# Mathematical Optimization in Industrial Processes Application to the design of bioreactors for water treatment.

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#### 19th of September, 2013

- <u>Postulate</u>: We consider water resources polluted with biochemical substances (glucose,sulfate,etc), called substrates.
- <u>Objective</u>: Clean this water by using biological microorganisms (bacteria,enzymes,etc), called biomasses, which develop by substrate consumption.
- <u>How</u>?: Connecting the water resource in a closed circuit with a bioreactor.



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ODE Approach for the bioreactor PDE Approach for the bioreactor

## **Bioreactors**

The bioreactors operating continuously are also called *chemostats*. The chemostat was studied by Jacques Monod, Aaron Novick and Leo Szilard in 1950.



The bioreactor is fed from the resource with a flow rate Q, and its output returns the cleaned water with the same flow rate Q.

ODE Approach for the bioreactor PDE Approach for the bioreactor

Mathematical Model for a bioreactor: An ODE Approach

Homogeneous distribution of the contaminant in the bioreactor:

$$\begin{cases} \frac{\mathrm{d}S_{\mathrm{r}}}{\mathrm{d}t}(t) = -\mu(S_{\mathrm{r}}(t))B_{\mathrm{r}}(t) + \frac{Q(t)}{V_{\mathrm{r}}}(S_{\mathrm{e}}(t) - S_{\mathrm{r}}(t)) & t > 0, \\ \frac{\mathrm{d}B_{\mathrm{r}}}{\mathrm{d}t}(t) = \mu(S_{\mathrm{r}}(t))B_{\mathrm{r}}(t) - \frac{Q(t)}{V_{\mathrm{r}}}B_{\mathrm{r}}(t) & t > 0, \end{cases} (1) \\ S_{\mathrm{r}}(0) = S_{\mathrm{e}}(0), \qquad B_{\mathrm{r}}(0) = B_{\mathrm{r},0}, \end{cases}$$

- -Q: flow rate  $(m^3/s)$ .
- $-S_r$ : substrate concentration inside the reactor (mol/m<sup>3</sup>).
- $-B_r$ : biomass concentration inside the reactor (mol/m<sup>3</sup>).
- - $S_e$ : substrate concentration that enters the bioreactor (mol/m<sup>3</sup>).
- - $V_r$ : reactor volume(m<sup>3</sup>).
- - $\mu$ : growth rate function (s<sup>-1</sup>).

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#### Assumption:

- Function  $\mu(\cdot)$  is increasing and  $\mu(0) = 0$ .
- **2** Function  $\mu(\cdot)$  es concave.

An example of growth rate function is given by the Monod Equation:



P. Gajardo, J. Harmand, H. Ramírez C., and A. Rapaport. Minimal time bioremediation of natural water resources. *Automatica*, 47(8), 2011.

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Particular case:  $S_e$  and Q are constant. System (1) has two fixed points:

- $E_1 = (S_e, 0) \Rightarrow No \text{ decontamination (leaching)}$
- $E_2=(S^{
  m Q}_{
  m r},S_{
  m e}-S^{
  m Q}_{
  m r})$ , where  $S^{
  m Q}_{
  m r}$  fulfills  $Q=V_{
  m r}\mu(S^{
  m Q}_{
  m r})$ .

#### Theorem (Stability analysis)

**1** If  $Q < V_r \mu(S_e) \Rightarrow E_1$  unstable,  $E_2$  is asymptotically stable.

- 2 If  $Q > V_r \mu(S_e) \Rightarrow E_1$  asymptotically stable,  $E_2$  is unstable.
- If  $Q = V_r \mu(S_e) \Rightarrow E_1 = E_2$  asymptotically stable.

#### Proof.

For (1) and (2) we use *Hartman-Grobman Theorem*. For (3) we use *Poincare-Bendixson Theorem* and *Dulac's Criterion*.

ODE Approach for the bioreactor PDE Approach for the bioreactor

(\*)

$$\begin{array}{l} x \doteq F(x) \\ x(0) = x_0 \end{array}$$

Let 
$$x^*$$
 a critical point of (\*), i.e,  $F(x^*) = 0$ .

Linearization: 
$$z' = DF(x^*)z$$
, where  $z = x - x^*$ .

#### Theorem (Hartman-Grobman)

Suppose  $x^*$  is a hyperbolic critical point (i,.e the real part of the eigenvalues of DF are not zero). Then the phase portrait of the linearization and the nonlinear equations are locally homeomorphic.

#### Theorem

- Real part of the eigenvalues of DF are negative ⇒ x<sup>\*</sup> is asymptotically stable.
- At least one eigenvalue of DF is positive  $\Rightarrow x^*$  is unstable.

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#### Theorem (Generalized Poincare Bendixson)

If the positive orbit  $\gamma^+(x)$  in (\*) is contained in a compact set K, where K only contains a finite number of critical points, then:

- $\omega(x)$  critical point, or
- $\omega(x)$  periodic orbit, or
- $\omega(x)$  connected set composed of finite number of fixed points.

### Theorem (Dulac Criterion - Only for $F = (F_1, F_2)$ )

Let D be a simply connected region in the phase plane. If there exists  $C^1$  function  $\psi(x, y)$  such that

$$\frac{\mathrm{d}}{\mathrm{d}x}(\psi(x,y)F_1(x,y)) + \frac{\mathrm{d}}{\mathrm{d}y}(\psi(x,y)F_2(x,y))$$

has constant sign in D, then (2) has no closed orbits lying in D.

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# Nondimensionalization of the system

#### Definition

Partial or full removal of units from an equation involving physical quantities by a suitable substitution of variables

**1** List all of the variables, parameters, and their dimensions:

Variable	Dimension	Parameter	Dimension
$B_{ m r}$	mol/m <sup>3</sup>	$V_{ m r}$	m <sup>3</sup>
$S_{ m r}$	mol/m <sup>3</sup>	$S_{ m e}(0)$	$mol/m^3$
t	S	$B_{\mathrm{r},0}$	mol/m <sup>3</sup>
Q	m <sup>3</sup> /s		
$S_{ m e}$	mol/m <sup>3</sup>		
$\mu$	$s^{-1}$		

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Take each variable and create a new variable dividing by a combination of parameters that has the same dimension:

$$egin{aligned} s_{ ext{e}} &= rac{S_{ ext{e}}}{S_{ ext{e}}(0)}, \quad s_{ ext{r}} &= rac{S_{ ext{r}}}{S_{e}(0)}, \quad b_{ ext{r}} &= rac{B_{ ext{r}}}{S_{e}(0)}, \quad ilde{\mu}(s_{ ext{r}}) &= rac{\mu(S_{ ext{r}})}{\mu(S_{ ext{e}}(0))} \ q &= rac{Q}{V_{ ext{r}}\mu(S_{ ext{e}}(0))}, \quad egin{aligned} t_{ ext{r}} &= rac{t}{rac{1}{\mu(S_{ ext{e}}(0))}} \end{aligned}$$

Sewrite the differential equation in terms of the new variables:

$$\begin{cases} \frac{\mathrm{d}s_{\mathrm{r}}}{\mathrm{d}t_{\mathrm{r}}}(t_{\mathrm{r}}) = -\tilde{\mu}(s_{\mathrm{r}}(t_{\mathrm{r}}))b_{\mathrm{r}}(t_{\mathrm{r}}) + q(t_{\mathrm{r}})(s_{\mathrm{e}}(t_{\mathrm{r}}) - s_{\mathrm{r}}(t_{\mathrm{r}})), & t_{\mathrm{r}} > 0, \\ \\ \frac{\mathrm{d}b_{\mathrm{r}}}{\mathrm{d}t_{\mathrm{r}}}(t_{\mathrm{r}}) = \tilde{\mu}(s_{\mathrm{r}}(t_{\mathrm{r}}))b_{\mathrm{r}}(t_{\mathrm{r}}) - q(t_{\mathrm{r}})b_{\mathrm{r}}(t_{\mathrm{r}}), & t_{\mathrm{r}} > 0, \\ \\ s_{\mathrm{r}}(0) = 1, & b_{\mathrm{r}}(0) = \alpha, \\ \end{cases}$$
where  $\alpha = \frac{B_{\mathrm{r},0}}{S_{\mathrm{e}}(0)}.$ 

ODE Approach for the bioreactor PDE Approach for the bioreactor

Dynamics in the water resource and coupled system

Natural water resource polluted with a substrate concentration  $S_1$ .  $\implies$  <u>Objective</u>: Reduce the concentration of the pollutant to a prescribed value  $S_{\text{lim}}$ .



Settler: Collect the biomass  $\Rightarrow$  avoids new contamination.

ODE Approach for the bioreactor PDE Approach for the bioreactor

The dynamics in the water resource can be described

$$\begin{cases} \frac{\mathrm{d}S_1}{\mathrm{d}t}(t) = \frac{Q(t)}{V}(S_{\mathrm{r}}(t) - S_1(t)), & t > 0, \\ S_1(0) = S_{\mathrm{l},0} \end{cases}$$
(2)

• Dimensional Analysis

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Variable	Dimension	Parameter	Dimension	
<i>S</i> <sub>1</sub>	mol/m <sup>3</sup>	V	m <sup>3</sup>	
t	S	$S_{\mathrm{l},0}$	$mol/m^3$	
$S_{ m r}$	mol/m <sup>3</sup>			
Q	m <sup>3</sup> /s			
$s_1 = rac{S_1}{S_1(0)},  s_{ m r} = rac{S_{ m r}}{S_1(0)},  q = rac{Q}{V_{ m r}\mu(S_1(0))},  t_1 = rac{t}{rac{V}{V_{ m r}\mu(S_1(0))}}$				
$\int rac{\mathrm{d} s_1}{\mathrm{d} t_1}(t_1) = -q(t_1)(s_1(t_1)-s_\mathrm{r}(t_1)),$				
$\left( \begin{array}{c} s_1(0)=1. \end{array} \right.$				

ODE Approach for the bioreactor PDE Approach for the bioreactor

# Quasi-Steady State Approximation

**Hypothesis:** 
$$V >> V_{\rm r} \Rightarrow \tau_1 \sim \frac{V}{V_{\rm r}} \frac{1}{\mu(S_1(0))} >> \tau_{\rm r} \sim \frac{1}{\mu(S_1(0))}$$

Reasonable time scale for the bioreactor  $\Rightarrow$   $S_1$ , Q negligibly changes.

Remember: When  $Q < V_r \mu(S_1)$ , the equilibrium point  $(S_r^Q, S_1 - S_r^Q)$  is asymptotically stable.

Reasonable time scale for the water resource  $\Rightarrow$  the bioreactor attains its equilibrium point.

Thus, System (2) can be approximated as:

$$\begin{cases} \frac{\mathrm{d}S_{1}}{\mathrm{d}t}(t) = \frac{Q(t)}{V}(S_{\mathrm{r}}^{\mathrm{Q}}(t) - S_{1}(t)), & t > 0, \\ S_{1}(0) = S_{1,0}. \end{cases}$$
(3)

where  $S^{ ext{Q}}_{ ext{r}}(t) \in [0, S_1(t)).$ 

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Q is constant in time ,  $\mu(\cdot)$  Monod Equation.  $(S_{r}^{*}, B_{r}^{*}) = (S_{r}^{Q}, S_{1}(t) - S_{r}^{Q})$  where  $S_{r}^{Q}$  fulfills  $Q = V_{r}\mu(S_{r}^{Q})$ .



ODE Approach for the bioreactor PDE Approach for the bioreactor

Mathematical Model for the Bioreactor: A PDE Approach

Spatial disparity of the contaminant in the bioreactor.



- Substrate and Biomass concentrations  $\Rightarrow$  Convection-Diffusion-Reaction equation.
- Fluid Flow  $\Rightarrow$  Vertical inflow:  $\mathbf{u} = (0, 0, -\bar{Q}(t))$  where  $\bar{Q}$  (m/s) is the flow rate per unit of area.

$$\begin{cases} \frac{\mathrm{d}S_{\mathrm{r}}}{\mathrm{d}t} = \nabla \cdot (D_{\mathrm{S}} \nabla S_{\mathrm{r}}) - \mathbf{u} \nabla S_{\mathrm{r}} - \mu(S_{\mathrm{r}}) B_{\mathrm{r}} & \text{in } \Omega^{*} , t > 0, \\ \frac{\mathrm{d}B_{\mathrm{r}}}{\mathrm{d}t} = \nabla \cdot (D_{\mathrm{B}} \nabla B_{\mathrm{r}}) - \mathbf{u} \nabla B_{\mathrm{r}} + \mu(S_{\mathrm{r}}) B_{\mathrm{r}} & \text{in } \Omega^{*} , t > 0. \end{cases}$$
(4)

 $D_{\rm S}, D_{\rm B}$  substrate and biomass diffusion coefficients (m<sup>2</sup>/s).

ODE Approach for the bioreactor PDE Approach for the bioreactor

We consider suitable initial conditions:

• 
$$S_{\mathrm{r}}(0,x)=S_{1}(0),$$
  $B_{\mathrm{r}}(0,x)=B_{\mathrm{init}}$   $orall x\in\Omega^{*},$ 

#### and boundary conditions:

Flux of the system:  $J = -D\nabla c + \mathbf{u}c$ 

D is the diffusion, **u** is the fluid flow and c is the concentration.

• 
$$\begin{cases} \mathbf{n} \cdot (-D_{\mathrm{S}} \nabla S_{\mathrm{r}} + \mathbf{u} S_{\mathrm{r}}) = -\bar{Q}(t) S_{\mathrm{e}}(t) \\ \mathbf{n} \cdot (-D_{\mathrm{B}} \nabla B_{\mathrm{r}} + \mathbf{u} B_{\mathrm{r}}) = 0 \end{cases} \quad \forall x \in \Gamma_{\mathrm{in}}^{*}, \ t > 0,$$
  
• 
$$\begin{cases} \mathbf{n} \cdot (-D_{\mathrm{S}} \nabla S_{\mathrm{r}}) = 0 \\ \mathbf{n} \cdot (-D_{\mathrm{B}} \nabla B_{\mathrm{r}}) = 0 \end{cases} \quad \forall x \in \Gamma_{\mathrm{wall}}^{*} \cup \Gamma_{\mathrm{out}}^{*}, \ t > 0. \end{cases}$$

ODE Approach for the bioreactor PDE Approach for the bioreactor

#### Simplification:



 $\Rightarrow$  Change to cylindrical coordinates (r, z) where r is the distance to the cylinder axis.

System (4) can be rewritten as:

$$\begin{cases} \frac{dS_{\rm r}}{dt} = \frac{1}{r} \frac{\mathrm{d}}{\mathrm{d}r} (rD_{\rm S} \frac{\mathrm{d}S_{\rm r}}{\mathrm{d}r}) + \frac{\mathrm{d}}{\mathrm{d}z} (D_{\rm S} \frac{\mathrm{d}S_{\rm r}}{\mathrm{d}z}) + \bar{Q}(t) \frac{\mathrm{d}S_{\rm r}}{\mathrm{d}z} - \mu(S_{\rm r})B_{\rm r} & \text{in } \Omega, t > 0, \\ \frac{\mathrm{d}B_{\rm r}}{\mathrm{d}t} = \frac{1}{r} \frac{\mathrm{d}}{\mathrm{d}r} (rD_{\rm B} \frac{\mathrm{d}B_{\rm r}}{\mathrm{d}r}) + \frac{\mathrm{d}}{\mathrm{d}z} (D_{\rm B} \frac{\mathrm{d}B_{\rm r}}{\mathrm{d}z}) + \bar{Q}(t) \frac{\mathrm{d}B_{\rm r}}{\mathrm{d}z} + \mu(S_{\rm r})B_{\rm r} & \text{in } \Omega, t > 0, \end{cases}$$

$$(5)$$

ODE Approach for the bioreactor PDE Approach for the bioreactor

We consider suitable initial conditions:

•  $S_{\mathrm{r}}(0,r,z)=S_{1}(0),$   $B_{\mathrm{r}}(0,r,z)=B_{\mathrm{init}}$   $\forall (r,z)\in\Omega,$ 

#### and boundary conditions:

• 
$$\begin{cases} D_{\rm S} \frac{\mathrm{d}S_{\rm r}}{\mathrm{d}z} + \bar{Q}(t)S_{\rm r} = \bar{Q}(t)S_{\rm e}(t) \\ D_{\rm B} \frac{\mathrm{d}B_{\rm r}}{\mathrm{d}z} + \bar{Q}(t)B_{\rm r} = 0 \end{cases} \quad \forall (r,z) \in \Gamma_{\rm in}, \ t > 0, \\ \frac{\mathrm{d}S_{\rm r}}{\mathrm{d}r} = 0 \quad \forall (r,z) \in \Gamma_{\rm wall} \cup \Gamma_{\rm sym} \cup \Gamma_{\rm out}, \ t > 0. \end{cases}$$

ODE Approach for the bioreactor PDE Approach for the bioreactor

The dynamics of the water resource can be described as follows:

$$\begin{cases} \frac{\mathrm{d}S_{1}}{\mathrm{d}t}(t) = \frac{Q(t)}{V}(S_{\mathrm{out}}(t) - S_{1}(t)), \quad t > 0, \\ S_{1}(0) = S_{1,0}. \end{cases}$$
(6)

where  $S_{out}(t) = \frac{\int_{\Gamma_{out}} S_r(t,r,z) dr}{L}$  and  $\Gamma_{out}$  is the outlet through

which treated water leaves the reactor.

Bioreactor with ODE model. Theoretical results Bioreactor with PDE model. Genetic Algorithms

# Optimization problem: Bioreactor with ODE model

Find 
$$Q^{\text{opt}} \in X$$
, such that  
 $T(Q^{\text{opt}}) = \min_{Q \in X} T(Q)$ , (7)

where T(Q) is the time required for achieving  $S_1(T(Q)) = S_{\lim}$ , with  $S_1$  being the solution of (2) (or 6).  $X = \{Q \text{ piecewise } C^1([0, +\infty)) : 0 \le Q(t) < V_r \mu(S_1(t)) \ \forall t \ge 0\}.$ 

## • Flux is constant in time

#### Theorem

If Q is constant, then the time required for the solution of (3) to attain the value  $S_{\rm lim}$  is:

$$\mathcal{T}(S^{\mathrm{Q}}_{\mathrm{r}}) = rac{1}{rac{V_{\mathrm{r}}}{V} \mu(S^{\mathrm{Q}}_{\mathrm{r}})} \ln \left( rac{S_{\mathrm{1,0}} - S^{\mathrm{Q}}_{\mathrm{r}}}{S_{\mathrm{lim}} - S^{\mathrm{Q}}_{\mathrm{r}}} 
ight)$$

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## • Q is a time variable

#### Definition

If a functional relation of the form

$$S^{\mathrm{Q}}_{\mathrm{r}}(t) = \omega(S_1(t)) \qquad orall t \in [0,+\infty)$$

can be found for the optimal control at time t for problem (7), then  $\omega$  is called optimal feedback.

#### Theorem

An optimal feedback  $S_{\rm r}^{\rm opt}$  must fulfill

$$S^{ ext{opt}}_{ ext{r}}(t) = rg\min_{S^{ ext{Q}}_{ ext{r}}(t) \in (0, \mathcal{S}_{1}(t))} \mu(S^{ ext{Q}}_{ ext{r}}(t))(S^{ ext{Q}}_{ ext{r}}(t) - \mathcal{S}_{1}(t)).$$

Moreover t 
ightarrow Q(t) is decreasing along any optimal trajectory.

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**Q** is constant in time  $\Rightarrow \Theta = [0, V_r \mu(S_{lim})).$ 

**Q** is a time variable: Q(t) is  $C^1$  and should be decreasing on time!

We work with 5 optimization parameters:

$$(Q_0, lpha_1, lpha_2, lpha_3, lpha_4) \in \Theta = [0, V_\mathrm{r}\mu(S_1(0))) imes [0, 1]^4$$

such that

$$Q(t) = \left\{egin{array}{ccc} Q_0 & t=0\ Q_0\cdotlpha_1 & t=t_1\ Q_0\cdotlpha_1\cdotlpha_2 & t=t_2\ Q_0\cdotlpha_1\cdotlpha_2\cdotlpha_3 & t=t_3\ Q_0\cdotlpha_1\cdotlpha_2\cdotlpha_3\cdotlpha_4 & t=t_4 \end{array}
ight.$$

for 4 different fixed times  $t_1$ ,  $t_2$ .  $t_3$  and  $t_4$ In the time intervals  $(t_i, t_{i+1})$  (i = 0, ..., 3), Q(t) is calculated with Cubic Hermite Spline

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It is reasonable to think that in practice it is easier to use Piecewise Constant Q(t).

$$Q(t)=\left\{egin{array}{ll} Q_0&t\in[0,t_1)\ Q_0\cdotlpha_1&t\in[t_1,t_2)\ Q_0\cdotlpha_1\cdotlpha_2&t\in[t_2,t_3)\ Q_0\cdotlpha_1\cdotlpha_2\cdotlpha_3&t\in[t_3,t_4)\ Q_0\cdotlpha_1\cdotlpha_2\cdotlpha_3\cdotlpha_4&t\in[t_4,t_5) \end{array}
ight.$$

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# Genetic Algorithms

Global optimization method based on a natural selection process that mimics biological evolution.

 $\min_{x\in\Theta}f(x)$ 

- $\Theta \subset R^{\mathbb{N}}$  is the search space.
  - Creation a random initial population in the search space.
  - Successive reproduction of the population by the stochastic steps Selection, Crossover, Mutation and Elitism.

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## Genetic Algorithm: Sketch



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• Selection: Individuals are selected according to their fitness value.

#### Roulette Wheel Selection:

Individual number	Individual	Fitness value	% of Total
1	(0,1)	169	22.72
2	(1, 0)	576	6.67
3	(0,0)	64	59.99
4	(1, 1)	361	10.63
Total		1170	100.0

Table: Roulette Wheel Selection.



• **Crossover:** Create a new solution candidate by combining the characteristics of two existing individuals Arithmetic Crossover

$$\mathsf{Chil} 1 = a \cdot \mathsf{Parent} 1 + (1 - a) \cdot \mathsf{Parent} 2$$
  
 $\mathsf{Chil} 2 = (1 - a) \cdot \mathsf{Parent} 1 + a \cdot \mathsf{Parent} 2$ 

• **Mutation:** Randomly modifies the value of one or more genes of an individual.

Non-Uniform mutation: To mutate an individual x, we compute

$$x' = \left\{egin{array}{l} x+\Delta(g,b-x) & ext{if } au=0 \ x-\Delta(g,b-x) & ext{if } au=1 \end{array}
ight.$$

where  $\Delta(g, y) = (1 - r^{(1 - \frac{g}{g_{\max}})^b}).$ 

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• Elitism: Ensures that at least one copy of the best individual(s) of the current generation is directly copied to the next generation.

#### Matrix formulation

$$X^{i+1} = (I_N - \mathcal{E}^i)(\mathcal{C}^i \mathcal{S}^i X^i + \mathcal{M}^i) + \mathcal{E}^i X^i$$

- - $X^i$ : Current generation
- - $\mathcal{E}^i$ : Elitism operator
- - $C^i$ : Crossover operator
- - $S^i$ : Selection operator
- - $\mathcal{M}^i$ : Mutation operator
- -*I<sub>N</sub>*: Identity matrix

## • Stopping criteria:

Generation number

No Improvement generation number

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#### Hybrid Genetic Algorithm

Coarse search of global minimum with GA



Prefine the local search by using a Gradient Descent method.

$$\mathbf{x}_{k+1} = \mathbf{x}_k - \nabla f(\mathbf{x}_k) \alpha_k,$$

where  $\alpha_k$  is the step length which must fulfill

$$f(x_{\mathrm{k}}-
abla f(x_{\mathrm{k}})lpha_{\mathrm{k}})\leq f(x_{\mathrm{k}}-
abla f(x_{\mathrm{k}})lpha) \quad ext{ for all } lpha\geq 0.$$



## Numerical Optimization Results: ODE Approach

### $\mu(\cdot)$ Monod Function



#### Optimal feedback can reduce the decontamination time by half!

## Numerical Optimization Results: PDE Approach

 $\mu(\cdot)$  Monod,  $\mathcal{S}_{\mathrm{lim}}=$  0.1,  $\mathcal{Q}(t)$  continuously differentiable.

$S_1(0) \; (mol/m^3)$	Time ${\it Q}_{ m HGA}^{ m opt}({\sf const})$ (s)	Time $\mathit{Q}_{ m HGA}^{ m opt}({\sf Tdep})$ (s)	
$D_{ m S} = D_{ m B} = 100 \; ({ m m}^2/{ m s})$			
5	72750	46650	
10	81840	48090	
20	90760	49360	
$D_{ m S} = D_{ m B} = 0.01~({ m m}^2/{ m s})$			
5	87710	55120	
10	100360	57870	
20	58870	26460	

#### Time dependent Q can reduce the decontamination time in 55%!

## Numerical Optimization Results: PDE Approach

 $\mu(\cdot)$  Monod,  $S_{
m lim}=$  0.1, Q(t) Piecewise constant.

$S_1(0) \; (mol/m^3)$	Time ${\it Q}_{ m HGA}^{ m opt}({\sf const})$ (s)	Time $\mathit{Q}_{ m HGA}^{ m opt}({\sf Tdep})$ (s)	
$D_{ m S} = D_{ m B} = 100 \; ({ m m}^2/{ m s})$			
5	72750	49390	
10	81840	51560	
20	90760	55010	
$D_{ m S} = D_{ m B} = 0.01~({ m m}^2/{ m s})$			
5	87710	60120	
10	100360	62990	
20	58870	23340	

#### Time dependent Q can reduce the decontamination time in 60%!

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# Comparison between ODE and PDE Approaches: Q(t) continuously differentiable

 $D_\mathrm{S}=D_\mathrm{B}=100~(\mathrm{m}^2/\mathrm{s})$ 

- Both fluxes are suitable.
- Decontamination times are comparable.



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$$D_{
m S} = D_{
m B} = 0.01 \; ({
m m}^2/{
m s})$$

 Optimal fluxes obtained with the ODE approach are not suitable ⇒ we are not able to achieve the target.



Mathematical Optimization in Industrial Processes

# Comparison between ODE and PDE Approaches: Q(t) piecewise constant

 $D_\mathrm{S} = D_\mathrm{B} = 100~(\mathrm{m^2/s})$ 

- Both fluxes are suitable.
- Decontamination times are comparable.



$$D_{\rm S} = D_{\rm B} = 0.01 \; ({\rm m}^2/{\rm s})$$

 Optimal fluxes obtained with the ODE approach are not suitable ⇒ we are not able to achieve the target.



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Mathematical Optimization in Industrial Processes

Numerical Optimization Results: Comparison between Q(t) continuously differentiable and piecewise constant

 $\mu(\cdot)$  Monod,  $S_{
m lim}=$  0.1

$S_1(0) \text{ (mol/m}^3)$	Time $Q_{ m HGA}^{ m opt}( m CD)$ (s)	Time $Q_{ m HGA}^{ m opt}( m PC)$ (s)	
$D_{ m S} = D_{ m B} = 100 \; ({ m m}^2/{ m s})$			
5	46650	49390	
10	48090	51560	
20	49360	55010	
$D_{ m S} = D_{ m B} = 0.01~({ m m}^2/{ m s})$			
5	55120	60120	
10	57870	62990	
20	26460	23340	

Q(t) continuously differentiable can reduce the decontamination time in 10%!

# Conclusions

- ODE approach for the bioreactor: We obtain the optimal flux with theoretical results.
- PDE approach for the bioreactor: We obtain the optimal flux using an Hybrid Genetic Algorithm.
- The decontamination time can be reduced by half if the optimal flux is chosen as a time variable rather than constant in time.
- For small diffusivities  $D_{\rm S}$  and  $D_{\rm B}$  the theoretical results obtained for the ODE approach are not suitable. We are not able to achieve the target.

## Future work

- Give an existence and uniqueness result for the system presented in the PDE approach for the bioreactor.
- Give a suitable nondimensionalization for the system presented in the PDE approach.
- More intensive study of HGA.
- Consider inhomogeneity in the water resource.

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