Motivation	Description of the method	biochemical process example	Conclusion	References
00	0000	000000		

# An Approximation of Minimax Control using Random Sampling and Symbolic Computation

### Jawher Jerray <sup>1</sup> Laurent Fribourg<sup>2</sup> Étienne André<sup>3</sup>

<sup>1</sup> Université Sorbonne Paris Nord, LIPN, CNRS, UMR 7030, F-93430, Villetaneuse, France and <sup>2</sup>Université Paris-Saclay, CNRS, ENS Paris-Saclay, LMF, F91190 Gif-sur-Yvette, France and <sup>3</sup>Université de Lorraine, CNRS, Inria, LORIA, F-54000 Nancy, France

Friday 9th July, 2021

ADHS 2021



Motivation	Description of the method	biochemical process example	Conclusion O	References
Outline				

#### 1 Motivation

2 Description of the method

3 biochemical process example

### 4 Conclusion



Motivation	Description of the method	biochemical process example	Conclusion	References
0				
Motivatio	n			

#### Dynamical systems:

- in which a function describes the time dependence of a point in a geometrical space.
- we only know certain observed or calculated states of its past or present state.
- dynamical systems have a direct impact on human development.

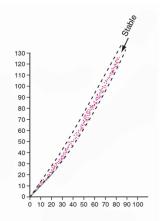
#### $\Rightarrow$ The importance of studying:

- synchronization
- behavior
- robust control



Motivation	Description of the method	biochemical process example	Conclusion	References
00				
Robusto	699			

- A control is considered robust if the dynamical system still stable, which means small perturbations to the solution using this control lead to a new solution that stays close to the original solution forever.
- A stable system produces a bounded output for a given bounded input.





Motivation 00	Description of the method ●000	biochemical process example	Conclusion O	References
Descript	ion of the method			

- We consider a dynamic system with control  $u(\cdot)$  and a bounded disturbance function  $d(\cdot)$  over a domain *D*). The control  $u(\cdot)$  is a piecewise constant function that changes of value only at times  $t = \tau, 2\tau, \ldots$ .
- The set of possible controls of the system for  $t \in [0, T]$  is finite and can be described by the set  $\mathcal{U} \equiv U^K$  where  $T = K\tau$  and  $K \in \mathbb{N}$ .



Motivation 00	Description of the method ●000	biochemical process example	Conclusion O	References
Descripti	on of the method			

- We consider a dynamic system with control  $u(\cdot)$  and a bounded disturbance function  $d(\cdot)$  over a domain *D*). The control  $u(\cdot)$  is a piecewise constant function that changes of value only at times  $t = \tau, 2\tau, \ldots$ .
- The set of possible controls of the system for  $t \in [0, T]$  is finite and can be described by the set  $\mathcal{U} \equiv U^K$  where  $T = K\tau$  and  $K \in \mathbb{N}$ .
- We suppose given a cost function  $C : \mathbb{R}^n \times U^K \to \mathbb{R}_{\geq 0}$ , which allows calculating the value  $\int_0^T C(x(t), u(t)) dt$  for any solution x(t) of  $\dot{x}(t) = f(x(t), u(t), d(t))$  for  $t \in [0, T]$ .



Motivation 00	Description of the method •••••	biochemical process example	Conclusion O	References
Descript	tion of the method			

- We consider a dynamic system with control  $u(\cdot)$  and a bounded disturbance function  $d(\cdot)$  over a domain *D*). The control  $u(\cdot)$  is a piecewise constant function that changes of value only at times  $t = \tau, 2\tau, \ldots$ .
- The set of possible controls of the system for  $t \in [0, T]$  is finite and can be described by the set  $\mathcal{U} \equiv U^K$  where  $T = K\tau$  and  $K \in \mathbb{N}$ .
- We suppose given a cost function  $C : \mathbb{R}^n \times U^K \to \mathbb{R}_{\geq 0}$ , which allows calculating the value  $\int_0^T C(x(t), u(t)) dt$  for any solution x(t) of  $\dot{x}(t) = f(x(t), u(t), d(t))$  for  $t \in [0, T]$ .
- The *minimax method* aims to find a control *v* defined by:

$$v = \operatorname*{arg\,min}_{u \in U^K} \max_{x(\cdot)} \mathcal{J}_{z_0,\varepsilon}(x(\cdot), u(\cdot))$$

with  $\mathcal{J}_{z_0,\varepsilon}(x(\cdot), u(\cdot)) \equiv \left\{ \int_0^T C(x(t), u(t)) dt \mid \exists d(\cdot) \in D : \dot{x}(t) = f(x(t), u(t), d(t)) \text{ for } t \in [0, T] \land x(0) \in B(z_0, \varepsilon) \right\}.$ 

Motivation 00	Description of the method •••••	biochemical process example	Conclusion O	References
Descript	tion of the method			

- We consider a dynamic system with control  $u(\cdot)$  and a bounded disturbance function  $d(\cdot)$  over a domain *D*). The control  $u(\cdot)$  is a piecewise constant function that changes of value only at times  $t = \tau, 2\tau, \ldots$ .
- The set of possible controls of the system for  $t \in [0, T]$  is finite and can be described by the set  $\mathcal{U} \equiv U^K$  where  $T = K\tau$  and  $K \in \mathbb{N}$ .
- We suppose given a cost function  $C : \mathbb{R}^n \times U^K \to \mathbb{R}_{\geq 0}$ , which allows calculating the value  $\int_0^T C(x(t), u(t)) dt$  for any solution x(t) of  $\dot{x}(t) = f(x(t), u(t), d(t))$  for  $t \in [0, T]$ .
- The *minimax method* aims to find a control *v* defined by:

$$v = \underset{u \in U^{K}}{\arg\min\max} \mathcal{J}_{z_{0},\varepsilon}(x(\cdot), u(\cdot))$$

with  $\mathcal{J}_{z_0,\varepsilon}(x(\cdot), u(\cdot)) \equiv \left\{ \int_0^T C(x(t), u(t)) dt \mid \exists d(\cdot) \in D : \dot{x}(t) = f(x(t), u(t), d(t)) \text{ for } t \in [0, T] \land x(0) \in B(z_0, \varepsilon) \right\}.$ 

• We propose here a simplified method composed of two steps.



Motivation	Description of the method	biochemical process example	Conclusion	References
	0000			

## Description of the method

# First Step:

■ Obtain an upper-bound  $\mathcal{K}_{z_0,\varepsilon}(u(\cdot))$  of  $\max_{x(\cdot)} \mathcal{J}_{z_0,\varepsilon}(x(\cdot), u(\cdot))$  using an Euler-based symbolic computation method, with:

$$\mathcal{K}_{z_0,\varepsilon}(u(\cdot)) \equiv \max_{x(\cdot)\in \mathcal{B}(\tilde{x}_{z_0}^{u(\cdot)}(\cdot),\delta_{\varepsilon,D}^{u(\cdot)}(\cdot))} \{\int_0^I \mathcal{C}(x(t),u(t))dt\},\$$

where

- $\tilde{x}_{z_0}^u(\cdot)$  denotes Euler's approximate solution of  $\dot{x}(t) = f(x(t), u(t), \mathbf{0})$  for  $t \in [0, T]$  with null perturbation (i. e.,  $d(\cdot) = 0$ ) and initial condition  $z_0 \in \mathbb{R}^n$ ,
- $\delta_{e,0}^{(r,j)}(\cdot)$  denotes the upper-bound of the distance between an exact solution and an Euler approximate solution,
- $\mathbf{x}(\cdot) \in B(\tilde{\mathbf{x}}_{z_0}^{u(\cdot)}(\cdot), \delta_{\varepsilon, D}^{u(\cdot)}(\cdot))$  means, for all  $t \in [0, T]$ :  $\mathbf{x}(t) \in B(\tilde{\mathbf{x}}_{z_0}^{u(\cdot)}(t), \delta_{\varepsilon, D}^{u(\cdot)}(t))$ . In particular  $\mathbf{x}(0) \in B(z_0, \varepsilon)$ .<sup>1</sup>



<sup>1</sup> $y \in B(z, a)$  with  $y, z \in \mathbb{R}^n$  and  $a \ge 0$  means  $||y - z|| \le a$  where  $|| \cdot ||$  denotes the Euclidean norm.

Motivation	Description of the method	biochemical process example	Conclusion	References
	0000			

## Description of the method

# Second Step:

- We will not consider the absolute minimum, but a *probable near-minimum* of  $\mathcal{K}_{z_0,\varepsilon}(u(\cdot))$  (see [Vid01]).
- The probably approximate near-minimum of  $\mathcal{K}_{z_0,\varepsilon}$  is obtained by drawing randomly *N* control  $u_1, \cdots, u_N$  of  $U^K$ , i.e., by generating *N* independent identically distributed (i.i.d.) samples  $u_1, \cdots, u_N$  of  $U^K$ , with a uniform probability (i. e., with probability  $1/|U|^N$ ) then by taking  $\mathcal{K}_{z_0,\varepsilon}(u_N^*)$  with  $u_N^* = \arg\min_{u_1, \cdots, u_N} \mathcal{K}_{z_0,\varepsilon}(u_i)$ .



Motivation 00	Description of the method ○○○●	biochemical process example	Conclusion O	References
Descript	tion of the method			

#### Description of the method

# Advantage of the method:

- Avoid the excessive complexity of minimax methods,
- Use of samples with large size, as is often the case in statistical learning.
- Take into account *constraints* on the state of the system during its evolution.



Motivation	Description of the method	biochemical process example ●000000	Conclusion O	References
Biochemica	al process example			

Consider a biochemical process model *Y* of continuous culture fermentation (see [HouskaCDC09]) and initial condition in  $B_0 = B(x_0, \varepsilon)$  for some  $x_0 \in \mathbb{R}^2$  and  $\varepsilon > 0$  (see [BQ20]):. Let  $Y = (X, S, P) \in \mathbb{R}^3$  satisfies the differential system:

$$\begin{cases} \frac{dX}{dt} = -DX(t) + \mu(t)X(t) \\ \frac{dS}{dt} = D(S_f(t) - S(t)) - \frac{\mu(t)X(t)}{Y_{X/S}} \\ \frac{dP}{dt} = -DP + (\alpha\mu(t) + \beta)X(t) \end{cases}$$

<sup>[</sup>BQ20] J. B. van den Berg and E. Queirolo, "A general framework for validated continuation of periodic orbits in systems of polynomial ODEs," Journal of Computational Dynamics, vol. 0, no. 2158-2491-2019-0-10, 2020, ISSN: 2158-2491-D01: 10, 3934/jcd. 2021004.



Motivation	Description of the method	biochemical process example	Conclusion O	References
Biochemi	cal process exampl	e		

Consider a biochemical process model *Y* of continuous culture fermentation (see [HouskaCDC09]) and initial condition in  $B_0 = B(x_0, \varepsilon)$  for some  $x_0 \in \mathbb{R}^2$  and  $\varepsilon > 0$  (see [BQ20]):. Let  $Y = (X, S, P) \in \mathbb{R}^3$  satisfies the differential system:

$$\begin{cases} \frac{dX}{dt} = -DX(t) + \mu(t)X(t) \\ \frac{dS}{dt} = D(S_{f}(t) - S(t)) - \frac{\mu(t)X(t)}{Y_{x/s}} \\ \frac{dP}{dt} = -DP + (\alpha\mu(t) + \beta)X(t) \end{cases}$$

The model is controlled by  $S_f \in [S_f^{min}, S_f^{max}]$  and the specific growth rate  $\mu : \mathbb{R} \to \mathbb{R}$  of the biomass is a function of the states:

$$\mu(t) = \mu_m \frac{\left(1 - \frac{P(t)}{P_m}\right)S(t)}{K_m + S(t) + \frac{S(t)^2}{K_i}}$$

[BQ20] J. B. van den Berg and E. Queirolo, "A general framework for validated continuation of periodic orbits in systems of polynomial ODEs," Journal of Computational Dynamics, vol. 0, no. 2158-2491-2019-0-10, 2020, ISSN: 2158-2491. DOI: 10.3934/jcd.2021004.



Motivation	Description of the method	biochemical process example	Conclusion O	References
Biochemica	al process example			

Maximize the average productivity presented by the cost function:

$$\mathcal{J}_{Z_0,\varepsilon}(x(\cdot),S_f(\cdot)) = \frac{1}{T}\int_0^T DP(t)dt$$

While satisfying the constraint on the state X:

$$\frac{1}{T}\int_0^T X(t)dt \le 5.8$$



Motivation 00	Description of the method	biochemical process example ○○●○○○	Conclusion O	References
Biochemi	cal process examp	ام		

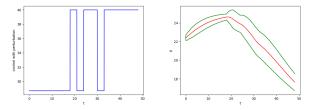
- The set  $S \subset \mathbb{R}^n \equiv [3, 8] \times [10, 28] \times [15.5, 25.5]$ .
- The codomain [28.7, 40] of the original continuous control function  $S_f(\cdot)$  is discretized into a finite set U. After discretization,  $S_t(\cdot)$  is a piecewise-constant function that takes its values in the finite set U made of 2 values uniformly taken in {28.7, 40}.
- We take:  $z_0 = (6.52, 12.5, 22.40), \tau = 3, \Delta t = \tau/100^2, T = 48, K = T/\tau = 16.$ We consider an additive disturbance *d* with  $d(\cdot) \in \mathcal{D} = [-0.05, 0.05]$ .
- In total, we have  $2^k = 2^{16}$  possible control cases.



 $<sup>^{2}\</sup>Delta t$  is the "sub-sampling' parameter of the Euler scheme.

Motivation 00	Description of the method	biochemical process example	Conclusion O	References
<b>—</b> · · ·				

### Biochemical process example



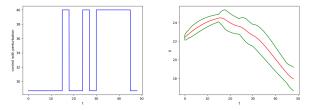
Left: control  $u^*$  satisfying the constraint on X, obtained by selection among 655 samples picked randomly; right: P(t) under  $u^*$  without perturbation (red curve) and with an additive perturbation  $d \in [-0.05, 0.05]$  (green curve) over 1 period (T = 48) for  $\Delta t = 1/400$  and initial condition (X(0), S(0), P(0)) = (6.52, 12.5, 22.4).

- We randomly pick one sample over every 100 possible controls, which gives  $2^{16}/100\approx 655$  samples.
- We get:  $\mathcal{K}_{z_0,\varepsilon}(u^*) = 3.1618$  (the constraint on the state *X* is satisfied since  $\frac{1}{T} \int_0^T X(t) dt = 5.782 \le 5.8$ ). The CPU computation time of this example is 7 seconds.



Motivation	Description of the method	biochemical process example	Conclusion O	References

## Biochemical process example



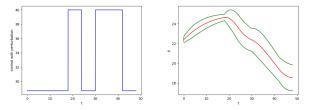
Left: control  $u^*$  satisfying the constraint on X, obtained by selection among 6554 samples picked randomly; right: P(t) under  $u^*$  without perturbation (red curve) and with an additive perturbation  $d \in [-0.05, 0.05]$  (green curve) over 1 period (T = 48) for  $\Delta t = 1/400$  and initial condition (X(0), S(0), P(0)) = (6.52, 12.5, 22.4).

- We randomly pick one sample over every 10 possible controls, which gives  $2^{16}/10\approx 6,554$  samples.
- We get:  $\mathcal{K}_{z_0,\varepsilon}(u^*) = 3.1667$ . (the constraint on the state X is satisfied since  $\frac{1}{T} \int_0^T X(t) dt = 5.794 \le 5.8$ ) The CPU computation time of this example is 18.69 seconds.



Motivation	Description of the method	biochemical process example	Conclusion	References
		000000		
<b>D</b> 1 1	· · · ·			

### Biochemical process example



Left: control  $u^*$  satisfying the constraint on X, obtained by selection among 65536 samples picked randomly; right: P(t) under  $u^*$  without perturbation (red curve) and with an additive perturbation  $d \in [-0.05, 0.05]$  (green curve) over 1 period (T = 48) for  $\Delta t = 1/400$  and initial condition (X(0), S(0), P(0)) = (6.52, 12.5, 22.4).

We consider all the possible controls, which gives  $2^{16} = 65,536$  samples. (The computation is tractable in this example because the set *U* contains only 2 modes, and because the length *K* of the horizon is moderate.)

We get:  $\mathcal{K}_{z_0,\varepsilon}(u^*) = 3.1677$  (the constraint is satisfied since  $\frac{1}{T} \int_0^T X(t) dt = 5.7995 \le 5.8$ ). The CPU computation time of this example is 200 seconds.



Motivation	Description of the method	biochemical process example	Conclusion	References
			•	

# Conclusion

#### Conclusion

- We showed that the simple combination of random sampling with a symbolic computation method allows to deal with robust optimization problems for nonlinear systems on non-convex domains.
- The method doesn't contain sophisticated theories such as analysis of viscosity solutions of the Hamilton-Jacobi-Bellman-Isaacs equation.



Motivation 00	Description of the method	biochemical process example	Conclusion O	References
[BQ20]	continuation of periodic	E. Queirolo, "A general frame orbits in systems of polynomi ics, vol. 0, no. 2158-2491-20 34/jcd.2021004.	ial ODEs," <mark>Journal</mark> d	
[Vid01]	statistical learning theor	mized algorithms for robust co y," <b>Automatica</b> , vol. 37, no. 1 .0.1016/S0005-1098(01)001	10, pp. 1515–1528, 2	

