

DM811

Heuristics for Combinatorial Optimization

Lecture 15

Methods for Experimental Analysis

Marco Chiarandini

Department of Mathematics & Computer Science
University of Southern Denmark

Course Overview

1. Combinatorial Optimization, Methods and Models
2. General overview
3. Solver System and Working Environment
4. Construction Heuristics
5. Local Search: Components, Basic Algorithms
6. Local Search: Neighborhoods and Search Landscape
7. Efficient Local Search: Incremental Updates and Neighborhood Pruning
8. Stochastic Local Search & Metaheuristics
9. Methods for the Analysis of Experimental Results
10. Configuration Tools: F-race
11. Very Large Scale Neighborhoods

Examples: GCP, CSP, TSP, SAT, MaxIndSet, SMTWP, Steiner Tree, p-median, set covering

Outline

1. Experimental Analysis
2. Descriptive Statistics
 - Performance Measures
 - Sample Statistics
 - Scenarios of Analysis
 - Guidelines for Presenting Data
3. Inferential Statistics
 - Statistical Tests
 - Experimental Designs
4. Race: Sequential Testing

Contents and Goals

Provide a view of issues in [Experimental Algorithmics](#)

- Exploratory data analysis
- Presenting results in a concise way with graphs and tables
- Organizational issues and Experimental Design

- Basics of inferential statistics
- Sequential statistical testing: race, a methodology for tuning

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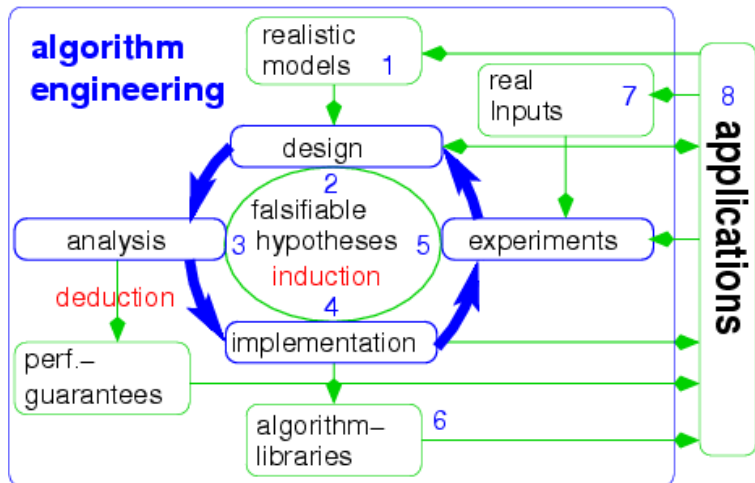
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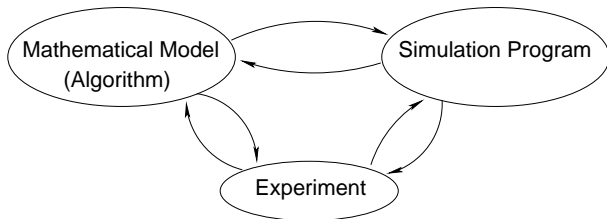
Experimental Algorithmics is an important part in the algorithm production cycle, which is referred to as [Algorithm Engineering](#)

The Engineering Cycle



from <http://www.algorithm-engineering.de/>

Experimental Algorithmics



In empirical studies we consider simulation programs which are the implementation of a mathematical model (the algorithm)

[McGeoch, 1996]

Experimental Algorithmics

Goals

- Defining standard methodologies
- Comparing relative performance of algorithms so as to identify the best ones for a given application
- Characterizing the behavior of algorithms
- Identifying algorithm separators, *i.e.*, families of problem instances for which the performance differ
- Providing new insights in algorithm design

Fairness Principle

Fairness principle: being completely fair is perhaps impossible but try to remove any possible bias

- possibly all algorithms must be implemented with the **same style**, with the **same language** and **sharing common subprocedures and data structures**
- the code must be **optimized**, e.g., using the best possible data structures
- running times must be comparable, e.g., by running experiments on the **same computational environment** (or redistributing them randomly)

Definitions

The most typical scenario considered in analysis of search heuristics

Asymptotic heuristics with time/quality limit decided *a priori*

The algorithm \mathcal{A}^∞ is halted when time expires or a solution of a given quality is found.

Deterministic case: \mathcal{A}^∞ on π
returns a solution of cost x .

The performance of \mathcal{A}^∞ on π is a
scalar $y = x$.

Randomized case: \mathcal{A}^∞ on π returns
a solution of cost X , where X is a
random variable.

The performance of \mathcal{A}^∞ on π is the
univariate $Y = X$.

[This is not the only relevant scenario: to be refined later]

Random Variables and Probability

Statistics deals with random (or stochastic) variables.

A variable is called random if, prior to observation, its outcome cannot be predicted with certainty.

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Discrete variables

Probability distribution:

$$p_i = P[x = v_i]$$

Cumulative Distribution Function (CDF)

$$F(v) = P[x \leq v] = \sum_i p_i$$

Mean

$$\mu = E[X] = \sum x_i p_i$$

Variance

$$\sigma^2 = E[(X - \mu)^2] = \sum (x_i - \mu)^2 p_i$$

Continuous variables

Probability density function (pdf):

$$f(v) = \frac{dF(v)}{dv}$$

Cumulative Distribution Function (CDF):

$$F(v) = \int_{-\infty}^v f(v) dv$$

Mean

$$\mu = E[X] = \int x f(x) dx$$

Variance

$$\sigma^2 = E[(X - \mu)^2] = \int (x - \mu)^2 f(x) dx$$

Generalization

For each general problem Π (e.g., TSP, GCP) we denote by C_Π a set (or class) of instances and by $\pi \in C_\Pi$ a single instance.

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$$Pr(Y = y | \pi)$$

It is often more interesting to generalize the performance on a class of instances C_Π , that is,

$$Pr(Y = y, C_\Pi) = \sum_{\pi \in \Pi} Pr(Y = y | \pi) Pr(\pi)$$

Sampling

In experiments,

1. we sample the population of instances and
2. we sample the performance of the algorithm on each sampled instance

If on an instance π we run the algorithm r times then we have r replicates of the performance measure Y , denoted Y_1, \dots, Y_r , which are independent and identically distributed (i.i.d.), i.e.

$$Pr(y_1, \dots, y_r | \pi) = \prod_{j=1}^r Pr(y_j | \pi)$$

$$Pr(y_1, \dots, y_r) = \sum_{\pi \in \mathcal{C}_\pi} Pr(y_1, \dots, y_r | \pi) Pr(\pi).$$

Instance Selection

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In *simulation studies* instances may be:

- real world instances
- random variants of real world-instances
- online libraries
- randomly generated instances

They may be grouped in classes according to some features whose impact may be worth studying:

- type (for features that might impact performance)
- size (for scaling studies)
- hardness (focus on hard instances)
- application (e.g., CSP encodings of scheduling problems), ...

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Within the class, instances are drawn with uniform probability $p(\pi) = c$

Statistical Methods

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- describe, summarizing, the data (descriptive statistics)
- make inference on those data (inferential statistics)

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(are the observed results enough to justify the claims?)
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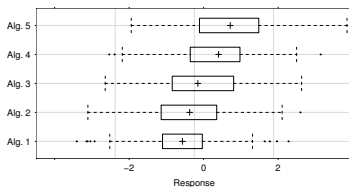
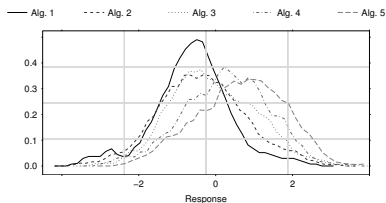
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In the **practical context** of heuristic design and implementation (i.e., **engineering**), statistics helps to take correct design decisions with the **least amount of experimentation**

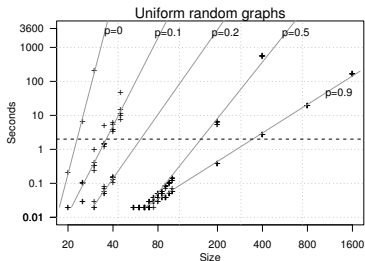
Objectives of the Experiments

- **Comparison:**
bigger/smaller, same/different,
Algorithm Configuration,
Component-Based Analysis
 - Standard statistical methods:
*experimental designs, test
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- **Comparison:**
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- **Characterization:**
Interpolation: fitting models to data
Extrapolation: building models of data, explaining phenomena
 - Standard statistical methods: *linear and non linear regression model fitting*



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Measures and Transformations

On a single instance

Design: Several runs on an instance

	Algorithm 1	Algorithm 2	...	Algorithm k
Instance 1	X_{11}	X_{21}		X_{k1}
⋮	⋮	⋮		⋮
Instance 1	X_{1r}	X_{2r}		X_{kr}

Measures and Transformations

On a single instance

Computational effort indicators

- number of elementary operations/algorithmic iterations
(e.g., search steps, objective function evaluations, number of visited nodes in the search tree, consistency checks, etc.)
- total CPU time consumed by the process
(sum of *user* and *system* times returned by `getrusage`)

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Solution quality indicators

- value returned by the cost function
- error from optimum/reference value
- (optimality) gap $\frac{UB-LB}{LB+\epsilon}$ (if max $\frac{UB-LB}{UB+\epsilon}$)
 ϵ is an infinitesimal for the case $LB = 0$ but $UB - LB \neq 0$
- ranks

Measures and Transformations

On a class of instances

Design A: One run on various instances

	Algorithm 1	Algorithm 2	...	Algorithm k
Instance 1	X_{11}	X_{12}		X_{1k}
⋮	⋮	⋮		⋮
Instance b	X_{b1}	X_{b2}		X_{bk}

Design B: Several runs on various instances

	Algorithm 1	Algorithm 2	...	Algorithm k
Instance 1	X_{111}, \dots, X_{11r}	X_{121}, \dots, X_{12r}		X_{1k1}, \dots, X_{1kr}
Instance 2	X_{211}, \dots, X_{21r}	X_{221}, \dots, X_{22r}		X_{2k1}, \dots, X_{2kr}
⋮	⋮	⋮		⋮
Instance b	X_{b11}, \dots, X_{b1r}	X_{b21}, \dots, X_{b2r}		X_{bk1}, \dots, X_{bkr}

Measures and Transformations

On a class of instances

Computational effort indicators

- no transformation if the interest is in studying scaling
- standardization if a fixed time limit is used
- geometric mean (used for a set of numbers whose values are meant to be multiplied together or are exponential in nature),
- otherwise, better to group homogeneously the instances

Solution quality indicators

Different instances imply different scales \Rightarrow need for an invariant measure

(However, many other measures can be taken both on the algorithms and on the instances [McGeoch, 1996])

Measures and Transformations

On a class of instances (cont.)

Solution quality indicators

- Distance or error from a reference value (assume minimization case):

$$e_1(x, \pi) = \frac{x(\pi) - \bar{x}(\pi)}{\sqrt{\hat{\sigma}(\pi)}} \quad \text{standard score}$$

$$e_2(x, \pi) = \frac{x(\pi) - x^{opt}(\pi)}{x^{opt}(\pi)} \quad \text{relative error}$$

$$e_3(x, \pi) = \frac{x(\pi) - x^{opt}(\pi)}{x^{worst}(\pi) - x^{opt}(\pi)} \quad \text{invariant [Zemel, 1981]}$$

- optimal value computed exactly or known by construction
- surrogate value such bounds or best known values
- Rank (no need for standardization but loss of information)

Outline

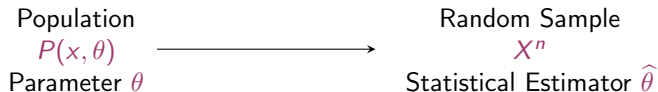
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Sampling

- We work with samples (instances, solution quality) drawn from populations

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Summary Measures

Measures to describe or characterize a population

- Measure of central tendency, location
- Measure of dispersion

One such a quantity is

- a **parameter** if it refers to the population (Greek letters)
- a **statistics** if it is an *estimation* of a population parameter from the sample (Latin letters)

Measures of central tendency

- Arithmetic Average (Sample mean)

$$\bar{X} = \frac{\sum x_i}{n}$$

- *Quantile*: value above or below which lie a fractional part of the data (used in nonparametric statistics)
 - Median

$$\mathcal{M} = x_{(n+1)/2}$$

- Quartile

$$Q_1 = x_{(n+1)/4} \quad Q_3 = x_{3(n+1)/4}$$

- q -quantile

q of data lies below and $1 - q$ lies above

- Mode

value of relatively great concentration of data
(*Unimodal* vs *Multimodal* distributions)

Measure of dispersion

- Sample range

$$R = x_{(n)} - x_{(1)}$$

- Sample variance

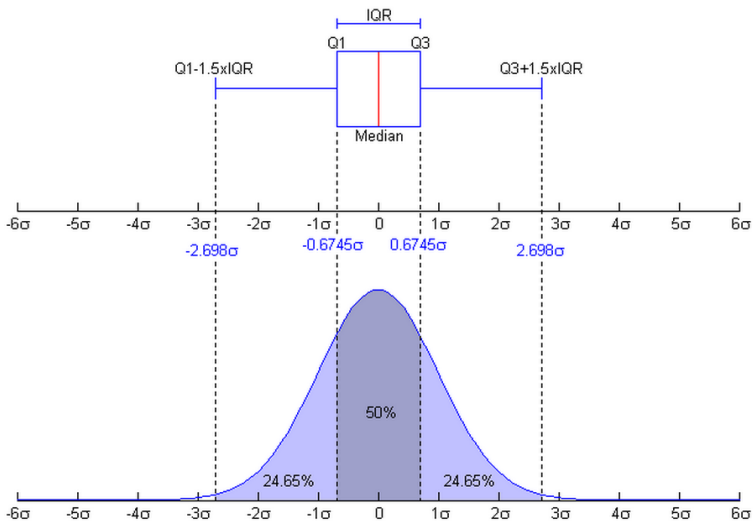
$$s^2 = \frac{1}{n-1} \sum (x_i - \bar{X})^2$$

- Standard deviation

$$s = \sqrt{s^2}$$

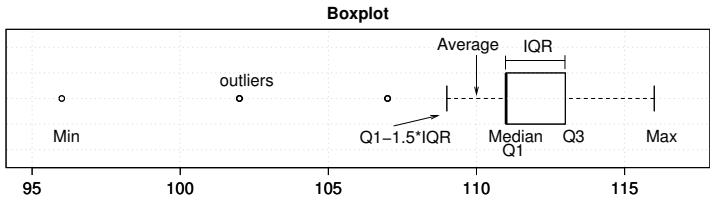
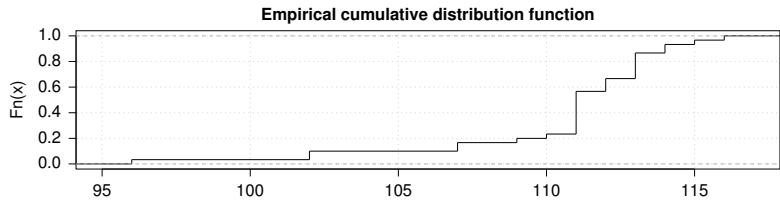
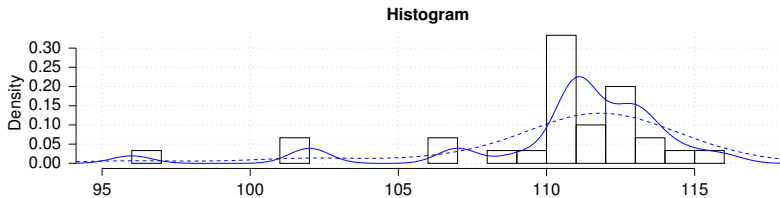
- Inter-quartile range

$$IQR = Q_3 - Q_1$$



Boxplot and a probability density function (pdf) of a Normal $N(0,1\sigma^2)$ Population.
 (source: Wikipedia)

[see also: <http://informationandvisualization.de/blog/box-plot>]



In R

```
> x<-runif(10,0,1)
  mean(x), median(x), quantile(x), quantile(x,0.25)
  range(x), var(x), sd(x), IQR(x)
> fivenum(x)
 #(minimum, lower-hinge, median, upper-hinge, maximum)
[1] 0.18672 0.26682 0.28927 0.69359 0.92343
> summary(x)
> aggregate(x,list(factors),median)
> boxplot(x)
```

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Scenarios

A. Single-pass heuristics

B. Asymptotic heuristics:

Two approaches:

1. Univariate

1.a Time as an external parameter decided *a priori*

1.b Solution quality as an external parameter decided *a priori*

2. Cost dependent on running time:

Scenario A

Single-pass heuristics

Deterministic case: \mathcal{A}^{-1} on class C_{Π} returns a solution of cost x with computational effort t (e.g., running time).

The performance of \mathcal{A}^{-1} on class C_{Π} is the vector $\vec{y} = (x, t)$.

Randomized case: \mathcal{A}^{-1} on class C_{Π} returns a solution of cost X with computational effort T , where X and T are random variables.

The performance of \mathcal{A}^{-1} on class C_{Π} is the bivariate $\vec{Y} = (X, T)$.

Example

Scenario:

- ▷ 3 heuristics \mathcal{A}_1^+ , \mathcal{A}_2^+ , \mathcal{A}_3^+ on class C_{Π} .
- ▷ homogeneous instances or need for data transformation.
- ▷ 1 or r runs per instance
- ▶ **Interest:** inspecting solution cost and running time to observe and compare the level of approximation and the speed.

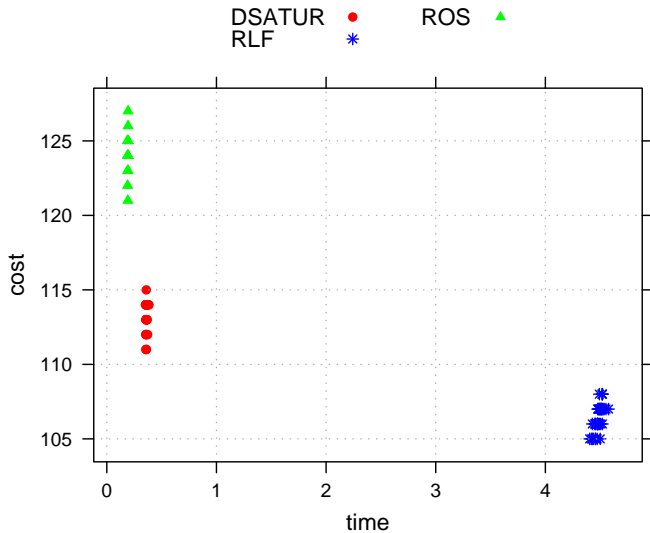
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Tools:

- Scatter plots of solution-cost and run-time



Multi-Criteria Decision Making

Needed some definitions on **dominance relations**

In **Pareto sense**, for points in \mathbf{R}^2

$\vec{x}^1 \preceq \vec{x}^2$	weakly dominates	$x_i^1 \leq x_i^2$ for all $i = 1, \dots, n$
$\vec{x}^1 \parallel \vec{x}^2$	incomparable	neither $\vec{x}^1 \preceq \vec{x}^2$ nor $\vec{x}^2 \preceq \vec{x}^1$

Scenario B

Asymptotic heuristics

There are two approaches:

- 1.a. Time as an external parameter decided *a priori*.

The algorithm is halted when time expires.

Deterministic case: \mathcal{A}^∞ on class C_Π returns a solution of cost x .

The performance of \mathcal{A}^∞ on class C_Π is the scalar $y = x$.

Randomized case: \mathcal{A}^∞ on class C_Π returns a solution of cost X , where X is a random variable.

The performance of \mathcal{A}^∞ on class C_Π is the univariate $Y = X$.

Example

Scenario:

- ▷ 3 heuristics A_1^∞ , A_2^∞ , A_3^∞ on class C_Π .
(Or 3 heuristics A_1^∞ , A_2^∞ , A_3^∞ on class C_Π without interest in computation time because negligible or comparable)
- ▷ homogeneous instances (no data transformation) or heterogeneous (data transformation)
- ▷ 1 or r runs per instance
- ▷ a priori time limit imposed
- ▶ **Interest:** inspecting solution cost

Example

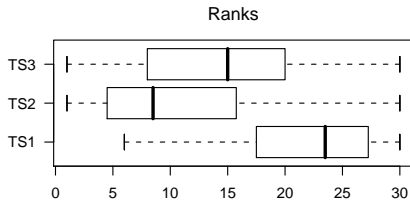
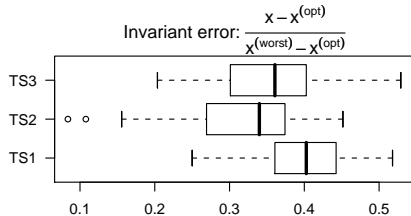
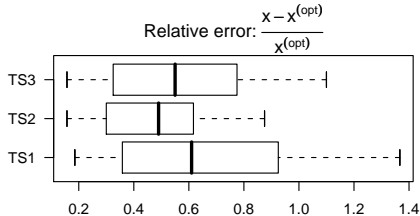
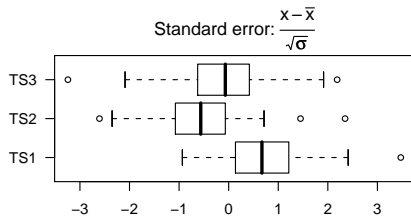
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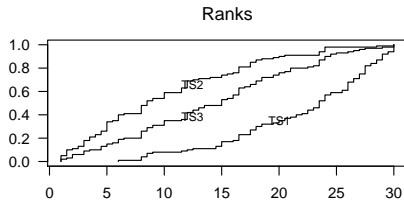
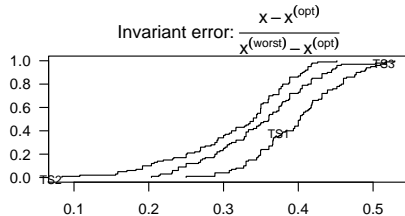
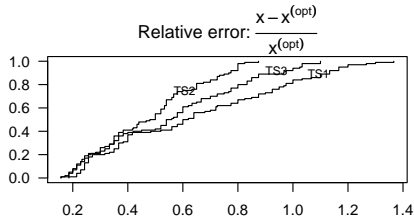
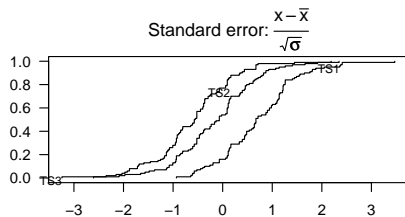
Tools:

- Histograms (summary measures: mean or median or mode?)
- Boxplots
- Empirical cumulative distribution functions (ECDFs)

On a class of instances



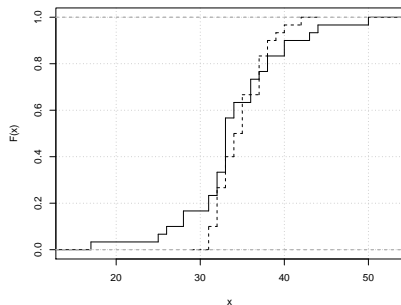
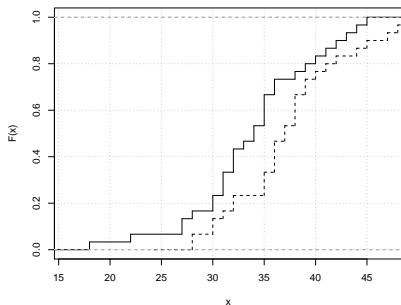
On a class of instances



Stochastic Dominance

Definition: Algorithm \mathcal{A}_1 probabilistically dominates algorithm \mathcal{A}_2 on a problem instance, iff its CDF is always "below" that of \mathcal{A}_2 , i.e.:

$$F_1(x) \leq F_2(x), \quad \forall x \in X$$



R code behind the previous plots

We load the data and plot the comparative boxplot for each instance.

```
> load("TS.class-G.dataR")
> G[1:5,]
  alg inst run sol time.last.imp tot.iter parz.iter exit.iter exit.time opt
1 TS1 G-1000-0.5-30-1.1.col 1 59 9.900619 5955 442 5955 10.02463 30
2 TS1 G-1000-0.5-30-1.1.col 2 64 9.736608 3880 130 3958 10.00062 30
3 TS1 G-1000-0.5-30-1.1.col 3 64 9.908618 4877 49 4877 10.03263 30
4 TS1 G-1000-0.5-30-1.1.col 4 68 9.948622 6996 409 6996 10.07663 30
5 TS1 G-1000-0.5-30-1.1.col 5 63 9.912620 3986 52 3986 10.04063 30
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> library(lattice)
> bwplot(alg ~ sol | inst,data=G)
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```

If we want to make an aggregate analysis we have the following choices:

- maintain the raw data,
- transform data in standard error,
- transform the data in relative error,
- transform the data in an invariant error,
- transform the data in ranks.

Maintain the raw data

```
> par(mfrow=c(3,2),las=1,font.main=1,mar=c(2,3,3,1))  
> #original data  
> boxplot(sol~alg,data=G,horizontal=TRUE,main="Original data")
```

Transform data in standard error

```

> #standard error
> T1 <- split(G$sol,list(G$inst))
> T2 <- lapply(T1,scale=center=TRUE,scale=TRUE)
> T3 <- unsplit(T2,list(G$inst))
> T4 <- split(T3,list(G$alg))
> T5 <- stack(T4)
> boxplot(values~ind,data=T5,horizontal=TRUE,main=expression(paste("Standard error: ",
  frac(x-bar(x),sqrt(sigma))))))
> library(latticeExtra)
> ecdfplot(~values,group=ind,data=T5,main=expression(paste("Standard error:
",frac(x-bar(x),sqrt(sigma))))))

> #standard error
> G$scale <- 0
> split(G$scale, G$inst) <- lapply(split(G$sol, G$inst), scale=center=TRUE,scale=TRUE)

```

Transform the data in relative error

```
> #relative error  
> G$err2 <- (G$sol - G$opt)/G$opt  
> boxplot(err2~alg, data=G, horizontal=TRUE, main=expression(paste("Relative error: ", frac(x  
- x^(opt), x^(opt))))))  
> ecdfplot(G$err2, group=G$alg, main=expression(paste("Relative error: ", frac(x - x^(opt), x^(  
opt))))))
```

Transform the data in an invariant error

We use as surrogate of x^{worst} the median solution returned by the simplest algorithm for the graph coloring, that is, the ROS heuristic.

```
> #error 3
> load("ROS.class-G.dataR")
> F1 <- aggregate(F$sol,list(inst=F$inst),median)
> F2 <- split(F1$x,list(F1$inst))
> G$ref <- sapply(G$inst,function(x) F2[[x]])
> G$err3 <- (G$sol - G$opt)/(G$ref - G$opt)
> boxplot(err3~alg,data=G,horizontal=TRUE,main=expression(paste("Invariant error: ",frac(
  x - x^(opt),x^(worst) - x^(opt))))))
> ecdfplot(G$err3,group=G$alg,main=expression(paste("Invariant error: ",frac(x - x^(opt),x^(
  worst) - x^(opt))))))
```

Transform the data in ranks

```
> #rank  
> G$rank <- G$sol  
> split(G$rank, G$inst) <- lapply(split(G$sol, D$inst), rank)  
> bwplot(rank~reorder(alg,rank,median),data=G,horizontal=TRUE,main="Ranks")  
> ecdfplot(rank,group=alg,data=G,main="Ranks")
```

Scenario B

Asymptotic heuristics

There are two approaches:

- 1.b. **Solution quality** as an external parameter decided *a priori*. The algorithm is halted when quality is reached.

Deterministic case: \mathcal{A}^∞ on class C_Π finds a solution in running time t .

The performance of \mathcal{A}^∞ on class C_Π is the scalar $y = t$.

Randomized case: \mathcal{A}^∞ on class C_Π finds a solution in running time T , where T is a random variable.

The performance of \mathcal{A}^∞ on class C_Π is the univariate $Y = T$.

Dealing with Censored Data

Asymptotic heuristics, Approach 1.b

- ▷ Heuristic \mathcal{A}^+ stopped before completion or \mathcal{A}^∞ truncated (always the case)
- ▶ **Interest:** determining whether a prefixed goal (optimal/feasible) has been reached

The computational effort to attain the goal can be specified by a cumulative distribution function $F(t) = P(T < t)$ with T in $[0, \infty)$.

If in a run i we stop the algorithm at time L_i then we have a **Type I right censoring**, that is, we know either

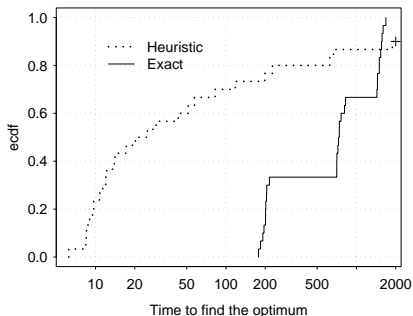
- T_i if $T_i \leq L_i$
- or $T_i \geq L_i$.

Hence, for each run i we need to record $\min(T_i, L_i)$ and the indicator variable for observed optimal/feasible solution attainment, $\delta_i = I(T_i \leq L_i)$.

Example

Asymptotic heuristics, Approach 1.b: Example

- ▷ An exact *vs* an heuristic algorithm for the *2-edge-connectivity augmentation problem*.
- ▶ **Interest:** time to find the optimum on different instances.



Uncensored:

$$F(t) = \frac{\# \text{ runs} < t}{n}$$

Censored:

$$F(t) = \frac{\# \text{ runs} < t}{n}$$

Scenario B

Asymptotic heuristics

There are two approaches:

2. Cost dependent on running time:

Deterministic case: \mathcal{A}^∞ on π returns a current best solution x at each observation in t_1, \dots, t_k .

The performance of \mathcal{A}^∞ on π is the **profile** indicated by the vector $\vec{y} = \{x(t_1), \dots, x(t_k)\}$.

Randomized case: \mathcal{A}^∞ on π produces a monotone stochastic process in solution cost $X(\tau)$ with any element dependent on the predecessors.

The performance of \mathcal{A}^∞ on π is the **multivariate** $\vec{Y} = (X(t_1), X(t_2), \dots, X(t_k))$.

Example

Scenario:

- ▷ 3 heuristics \mathcal{A}_1^∞ , \mathcal{A}_2^∞ , \mathcal{A}_3^∞ on instance π .
- ▷ single instance hence no data transformation.
- ▷ r runs
- ▶ **Interest:** inspecting solution cost over running time to determine whether the comparison varies over time intervals

Example

Scenario:

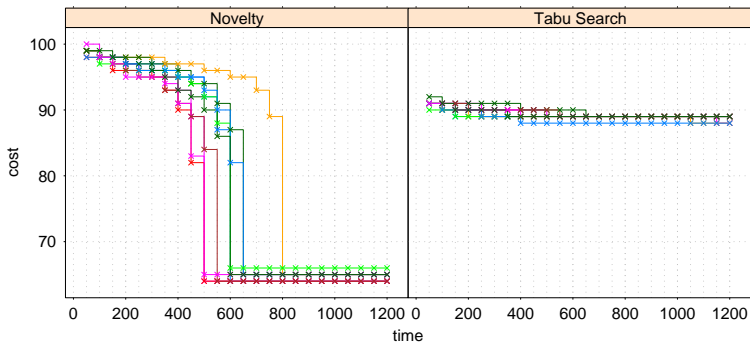
- ▷ 3 heuristics \mathcal{A}_1^∞ , \mathcal{A}_2^∞ , \mathcal{A}_3^∞ on instance π .
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Tools:

- Quality profiles

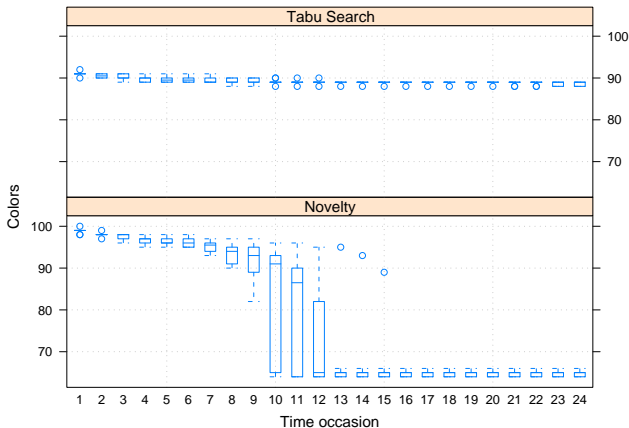
The performance is described by **multivariate random variables** of the kind $\vec{Y} = \{Y(t_1), Y(t_2), \dots, Y(t_k)\}$.

Sampled data are of the form $\vec{Y}_i = \{Y_i(t_1), Y_i(t_2), \dots, Y_i(t_k)\}$, $i = 1, \dots, 10$ (10 runs per algorithm on one instance)



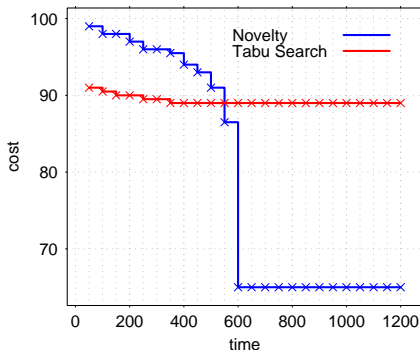
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The median behavior of the two algorithms

Summary

Visualize your data for your **analysis** and for **communication** to others

Explore your data:

- make plots: histograms, boxplots, empirical cumulative distribution functions, correlation/scatter plots
- look at the numerical data and interpret them in practical terms: computation times, distance from optimum
- look for patterns

All the above both at a single instance level and at an aggregate level.

Outline

1. Experimental Analysis
2. Descriptive Statistics
 - Performance Measures
 - Sample Statistics
 - Scenarios of Analysis
 - Guidelines for Presenting Data**
3. Inferential Statistics
 - Statistical Tests
 - Experimental Designs
4. Race: Sequential Testing

Making Plots

<http://algo2.iti.uni-karlsruhe.de/sanders/courses/bergen/bergenPresenting.pdf>

[Sanders, 2002]

- Should the experimental setup from the exploratory phase be redesigned to increase conciseness or accuracy?
- What parameters should be varied? What variables should be measured?
- How are parameters chosen that cannot be varied?
- Can tables be converted into curves, bar charts, scatter plots or any other useful graphics?
- Should tables be added in an appendix?
- Should a 3D-plot be replaced by collections of 2D-curves?
- Can we reduce the number of curves to be displayed?
- How many figures are needed?
- Should the x-axis be transformed to magnify interesting subranges?

- Should the x-axis have a logarithmic scale? If so, do the x-values used for measuring have the same basis as the tick marks?
- Is the range of x-values adequate?
- Do we have measurements for the right x-values, i.e., nowhere too dense or too sparse?
- Should the y-axis be transformed to make the interesting part of the data more visible?
- Should the y-axis have a logarithmic scale?
- Is it misleading to start the y-range at the smallest measured value? (if not too much space wasted start from 0)
- Clip the range of y-values to exclude useless parts of curves?
- Can we use banking to 45° ?
- Are all curves sufficiently well separated?
- Can noise be reduced using more accurate measurements?
- Are error bars needed? If so, what should they indicate? Remember that measurement errors are usually not random variables.

- Connect points belonging to the same curve.
- Only use splines for connecting points if interpolation is sensible.
- Do not connect points belonging to unrelated problem instances.
- Use different point and line styles for different curves.
- Use the same styles for corresponding curves in different graphs.
- Place labels defining point and line styles in the right order and without concealing the curves.
- Give axis units
- Captions should make figures self contained.
- Give enough information to make experiments reproducible.
- Golden ratio rule: make the graph wider than higher [Tufté 1983].
- Rule of 7: show at most 7 curves (omit those clearly irrelevant).
- Avoid: explaining axes, connecting unrelated points by lines, cryptic abbreviations, microscopic lettering, pie charts

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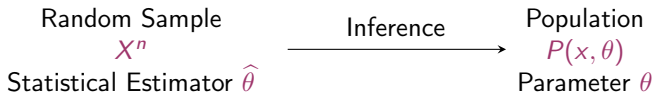
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Inferential Statistics

- We work with samples (instances, solution quality)
- But we want sound conclusions: generalization over a given population (all possible instances)
- Thus we need **statistical inference**

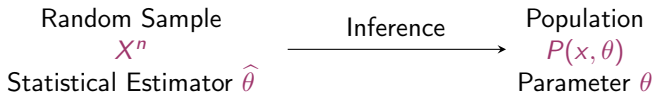
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- But we want sound conclusions: generalization over a given population (all possible instances)
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Since the analysis is based on finite-sized sampled data, statements like
“the cost of solutions returned by algorithm A is smaller than that of algorithm B ”

must be completed by

“at a level of significance of 5%”.

A Motivating Example

- There is a competition and two stochastic algorithms \mathcal{A}_1 and \mathcal{A}_2 are submitted.
- We run both algorithms once on n instances.
On each instance either \mathcal{A}_1 wins (+) or \mathcal{A}_2 wins (-) or they make a tie (=).

Questions:

1. If we have only 10 instances and algorithm \mathcal{A}_1 wins 7 times how confident are we in claiming that algorithm \mathcal{A}_1 is the best?
2. How many instances and how many wins should we observe to gain a confidence of 95% that the algorithm \mathcal{A}_1 is the best?

A Motivating Example

- p : probability that \mathcal{A}_1 wins on each instance (+)
- n : number of runs without ties
- Y : number of wins of algorithm \mathcal{A}_1

If each run is independent and consistent:

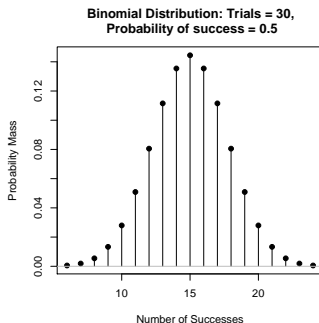
$$Y \sim B(n, p) : \quad \Pr[Y = y] = \binom{n}{y} p^y (1 - p)^{n-y}$$

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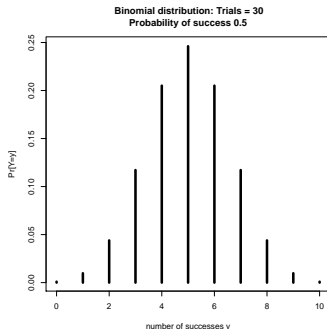
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1 If we have only 10 instances and algorithm \mathcal{A}_1 wins 7 times how confident are we in claiming that algorithm \mathcal{A}_1 is the best?

Under these conditions, we can check how unlikely the situation is if it were $p(+)\leq p(-)$.

If $p = 0.5$ then the chance that algorithm \mathcal{A}_1 wins 7 or more times out of 10 is 17.2%: quite high!



2 How many instances and how many wins should we observe to gain a confidence of 95% that the algorithm \mathcal{A}_1 is the best?

To answer this question, we compute the 95% quantile, *i.e.*, $y : \Pr[Y \geq y] < 0.05$ with $p = 0.5$ at different values of n :

n	10	11	12	13	14	15	16	17	18	19	20
y	9	9	10	10	11	12	12	13	13	14	15

2 How many instances and how many wins should we observe to gain a confidence of 95% that the algorithm \mathcal{A}_1 is the best?

To answer this question, we compute the 95% quantile, *i.e.*, $y : \Pr[Y \geq y] < 0.05$ with $p = 0.5$ at different values of n :

n	10	11	12	13	14	15	16	17	18	19	20
y	9	9	10	10	11	12	12	13	13	14	15

This is an application example of **sign test**, a special case of binomial test in which $p = 0.5$

Statistical tests

General procedure:

- Assume that data are consistent with a **null hypothesis** H_0 (e.g., sample data are drawn from distributions with the same mean value).
- Use a statistical test to compute how likely this is to be true, given the data collected. This “likely” is quantified as the **p-value**.
- Do not reject H_0 if the **p-value** is larger than an user defined threshold called **level of significance** α .
- Alternatively, (**p-value** $< \alpha$), H_0 is rejected in favor of an **alternative hypothesis**, H_1 , at a level of significance of α .

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Preparation of the Experiments

Variance reduction techniques

- Same pseudo random seed

Sample Sizes

- If the sample size is large enough (infinity) any difference in the means of the factors, no matter how small, will be significant
- Real vs Statistical significance
Study factors until the improvement in the response variable is deemed small
- Desired statistical power + practical precision \Rightarrow sample size

Note: If resources available for N runs then the optimal design is **one run on N instances** [Birattari, 2004]

Experimental Design

Algorithms \Rightarrow Treatment Factor; Instances \Rightarrow Blocking/Random Factor

Design A: One run on various instances (Unreplicated Factorial)

	Algorithm 1	Algorithm 2	...	Algorithm k
Instance 1	X_{11}	X_{12}		X_{1k}
\vdots	\vdots	\vdots		\vdots
Instance b	X_{b1}	X_{b2}		X_{bk}

Design B: Several runs on various instances (Replicated Factorial)

	Algorithm 1	Algorithm 2	...	Algorithm k
Instance 1	X_{111}, \dots, X_{11r}	X_{121}, \dots, X_{12r}		X_{1k1}, \dots, X_{1kr}
Instance 2	X_{211}, \dots, X_{21r}	X_{221}, \dots, X_{22r}		X_{2k1}, \dots, X_{2kr}
\vdots	\vdots	\vdots		\vdots
Instance b	X_{b11}, \dots, X_{b1r}	X_{b21}, \dots, X_{b2r}		X_{bk1}, \dots, X_{bkr}

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Unreplicated Designs

Procedure Race [Birattari 2002]:

repeat

Randomly select an unseen instance and run all candidates on it

Perform *all-pairwise comparison* statistical tests

Drop all candidates that are significantly inferior to the best algorithm

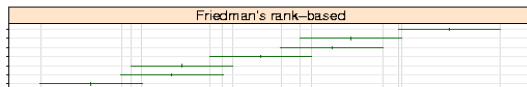
until only one candidate left or no more unseen instances ;

- F-Race use Friedman test
- Holm adjustment method is typically the most powerful

```
race(wrapper.file, maxExp=0,
      stat.test=c("friedman","t.bonferroni","t.holm","t.none"),
      conf.level=0.95, first.test=5, interactive=TRUE,
      log.file="", no.slaves=0,...)
```

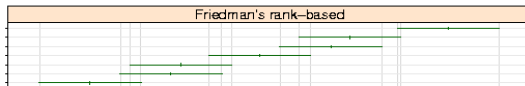
Sequential Testing

⇒ 7 algorithms, 5 instances, 3 runs

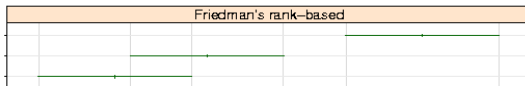


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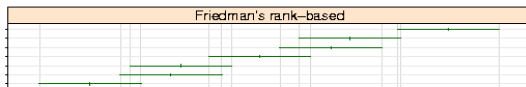


⇒ 3 algorithms, 5 instances, 9 runs

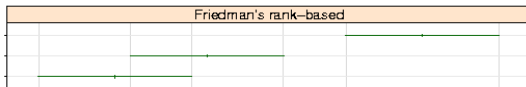


Sequential Testing

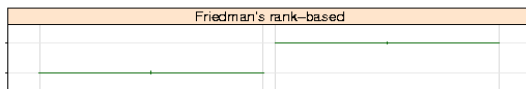
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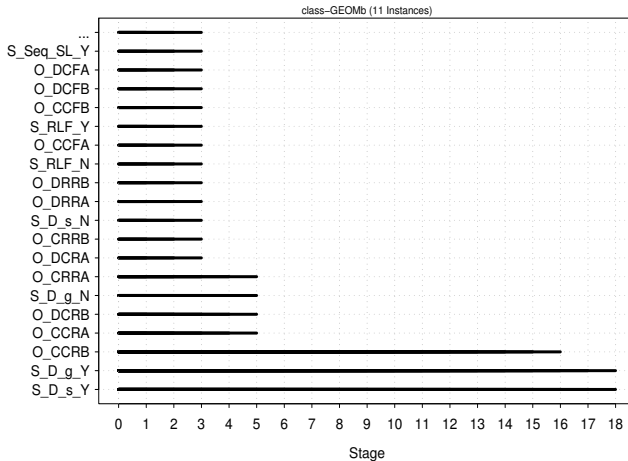
⇒ 3 algorithms, 5 instances, 9 runs



⇒ 2 algorithms, 5 instances, 30 runs



Sequential Testing



References

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- Chiarandini M. (2009). **Experimental analysis of optimization heuristics using R**. Lecture notes available at <http://www.imada.sdu.dk/~marco/Teaching/Files/Rnotes.pdf>.
- Sanders P. (2002). **Presenting data from experiments in algorithmics**. In *Experimental Algorithmics – From Algorithm Design to Robust and Efficient Software*, vol. 2547 of **LNCS**, pp. 181–196. Springer.