# Robust statistics and some non-parametric statistics

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Statistics Course for Astrophysicists, 2010–2011

**Robust statistics** 

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Why robust statistics? Removal of data L-estimators R-estimators M-estimators

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L-estimators

**R**-estimators

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## Caveat

The presentation aims at being user-focused and at presenting usable recipes

Do not expect a fully mathematically rigorous description!

This has been prepared in the hope to be useful even in the stand-alone version

Please provide me feedback with any misconception or error you may find, and with any topic not addressed here but which should be included

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## Example: The average weight of opossums



- Find a group of N opossums
- Weight them:  $w_i$ , i = 1, ..., N
- Calculate the average weight as  $1/N \sum_i w_i$
- You should get about 3.5 kg
- But you better not use this group!

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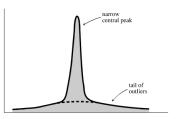
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## General concepts



Robust statistics:

- are not (less) affected by the presence of outliers or deviations from model assumptions
- are related, but not identical to non-parametric statistics, where we drop the hypothesis of underlying Gaussian distribution

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## Main approaches

Four main approaches:

- Discard data
- L-estimators: Use linear combinations of order statistics
- R-estimators: Use rank instead of values
- M-estimators: Estimators based on Maximum-likelihood argument

The breakdown point is the fraction of outliers above which the robust statistics fails. It can reach 50%

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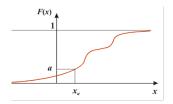
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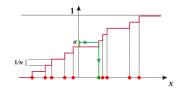
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## Quantiles



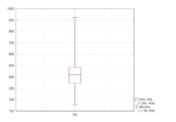


- *q*-quantiles (*q* ∈ ℕ) are the set of value
   *x*<sub>[*a*]</sub>, *a* = *k*/*q*, *k* = 1,..., *q* − 1 from a probability distribution for which *P*(*x* < *x*<sub>[*a*]</sub>) = *a*
- 4-quantiles, or quartiles are then  $x_{[0.25]}$ ,  $x_{[0.5]}$ ,  $x_{[0.75]}$
- *q*-quantiles from a sorted sample  $\{x_i\}, i = 1, ..., N$ :  $x_{[a]} = \min_i x_i \mid i/N \ge a$
- Example: if N = 12, quartiles are: x<sub>3</sub>, x<sub>6</sub>, x<sub>9</sub>; if N = 14, quartiles are: x<sub>4</sub>, x<sub>7</sub>, x<sub>11</sub>

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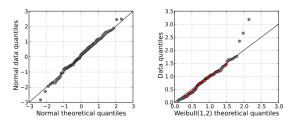
## Box plot



### Robust statistics Stéphane Paltani Why robust

- A box plot is a synthetic way to look at the sample properties
- It represents in a single plot, the minimum, the quartiles (hence also the median) and the maximum
- ▶ that is, the quantiles  $x_{[0]}$ ,  $x_{[0.25]}$ ,  $x_{[0.5]}$ ,  $x_{[0.75]}$ ,  $x_{[1.0]}$

## Q-Q plot



- A Q-Q plot (quantile-quantile plot) is powerful way to compare sample properties with an assumed underlying distribution
- ► The QQ-plot is the plot of x<sub>i</sub> vs f<sub>i</sub>, where f<sub>i</sub> are the values for which P(x < f<sub>i</sub>) = i/(N + 1), when x<sub>i</sub> is sorted in increasing order
- One usually also draw f<sub>i</sub> vs f<sub>i</sub> for comparison with expected values
- Outliers appear quite clearly

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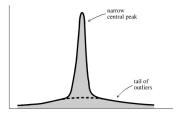
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## What are these outliers?



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- They can be due to errors in the measurement
- They may be bona fide measurements, but the distribution is heavy-tailed, so these points may be the black swan and contain important information
- Outliers can be difficult to identify with confidence, and they can hide each other
- There is the risk of data manipulation

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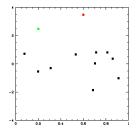
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## Chauvenet's criterion



## • Assuming a data set $\{x_i\} \sim \mathcal{N}(\mu, \sigma), i = 1, \dots, N$

- Calculate for each  $i P(x_i) \equiv P(|x| > |x_i|)$
- ► Discard the point if N · P(x<sub>i</sub>) < 0.5, i.e. if the probability to have such an extreme value is less than 50%, taking the number of trials into account</p>
- Moderate outliers may mask more extreme outliers
- Grubbs's test uses absolute maximum deviation  $G = \max_{i} \frac{|x_i \mu|}{\sigma}$

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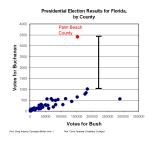
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## Dixon's Q test



- Dixon's Q test: Find the largest gap in a sorted sample, and divide it by the total range
- In this case:  $Q = \frac{2500}{3500} \simeq 0.71$

### Critical values:

Number of values:	3	4	5	6	7	8	9	10
Q90%:	0.941	0.765	0.642	0.560	0.507	0.468	0.437	0.412
Q95%:	0.970	0.829	0.710	0.625	0.568	0.526	0.493	0.466
Q99%:	0.994	0.926	0.821	0.740	0.680	0.634	0.598	0.568

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## **Trimmed estimators**

- Trimming is a generic method to to make an estimator robust
- The n% trimmed estimator is obtained by calculating the estimator on the sample limited to the range [x<sub>[n%]</sub>, x<sub>[1-n%]</sub>]
- This is not equivalent to removing outliers, as the trimmed estimators have the same expectations if there are no outliers
- The trimmed mean (or truncated mean) is a robust alternative to the mean, which is more dependent on the distribution than the median

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## Winsorized estimators

- Winsorizing is another generic method to to make an estimator robust which is very similar to trimming
- The n% winsorizing estimator is obtained by replacing in the sample all values below x<sub>[n%]</sub> by x<sub>[n%]</sub> and all values above x<sub>[1-n%]</sub>] by x<sub>[1-n%]</sub>
- I have no idea why you would want to do that...

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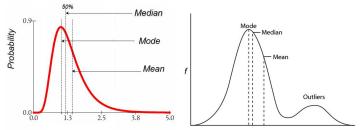
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## Median



- The median x̃ is the 2-quantile, i.e. x<sub>[0.5]</sub>, i.e. x<sub>N/2</sub> if N is even or x<sub>N+1/2</sub> if N is odd
- If we have 10 "opossums", the mean will be about one ton! It has a breakdown point of 0 %
- The median has a breakdown point of 50%, i.e. you would get a roughly correct weight estimations even if there are 5 mammoths in your populations of 10 "opossums"

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## Median average deviation

- ► Assuming a sample {x<sub>i</sub>}, i = 1,..., N, and its median x̃, the median average deviation is MAD({x<sub>i</sub>}) = median(|x<sub>i</sub> x̃|)
- Example:  $\{x_i\} \equiv \{1, 1, 2, 2, 4, 6, 9\}$  $\tilde{x} = 2, \{|x_i - \tilde{x}|\} \equiv \{0, 0, 1, 1, 2, 4, 7\}$ , so MAD $(\{x_i\}) = 1$
- Note that 6 and 9 are completely ignored
- Relation to standard deviation is distribution-dependent:
  - For a Gaussian distribution,  $\sigma_x \simeq 1.4826 \operatorname{MAD}(\{x_i\})$
  - For a uniform distribution,  $\sigma_x = \sqrt{4/3} \operatorname{MAD}(\{x_i\})$

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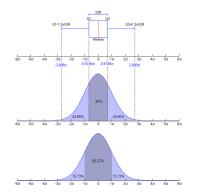
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## Inter-quartile range



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- Assuming a sample  $\{x_i\}, i = 1, \dots, N$
- ► The inter-quartile range is the difference between the third and first quartiles: IQR({x<sub>i</sub>}) = x<sub>[0.75]</sub> x<sub>[0.25]</sub>
- ▶ For a Gaussian distribution,  $\sigma_x \simeq IQR(\{x_i\})/1.349$
- This can be used as a (non-robust) normality test

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## Ranks

- R-estimators are based on rank
- Given a sample  $\{x_i\}, i = 1, \ldots, N$
- The rank of x<sub>i</sub> is i if x<sub>i</sub> is sorted in increasing order
- Example:  $\{x_i\} \equiv \{4, 9, 6, 21, 3, 11, 1\}$
- ► The ranks are: {3,5,4,7,2,6,1}, because the sorted {*x<sub>i</sub>*} is {1,3,4,6,9,11,21}

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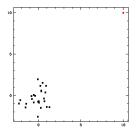
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## Pearson's r: a non-robust estimator



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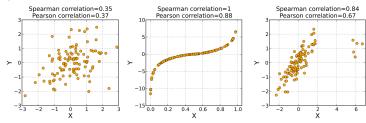
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- ► Pearson's *r* correlation coefficient between two random variables {*x<sub>i</sub>*, *y<sub>i</sub>*}, *i* = 1,..., *N* is:  $r = \frac{\overline{xy}}{\sigma_x \sigma_y}$
- But it is not robust. In this case, r = 0.85
- Of course, on can use L-estimators to compute xy, *σ<sub>x</sub>*, *σ<sub>y</sub>*
- But we would have to figure out the critical values

## Spearman's correlation coefficient



- Spearman's coefficient: Replace {x<sub>i</sub>} and {y<sub>i</sub>} with their ranks, and calculate s as the the Pearson's correlation coefficient of the ranks
- Consistent with Pearson's r in "good" conditions
- It is insensitive to the shape of y vs x and it is robust
- ► Significance: t = s√(N-2)/(1-s<sup>2</sup>) is approximately Student-t distributed with N 2 degrees of freedom
- Ties should have the same (average) rank, i.e. if x<sub>6</sub> = x<sub>7</sub>, give them a rank of 6.5

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## Kendall's $\tau$ rank correlation

- Rank still contain quantitative values
- Form N(N-1)/2 pairs  $\{\{x_i, y_i\}; \{x_j, y_j\}\}, j > i$

$$\tau = \frac{\sum_{i,j} \operatorname{sgn}(x_i - x_j) \operatorname{sgn}(y_i - y_j) - \sum_{i,j} \operatorname{sgn}(x_i - x_j) \operatorname{sgn}(y_j - y_i)}{N(N-1)/2}$$

- For relatively large sample,  $\tau \sim \mathcal{N}\left(0, \frac{2(2N+5)}{9N(N-1)}\right)$
- No single recipe in case of ties

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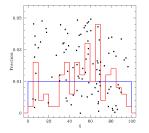
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## Do two distributions differ?



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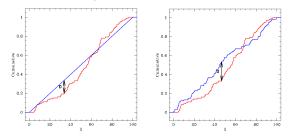
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- Assuming a sample {x<sub>i</sub>}, i = 1,..., N, is it a probable outcome from a draw of N random variables from a given distribution (say U(a, b))?
- Similarly, assuming two samples {*x<sub>i</sub>*}, *i* = 1,..., *N* and {*y<sub>j</sub>*}, *j* = 1,..., *M*, how probable is it that they have the same parent population?

## Cumulative comparison



### A (good) idea is to compare cumulative distributions

- $F(x) = \int_a^b f(x) dx$  for a continuous distribution f(x)
- $C_{\{x_i\}}(x) = \frac{1}{N} \sum_i H(x x_i)$  for a sample (H(x) is the Heaviside function, i.e., 1 if  $x \ge 0$ , 0 if x < 0)
- ► The KS test is the simplest quantitative comparison:  $D = \max_{x} |C_{\{x_i\}}(x) F(x)| \text{ or }$   $D = \max_{x} |C_{\{x_i\}}(x) C_{\{y_i\}}(x)|$

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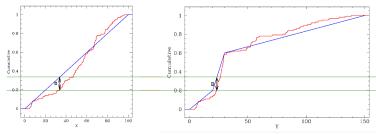
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# Independence on underlying distribution



- *D* is preserved under any transformation  $x \rightarrow y = \psi(x)$ , where  $\psi(x)$  is an arbitrary strictly monotonic function
- Thus KS test works with any underlying distribution
- ► The null-hypothesis distribution of *D* is:  $P(\lambda > D) = 2 \sum_{k=1}^{\infty} (-1)^{k-1} e^{-2k^2 \mu^2}, \text{ with } \mu = \left(\sqrt{N} + 0.12 + 0.11/\sqrt{N}\right) \cdot D$
- When comparing two samples, use:  $N_e = \frac{N \cdot M}{N + M}$

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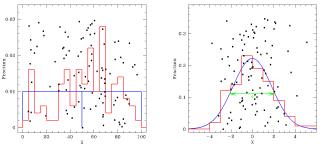
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## Caveats



In weird cases, KS-test might be extremely inefficient. KS test makes hidden assumptions

 KS-test will not work if you derive parameters from the data (see "Monte-Carlo methods" course)

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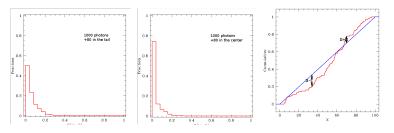
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# Kuiper test



- KS test is more sensitive in the tails than in the center
- Among several solutions, the simplest: Kuiper test!
  V = D<sup>+</sup> + D<sup>-</sup>
- $P(\lambda > V) = 2 \sum_{k=1}^{\infty} (4k^2 \mu^2 1) e^{-2k^2 \mu^2}$ , with  $\mu = (\sqrt{N} + 0.155 + 0.24/\sqrt{N}) \cdot V$
- Kuiper test can be used for distributions on a circle (see Paltani 2004)

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M-estimators Concepts Maximum-likelihood estimation So, these M-estimators...

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Concepts

Maximum-likelihood estimation So, these M-estimators..

## Concepts

- M-estimators are a generalization of maximum-likelihood estimators
- So, maximum likelihood is an M-estimator
- It allows to give arbitrary weight to data points
- Weights can be chosen so that they decrease for too far-off points, opposite to Gaussian weights in least square-estimation

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#### Concepts

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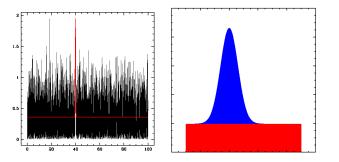
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Concepts

Maximum-likelihood estimation

# A case of known outlier distribution



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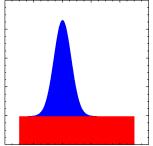
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Concepts

Maximum-likelihood estimation

- We search for a Gaussian peak in a very noisy environment
- In a fraction f of the cases the right peak is found. It has a Gaussian uncertainty
- ► In a fraction 1 f of the cases a spurious peak is found. It can be anywhere in the searched range
- We know the real distribution of our measurements

# Maximum-likelihood estimation



## **Robust statistics**

## Stéphane Paltani

Why robust statistics?

Removal of data

\_-estimators

**R-estimators** 

**/**-estimators

Concepts

Maximum-likelihood estimation

- We obtain a sample  $\{x_i\}, i = 1, \ldots, N$
- The distribution of  $x_i$  is  $f \mathcal{N}(\mu, \sigma) + (1 f) \mathcal{U}(0, 100)$
- One can use a maximum likelihood to find parameters  $\mu$ ,  $\sigma$  and f!

$$L = \prod_{i} \frac{f}{\sqrt{2\pi\sigma}} \exp\left(-(x_i - \mu)^2/2\sigma^2\right) + \frac{1 - f}{100}$$

# Outline

Why robust statistics?

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**L**-estimators

**R**-estimators

**M**-estimators

Concepts Maximum-likelihood estimation So, these M-estimators...

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**/**-estimators

Concepts Maximum-likelihood estimation

# Model fitting with M-estimators

Let's take sample {x<sub>i</sub>; y<sub>i</sub>}, i = 1,..., N, whose errors on y<sub>i</sub> we know are not normally distributed, and a model y(x; p), where p is the parameters of the model. As above, we have:

 $L = \prod_{i} \exp\left(-\rho(y_i, y(x_i; \mathbf{p}))\right)$ 

 $\rho$  is the negative logarithm of the probability

We want then to minimize:

 $\sum_{i} \rho(\mathbf{y}_i, \mathbf{y}(\mathbf{x}_i; \mathbf{p}))$ 

Let's assume that ρ is local, i.e. ρ(y<sub>i</sub>, y(x<sub>i</sub>; p)) ≡ ρ(z), with z = (y<sub>i</sub> − y(x<sub>i</sub>; p))/λ<sub>i</sub>, i.e. ρ depends only on the difference with the model scaled by a factor λ<sub>i</sub>

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# Parameter estimation (cont.)

• Let's write 
$$\psi(z) \equiv \frac{d\rho(z)}{dz}$$

The minimum of L is obtained when:

$$0 = \sum \frac{1}{\lambda_i} \psi \left( \frac{y_i - y(x_i; \mathbf{p})}{\lambda_i} \right) \left( \frac{\partial y(x_i; \mathbf{p})}{\partial \mathbf{p}_k} \right)$$

- We can solve this equation, or we can minimize  $\sum_{i} \rho\left(\frac{y_{i}-y(x_{i};\mathbf{p})}{\lambda_{i}}\right)$
- $\psi(z)$  acts as a weight in the above equation

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## Some weights one can think of

- In the Gaussian case, put λ<sub>i</sub> = σ<sub>i</sub>, and we get the least-squares estimates
- But we have then:  $\rho(z) = z^2/2$  and  $\psi(z) = z$
- Two-sided exponential, P(z) ~ exp(−z):
  ρ(z) = |z| and ψ(z) = sgn(z)
- Lorentzian,  $P(z) \sim \frac{1}{1+z^2/2}$ :  $\rho(z) = \log(1+z^2/2)$  and  $\psi(z) = \frac{z}{1+z^2/2}$

#### Robust statistics

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