

An Introduction to Scientific Experimentation and Benchmarking

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Full set of slides: http://www.cmap.polytechnique.fr/~nikolaus.hansen/gecco-2023-benchmarking-tutorial.pdf

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Please follow this link to access the last version of the slides : <u>http://www.cmap.polytechnique.fr/~nikolaus.hansen/gecco-2023-benchmarking-tutorial.pdf</u>

... feel free to ask questions...





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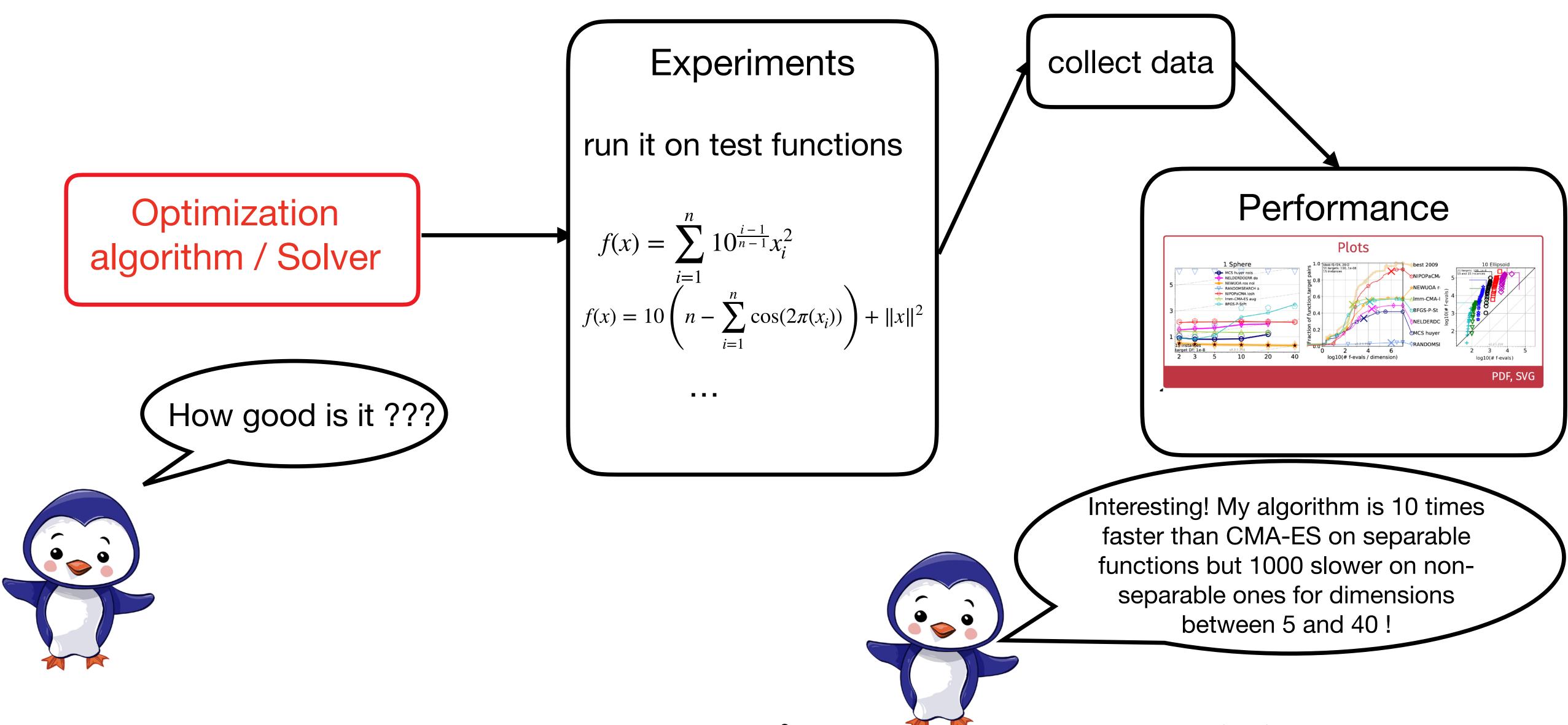
Scientific Experimentation

- Numerical experiment to answer some (scientific) questions
 - test/validate an hypothesis
 - understand a mechanism
- Often done when designing a new algorithm, but not only

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Benchmarking



Overview

- I Scientific experimentation
 - demo in Python
- II Benchmarking

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The first principle is not to fool yourself – and you are the easiest person to fool.

Richard Feynman



Why Experimentation?

- The behaviour of many if not most practically relevant algorithms is
 - problems

 \mathcal{O} -results are not sufficient in practice often, complex algorithms are truthfully represented only by source code implementation details matter (at least sometimes) hence we need an *alternative to theory* for investigation

even on simple problems (the algorithm is the only source of complexity) comprehension is the main driving force for scientific progress If it disagrees with experiment, it's wrong. [...] And that simple statement is the key to science. — R. Feynman

Virtually all algorithms have parameters

• We are interested in solving *black-box* optimisation problems

not amenable to a (full) theoretical analysis even when applied to simple

not comprehensible or predictable without (extensive) empirical examinations

like most (physical/biological/...) models in science we rarely have explicit knowledge about the "right" choice prevent overfitting is a particular challenge this is a *big* obstacle in designing and benchmarking algorithms

which may be "arbitrarily" complex and (by definition) not well-understood



Why is Experimentation Important?

"In the course of your work, you will from time to time encounter the situation where the facts and the theory do not coincide. In such circumstances [...], it is my earnest advice to respect the facts."

"If it disagrees with experiment, it's wrong. And that simple statement is the key to science. [...] That's all there is to it."

— Igor Sikorsky

— Richard P. Feynman https://youtu.be/b240PGCMwV0



Scientific Experimentation in a Nutshell

- start from a (good) scientific question
- design an experiment to test (try to "falsify"/"rule out"/render unlikely) one or several answers
- visualize (all) what make sense and interpret
- iterate ...

creativity and technique, comparable to the arts.

Effective scientific experimentation requires a healthy mixture of



Scientific Experimentation (dos and don'ts)

- with little ambiguity

to develop a mental model of the object of interest quantitative in a) effect size and b) error probability of the prediction to practice to make the clear distinction between a guess from intuition and observations to practice being proven wrong by observations and overcoming *confirmation bias*

reasons

check/test "everything" yourself, practice stress testing (e.g. a weird parameter setting) which (also) boosts understanding this is a key element for success interpreted/scripted languages have an advantage (quick test of code snippets) Why Most Published Research Findings Are False [loannidis 2005]

experimentation (see demonstration)

What is the aim? Answer a question, ideally quickly (minutes, seconds) and

consider in advance what the question is and in which way the/an experiment could answer the question

practice to make *quantitative* predictions of the possible/expected outcomes

do not (blindly) trust in code, claims, ... that you rely upon without good

run *rather many than fewer* (different) experiments *iteratively*, practice *online*

to run many experiments they must be quick to implement and run, ideally seconds rather than minutes (start with small dimension/budget) develops a sense for the effect of setup changes



Scientific Experimentation (dos and don'ts)

run any experiment at least twice

assuming that the outcome is stochastic or with a different initialization thereby getting an estimator of variation/dispersion/variance

display: the more the better, the more polished the better

figures are *intuition pumps* (not only for presentations or publications) it is hard to overestimate the value of a good figure data is the only way experimentation can help to answer questions, therefore look at the data, stare at them, study them carefully!

don't make minimising CPU-time a primary objective

avoid spending time in implementation *details* to tweak performance prioritize code clarity (minimize time to *change* code, to debug code, to maintain code) yet code optimization may be necessary to run experiments efficiently

Testing Heuristics: We Have it All Wrong [Hooker 1995] "The emphasis on competition is fundamentally anti-intellectual and does not build the sort of insight that in the long run is conducive to more effective algorithms"



Scientific Experimentation (dos and don'ts)

- It is usually (much) more important to understand why algorithm A unknown, unclear or trivial reasons)
- Remain aware: many devils are in the details, results or their or bugs

yet another reason to run many (slightly) different experiments check limit settings to give consistent results

Invariance is a very powerful, almost indispensable tool

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performs badly on function f, than to make algorithm A faster (for

mainly because an algorithm is applied to *unknown* functions, not to *f*, and the "why" allows to predict the effect of design decisions

interpretation may crucially depend on (simple or intricate) subtleties



Scientific Questions and Objectives (bad and good)

algorithm developed for 20 years (e.g. CMA-ES) in the domain where it excels (unconstrained, difficult problems)

likely to be very difficult, not very specific on the class of problems (where do we start?), given the algorithm framework (comparison-based, derivative-free) isn't the algorithm already close to optimal performance?

functions

not very specific: which functions do we want to solve better ? Why do we expect to solve problems better than CMA-ES with restarts? How can the algorithm scale with the number of local optima? why should it help to maintain several modes to approach a global optimum?

- a few functions with few local optima
- a paper using that in our domain as it will look good and cool

Invent a new algorithm that is better than a well-established one, say outperforms an

Have a CMA-ES version with a mixture of Gaussians to perform well on multimodal

what people do afterwards: do not compare with baseline (algorithm with restart), test on

• This new concept used in physics (or ...) is fancy, many famous people use it, I want to write



Scientific Questions and Objectives (bad and good)

- I want to understand why the algorithm fails in solving this problem
- problem with sparse structure
- class of problems

specific, easy to design an experiment that will give a clear answer

•

specific, leads to subquestions like what is the difficulty of the problem, can another method solve it ...

understanding why it fails is often the first step to solve the problem

I want to design a large-scale CMA-ES variant scaling linearly that solves

not widely explored, possible to identify classes of problems to experiment with

I want to know how this algorithm scales with the dimension to solve this

How do I add errors bars that are meaningful when plotting ECDF graphs? specific, unexplored, useful...



Desirable properties of scientific questions

What are good scientific questions / objectives:

- its optimal performance)
- realistic
- not too general / specific enough
- parts can be answered (quickly)
- related to an unexplored subdomain •
- not motivated by fashion
- relate to improve methods where it is needed

question/objective should be solvable (e.g., an algorithm is *not* already close enough to

have a mechanism why it could succeed / be answered (before to start experimentation)



Demo on scientific experimentation



What did we illustrate - Take home messages

- Interpretation of data can be difficult •
- trying various ideas may be necessary



Overview

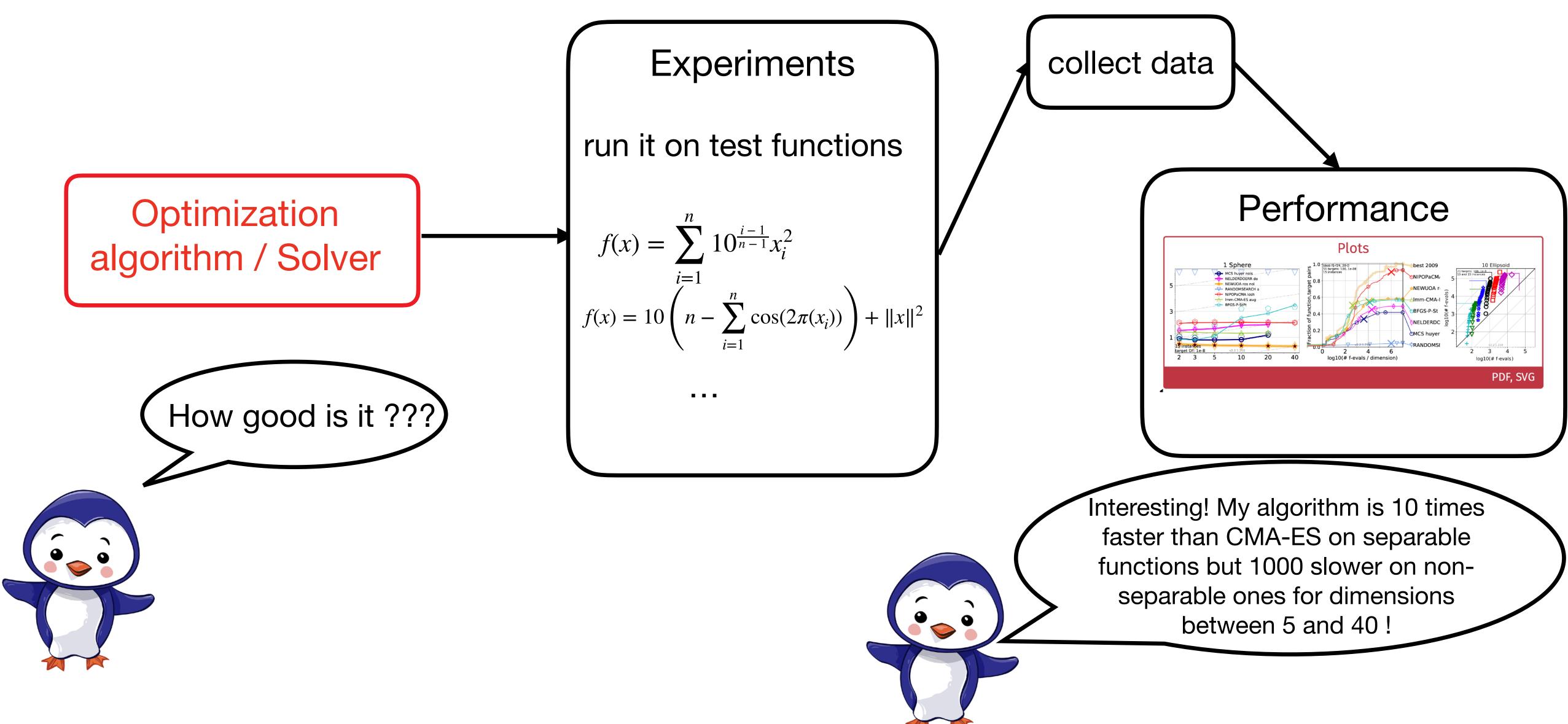
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Benchmarking



About Generalization

Does benchmarking make sense at all?

NFL theorems hold on sets of functions that are "closed under permutation". Whether all functions in such set are (equally often) observed in reality is an empirical question. Practical evidence suggests: some algorithms are vastly worse than others.

• The function or instance ID can not be input to the algorithm

We shall not set algorithm parameters depending on each function! AKA overfitting. The benchmarking setup: an algorithm that needs to repeatedly solve "new" problems or instances. possible but not recommended: Crafting Effort correction for using different parameter settings on different functions¹.

- Comparable data

1: Price KV. Differential evolution vs. the functions of the 2nd ICEO. In Proceedings of 1997 IEEE International Conference on Evolutionary Computation (ICEC'97) 1997 (pp. 153-157). IEEE. 21 Anne Auger and Nikolaus Hansen, Inria, IP Paris

After all there is no free lunch, right? Or is there?

 A benchmark should attempt to model commonplace and relevant "real-world" optimization problems The set of all observable optimization problems is WAY smaller than most sets of mathematically constructible problems.

• Invariance of algorithms is a relevant aspect to interpret (generalizability of) benchmarking results

depends on the benchmarking setup across publications across functions (e.g. speedup factor)



What about Competitions?

"The emphasis on competition is fundamentally anti-intellectual and does not build the sort of insight that in the long run is conducive to more effective algorithms". Hooker (1995) distinguishes "scientific testing" from "competitive testing"

in Testing Heuristics: We Have it All Wrong.



Miscellaneous

- A trivial (serial) algorithm portfolio: K algorithms can solve each and
- What differences are we interested in?
- Function/problem instances
- Search domain: discrete and continuous

every problem as fast as the fastest of these algorithms multiplied by K. Run in parallel, they become as fast as the fastest algorithm

2%, 20%, 200%, 200%,...

versus different functions

Examples come from the continuous domain.

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(Specific) Goals of Benchmarking

way

Specific goals can be:

- 1. Comparing against the "state-of-the-art" or against a baseline
- Understanding algorithms 2.
- 3. Predict performance on real-world problems
- Selecting algorithms to solve a given problem 4.
- Regression testing after changes of an algorithm or an implementation 5.
- Running a competition 6.

a competition setup needs to hide information from the competitor/experimenter

"Everybody" has to do it and it is tedious: choosing (and implementing) problems, performance measures, visualization, statistical tests, ...

We may think of benchmarking as measuring algorithm performance in a systematic and standardized

thereby creating a performance "profile" of an algorithm for a standardized assessment and for simplified comparison

any comparison between two or more algorithms

benchmarking usually raises questions, dedicated experimentation is often necessary to answer them

which suggests to consider using tools

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Benchmarking How To: The Global Picture

Two surprisingly (but not completely) independent components:

- the algorithms on?
- How to assess performance?
 - experimental setup
 - data collection
 - measures used and presented

Which benchmark, which suite of functions/problems do we run

For example and in particular, which collection of test problems?



COCO/BBOB: The Global Picture

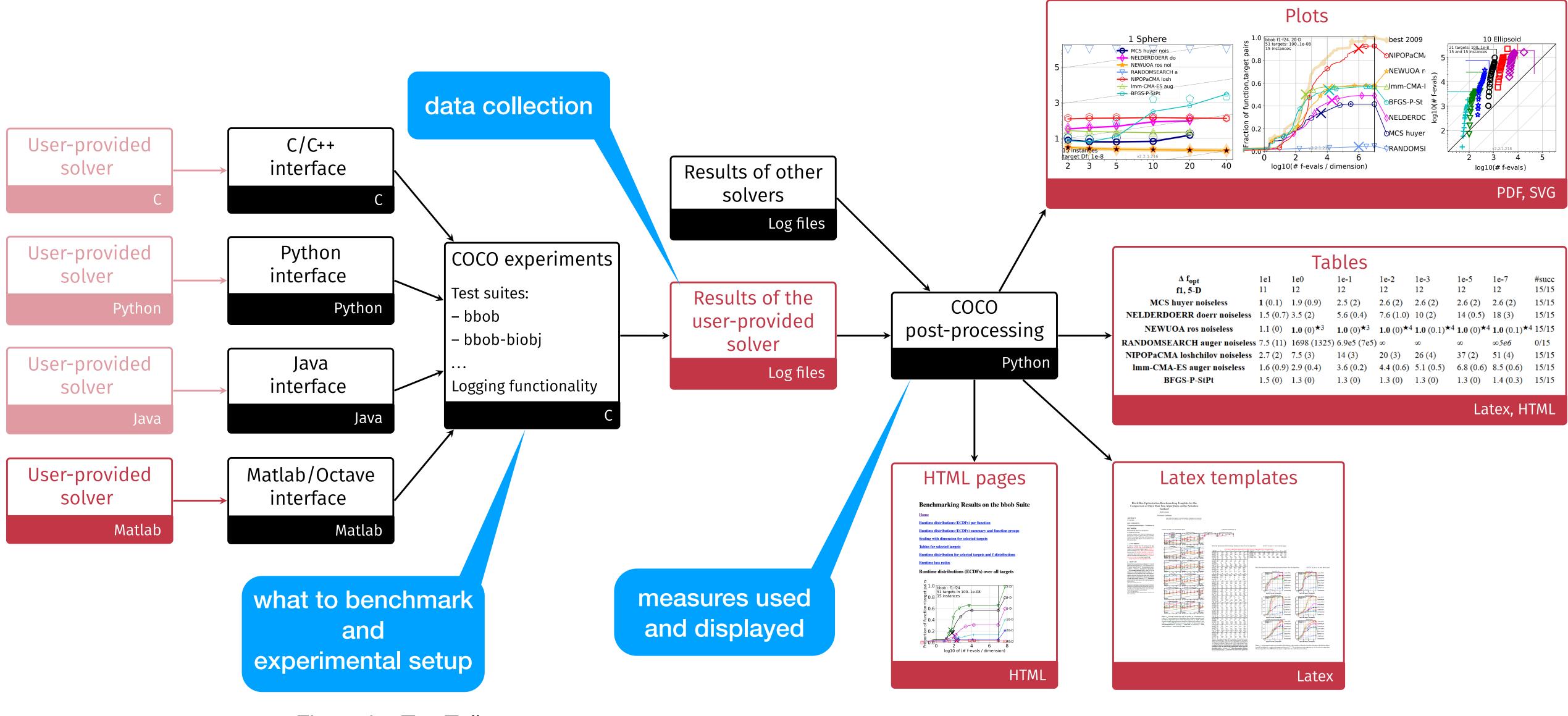


Figure by Tea Tušar in Hansen et al (2021), COCO: A platform for comparing continuous optimizers in a black-box setting. Optimization Methods and Software, 36(1), 114-174-duction to Scientific Experimentation and Benchmarking



What is the Benchmark?

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Choice of Test Problems



What to Benchmark?

- Taking all possible functions from a repository?
- Bad idea if •
 - function difficulties are unbalanced
 - and performance is aggregated
- Leads to bias in the performance assessment

Furious activity is no substitute for understanding (H.H. Williams)

too many small dimensional problems, convex problems...



What to Benchmark?

- test functions should be representative of difficulties we want to test
- related to real-word difficulties
- scalable •

- comprehensible but not too easy
- we should still hide properties from the solver (hide optimum, \dots)

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therefore NFL has no relevance as assumption of being closed under permutation has no relevance wrt real world problems

for performance to be generalizable to RW

dimension plays a big role in performance curse of dimensionality

BB optimization does not mean BB benchmarking

solvers should not be able to exploit the benchmark intentionally or not



Experimental Setup

- should allow as many algorithm types/interfaces as possible
- defines the information an algorithm is allowed to use

- pure repetitions only work for randomized algorithms
- may define a budget (or not)

bounded, unbounded, different input options, deterministic, randomized,...

search domain (and hence dimension), initial solution, regions of "interest", function as back-box not: function name/ID

should define what is recorded to afterwards measure performance

anytime vs targeted budget



Handling and Displaying Empirical Data



Displaying (Performance) Results

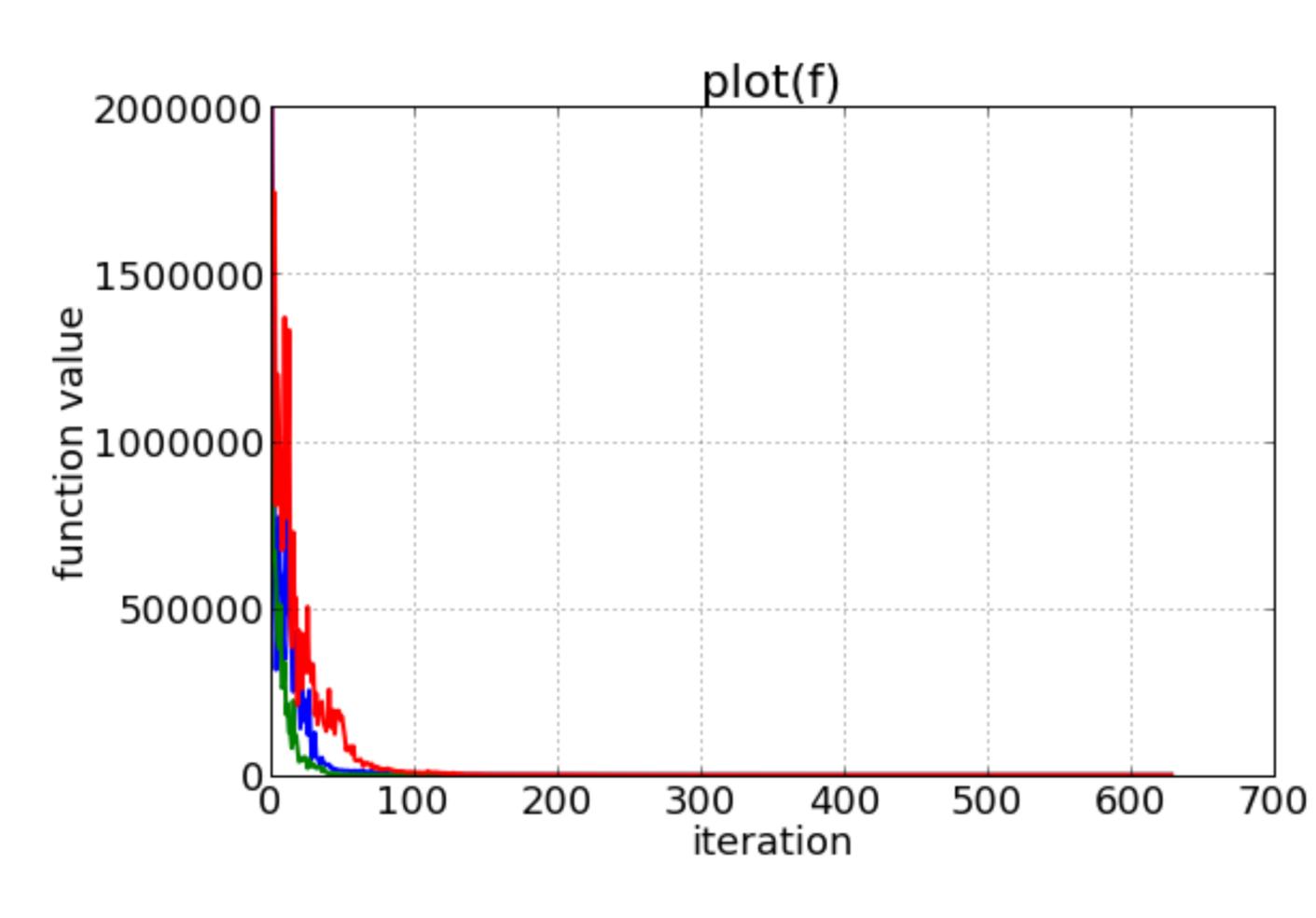
Empirically

convergence graphs is all we have to start with

the right presentation is important!



Displaying Three Runs

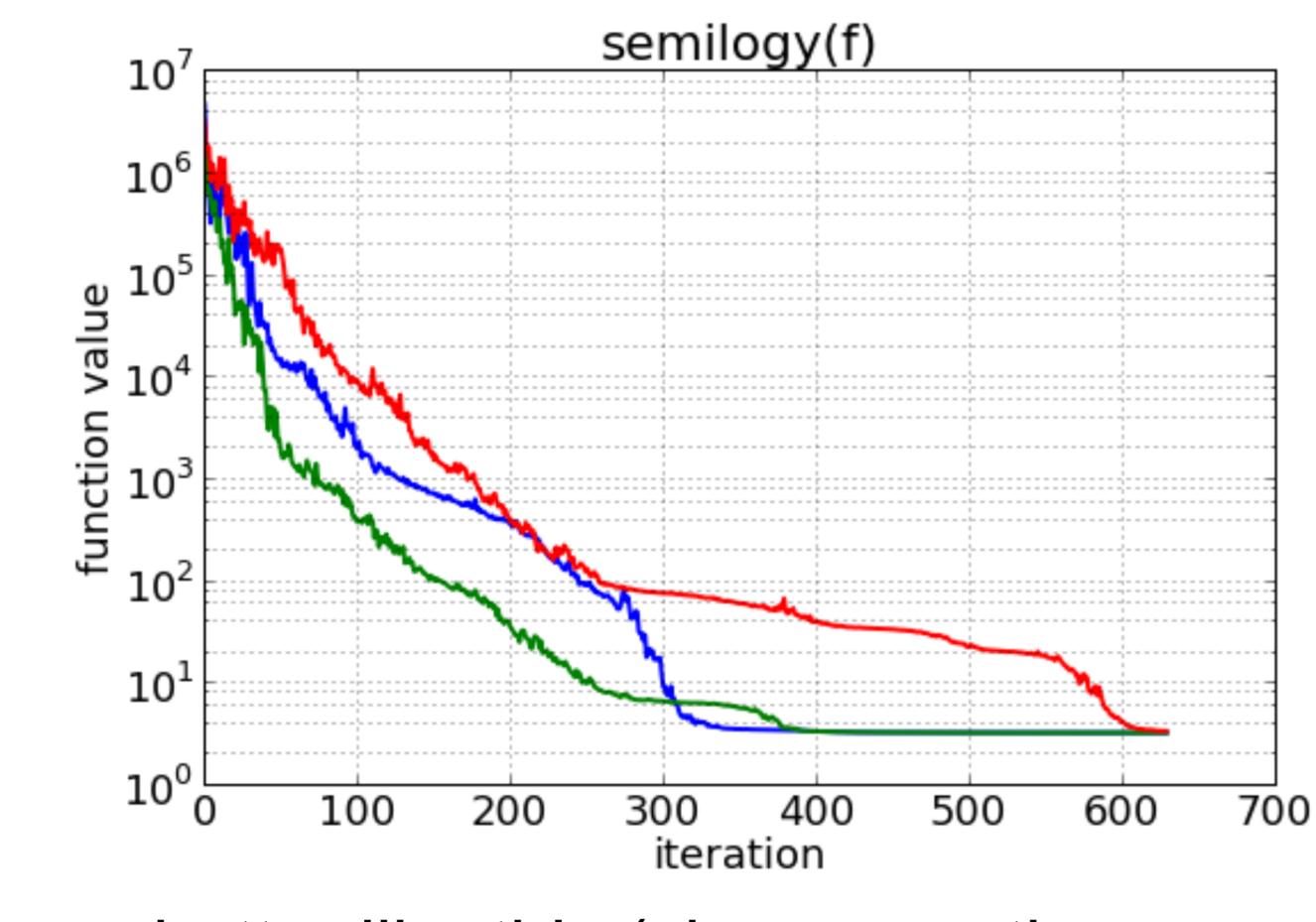


not like this (it's unfortunately not an extremely uncommon picture)

why not, what's wrong with it?

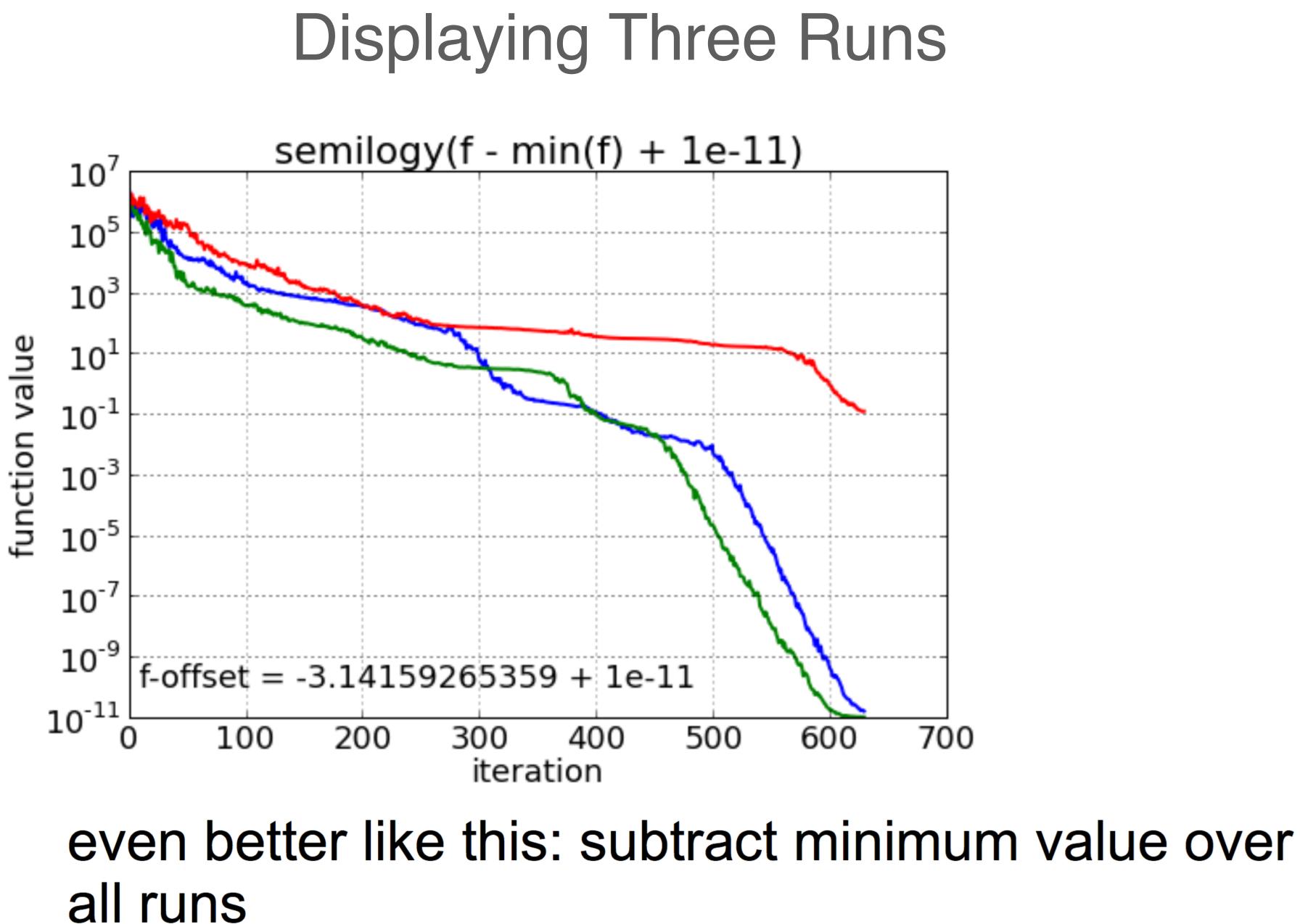


Displaying Three Runs



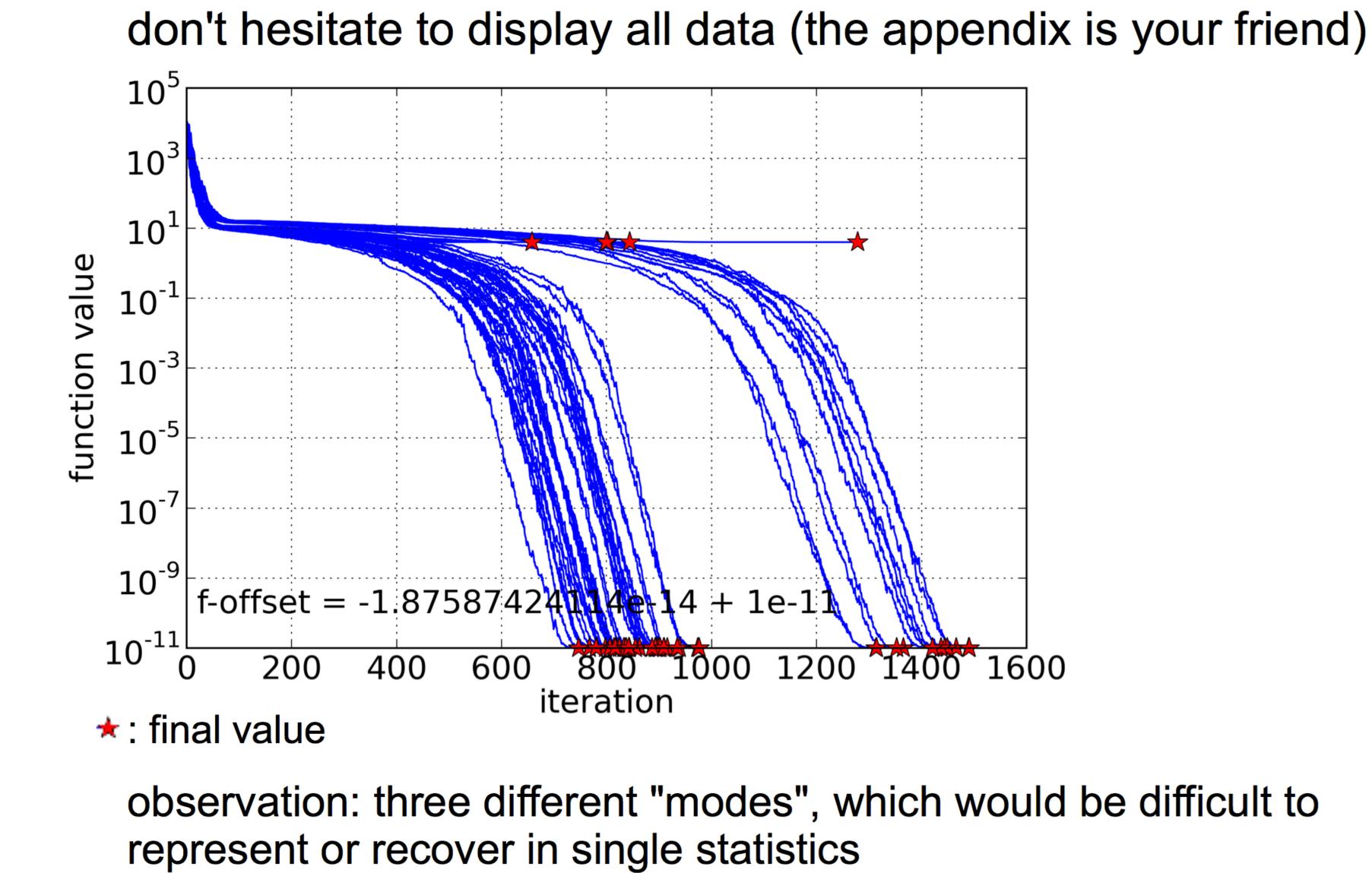
better like this (shown are the same data),







Displaying 51 Runs

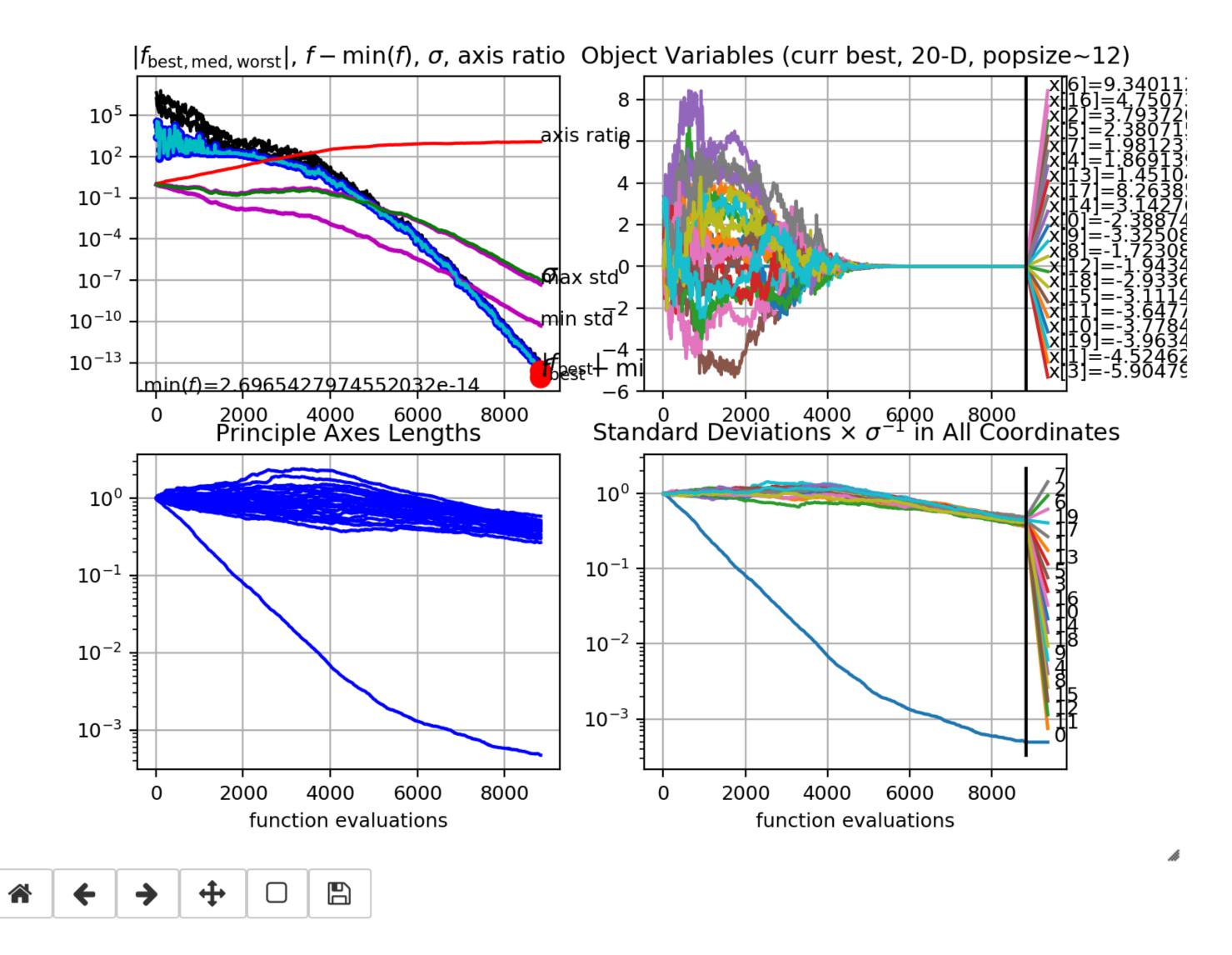




4.04686177e-08

cma.plot()

There is more to display than convergence graphs

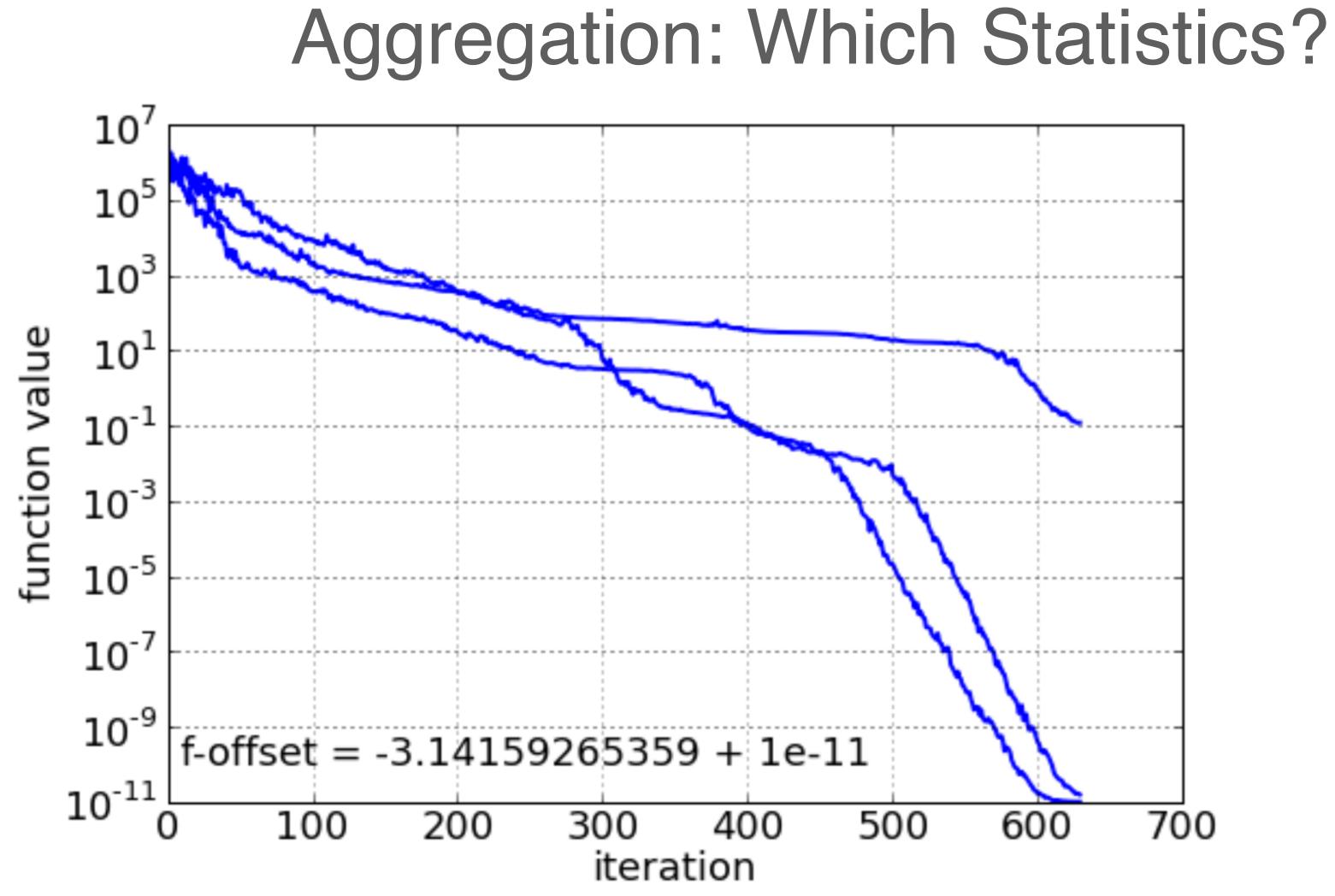


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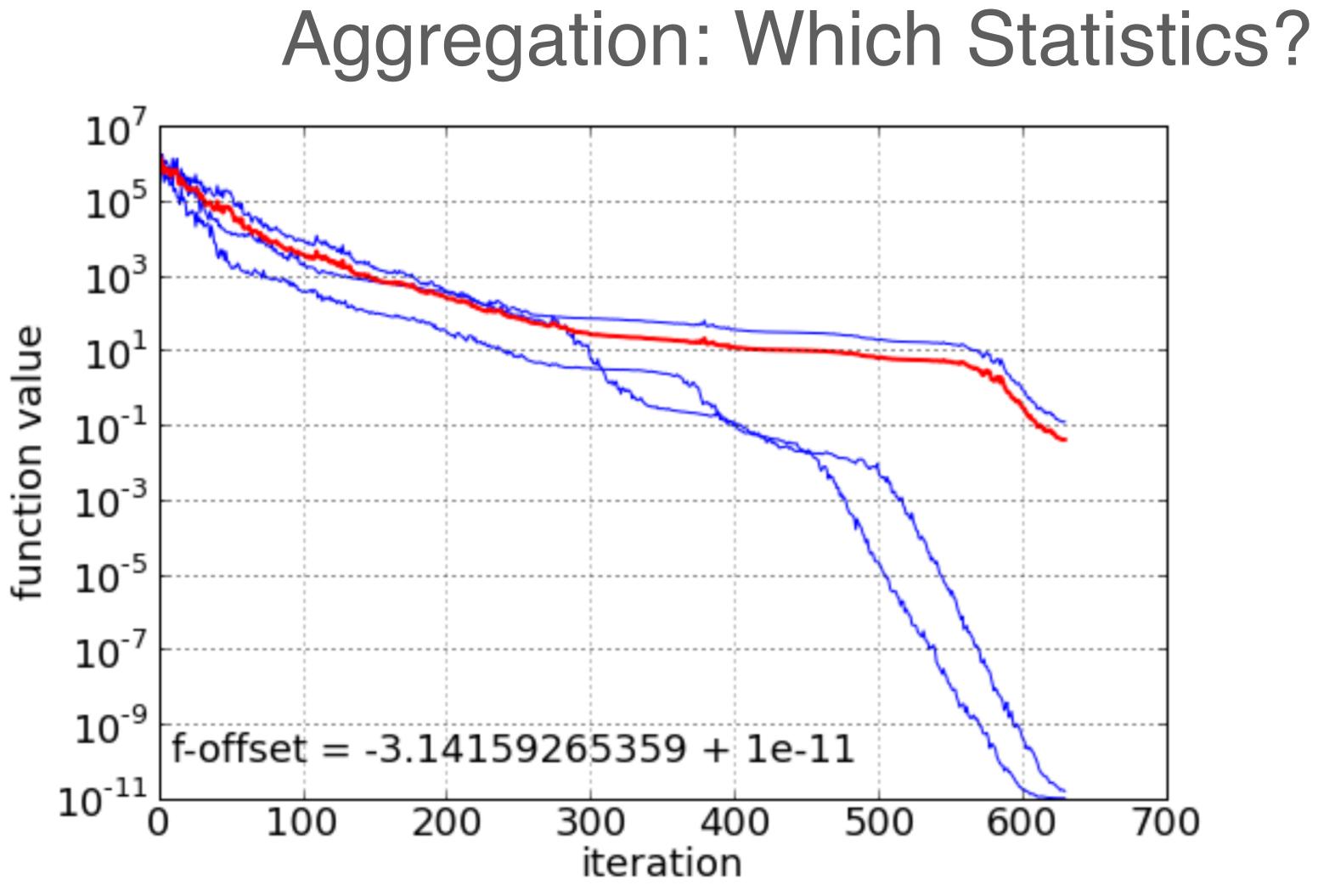
Figure 328





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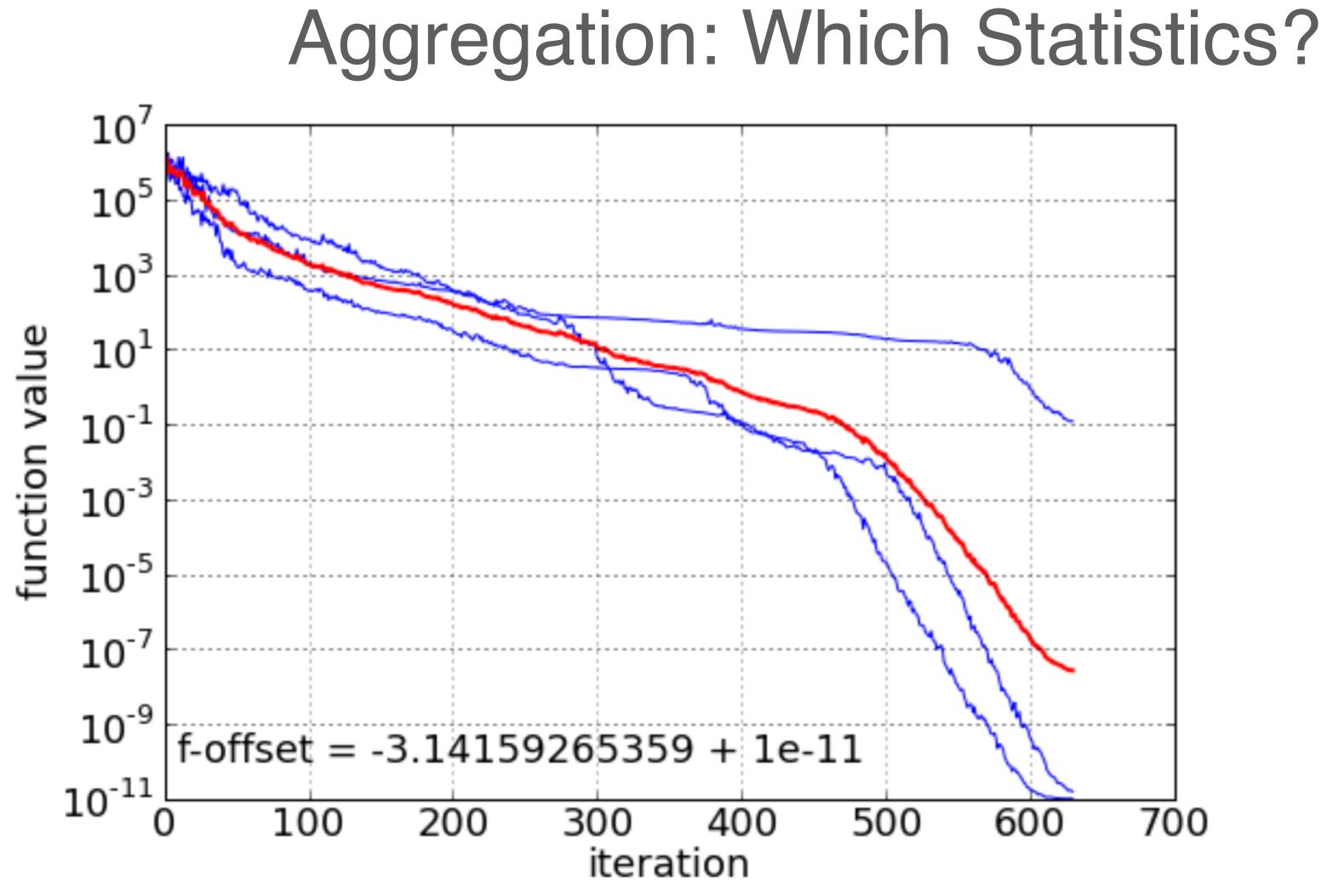




mean/average function value

tends to emphasize large values

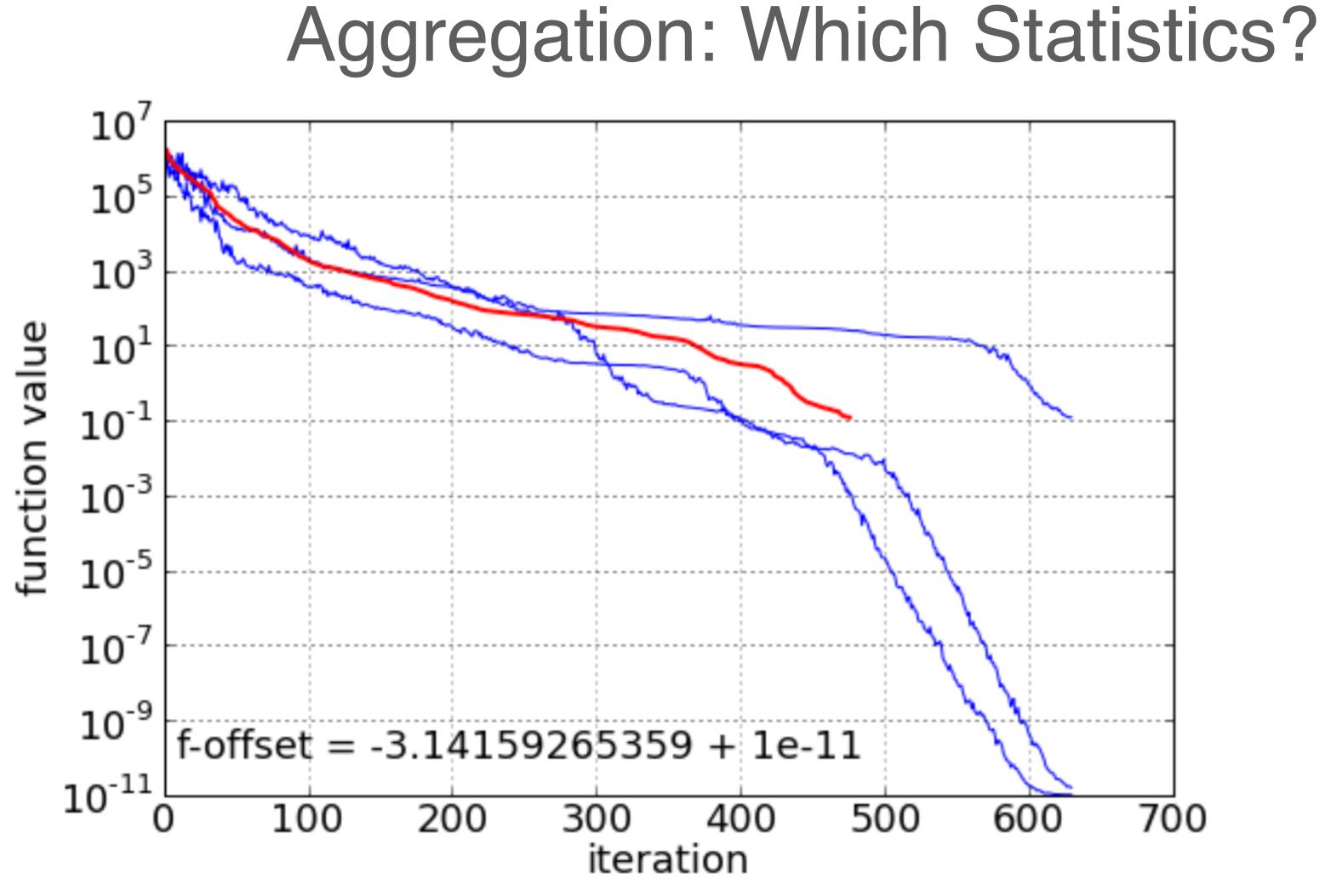




geometric average function value $exp(mean_i(log(f_i)))$:

- reflects "visual" average
- depends on offset

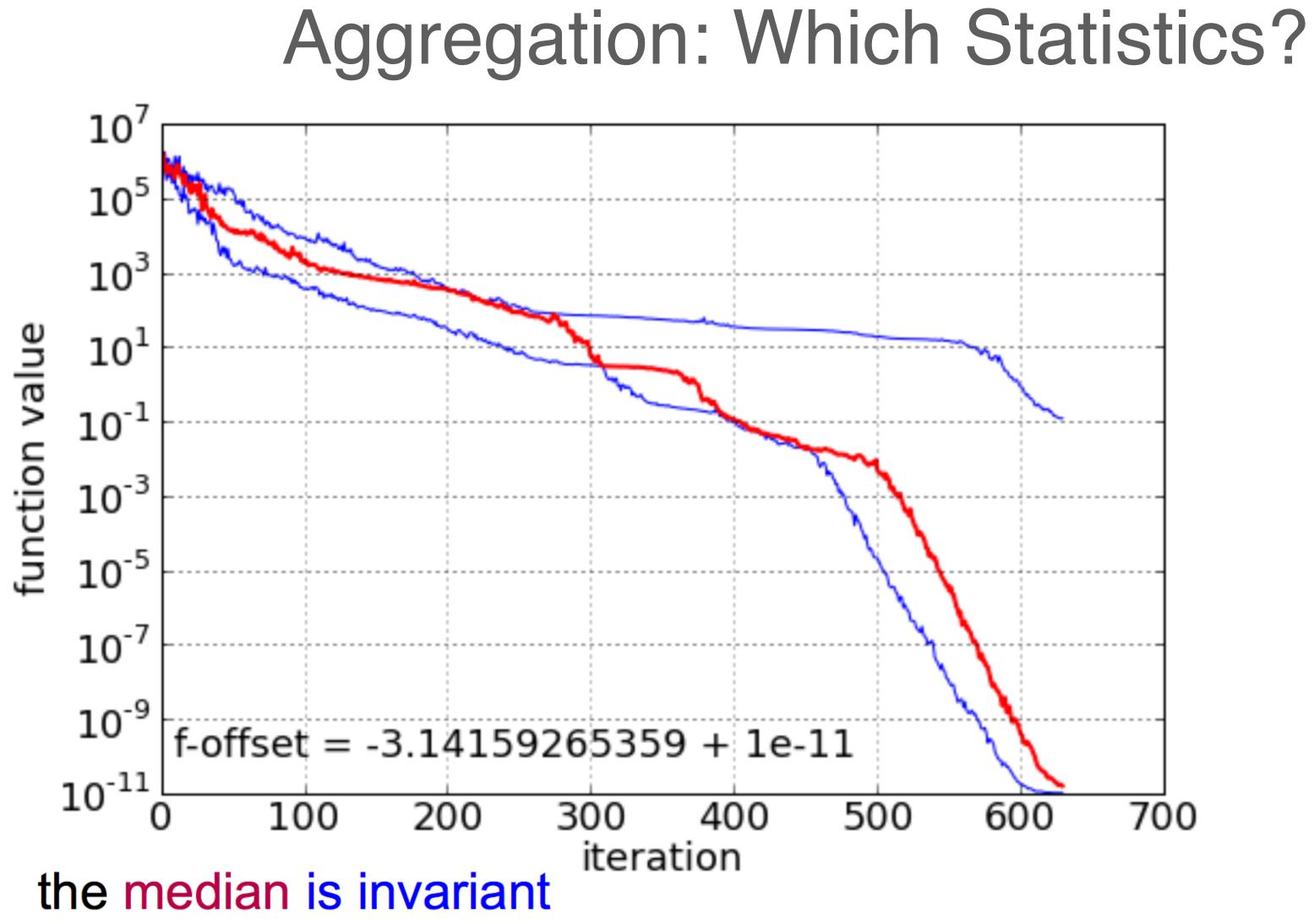




average iterations

- reflects "visual" average
- here: incomplete





- unique for uneven number of data
- independent of log-scale, offset...
- Anne Auger and Nikolau:
- same when taken over x- or y-direction

median(log(data))=log(median(data))

tion and Benchmarking



Implications

preferably, use the median as summary datum

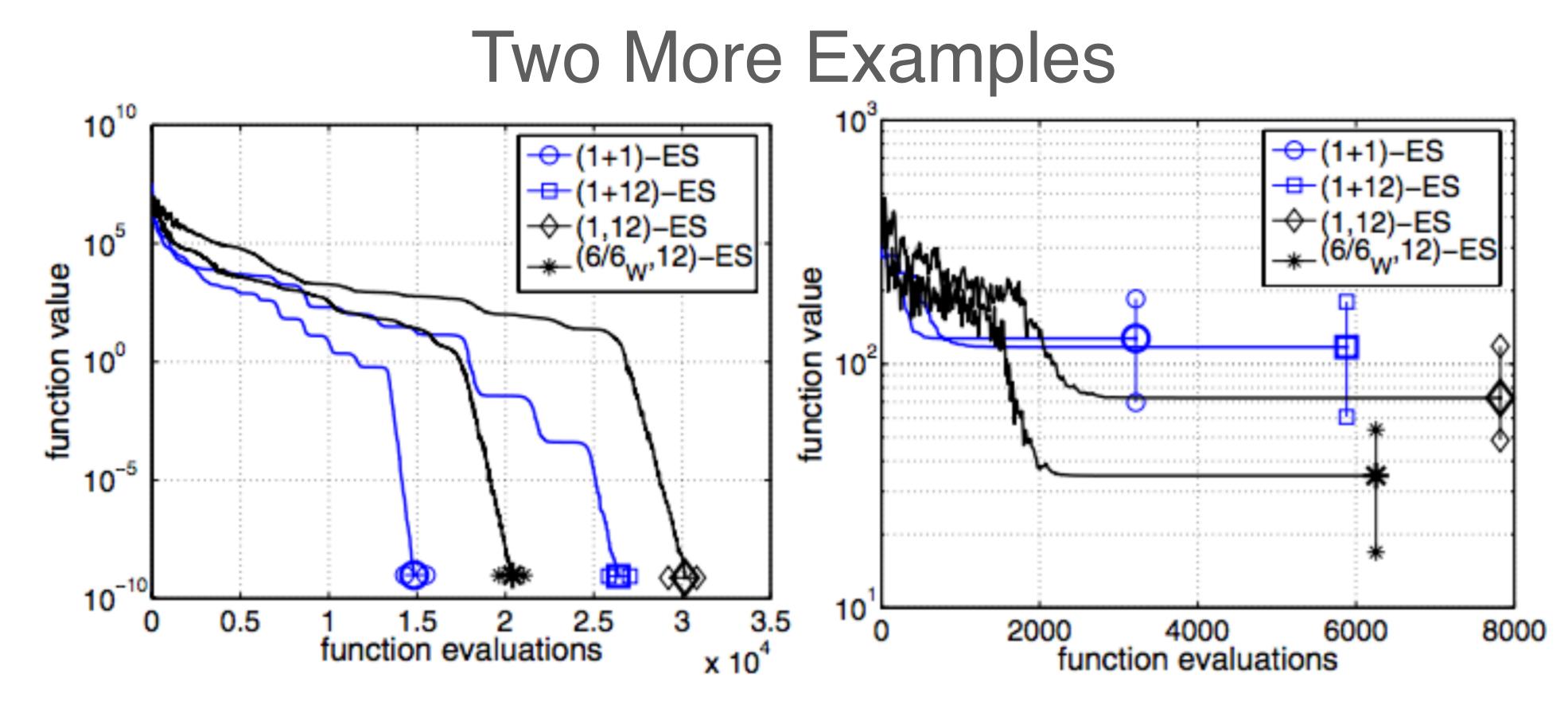
more general: use quantiles as summary data

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unless there are good reasons for a different statistics out of practicality: use an odd number of repetitions

for example out of 15 data: 2nd, 8th, and 14th value represent the 10%, 50%, and 90%-tile





Comparison of 4 algorithms using the "median run" and the 90% central range of the final value on two different functions (Ellipsoid and Rastrigin)

caveat: this range display with simple error bars fails, if, e.g., 30% of all runs "converge"



Cumulative Distribution Function (CDF)

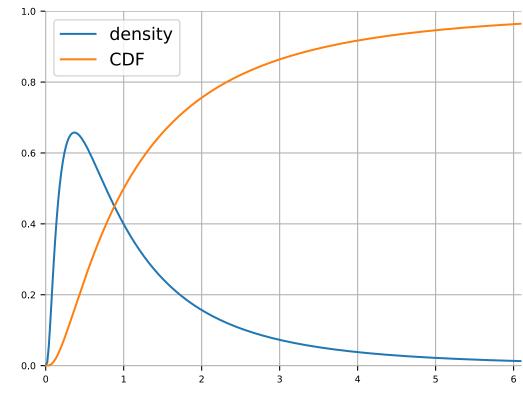
(CDF) is defined as

It characterizes the probability distribution of T

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Given a random variable T, the cumulative distribution function

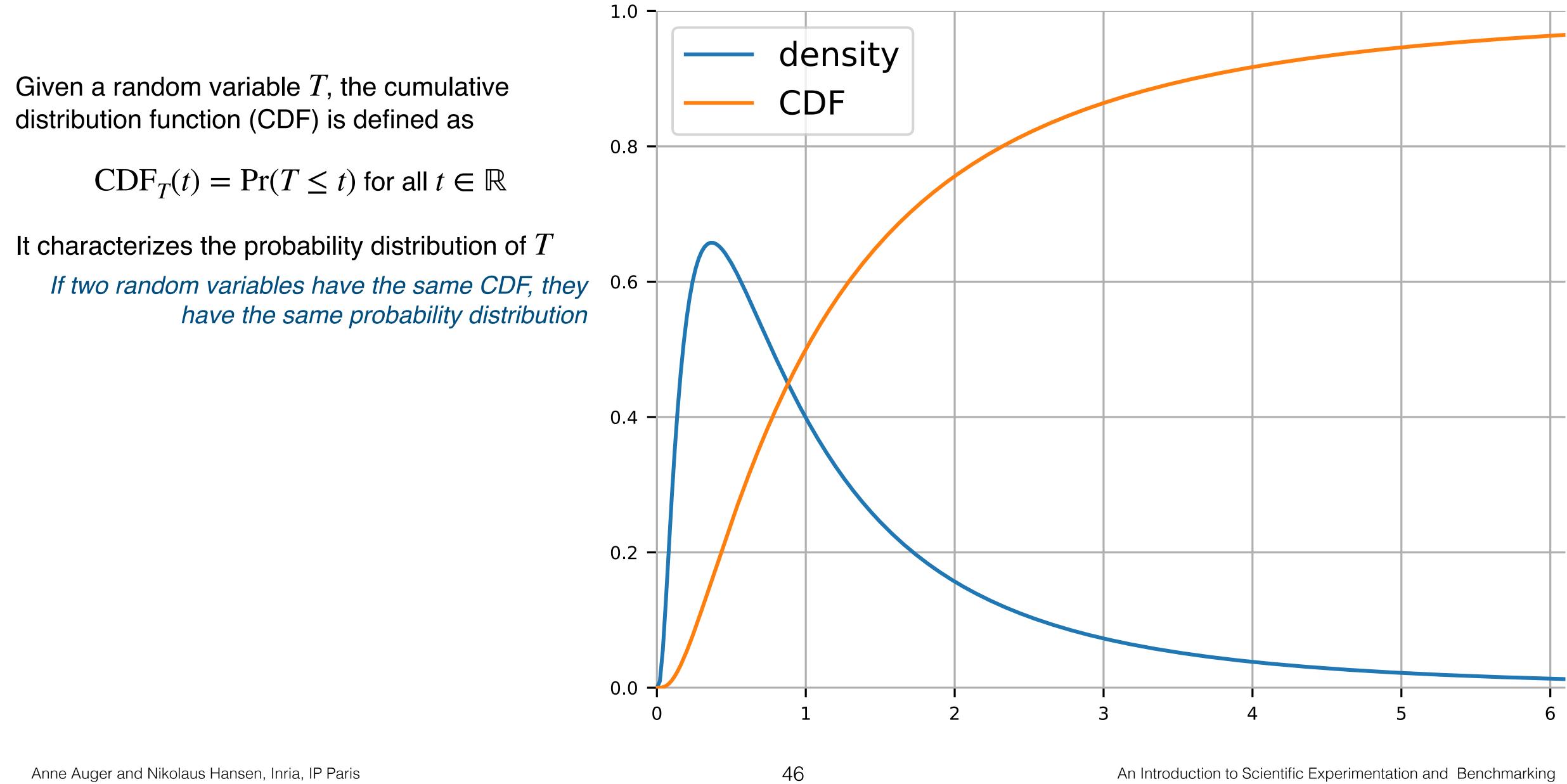
- $\text{CDF}_T(t) = \Pr(T \le t) \text{ for all } t \in \mathbb{R}$



If two random variables have the same CDF, they have the same probability distribution

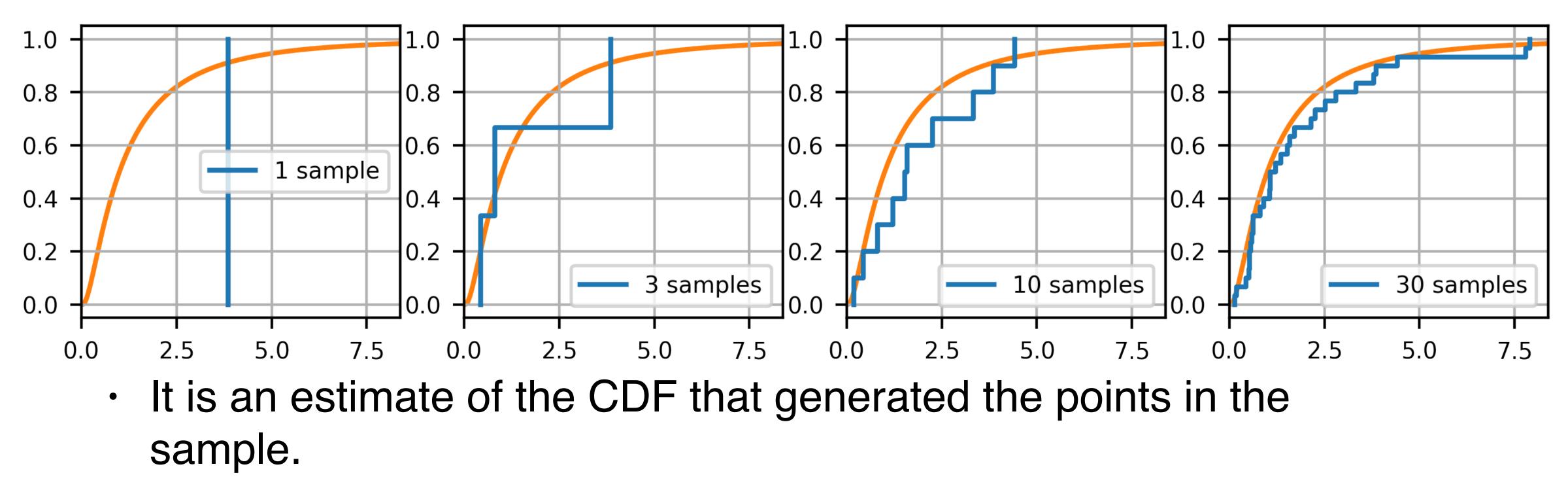


Cumulative Distribution Function (CDF)



Empirical Cumulative Distribution Function

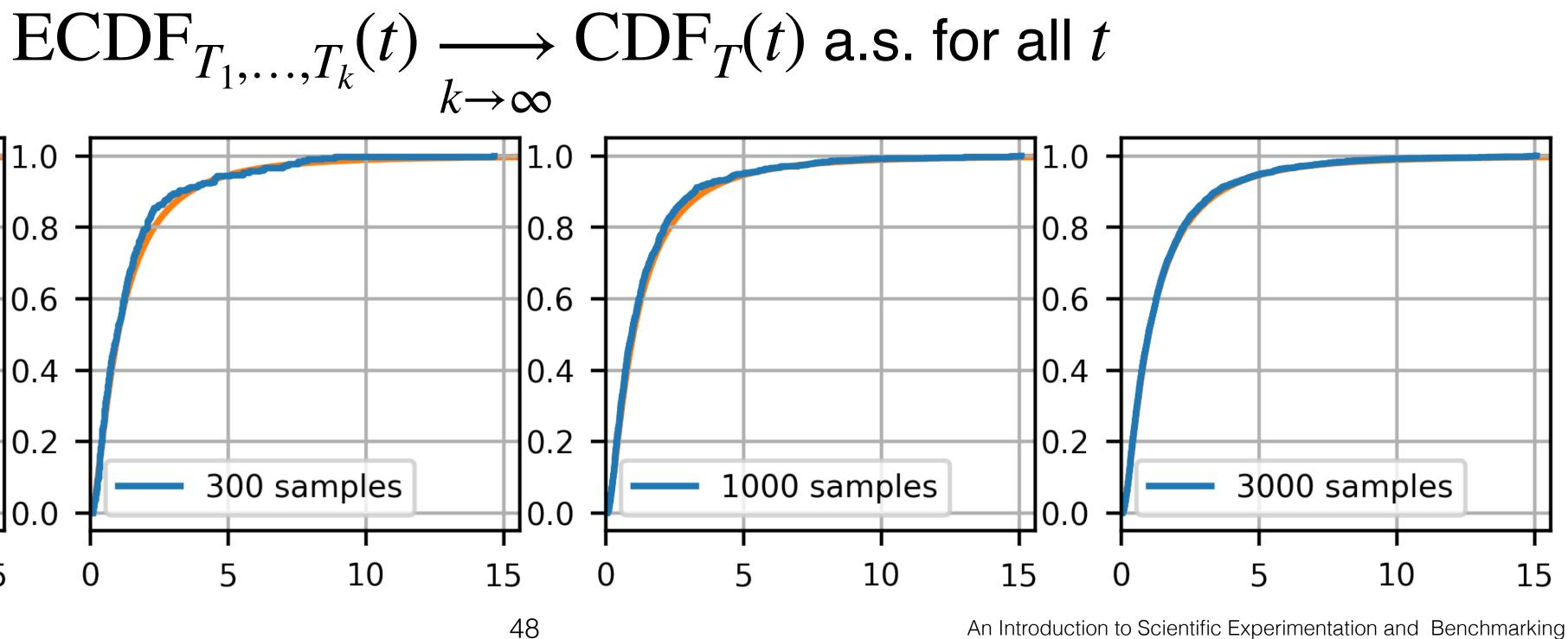
data.

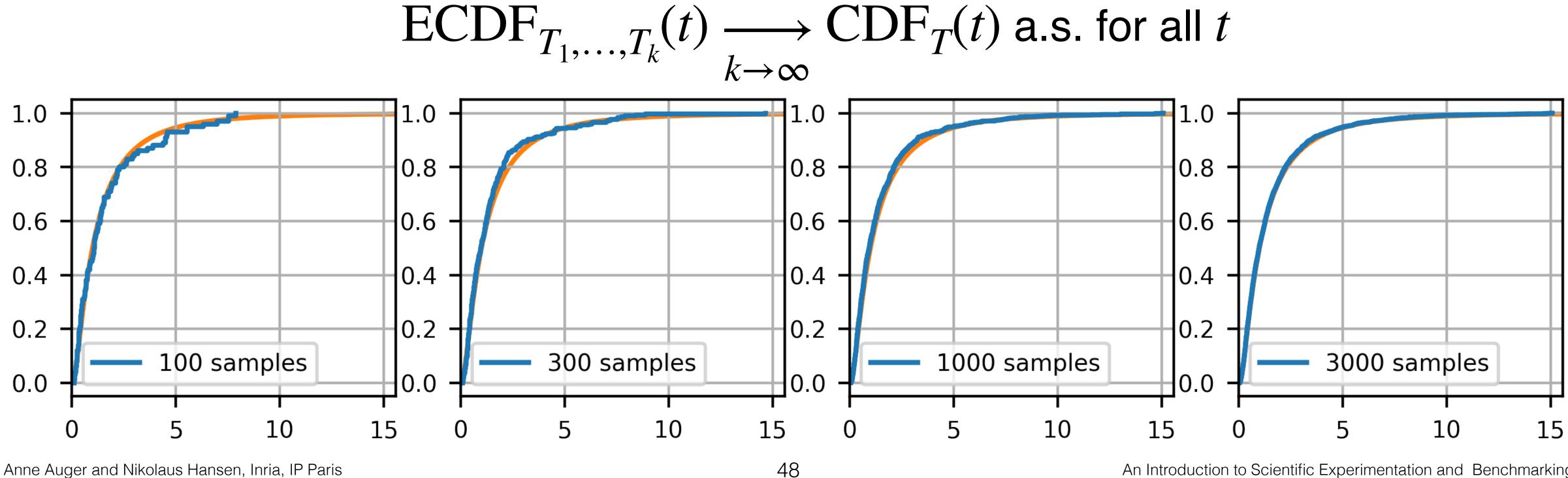


• Given a collection of data $T_1, T_2, ..., T_k$ (e.g. an empirical sample of a random variable) the *empirical* cumulative distribution function (ECDF) is a step function that jumps by 1/k at each value in the

Empirical Cumulative Distribution Function $ECDF_{(T_1,...,T_k)}(t) = \frac{\text{number of } T_i \le t}{k} = \frac{1}{k} \sum_{i=1}^k 1_{\{T_i \le t\}}$

For $\{T_i : i \ge 1\}$ i.i.d. realization of a random variable T, by the LLN





On Performance Measure

- When comparing algorithms:
 - Algorithm A is better than Algorithm B?
 - Algorithm A is 100 times faster than Algorithm B

- Requires •
 - adequate performance measure
 - adequate data collection

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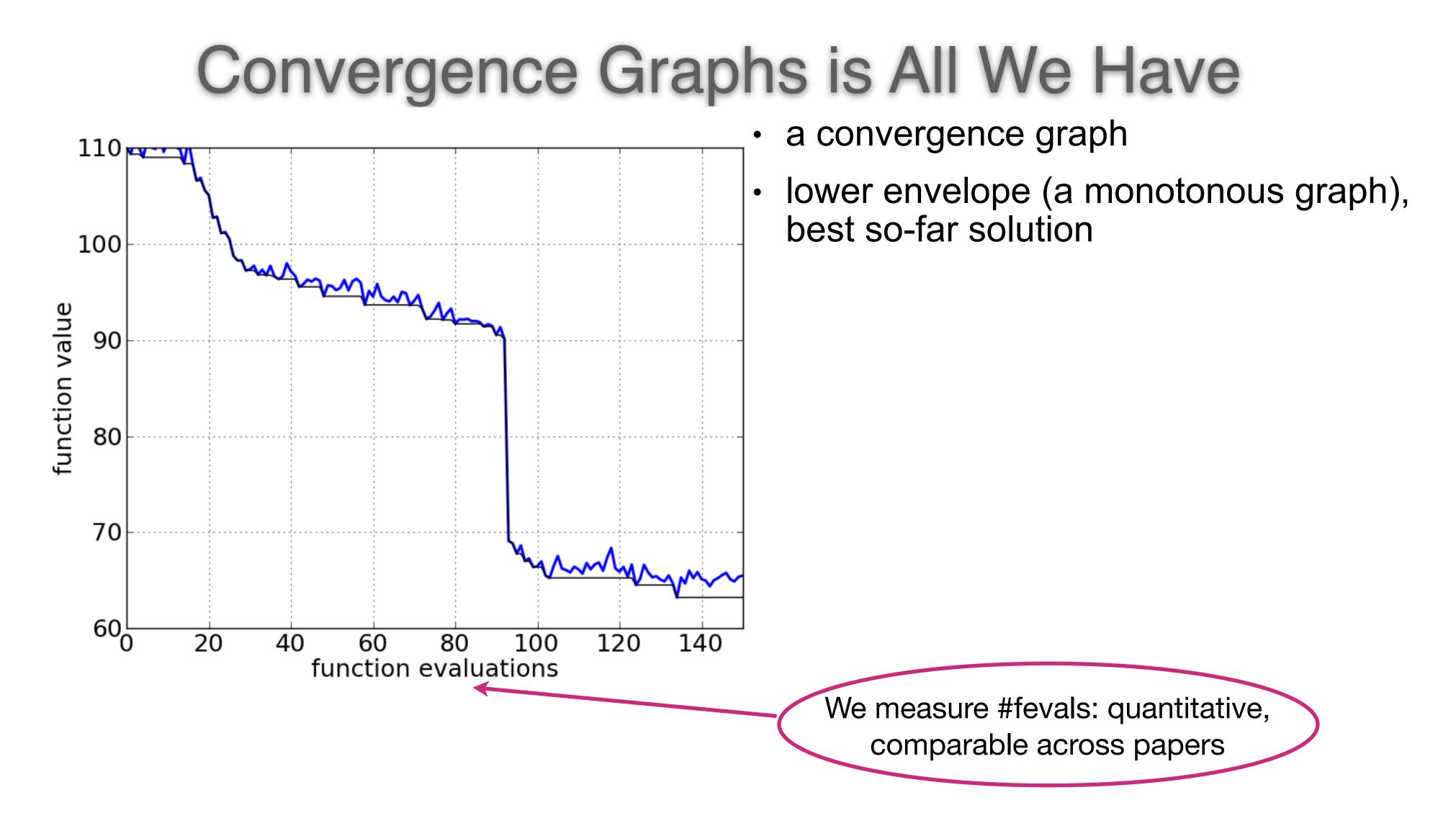
we want more than that

We want quantitative statements



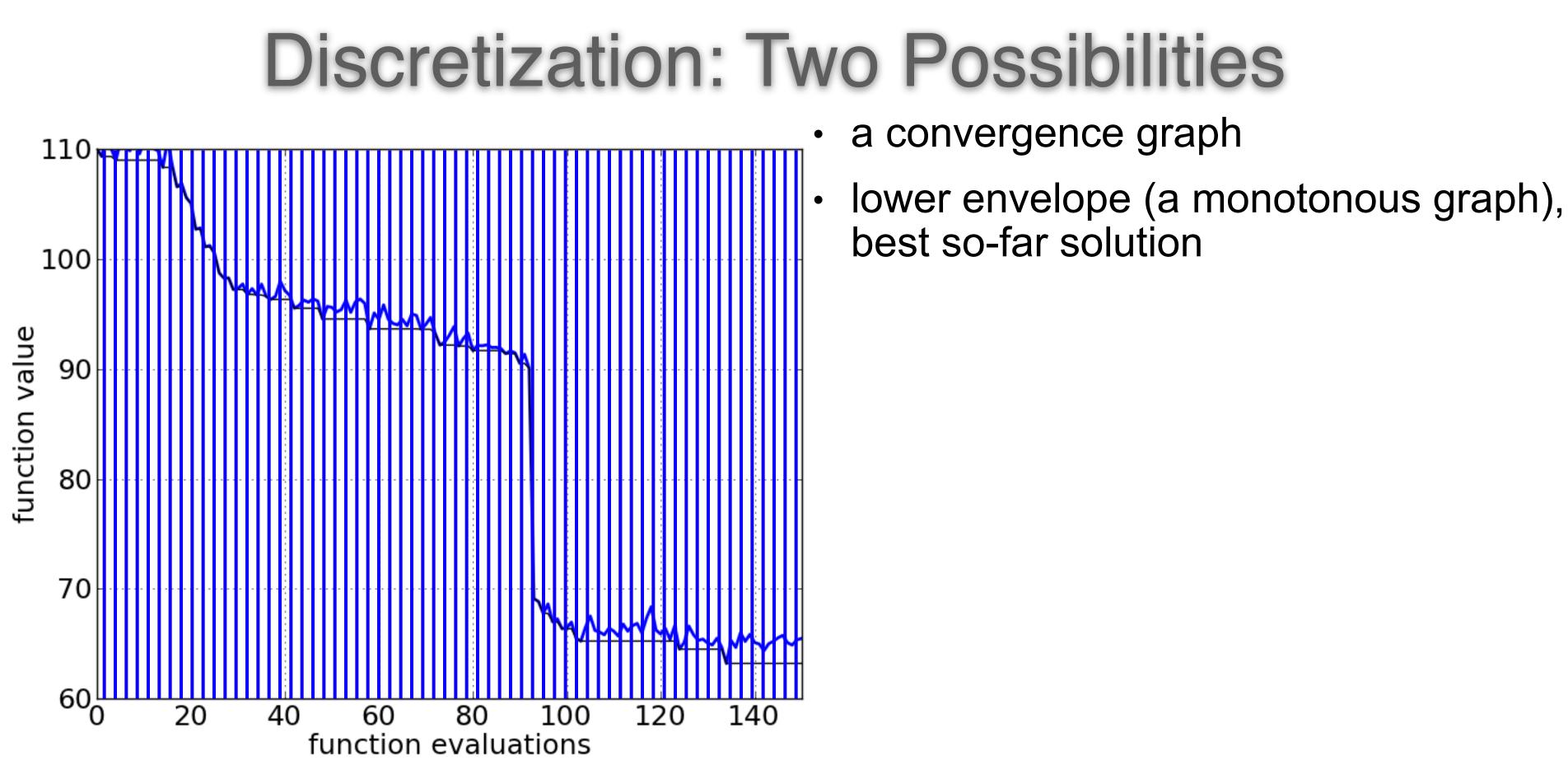
Collecting Empirical Data





using the lower envelope is a practical choice that relates to the first hitting time



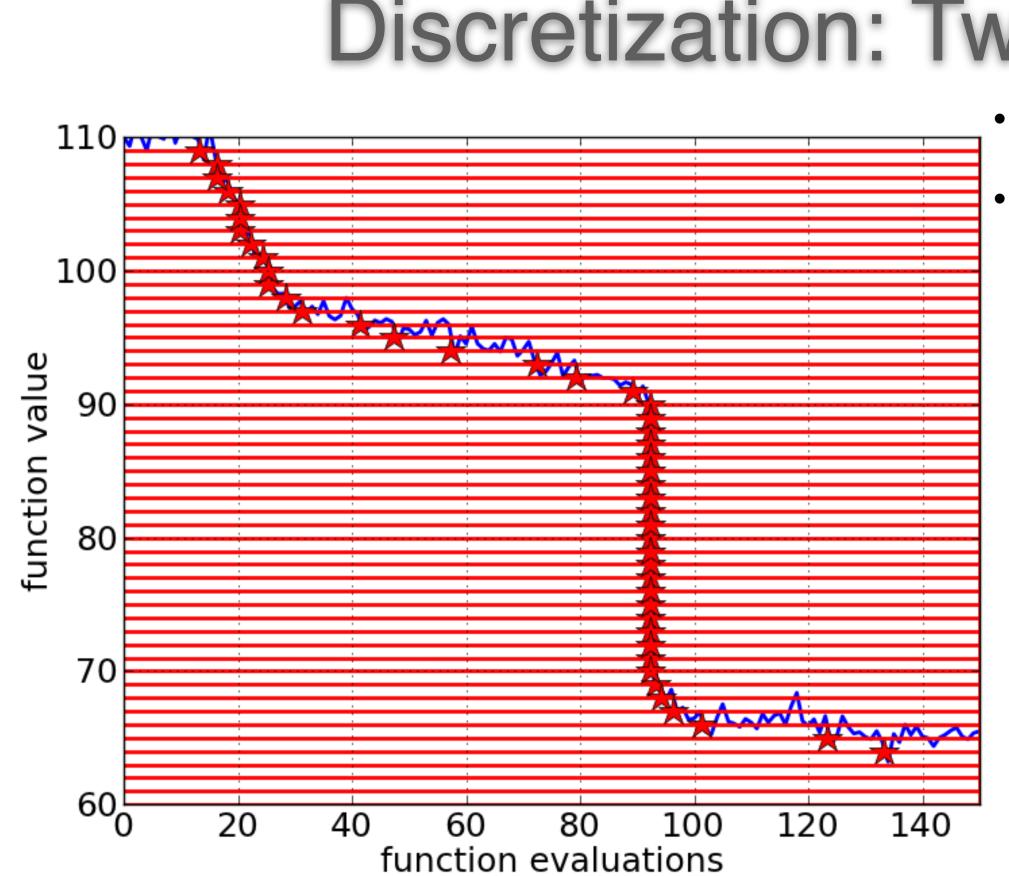


vertical: by evaluation is a natural discretization • for wall clock or CPU time we would need to determine discretization intervals

evaluations are the independent variable

function value is the dependent variable, the measurement





- horizontal: not a "natural" discretization
- function "target" values are the independent variable
- still recovers the original data

Discretization: Two Possibilities

- a convergence graph
- lower envelope (a monotonous graph), best so-far solution

we need to determine discretization intervals

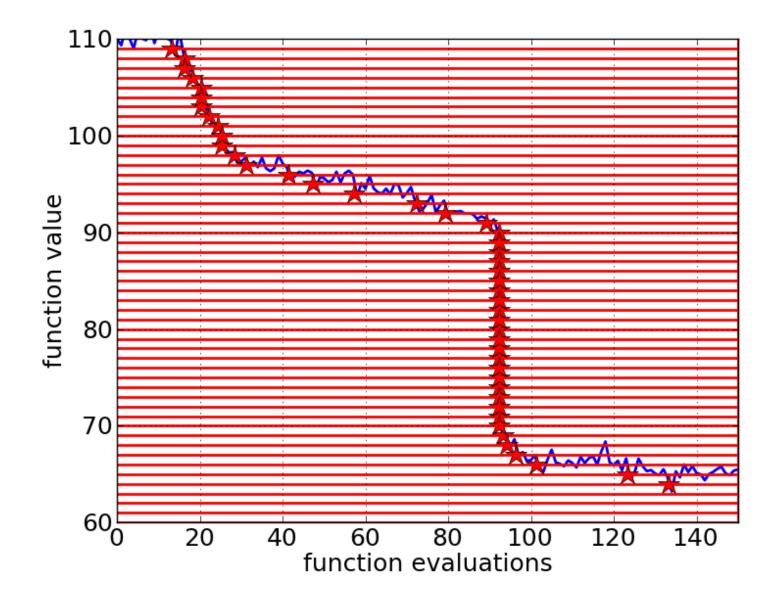
time is the dependent variable, the measurement

a time measurement for each discretization function value, these measurements can be plotted as ECDF



horizontal discretization

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using the

a technical subtlety

because it crucially determines the measurement we are looking at in the end

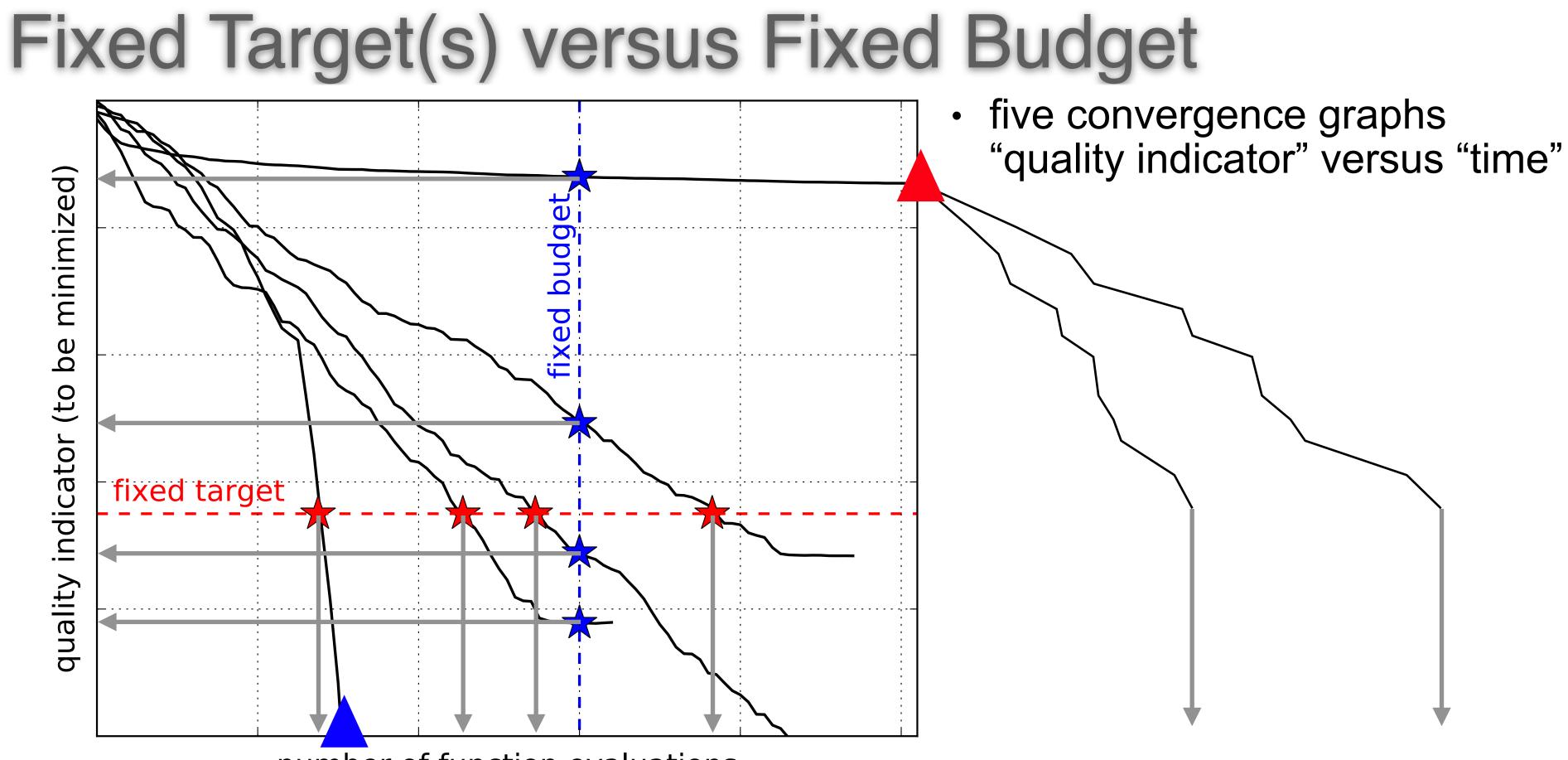
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is

not

just





number of function evaluations

- **Both** can lead to *imprecise data (a bound)* in some cases \bullet
 - "too" good performance (often overlooked)
 - "too" bad performance

then the data only provide a lower bound estimate for the runtime (and a fixed budget measure at the maximum budget) quick and dirty fix: assign 10 x time_out_budget

(reached global optimum up to the relevant or numerical precision before the given budget) quick and dirty fix: assign best possible (or measured) function value



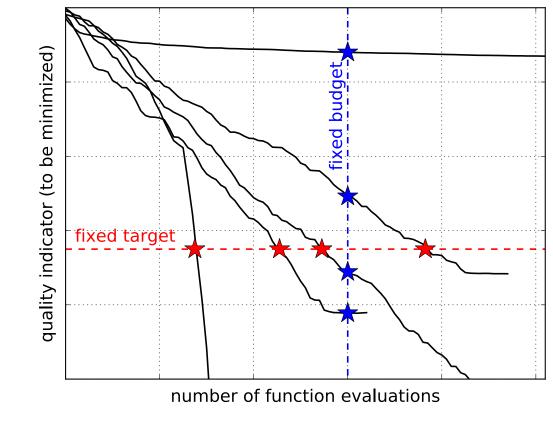
Fixed Target(s) versus Fixed Budget

The resulting measurement

- •

Does this make a difference?

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Fixed budget (vertical, target-free) design: function values (quality)

Fixed target design (budget-free) design: evaluations (runtime)

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Scales of Measurement ("Level" of Data)

- Nominal categorial, define a classification
- Ordinal define an order, ranks, function values (fixed budget)
- Interval differences are meaningful
- Rational ratios are meaningful, we usually can take the logarithm, function *evaluations* (fixed target)

CAVEAT: mathematical and semantic treatment of data is not the same. From a classification with values {1, 2} we can *mathematically* take differences and ratios of the values, but they have no meaningful *semantic interpretation*. Fahrenheit or Celsius versus Kelvin describe temperatures, however only Kelvin is on a rational scale of measurement.

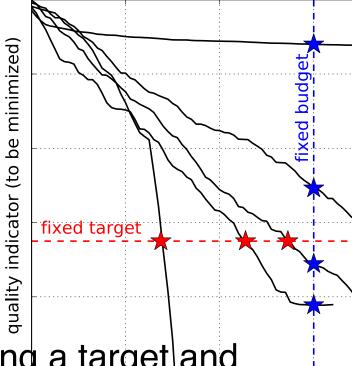


Summarizing Fixed Target(s) versus Fixed Budget

- The fixed budget (vertical) design is (much) easier to set up
- Runtimes have a quantitative interpretation
- geometric averages

whereas function values from different functions are in general not commensurable

Fixed target results are "budget-free"



target-free: choosing a budget is simpler than choosing a target and we need to choose a maximal "timeout" budget either Way

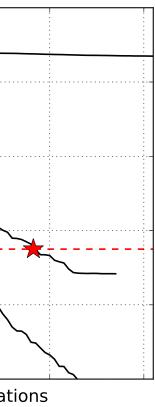
• For the (very) same reason, results from the fixed target (horizontal) design are (much) simpler to interpret and usually more conclusive

without specific knowledge/insight, a function value is impossible to interpret beyond ordering

"Algorithm A is 100 times faster than Algorithm B"

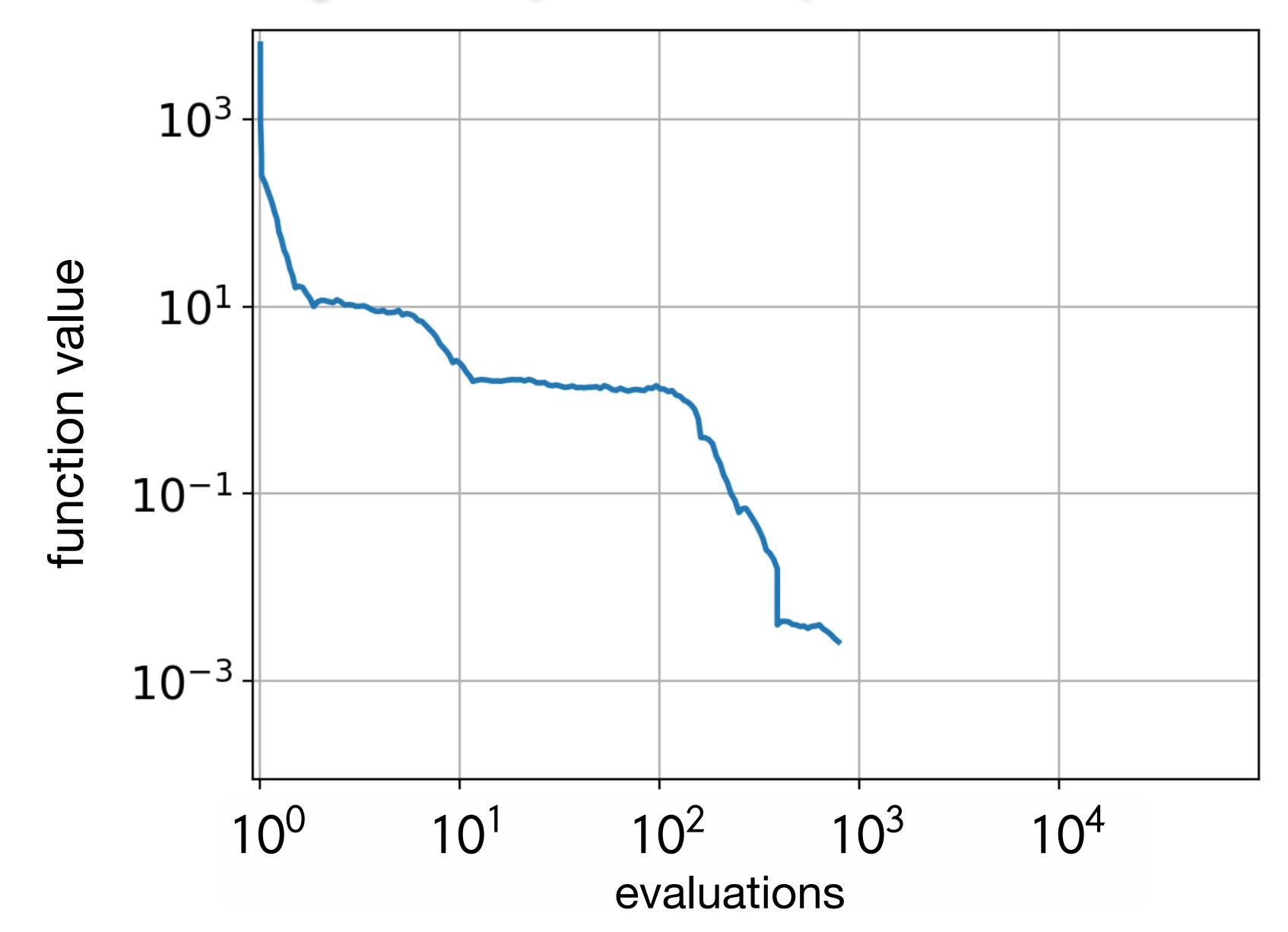
Fixed target results can be meaningfully aggregated in ECDFs and

we can compare results with different maximal "timeout" budgets

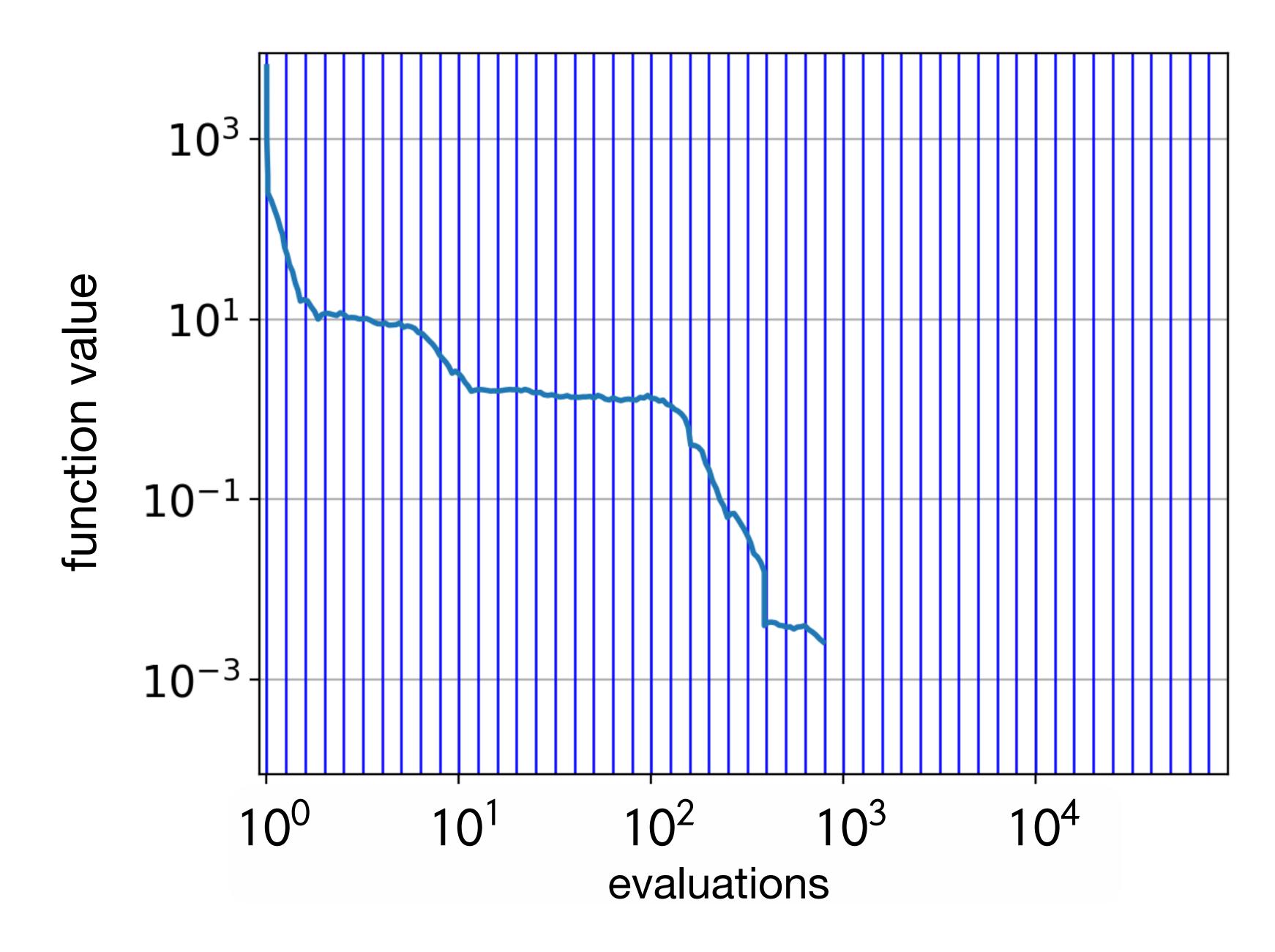




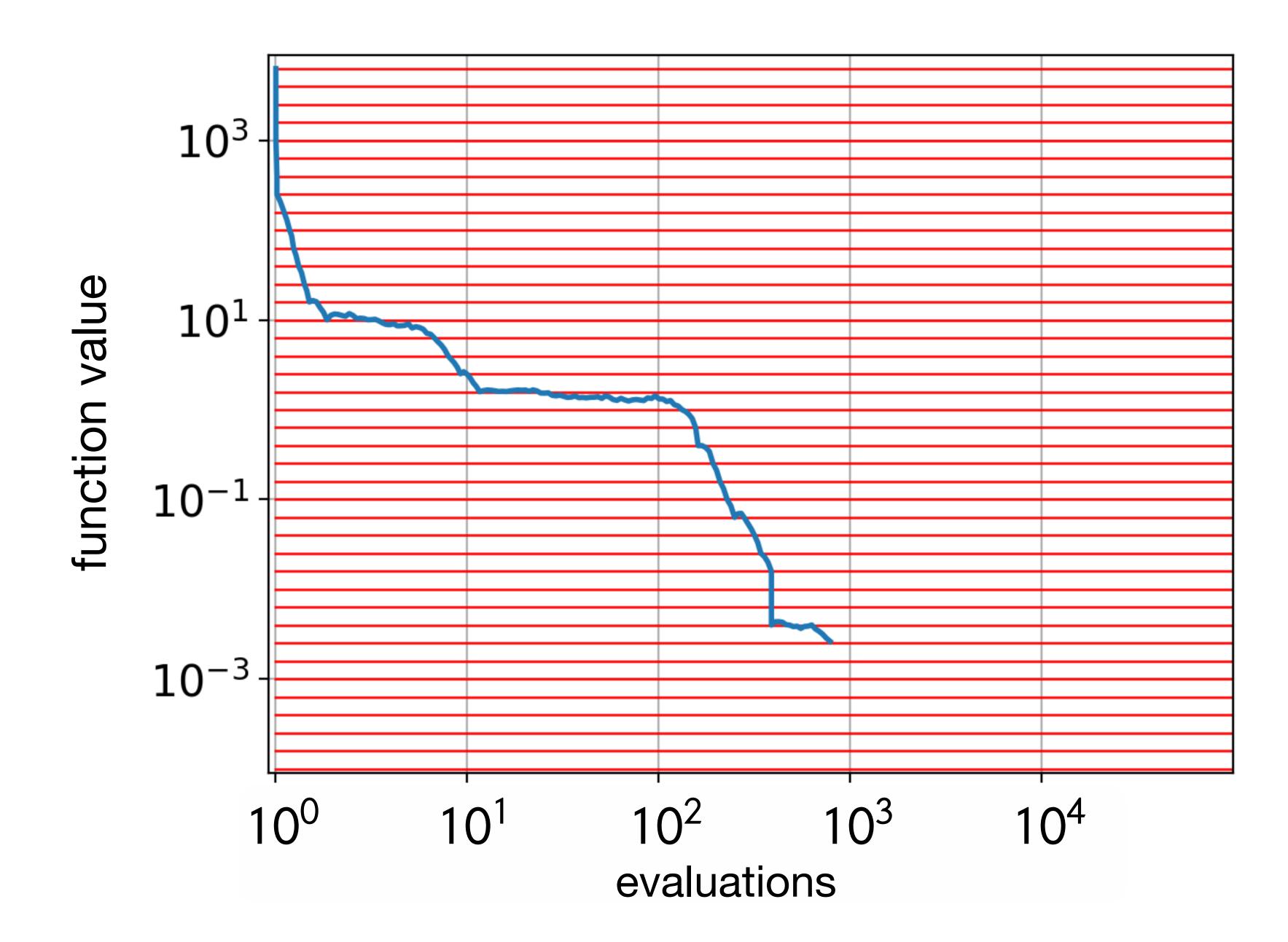
From a Convergence Graph to the Empirical Runtime Distribution



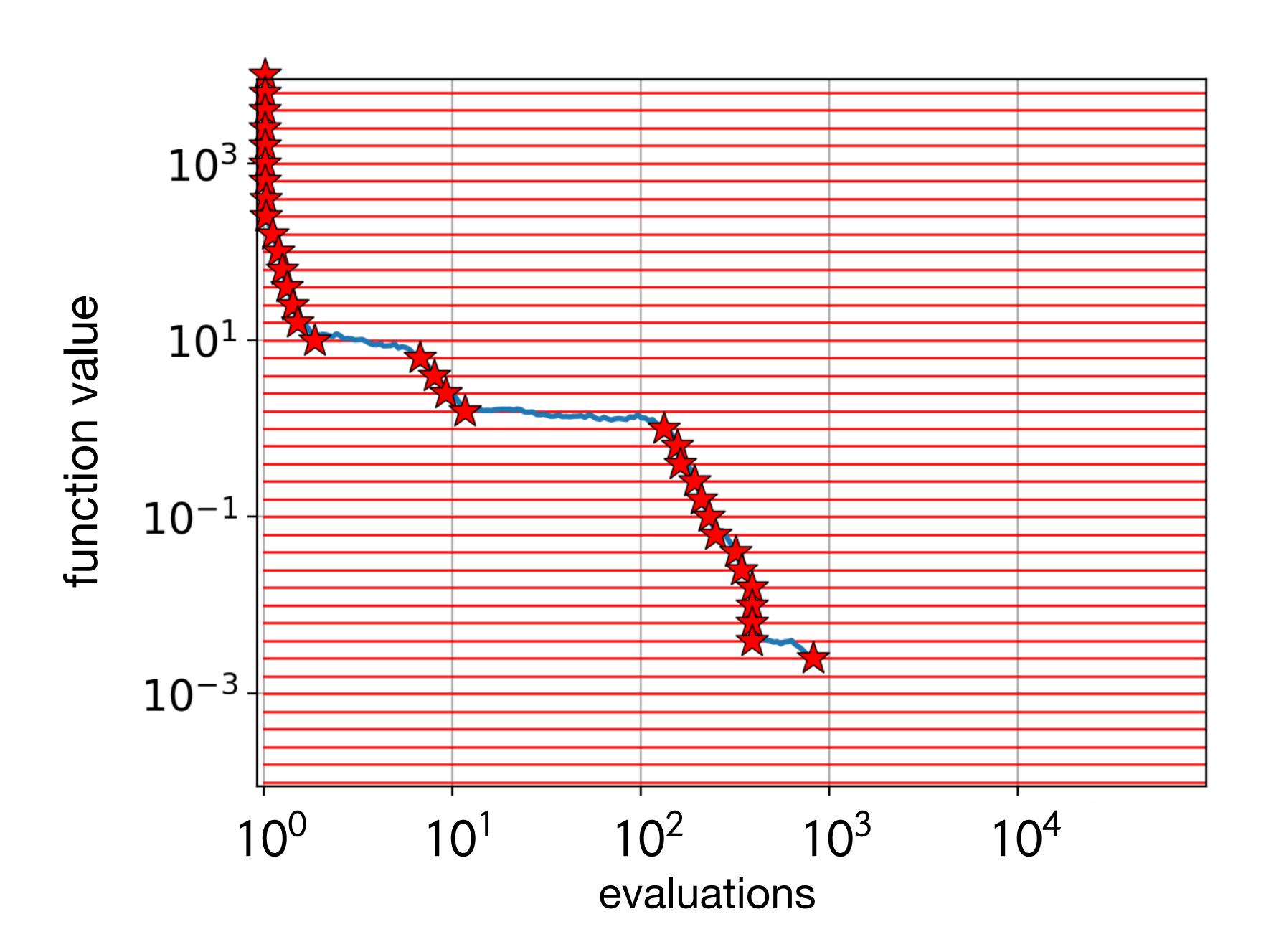




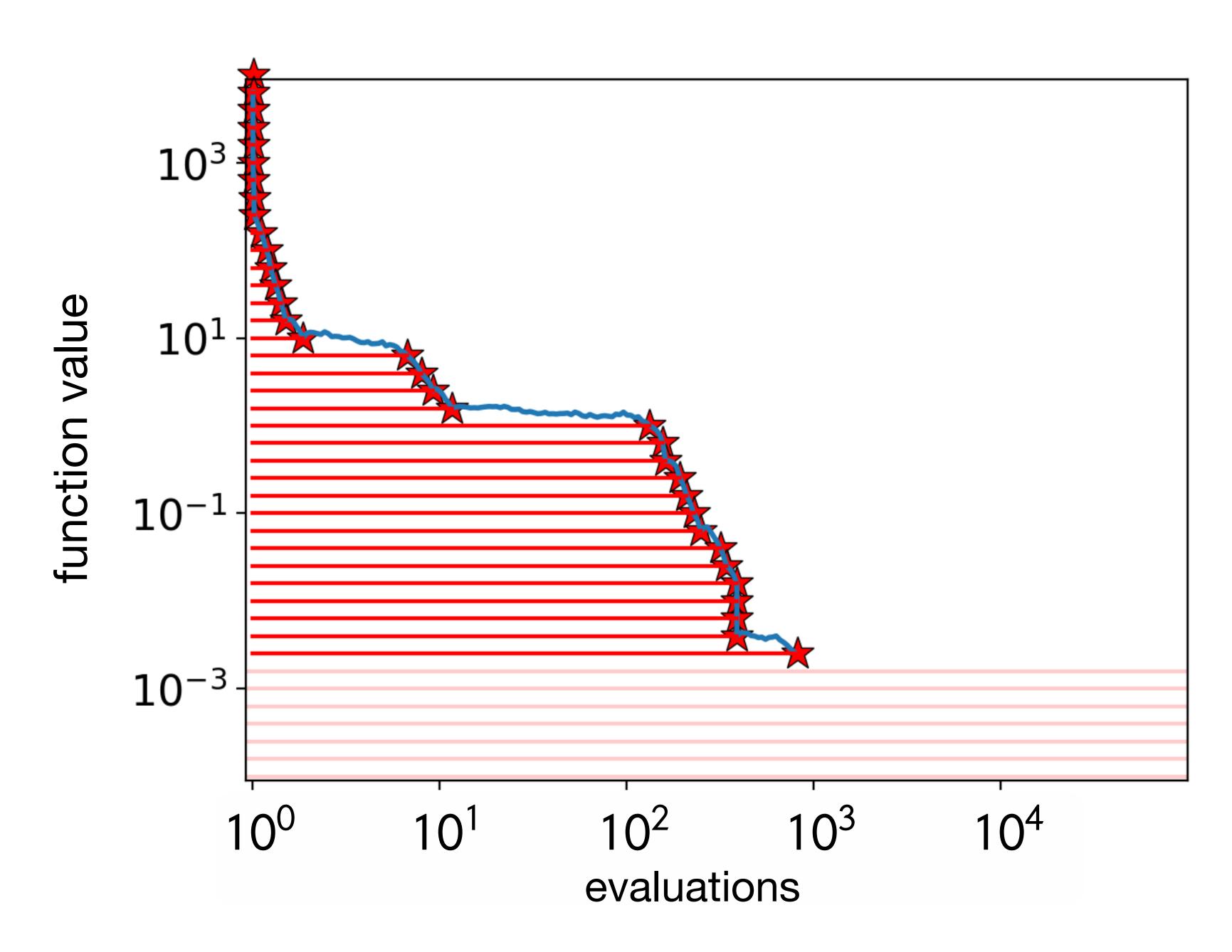




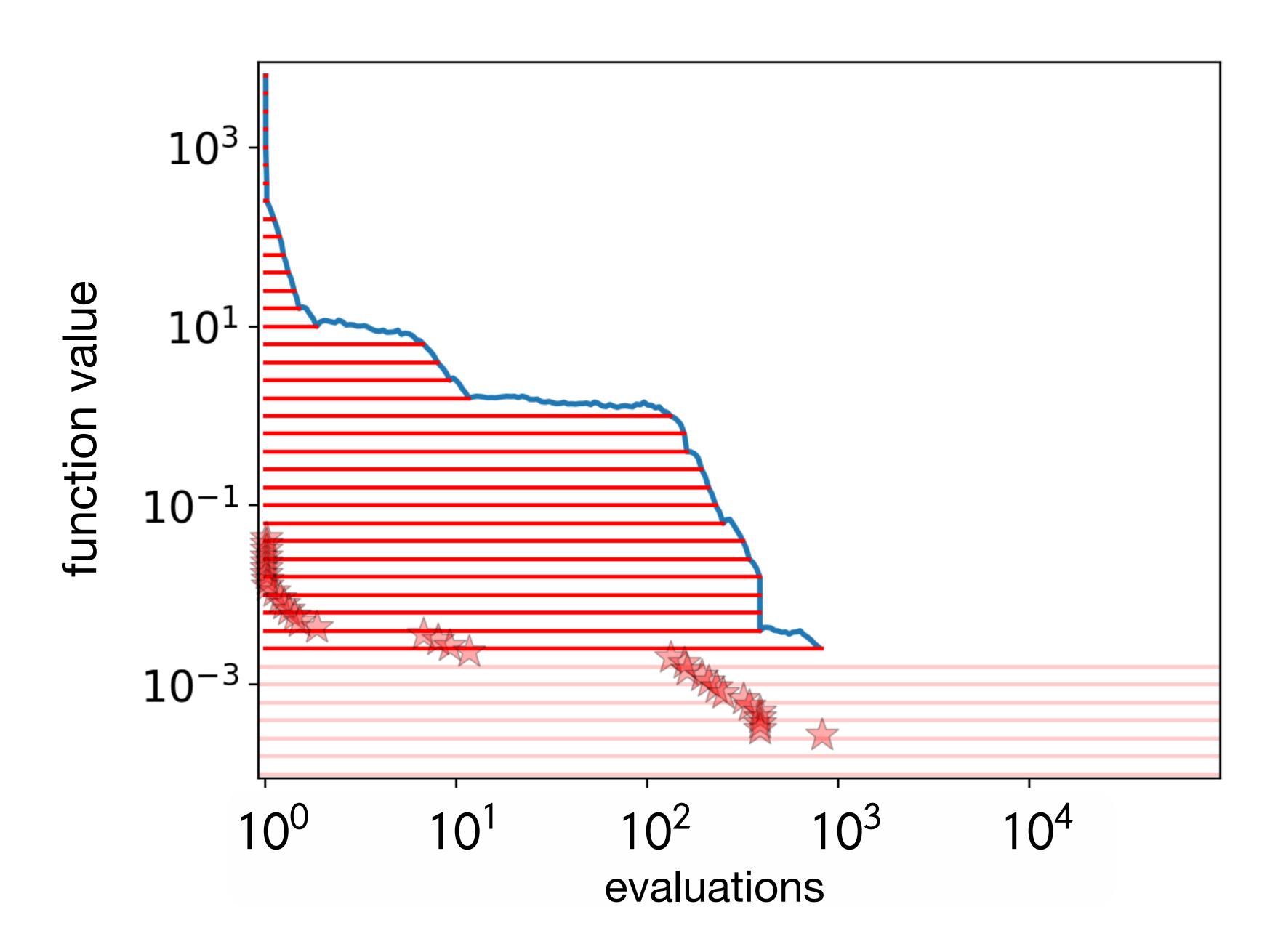




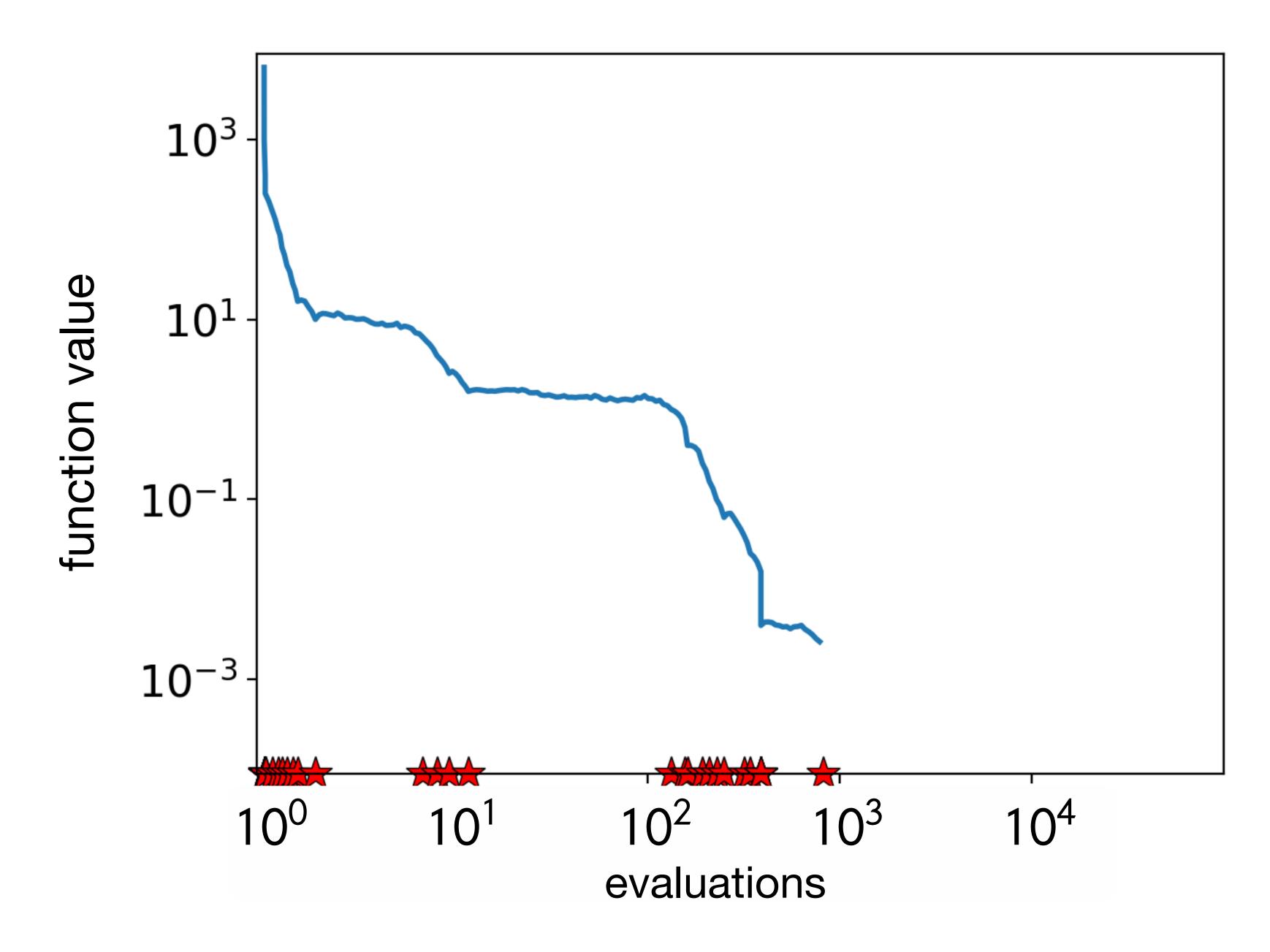




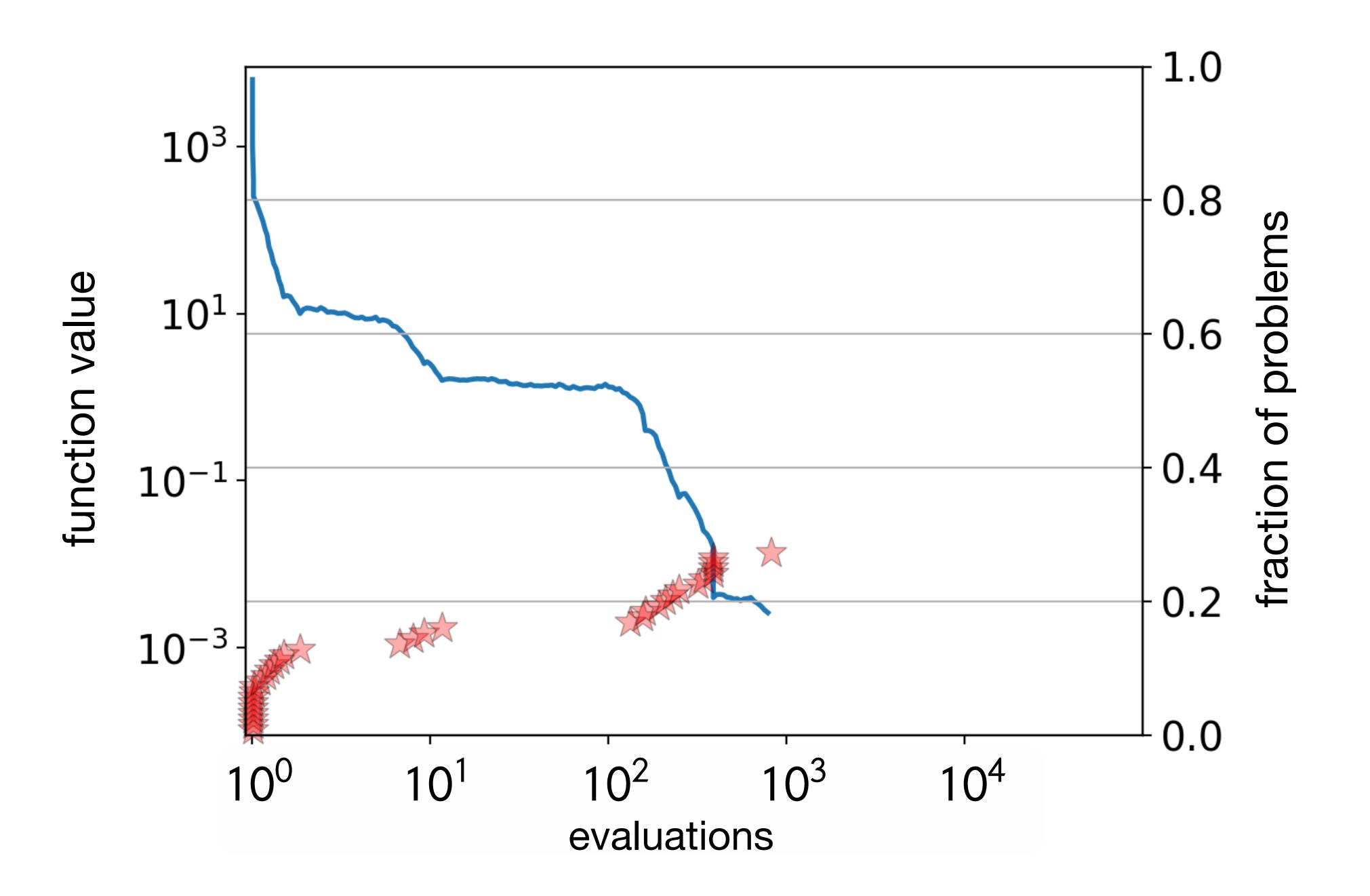




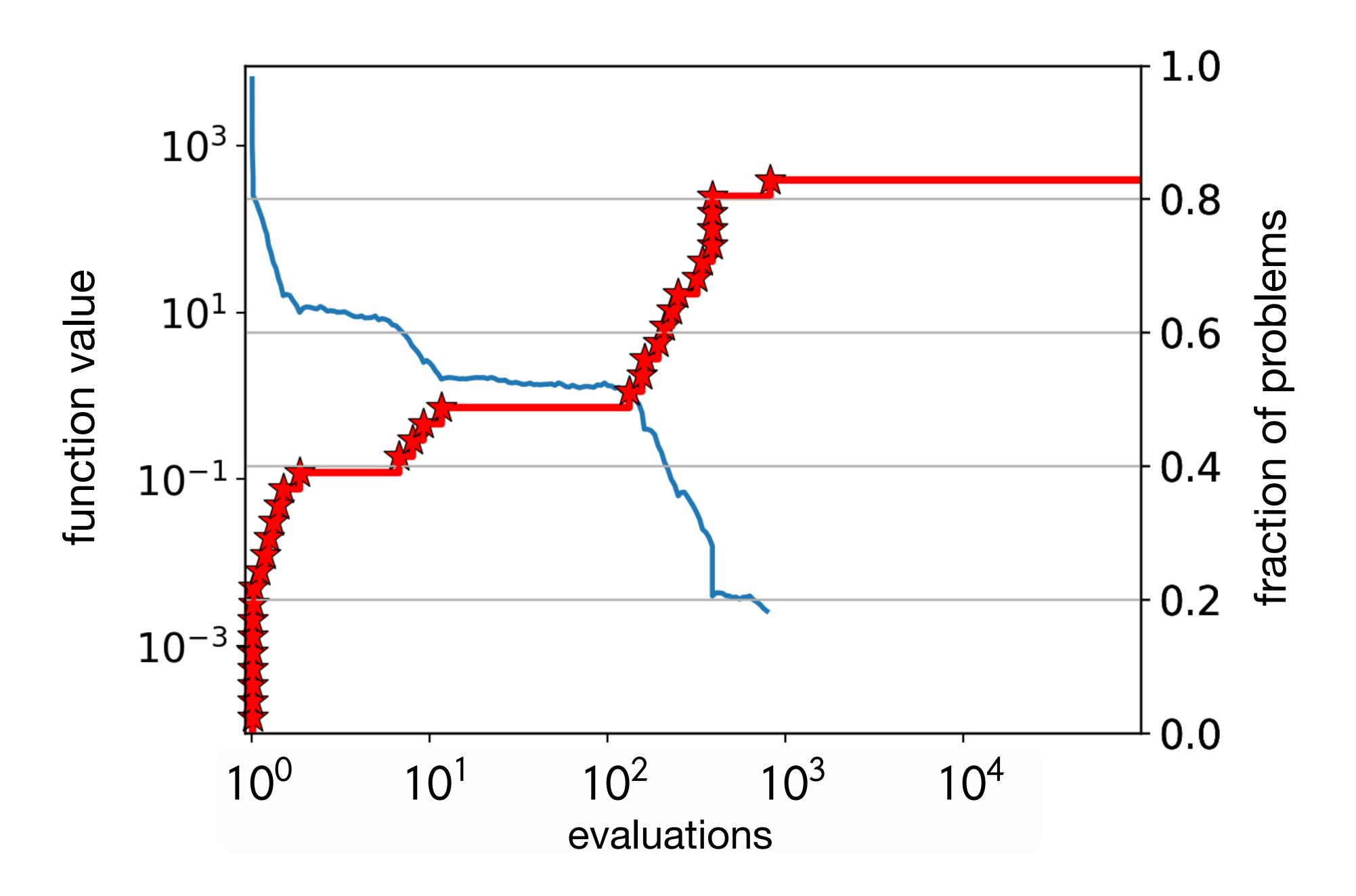




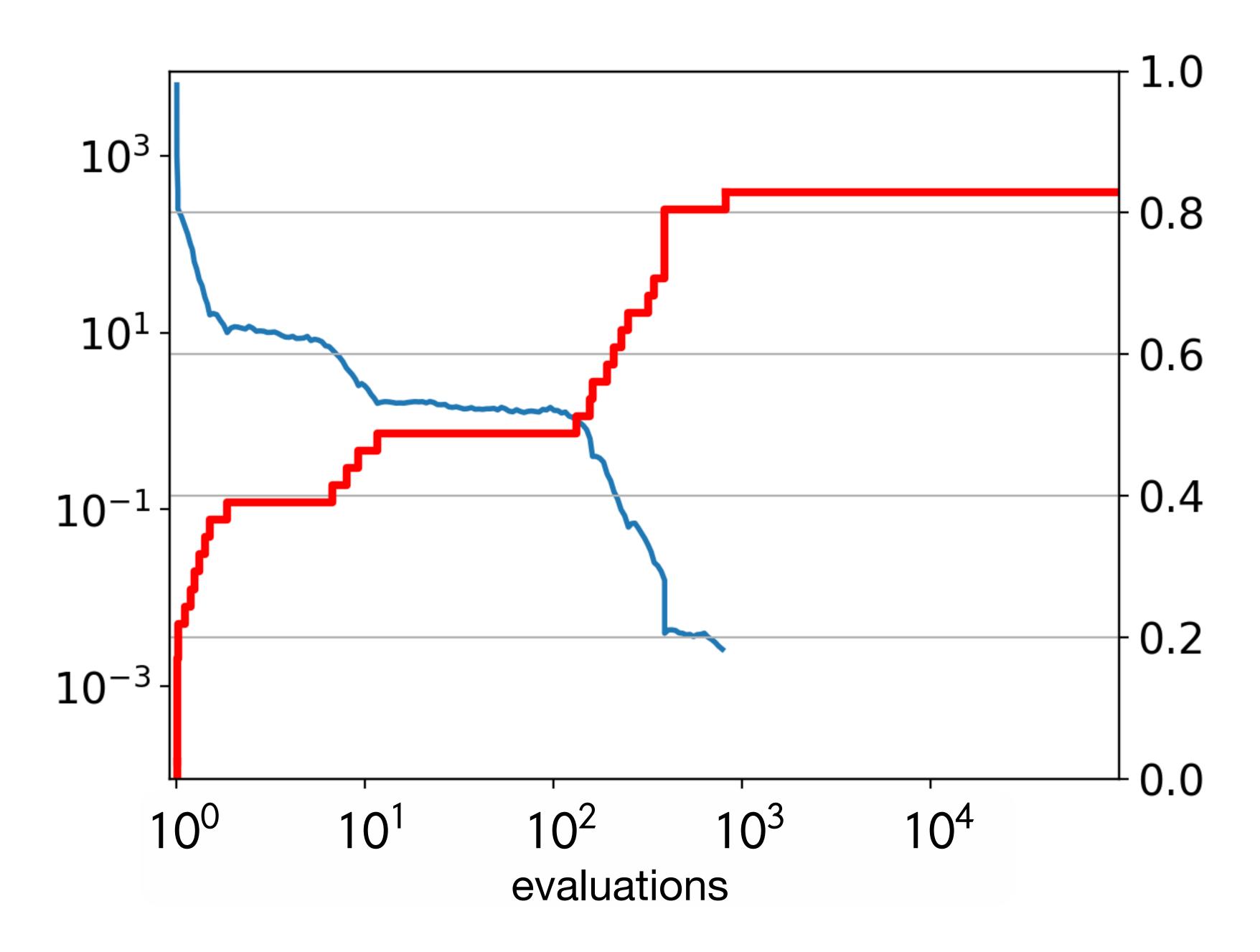




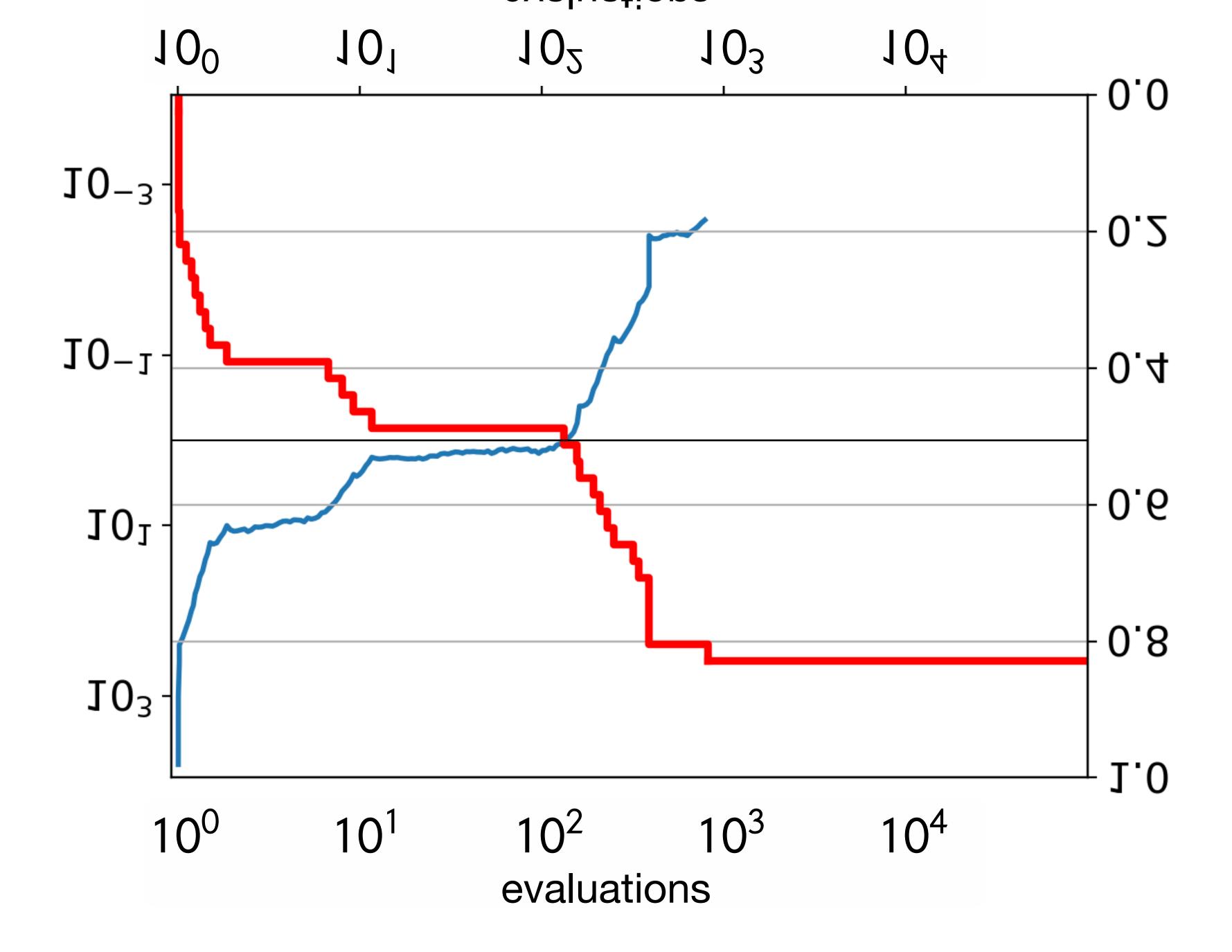




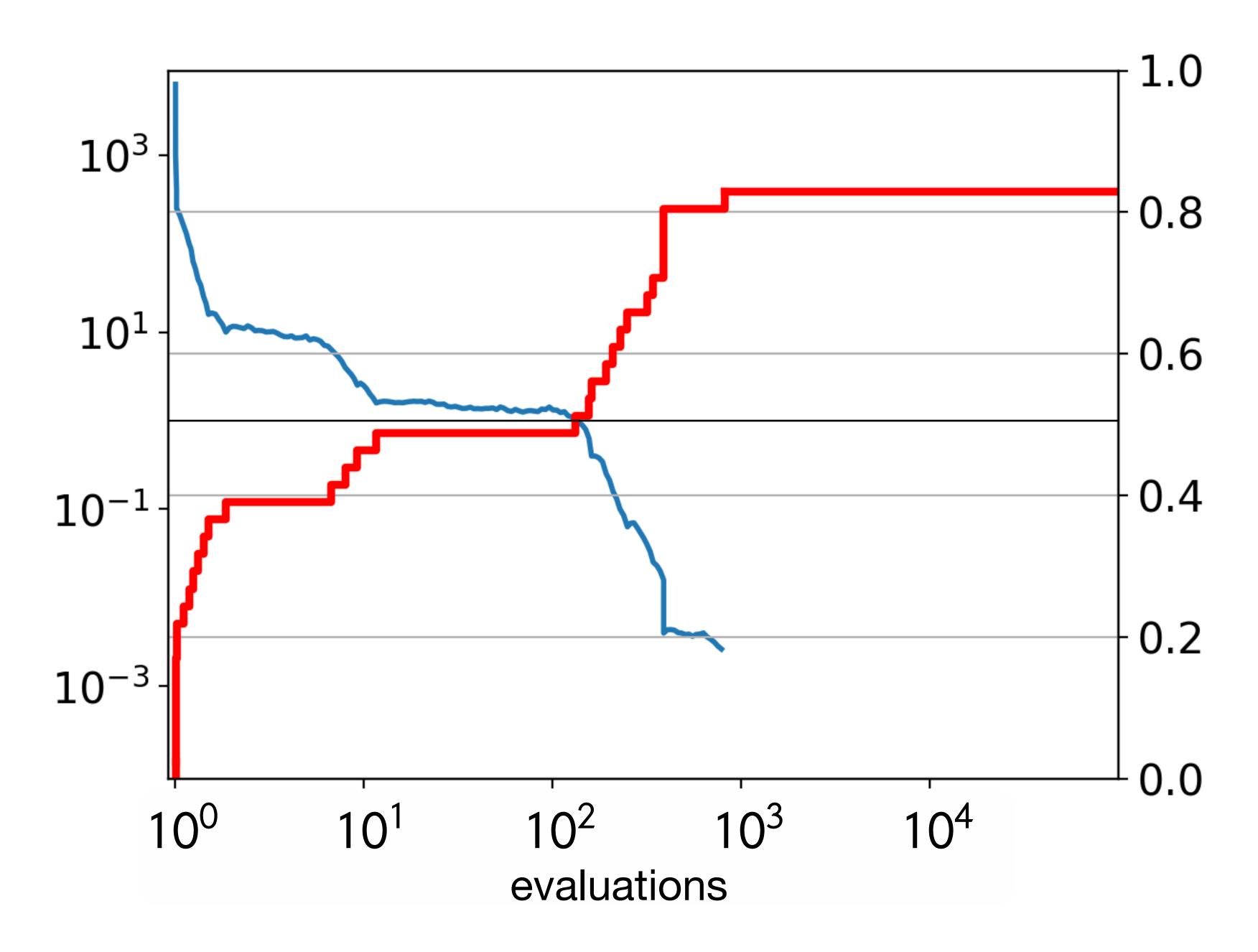




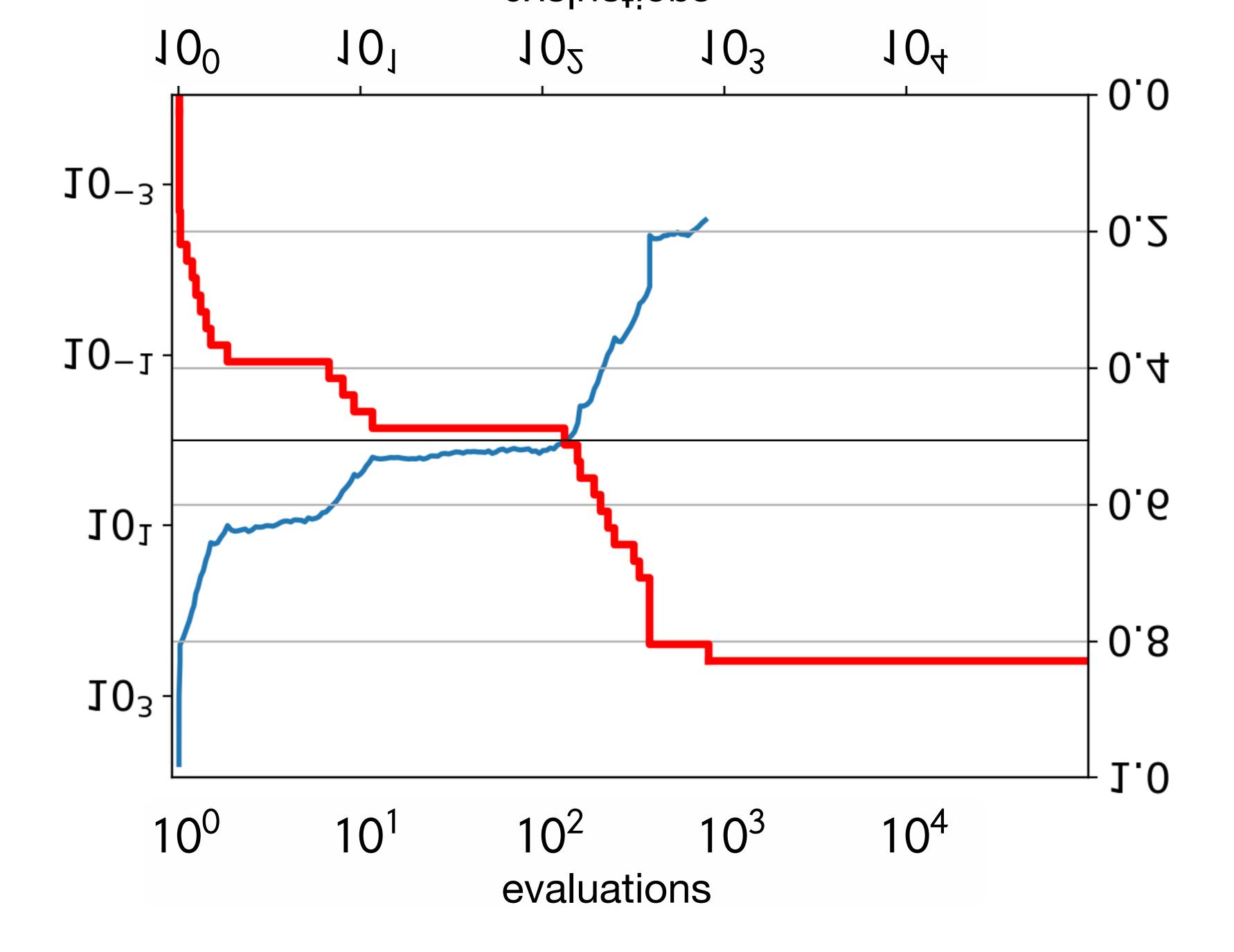




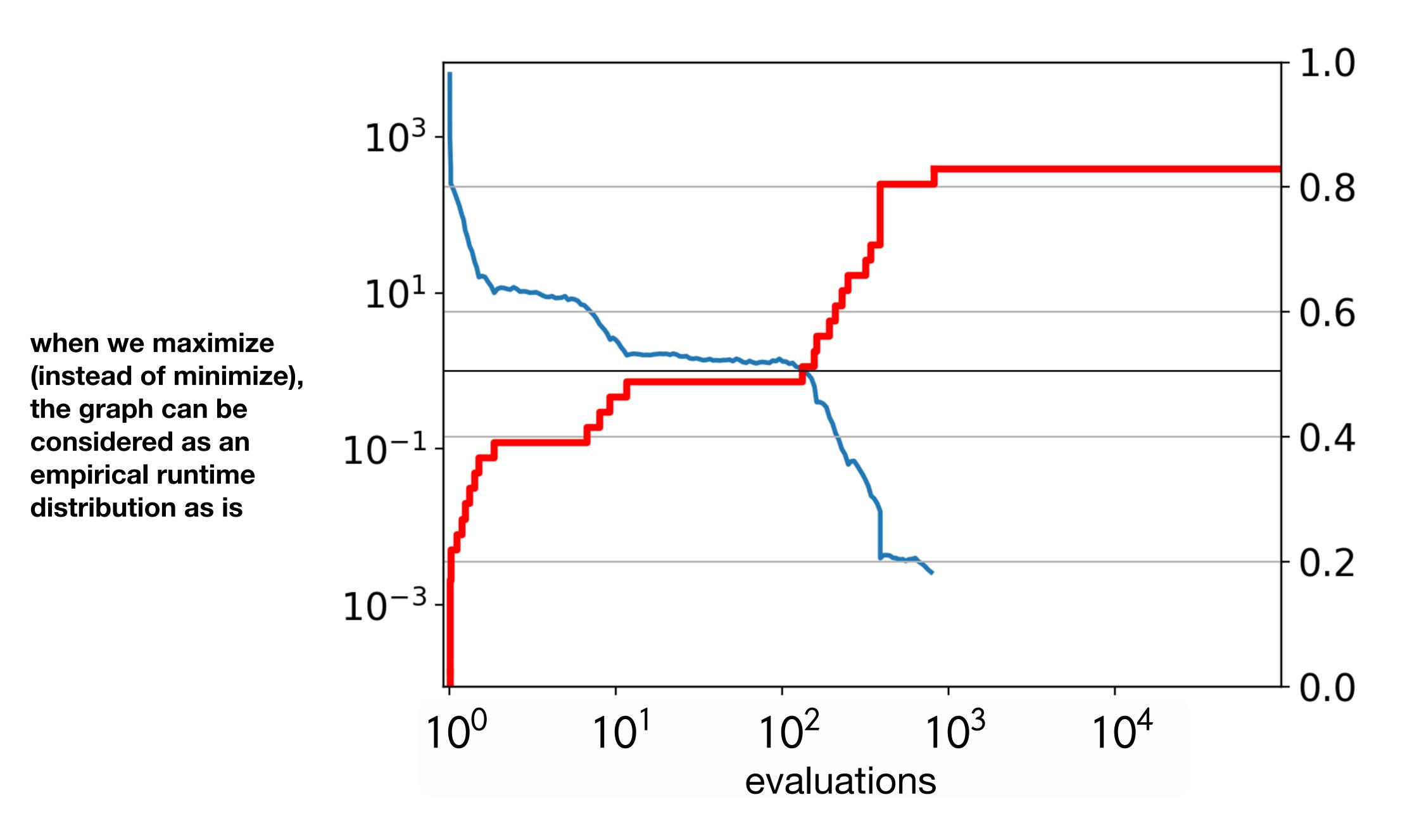




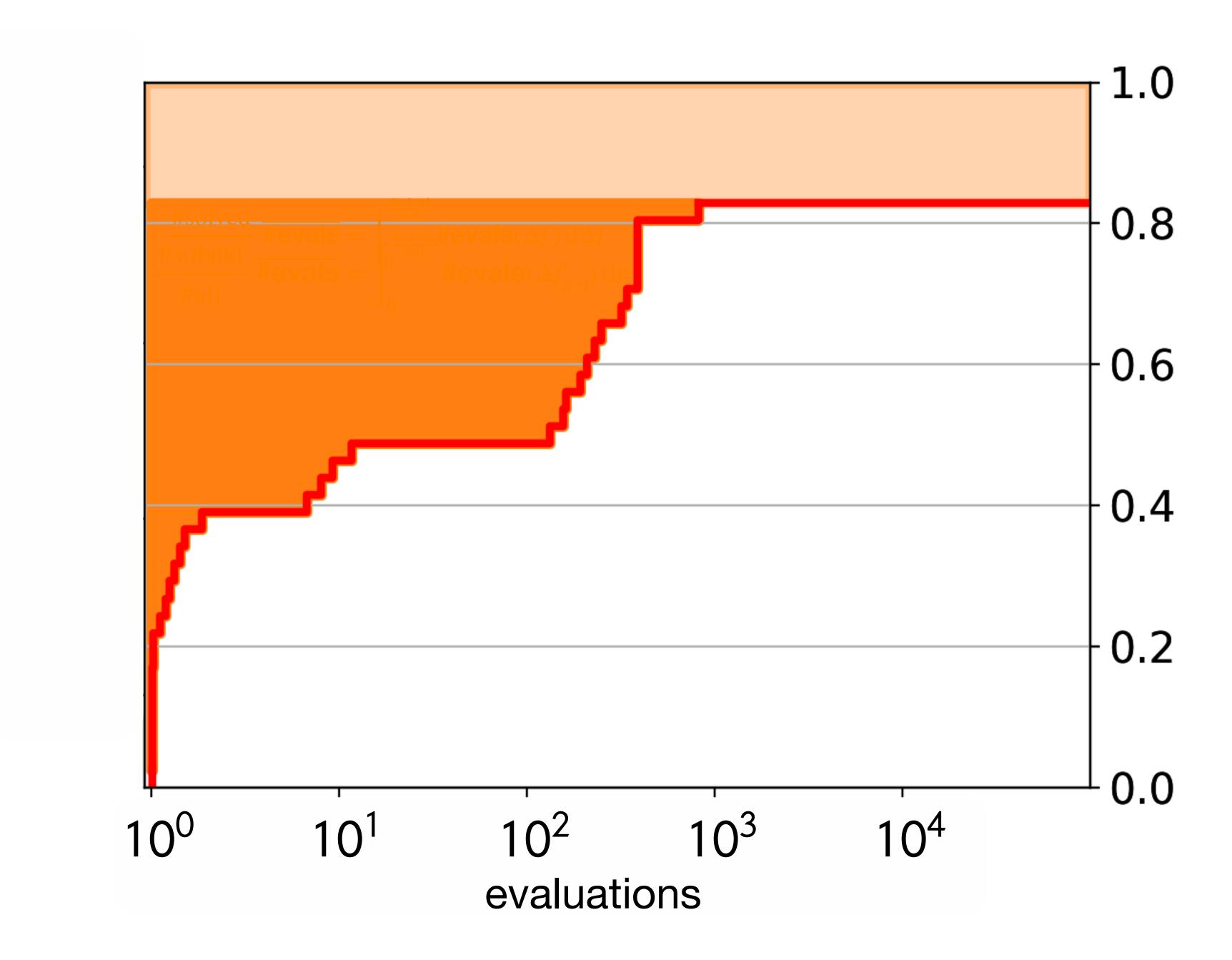












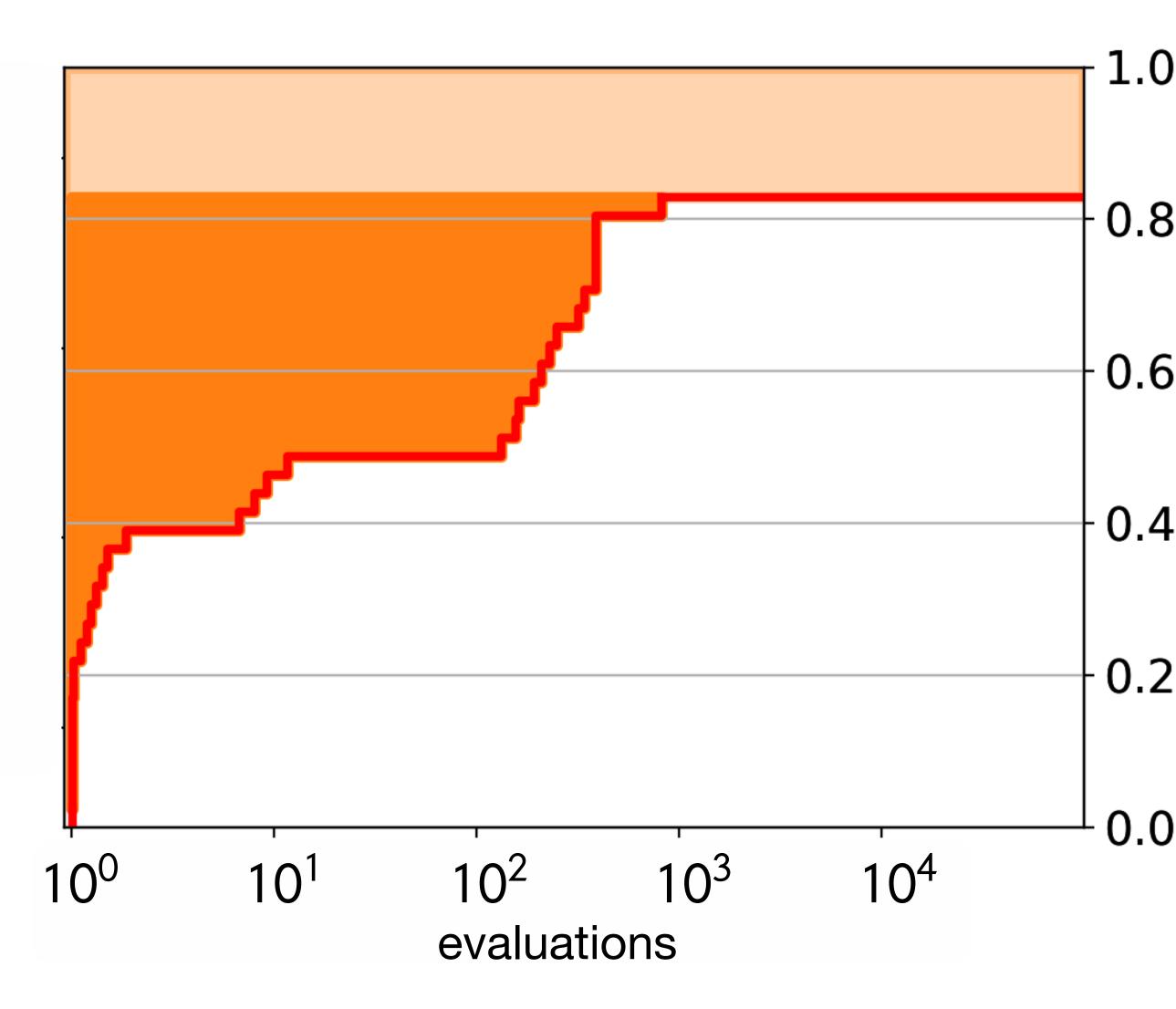


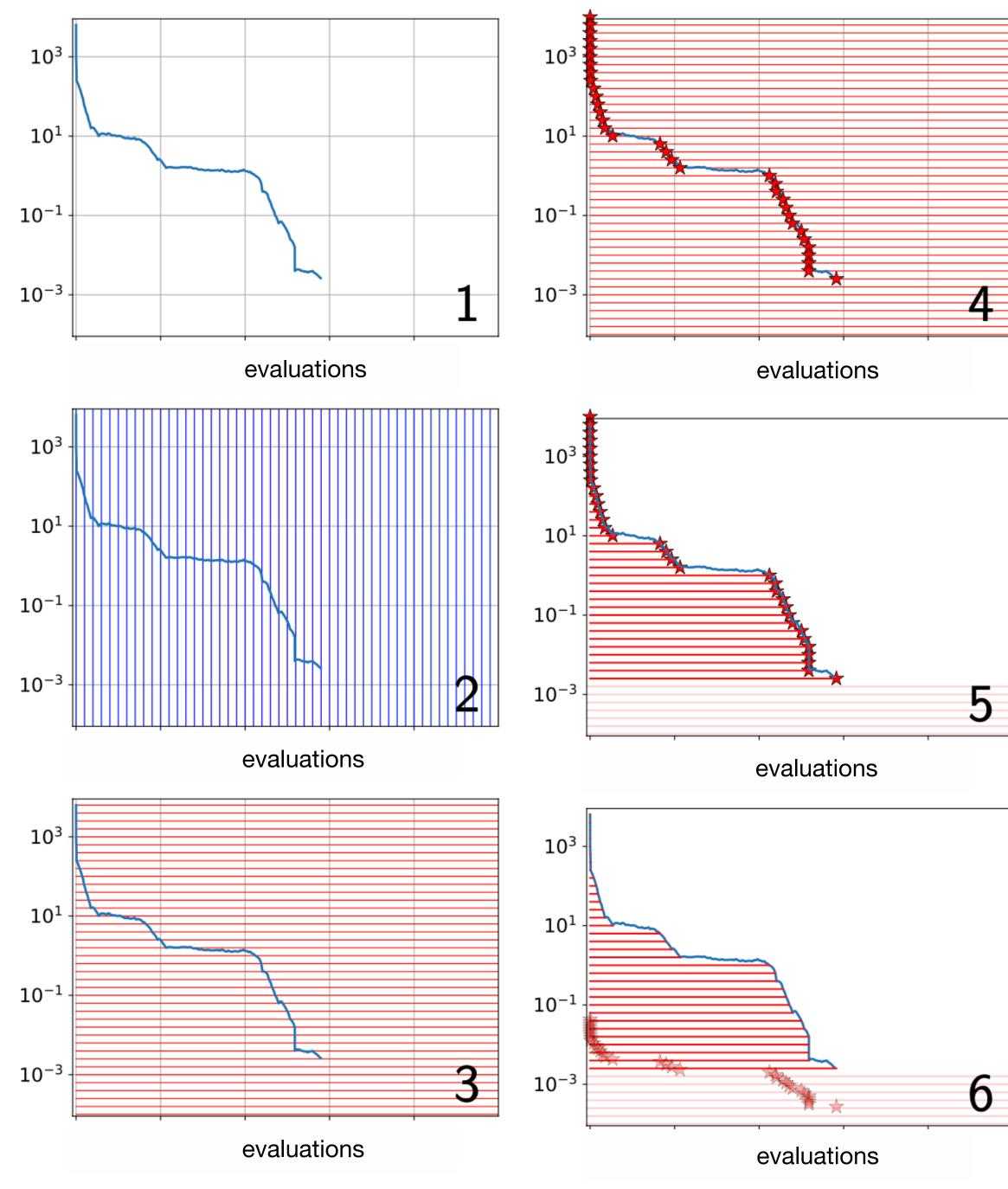
$$\overline{\text{\#evals}} = \frac{\#all}{\#solved} \int_{0}^{\frac{\#solved}{\#all}} \#evals(\Delta f_{i(r)})$$

When the x-axis is in log-scale, it is the geometric average

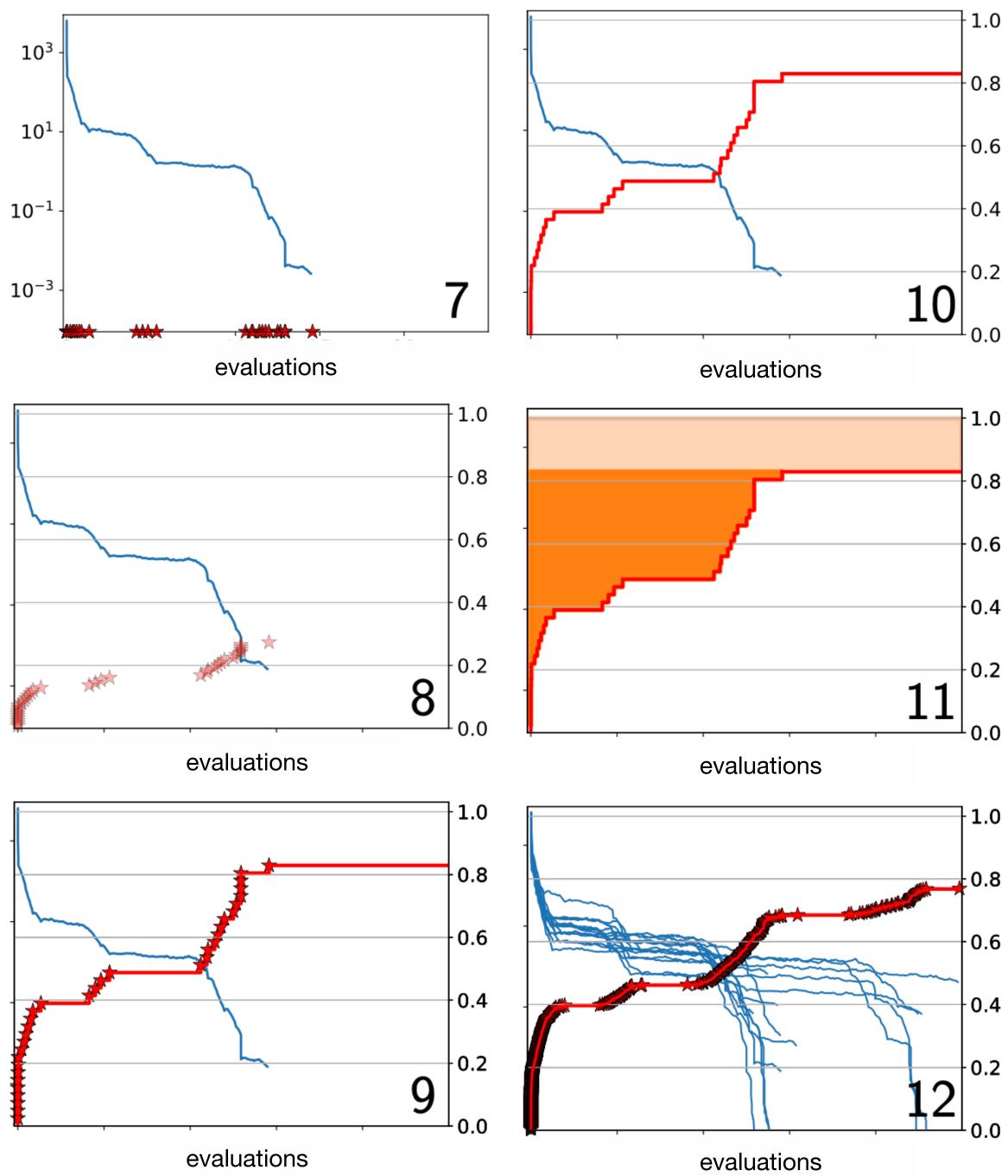
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dr



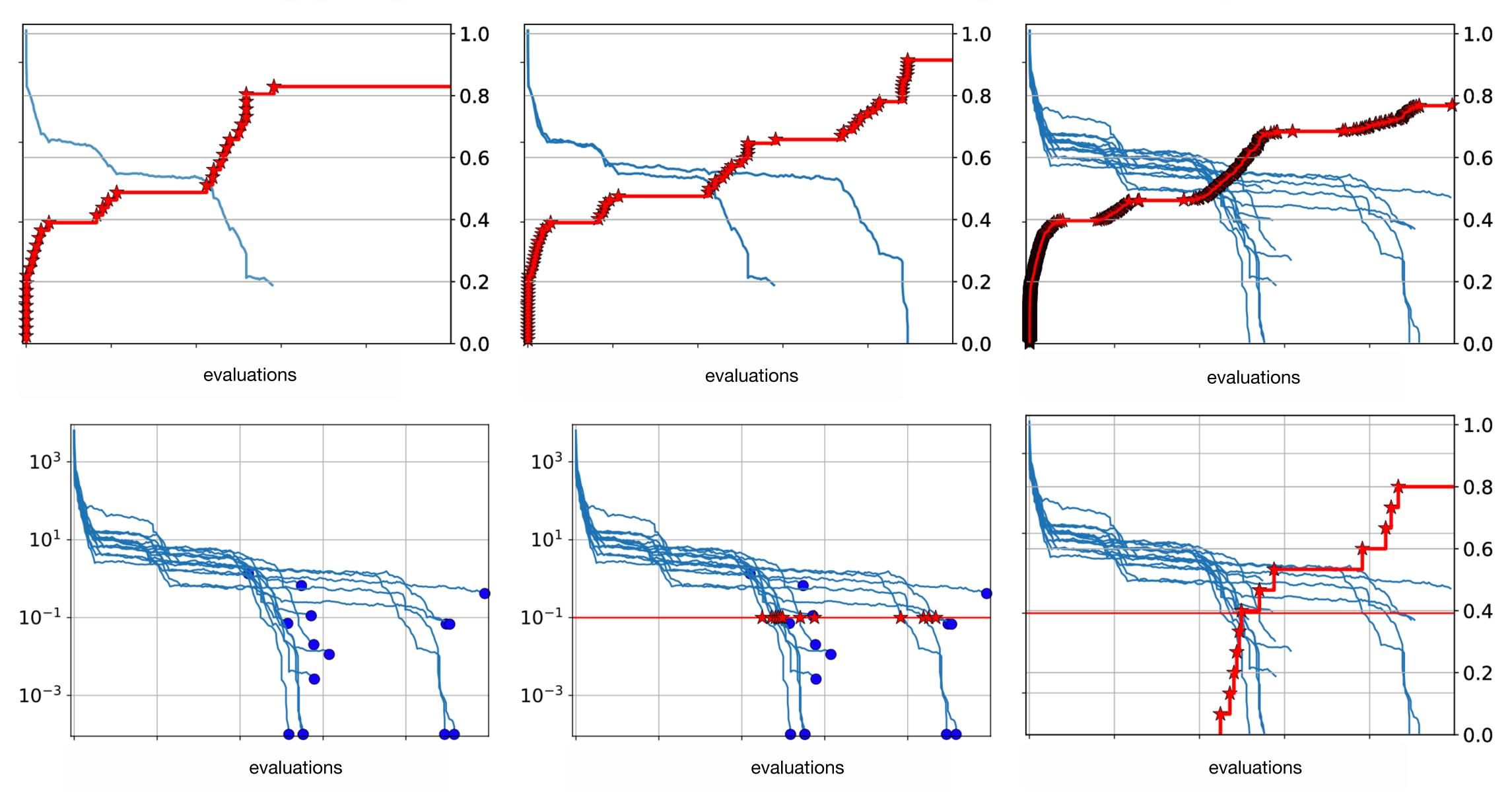


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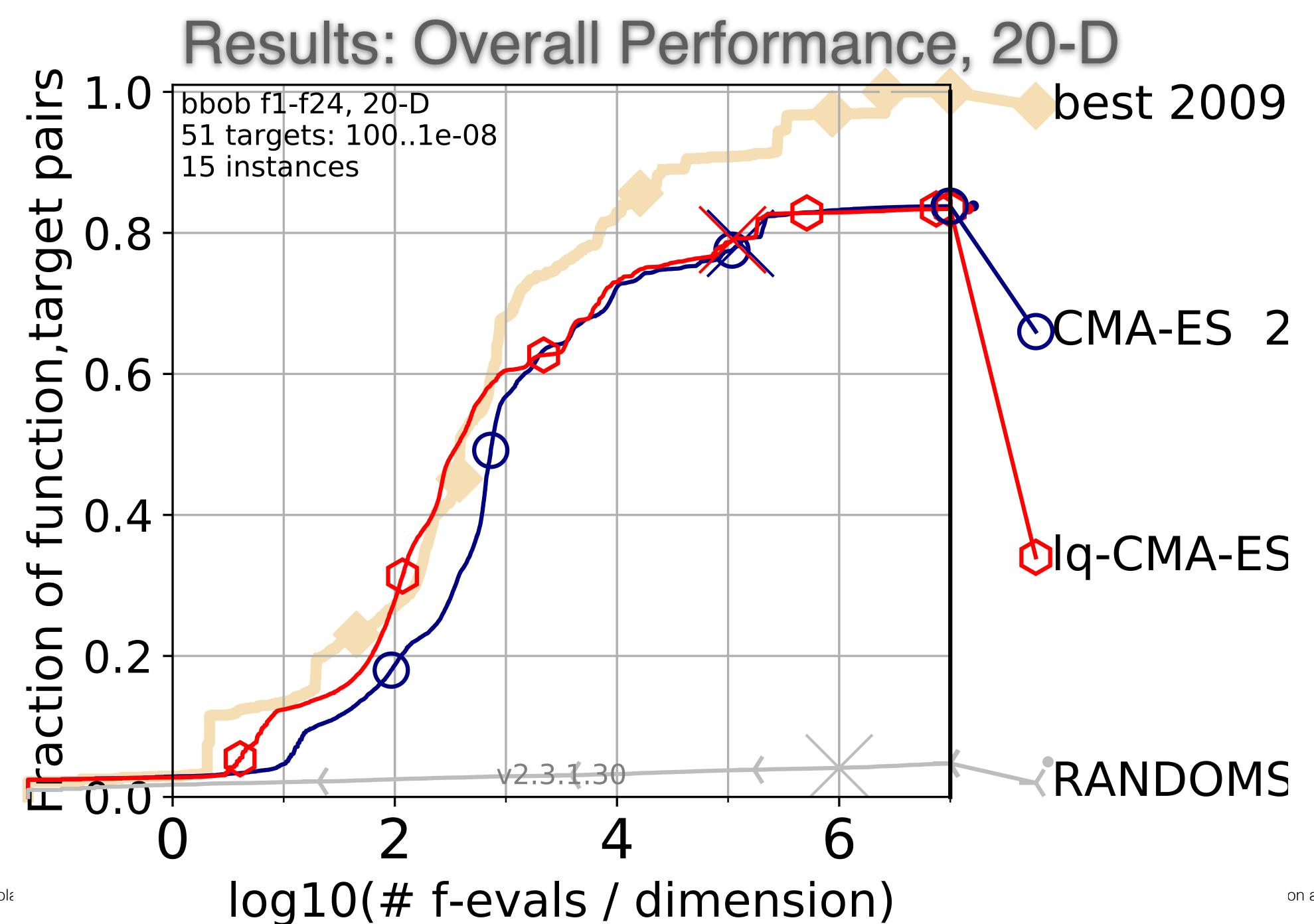
An Introduction to Scientific Experimentation and Benchmarking

Aggregation of Several Convergence Graphs



Anne Auger and Nikolaus Hansen, Inria, IP Paris





Anne Auger and Nikola



Data and Performance Profiles

Anne Auger and Nikolaus Hansen, Inria, IP Paris



Data Profile

certain target on a problem $p \in \mathscr{P}$.

The data profile is the ECDF of $\{T_{p,s}/(n+1), p \in \mathscr{P}\}$:

Benchmarking Derivative-Free Optimization Algorithms by J. Moré and S. Wild. SIAM J. Optimization, Vol. 20 (1), pp.172-191, 2009.

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•

Given $T_{p,s}$ a collection of runtime (#of f-evals) for a solver s to reach a

Normalization is done because runtime associated to different dimensions are put together



Data Profile

Given $T_{p,s}$ a collection of runtime (#of f-evals) for a solver s to reach a certain target on a problem $p \in \mathscr{P}$.

The data profile is the ECDF of {

 $\mathbf{ECDF}_{\{T_{p,s}/(n+1), p \in \mathcal{G}\}}$

Benchmarking Derivative-Free Optimization Algorithms by J. Moré and S. Wild. SIAM J. Optimization, Vol. 20 (1), pp.172-191, 2009.

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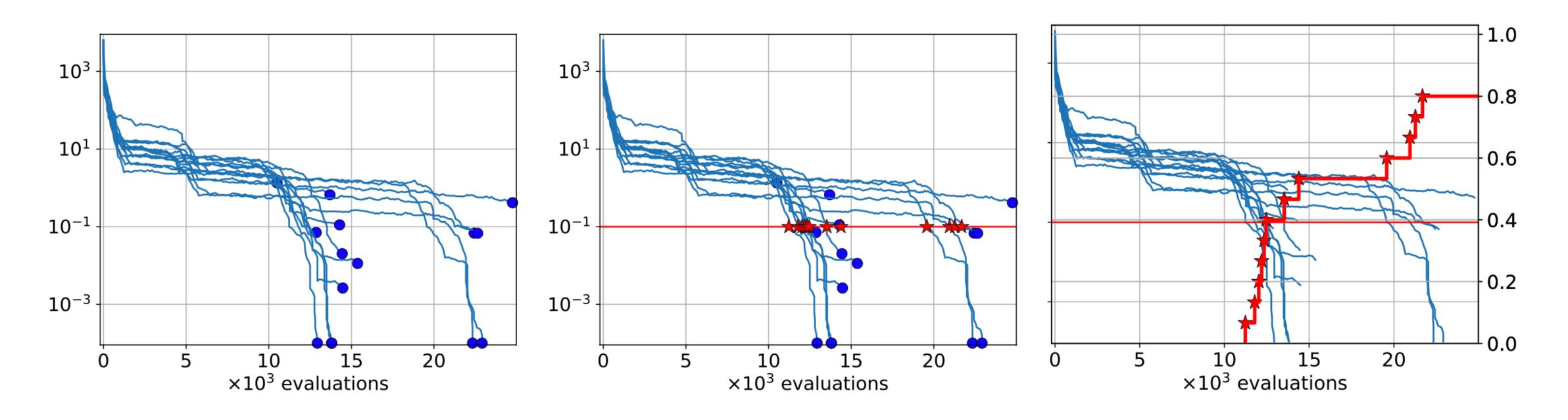
$$\{T_{p,s}/(n+1), p \in \mathcal{P}\}$$
:

Normalization is done because runtime associated to different dimensions are put together

$$\mathcal{P}_{f}(\mathbf{t}) = \frac{1}{|\mathcal{P}|} \sum_{p=1}^{|\mathcal{P}|} 1_{\left\{\frac{T_{p,s}}{n+1} \leq t\right\}}$$



Data Profile



target or shifting the respective graph vertically is the same

Targets may be different for each function, but choosing a different



Performance Profile

set of solvers \mathcal{S}

 $r_{p,s} = -$ m

The performance profile of a solver s is the ECDF of $\{r_{p,s}, p \in \mathscr{P}\}$:

$$\mathrm{ECDF}_{\{r_{p,s}, p \in \mathcal{P}\}}(\mathsf{t}) = \frac{1}{|\mathcal{P}|} \sum_{p=1}^{|\mathcal{P}|} \mathbf{1}_{\{r_{p,s} \leq t\}}$$

E. D. Dolan and J. J. Moré, Benchmarking optimization software with performance profiles, Math. Program., 91 (2002), pp. 201–213.

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Normalize runtime by performance of best solver: Define the performance on a problem p by a solver s as the runtime divided by the runtime of best solver among a

$$T_{p,s}$$
$$in\{T_{p,s}: s \in \mathcal{S}\}$$

It "Removes" the order of magnitude of $T_{p,s}$ and thus the information of difficulty





Data and Performance Profile: Discussion

- Performance and Data profiles are just ECDF of (normalized) runtime associated to a single target per problem
- Performance profile
 - normalized by the smallest (best) runtime
 - relative to the set of solvers benchmarked

 we do not see the problem difficulty anymore: normalization removes absolute value

difficult to compare across papers





Aggregation of Data

is necessary ullet

> e.g., BBOB takes about 24 x 15 x 30 \approx 10,000 single measurements for each algorithm in each dimension

- over different values
- any runtimes can be meaningfully aggregated
- successful and unsuccessful runs can be meaningfully aggregated,

 implicit assumption: uniform distribution over all problems we aggregate over shall somewhat reflect the problem distribution in reality

properties that can be inexpensively probed should not (never) be aggregated

For example: dimensionality. Why?

Assuming they come in the same unit of measurement (here evaluations). However: not all ways to aggregate runtimes are meaningful. We should use a log scale when they come from different distributions.

solving the fast vs successful comparison "dilemma" once and for all. Using simulated restarts or Enes/ERT/SP2, see "Treating success probabilities".



Aggregation of Data

1. aggregating repetitions over the same (or very similar) problem(s) in particular with unsuccessful trials

2. aggregating data from different problems

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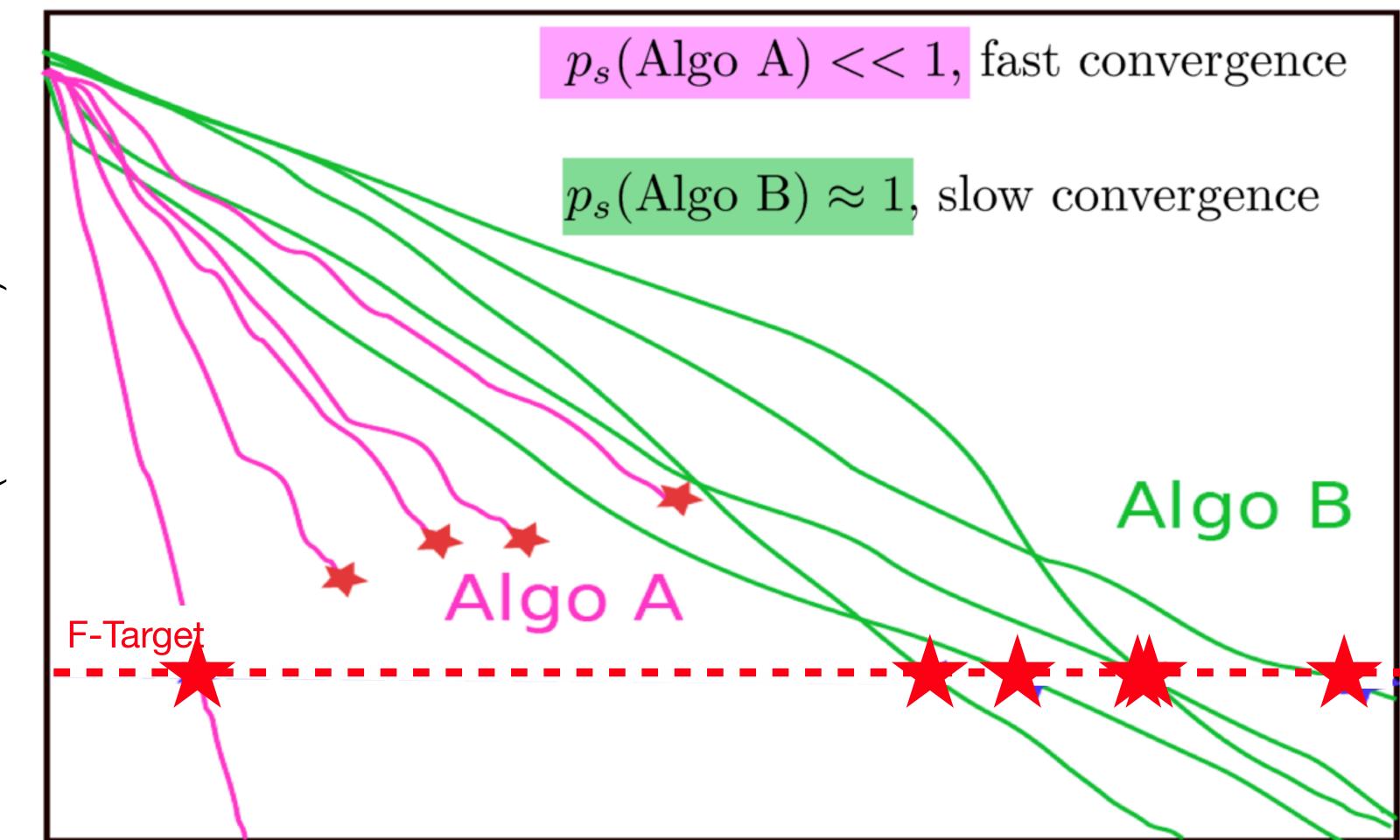
Expected RunTime (ERT)

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Aggregated measurement of repetitions



Treating Success Probabilities Solving the fast-versus-successful comparison dilemma



function (or indicator) value

number of evaluations



Treating Success Probabilities Solving the fast-versus-successful comparison dilemma

(artificial) restarts using the given independent runs

Algo Restart B:

Algo Restart A:

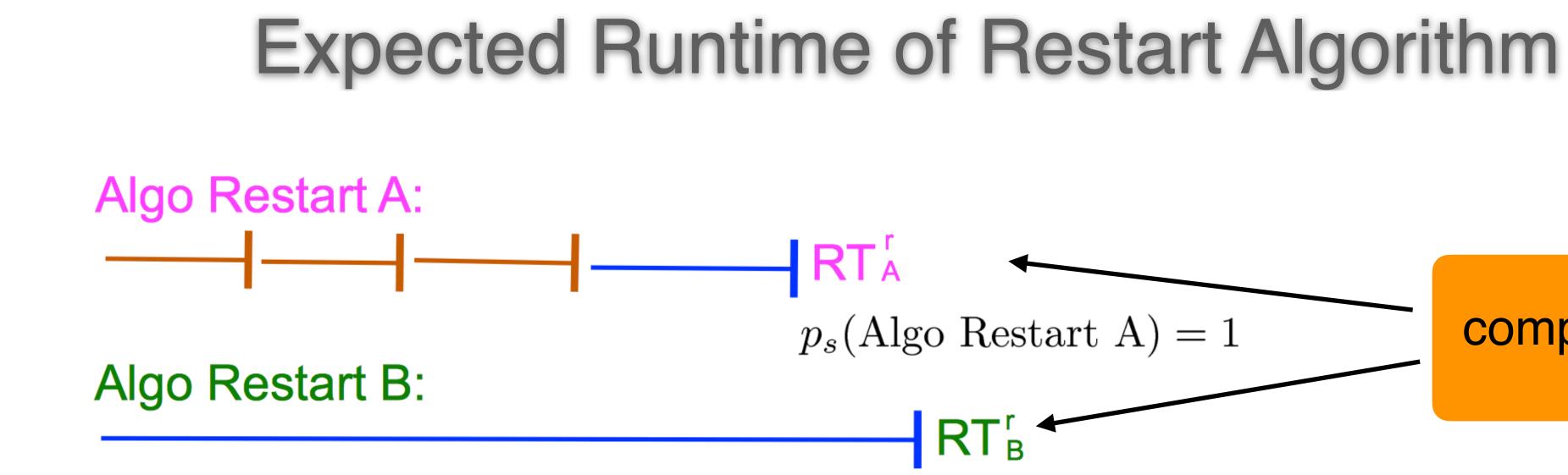
Caveat: the performance of algorithm A critically depends on termination methods (before to hit the target) which reflects the situation on a practical problem unless many runs can be done in parallel

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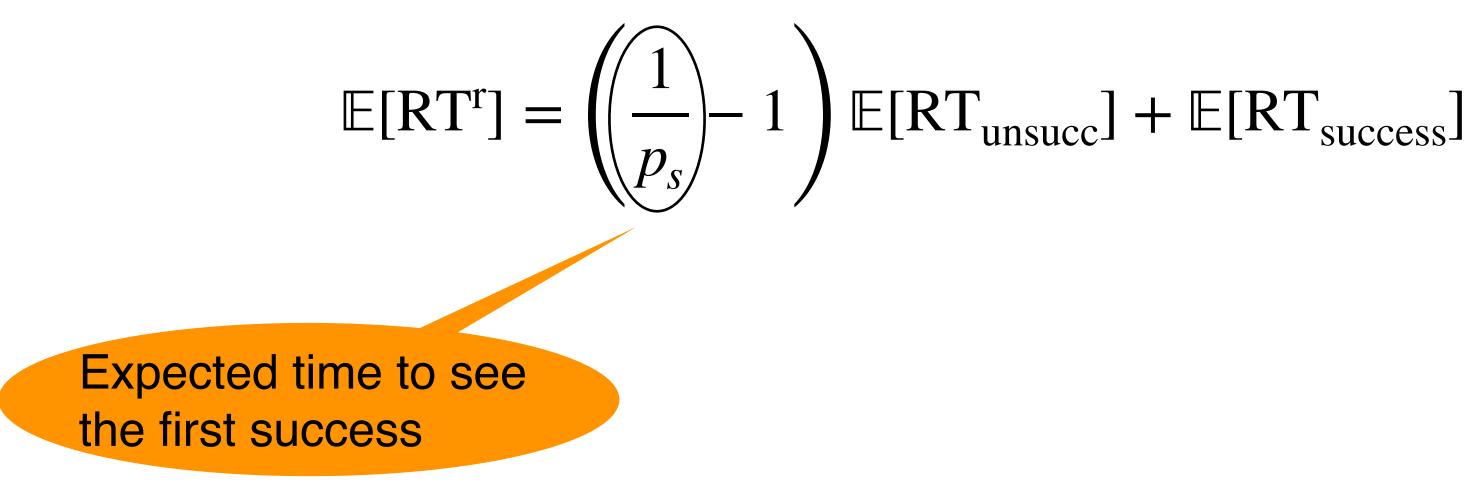
We can **simulate a runtime distribution** by simulated

$p_s(\text{Algo Restart A}) = 1$ **RT**^r_B $p_s(\text{Algo Restart B}) = 1$





Expected Runtime of Restart Algorithm:



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comparable runtimes

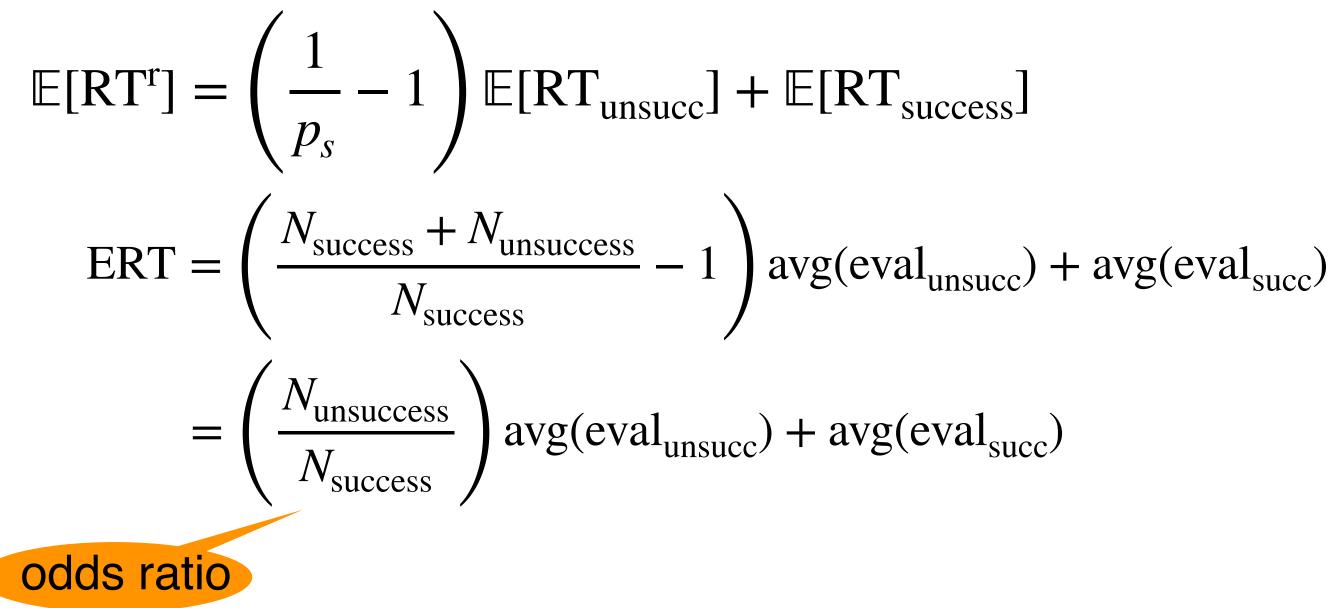
 $p_s(\text{Algo Restart B}) = 1$





Expected RunTime - ERT

Expected runtime (ERT, aka Enes, SP2, aRT) estimates $\mathbb{E}[RT]$



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 $ERT = \frac{\text{#evaluations(until to hit target or stop)}}{\text{#successes}}$

unsuccessful runs count (only) in the nominator

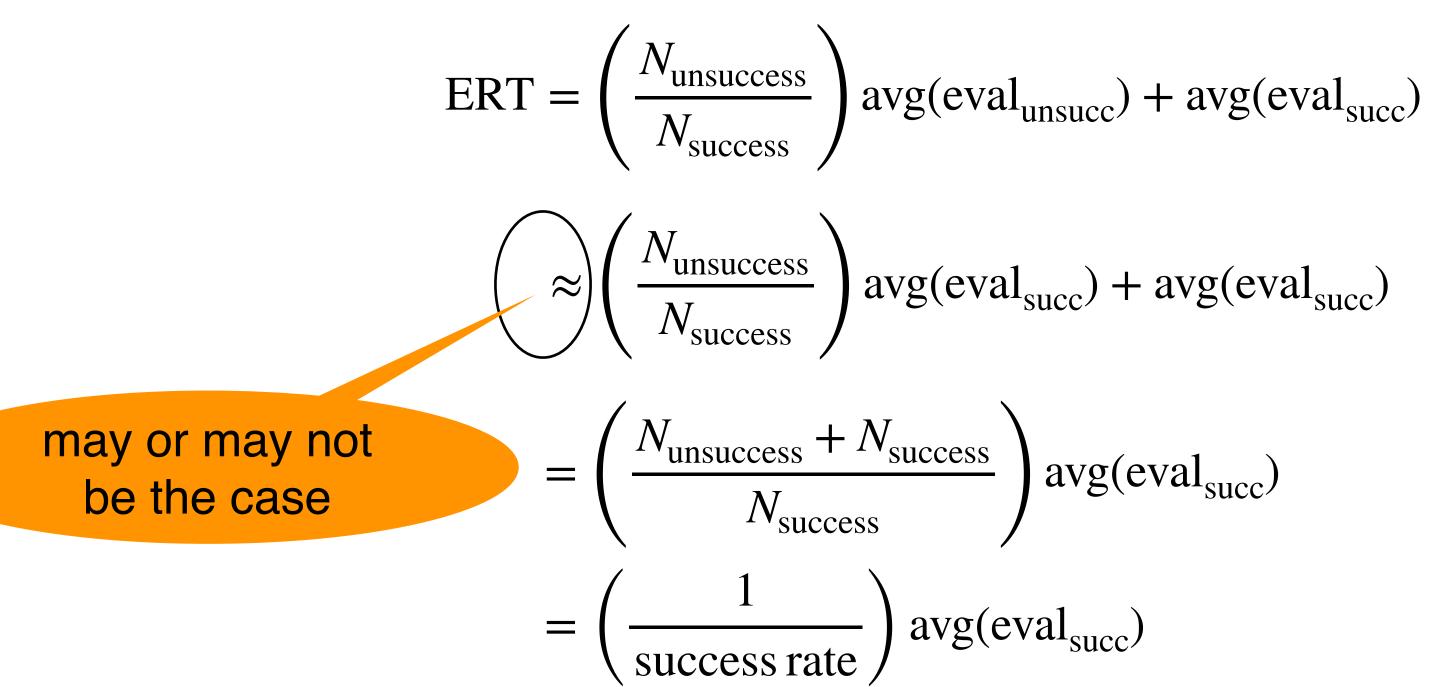
defined (only) for #successes > 0







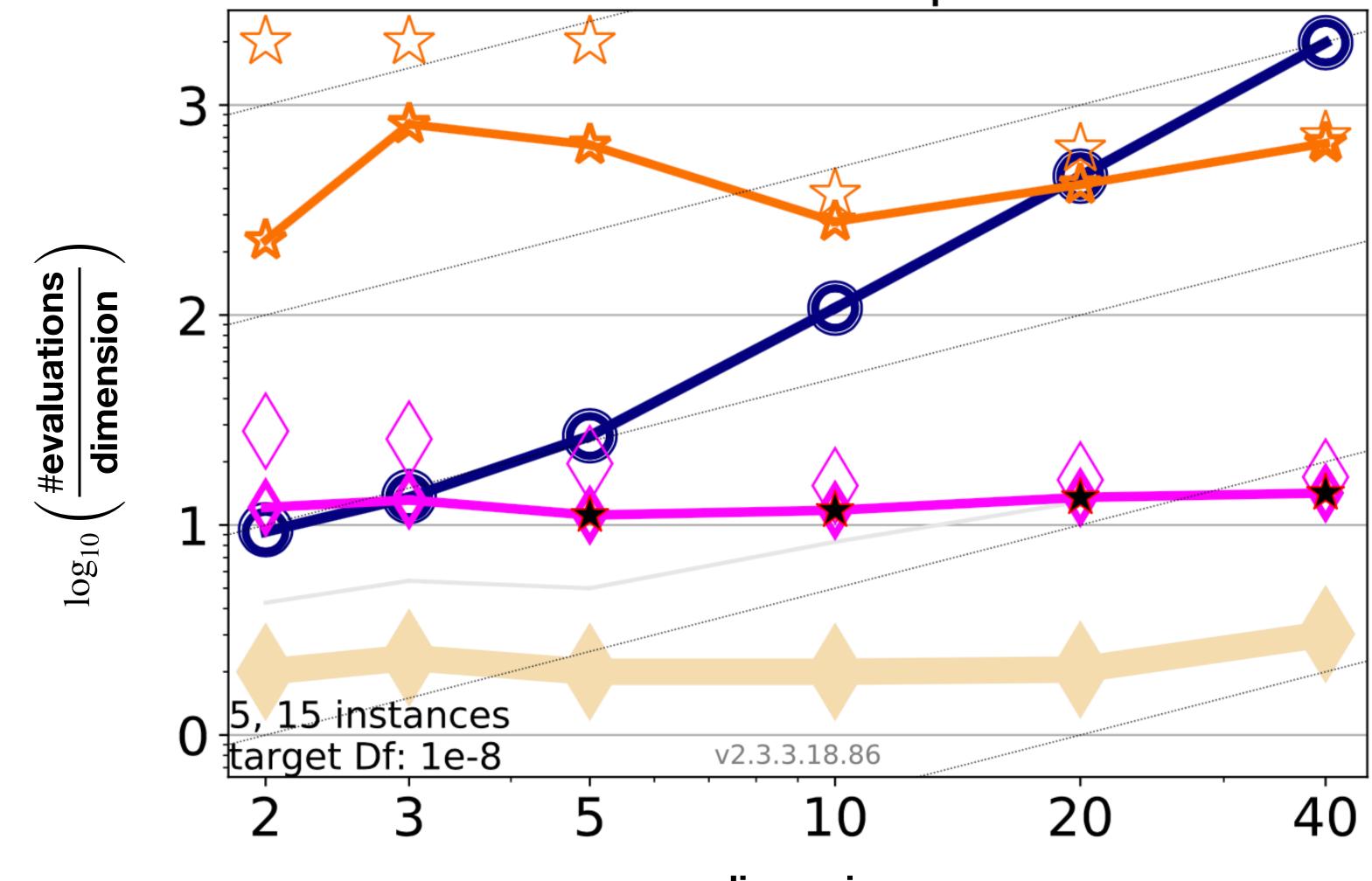
ERT Related Performance Measures



The last three lines are AKA Q-measure or SP1 (success performance). See [Price 1997] and [Auger&Hansen 2005]

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On Scaling

dimension



Aggregation of Data

1. aggregating repetitions over the same (or very similar) problem(s) in particular with unsuccessful trials

2. aggregating data from different problems

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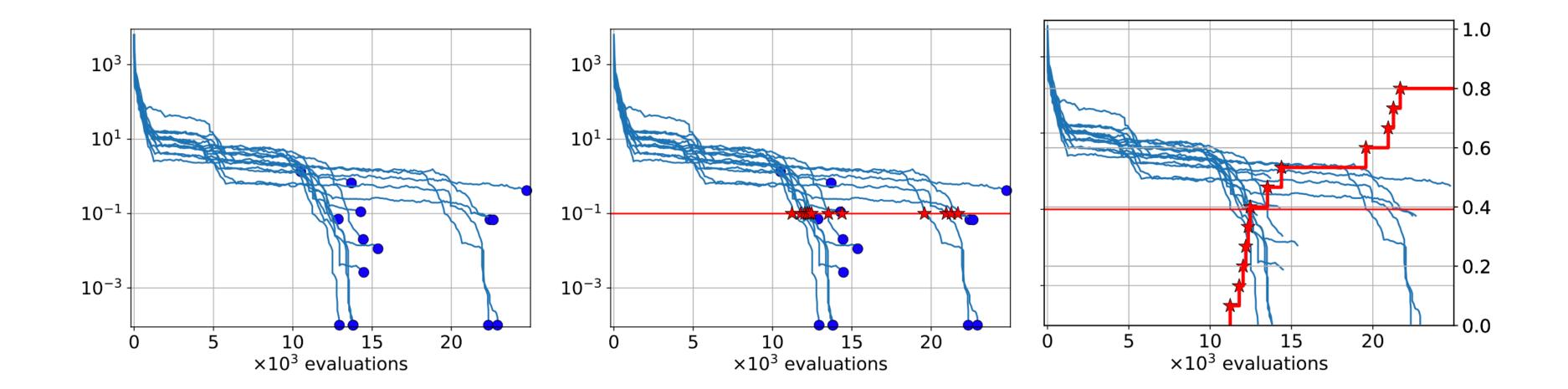
already a single convergence graph contains different problems



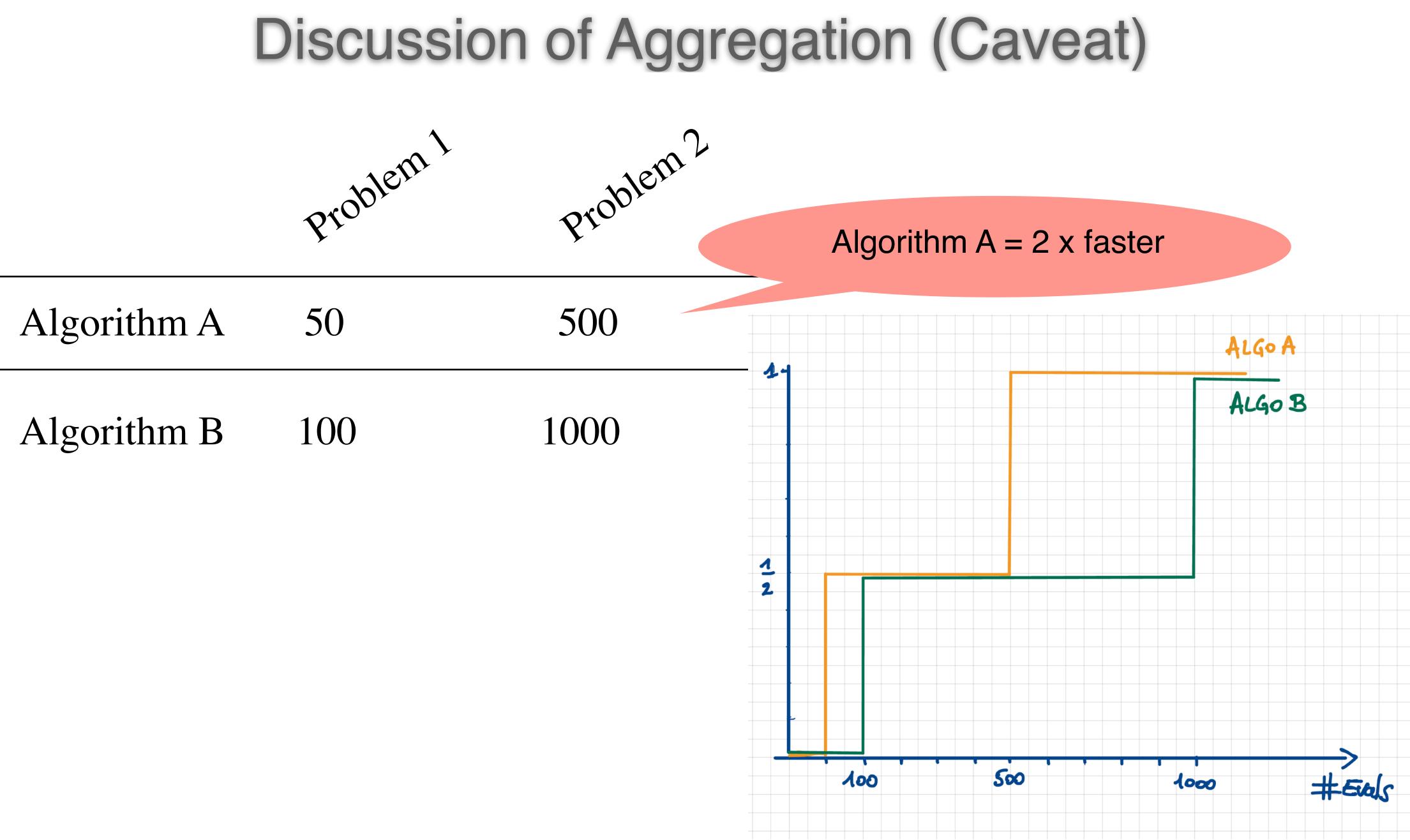
Aggregation of Data: ECDFs With Different Problems

• ECDFs (re-)order the data (sort the data)

hence we lose the problem label single convergence graph ECDFs are not affected

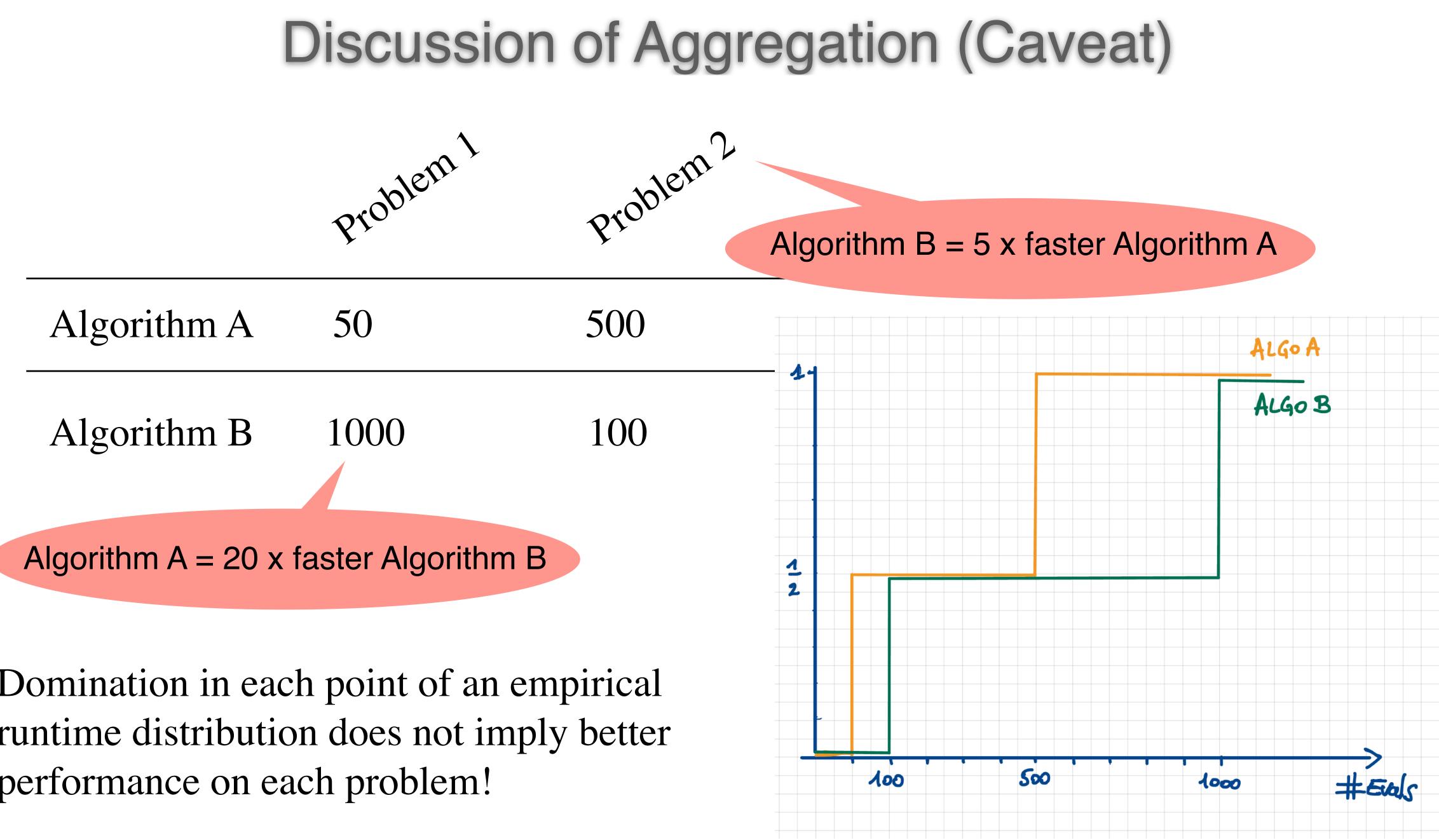






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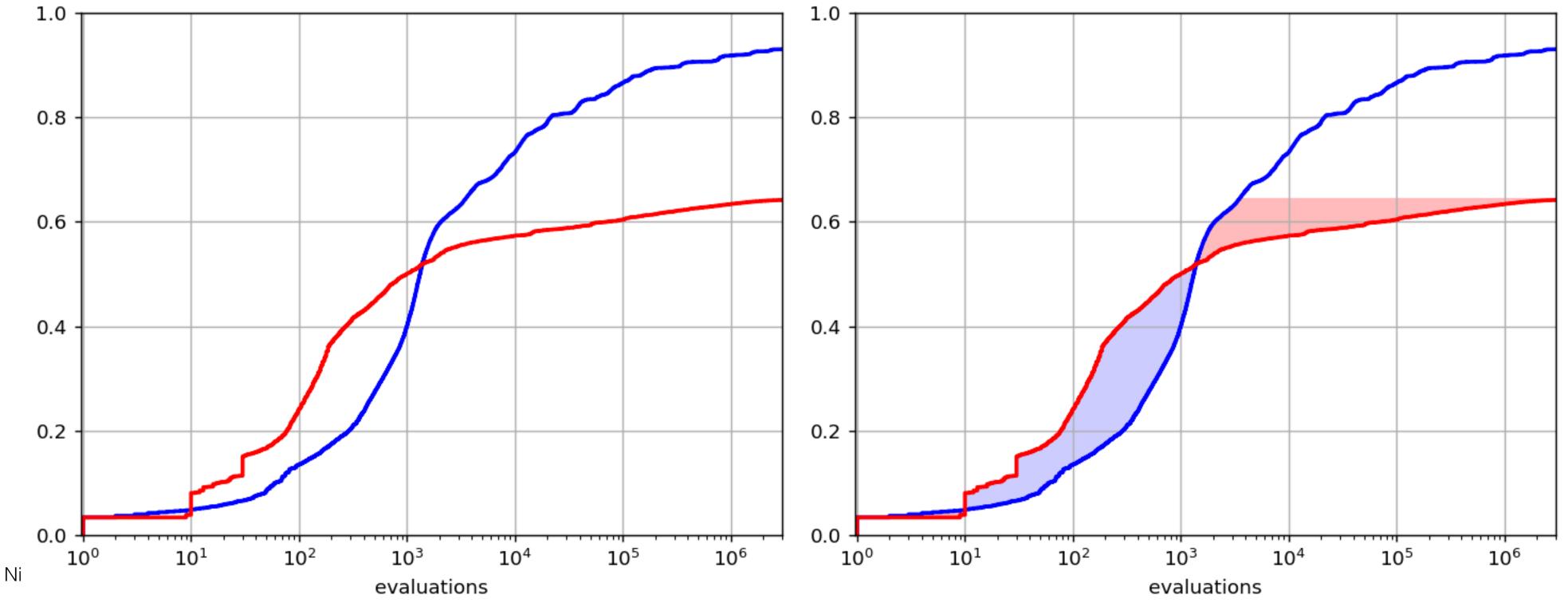
Domination in each point of an empirical runtime distribution does not imply better performance on each problem!



The average runtime ratio

$$\frac{\exp\left(\frac{1}{k}\sum_{i}\log(B_{i})\right)}{\exp\left(\frac{1}{k}\sum_{i}\log(A_{i})\right)} = \exp\left(\frac{1}{k}\sum_{i}\log(B_{i}) - \frac{1}{k}\sum_{i}\log(A_{i})\right) = \exp\left(\frac{1}{k}\sum_{i}\log\left(\frac{B_{i}}{A_{i}}\right)\right)$$

is the area between the runtime distribution graphs of two algorithms A,B is the geometric average when the x-axis is in log-scale with the geometric average it is invariant under the exchange of operators: the ratio of the averages equals the average of the ratios



Anne Auger and Ni



Take Home Messages

Select a balanced testbed

- Use quantitative measurements
- Don't aggregate over attributes that are simple to determine •
- Benchmarking is tedious but necessary •

furious activity is no substitute for understanding using "all functions" is likely to introduce a bias (too many simple or low dimensional problems)

which should preferably be comparable across publications empirical CDFs are a very useful tool

like dimension

use a provided software platform?



Statistical Analysis

"The first principle is that you must not fool yourself, and you are the easiest person to fool. So you have to be very careful about that. After you've not fooled yourself, it's easy not to fool other[scientist]s. You just have to be honest in a conventional way after that."

- Statistical analysis is secondary!
- small p-value, HARKing, hypothesis generating)

- Richard P. Feynman

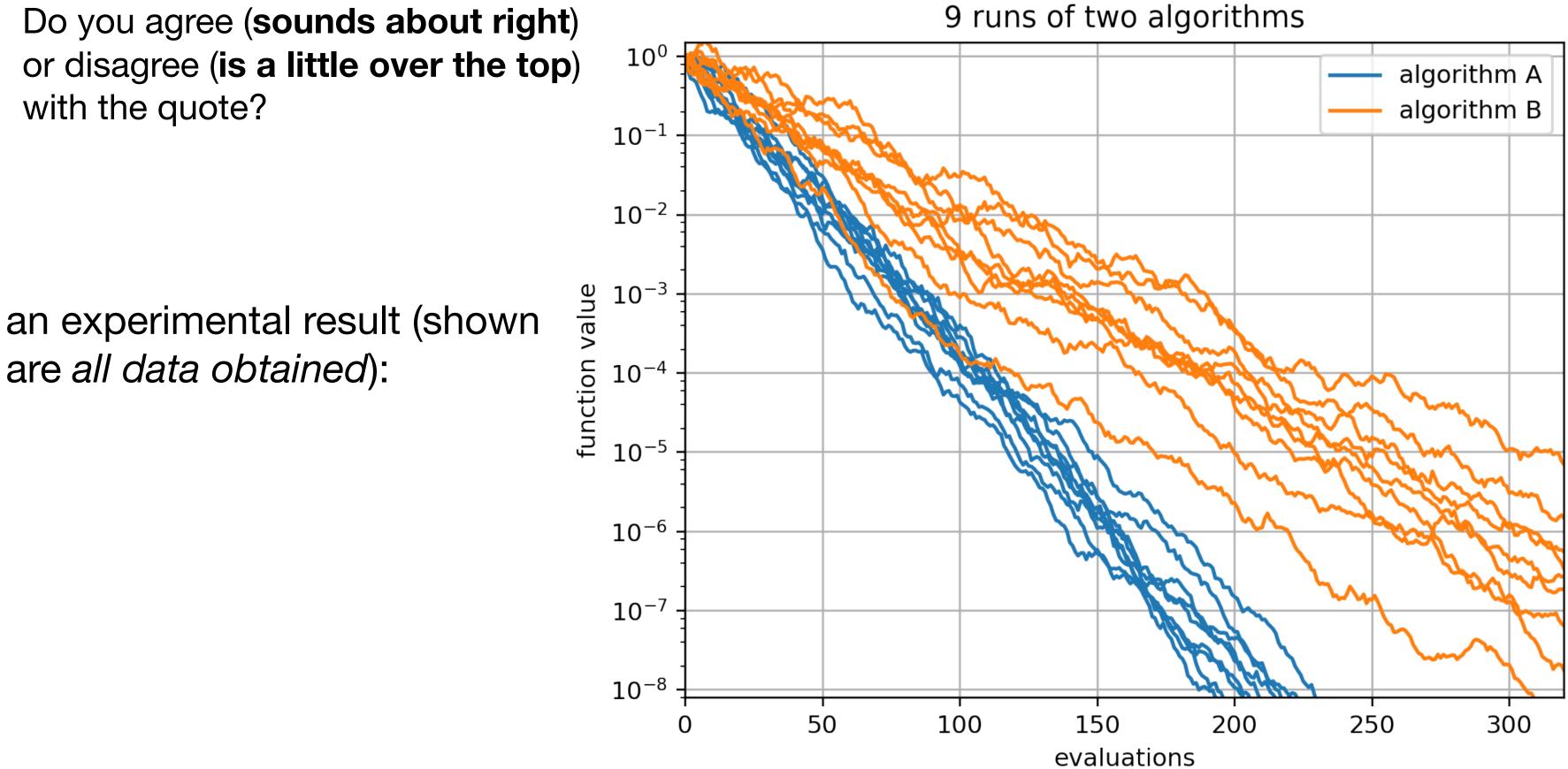
useful only after a (quantitative) conclusion has been drawn from the data.

• Don't confuse statistical (significance) testing (hypothesis testing) with statistical data exploration (searching for which test of many has a



Statistical Analysis

"[...] experimental results lacking proper statistical analysis must be considered anecdotal at best, or even wholly inaccurate."

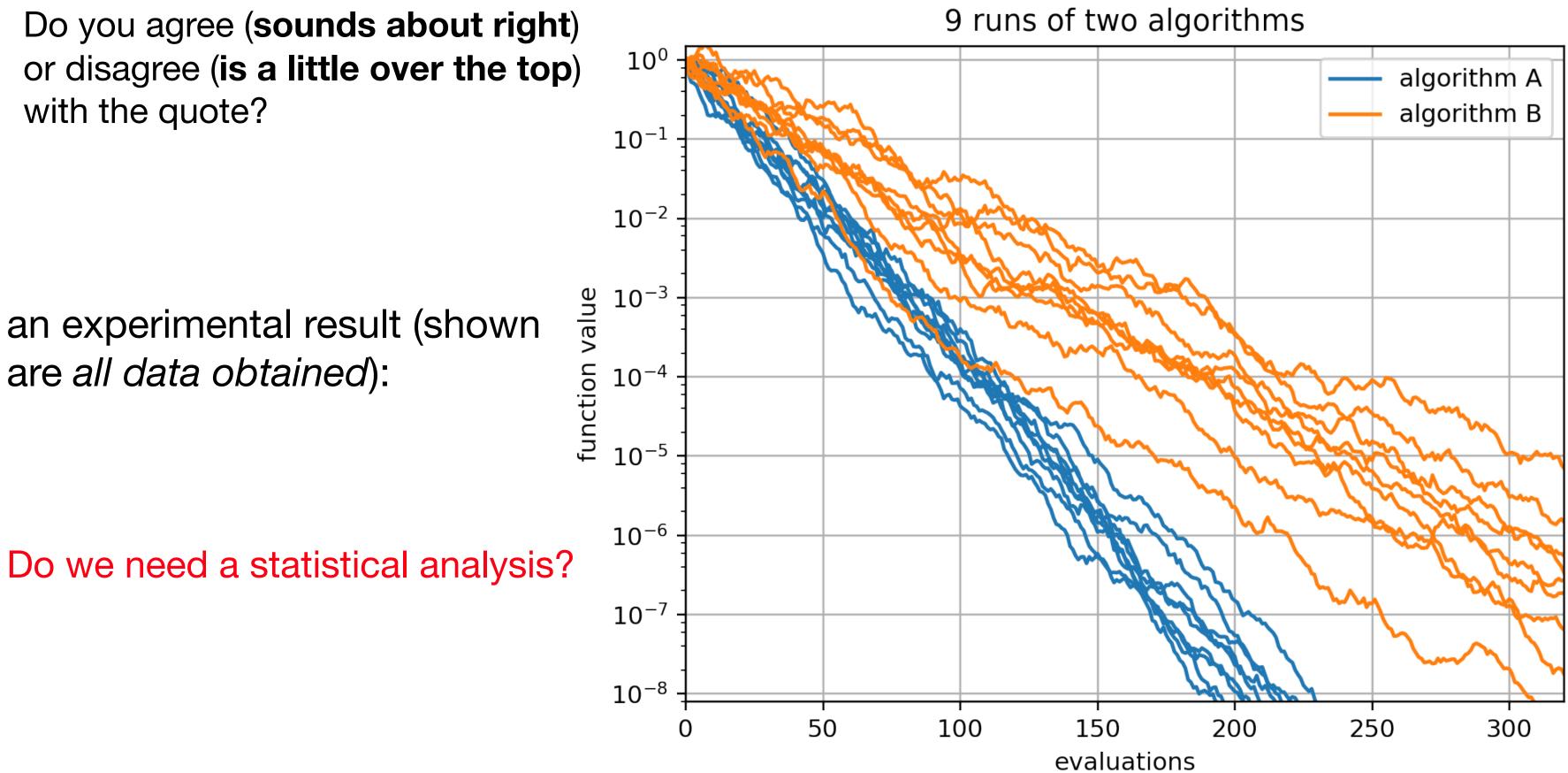


— M. Wineberg, 2016



Statistical Analysis

"[...] experimental results lacking proper statistical analysis must be considered anecdotal at best, or even wholly inaccurate."



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— M. Wineberg, 2016



What about the *p*-value?

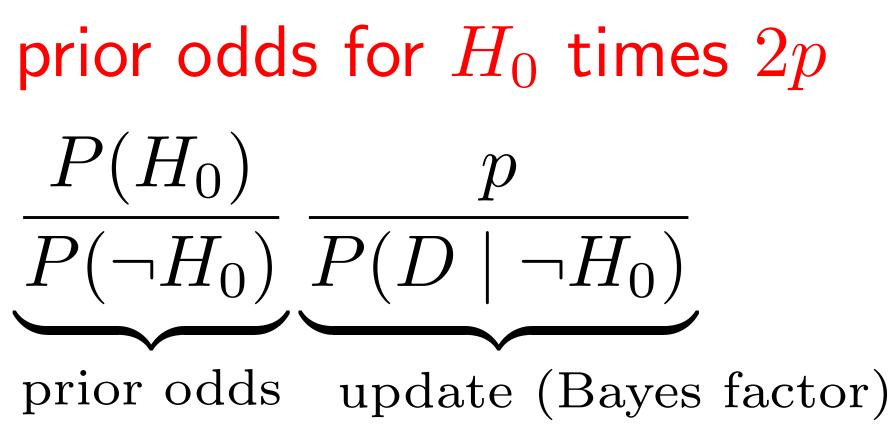
The "p-value of statistical significance" is the probability that we will observe a "false positive" outcome, $P(D \mid H_0)$ i.e., the probability that the observed (or more extreme "favorable") data D would be observed under the null hypothesis H₀ "by chance" (while in reality there is no "effect" to be seen).

It is straight forward to compute the posterior odds that H_0 is true from the observed p-value:

posterior odds for
$$H_0 \approx$$

$$\frac{P(H_0 \mid D)}{P(\neg H_0 \mid D)} =$$
posterior odds

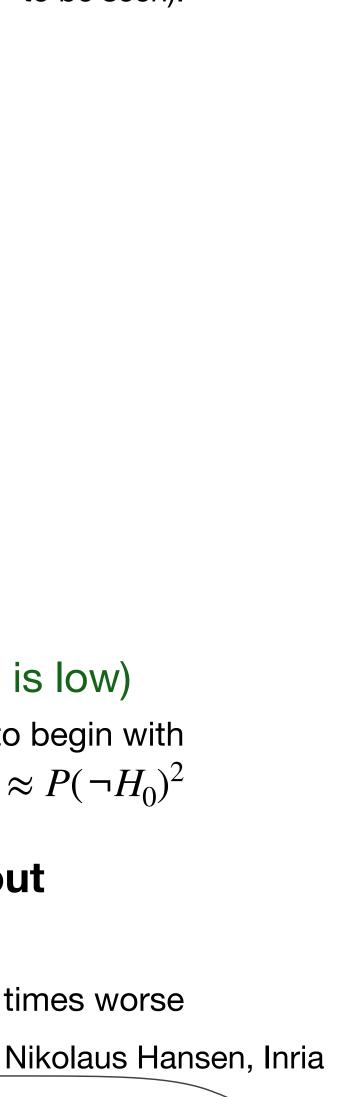
whether rejecting H_0 is a reasonable decision



• if we try to improve a state-of-the-art algorithm, the prior odds of H_0 is usually high (the odds of success is low) as a rule of thumb, the higher the prior odds for H_0 , the more interesting is the scientific question to begin with to accept $\neg H_0$ with the same confidence we had before in H_0 , we need $p \approx P(\neg H_0)^2$

• if we are not willing to estimate the prior odds of H_0 , we are not justified to conclude anything about

we can only conclude that the odds for H_0 became about 2p times worse



Statistical Testing: General Prodecure

- which is tested for statistical significance
- we prefer "nonparametric" methods
- •
- given H0 is true = probability "H0 is rejected" given H0 is true
- *given* an observed p-value, *fewer data are better*

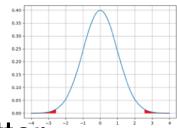
first, check *the relevance* of the result, for example, of the difference

this also means: preferably do not *explorative testing* (e.g. test *all* pairwise combinations), any ever so small difference can be made *statistically* significant with a simple trick, but *not made* significant in the sense of *relevant* or *important* or *meaningful*

not assuming that the statistics come from a *parametrised* family of probability distributions

Null hypothesis (H0) = both/all data come from the same distribution

p-value = significance level = probability of a *false* positive outcome

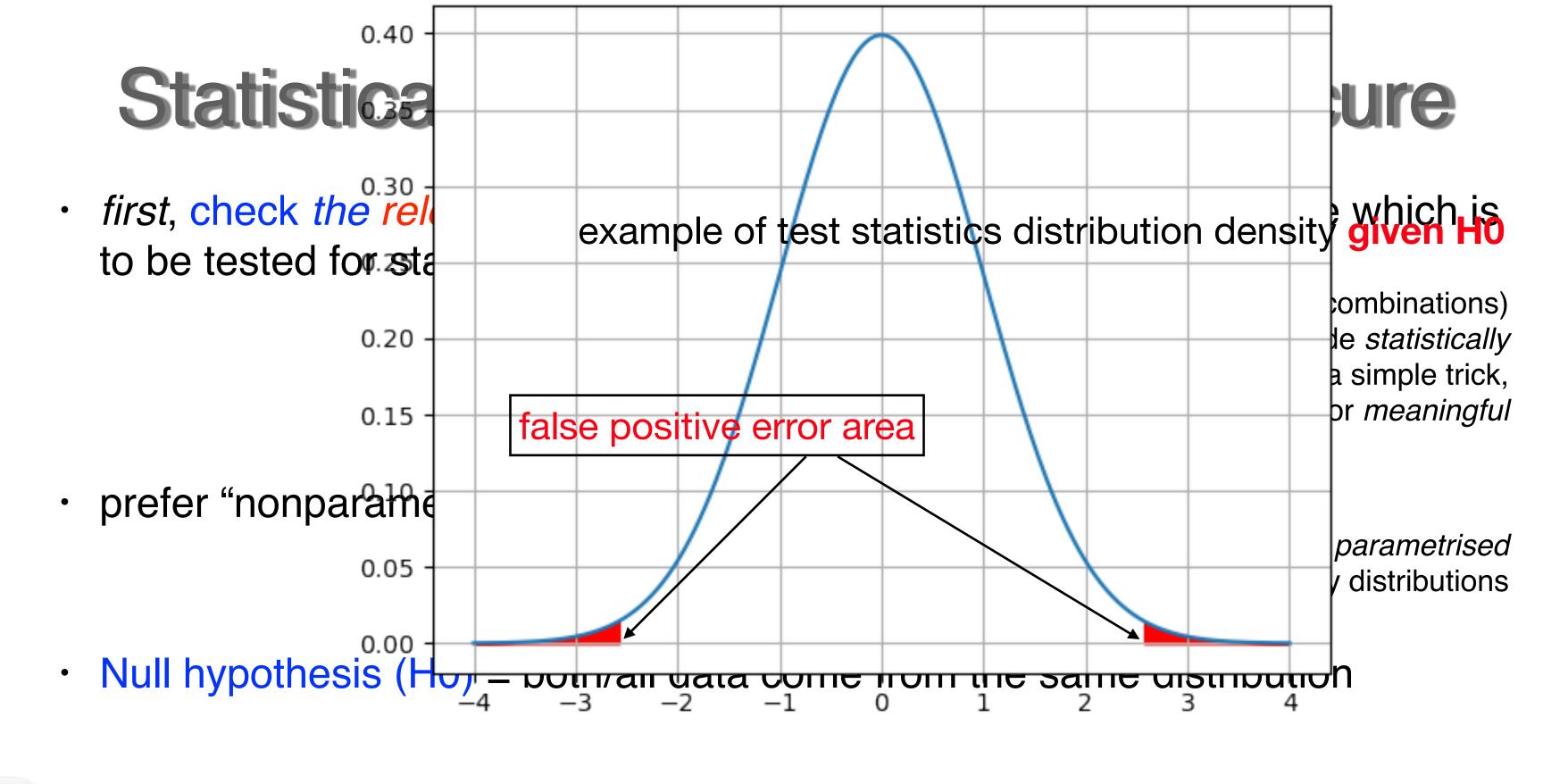


smaller p-values are better

obsolete practice: <0.1% or <1% or <5% is usually considered as *statistically significant*

more data (almost inevitably) lead to smaller p-values, hence to achieve the same p-value with fewer data, the *between*-difference must be larger compared to the *within*-variation





H0 is true = probability H0 is rejected given H0 is true

• *given* a found/observed p-value, *fewer data are better*

p-value = significance level = probability of a *false* positive outcome given

smaller p-values are better <0.1% or <1% or <5% is usually considered as *statistically significant*

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Statistical Testing: General Prodecure

- to be tested for statistical significance
- prefer "nonparametric" methods
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given an observed p-value, *fewer data are better*

first, check *the relevance* of the result, for example of the difference which is

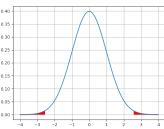
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Statistical Testing: Methods

- Whitney U test)
 - values are observed

The "*lack" of necessary preconditions* is the main reason to use the rank-sum test. even a few equal values are not detrimental

the rank-sum test is *nearly as efficient* as the t-test which requires certain distributions for empirical mean and variance

for discrete data with ties: scipy.stats.mannwhitneyu(..., alternative='two-sided')

- Outcome: a *p*-value

the probability that the observed or a more extreme data set was generated under the null hypothesis; the probability to *mistakenly* reject the null hypothesis

we use the rank-sum test (scipy.stats.ranksums, aka Wilcoxon or Mann-

Assumption: all observations (data values) are obtained independently and no equal

Null hypothesis (nothing relevant is observed if): Pr(x < y) = Pr(y < x)

H0: the probability to be greater or smaller (better or worse) is the same the aim is to be able to reject the null hypothesis

Procedure: computes subsums of ranks in the ranking of all (combined) data values $Alg1 = [400, 422, 440] vs Alg2 = [444, 490, 555] \implies ranks: Alg1 = [1, 2, 3] vs Alg2 = [4, 5, 6]$



Statistical Testing: Methods

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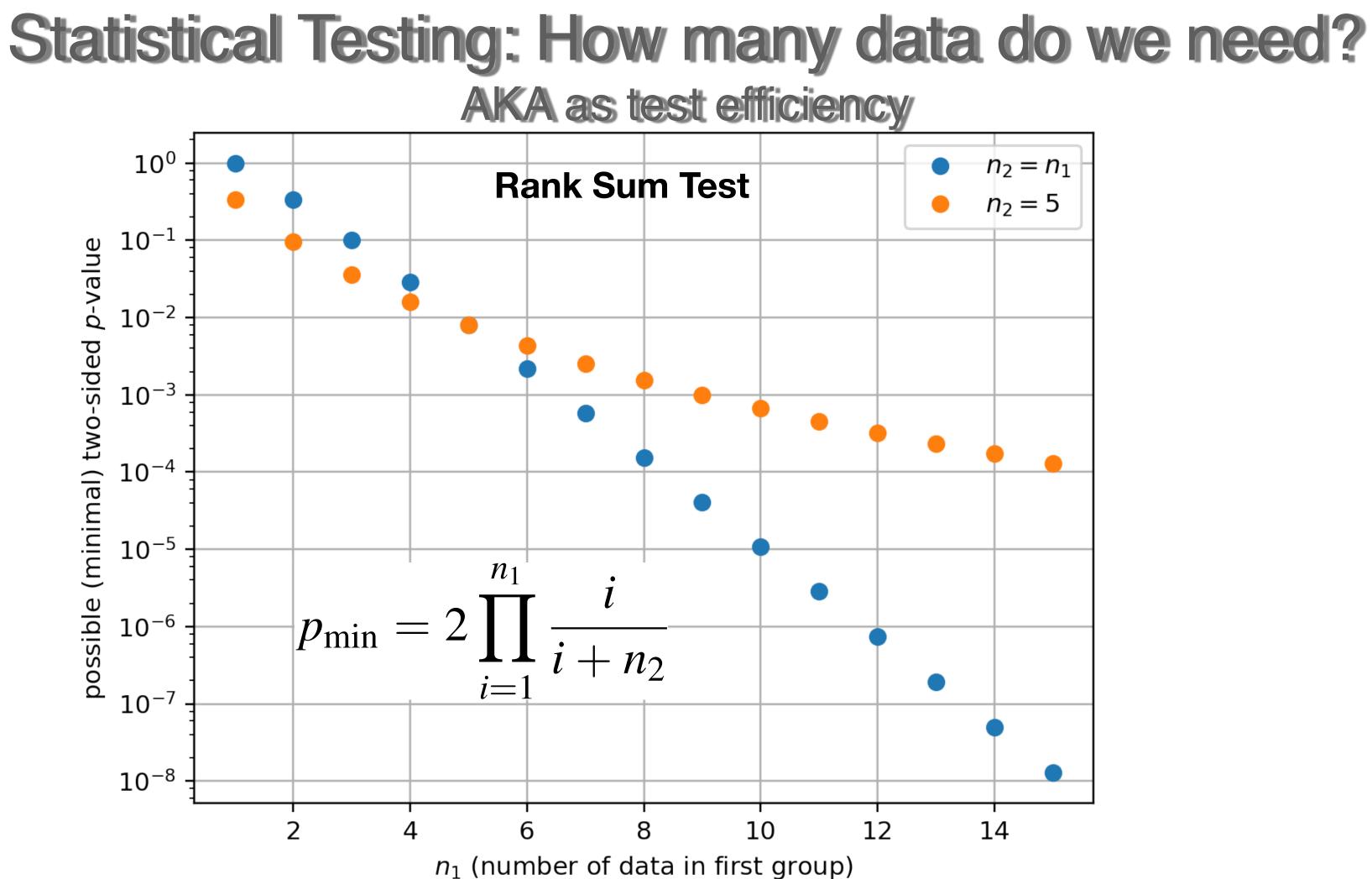
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 $Alg1 = [400, 422, 440] vs Alg2 = [444, 490, 555] \implies ranks: Alg1 = [1, 2, 3] vs Alg2 = [4, 5, 6]$







- assumption: data are fully "separated", that is,
- use the Bonferroni correction for multiple tests

 $\forall i, j : x_i < y_j \text{ or } \forall i, j : x_i > y_j \text{ (two-sided)}$ observation: adding 2 data points in each group gives about one additional order of magnitude

simple and conservative: multiply the computed p-value by the number of tests



Statistical Testing: How many data do we need?

is plenty

• We often take two times 11 or 31 or 51

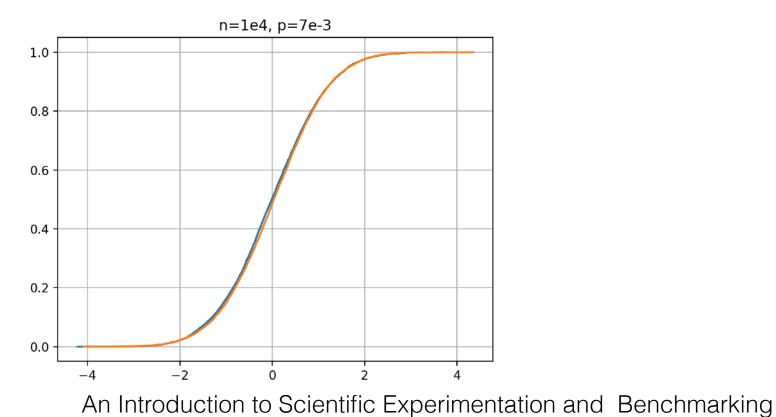
Too many data make statistical significance meaningless

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In the best case: at least ten (two times five) and two times nine

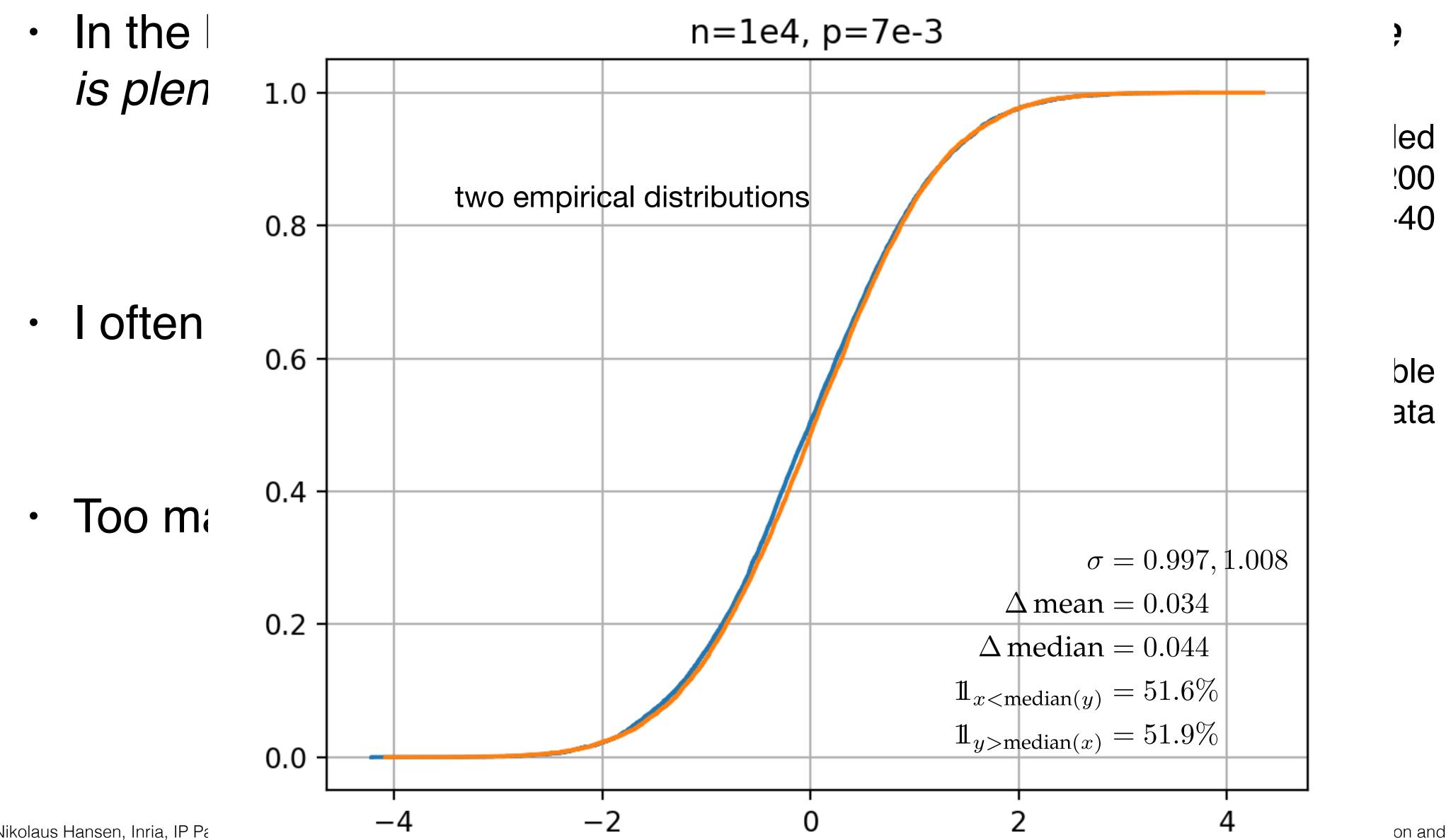
minimum number of data to possibly get two-sided *p* < 1%: 5+5 or 4+6 or 3+9 or 2+19 or 1+200 and p < 5%: 4+4 or 3+5 or 2+8 or 1+40

median, 5%-tile and 95%-tile are easily accessible with 11 or 31 or 51... data





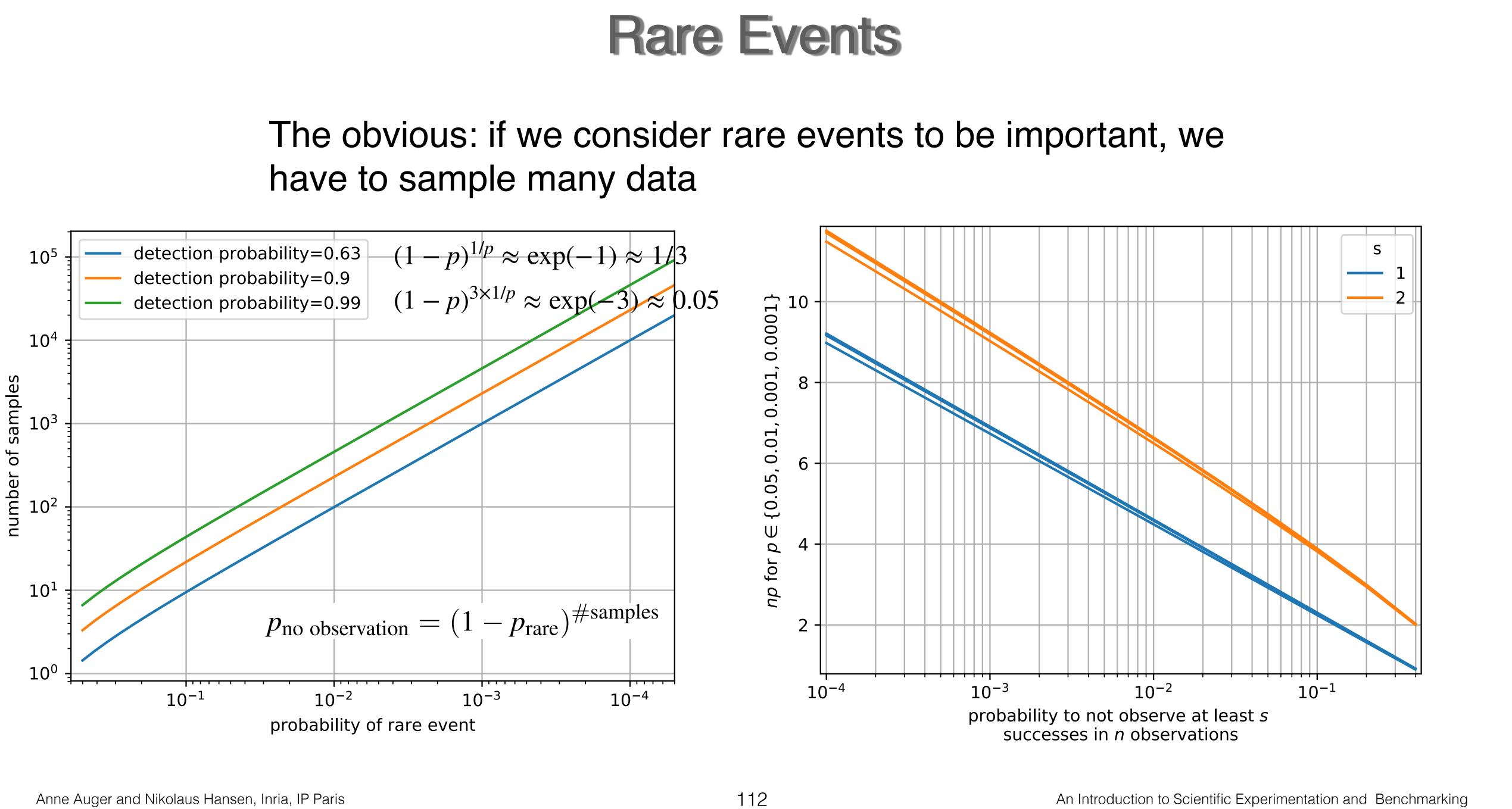
Statistical Testing: How many data do we need?



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have to sample many data





```
import scipy.stats
1
 p1, n1 = 9, 90 # 10% success
2
 p2, n2 = 20, 100 # 20% success
3
  scipy.stats.chi2_contingency([[p1, n1 - p1],
4
5
```

(2.930076628352492) χ^2 -statistics 0.0869433657247775 p-value [[13.73684211 76.26315789][15.26315789 84.73684211])

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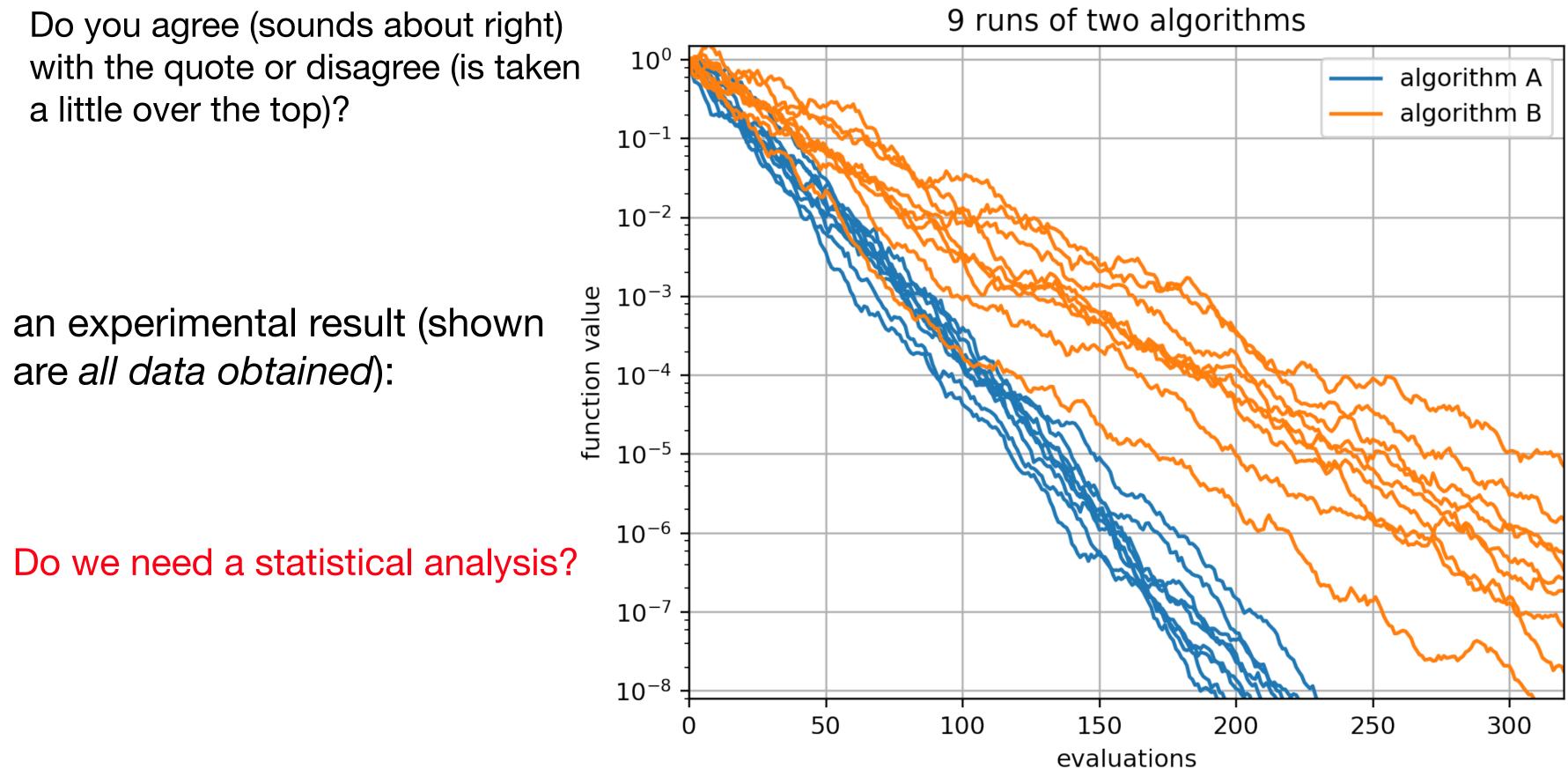
Testing Frequencies

[p2, n2 – p2]])



Statistical Analysis

"[...] experimental results lacking proper statistical analysis must be considered anecdotal at best, or even wholly inaccurate."



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— M. Wineberg, 2016



Using COCO

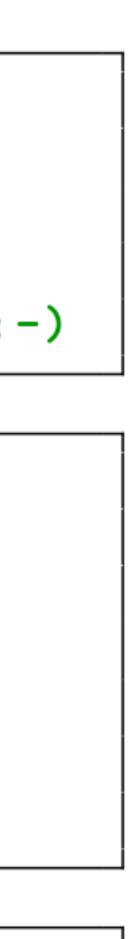
Running an experiment

\$ ### get and install the code
\$ git clone https://github.com/numbbo/
\$ cd coco
\$ <pre>python do.py run-python # install H</pre>
\$ <pre>python do.py install-postprocessing</pre>

```
#!/usr/bin/env python
"""Python script to benchmark fmin of scipy.optimize"""
from __future__ import division # not needed in Python 3
import cocoex, cocopp # experimentation and post-processing modules
import scipy.optimize # to define the solver to be benchmarked
### input
suite_name = "bbob"
output folder = "scipy-optimize-fmin"
```

/coco.git # get coco using git

Python experimental module cocoex install-user # install postprocessing :-)



#!/usr/bin/env python """Python script to benchmark fmin of scipy.optimize""" from __future__ import division # not needed in Python 3 import cocoex, cocopp # experimentation and post-processing modules import scipy.optimize # to define the solver to be benchmarked

input suite_name = "bbob" output_folder = "scipy-optimize-fmin" fmin = scipy.optimize.fmin budget_multiplier = 2 # increase to 10, 100, ...

prepare

suite = cocoex.Suite(suite_name, "", "") observer = cocoex.Observer(suite_name, "result_folder: " + output_folder)

go

for problem in suite: # this loop will take several minutes or longer problem.observe_with(observer) # will generate the data for cocopp # restart until the problem is solved or the budget is exhausted while (not problem.final_target_hit and problem.evaluations < problem.dimension * budget_multiplier): fmin(problem, problem.initial_solution_proposal()) # we assume that 'fmin' evaluates the final/returned solution



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post-process data cocopp.main(observer.result_folder) # re-run folders look like "...-001" etc

observer = cocoex.Observer(suite_name, "result_folder: " + output_folder)



Selecting algorithms for comparison

Using COCO



Home - coco-data-archive

numbbo.github.io/data-archive/

coco-data-archive

Q Search coco-data-archive

× +

Home

bbob data archive

bbob-noisy data archive

bbob-biobj data archive

bbob-largescale data archive

bbob-mixint data archive

COCO data archives

Welcome to the website of the benchmarking data archives of the COCO (Comparing Continuous Optimizers) platform.

Besides providing the official archives from numerical benchmarking experiments, run on the COCO platform and available through its cocopp Python module, this website also provides lists of these official data sets with additional information like authors, links to papers, source code etc. Right now, data sets for the following test suites are available:

- bbob: 24 single-objective noiseless functions
- bbob-noisy: 24 single-objective noisy functions •
- bbob-biobj: 55 bi-objective functions •
- bbob-largescale: large-scale version of the 24 bbob functions (dimension up to 640)
- bbob-mixint: mixed-integer versions of the 24 bbob functions

Related links

- postprocessed data of these archives for browsing
- how to submit a data set
- how to create and use COCO data archives with the cocopp.archiving Python module

This site uses Just the Docs, a documentation theme for Jekyll. Q **公** M

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Using COCO

🌽 bbob data archive - coco-data- 🗙 🕂

numbbo.github.io/data-archive/bbob/

Q Search coco-data-archive

Home

bbob data archive

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bbob-biobj data archive

bbob-largescale data archive

coco-data-archive

bbob-mixint data archive

Algorithm data sets for the bbob test suite

In the first table below, you will find all official algorithm data sets on the bbob test suite, together with their year of publication, the authors, and related PDFs for each data set. Links to the source code to run the corresponding experiments/algorithms are provided whenever available.

A second table mentions data sets that have been collected on the bbob suite, but which are not complete in the sense that they miss at least one of the requested dimensions 2, 3, 5, 10, 20.

To sort the tables, simply click on the table header of the corresponding column.

	Number	Algorithm Name	Year	Author(s)	link to data	related PDFs, source code, a
	000	ALPS	2009	Hornby	data	pdf
	001	AMALGAM	2009	Bosman et al.	data	pdf noiseless - pdf noisy
	002	BAYEDA	2009	Gallagher	data	pdf noiseless - pdf noisy
	003	BFGS	2009	Ros	data	pdf noiseless - pdf noisy
This site uses Just the Docs, a documentation theme for Jekyll.	004	BIPOP-CMA-ES	2009	Hansen	data	pdf noiseless - pfd noisy



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Visit <u>https://</u> numbbo.github.io/dataarchive/



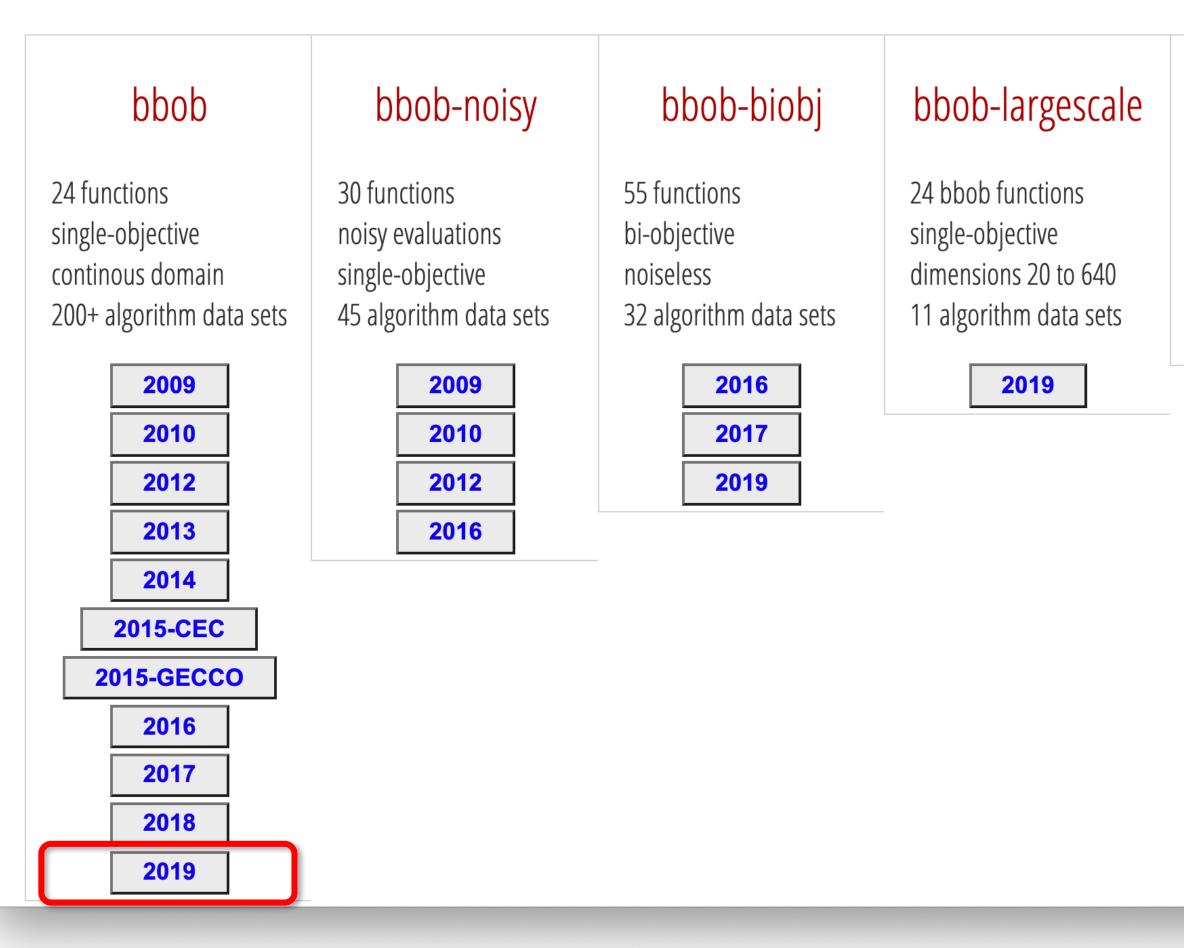


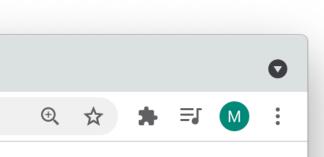


numbbo.github.io/ppdata-archive/

COCO ppdata-archive

This archive contains *postprocessed* data displaying benchmarking experiments of various numerical optimization algorithms on the various test suites provided by the Comparing Continuous Optimizers platform. Experiments are conducted in a blackbox setting and data are collected by year.





bbob-mixint

24 functions 80% discrete variables single-objective no official data yet

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