Constrained Evolutionary Search for Model Parameters. Case Studies in Thymocyte Dynamics

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Constrained Evolutionary Search

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Outline



- 2 Constrained evolutionary search of the parameter space
- 3 Case study: modelling a perturbed thymocyte dynamics

Exploring the output of the evolutionary search

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A parameterized model in pharmacology

Allosteric two-state model of receptor activation



- L receptor isomerization constant
- *K* (*M*) equilibrium dissociation constant for *A* (*B*)
- γ binding cooperativity for A and B
- δ activation cooperativity A and B
- α , β intrinsic efficacies

- [A] ([B])-orthosteric (allosteric) ligand concentration
- $[R]_a$ ($[R]_t$)- activated (total) receptor concentration



A parameterized model in pharmacology

Allosteric two-state model of receptor activation



What is known?

 Experimental values for [A], [B], [R]_a/[R]_t

What is required?

K, L, M, α, β, γ, δ

Typical approach:

minimize the mean squared error (MSE)

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A parameterized model in pharmacology

Allosteric two-state model of receptor activation



What is known?

 Experimental values for [A], [B], [R]_a/[R]_t

What is required?

K, L, M, α, β, γ, δ

Typical approach:

minimize the mean squared error (MSE)

Particularities:

- explicit relationship between MSE and the parameters
- not easy to establish initial approximations for the parameters
- possible several equally good sets of parameters optimization

= [D.Roche et al, 2013] ~

Compartmental models of the thymus

- the thymus is an import organ of the immune system
- its main role is to produce various types of thymocytes (T cells):
 - double negative T cells (N population)
 - double positive T cells (P population)
 - single positive T cells (M4 and M8 populations)
- there are several processes involving the T cell populations:
 - cell proliferation (growth)
 - cell death (involution)
 - cell maturation and differentiation (transfer)



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Compartmental models of the thymus



Parameters:

- proliferation rates: r_n, r_p, r_4, r_8
- death rates: d_n, d_p, d_4, d_8
- transfer rates: $s_n, s_4, s_8, s_{o4}, s_{o8}$
- bone marrow inflow rate: b
- carrying capacities: *K*, *K*_n Example: Mehr's model [Mehr et al, 1996]

$$\begin{cases} \dot{N}(t) &= r_n N(t)(1 - N(t)/K_n) - d_n N(t) - s_n N(t) + b(1 - N(t)/K_n) \\ \dot{P}(t) &= r_p P(t)(1 - Z(t)/K) - d_p P(t) - (s_4 + s_8)P + s_n N(t) \\ \dot{M}_4(t) &= r_4 M_4(t)(1 - Z(t)/K) - d_4 M_4(t) - s_{04} M_4(t) + s_4 P(t) \\ \dot{M}_8(t) &= r_8 M_8(t)(1 - Z(t)/K) - d_8 M_8(t) - s_{08} M_8(t) + s_8 P(t) \\ Z(t) &= N(t) + P(t) + M_4(t) + M_8(t) \end{cases}$$

Compartmental models of the thymus



What is required?

- proliferation/ death/ transfer/ inflow rates
- carrying capacities (K_n, K)





Experimental estimates of the number of cells during their evolution (N-red, P-blue, M_4 -magenta, M_8 -green)

Compartmental models of the thymus

Particularities:

- search for parameters minimizing the mean squared error betwen the numerically estimated solutions of the ODE and the experimental data
- "semi-transparent" model: no explicit analytical relationship between *MSE* and the parameters to be estimated
- constraints:
 - $K_n < K$ (relationship between the carrying capacities)
 - desired evolution of the populations of cells (e.g. steady-state or involution)

[PNII-ID-PCE: REVISAL - Modeling and simulation of the dynamics of thymocyte populations and cells of the thymus medulla under normal and pathological situations, 2012-2014]

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Some problems requiring parameter estimation

A parameterized software module Image registration





 Registration: Find a transformation which maps pixels of one image to corresponding pixels of the other image

[D. Gil et al., 2013]

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A parameterized software module

Image registration

Particularities:

- What is available: a software module implementing a registration algorithm and maximal error threshold
- What is required: parameters of the registration algorithm such that the error is below the threshold and the execution time of the algorithm is minimized
- "opaque" model: almost nothing is known about the algorithm inside the software module (it receives the parameters and provides the error and the running time)
- constraints on parameters (e.g $p_1 < p_2$) and on the error ($err < \epsilon$)
- the values provided for the execution time can be uncertain

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Summary: types of parameterized models

"Transparent" models

- the search criterion depends in an explicit way on the parameters
- gradient information could be available

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"Semi-transparent" models

- the search criterion depends only in an implicit way on the parameters
- its evaluation requires numerical solutions of some equations

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Summary: types of parameterized models

"Transparent" models

- the search criterion depends in an explicit way on the parameters
- gradient information could be available

"Semi-transparent" models

- the search criterion depends only in an implicit way on the parameters
- its evaluation requires numerical solutions of some equations

"Opaque" models

- the evaluation of the search criterion is based on a "black box" module
- nothing is known about the influence of the parameters on the search criterion

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Outline



2 Constrained evolutionary search of the parameter space

3 Case study: modelling a perturbed thymocyte dynamics

Exploring the output of the evolutionary search

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Why using an evolutionary search?

- it requires minimal knowledge on the model
- if properly designed it ensures a good exploration of the parameter space
- it can be used for:
 - multimodal/ multiobjective/ constrained optimization

and can provide several results of similar quality



Population-based stochastic search

- Exploration: mutation and crossover (reproduction)
- Exploitation: selection

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Which search method to use?

Many options:

- Evolution Strategies (ES)
- Covariance Matrix Adaptation ES (CMA-ES)
- Particle Swarm Optimization (PSO)
- Differential Evolution (DE)
- Ant Systems (AS)
- Harmony Search (HS)
- Artificial Bees Colonies (ABC)
- other nature inspired metaheuristics (e.g. firefly, cuckoo, bacterial foraging, gravitational search)

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Selection criteria:

- appropriateness ability to deal with the problem characteristics
- simplicity easy to understand/ implement
- competitiveness good behavior for test functions
- availability easy to find implementations

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Common types of constraints

From simple to complex constraints

• Bounding box constraints: $x \in [a, b]$

Repairing rules:

- iterate the reproduction operator until the offspring satisfies the constraint
- use a symmetry based rule, i.e. when $x \notin [a, b]$ iterate:

$$\mathbf{x}' = \left\{egin{array}{cc} b - (\mathbf{x} - b) & ext{if } \mathbf{x} > b \ a + (a - \mathbf{x}) & ext{if } \mathbf{x} < a \end{array}
ight.$$

until $x' \in [a, b]$.

select randomly an element in the search range

Common types of constraints

From simple to complex constraints

Constraints on the set of feasible values: $x \in [a, b] \cap \mathbb{Z}$

- use specific operators to generate new elements
- use operators for search in continuous spaces + apply a rounding function

Simple inequality constraints: x < y

- instead of searching for x and y
- search for x and $\delta > 0$ (such that $y = x + \delta$)

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Common types of constraints

From simple to complex constraints

General form of a constrained optimization problem

Find x which minimizes f(x) subject to

•
$$g_i(x) \leq 0, \quad i=1,\ldots,m$$

• $h_i(x) = 0$, j = 1, ..., p (usually transformed in $|h_i(x)| < \epsilon$)

Types of constraints

- g_i and h_i depend explicitly on x easy to check if the constraint is satisfied
- there is no explicit dependence between q_i , h_i and x (e.g. the constraint could just say that the estimated model behaves in a given way) - only a degree of constraint satisfaction can be estimated

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Dealing with constraints in evolutionary search Penalty method

Main idea:

combine the objective function with a penalty function measuring the degree of violating the constraints

$$F(x) = f(x) + \sum_{i=1}^{m} r_i \max(0, g_i(x)) + \sum_{j=1}^{p} c_j |h_j(x)|$$

Advantage:

easy to implement

Disadvantage:

the choice of the penalty weights is not obvious

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Dealing with constraints in evolutionary search Feasibility rules

Main idea (Deb's feasibility rule):

use separate objective value (*f*) and penalty value (degree of constraint violation - ϕ) when compare two elements; *x* is better than *x*' if:

- x and x' are both feasible and f(x) < f(x')
- x is feasible and x' is not feasible
- x and x' are both unfeasible and $\phi(x) < \phi(x')$

Advantages:

- easy to implement and to couple with various search algorithms
- it does not require parameters

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Dealing with constraints in evolutionary search Feasibility rules

Main idea (Deb's feasibility rule):

use separate objective value (*f*) and penalty value = degree of constraint violation (ϕ) when compare two elements; *x* is better than *x'* if:

- x and x' are both feasible and f(x) < f(x')
- x is feasible and x' is not feasible
- *x* and *x'* are both unfeasible and $\phi(x) < \phi(x')$

Disadvantage:

 separating the constraints and the objective function can lead to diversity loss (because they strongly favour the feasible solutions)

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Dealing with constraints in evolutionary search Stochastic ranking

Main idea:

decides randomly which selection criterion to use (objective or penalty function)

x is better than x' if

$$\left\{ \begin{array}{l} ((\phi(x) = \phi(x') = 0) \text{ or } (\operatorname{rand}(0, 1) < P_f)) \text{ and } (f(x) < f(x'))) \\ \phi(x) < \phi(x') \end{array} \right.$$

Advantages:

it limits the diversity loss (by accepting promising but unfeasible candidates)

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Dealing with constraints in evolutionary search Stochastic ranking

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decides randomly which selection criterion to use (objective or penalty function)

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Disadvantages:

• it requires the specification of a parameter (P_f , e.g. $P_f = 0.45$)

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Constrained Evolutionary Search

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Context

- the dynamics of thymocytes is very sensitive to any pathological situation associated with an increase of glucocorticoids (e.g. diabetes,obesity, infections)
- by administrating a particular glucocorticoid (dexamethasone -DXM) one can induce a transient involution of thymus activity similar with that common in pathological situations
- it would be useful to model this transient perturbation and to extract information on the impact DXM has on various mechanisms (cell proliferation, death and differentiation)

Experimental data [dr. F. Mic, UMF Timisoara]

 more than 70 experiments on young and adult mice (before and after a treatment with DXM)



What is known ?

- DXM induces a significant depletion of thymocytes, especially of DP cells
- after treatment the thymus rebounds and the pre-treatment level is almost reached after 14 days

What would be useful to know?

- can the perturbation induced by DXM be modelled through transiently perturbed rates/ mechanisms?
- which of the mechanisms (proliferation, death, differentiation) is most affected?
- when does the perturbation on each mechanism reaches the maximal value?

Constrained Evolutionary Search

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Possible approach

- start from a model describing the normal dynamics (e.g. a multi-compartmental model)
- introduce transient perturbations into the model (e.g perturbed rates, transient inhibition of proliferation)



Perturbed rates

- Main idea: additive perturbation of rates
- Perturbing functions family

$$\xi(C;t) = rac{c_1}{t^{c_3}+c_2} - rac{c_1c_4/c_2}{t^{c_5}+c_4}, c_i > 0$$

• Perturbed rates: $r + \xi(C; t)$



Parameters estimation

• Optimization problem: find the parameters which

- minimize the mean squared error (MSE)
- satisfy constraints concerning the positivity of all perturbed rates and the vanishing of the perturbation

$$MSE(x) = \frac{1}{4n} \sum_{\pi \in \{N, P, M_4, M_8\}} \left(\frac{1}{\max_{j = \overline{1, n}} \{\overline{\pi}_j^2\}} \sum_{j = 1}^n (\pi(x; t_j) - \overline{\pi}_j)^2 \right)$$

- n = number of experimental values
- $\bar{\pi}$ = experimental values corresponding to each of the four populations
- $\pi(x; t)$ = numerically estimated solutions

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• For each perturbed rate, two constraints should be satisfied

$$r + \xi(C; t) \ge 0$$
 for all $t \in [t_a, t_f]$

 $|\xi(\mathbf{C}; \mathbf{t}_f)| < \epsilon_f$

- t_a = time moment when the perturbation starts (treatment administration moment)
- t_f = time moment when the perturbation should be small enough (the effect of treatment is passed)
- ϵ_f = small enough value (negligible perturbation)

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Main difficulty in dealing with the constraints

- the constraints on positivity of rates cannot always be checked exactly
- in some cases, sufficient conditions for positivity can be found but usually they are not also necessary:

$$r \ge \max\{c_1/(c_2^2+c_2), c_1c_4/(c_2^2+c_2c_4+c_2)\}$$

- need to define a degree of constraint satisfaction
- $S_{p}^{i}(C_{j}) = 1$ if the sufficient condition is satisfied, otherwise

$$S_{p}^{j}(C_{j}) = \frac{\operatorname{card}\{t \in T_{h} | r_{j} + \xi(C_{j}; t) > 0\} - \delta}{\operatorname{card}(T_{h})}$$

 $T_h = \{t_a, t_a + h, \dots, t_f\}, \quad h > 0 - discretization step, \quad \delta > 0$

Positivity constraint:

$$\mathcal{S}^{j}_{\mathcal{P}}(\mathcal{C}_{j}) = rac{ ext{card}\{t \in \mathcal{T}_{h} | r_{j} + \xi(\mathcal{C}_{j}; t) > 0\} - \delta}{ ext{card}(\mathcal{T}_{h})}$$

Perturbation vanishing constraint:

$$S_{\nu}^{j}(C_{j}) = \begin{cases} 1 & \text{if } |\xi(C_{j}; t_{f})| \le \epsilon_{f} \\ 1 - \min\{1, |\xi(C_{j}; t_{f})|\} & \text{otherwise} \end{cases}$$

Combined constraints satisfaction degree:

$$\mathcal{S}(\mathcal{C}) = \prod_{j=1}^q \mathcal{S}^j_{
ho}(\mathcal{C}_j) \mathcal{S}^j_{
ho}(\mathcal{C}_j) \in [0,1]$$

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Constraints handling: different ways of combining the satisfaction degree with MSE

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Various comparison rules

x is considered better than x' if one condition is satisfied

Rule A (θ - threshold for the satisfaction degree, e.g. θ = 0.99):

•
$$S(x) \ge \theta$$
 and $S(x') < \theta$;

- $S(x) \ge \theta$ and $S(x') \ge \theta$ and MSE(x) < MSE(x');
- $S(x) < \theta$ and $S(x') < \theta$ and $S(x) \ge S(x')$

(similar to Deb's rule)

Rule B:

•
$$S(x) \ge \theta$$
 and $S(x') < \theta$

- S(x)S(x') = 0 and $MSE(x) \le MSE(x')$
- $S(x) \neq 0$, $S(x') \neq 0$ and $MSE(x)/S(x) \leq MSE(x')/S(x')$

Various comparison rules

x is considered better than x' if one condition is satisfied

Rule C:

- S(x) > 0 and S(x') = 0
- S(x) = 0, S(x') = 0 and $MSE(x) \le MSE(x')$
- $S(x) \neq 0$, $S(x') \neq 0$ and $MSE(x)/S(x) \leq MSE(x')/S(x')$.

Various comparison rules

x is considered better than x' if one condition is satisfied

Rule D (S interpreted as a probability that the constraint is satisfied):

- $U_1 \le S(x)$ and $U_2 > S(x')$;
- $U_1 > S(x)$ and $U_2 > S(x')$ and $MSE(x) \le MSE(x')$;
- $U_1 \leq S(x)$ and $U_2 \leq S(x')$ and $MSE(x)/S(x) \leq MSE(x')/S(x')$

 $(U_1 \text{ and } U_2 \text{ are random values uniformly distributed on } [0, 1])$

Rule E (inspired by stochastic ranking):

- $S(x) \ge \theta$, $S(x') \ge \theta$ and MSE(x) < MSE(x')
- $U < P_f$, $S(x)S(x') \neq 0$, MSE(x)/S(x) < MSE(x')/S(x')
- $S(x) \geq S(x')$

Evolutionary search

Initialization: $x_i = U(a_i, b_i), \quad i = \overline{1, m}$ while $\langle \text{ NOT termination } \rangle$ do • Mutation:

$$y_i = x_{r_1} + F \cdot (x_{r_2} - x_{r_3}), \quad i = \overline{1, m}$$

• Crossover:

$$z_i^j = \left\{ egin{array}{ll} y_i^j & ext{if } rand(0,1) < CR ext{ or } j = j_{0^*} \ x_i^j & ext{otherwise} \end{array}
ight.$$

$$i = \overline{1, m}, j = \overline{1, n}$$

• Selection:

$$egin{aligned} & x_i(g\!+\!1) = \left\{ egin{aligned} & z_i & ext{if } f(z_i) \leq f(x_i(g)) \ & x_i^j & ext{if } f(z_i) > f(x_i(g)) \end{aligned}
ight. \end{aligned}$$

Differential Evolution [Storn&Price,



m - population size $F \in (0, 2)$ - scale factor $CR \in [0, 1]$ - crossover rate j_0 - randomly selected component

Evolutionary search

JADE algorithm [Zhang&Sanderson, 2009]:

self-adaptive version of Differential Evolution

$$z'_{i} = \begin{cases} x'_{i} + F_{i} \cdot (x'_{rbest} - x'_{i}) + F_{i} \cdot (x'_{r1} - x'_{r2}) & \text{if } rand() \le CR_{i} \\ x'_{i} & \text{otherwise} \end{cases}$$

JADE particularities

- x_{rbest} chosen from the p% best elements in the population
- *x*_{r2} chosen from an archive consisting of elements discarded by selection
- F_i generated using a Gaussian distribution
- *CR_i* generated using a Cauchy distribution
- the parameters of these distributions are adjusted during the evolutionary process

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Numerical experiments

- Experimental dataset:
 - 232 values (number of cells in each of the four thymocyte populations)
 - collected from young and adult mice thymus either before or after a treatment administration
- number of parameters: k = 71
- number of constraints: q = 26
- number of independent runs: 30

JADE parameters:

- population size: 20
- generations: 5000
- percent of best elements: p = 10%

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Comparison between ranking rules

Quality of fit (*MSE*), constraints satisfaction degree (*S*), estimated feasibility probability (*FP*(θ) for $\theta = 0.99$).

Rule	MSE	S	$FP(\theta)$
A ($\theta = 1$)	0.0338 ± 0.0012	1 ± 0	1
A ($ heta=0.99$)	0.0270 ± 0.0010	0.9966 ± 0.0033	1
$B~(\theta=0.99)$	0.0268 ± 0.0014	$0.9999 \pm 5 \cdot 10^{-6}$	1
С	0.0261 ± 0.0009	0.9878 ± 0.0119	0.45
D	0.0290 ± 0.0017	$0.9999 \pm 3 \cdot 10^{-6}$	1
$E(P_f = 0.45)$	0.0250 ± 0.0005	0.9935 ± 0.0011	1
Unconstrained	0.0208 ± 0.0022	0.0468 ± 0.0776	0

Remark: Rule E is better than the other ones (Mann-Whitney statistical test, p-value< 10^{-5})

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Simulated dynamics



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Simulated dynamics



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Perturbed rates



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Perturbed rates



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[Thomas-Vaslin et al., 2008]

- the proliferation process consists of several stages
- differentiation arises at each stage but with different rates
- used to model the impact of ganciclovir on thymus dynamics (when continuously administrated for 7 days)
- perturbation modelled by triggering off the proliferation process

$$\begin{split} \dot{N}_0(t) &= \sigma_N - (r_N + d_N) N_0(t) \\ \dot{N}_i(t) &= 2\gamma(t) r_N N_{i-1}(t) - (r_N + d_N + \mu_N(i)) N_i(t), \quad i = \overline{1, n_N} \end{split}$$

$$\begin{aligned} \dot{P}_{0}(t) &= \sum_{i=1}^{n_{N}} \mu_{N}(i) N_{i}(t) + 2\gamma(t) r_{N} N_{n_{N}}(t) - (r_{P} + d_{P}) P_{0}(t) \\ \dot{P}_{i}(t) &= 2\gamma(t) r_{P} P_{i-1}(t) - (r_{P} + d_{P} + \mu_{P}(i)) P_{i}(t), \quad i = \overline{1, n_{P} - 1} \\ \dot{P}_{n_{P}}(t) &= \sum_{i=1}^{n_{P} - 1} \mu_{P}(i) P_{i}(t) + 2\gamma(t) r_{P} P_{n_{P} - 1}(t) - \mu_{LP} P_{n_{P}}(t) \end{aligned}$$

Parameters to be estimated

- Proliferation rates: r_N , r_P , r_4 and r_8
- Death rates: d_N , d_P , d_4 and d_8

• Transfer rates:
$$\mu_N(i) = (\alpha_N \cdot i)^n$$
, $\mu_P(i) = (\alpha_P \cdot i)^n$, μ_{LP} ,
 $e_4(i) = (\alpha_{e4} \cdot i)^n$ and $e_8(i) = (\alpha_{e8} \cdot i)^n$

Number of stages: n_N, n_P

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$$\begin{split} M_{80}(t) &= \alpha_8 \mu_{LP} P_{n_P}(t) - (r_8 + d_8) M_{80}(t) \\ \dot{M}_{8i}(t) &= 2\gamma(t) r_8 M_{8,i-1}(t) - (r_8 + d_8 + e(i)) M_{8i}(t), \qquad i = \overline{1, n_8 - 1} \\ \dot{M}_{8n_8}(t) &= 2\gamma(t) r_8 M_{8,n_8 - 1}(t) - (d_8 + e_8(n_8)) M_{8n_8}(t) \end{split}$$

Parameters to be estimated

- Proliferation rates: r_N, r_P, r₄ and r₈
- Death rates: d_N , d_P , d_4 and d_8 .

• Transfer rates:
$$\mu_N(i) = (\alpha_N \cdot i)^n$$
, $\mu_P(i) = (\alpha_P \cdot i)^n$, μ_{LP} ,
 $e_4(i) = (\alpha_{e4} \cdot i)^n$ and $e_8(i) = (\alpha_{e8} \cdot i)^n$

• Number of stages: *n*₄, *n*₈

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On/off proliferation control

$$\gamma(t) = \left\{ egin{array}{cc} \mathsf{0} & ext{if } t < au_0 \ \mathsf{1} & ext{if } t \geq au_0 \end{array}
ight.$$

Continuous inhibition function

$$\gamma(t) = \begin{cases} \exp(-\delta_0 t) & \text{if } t < \tau_0 \\ 1/(1 + \exp(-\delta_1(t - \tau_1))) & \text{if } t \ge \tau_0 \end{cases}$$

- \(\tau_0\) is estimated
- exponential decrease of the proliferation rate and logistic growth of the proliferation rate
- different parameters for different populations

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Outline

Some problems requiring parameter estimation

- 2 Constrained evolutionary search of the parameter space
- 3 Case study: modelling a perturbed thymocyte dynamics
- Exploring the output of the evolutionary search

Exploring the output of the evolutionary search



Evolved solutions of similar quality ($MSE = 0.029 \pm 0.002$)

D. Zaharie (UVT)

Constrained Evolutionary Search

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Exploring the output of the evolutionary search



Summary

- in the case of "semi-transparent" or "opaque" models, estimating the parameters can lead to difficult constrained optimization problems
- for some real problems it is not easy the check if a constraint involving the parameters is satisfied or not; estimating a constraint satisfaction degree could be useful
- combining the MSE value with the constraint satisfaction degree could be beneficial
- an evolutionary search for parameters can provide several possible solutions => information about the distribution of parameters

Invitation to Timisoara - Romania

- ... at the High Performance Research Center (http://hpc.uvt.ro)
 - hybrid CPU+GPU cluster (450 cores, 7 Nvidia Tesla based blades, 40Gbps Infiniband, 750 GB RAM, 30TB storage)
 - IBM BlueGene/P (4096 cores, 11.7 Tflops, 4TB RAM, 28TB storage)
- ... open acces offered through FP7-REGPOT project HOST (http://host.hpc.uvt.ro)
- ... current research topics
 - Cloud computing technologies for HPC service exposure
 - Scheduling algorithms and techniques
 - Parallel computing in remote data processing
 - Large scale numerical computations
 - HPC-based intelligent services

Invitation to Timisoara - Romania

... at SYNASC 2013 - 15th Symposium on Symbolic and Numeric Algorithms for Scientific Computation - 23-26 September 2013 (http://synasc13.info.uvt.ro)

Program chair: Nikolaj Bjorner, Microsoft Research Tracks:

- Symbolic Computation
- Numeric Computing
- Logic and Programming
- Distributed Computing
- Artificial Intelligence
- Advances in the Theory of Computing

Invited speakers

- Ivona Brandic, Vienna University of Technology
- Gabriel Ciobanu, Romanian Academy, Iasi
- Leonardo de Moura, Microsoft Research, USA
- Grigore Rosu, University of Illinois at Urbana-Champaign
- Dan Simovici, University of Massachusetts, Boston

Invitation to Timisoara - Romania

... at SYNASC 2013 - 15th Symposium on Symbolic and Numeric Algorithms for Scientific Computation - 23-26 September 2013 (http://synasc13.info.uvt.ro) Workshops:

- ACSys: Agents for Complex Systems
- NCA: Natural Computing and Applications
- HPCSP: High Performance Computing for Scientific Problems
- MICAS: Management of Resources in Sky and Cloud Computing
- IAFP: Iterative Approximation of Fixed Points

Submission deadline for workshops: 15-30 July 2013 (depending on the workshop)