



Challenges in the Optimization of Bio-Systems

Gerhard-Wilhelm Weber

Institute of Applied Mathematics, Middle East Technical University, Ankara, Turkey

<http://www.iam.metu.edu.tr/research/groups/compbio/index.html>

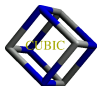


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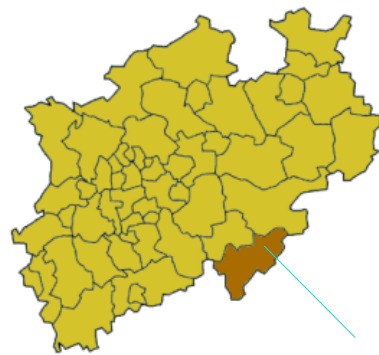


Content

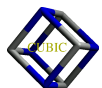
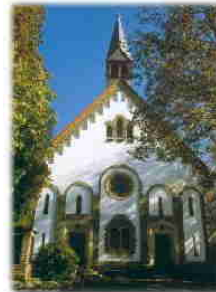
- Motivation: Bio-Systems
- Computational Biology
- Modeling and Prediction of Gene Expression Patterns
- Population Dynamics, Gene Dynamics and Variation
- Optimization
- Computational Metabolism, Medicine and Tomography
- Earth Warming and Sustainable Development
- Conclusion and Perspectives



Bio-Systems



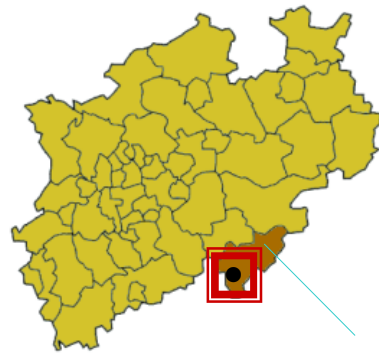
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Bio-Systems



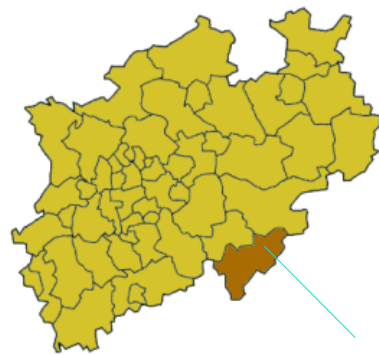
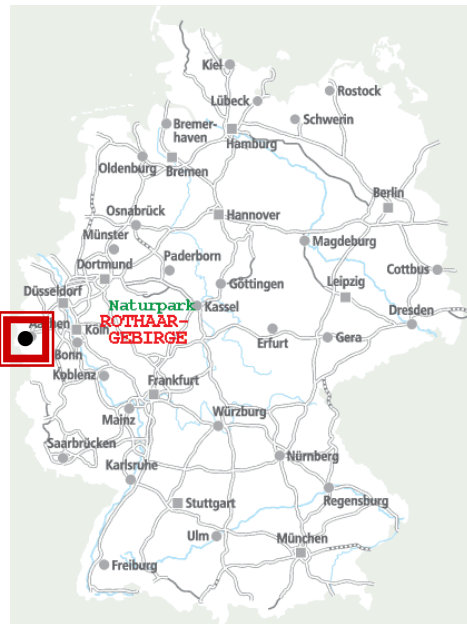
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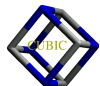
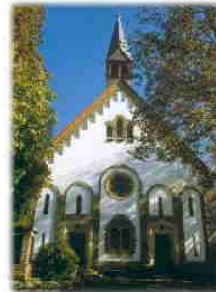
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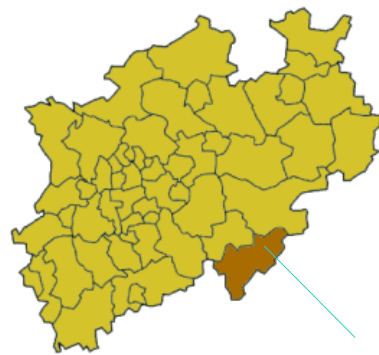
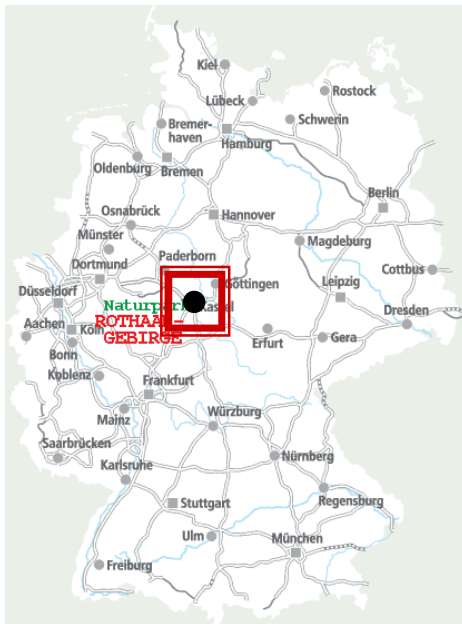
Bio-Systems



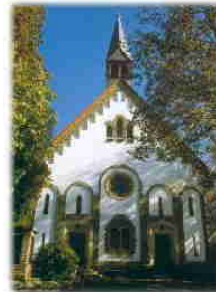
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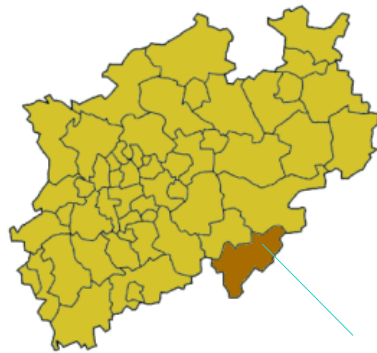
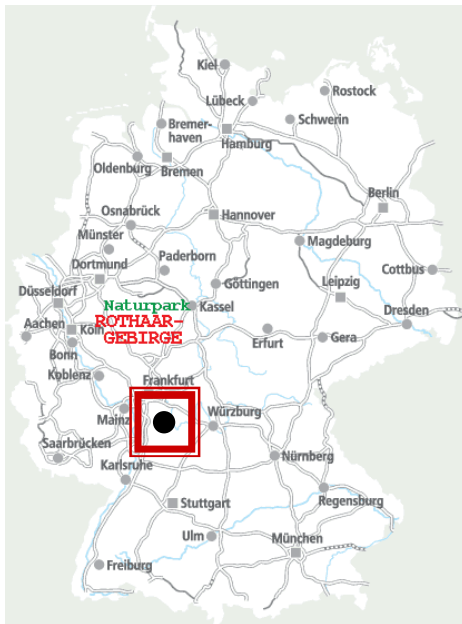
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Wemlighausen



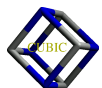
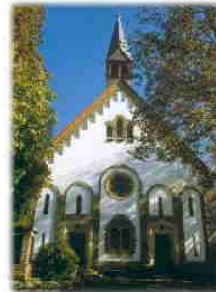
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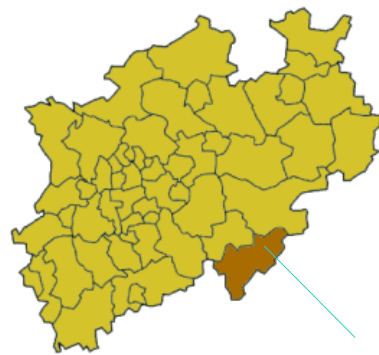
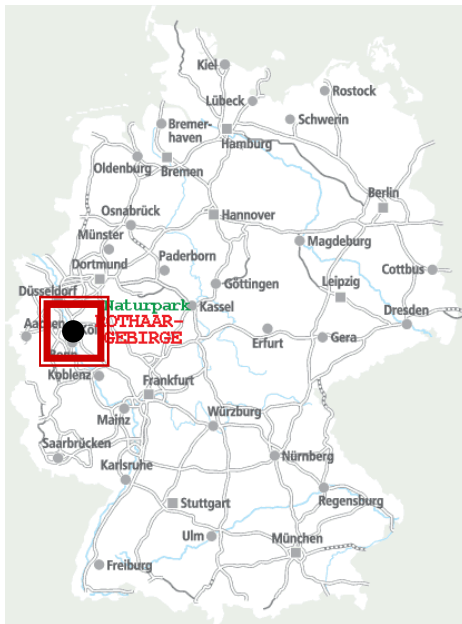
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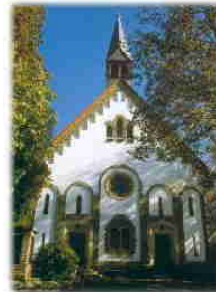
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Bio-Systems



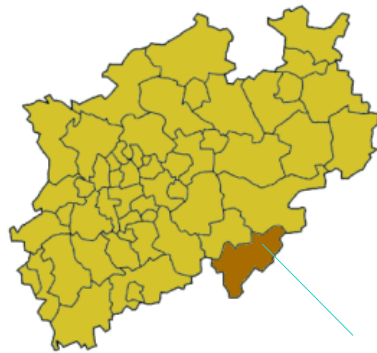
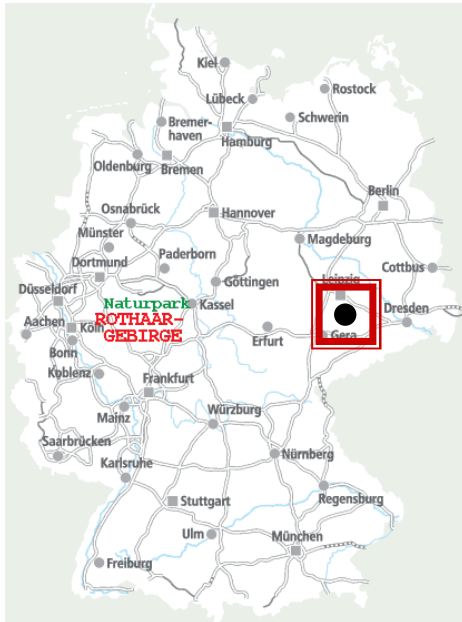
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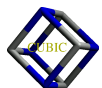
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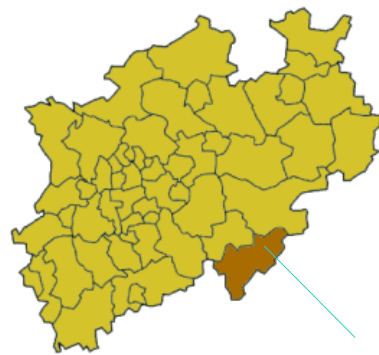
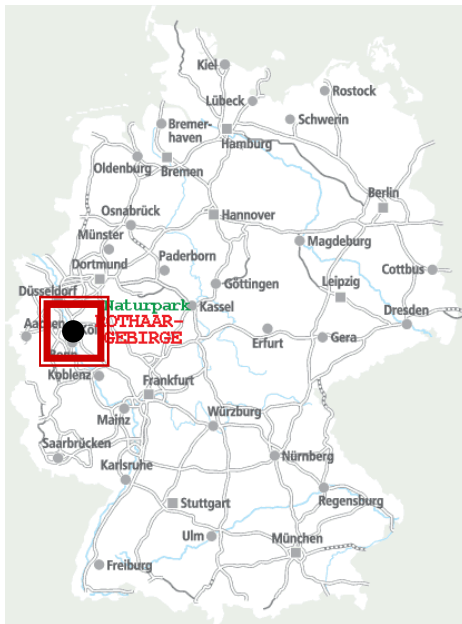
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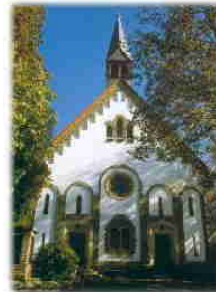
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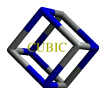
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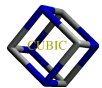
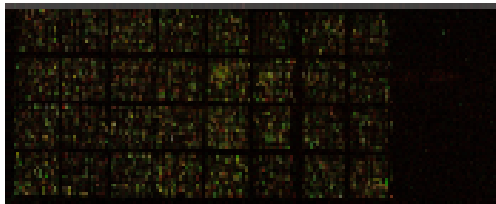
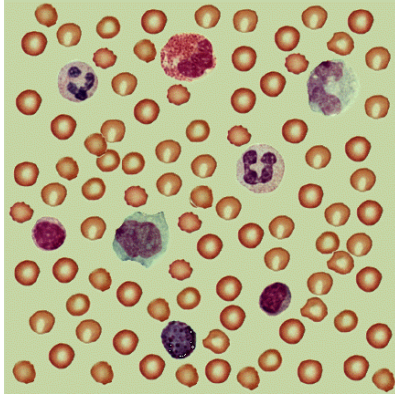
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Bio-Systems



Bio-Systems

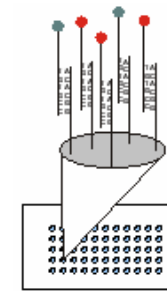
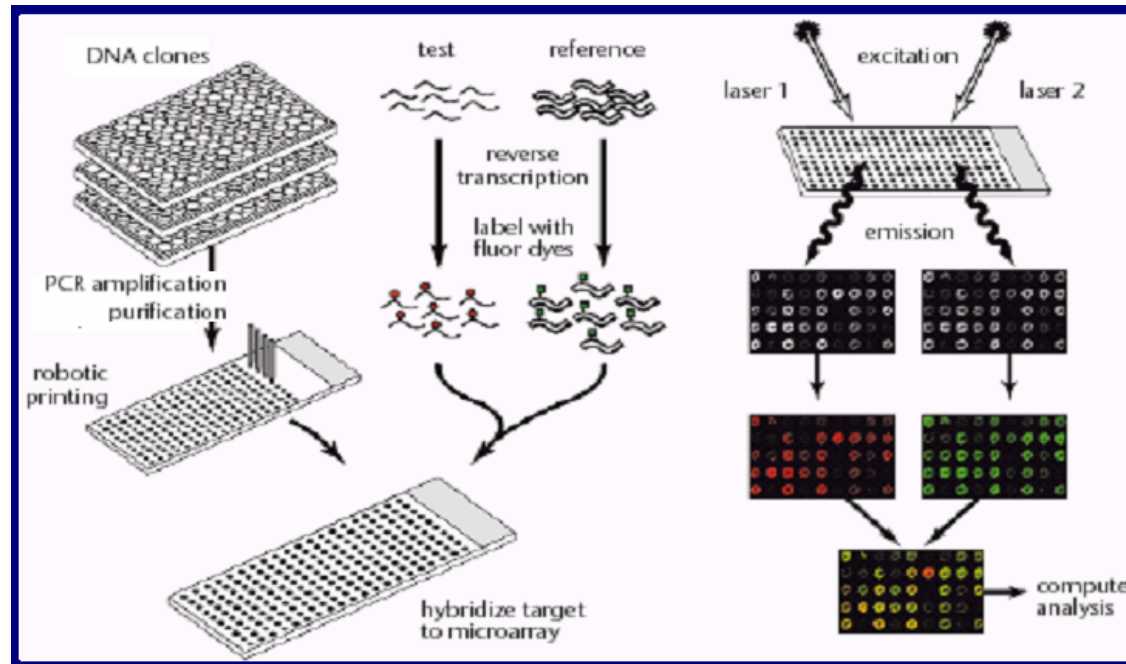


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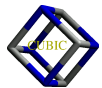


anticipation of gene patterns based on

DNA microarray chip experiments



M.U. Akhmet, H. Öktem
S.W. Pickl, E. Quek Ming Poh
T. Ergenç, B. Karasözen
J. Gebert, N. Radde, W.
Ö. U•ur, R. Wünschiers
M. Ta•tan, F.B. Y•maz



Gene Patterns Modeling & Prediction

least squares – ML
statistical learning

time-contin.

$$\dot{E} = M(E) E$$

Expression data

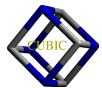
$$E(0) = E_0$$

time-discr.

Ex.: Euler, Runge-Kutta

$$E_{k+1} = M_k E_k$$

$$M_k = (em_{ij}) \in M$$



Gene Patterns Model. & Pred.

For which parameters, i.e., for which set M (or: dynamics),
is *stability* guaranteed ?

development of process
its feasibility
goodness-of-fit (model) test

Def.: M is *stable* : $\Leftrightarrow$ $\exists B : (\text{compl.}) \text{ bounded neighbourhood of } O_n,$
 $\forall m \in \bullet\bullet, M_0, M_1, \dots, M_{m-1} \in M :$
 $(M_{m-1} M_{m-2} \dots M_0) B \subseteq B .$



Analysis with Polytopes

Theorem (Brayton, Tong 1979) :

Given a set $M := \{M_0, \dots, M_{m-1}\}$ of m distinct complex matrices.

Then,

$$M \text{ is stable} \iff B^* = \bigcup_{k=0}^{\infty} B_k \text{ is bounded.}$$

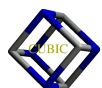
Here, B_0 is a bounded neighbourhood of 0_n ,

and for $k > 0$

$$B_k := H \left(\bigcup_{i=0}^{\infty} M_{k'}^i B_{k-1} \right),$$

where

$$k' := (k-1) \bmod m, \quad H : \text{convex hull.}$$



Extremal Points

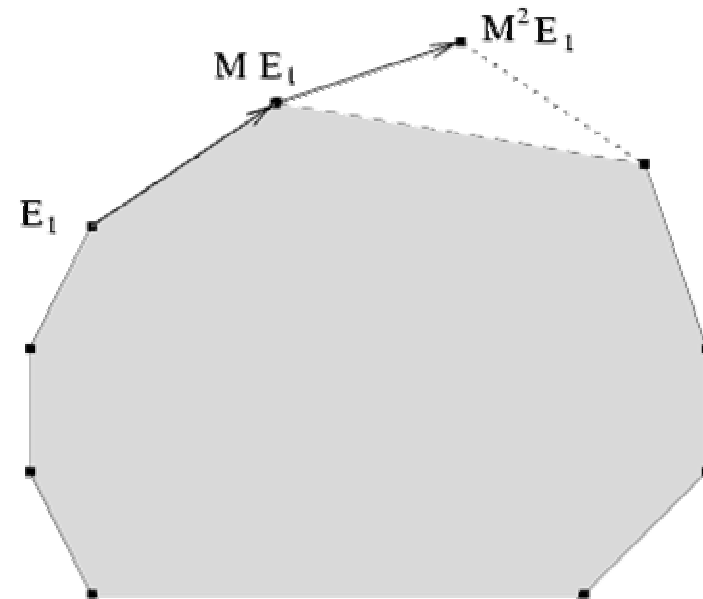
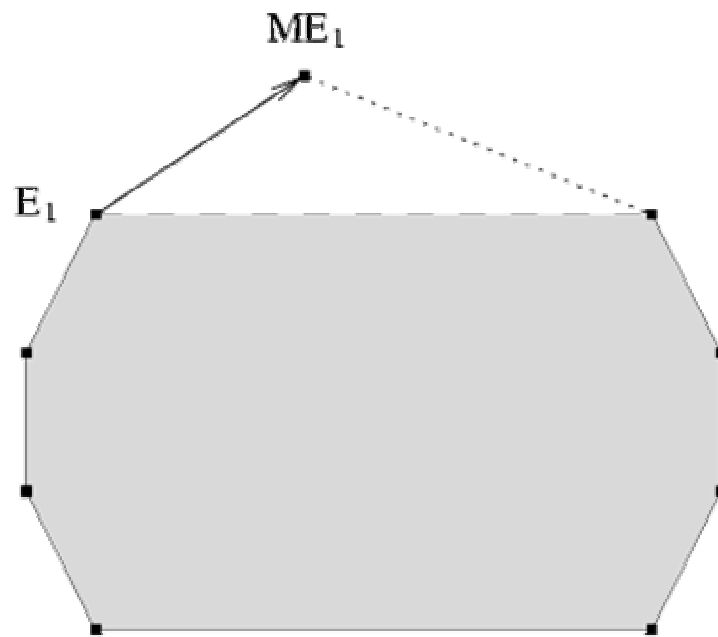
The “discrete” power of the algorithm is based on using **polyhedra** B_k and focussing on the **extremal points** of the sets B_k .

Theorem 1 : If Z is an extremal point of B_k , then there exist $j \in \mathbb{N}_0$ and an extremal point u of B_{k-1} , such that

$$Z = M_k^j u .$$



Construction Principle



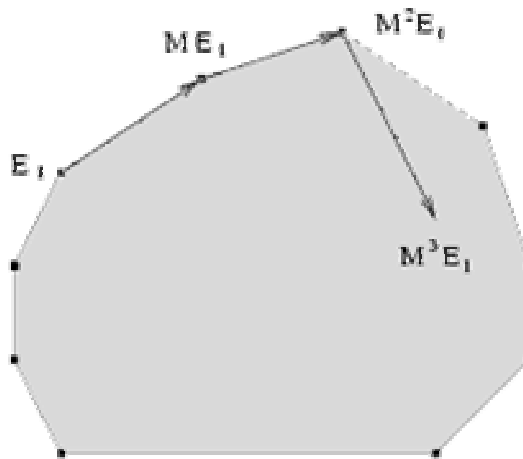
Stopping Criterion

Theorem 2 :

Let $z_i = M_{k'}^j u_i$ (as above, $i \in \{1, 2, \dots, r\}$).

Then,

$$H \{z_1, \dots, z_r\} = B_k \iff M_{k'} z_i \in H \{z_1, \dots, z_r\} \quad \forall \quad i=1, 2, \dots, r.$$



Construction Principle

- $B_k = B_{k_0} \quad (k \geq k_0) \Rightarrow \left\{ \begin{array}{l} \text{"stopping" (at step } k_0), \\ B^* = \bigcup_{i=0}^{\infty} B_i \text{ bounded,} \\ \text{stability of matrices / dynamics} \end{array} \right.$
- $\partial B_0 \cap \partial B_k = \emptyset \Rightarrow \left\{ \begin{array}{l} B^* \text{ unbounded} \\ \text{instability} \end{array} \right.$

Gebert, Laetsch, Pickl, W., Wünschiers 2005

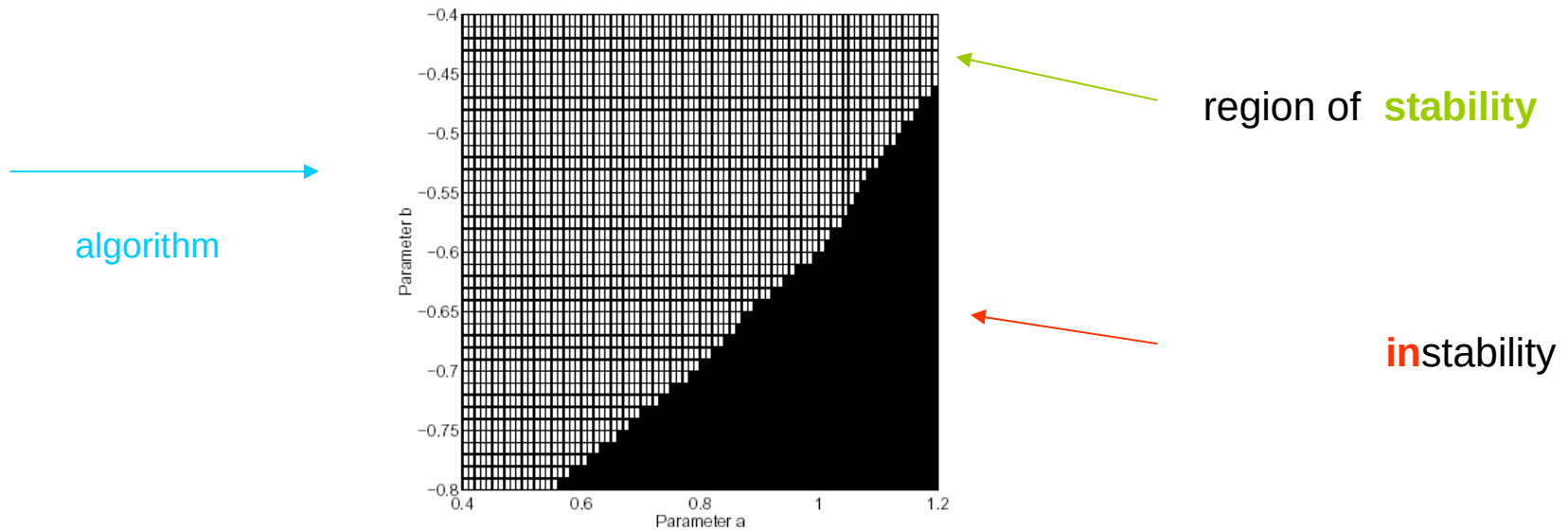
Ergenç, W. 2004



Numerical Example

Ex. (Pickl 1999) :

$$M = \{M_0, M_1\} \quad M_0 = \begin{pmatrix} a & 1 \\ b & 0 \end{pmatrix}, \quad M_1 = \begin{pmatrix} 0 & 1 \\ b & 0 \end{pmatrix}$$

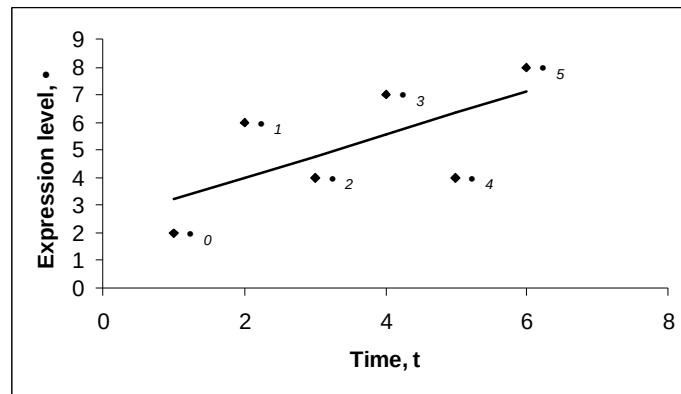


Gene Network

Ex. :

$$\dot{E} = M E \quad h_k = 1$$

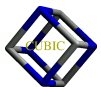
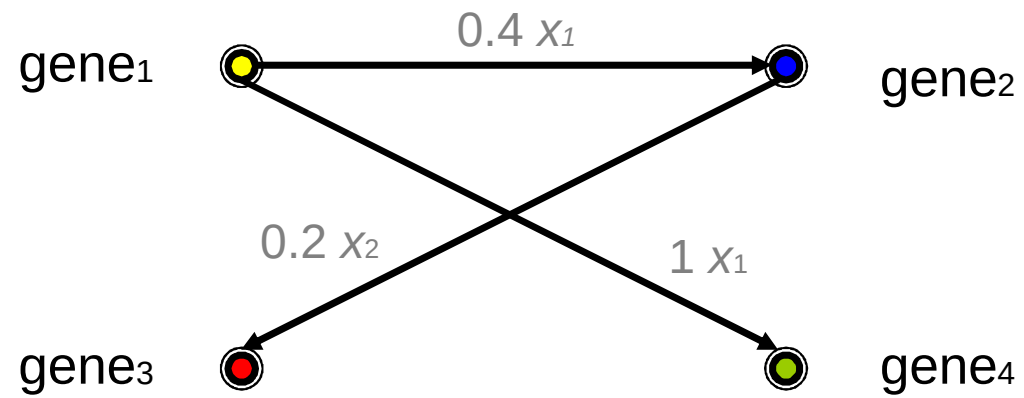
$$A = \begin{pmatrix} E_1(t_0) & E_2(t_0) & E_3(t_0) & E_4(t_0) \\ E_1(t_1) & E_2(t_1) & E_3(t_1) & E_4(t_1) \\ E_1(t_2) & E_2(t_2) & E_3(t_2) & E_4(t_2) \\ E_1(t_3) & E_2(t_3) & E_3(t_3) & E_4(t_3) \end{pmatrix} = \begin{pmatrix} 255 & 250 & 0 & 255 \\ 255 & 200 & 50 & 0 \\ 255 & 180 & 70 & 255 \\ 255 & 170 & 80 & 0 \end{pmatrix}$$



$$M = \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0.4 & -0.61 & 0 & 0 \\ 0 & 0.2 & -0.39 & 0 \\ 1 & 0 & 0 & -2 \end{pmatrix}$$



Gene Network



$$\dot{E} = F(E)$$

nonlinearities

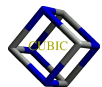
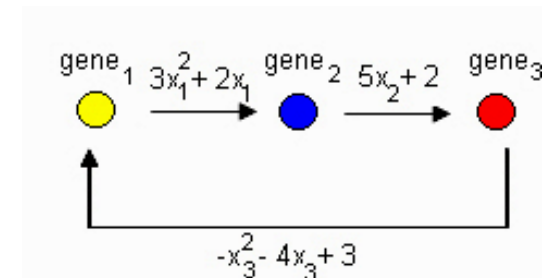
Ex.:

$$F_j(E) = \sum_{i=1}^n f_{j,i}(E_i) = \sum_{i=1}^n (a_{j,i}(E_i)^2 + b_{j,i}(E_i) + c_{j,i}).$$

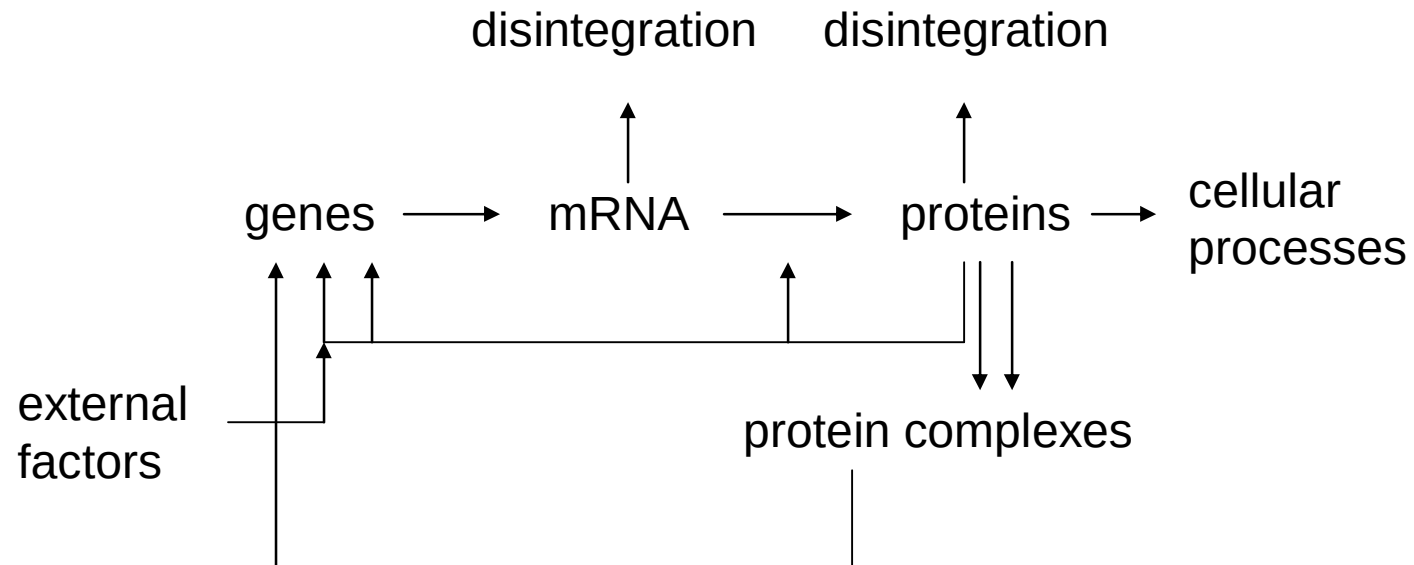
$\bar{E}_k$: exper. data

$\dot{E}_k$: approx. increase/decrease

$$\min_{\substack{c_{j,1}, b_{j,1}, a_{j,1}, \\ \dots \\ c_{j,n}, b_{j,n}, a_{j,n}}} \sum_{m=0}^{t-1} || F(\bar{E}_m) - \dot{E}_m ||^2$$



Inference of Gene Regulatory Networks



Gene transcription also regulates the transcription factors defining a **dynamic regulatory network** involving highly *nonlinear* **feedback mechanisms**.



Proposed Model Class

piecewise linear:

$$E(k+1) = M_{s(k)}E(k) + b_{s(k)}$$

$$s(k) = F_B Q(E(k-1))$$

$$Q_i(E(k)) = \begin{cases} 1 & \text{if } E_i(k) > \Omega_i \\ 0 & \text{else} \end{cases}$$

Akhmet, Gebert, Pickl, Öktem, W. 2004

Gebert, Radde, W. 2004

$$\dot{E}(t) = M_{s(t)}E(t) + b_{s(t)}$$

$$s(t) = F(Q(E(t), E(t-q_1), \dots, E(t-q_p)))$$

$$Q_i(E^1, E^2, \dots, E^{p+1}) \in \{0, 1, \dots, q_i\}$$



Dynamics and Variation

- population **dynamics**

M.U. Akhmet , V. Tkachenko
H. Öktem, S.W. Pickl, W., et al.



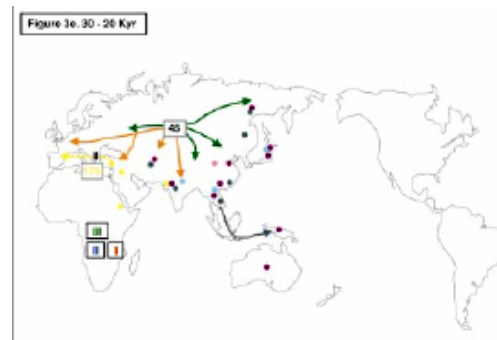
with **delay** and **impulse**

$$\dot{E}(t) = f(t, E(t), E([t + t]))$$

anticipation

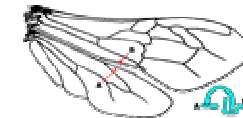
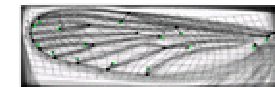
- gene **dyn.**

I. Togan
C. Berkman
et al.



& **variation**

A. Kence
et al.



←
time

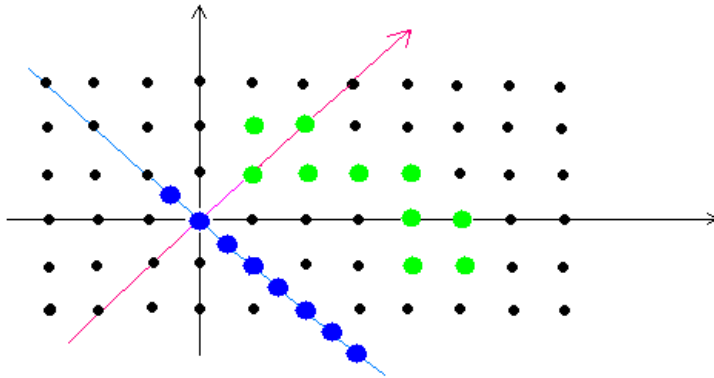


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... Tomography

Discrete ...



Ö. Ya•ar, O. Özgür, W.

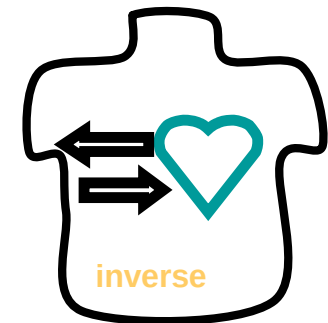
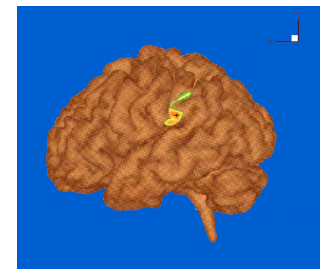
C. Diner, A. Do•an
F. Özbudak, A. Tiefenbach

M. Eyübo•lu, N. Gençer
Y. Serina•o•lu, A. Kurt
S. Sar•kaya, W.

VLSI chip design

medicial imaging

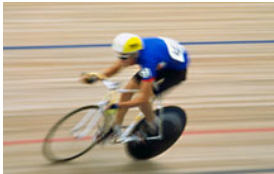
Brain / Heart ...



inverse problems

Metabolic Engineering

sports (and general) medicine



object oriented

modules

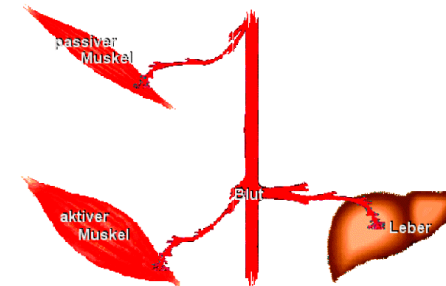
DAE

ODE

training program

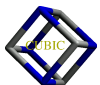
math. modeling

parameter estimation



A. Schulte Thomas 2004

S. Özöür 2005



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Generalized Semi-Infinite Opt.

$$\mathcal{P}_{\mathcal{GSI}}(f, h, g, u, v) \quad \begin{cases} \text{Minimize } f(x) \text{ on } M_{\mathcal{GSI}}[h, g], \text{ where} \\ M_{\mathcal{GSI}}[h, g] := \{ x \in \mathbb{R}^n \mid h_i(x) = 0 \ (i \in I), \\ g(x, y) \geq 0 \ (y \in Y(x)) \}. \end{cases}$$

C^2

$$Y(x) = M_{\mathcal{F}}[u(x, \cdot), v(x, \cdot)]$$

$$:= \{ y \in \mathbb{R}^q \mid u_k(x, y) = 0 \ (k \in K), \ v_\ell(x, y) \geq 0 \ (\ell \in L) \}$$

I, K, L finite

W. 2003



Reverse Chebychev Approx.

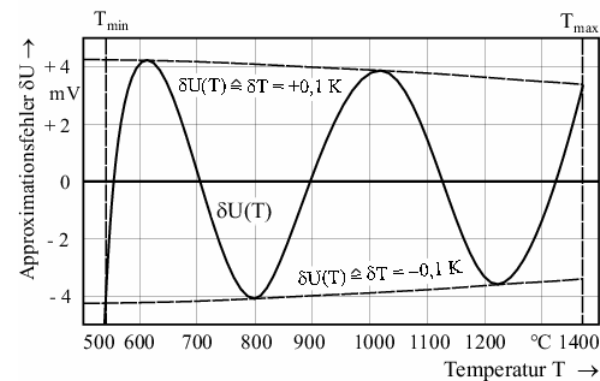
Ex.: approx. of a **thermo-couple characteristic**

Hoffmann, Reinhard

thermo-couple $f(y)$: *spline* of polynomials with deg. 3 – 13, on $[a,b]$

to be approx. by $\Psi(x_o, y) := \sum_{k=0}^{n^o} x_{k+1} y^k$:

- **bounds** on *error* $\delta(x_o, y) := \Psi(x_o, y) - f(y)$



Bernhard

$\mathcal{P}_{GSI}^1$

- some **interpol.**

(= y) →



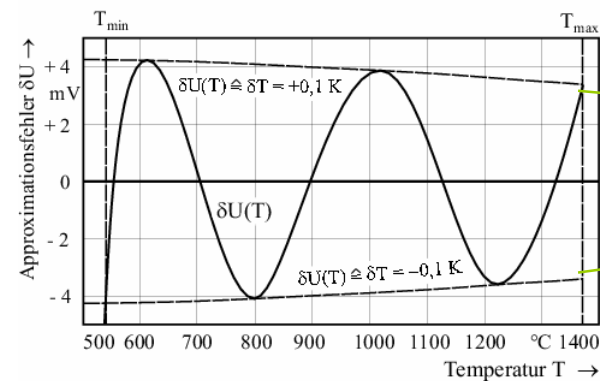
Reverse Chebyshev Approx.

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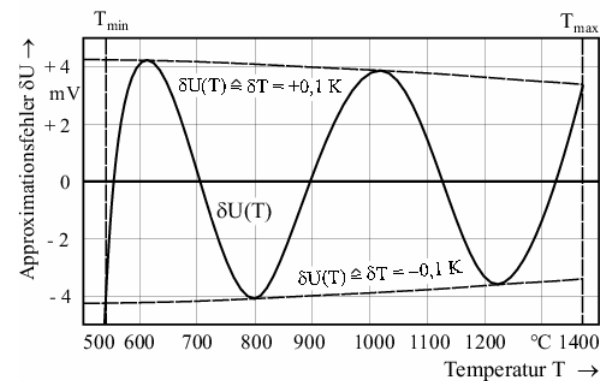
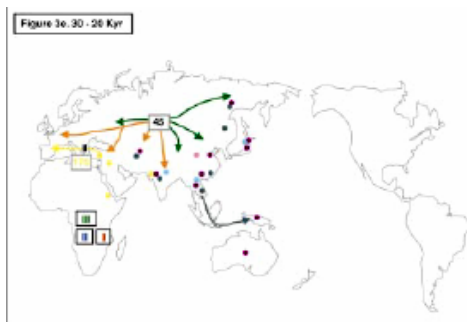
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← time

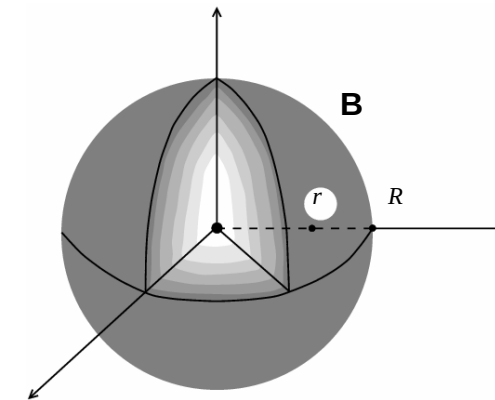
- some **interpol.**

(= y) →



Time-Optimal Control

Ex.: time-minimal cooling (or heating) of



$$\mathcal{P}^{\text{tm}} \left\{ \begin{array}{l} \text{Min } l(T, u) := T \text{ such that} \\ \\ \text{there is a bounded function } \theta : [0, R] \times [0, \infty) \rightarrow \mathbb{R}, \text{ where} \\ \theta|_{(0, R] \times (0, \infty)} \text{ is partially differentiable,} \\ u = \theta(R, \cdot)|_{[0, T]} \text{ is continuous, and} \\ \\ \theta_t(r, t) = a \Delta \theta(r, t) = \frac{a}{r^2} \frac{\partial}{\partial r} (r^2 \theta_r(r, t)) \quad ((r, t) \in (0, R] \times (0, \infty)), \\ \\ \theta(r, 0) = \theta_0 \quad (r \in [0, R]), \\ \\ \theta(R, T) = \theta_E, \\ \\ T \geq 0, \\ \\ |\sigma_u(R, t)| \leq \sigma^* \quad (t \in [0, T]). \end{array} \right.$$

$$\mathcal{P}_T^{\text{nm}} \left\{ \begin{array}{l} \text{Min}_{u_T} \|\sigma_{u_T}^{\theta_0}(R, \cdot)\|_{\infty, T}, \\ \\ \text{where } u_T \in C([0, T], \mathbb{R}) \text{ fulfills} \\ \\ u_T(T) = \theta_E. \end{array} \right.$$

$$\forall T > 0 \quad \exists! \hat{u}_T$$

$$\mathcal{P}_{\text{GSI}}^2$$



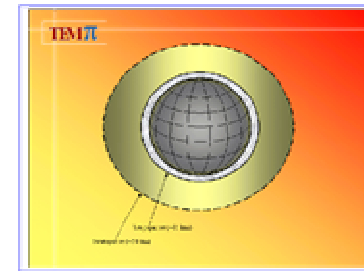
Anticipation

Ex.'s.:

- **thermo-regulation**
- control of **earth warming**

Pickl, W.

$q(r, x, t)$
local-global



- *maximization* of time-horizon *longest* term approximation / description

anticipation



TEM Model

$$E_i(t+1) = E_i(t) + \sum_{j=1}^n em_{ij} M_j(t)$$

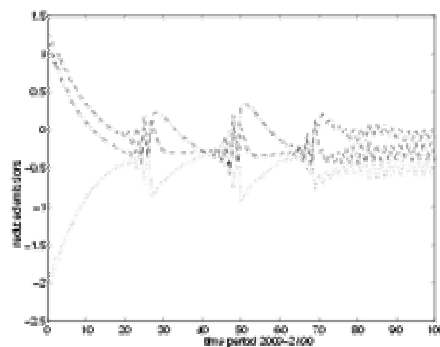
S.W. Pickl

$$M_i(t+1) = M_i(t) - \lambda_i M_i(t) [M_i^* - M_i(t)] * [E_i(t) + \varphi_i \Delta E_i(t)] + u_i(t)$$

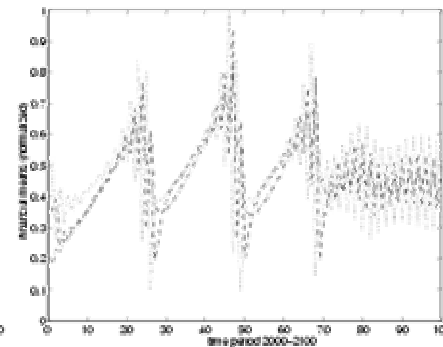
Z. Alparslan

$u_i(t)$ control parameter

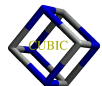
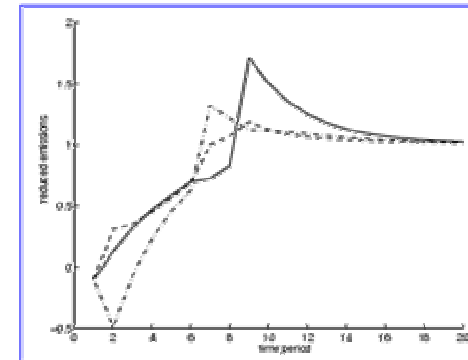
B. K., W.



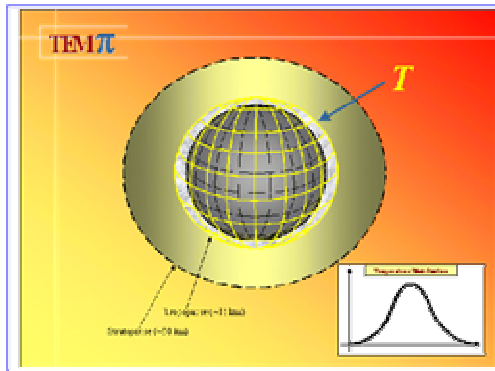
emissions reduced



financial means



JET

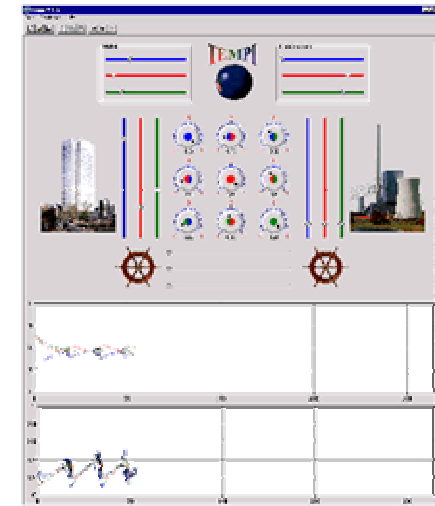


EU project

application

“Joint International Emissions Trading,
Analysis,
Forecast,
and Optimal Strategies
in
**Sustainable
Low Carbon Energy Management**”

common learning



U. Leopold-W., B. K., S.W. P., M. Türkay, W.

Risk Management

H. Körezlioğlu et al.

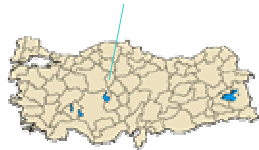


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Sustainable Living

Balaban Valley



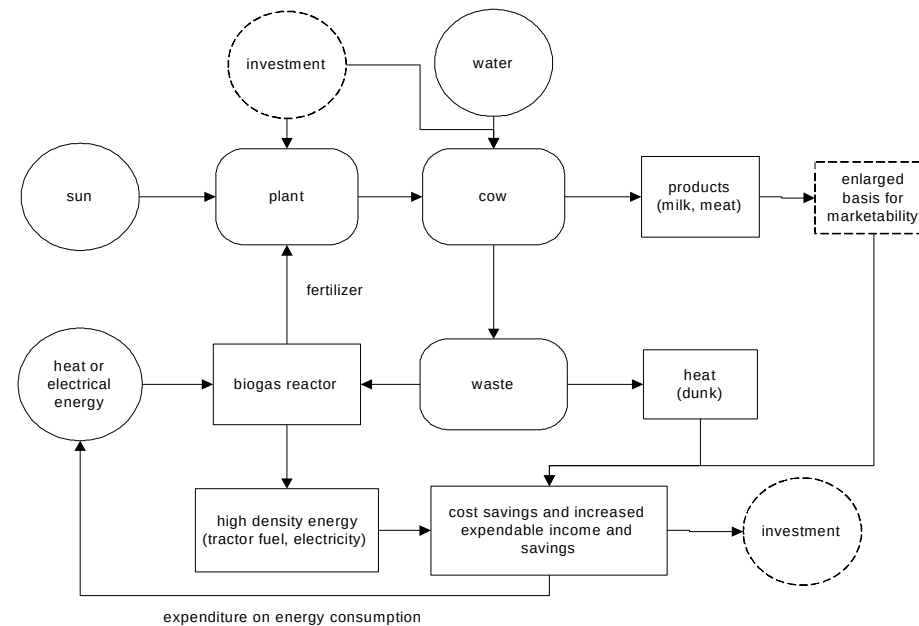
improvement of living conditions

learning

modeling

optimization

data mining



D. DeTombe, A., I., H. Gökmen

S. Kayal, M., Y. Ecevit

T. Bali, W. et al.

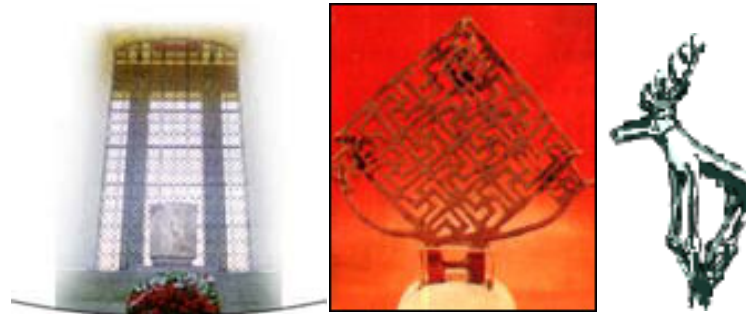


Turkey

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Conclusions and Perspectives



Operations Research Society
of Israel

Magyar Operációkutatási Társaság

Operations Research Society
of Hungary



Association of European
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Operations Research
Society of Turkey

EUROPT

EURO Working Group
on Continuous Optimization

GOR

Operations Research Society
of Germany



IAM, METU Ankara
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Conclusions and Perspectives



5th EUROPT Workshop "Advances in Continuous Optimization"

Reykjavik, Iceland
June 29 – July 1, 2006

EURO Summer Institute

"Optimization Challenges in Engineering: Methods, Software, and Applications"

Wittenberg, Germany
August 18 – September 2, 2006



EURO XXI in Iceland July 2-5, 2006

21st European Conference on Operational Research



OR for Better Management of Sustainable Development



Earth Institute

Young People for OR in Developing Countries



Balaban Valley Project



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Sincere invitation

to

■■■
EUROPT

EURO Working Group

in Continuous Optimization

<http://www.iam.metu.edu.tr/EUROPT/>

two new EURO WGs planned:
Computational Biology & Bioinformatics
OR for Development

and **Computational Biology and Medicine Group**

<http://www.iam.metu.edu.tr/research/groups/compbio/index.html>

gweber@metu.edu.tr



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