Simulation-based Test Functions for Optimization Algorithms GECCO 2017, GECH Track

> Martin Zaefferer, Andreas Fischbach, Boris Naujoks, Thomas Bartz-Beielstein

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Technology Arts Sciences TH Köln





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4 Summary and Outlook

We need test functions to ...

- analyze
- understand
- compare
- design / configure / tune

... algorithms

Test functions should be (sufficiently) ...



Simulation-based Test Functions

Existing Approach: Model-based

1 Collect data from real-world problem

2 Learn structure via model (e.g., Kriging / Gaussian processes)

3 Vary model to generate problem instances

4 Use estimation / prediction as test functions

Model-based Test Functions

Advantages

- Relevance due to real-world data
- Data more easily accessible
- Nonlinear models yield flexibility
- Variation yields diversity

Disadvantages

- Bias due to data and model
- Unknown characteristics
- Method for variation?
- and ...

Problem

- Most predictors are smoothing
- Desirable, e.g., for surrogate model-based optimization
- Undesirable for test function generation
- Too easy
- \rightarrow requirement: non-smoothing



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Proposed Remedy

- Simulation e.g., $\hat{\mathbf{y}}_{nc} = \mathbf{1}\hat{\mu} + C_s^{1/2} \boldsymbol{\epsilon}$ instead of estimation e.g., $\hat{y}(x) = \hat{\mu} + \mathbf{k}^T \mathbf{K}^{-1} (\mathbf{y} - \mathbf{1}\hat{\mu})$
- Goal of *estimation*:
 - Close to the "true" value
- Goal of *simulation*:
 - Close to the "true" process behavior
- Potentially conflicting!

Simulation Example: Undersea Cable



- data
 - —— estimation
- - conditional simulation

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Simulation-based Test Functions:

Advantages

- Avoid / reduce smoothing
- Reproduces *behavior* of the real-world process
- *Conditional* simulation can reproduce the training data
- Principled approach to generate diverse instances

Disadvantages

- Between simulated samples: interpolation (smoothing, less problematic than with estimation)
- Required number of simulated samples unknown
- Complexity in case of large number of samples



- Training data: 6 samples
- Model simulated at m = 100 samples

Example code available at:

https://martinzaefferer.de

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Case Study: Protein Sequence Optimization

- Freely available data set
- All DNA sequences of length 10 ACGTAACGGT, CGTAAGATTC, ...
- Objective/Fitness: maximize affinity to fluorescent protein (APC)
- Kriging model trained with 100 sequences
- Simulated for m = 1000 sequences

Results: Landscape Analysis

- Correlation length
 - True: 4.5
 - Model: 4.48
- Fitness distance correlation
 - True: -0.32
 - Model: -0.37 (+/- 0.09)
- Number of local optima
 - True: 6805
 - Model: \leq 49



Results: Landscape Analysis

- Interpolating m = 1000 samples yields too much smoothness
 - but: better than interpolating the 100 training samples
- Larger m required, computational issues
- Workaround:
 - Restrict to subspace: fix end of sequences to ACGTA
 - Simulate all sequences in the subspace
 - True: 16 local optima
 - Estimation: 2 local optima
 - Simulation: 10 19 local optima 🗸

Performance: True vs. Estimation



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Performance: True vs. Simulation



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Take Home Message

- Model-based test functions can produce difficult, diverse, flexible, relevant and cheap to evaluate test functions
- Use (conditional) simulation not estimation
- If performance on similar problems is of interest: simulation
- If performance on **potential realizations of the same problem** is of interest: conditional simulation

Open Issues

• How many simulated samples (*m*) are needed for a certain problem?

• What to do if m grows too large?

Thanks for Listening

• Questions? Remarks?

PS: You can find the employed modeling tools in the package CEGO on CRAN: https://cran.r-project.org/package=CEGO. Check the earlier described 1-dimensional example to see how it works.