

# **Evaluation of Derivative-Free Optimizers for Parameter Estimation in Systems Biology**

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# **Motivation**

#### **Parameter estimation**



For an objective  $J : \mathbb{R}^n \to \mathbb{R}$  find

 $\min_{\theta} J(\theta)$ 

subject to  $\ell \leq \theta \leq u$ .

## Derivative-free optimization: Why?

- J is given as a black-box
- J is noisy, costly, or non-smooth
- J is multi-modal
- simple to use



→ J(θ)



θ

model

# Optimizers

- FMINCON
- IMFIL
- BOBYQA
- PATTERNSEARCH-GPS
- PATTERNSEARCH-MADS
- RCS
- DHC
- MEIGO-DHC
- FMINSEARCHBND
- SIMULANNEALBND



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#### Multi-start local optimization



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- DIRECT
- MCS
- CMAES
- GA
- PARTICLESWARM
- PSWARM
- MEIGO-ESS-BOBYQA
- MEIGO-ESS-DHC
- MEIGO-ESS-MEIGO-DHC

## **Global methods**



- DIRECT
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## **Evaluation on classic test problems**

- cover issues optimizers typically have problems with
- parameters: 2, 3, 4, 5, 10, 15, 20, 25, 30, 35, 40, 50, 75, 100, 200, 300

	smooth	non-smooth	total
convex	130	65	195
non-convex	157	114	271
total	287	179	466



- A1 Run all optimizers k = 20 with budgets of m = 2000 function calls. A problem is regarded solved if the best function value J found by the k satisfies  $J \le J^* + \varepsilon$ .
- A2 Like A1, but consider every of the *k* runs on its own to assess reliability.
- A3 Like A1, but run global optimizers k = 1 time with a budget of m = 40000 function calls.
- A4 Like A1, but Gaussian noise added to every function call.





- overall, MCS and DIRECT performed best
- of the local optimizers, DHC, FMINCON and BOBYQA performed best
- DHC more reliable than BOBYQA on non-smooth problems





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- DFO methods can be an alternative to gradient-based optimization with finite differences
- global optimizers considerably better with a higher budget, in particular MEIGO-ESS
- then, its global search makes MEIGO-ESS superior to simple multi-start





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MEIGO-ESS-DHC

# Performance on biological ODE models

ID	Description	Parameters	Source
M1	conversion reaction	2	-
M2	enzyme-catalyzed reaction	4	-
М3	mRNA transfection	5	Leonhardt et al. 2014
M4	Pom1 gradient formation	7	Hross et al. 2016
M5	Hopf bifurcation	11	Ballnus et al. 2017
M6	JAK-STAT signaling	17	Swameye et al. 2003
M7	RAF-MEK-ERK signaling	28	Fiedler et al. 2016
M8	histone methylation	48	Zheng et al. 2012

• J: negative log-likelihood assuming Gaussian measurement noise

- best optimizers from first study
- FMINCON+g for comparison, i.e. with analytic gradients
- k = 100 starts, function call budgets based on prior experience
- used the toolboxes  $\mathsf{AMICI}^1$  and  $\mathsf{PESTO}^2$

 <sup>1</sup>Fröhlich et al.. Scalable parameter estimation for genome-scale biochemical reaction networks. Plos Computational Biology 2017
 <sup>2</sup>Stapor et al.. PESTO: Parameter EStimation TOolbox. Bioinformatics 2018

## Performance on biological ODE models



	solved problems	
	BOBYQA	5
	DHC	4
	DIRECT	1
	MCS	3
	CMAES	5
	PSWARM	6
MEI	GO-ESS-BOBYQA	4
	MEIGO-ESS-DHC	4
	FMINCON	7
	FMINGON+g	7

- FMINCON, FMINCON+g performed best
- marked contrast to classic test functions: DFO methods only reliable on simple models
- analytic gradients improved convergence considerably

#### Waterfall plots



#### ... analytic gradients improve convergence

• idea: combine global DFO methods with local derivative-based searches



• study: profound comparison of MEIGO-ESS and multi-start<sup>1</sup>

<sup>1</sup>Villaverde et al.. Benchmarking optim. methods for parameter estimation in large kinetic models. 2018 (submitted)

# Conclusion

## Conclusion

- know your problem
- derivative-based optimization superior to DFO for ODE models
- if possible, compute derivatives
- exceptions confirm the rule



Not everything is a nail.

#### github.com/icb-dcm

Schälte, Stapor, Hasenauer. Evaluation of Derivative-Free Optimizers for Parameter Estimation in Systems Biology. FOSBE 2018

- ICB-DCM:
  - Paul Stapor
  - Fabian Fröhlich
  - Daniel Weindl
  - Jan Hasenauer
- IIM-CSIC:
  - Julio Banga
  - Jose Egea
- the toolbox authors
- ... and many more



$$ar{y}(t) = y(t, heta) + arepsilon( heta)$$
 $J( heta) = -\log(p( heta|ar{y})) - \log(p( heta))$ 

- ∇<sub>θ</sub>J can be computed for ODE models using forward (FSA) or adjoint sensitivity analysis (ASA)
- implemented in the toolbox AMICI
- in particular valuable for large-scale models

#### Forward sensitivity analysis for ODE models

State and output sensitivities

$$s^{x}(t, heta):=
abla_{ heta}x(t, heta)\quad s^{y}(t, heta):=
abla_{ heta}y(t, heta)$$

Forward sensitivity equation

$$\begin{aligned} \dot{s}^{x} &= \frac{\partial f}{\partial x} s^{x} + \frac{\partial f}{\partial \theta}, \qquad s^{x}(0,\theta) = \frac{\partial x_{0}}{\partial \theta}(\theta) \\ \dot{s}^{y} &= \frac{\partial g}{\partial x} s^{x} + \frac{\partial g}{\partial \theta} \end{aligned}$$

Augment and solve ODE of dim  $n_x(1 + n_\theta)$ 

#### Adjoint sensitivity analysis for ODE models

1. Calculation of state via forward simulation

$$\dot{x}(t,\theta) = f(x(t,\theta),\theta)$$
  
 $\dot{y}(t,\theta) = g(x(t,\theta),\theta)$ 

2. Calculation of **adjoint state**  $p(t) \in \mathbb{R}^{n_{\chi}}$  as solution to backward ODE

$$\begin{split} &\lim_{t \to t_N^+} p(t) = 0\\ &\text{for } j = n_t : -1 : 1\\ &\dot{p}(t) = -\frac{\partial f}{\partial x} |_{x(t,\theta),\theta}^T p(t), \quad t \in (t_{j-t}, t_j)\\ &\text{with } p(t_j) = \lim_{t \to t_j^+} + \sum_{k=1}^{n_y} \frac{\partial g_k}{\partial x} |_{x(t_j,\theta),\theta} \frac{\bar{y}_{jk} - y_k(t_j,\theta)}{\sigma_{jk}^2} \end{split}$$

3. Calculation of gradient via 1-dim integral

$$\nabla_{\theta} J(\theta) = -\int_{0}^{T_{N}} p(t)^{T} \frac{\partial f}{\partial \theta}|_{x(t,\theta),\theta} dt - p(t_{0})^{T} \frac{\partial x_{0}}{\partial \theta}|_{\theta} - \sum_{j,k} \frac{\partial g_{k}}{\partial \theta}|_{x(t_{j},\theta),\theta}^{T} \frac{\tilde{y}_{jk} - y_{k}(t_{j},\theta)}{\sigma_{jk}^{2}}$$