  in the previous example)?

### Consider the following:



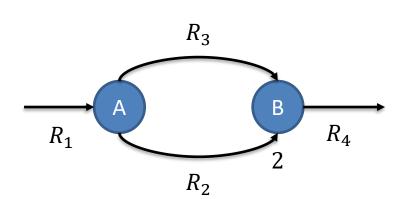
$$\frac{dc_A}{dt} = v_1 - v_2 - v_3$$

$$\frac{dc_B}{dt} = v_3 + 2v_2 - v_4$$

#### Partial coupling and multisubstrate reactions

Does partial coupling require multisubstrate reactions (like  $R_4$  in the previous example)?

There is a subtle network difference which we will tackle in the next lectures that distinguishes the two cases.



$$\frac{dc_A}{dt} = v_1 - v_2 - v_3$$

$$\frac{dc_B}{dt} = v_3 + 2v_2 - v_4$$

$$v_1 \neq 0$$
 if and only if  $v_4 \neq 0$ 

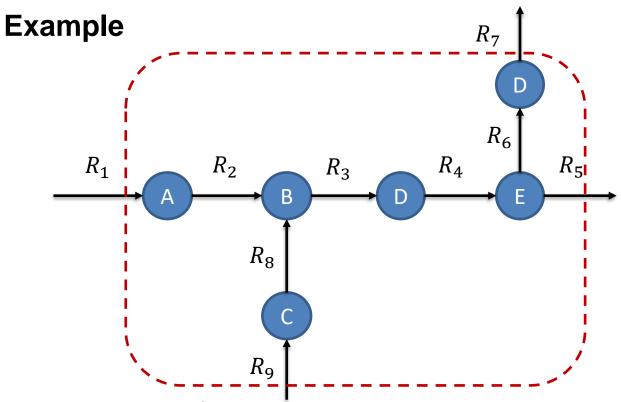
#### **Definition**

A reaction  $R_i$  is fully coupled to reaction  $R_j$  if non-zero flux for  $R_i$  implies not only a non-zero but also a fixed flux for  $R_j$  and vice versa.

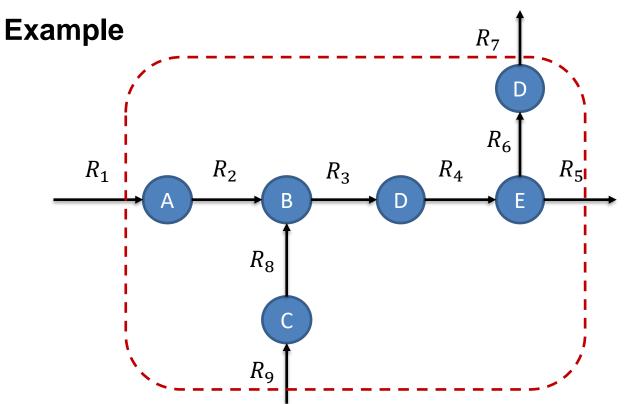
#### Mathematical

A reaction  $R_i$  is directionally coupled to reaction  $R_j$  if for every v with Nv = 0,  $v_i \neq 0$  implies  $v_j \neq 0$ , with  $\frac{v_i}{v_j} = const$  and vice versa.

**Notation**  $R_i \Leftrightarrow R_j$  or  $i \Leftrightarrow j$ 



Reaction  $R_3$  is fully coupled to  $R_4$ Denoted by  $R_3 \Leftrightarrow R_4$ 

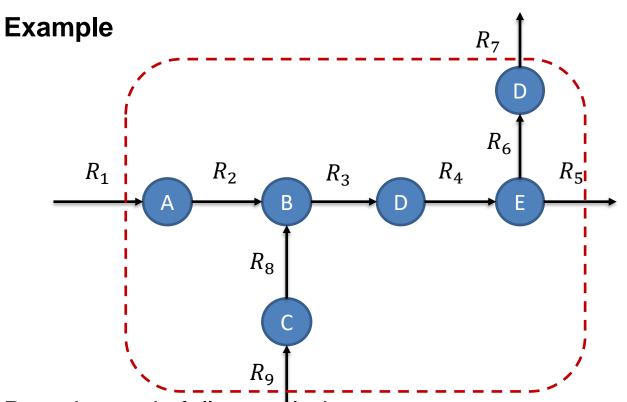


$$\frac{dc_D}{dt} = v_3 - v_4$$

$$v_3 - v_4 = 0$$

Reaction  $R_3$  is fully coupled to  $R_4$ Denoted by  $R_3 \Leftrightarrow R_4$ 

$$\frac{v_3}{v_4} = 1$$
 (fixed constant)



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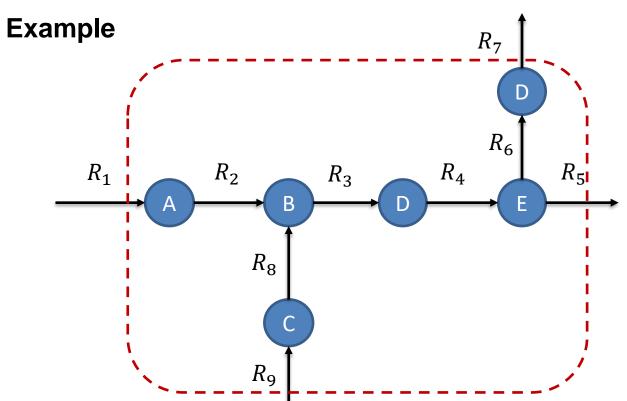
Reaction  $R_3$  is fully coupled to  $R_4$ 

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Other fully coupled pairs?

Hence

$$\frac{v_3}{v_4} = 1$$
 (fixed constant)



 $\frac{ac_D}{dt} = v_3 - v_4$ 

 $v_3 - v_4 = 0$ 

Reaction  $R_3$  is fully coupled to  $R_4$ 

Denoted by  $R_3 \Leftrightarrow R_4$ 

Other fully coupled pairs?

Hence

 $\frac{v_3}{v_4} = 1$  (fixed constant)

Main goal
Classify reaction pairs based on their couplings.

Three types of reaction couplings

Directional coupling vi!= 0 => vj!=0

**Partial coupling** 

vi != 0 => vj !=0, with vi/vj != const

Full coupling vi/vj = vij, s.t. for every v, Nv=0

Reaction pairs which are not directionally, partially, or fully coupled will be called **uncoupled**.

#### **FCA** questions

#### **Questions**

1. Is there a more efficient way to determine reactions whose fluxes show correlation of 1? Sor 12 보기 위해 샘플링을 하는것말고 방법이 없는가?

do we have to sample or one LP is sufficient?

- 2. Can we specify how the structure of the metabolic network leads to dependencies between reaction fluxes?
- 3. What are the biological implications of such dependencies?

These questions can be answered by

Flux Coupling Analysis (FCA)

vi != 0 => vj !=0 vi != 0 => vj !=0, with vi/vj != const vi/vj = vij(const), s.t. for every v, Nv=0

Consider the following program

$$\max_{v} (\min) \frac{v_i}{v_i}$$
 ratio  $\exists t$ .

s.t.

$$Nv=0$$
 steady state  $\forall i, 1 \leq i \leq m, v_i^{min} \leq v_i \leq v_i^{max}$  capacity constraints

The objective is **not linear** function but a **fraction of linear** functions

this is referred to as linear fractional program

이미 min이 0이라는것을 알고, max는 fluxrange()로 구할 수 있다 네가티브는 없다.

What are the possible outcomes for  $\max_{v} (\min) \frac{v_i}{v_j}$  assuming that all reactions in the network are irreversible?

**Note**: reversible reactions can be divided into two irreversible reactions (forward and backward direction)

Clearly  $\frac{v_i}{v_j}$  lies in the interval  $[0, \infty)!$ 

There are five possibilities

#### There are five possibilities

all reactions are necessaarily irreversable

- 1.  $0 \le \frac{v_i}{v_j} \le c$  for some constant c > 0
- 2.  $c_1 \le \frac{v_i}{v_j} \le c_2$  for some constant  $c_1 > 0$  and  $c_2 > 0$ ,  $c_1 \ne c_2$
- 3.  $\frac{v_i}{v_j} = c$  for some constant c > 0
- 4.  $c \leq \frac{v_j}{v_i}$  for some constant c > 0, without an upper bound
- 5.  $0 \le \frac{v_i}{v_j}$  without an upper bound

#### There are five possibilities

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- 3.  $\frac{v_i}{v_j} = c$  for some constant c > 0 full coupling  $i \Leftrightarrow j$
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#### There are five possibilities

vi가 0이 아니라는것을 의미

1번의 경우 ub를 가짐

- 1.  $0 \le \frac{v_i}{v_j} \le c$  for some constant c > 0 directional coupling  $i \to j$
- 2.  $c_1 \le \frac{v_i}{v_j} \le c_2$  for some constant  $c_1 > 0$  and  $c_2 > 0$ ,  $c_1 \ne c_2$
- 3.  $\frac{v_i}{v_j} = c$  for some constant c > 0 full coupling  $i \iff j$
- 4.  $c \leq \frac{v_j}{v_i}$  for some constant c > 0, without an upper bound directional coupling  $i \to i$
- 5.  $0 \le \frac{v_i}{v_i}$  without an upper bound

#### There are five possibilities

- 1.  $0 \le \frac{v_i}{v_j} \le c$  for some constant c > 0 directional coupling  $i \to j$
- $2. \ c_1 \leq \frac{v_i}{v_j} \leq c_2$  for some constant  $c_1 > 0$  and  $c_2 > 0$ ,  $c_1 \neq c_2$  partial coupling  $i \leftrightarrow j$
- 3.  $\frac{v_i}{v_j} = c$  for some constant c > 0 full coupling  $i \iff j$
- 4.  $c \le \frac{v_j}{v_i}$  for some constant c > 0, without an upper bound directional coupling  $j \to i$
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- 3.  $\frac{v_i}{v_j} = c$  for some constant c > 0 full coupling  $i \Leftrightarrow j$
- 4.  $c \le \frac{v_j}{v_i}$  for some constant c > 0, without an upper bound directional coupling  $j \to i$
- 5.  $0 \le \frac{v_i}{v_i}$  without an upper bound **uncoupled**

Solving the linear fractional program (LFP)

$$\max_{v} (\min) \frac{v_i}{v_j}$$

s.t.

$$Nv=0$$
 steady state  $\forall i, 1 \leq i \leq m, 0 \leq v_i \leq v_i^{max}$  capacity constraints

provides the means to either classify each pair of reactions into the introduced coupling types or to state that it is uncoupled

However, we do not know how to solve LFPs

#### **Solving Linear Fractional Program**

To solve the linear fractional program (LFP), let us multiply numerator, denominator and all constraints by some positive t

특정 값 t를 곱해서, t에대한 스케일링을 한다

$$\max_{vt} (\min) rac{v_i t}{v_j t}$$
위아래에 t를 곱해도 objective function은 변하지 않는다. 즉, 정답이 변하지는 않음!

s.t.

t를 곱함으로써 스케일된 flux distribution또한 steady state일것이 분명함!

$$N(vt) = 0$$
 steady state

$$\forall i, 1 \leq i \leq m, 0 \leq v_i t \leq v_i^{max} t$$
 capacity constraints

 $\begin{array}{c} \text{vit} \succeq \text{flux distribution} \\ t \geq 0 \end{array}$ 

+ v' 가 스케일된 distribution vt라고 할때, vj'은?

vmint <= vt <= vmaxt

+ vj' = vjt = vj \* 1/vj = 1

+ 즉, vit/vjt = vit = vi'

If  $v_j > 0$ , then we can take  $t = 1/v_j$ 

t = 1/vj라고 가정해보자. 왜냐면 t = 0보다 큰 어떤 값이든 될 수 있으니까!

#### Solving Linear Fractional Program

To solve the linear fractional program (LFP), let us multiply numerator, denominator and all constraints by some positive t

$$\max_{vt} (\min) \frac{v_i t}{v_i t}$$

fluxrange가 min/max Vi'를 구한다면. 

s.t.

$$N(vt) = 0 \text{ steady state}$$
 
$$\forall i, 1 \leq i \leq m, 0 \leq v_i t \leq v_i^{max} t \text{ capacity constraints}$$
 
$$t \geq 0$$

For the vector v' = vt it then holds that  $v'_i = 1$ , so the LFP can be transformed into

#### **Solving Linear Fractional Program**

To solve the linear fractional program (LFP), let us multiply numerator, denominator and all constraints by some positive t

$$\max_{v'}$$
 (min)  $v'_i$ 

s.t.

$$Nv'=0$$
 steady state  $\forall i, 1 \leq i \neq j \leq m, 0 \leq v'_i \leq v_i^{max} t$  capacity constraints  $v'_j = 1_{t=1/V_j}$   $t>0$ 

#### This is a linear program which we encountered in FVA

The transformation is known as Charnes-Cooper transformation!

To solve the linear fractional program (LFP), let us multiply numerator, denominator and all constraints by some positive t

$$\max_{v'}$$
 (min)  $v'_i$ 

s.t.

$$Nv'=0$$
 steady state  $orall i, 1 \leq i \neq j \leq m, 0 \leq {v'}_i \leq v_i^{max} t$  capacity constraints  $v'_j=1$   $t>0$ 

Conclusion: For one reaction, we need to solve 2(m-1) linear programs to determine the couplings to the other (m-1) reactions

For all reactions – we need to solve 2m(m-1) LPs

모든 커플링을 조사하려면 굉장히 많은 수의 계산이 필요하다.. 너무 낭비임.

### The Charnes-Cooper transformation

$$\max_{x} \frac{c^{T}x + \alpha}{d^{T}x + \beta}$$

예를 들어서, cT = [ 1,1,0,0,0] dT = [0,0,1,1,0] alpha = 0 beta = 0

s.t.

$$Ax \leq b$$

vi/vj = i(), s.t. for every v, Nv=0

Assumption:

The feasible region is such that  $d^Tx + \beta > 0$ 

### The Charnes-Cooper transformation

두개로 나눌 수 있다

$$\max_{x} \frac{c^{T}x + \alpha}{d^{T}x + \beta} = \max_{x} \frac{c^{T}}{d^{T}x + \beta}x + \frac{\alpha}{d^{T}x + \beta}$$

s.t.

dTx + beta는 Vj이고, t는 1/Vj라고 가정해보자.

$$Ax \leq b$$

Introduce the substitution

$$y = \frac{1}{d^T x + \beta} x$$
 and  $t = \frac{1}{d^T x + \beta}$ 

dTx + beta = dTy + beta\*t = 1..? 이건 좀더 공부해보자

이 둘을 이렇게 정의해보고 써보자.

### The Charnes-Cooper transformation

cT가 이라면? 그냥 0이다.

$$\max_{x} c^{T}y + \alpha t$$

s.t.

$$Ay \le bt$$

$$d^T y + \beta t = 1$$

$$t \ge 0$$

Introduce the following

$$y = \frac{1}{d^T x + \beta} x$$
 and  $t = \frac{1}{d^T x + \beta}$ 

For all reactions – we need to solve m(m-1) LPs (expensive!)

For instance, calculation of coupling in the large-scale network of E. coli (iJR904) would take more than 24 hours

Recent methods and tools are based on assessing feasibility programs, rather than linear programs মান্টা বুবা, মান্টান.

This allows calculations in a matter of few hours for the same network

# **FCA** questions

#### **Questions**

1. Is there a more efficient way to determine reactions whose fluxes show correlation of 1?

do we have to sample or one LP is sufficient?

- 2. Can we specify how the structure of the metabolic network leads to dependencies between reaction fluxes?
- 3. What are the biological implications of such dependencies?

These questions can be answered by

Flux Coupling Analysis (FCA)

### Couplings in real-world networks

All coupling types can be found in metabolic network reconstructions across all kingdoms of life

Reaction couplings can be identified in well-studied pathways

**Figure 6** Coupled reaction set identified for purine biosynthesis in *E. coli* on a glucose-minimal medium, assuming a constant biomass composition. The numbers indicate the relative values or range of values for each flux in any particular flux distribution for given growth condition. Secondary metabolites and cofactors are omitted for simplicity.

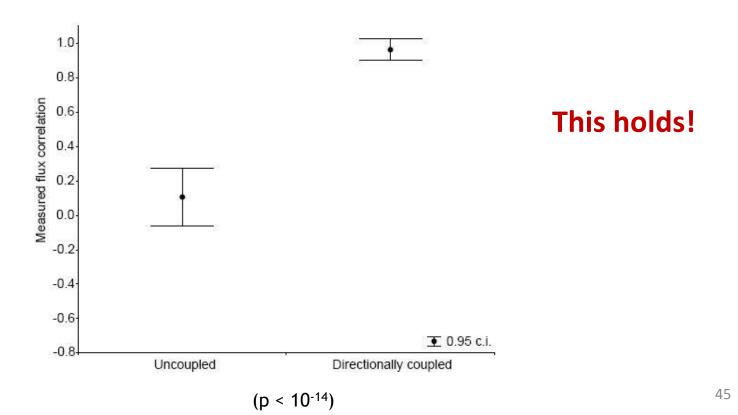
Burgard et al. (2004) Genome Research

# **Couplings and flux data**

Do directionally coupled reactions exhibit higher correlation of their fluxes in comparison to uncoupled reactions based on measured flux distributions?

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# Couplings and gene regulation

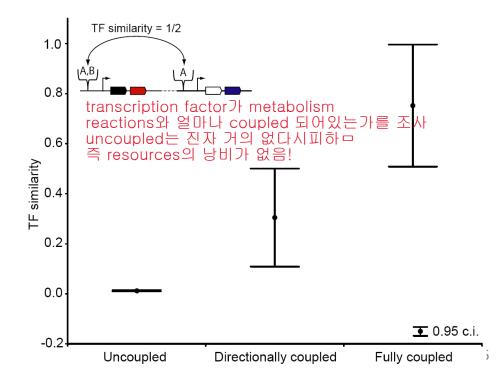
# Does flux coupling relate to transcriptional coregulation of genes?

For fully coupled fluxes, this would imply that transcriptional regulation is organized in a way that no resources are

wasted!

Notebaart et al. (2007)

PloS Comp. Biol.



# **Couplings and evolution**

Some genes are lost during evolution (for instance, if they have a deleterious effect on fitness)

Pal et al. (2005) *Nature Genetics* found that genes associated to reactions with coupled fluxes are more likely lost together over evolution!

Coupling type	Event	#Events	OR* (95% c.i.)
Fully coupled	Transfer	59	64.6 (24.2- 168.8)
Fully coupled	Loss	1,624	50.0 (41.8-59.6)
Directionally coupled	Transfer	78	60.3 (24.3- 147.2)
Directionally coupled	Loss	2,833	9.6 (8.3-11.1)

Odds-ratio

### Recap

```
Definitions of three coupling types
       directional
       partial
       full
Computational ways to determine reaction couplings
       linear fractional program (LFP)
                                                Type text here
       Charnes-Cooper transformation to LP
Implications of reactions couplings
       flux correlations
       co-regulation of fully coupled reactions
       gene gain or loss in evolution
```