

Statistische Methoden der Datenanalyse

Hans Dembinski
IEKP, KIT Karlsruhe

Topics for today

- Confidence limits
- Monte-Carlo and resampling methods
- Testing of hypotheses

Confidence limits

Confidence intervals and limits

Confidence intervals from likelihood ratios (see Thursday's lecture) are always two-sided
What about one-sided limits? Fundamental way of constructing an interval?

Two-sided intervals are not unique

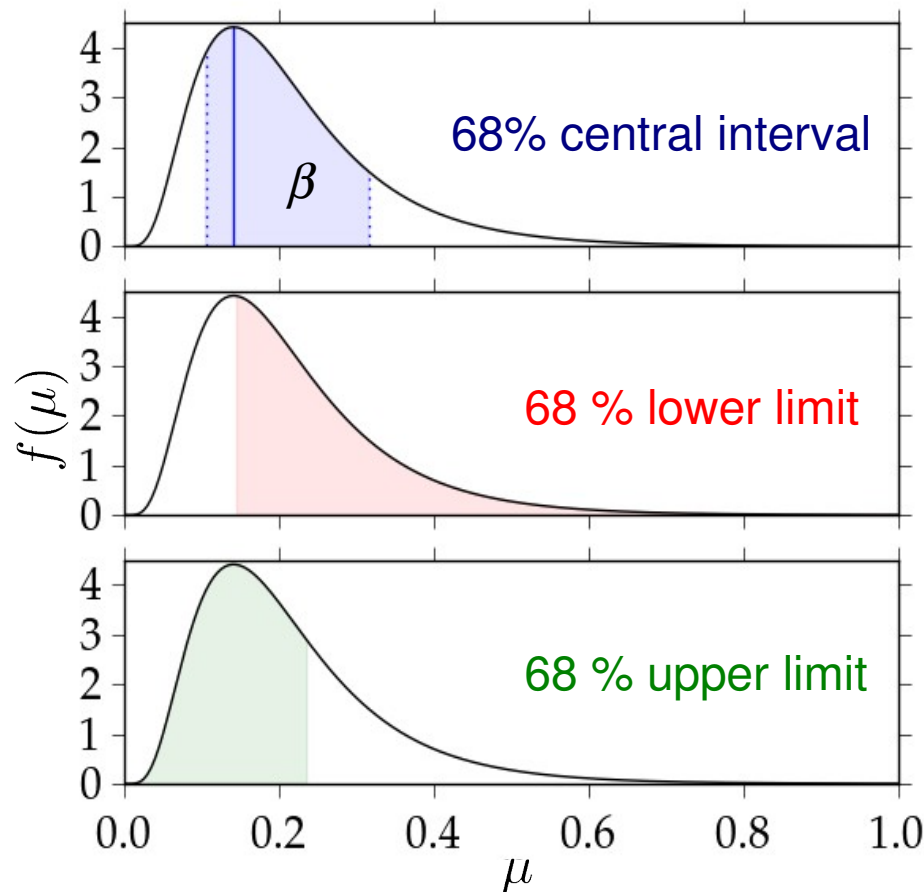
$$\beta = \int_{\mu_l}^{\mu_u} f(\mu) d\mu$$

Many p_l, p_u give same coverage β

Usual choice: **central interval**

$$\int_{-\infty}^{\mu_l} f(\mu) d\mu = \int_{\mu_u}^{\infty} f(\mu) d\mu = (1 - \beta)/2$$

No freedom of choice
for upper or lower limit

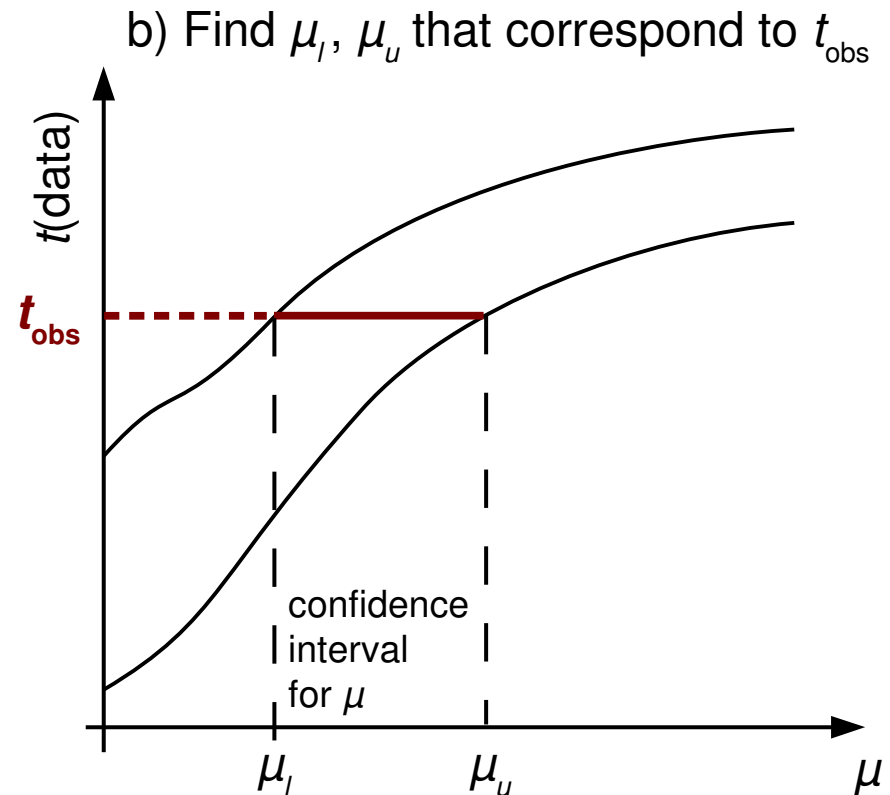
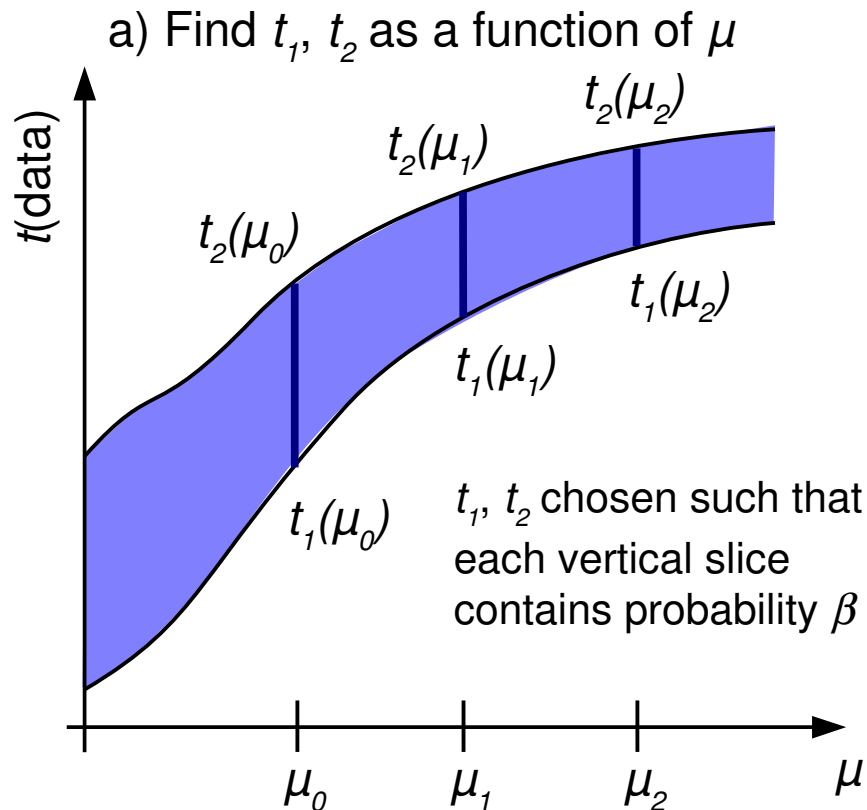


Neyman construction

Let us assume we have an estimator $t(\vec{x})$ of the data \vec{x} with known p.d.f. $f(t|\mu)$

We want to know the confidence interval for μ with coverage $C = \beta$

$$\beta = P[t_1 \leq t \leq t_2 | \mu] = P[t_1(\mu) \leq t \leq t_2(\mu)] = \int_{t_1}^{t_2} f(t|\mu) d\mu$$



Neyman construction

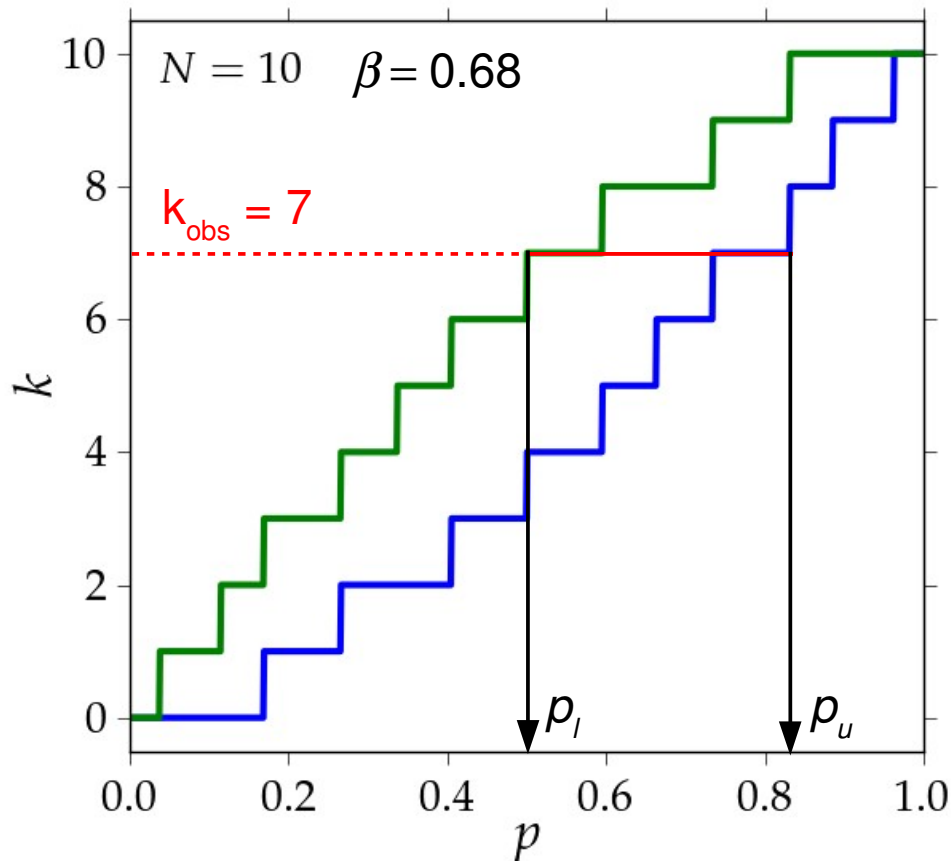
Neyman construction of confidence intervals for parameter p of binomial distribution

$$P(k|p, N) = \binom{N}{k} p^k (1 - p)^{N-k}$$

Two event classes A, B

Probability $p = P[A] = 1 - P[B]$

$P(k | p, N)$ probability of getting k events A out of N total



Neyman-constructed intervals

Note that $p_u > 0$ for $k = 0$

and $p_l < 1$ for $k = N$

Compare with **usual method**

$$\sigma[p] \approx \frac{\sqrt{V[k]}}{N} = \sqrt{\frac{\frac{k}{N}(1 - \frac{k}{N})}{N}}$$

where $\sigma[p] = 0$ for $k = 0$ and $k = N$

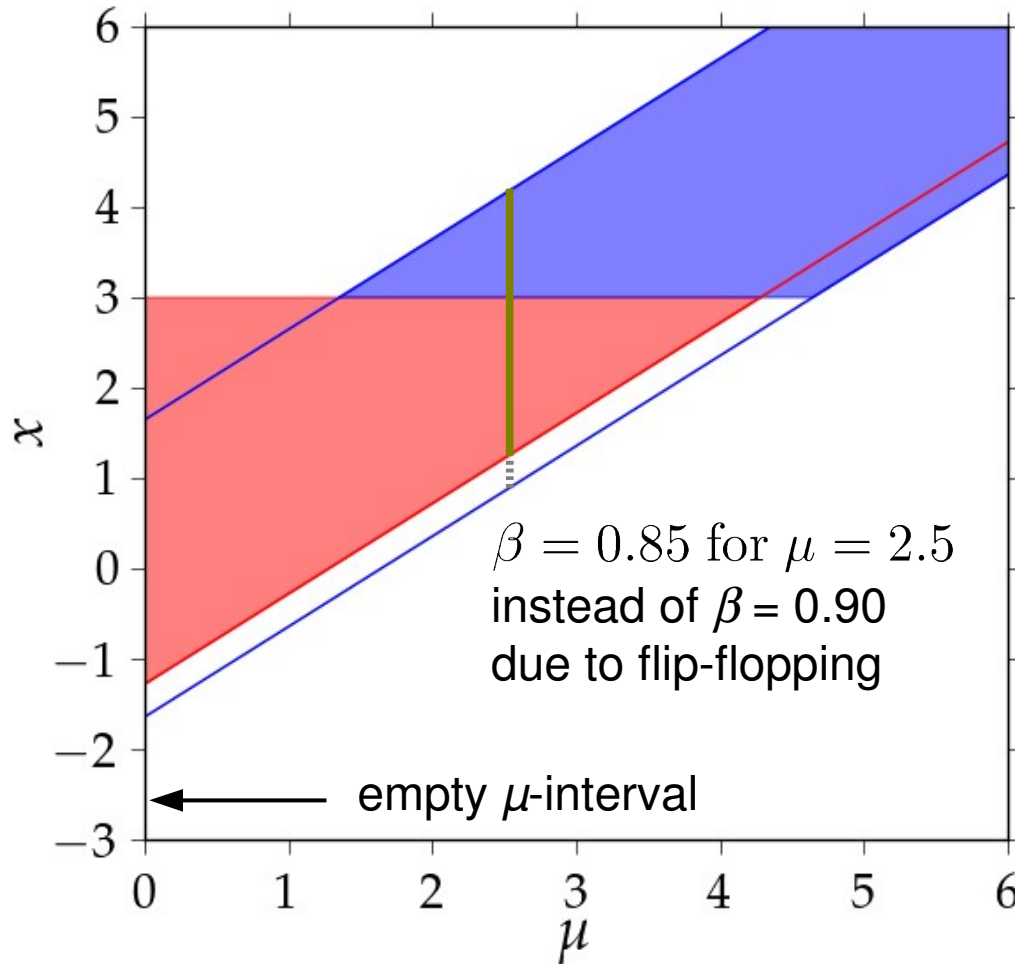
Beware:

Discrete distributions $C \geq \beta$

Continuous distributions $C = \beta$

Flip-flopping and empty intervals

Let's regard observation x from Normal distribution with $\mu > 0$ (physical constraint)
 $\sigma = 1$



Flip-flopping

Two choices for confidence interval, typical approach:

- Give two-sided limit if $x \gg 0$
- Give upper limit if $x \approx 0$

But: Switching method depending on data leads to **under-coverage**

Empty intervals

μ -intervals can be empty for $x \ll \mu$
due to constraint on $\mu > 0$



Solution: Feldman-Cousins limits

Feldman-Cousins limits

Unifies construction of two-sided limits and one-sided limits

Avoids empty intervals

Neyman construction + growth rule

Successively grow x -interval at the end with the largest likelihood ratio $L(x|\mu)/L(x|\hat{\mu})$

$$\frac{L(x_1|\mu)}{L(x_1|\hat{\mu})} \quad \hat{\mu} \quad \frac{L(x_2|\mu)}{L(x_2|\hat{\mu})}$$

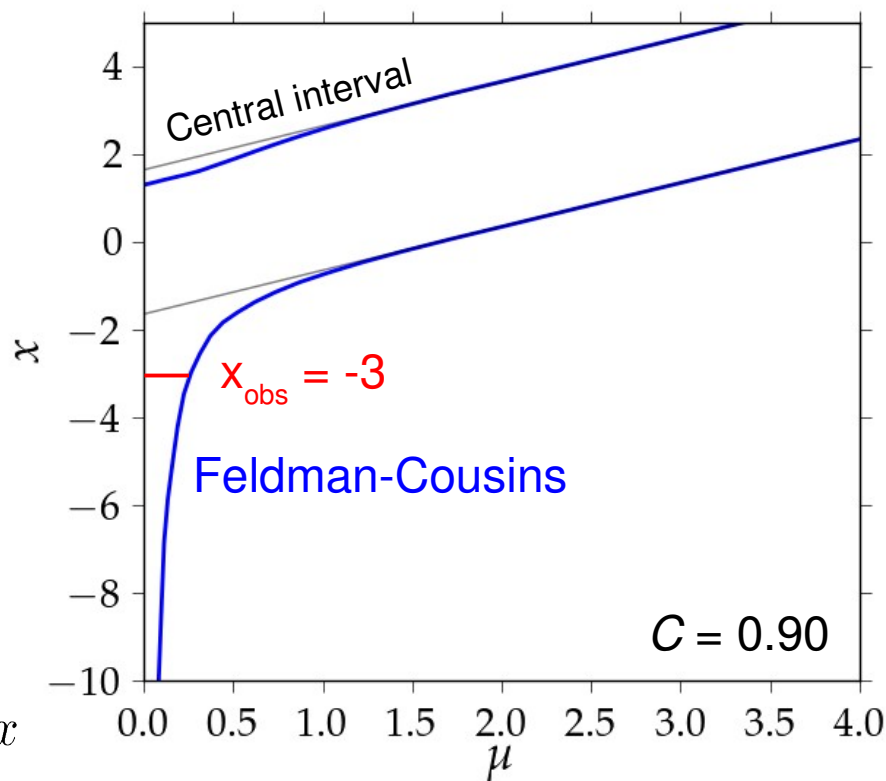
— — —
 $x_1, x_1 + dx \quad \quad \quad x_2, x_2 + dx$

$\hat{\mu}$ is the maximum likelihood estimate of μ given x under the condition $\hat{\mu} \geq 0$

Example: Normal distribution $\mu > 0, \sigma = 1$

$$L(x|\hat{\mu}) = \begin{cases} \frac{1}{\sqrt{2\pi}}, & x \geq 0, \hat{\mu} = x \\ \frac{1}{\sqrt{2\pi}} \exp(-x^2/2), & x < 0, \hat{\mu} = 0 \end{cases}$$

$$\frac{L(x|\mu)}{L(x|\hat{\mu})} = \begin{cases} \exp(-(x - \mu)^2/2), & x \geq 0 \\ \exp(x\mu - \mu^2/2), & x < 0 \end{cases}$$



Feldman-Cousins construction is recommended if you want to report a result close to a physical boundary

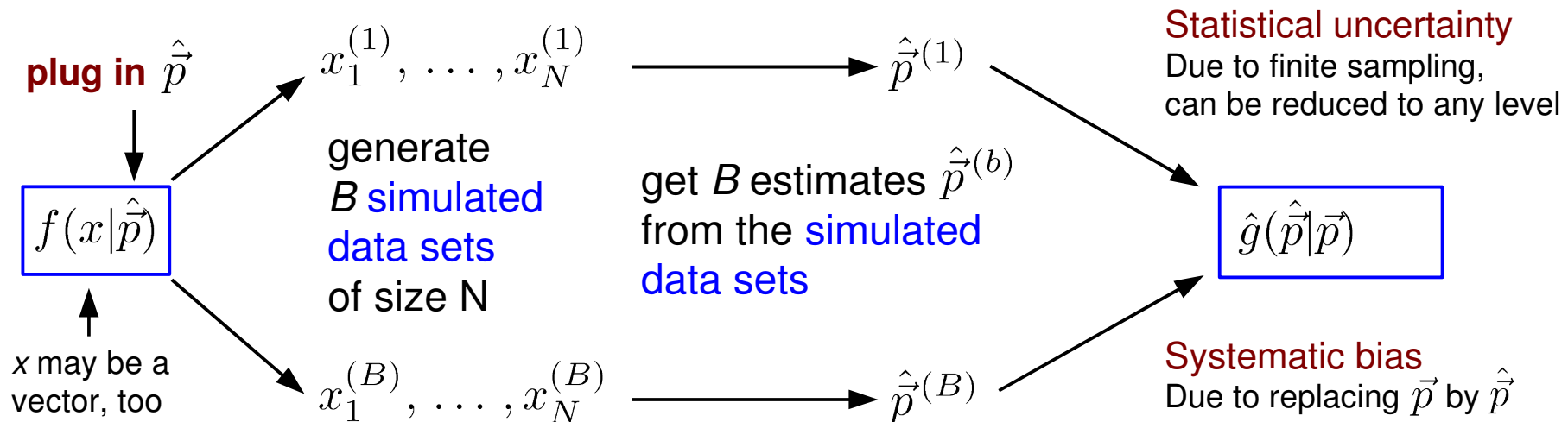
Monte-Carlo and resampling methods

Parametric bootstrap

Let's assume we have the p.d.f. $f(x|\vec{p})$ for an observation x given parameters \vec{p} and an estimate $\hat{\vec{p}}$ obtained from N observations x_i

We want to know $g(\hat{\vec{p}}|\vec{p})$ or a summary statistic like bias and variance of $\hat{\vec{p}}$

Monte-Carlo method (= parametric bootstrap)



Bias of $\hat{\vec{p}}$

$$\hat{E}[\hat{\vec{p}} - \vec{p}] = \frac{1}{B} \sum_b \hat{\vec{p}}^{(b)} - \hat{\vec{p}}$$

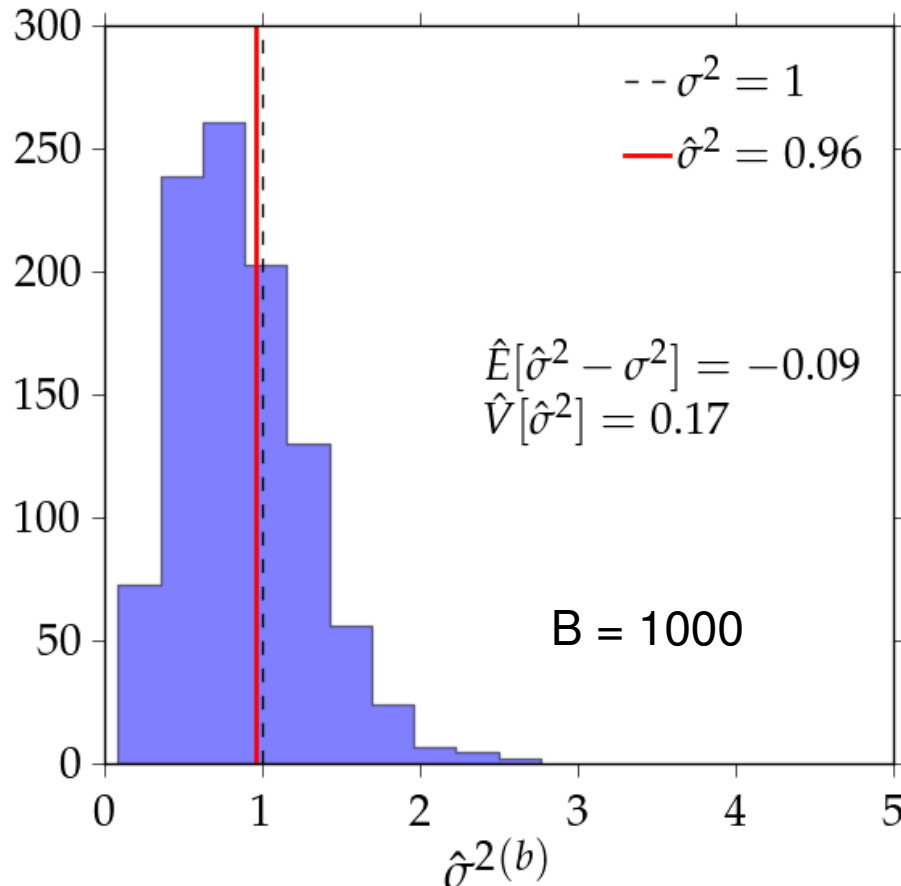
Variance of $\hat{\vec{p}}$

$$\widehat{\text{cov}}[\hat{\vec{p}}]_{ij} = \frac{1}{B-1} \sum_b \hat{p}_i^{(b)} \hat{p}_j^{(b)} - \frac{1}{B(B-1)} \left(\sum_b \hat{p}_i^{(b)} \right)^2$$

Parametric bootstrap

Example: Normal distribution $\mu = 0$, $\sigma = 1$, $N = 100$

Study biased estimator
$$\hat{\sigma}^2 = \frac{1}{N} \sum_i x_i^2 - \frac{1}{N^2} \left(\sum_j x_j \right)^2$$



Analytical results for normal distribution

$$E[\hat{\sigma}^2 - \sigma^2] = -\frac{\sigma^2}{N} = -0.1$$

$$V[\sigma^2] = 2\sigma^4 \frac{N-1}{N^2} \approx 0.18$$

Parametric bootstrap

- + Bias and variance without analytical effort
- + Works with arbitrarily complex estimators
 - o Computationally intensive
- Systematic bias can be important if \hat{p} is far away from p

Better performance for large N

Random number generation

Scientific programming libraries provide excellent pseudo random number generators

Pseudo random numbers have uniform (flat) distribution, how to get arbitrary $f(\vec{x})$?

a) Transformation method $y = F(x) = \int_{-\infty}^x dx f(x) \rightarrow x = F^{-1}(y)$

↑
follows uniform distribution

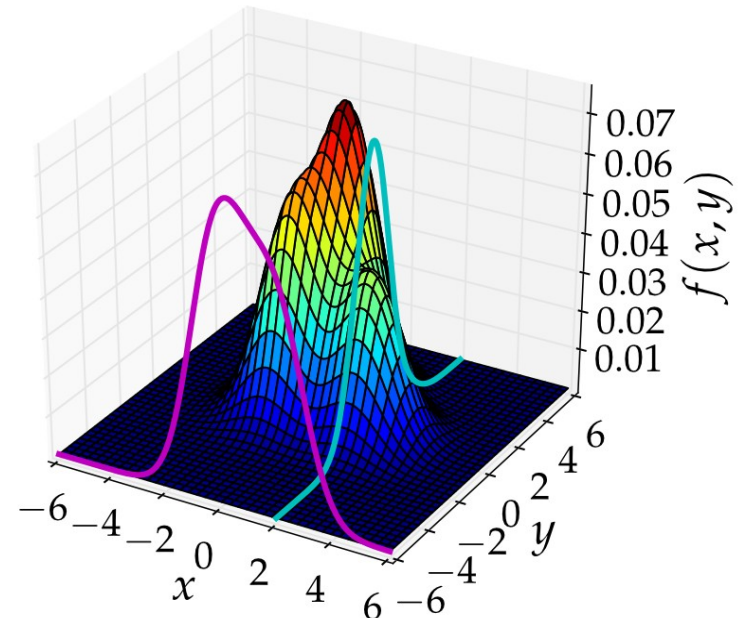
Practical only if $F^{-1}(y)$ or a suitable approximation to it is available

Multivariate case complex, an example in 2d:

Solve in order

$$\int_{-\infty}^{x_0} dx'_0 \int_{-\infty}^{\infty} dx'_1 f(x'_0, x'_1) = y_0$$
$$\frac{\int_{-\infty}^{x_1} dx'_1 f(x_0, x'_1)}{\int_{-\infty}^{\infty} dx'_1 f(x_0, x'_1)} = y_1$$

Analog in case of more dimensions



Random number generation

Scientific programming libraries provide excellent pseudo random number generators
Pseudo random numbers have uniform (flat) distribution, how to get arbitrary $f(\vec{x})$?

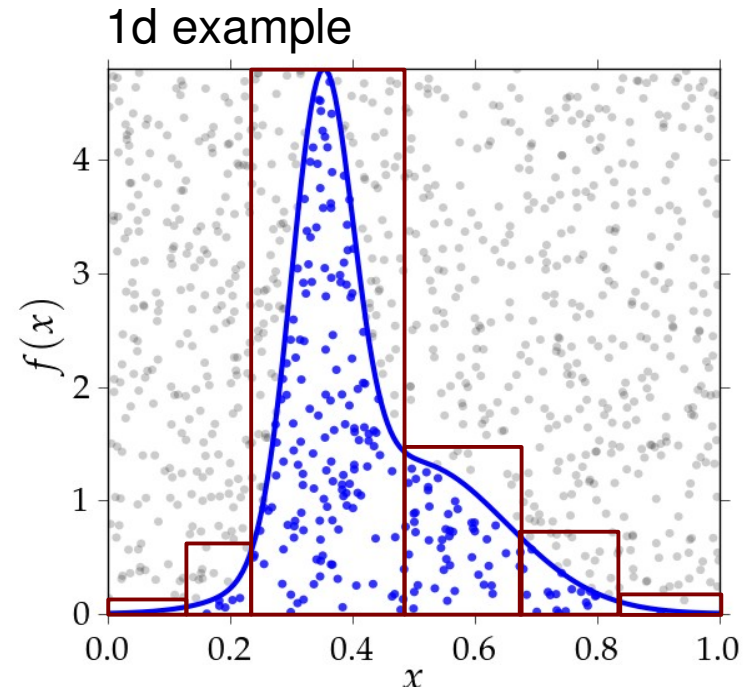
b) Accept-Reject method

Construct a (hyper-)rectangle around $f(\vec{x})$ that completely encloses it

Uniformly draw points (\vec{x}, f') from inside the (hyper-)rectangle and accept \vec{x} if $f' < f(\vec{x})$

- + Very general method
- + Simple to set up
- o Need to know $\max[f(x)]$
- Inefficient/slow: many points are wasted

Efficiency is greatly improved by
sampling from several local boxes



Full bootstrap

What to do if $f(x|\vec{p})$ is unknown?

We could still use the Monte-Carlo method to study an estimator $t(\vec{x})$ of the data \vec{x} if we had an estimate of $f(x)$

Non-parametric maximum-Likelihood estimate of $f(x)$

maximize $\ln \hat{f}(x) = \sum_i \ln \hat{f}(x_i)$ without any further knowledge except $\int_{-\infty}^{\infty} \hat{f}(x) = 1$

non-parametric
ML estimate

$$\hat{f}_B(x) = \frac{1}{N} \sum_i \delta(x - x_i)$$

Assumptions: x_i from same $f(x)$, x_i are independent

No proof, but...

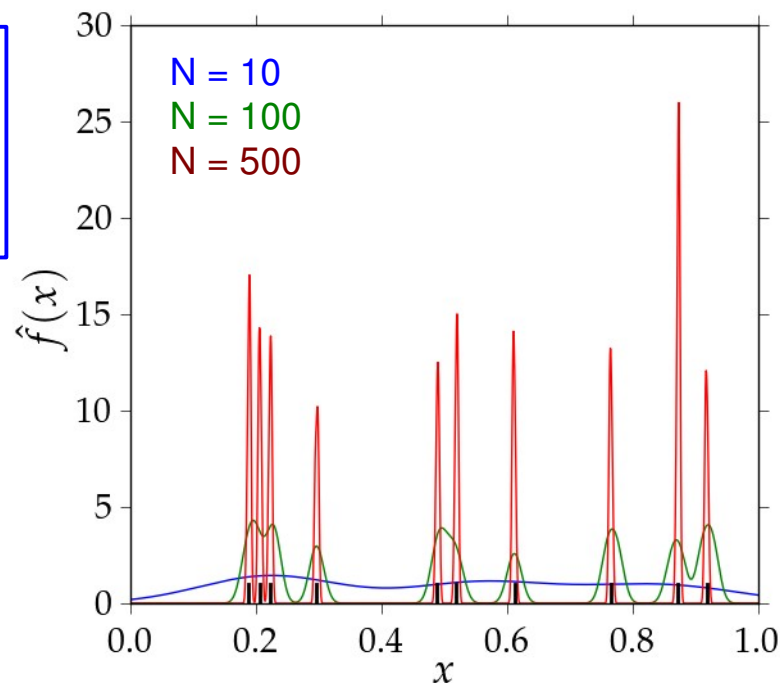
$$f(x|\vec{a}) = \frac{1}{\sum_j a_j} \sum_{k=0}^K a_k g(x|\mu_k, \sigma)$$

$$\mu_k = \frac{k}{N-1} \Delta x$$

$$\sigma = \frac{1}{N-1} \Delta x$$

$g(x|\mu, \sigma)$ Normal p.d.f.

converges to $\hat{f}_B(x)$ for $K \rightarrow \infty$ (infinite flexibility)



Analytic bootstrap estimates

Plugin principle

Construct bootstrap estimate by replacing true variable in formula by empirical one

$$E_B[x] = \int dx x \hat{f}(x) = \int dx x \frac{1}{N} \sum_i \delta(x - x_i) = \frac{1}{N} \sum_i x_i \quad \text{sample mean}$$

$$V_B[x] = E_B[x^2] - E_B[x]^2 = \frac{1}{N} \sum_i x_i^2 - \left(\frac{1}{N} \sum_j x_j \right)^2 \quad \text{sample variance (biased)}$$

Like any estimator, a bootstrap estimator can be biased if the sample size is small (Bias can be detected and corrected by a double bootstrap, i.e. bootstrapping the bootstrap)

Two other bootstrap estimates are well known to physicists

Uncertainty of a Poisson count $k \pm \sqrt{k}$: $V[\lambda] = \lambda \rightarrow V_B[\lambda] = k$

Uncertainty of a binomial proportion (e.g. efficiency of a detector):

$$V[k/N] = \frac{p(1-p)}{N} \rightarrow V_B[k/N] = \frac{k/N(1-k/N)}{N}$$

Monte-Carlo bootstrap estimates

Use in Monte-Carlo estimation

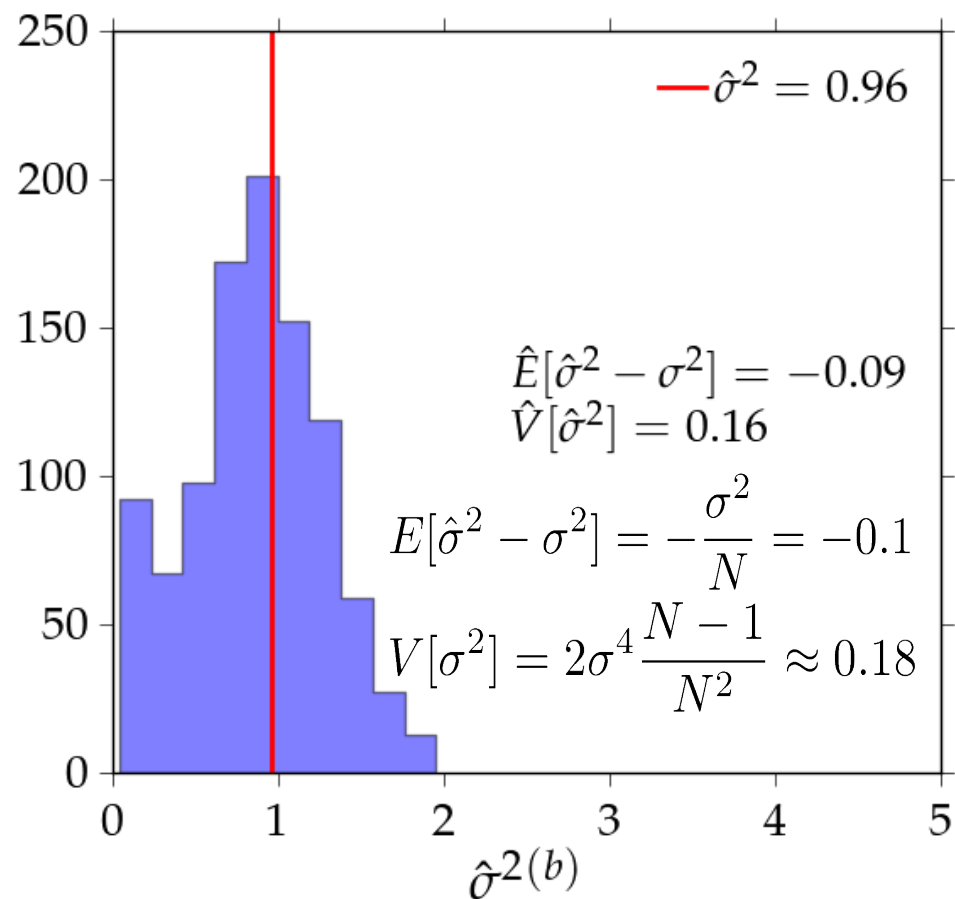
Draw random numbers from $\hat{f}(x)$
Pick x_i with equal probability with replacement

Re-examination

Normal distribution $\mu = 0$, $\sigma = 1$, $N = 100$
and biased estimator

$$\hat{\sigma}^2 = \frac{1}{N} \sum_i (x_i - \frac{1}{N} \sum_j x_j)^2$$

- o Same pros/cons as parametric bootstrap
- + Effortless to apply
- Biased for estimators that depend strongly on distribution tails



Other resampling methods

Jackknife – fast approximation to full bootstrap

$$\hat{E}_{\text{jack}}[\hat{p} - p] = \frac{N-1}{N} \sum_j (\hat{p}_{(j)} - \hat{p}) \quad \hat{V}_{\text{jack}}[\hat{p}] = \frac{N-1}{N} \sum_i \hat{p}_{(i)}^2 - \frac{N-1}{N^2} \left(\sum_j \hat{p}_{(j)} \right)^2$$

$\hat{p}_{(j)} = t(x_1, \dots, x_{j-1}, x_{j+1}, \dots, x_N)$ estimate of p without observation x_j

Only needs N additional evaluations of $t(\vec{x})$, but less precise

Leave-one-out cross-validation – compare prediction power of models

Can only be used with (x, y) pairs, $y = f(x)$

$$\text{LOOCV} = \sum_i (y_i - f_{(i)}(x_i))^2 \propto \text{mean squared error} = \text{bias}^2 + \text{variance}$$

bias² is large if model is not flexible enough, i.e. is missing effects in the data

variance is large if model is too flexible, i.e. “overfitting” the data

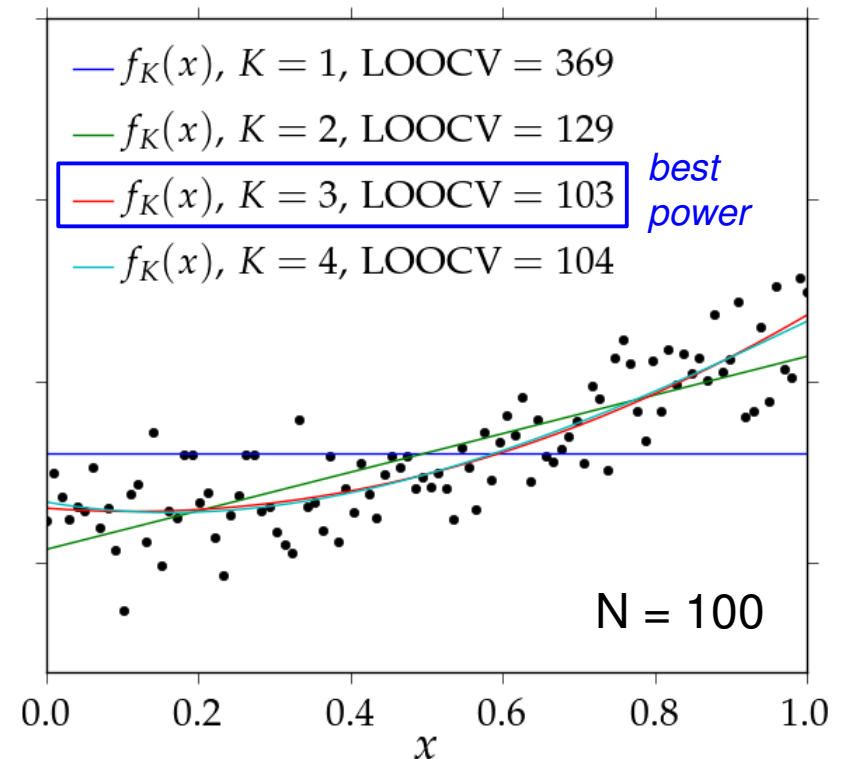
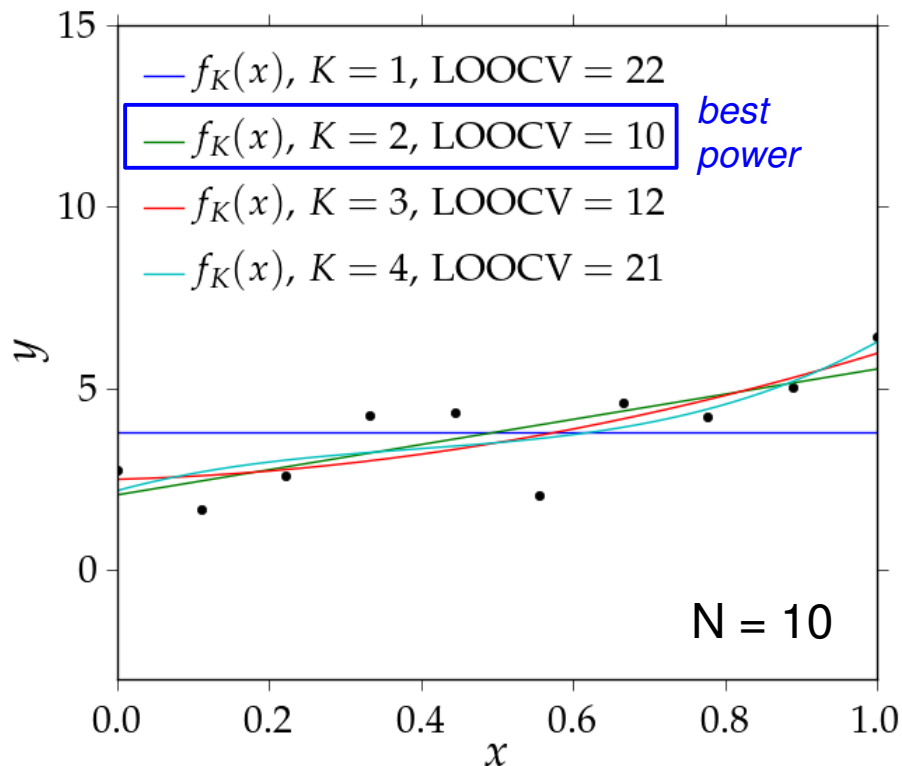
Model with best prediction power has smallest LOOCV value

Maximizing prediction power

Example: fit of a polynomial model

True model $f(x) = 1 + 2x + 3x^2$, $y_{obs} = f(x) + \text{Normal fluctuation with } \mu = 0, \sigma = 1$

Fitting model $f_K(x) = \sum_{k=0}^K p_k x^k$, what K to choose if K is unknown?



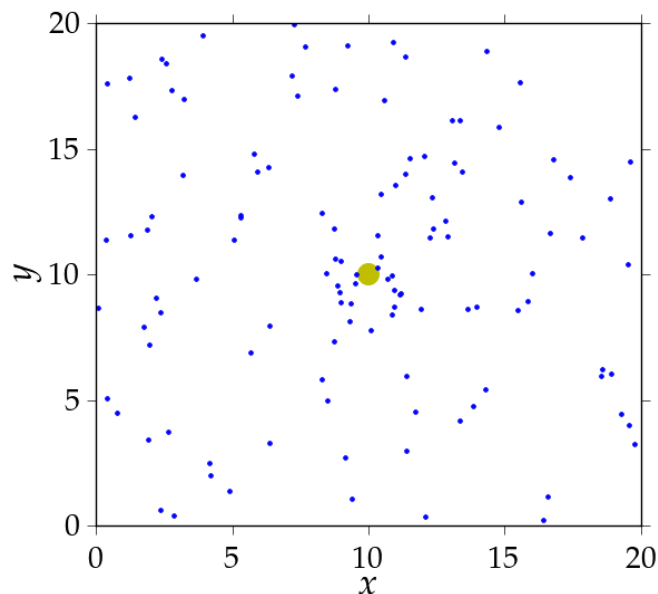
Testing hypotheses

Humor



<http://xkcd.com/892>

Testing hypotheses



Introductory example

Does a fraction of the sky contain a source of cosmic rays?

Hypothesis H_0 (“background hypothesis”)

There is only background and no source

Hypothesis H_1 (“signal hypothesis”)

There is background and a source!

Lots of special cases (see literature), most common one for Physicists:

$$f(\vec{x}) = (1 - s)f_B(\vec{x}) + sf_S(\vec{x}|\vec{p}_S)$$

$$H_0 : s = 0 \quad H_1 : s > 0$$

→ *Continuous family of hypotheses*

We need a **test statistic** that discriminates between H_0 and H_1

$$-2 \ln \lambda = -2 \ln \left[\frac{\max L(s = 0, \vec{p}_S = 0)}{\max L(s, \vec{p}_S)} \right]$$

likelihood ratio is asymptotically the **most powerful test statistic**

Type I and type II errors

Hypothesis tests are fully characterized by their Type I and Type II errors

Type II error β

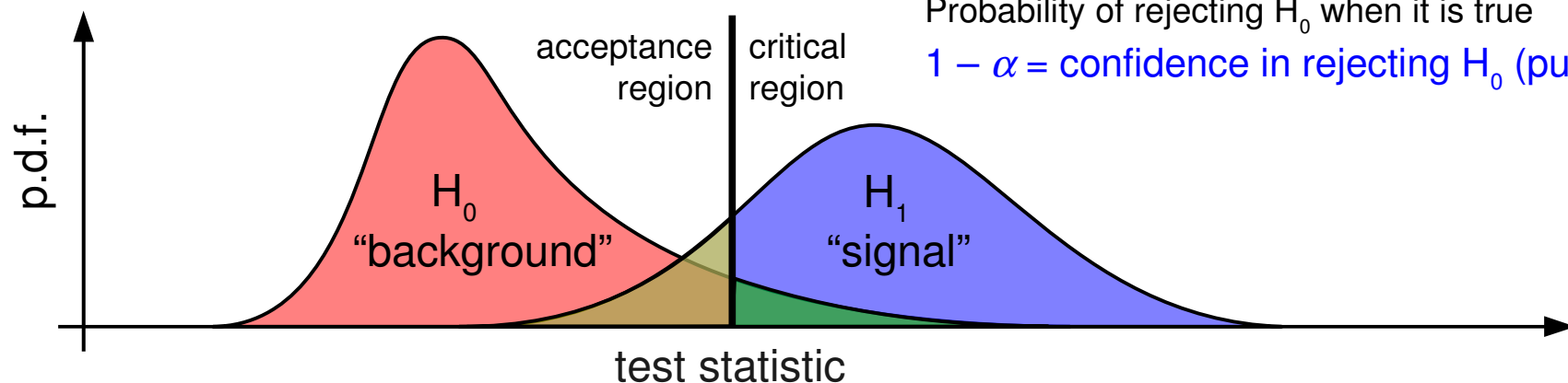
Probability of accepting H_0 when it is false

$1 - \beta = \text{power of the test (efficiency)}$

Type I error α

Probability of rejecting H_0 when it is true

$1 - \alpha = \text{confidence in rejecting } H_0 \text{ (purity)}$



Desired confidence $1 - \alpha$ defines the critical region, so tests are compared by their power $1 - \beta$

In our case, $1 - \beta$ cannot be calculated, since H_1 is not fully determined

Fortunately, only H_0 is needed to determine the critical region

Test must be completely defined **before** seeing the data

Confidence of rejecting H_0 **is not** confidence in choosing H_1 ($\alpha \neq \beta$)

Critical region

How to determine critical region for given confidence $1-\alpha$?

$$f(\vec{x}) = (1-s)f_B(\vec{x}) + sf_S(\vec{x}|\vec{p}_S)$$

$$H_0 : s = 0$$

$$H_1 : s > 0$$

$$-2 \ln \lambda = -2 \ln \left[\frac{\max L(s=0, \vec{p}_S=0)}{\max L(s, \vec{p}_S)} \right]$$

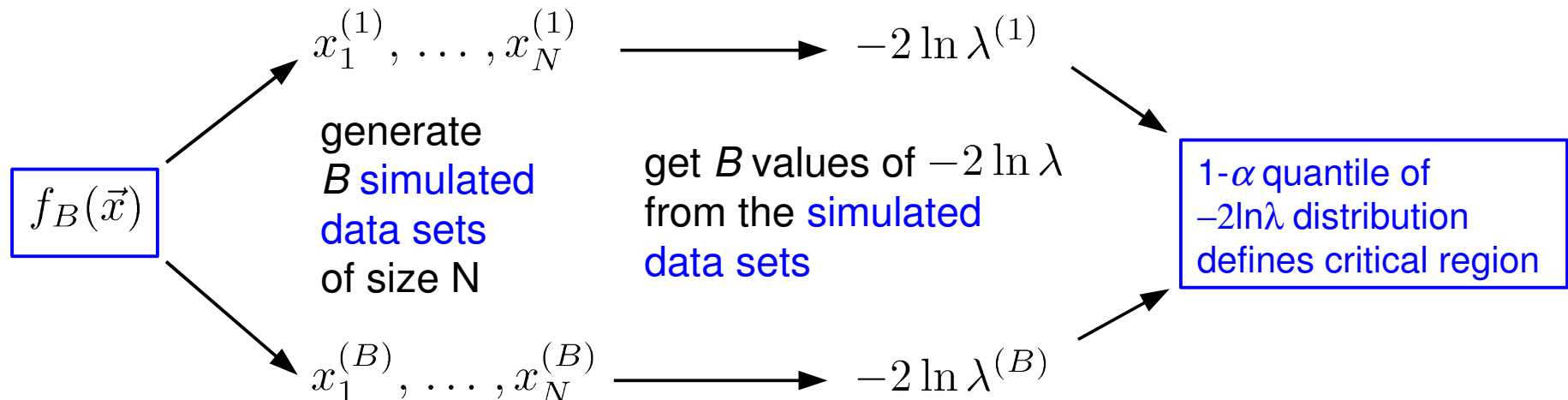
is asymptotically distributed as $\chi^2(r)$

r = number of parameters

fixed by H_0 but left free by H_1

Asymptotic properties are nice, but test usually used with small data sets...

Recommended: Monte-Carlo-based determination of critical region



Critical region – example

In our example

Background: 2d uniform distribution

Signal: 2d normal distribution

$$f(x, y) = (1 - s)f_B(x, y) + sf_S(x, y)$$

$$f_B(x, y) = \frac{1}{\Delta x \Delta y}$$

$$f_S(x, y) = \frac{1}{2\pi} \exp \left[-\frac{1}{2}(x^2 + y^2) \right]$$

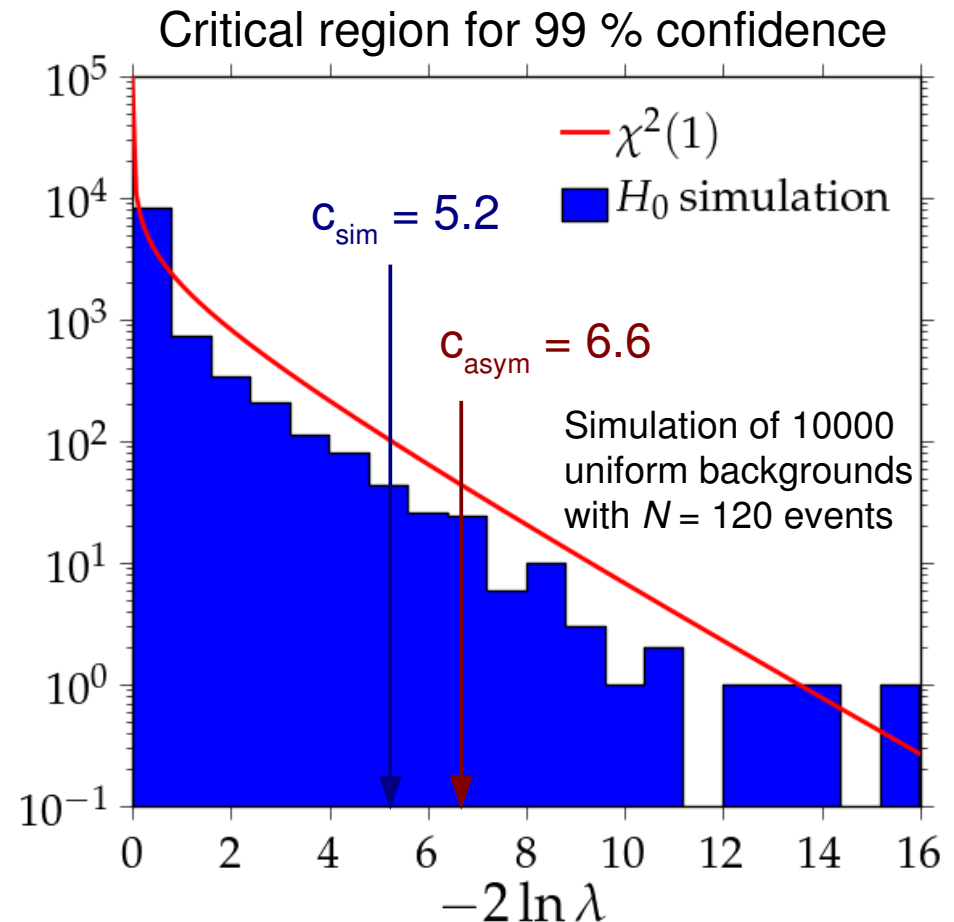
$$H_0 : \lambda_S = 0 \quad H_1 : \lambda_S > 0$$

s is free in H_1 , but fixed in H_0

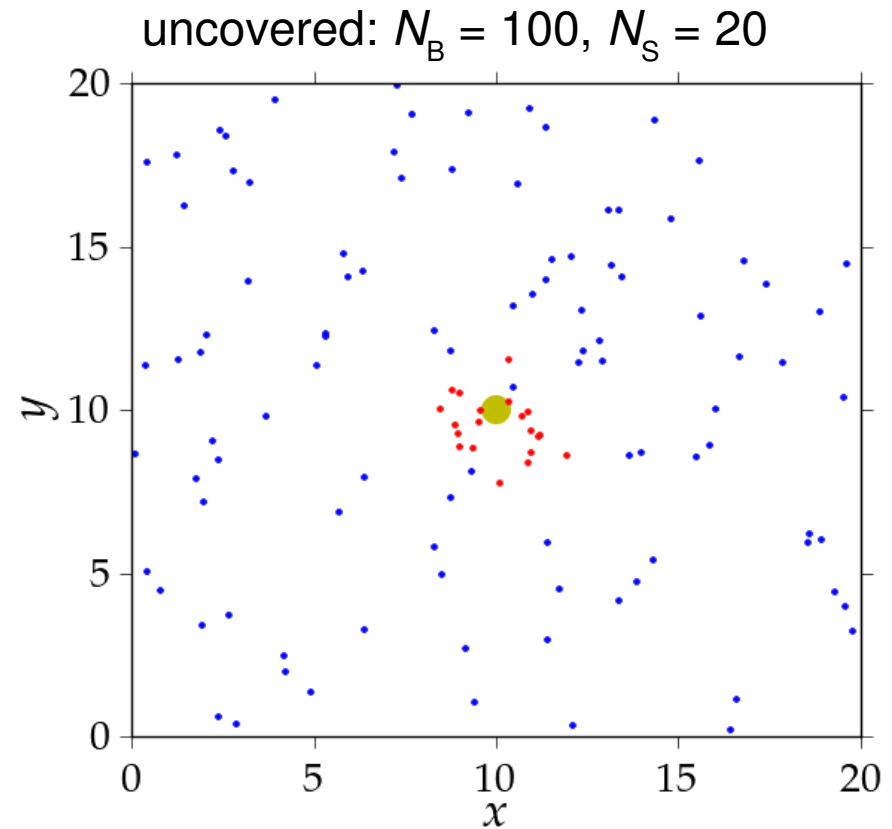
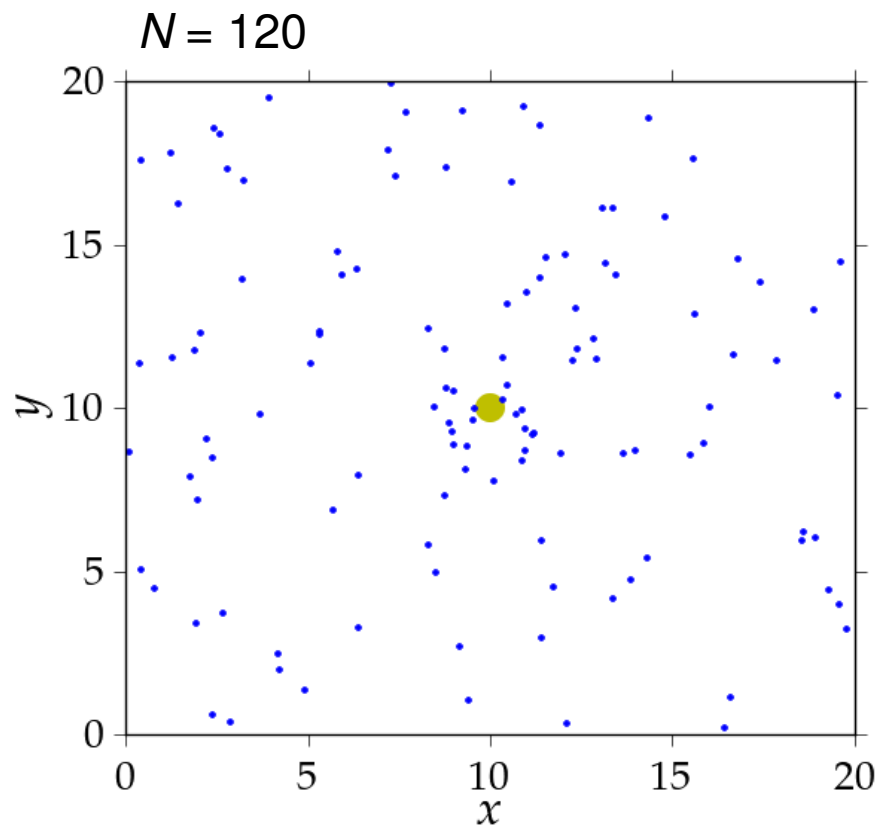
no other free parameters

→ asymptotic distribution of $-2\ln\lambda$ is $\chi^2(1)$

Reject H_0 if in real data set $-2\ln\lambda > c$



Hypothesis test – example



Test statistic
in real data $-2 \ln \lambda = 38.3 > c$



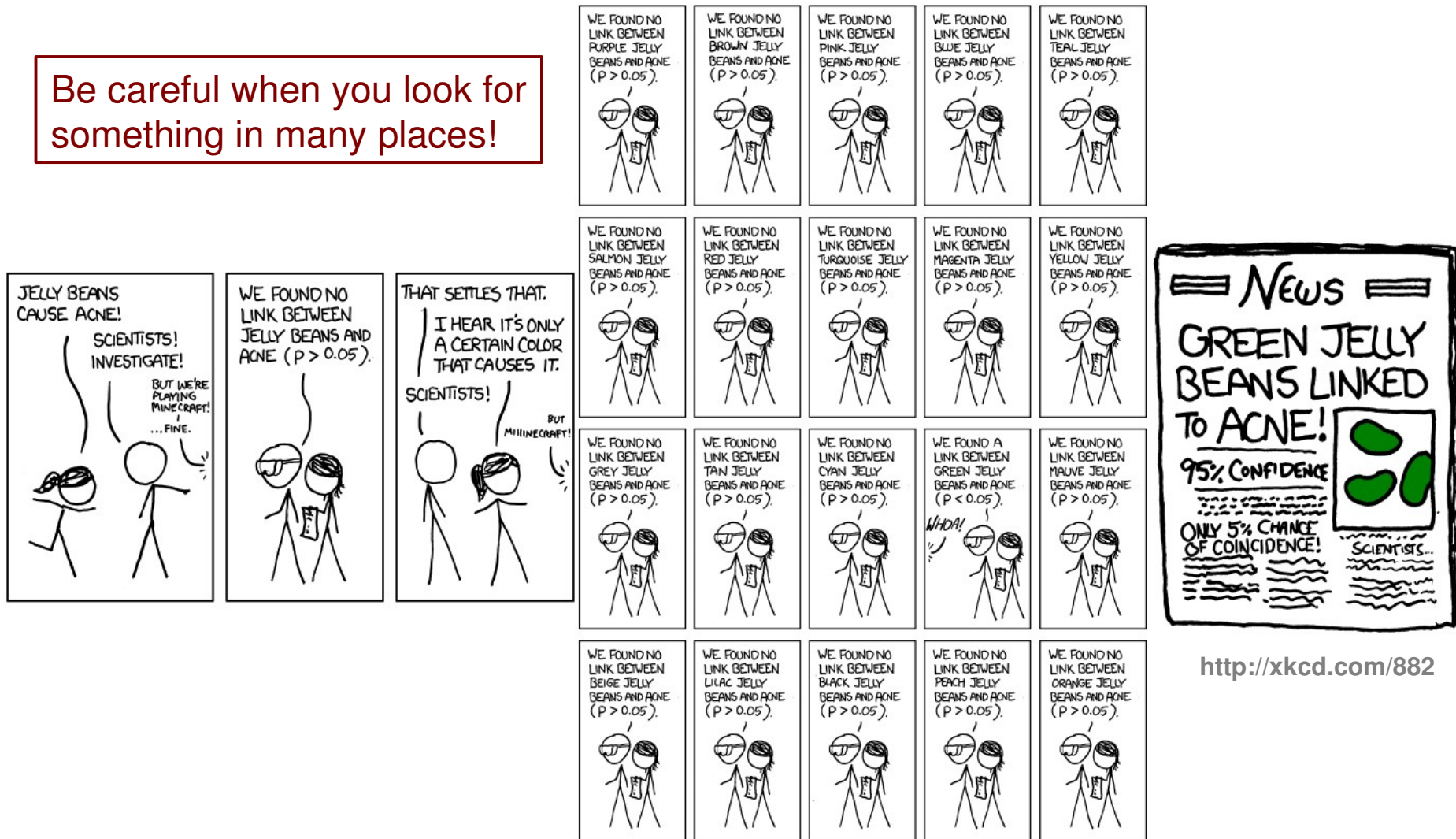
We reject the background-only hypothesis H_0
with a confidence of at least 99 %

Confidence does **not** increase even if $-2 \ln \lambda \gg c$! (property of test, not of data)

Trial factors

a.k.a. Look-Elsewhere-Effect

Be careful when you look for something in many places!



Trial factors – example

Our example revisited with **signal location unknown**

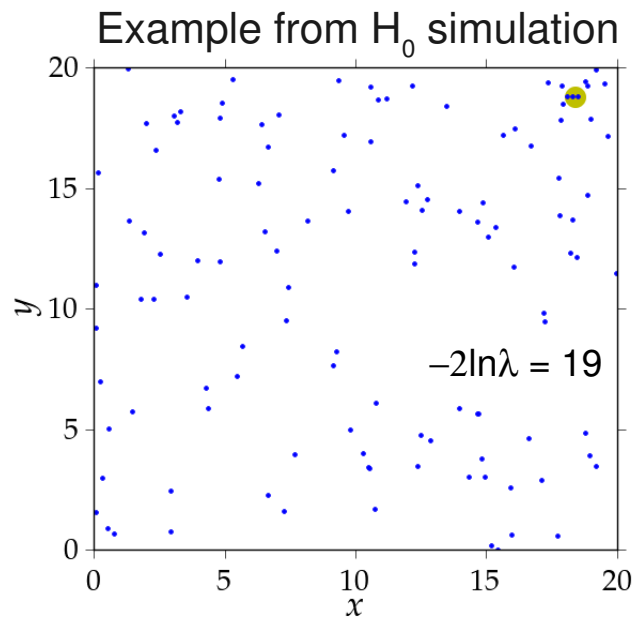
$$f_S(x, y | \mu_x, \mu_y) = \frac{1}{2\pi} \exp \left[-\frac{1}{2} \left((x - \underline{\mu_x})^2 + (y - \underline{\mu_y})^2 \right) \right]$$

source position free
in signal model

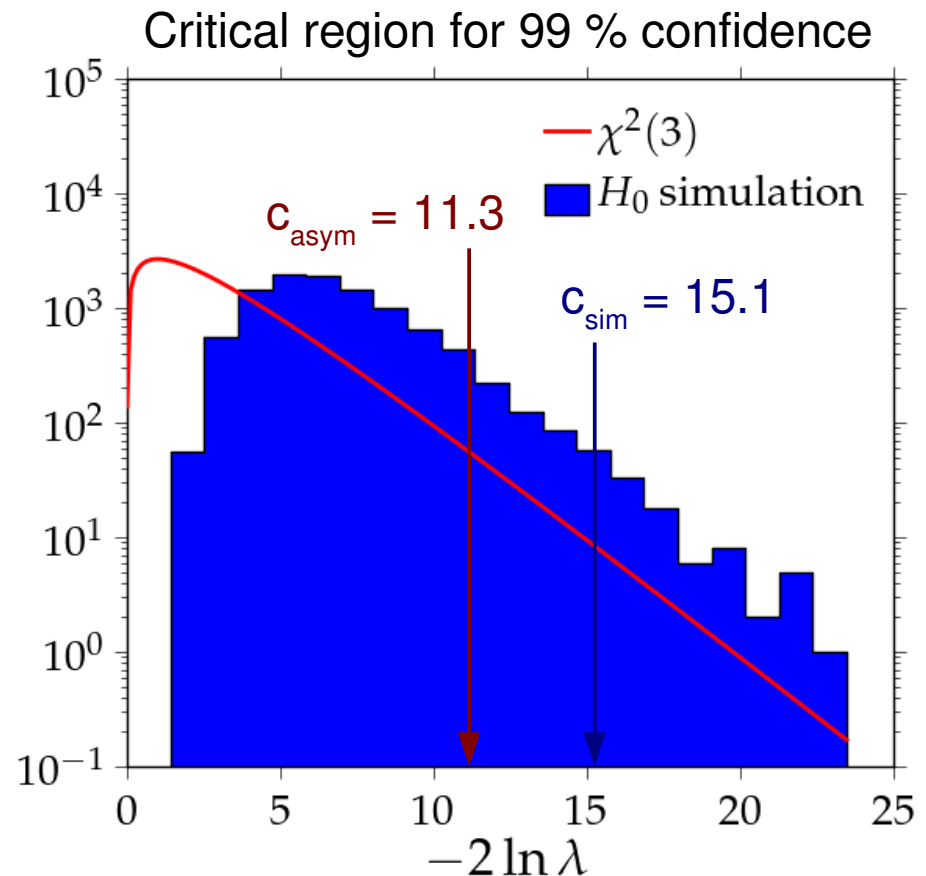
Samples from uniform distribution
tend to form clusters

→ Clusters appear like sources

→ large $-2\ln\lambda$ more frequent



Natural form of apophenia!



Goodness-of-fit tests

Goodness-of-fit (GOF) test = Lesser form of Hypothesis test

Test of H_0 with against all possible other hypotheses: H_1 completely unspecified

→ Power $1-\beta$ unknown

Components of a GOF test

Test statistic t and c.d.f. $F(t) = \int_t^{-\infty} dt' f(t'|H_0)$ to convert t into P-value

Small P-values indicate “bad fit” of model to data

P-value = $P(\text{data}|H_0)$ **is not** $P(H_0|\text{data})$

Large P-values are not evidence in favor of H_0 !

Some GOF test statistics are independent of H_0 (distribution-free) $f(t|H_0) = f(t)$

e.g. Pearson's **Chi-square test** and Smirnov-Cramér-von Mises' test

(for data pairs and binned data)

(for unbinned data)

For combined tests calculate P-value from Monte-Carlo simulations of H_0

Pearson's Chi-square test

Idea: sum up squares of normalized residuals of data points around model

$$t = (\vec{y} - \vec{f}(\vec{x}))^T \tilde{V}^{-1} (\vec{y} - \vec{f}(\vec{x})) = \sum_{i=1}^N \left(\frac{y_i - f(x_i)}{\sigma_i} \right)^2 = \sum_{i=1}^N z_i^2$$

↑
if y_i are uncorrelated

z_i have normal distribution with $\mu = 0$, $\sigma = 1$ independent of H_0

If y_i are correlated, one can find transformation to decorrelate them and get same result

$$E[t] = \sum_{i=1}^N E[z_i^2] = N \quad V[t] = \sum_{i=1}^N V[z_i^2] = 2N \quad f(t) = \frac{\frac{1}{2} \left(\frac{t}{2}\right)^{N/2-1} e^{-t/2}}{\Gamma\left(\frac{N}{2}\right)}$$

If $f(x)$ has k free parameters fitted to the y_i , replace N by $N - k$

No formal proof here, but intuition:

Due to fit of $f(x)$, z_i are no longer independent $\rightarrow k$ "degrees of freedom" lost

Smirnov-Cramér-von Mises test

$$t = \int_{-\infty}^{\infty} dx [\hat{F}(x) - F(x)]^2 f(x) \quad \text{is independent of } f(x) (= H_0)$$

$$\hat{F}(x) = \int_{-\infty}^x dx' \hat{f}(x) = \frac{1}{N} \int_{-\infty}^x dx' \sum_i \delta(x - x_i) = \frac{1}{N} \sum_i H(x - x_i)$$

$H(x)$ Heaviside step function

Proof: insert substitution $y = F(x) \longrightarrow t = \int_{-\infty}^{\infty} dy [\hat{F}(y) - y]^2$

$$E[t] = \frac{1}{6N} \quad V[t] = \frac{4N - 3}{180N^3} \quad \text{no } f(t) \text{ in closed form} \rightarrow \text{tables}$$

Based on
asymptotic
distribution,
reached for $N \geq 3$

Confidence level $1-\alpha$	Critical value of $N t$
0.90	0.347
0.95	0.461
0.99	0.743
0.999	1.168

Backup

General formula for any distribution

$$V[\hat{\sigma}^2] = \frac{1}{N} \left(\mu_4 - \frac{N-3}{N-1} \sigma^4 \right)$$

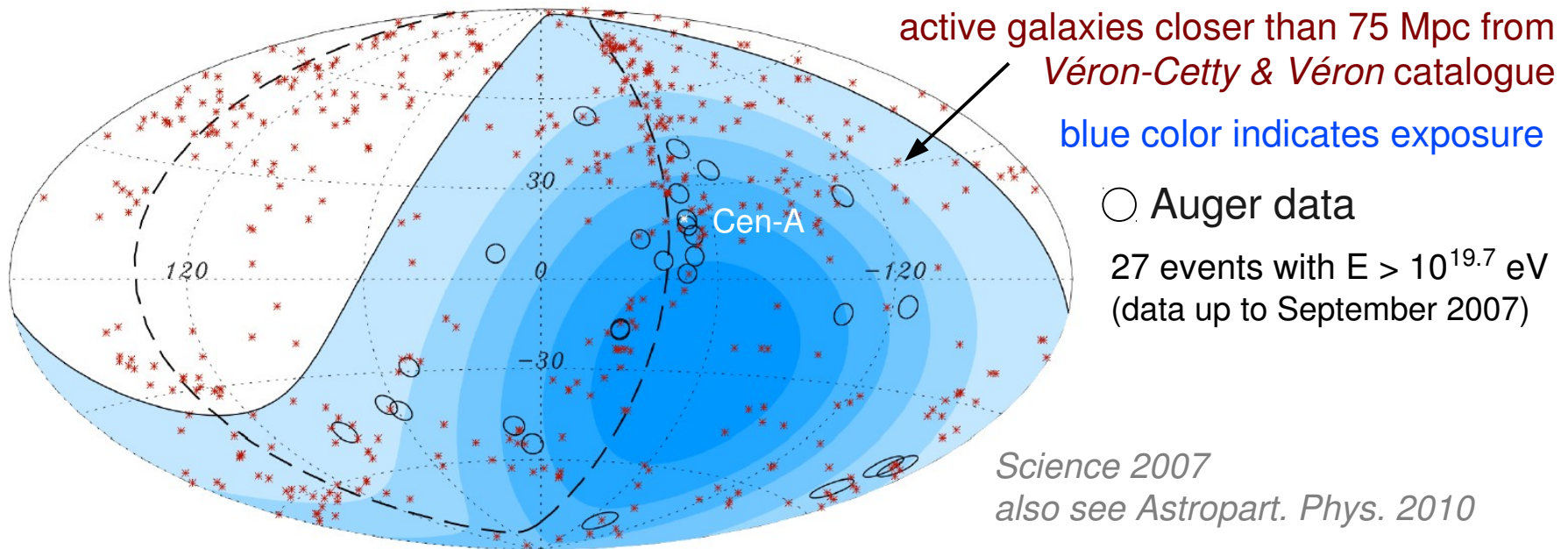
$$\text{with } \mu_4 = \frac{1}{N} \sum_i (x_i - \mu)^4$$

Hypothesis probability after seeing data

Testing hypotheses

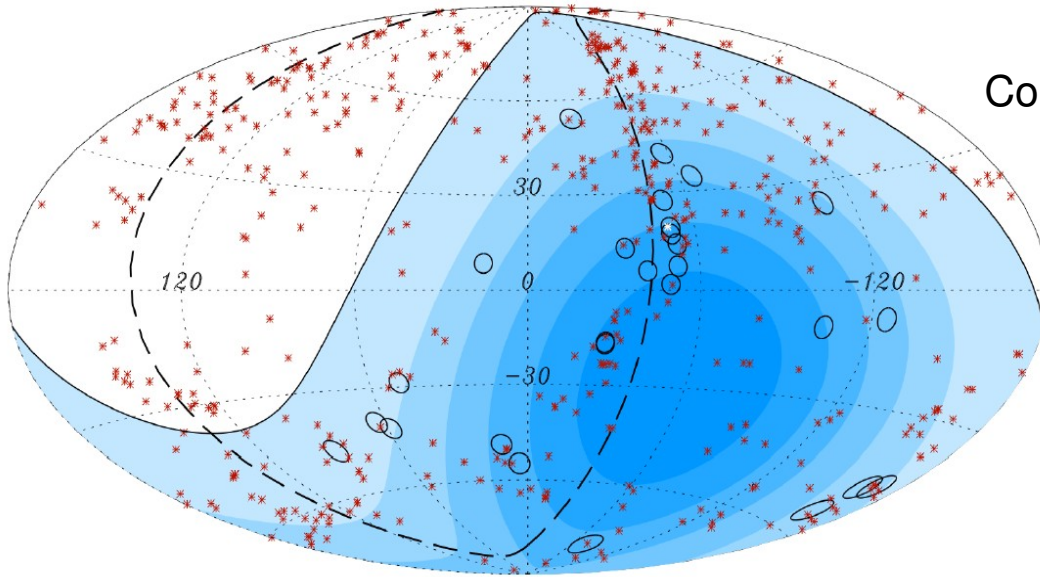
Recent example: structures in cosmic ray sky found by the Pierre Auger Observatory

sky map of CR arrival directions in galactic coordinates



UHECR sky seems anisotropic, let's reject the hypothesis H_0 [CRs are isotropic]!
With what confidence can we do it? or What is the **probability to be mistaken**?

Testing hypotheses



Correlation = $\text{angle}(\text{cosmic ray, AGN}) < 3.2^\circ$

Test statistic

Number of correlating events k out of N
(k follows binomial distribution)

H_0 prediction (isotropy)

21 % of cosmic rays correlate $\rightarrow p = 0.21$
(AGN coverage of the sky)

H_1 prediction (anisotropy)

$p \geq 0.21$

$N = 13$ (first 14 events were used to define test)
 $k = 8$