# Statistical Methods for Particle Physics Lecture 1: intro, parameter estimation, tests http://indico.ihep.ac.cn/event/4902/



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#### Outline

Lecture 1: Introduction and review of fundamentals Probability, random variables, pdfs Parameter estimation, maximum likelihood Statistical tests for discovery and limits

Lecture 2: Multivariate methods

Neyman-Pearson lemma Fisher discriminant, neural networks Boosted decision trees

Lecture 3: Systematic uncertainties and further topics Nuisance parameters (Bayesian and frequentist) Experimental sensitivity The look-elsewhere effect

#### Some statistics books, papers, etc.

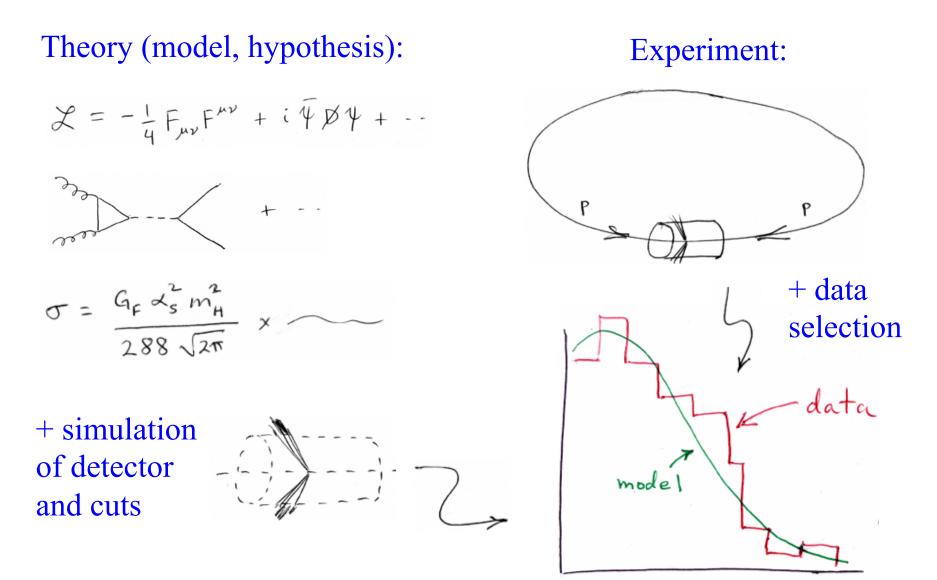
- G. Cowan, *Statistical Data Analysis*, Clarendon, Oxford, 1998 R.J. Barlow, *Statistics: A Guide to the Use of Statistical Methods in the Physical Sciences*, Wiley, 1989
- Ilya Narsky and Frank C. Porter, *Statistical Analysis Techniques in Particle Physics*, Wiley, 2014.
- L. Lyons, Statistics for Nuclear and Particle Physics, CUP, 1986
- F. James., *Statistical and Computational Methods in Experimental Physics*, 2nd ed., World Scientific, 2006
- S. Brandt, *Statistical and Computational Methods in Data Analysis*, Springer, New York, 1998 (with program library on CD) K.A. Olive et al. (Particle Data Group), *Review of Particle Physics*, Chin. Phys. C, 38, 090001 (2014).; see also pdg.lbl.gov sections on probability, statistics, Monte Carlo

#### More statistics books (中文)

朱永生, 实验物理中的概率和统计(第二版), 科学出版社, 北京, 2006。

朱永生(编著),实验数据多元统计分析,科学出版社, 北京,2009。

#### Theory ↔ Statistics ↔ Experiment



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# Data analysis in particle physics

Observe events (e.g., pp collisions) and for each, measure a set of characteristics:

particle momenta, number of muons, energy of jets,... Compare observed distributions of these characteristics to predictions of theory. From this, we want to:

Estimate the free parameters of the theory:  $m_{\mu} = 125.4$ 

Quantify the uncertainty in the estimates:  $\pm 0.4$  GeV

Assess how well a given theory stands in agreement with the observed data:  $O^+$  good,  $2^+$  bad

To do this we need a clear definition of PROBABILITY

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# A definition of probability

Consider a set S with subsets A, B, ...

For all  $A \subset S, P(A) \ge 0$  P(S) = 1If  $A \cap B = \emptyset, P(A \cup B) = P(A) + P(B)$ 



Kolmogorov axioms (1933)

Also define conditional probability of *A* given *B*:

$$P(A|B) = \frac{P(A \cap B)}{P(B)}$$

Subsets A, B independent if:  $P(A \cap B) = P(A)P(B)$ 

If A, B independent, 
$$P(A|B) = \frac{P(A)P(B)}{P(B)} = P(A)$$

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# Interpretation of probability

I. Relative frequency

A, B, ... are outcomes of a repeatable experiment

 $P(A) = \lim_{n \to \infty} \frac{\text{times outcome is } A}{n}$ 

cf. quantum mechanics, particle scattering, radioactive decay...

- II. Subjective probability

  A, B, ... are hypotheses (statements that are true or false)
  P(A) = degree of belief that A is true

  Both interpretations consistent with Kolmogorov axioms.
- In particle physics frequency interpretation often most useful, but subjective probability can provide more natural treatment of non-repeatable phenomena:

systematic uncertainties, probability that Higgs boson exists,...

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# Bayes' theorem

From the definition of conditional probability we have,

$$P(A|B) = \frac{P(A \cap B)}{P(B)}$$
 and  $P(B|A) = \frac{P(B \cap A)}{P(A)}$ 

but  $P(A \cap B) = P(B \cap A)$ , so

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

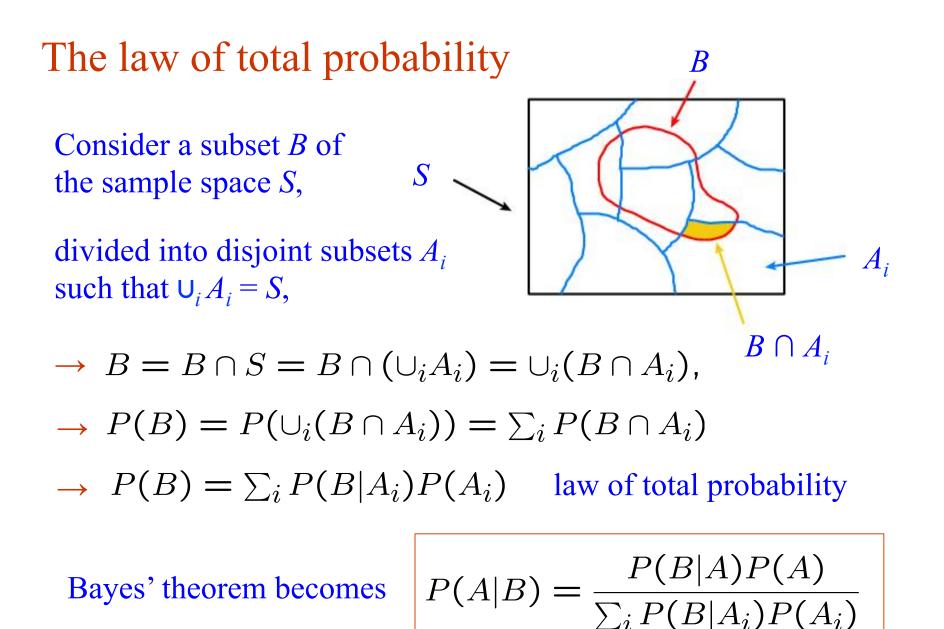
First published (posthumously) by the Reverend Thomas Bayes (1702–1761)

An essay towards solving a problem in the doctrine of chances, Philos. Trans. R. Soc. 53 (1763) 370; reprinted in Biometrika, 45 (1958) 293.

Bayes' theorem



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# An example using Bayes' theorem

Suppose the probability (for anyone) to have a disease D is:

 $P(D) = 0.001 \leftarrow \text{prior probabilities, i.e.,}$  $P(\text{no } D) = 0.999 \leftarrow \text{before any test carried out}$ 

Consider a test for the disease: result is + or -

- P(+|D) = 0.98 P(-|D) = 0.02  $\leftarrow$  probabilities to (in)correctly identify a person with the disease
- P(+|no D) = 0.03 P(-|no D) = 0.97 P(-|no D) = 0.97 P(-|no D) = 0.97

Suppose your result is +. How worried should you be?

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Bayes' theorem example (cont.)

The probability to have the disease given a + result is

$$p(\mathbf{D}|+) = \frac{P(+|\mathbf{D})P(\mathbf{D})}{P(+|\mathbf{D})P(\mathbf{D}) + P(+|\mathrm{no} \ \mathbf{D})P(\mathrm{no} \ \mathbf{D})}$$

# $= \frac{0.98 \times 0.001}{0.98 \times 0.001 + 0.03 \times 0.999}$

 $= 0.032 \leftarrow \text{posterior probability}$ 

#### i.e. you're probably OK!

Your viewpoint: my degree of belief that I have the disease is 3.2%. Your doctor's viewpoint: 3.2% of people like this have the disease.

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# Frequentist Statistics – general philosophy

In frequentist statistics, probabilities are associated only with the data, i.e., outcomes of repeatable observations (shorthand:  $\vec{x}$ ).

Probability = limiting frequency

Probabilities such as

*P* (Higgs boson exists), *P* (0.117 <  $\alpha_{\rm s}$  < 0.121),

etc. are either 0 or 1, but we don't know which.

The tools of frequentist statistics tell us what to expect, under the assumption of certain probabilities, about hypothetical repeated observations.

A hypothesis is is preferred if the data are found in a region of high predicted probability (i.e., where an alternative hypothesis predicts lower probability).

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# Bayesian Statistics – general philosophy

In Bayesian statistics, use subjective probability for hypotheses:

probability of the data assuming hypothesis *H* (the likelihood)  $P(H|\vec{x}) = \frac{P(\vec{x}|H)\pi(H)}{\int P(\vec{x}|H)\pi(H) dH}$ posterior probability, i.e., after seeing the data  $P(H|\vec{x}) = \frac{P(\vec{x}|H)\pi(H)}{\int P(\vec{x}|H)\pi(H) dH}$ normalization involves sum over all possible hypotheses

Bayes' theorem has an "if-then" character: If your prior probabilities were  $\pi(H)$ , then it says how these probabilities should change in the light of the data.

No general prescription for priors (subjective!)

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Random variables and probability density functions A random variable is a numerical characteristic assigned to an element of the sample space; can be discrete or continuous.

Suppose outcome of experiment is continuous value *x* 

$$P(x \text{ found in } [x, x + dx]) = f(x) dx$$

 $\rightarrow f(x) =$  probability density function (pdf)

$$\int_{-\infty}^{\infty} f(x) \, dx = 1 \qquad x \text{ must be somewhere}$$

Or for discrete outcome  $x_i$  with e.g. i = 1, 2, ... we have

$$P(x_i) = p_i$$
probability mass function $\sum_i P(x_i) = 1$ x must take on one of its possible values

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# Other types of probability densities

Outcome of experiment characterized by several values, e.g. an *n*-component vector,  $(x_1, ..., x_n)$ 

$$\rightarrow$$
 joint pdf  $f(x_1, \ldots, x_n)$ 

Sometimes we want only pdf of some (or one) of the components  $\rightarrow$  marginal pdf  $f_1(x_1) = \int \cdots \int f(x_1, \dots, x_n) dx_2 \dots dx_n$  $x_1, x_2$  independent if  $f(x_1, x_2) = f_1(x_1) f_2(x_2)$ 

Sometimes we want to consider some components as constant  $f(x_1, x_2)$ 

$$\rightarrow$$
 conditional pdf  $g(x_1|x_2) = \frac{f(x_1, x_2)}{f_2(x_2)}$ 

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# Expectation values

Consider continuous r.v. x with pdf f(x). Define expectation (mean) value as  $E[x] = \int x f(x) dx$ Notation (often):  $E[x] = \mu$  ~ "centre of gravity" of pdf. For a function y(x) with pdf g(y),

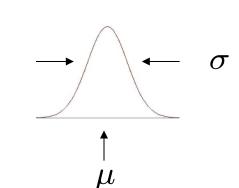
$$E[y] = \int y g(y) dy = \int y(x) f(x) dx$$
 (equivalent)

Variance:  $V[x] = E[x^2] - \mu^2 = E[(x - \mu)^2]$ 

Notation:  $V[x] = \sigma^2$ 

Standard deviation:  $\sigma = \sqrt{\sigma^2}$ 

 $\sigma$  ~ width of pdf, same units as *x*.



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#### Covariance and correlation

Define covariance cov[x,y] (also use matrix notation  $V_{xy}$ ) as

$$cov[x, y] = E[xy] - \mu_x \mu_y = E[(x - \mu_x)(y - \mu_y)]$$

Correlation coefficient (dimensionless) defined as

$$\rho_{xy} = \frac{\operatorname{cov}[x, y]}{\sigma_x \sigma_y}$$

If x, y, independent, i.e.,  $f(x, y) = f_x(x)f_y(y)$ , then

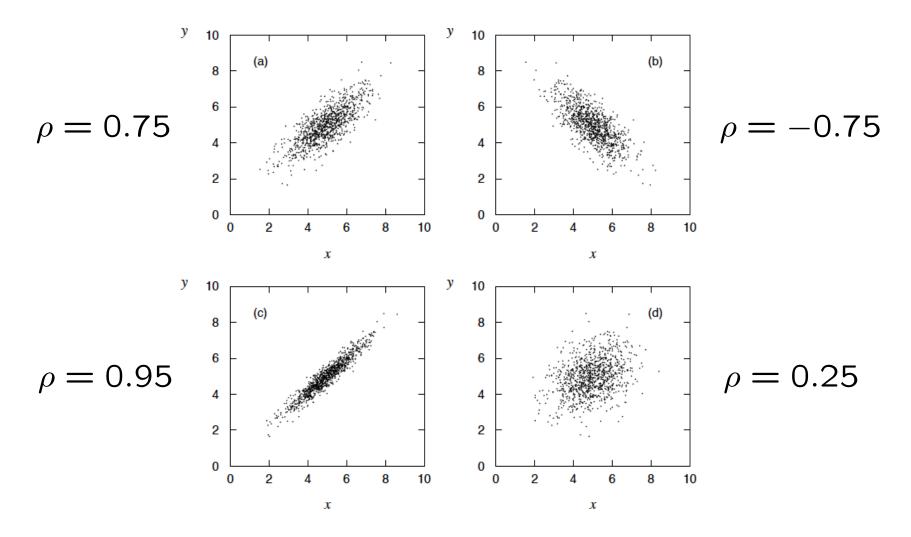
$$E[xy] = \int \int xy f(x, y) \, dx \, dy = \mu_x \mu_y$$
  

$$\rightarrow \operatorname{COV}[x, y] = 0 \qquad x \text{ and } y, \text{ `uncorrelated'}$$

N.B. converse not always true.

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# Correlation (cont.)



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## Review of frequentist parameter estimation

Suppose we have a pdf characterized by one or more parameters:

$$f(x;\theta) = \frac{1}{\theta}e^{-x/\theta}$$

random variable

parameter

Suppose we have a sample of observed values:  $\vec{x} = (x_1, \ldots, x_n)$ 

We want to find some function of the data to estimate the parameter(s):

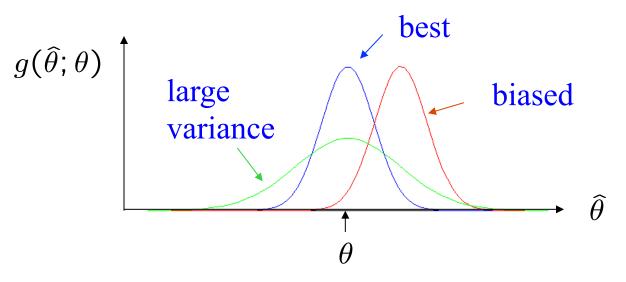
 $\hat{\theta}(\vec{x}) \leftarrow \text{estimator written with a hat}$ 

Sometimes we say 'estimator' for the function of  $x_1, ..., x_n$ ; 'estimate' for the value of the estimator with a particular data set.

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# Properties of estimators

If we were to repeat the entire measurement, the estimates from each would follow a pdf:



We want small (or zero) bias (systematic error):  $b = E[\hat{\theta}] - \theta$ 

→ average of repeated measurements should tend to true value.
 And we want a small variance (statistical error): V[θ̂]
 → small bias & variance are in general conflicting criteria

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# Distribution, likelihood, model

Suppose the outcome of a measurement is *x*. (e.g., a number of events, a histogram, or some larger set of numbers).

The probability density (or mass) function or 'distribution' of x, which may depend on parameters  $\theta$ , is:

 $P(x|\theta)$  (Independent variable is x;  $\theta$  is a constant.)

If we evaluate  $P(x|\theta)$  with the observed data and regard it as a function of the parameter(s), then this is the likelihood:

 $L(\theta) = P(x|\theta)$  (Data x fixed; treat L as function of  $\theta$ .)

We will use the term 'model' to refer to the full function  $P(x|\theta)$  that contains the dependence both on *x* and  $\theta$ .

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#### Bayesian use of the term 'likelihood'

We can write Bayes theorem as

$$p(\theta|x) = \frac{L(x|\theta)\pi(\theta)}{\int L(x|\theta)\pi(\theta) \, d\theta}$$

where  $L(x|\theta)$  is the likelihood. It is the probability for x given  $\theta$ , evaluated with the observed x, and viewed as a function of  $\theta$ .

Bayes' theorem only needs  $L(x|\theta)$  evaluated with a given data set (the 'likelihood principle').

For frequentist methods, in general one needs the full model. For some approximate frequentist methods, the likelihood is enough.

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#### The likelihood function for i.i.d.\*. data

\* i.i.d. = independent and identically distributed

Consider *n* independent observations of *x*:  $x_1, ..., x_n$ , where *x* follows  $f(x; \theta)$ . The joint pdf for the whole data sample is:

$$f(x_1,\ldots,x_n;\theta) = \prod_{i=1}^n f(x_i;\theta)$$

In this case the likelihood function is

$$L(\vec{\theta}) = \prod_{i=1}^{n} f(x_i; \vec{\theta}) \qquad (x_i \text{ constant})$$

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# Maximum likelihood

The most important frequentist method for constructing estimators is to take the value of the parameter(s) that maximize the likelihood:  $\hat{\theta} = \operatorname{argmax} L(x|\theta)$ 

The resulting estimators are functions of the data and thus characterized by a sampling distribution with a given (co)variance:

In general they may have a nonzero bias:

Under conditions usually satisfied in practice, bias of ML estimators is zero in the large sample limit, and the variance is as small as possible for unbiased estimators.

ML estimator may not in some cases be regarded as the optimal trade-off between these criteria (cf. regularized unfolding).

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$$V_{ij} = \operatorname{cov}[\hat{\theta}_i, \hat{\theta}_j]$$

$$b = E[\hat{\theta}] - \theta$$

ML example: parameter of exponential pdf

Consider exponential pdf, 
$$f(t; \tau) = \frac{1}{\tau}e^{-t/\tau}$$

and suppose we have i.i.d. data,  $t_1, \ldots, t_n$ 

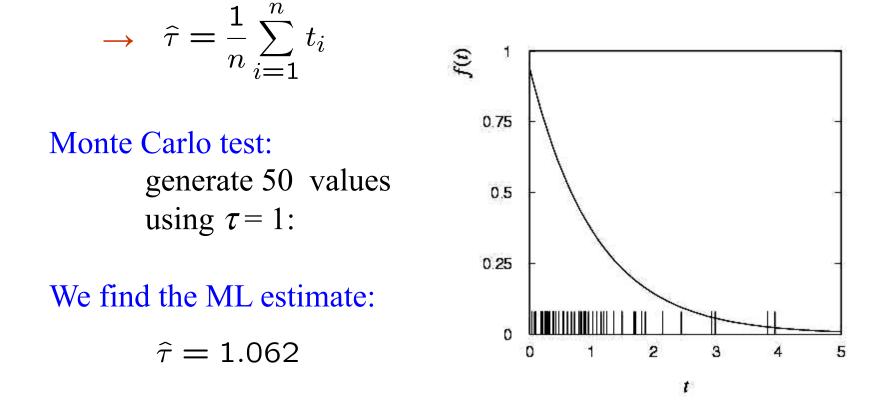
The likelihood function is 
$$L(\tau) = \prod_{i=1}^{n} \frac{1}{\tau} e^{-t_i/\tau}$$

The value of  $\tau$  for which  $L(\tau)$  is maximum also gives the maximum value of its logarithm (the log-likelihood function):

$$\ln L(\tau) = \sum_{i=1}^{n} \ln f(t_i; \tau) = \sum_{i=1}^{n} \left( \ln \frac{1}{\tau} - \frac{t_i}{\tau} \right)$$

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# ML example: parameter of exponential pdf (2) Find its maximum by setting $\frac{\partial \ln L(\tau)}{\partial \tau} = 0$ ,



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ML example: parameter of exponential pdf (3) For the exponential distribution one has for mean, variance:

$$E[t] = \int_0^\infty t \, \frac{1}{\tau} e^{-t/\tau} \, dt = \tau$$

$$V[t] = \int_0^\infty (t - \tau)^2 \frac{1}{\tau} e^{-t/\tau} dt = \tau^2$$

For the ML estimator  $\hat{\tau} = \frac{1}{n} \sum_{i=1}^{n} t_i$  we therefore find

$$E[\hat{\tau}] = E\left[\frac{1}{n}\sum_{i=1}^{n}t_i\right] = \frac{1}{n}\sum_{i=1}^{n}E[t_i] = \tau \quad \longrightarrow \quad b = E[\hat{\tau}] - \tau = 0$$

$$V[\hat{\tau}] = V\left[\frac{1}{n}\sum_{i=1}^{n} t_i\right] = \frac{1}{n^2}\sum_{i=1}^{n} V[t_i] = \frac{\tau^2}{n} \longrightarrow \quad \sigma_{\hat{\tau}} = \frac{\tau}{\sqrt{n}}$$

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# Variance of estimators: Monte Carlo method

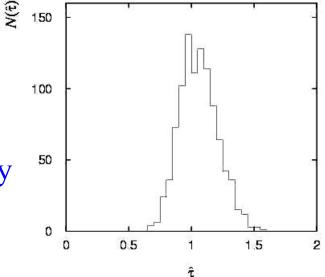
Having estimated our parameter we now need to report its 'statistical error', i.e., how widely distributed would estimates be if we were to repeat the entire measurement many times.

One way to do this would be to simulate the entire experiment many times with a Monte Carlo program (use ML estimate for MC).

For exponential example, from sample variance of estimates we find:  $\hat{a} = 0.151$ 

 $\hat{\sigma}_{\hat{\tau}} = 0.151$ 

Note distribution of estimates is roughly Gaussian – (almost) always true for ML in large sample limit.



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Variance of estimators from information inequality

The information inequality (RCF) sets a lower bound on the variance of any estimator (not only ML):

$$V[\hat{\theta}] \ge \left(1 + \frac{\partial b}{\partial \theta}\right)^2 / E\left[-\frac{\partial^2 \ln L}{\partial \theta^2}\right] \qquad \text{Bound (MVB)} \\ (b = E[\hat{\theta}] - \theta)$$

Often the bias b is small, and equality either holds exactly or is a good approximation (e.g. large data sample limit). Then,

$$V[\hat{\theta}] \approx -1 \left/ E\left[\frac{\partial^2 \ln L}{\partial \theta^2}\right] \right.$$

Estimate this using the 2nd derivative of  $\ln L$  at its maximum:

$$\widehat{V}[\widehat{\theta}] = -\left(\frac{\partial^2 \ln L}{\partial \theta^2}\right)^{-1} \bigg|_{\theta = \widehat{\theta}}$$

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Variance of estimators: graphical method Expand  $\ln L(\theta)$  about its maximum:

$$\ln L(\theta) = \ln L(\hat{\theta}) + \left[\frac{\partial \ln L}{\partial \theta}\right]_{\theta = \hat{\theta}} (\theta - \hat{\theta}) + \frac{1}{2!} \left[\frac{\partial^2 \ln L}{\partial \theta^2}\right]_{\theta = \hat{\theta}} (\theta - \hat{\theta})^2 + \dots$$

First term is  $\ln L_{max}$ , second term is zero, for third term use information inequality (assume equality):

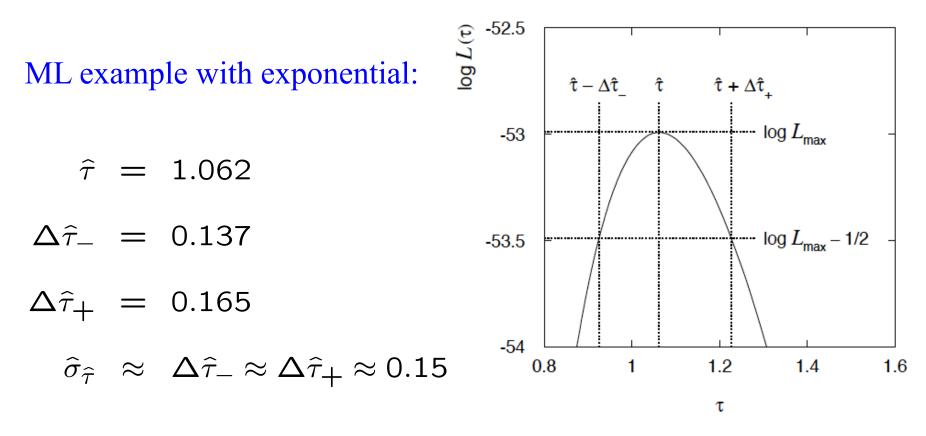
$$\ln L(\theta) \approx \ln L_{\max} - \frac{(\theta - \widehat{\theta})^2}{2\widehat{\sigma^2}_{\widehat{\theta}}}$$

i.e., 
$$\ln L(\hat{\theta} \pm \hat{\sigma}_{\hat{\theta}}) \approx \ln L_{\max} - \frac{1}{2}$$

 $\rightarrow$  to get  $\hat{\sigma}_{\hat{\theta}}$ , change  $\theta$  away from  $\hat{\theta}$  until ln *L* decreases by 1/2.

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## Example of variance by graphical method



Not quite parabolic  $\ln L$  since finite sample size (n = 50).

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Information inequality for *n* parameters Suppose we have estimated *n* parameters  $\vec{\theta} = (\theta_1, \dots, \theta_n)$ . The (inverse) minimum variance bound is given by the

Fisher information matrix:

$$I_{ij} = E\left[-\frac{\partial^2 \ln L}{\partial \theta_i \partial \theta_j}\right] = -n \int f(x; \vec{\theta}) \frac{\partial^2 \ln f(x; \vec{\theta})}{\partial \theta_i \partial \theta_j} dx$$

The information inequality then states that  $V - I^{-1}$  is a positive semi-definite matrix, where  $V_{ij} = \text{cov}[\hat{\theta}_i, \hat{\theta}_j]$ . Therefore

$$V[\hat{\theta}_i] \ge (I^{-1})_{ii}$$

Often use  $I^{-1}$  as an approximation for covariance matrix, estimate using e.g. matrix of 2nd derivatives at maximum of L.

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#### Two-parameter example of ML

Consider a scattering angle distribution with  $x = \cos \theta$ ,

$$f(x; \alpha, \beta) = \frac{1 + \alpha x + \beta x^2}{2 + 2\beta/3}$$

Data:  $x_1, ..., x_n, n = 2000$  events.

As test generate with MC using  $\alpha = 0.5$ ,  $\beta = 0.5$ 

From data compute log-likelihood:

$$\ln L(\alpha,\beta) = \sum_{i=1}^{n} \ln f(x_i;\alpha,\beta)$$

Maximize numerically (e.g., program MINUIT)

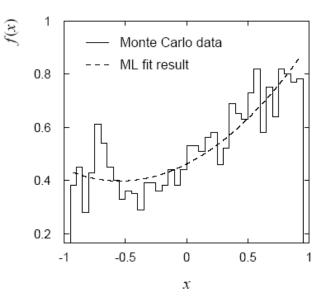
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#### Example of ML: fit result Finding maximum of $\ln L(\alpha, \beta)$ numerically (MINUIT) gives

$$\hat{\alpha} = 0.508$$

$$\hat{\beta} = 0.47$$

**N.B.** Here no binning of data for fit, but can compare to histogram for goodness-of-fit (e.g. 'visual' or  $\chi^2$ ).



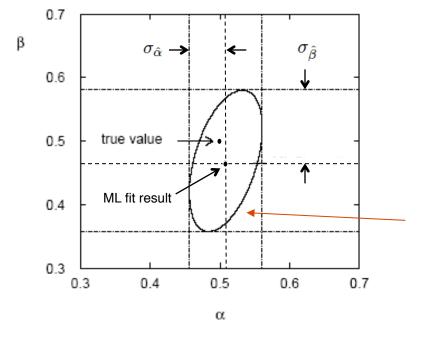
(Co)variances from 
$$(\widehat{V^{-1}})_{ij} = -\frac{\partial^2 \ln L}{\partial \theta_i \partial \theta_j}\Big|_{\vec{\theta} = \hat{\vec{\theta}}}$$

(MINUIT routine HESSE)

 $\hat{\sigma}_{\hat{\alpha}} = 0.052 \quad \operatorname{cov}[\hat{\alpha}, \hat{\beta}] = 0.0026$  $\hat{\sigma}_{\hat{\alpha}} = 0.11 \quad r = 0.46$ 

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Variance of ML estimators: graphical method Often (e.g., large sample case) one can approximate the covariances using only  $\hat{V}_{ij}^{-1} \approx -\frac{\partial^2 \ln L}{\partial \theta_i \partial \theta_j}\Big|_{\theta=0}$ the likelihood  $L(\theta)$ :



This translates into a simple graphical recipe:

$$n L(\alpha, \beta) = ln L_{max} - 1/2$$

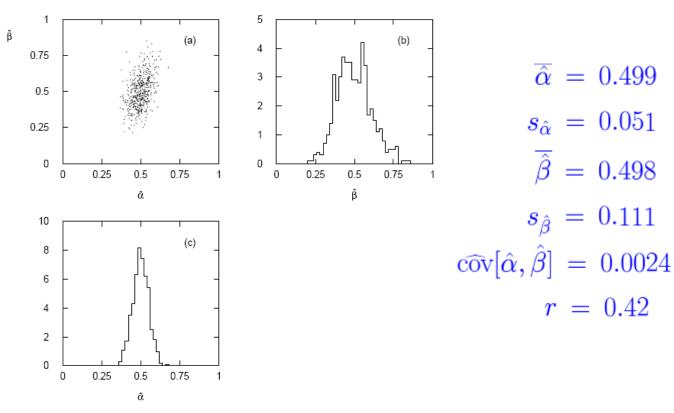
 $\rightarrow$  Tangent lines to contours give standard deviations.

 $\rightarrow$  Angle of ellipse  $\phi$  related to correlation:  $\tan 2\phi = \frac{2\rho\sigma_{\hat{\alpha}}\sigma_{\hat{\beta}}}{\sigma^2 - \sigma^{\hat{\beta}^2}}$ 

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#### Variance of ML estimators: MC

To find the ML estimate itself one only needs the likelihood  $L(\theta)$ . In principle to find the covariance of the estimators, one requires the full model  $P(x|\theta)$ . E.g., simulate many times independent data sets and look at distribution of the resulting estimates:



#### Frequentist statistical tests

Consider a hypothesis  $H_0$  and alternative  $H_1$ .

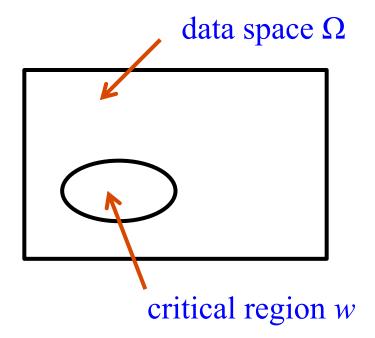
A test of  $H_0$  is defined by specifying a critical region *w* of the data space such that there is no more than some (small) probability  $\alpha$ , assuming  $H_0$  is correct, to observe the data there, i.e.,

$$P(x \in w \mid H_0) \le \alpha$$

Need inequality if data are discrete.

 $\alpha$  is called the size or significance level of the test.

If x is observed in the critical region, reject  $H_0$ .

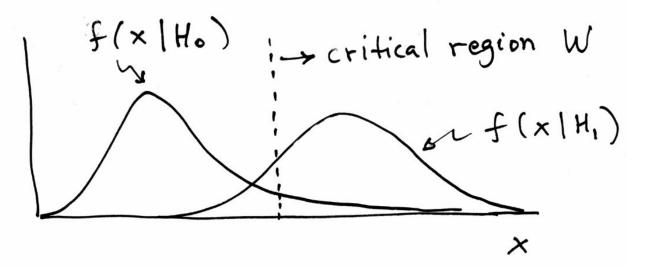


### Definition of a test (2)

But in general there are an infinite number of possible critical regions that give the same significance level  $\alpha$ .

So the choice of the critical region for a test of  $H_0$  needs to take into account the alternative hypothesis  $H_1$ .

Roughly speaking, place the critical region where there is a low probability to be found if  $H_0$  is true, but high if  $H_1$  is true:



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#### Type-I, Type-II errors

Rejecting the hypothesis  $H_0$  when it is true is a Type-I error. The maximum probability for this is the size of the test:

$$P(x \in W \mid H_0) \le \alpha$$

But we might also accept  $H_0$  when it is false, and an alternative  $H_1$  is true.

This is called a Type-II error, and occurs with probability

$$P(x \in \mathbf{S} - W | H_1) = \beta$$

One minus this is called the power of the test with respect to the alternative  $H_1$ :

Power = 
$$1 - \beta$$

#### *p*-values

Suppose hypothesis *H* predicts pdf  $f(\vec{x}|H)$  for a set of observations  $\vec{x} = (x_1, \dots, x_n)$ .

We observe a single point in this space:  $\vec{x}_{ODS}$ 

What can we say about the validity of *H* in light of the data?

Express level of compatibility by giving the *p*-value for *H*:

p = probability, under assumption of H, to observe data with equal or lesser compatibility with H relative to the data we got.

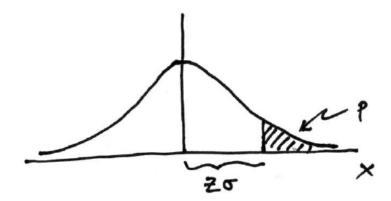


This is not the probability that *H* is true!

Requires one to say what part of data space constitutes lesser compatibility with *H* than the observed data (implicitly this means that region gives better agreement with some alternative).

#### Significance from *p*-value

Often define significance Z as the number of standard deviations that a Gaussian variable would fluctuate in one direction to give the same p-value.



$$p=\int_Z^\infty rac{1}{\sqrt{2\pi}}e^{-x^2/2}\,dx=1-\Phi(Z)$$
 1 - TMath::Freq

 $Z = \Phi^{-1}(1-p)$  TMath::NormQuantile

E.g. Z = 5 (a "5 sigma effect") corresponds to  $p = 2.9 \times 10^{-7}$ .

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#### Using a *p*-value to define test of $H_0$

One can show the distribution of the *p*-value of H, under assumption of H, is uniform in [0,1].

So the probability to find the *p*-value of  $H_0$ ,  $p_0$ , less than  $\alpha$  is

$$P(p_0 \le \alpha | H_0) = \alpha$$

We can define the critical region of a test of  $H_0$  with size *a* as the set of data space where  $p_0 \le \alpha$ .

Formally the *p*-value relates only to  $H_0$ , but the resulting test will have a given power with respect to a given alternative  $H_1$ .

The Poisson counting experiment

Suppose we do a counting experiment and observe *n* events.

Events could be from *signal* process or from *background* – we only count the total number.

Poisson model:

$$P(n|s,b) = \frac{(s+b)^n}{n!}e^{-(s+b)}$$

s = mean (i.e., expected) # of signal events

b = mean # of background events

Goal is to make inference about *s*, e.g.,

test s = 0 (rejecting  $H_0 \approx$  "discovery of signal process")

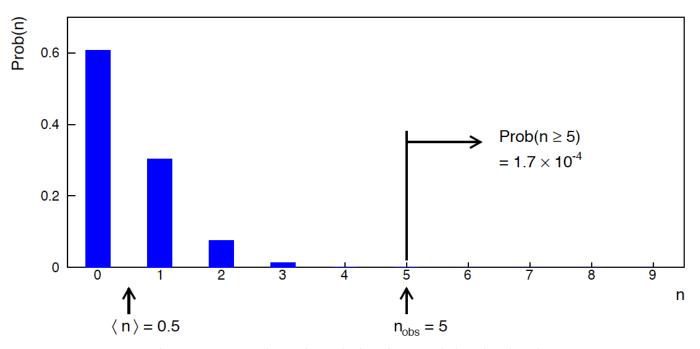
test all non-zero *s* (values not rejected = confidence interval)

In both cases need to ask what is relevant alternative hypothesis.G. CowaniSTEP 2015, Jinan / Statistics for Particle Physics / Lecture 1

Poisson counting experiment: discovery *p*-value Suppose b = 0.5 (known), and we observe  $n_{obs} = 5$ . Should we claim evidence for a new discovery?

Give *p*-value for hypothesis *s* = 0:

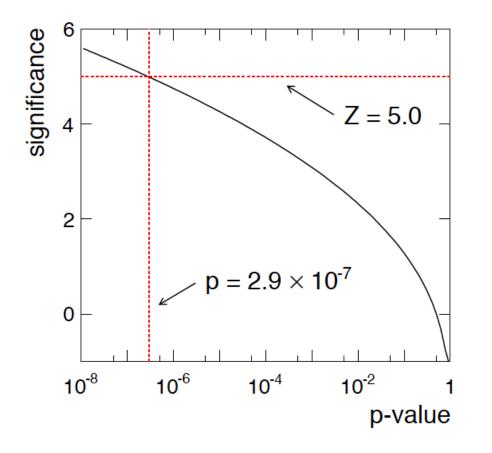
$$p$$
-value =  $P(n \ge 5; b = 0.5, s = 0)$   
=  $1.7 \times 10^{-4} \ne P(s = 0)!$ 



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# Poisson counting experiment: discovery significance Equivalent significance for $p = 1.7 \times 10^{-4}$ : $Z = \Phi^{-1}(1-p) = 3.6$ Often claim discovery if Z > 5 ( $p < 2.9 \times 10^{-7}$ , i.e., a "5-sigma effect")



In fact this tradition should be revisited: *p*-value intended to quantify probability of a signallike fluctuation assuming background only; not intended to cover, e.g., hidden systematics, plausibility signal model, compatibility of data with signal, "look-elsewhere effect" (~multiple testing), etc.

Confidence intervals by inverting a test Confidence intervals for a parameter  $\theta$  can be found by defining a test of the hypothesized value  $\theta$  (do this for all  $\theta$ ):

Specify values of the data that are 'disfavoured' by  $\theta$  (critical region) such that  $P(\text{data in critical region}) \le \alpha$  for a prespecified  $\alpha$ , e.g., 0.05 or 0.1.

If data observed in the critical region, reject the value  $\theta$ .

Now invert the test to define a confidence interval as:

set of  $\theta$  values that would not be rejected in a test of size  $\alpha$  (confidence level is  $1 - \alpha$ ).

The interval will cover the true value of  $\theta$  with probability  $\geq 1 - \alpha$ .

Equivalently, the parameter values in the confidence interval have p-values of at least  $\alpha$ .

47

To find edge of interval (the "limit"), set  $p_{\theta} = \alpha$  and solve for  $\theta$ . G. Cowan iSTEP 2015, Jinan / Statistics for Particle Physics / Lecture 1

#### Frequentist upper limit on Poisson parameter

Consider again the case of observing  $n \sim \text{Poisson}(s + b)$ . Suppose b = 4.5,  $n_{\text{obs}} = 5$ . Find upper limit on *s* at 95% CL. Relevant alternative is s = 0 (critical region at low *n*) *p*-value of hypothesized *s* is P( $n \le n_{\text{obs}}$ ; *s*, *b*)

Upper limit  $s_{up}$  at  $CL = 1 - \alpha$  found from

$$\alpha = P(n \le n_{\text{obs}}; s_{\text{up}}, b) = \sum_{n=0}^{n_{\text{obs}}} \frac{(s_{\text{up}} + b)^n}{n!} e^{-(s_{\text{up}} + b)}$$

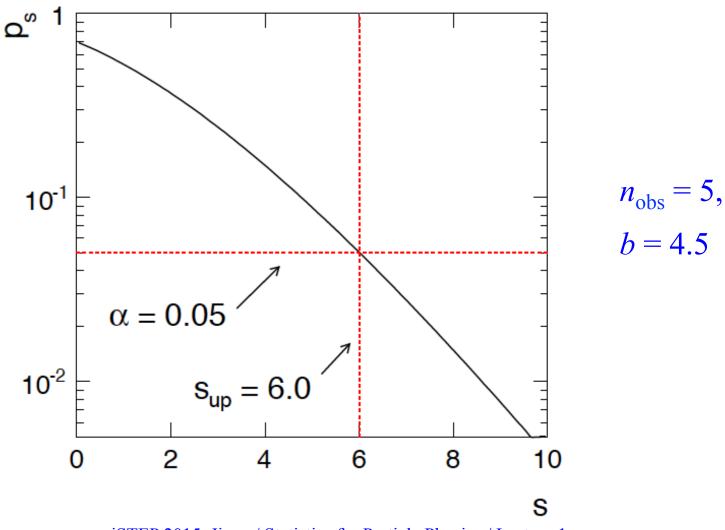
$$s_{\rm up} = \frac{1}{2} F_{\chi^2}^{-1} (1 - \alpha; 2(n_{\rm obs} + 1)) - b$$

$$=\frac{1}{2}F_{\chi^2}^{-1}(0.95;2(5+1)) - 4.5 = 6.0$$

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#### Frequentist upper limit on Poisson parameter

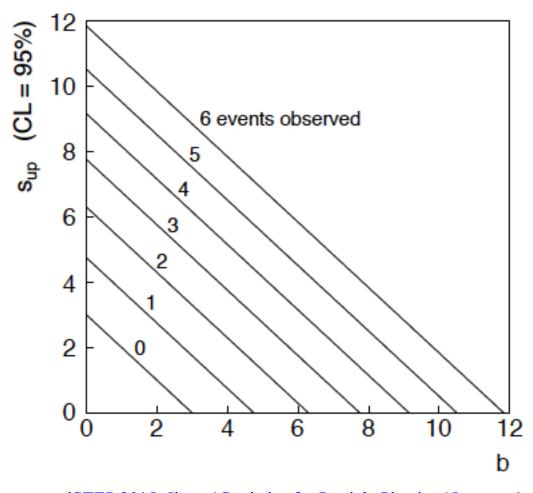
Upper limit  $s_{up}$  at  $CL = 1 - \alpha$  found from  $p_s = \alpha$ .



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# $n \sim \text{Poisson}(s+b)$ : frequentist upper limit on *s* For low fluctuation of *n* formula can give negative result for $s_{up}$ ; i.e. confidence interval is empty.



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Limits near a physical boundary

Suppose e.g. b = 2.5 and we observe n = 0.

If we choose CL = 0.9, we find from the formula for  $s_{up}$ 

 $s_{\rm up} = -0.197$  (CL = 0.90)

Physicist:

We already knew  $s \ge 0$  before we started; can't use negative upper limit to report result of expensive experiment!

Statistician:

The interval is designed to cover the true value only 90% of the time — this was clearly not one of those times.

Not uncommon dilemma when testing parameter values for which one has very little experimental sensitivity, e.g., very small *s*.

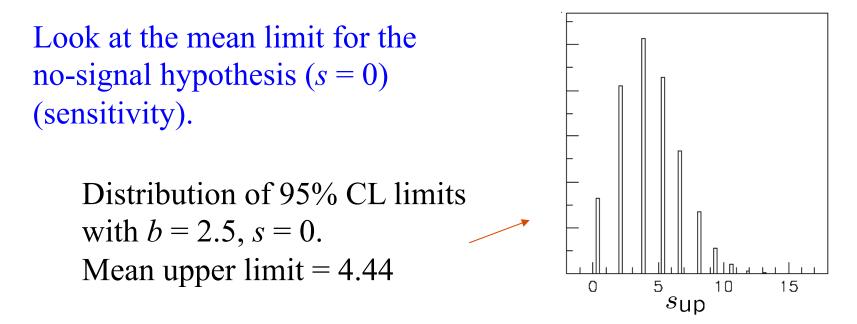
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#### Expected limit for s = 0

Physicist: I should have used CL = 0.95 — then  $s_{up} = 0.496$ 

Even better: for CL = 0.917923 we get  $s_{up} = 10^{-4}!$ 

Reality check: with b = 2.5, typical Poisson fluctuation in *n* is at least  $\sqrt{2.5} = 1.6$ . How can the limit be so low?



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#### The Bayesian approach to limits

In Bayesian statistics need to start with 'prior pdf'  $\pi(\theta)$ , this reflects degree of belief about  $\theta$  before doing the experiment.

Bayes' theorem tells how our beliefs should be updated in light of the data *x*:

$$p(\theta|x) = \frac{L(x|\theta)\pi(\theta)}{\int L(x|\theta')\pi(\theta') d\theta'} \propto L(x|\theta)\pi(\theta)$$

Integrate posterior pdf  $p(\theta | x)$  to give interval with any desired probability content.

For e.g.  $n \sim \text{Poisson}(s+b)$ , 95% CL upper limit on *s* from

$$0.95 = \int_{-\infty}^{s_{\rm up}} p(s|n) \, ds$$

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Bayesian prior for Poisson parameter

Include knowledge that  $s \ge 0$  by setting prior  $\pi(s) = 0$  for s < 0.

Could try to reflect 'prior ignorance' with e.g.

$$\pi(s) = \begin{cases} 1 & s \ge 0\\ 0 & \text{otherwise} \end{cases}$$

Not normalized but this is OK as long as L(s) dies off for large s.

Not invariant under change of parameter — if we had used instead a flat prior for, say, the mass of the Higgs boson, this would imply a non-flat prior for the expected number of Higgs events.

Doesn't really reflect a reasonable degree of belief, but often used as a point of reference;

or viewed as a recipe for producing an interval whose frequentist properties can be studied (coverage will depend on true *s*).

Bayesian interval with flat prior for s

Solve to find limit  $s_{up}$ :

$$s_{\rm up} = \frac{1}{2} F_{\chi^2}^{-1} [p, 2(n+1)] - b$$

where

$$p = 1 - \alpha \left( 1 - F_{\chi^2} \left[ 2b, 2(n+1) \right] \right)$$

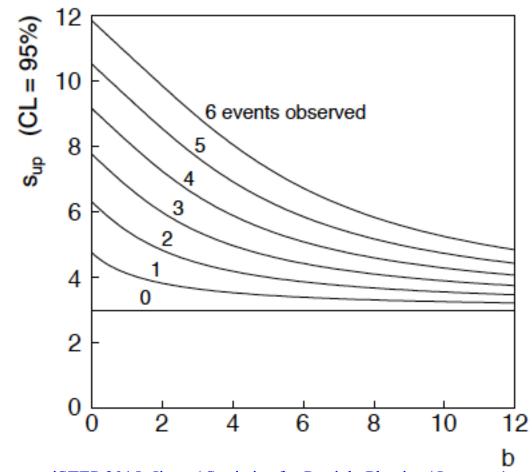
For special case b = 0, Bayesian upper limit with flat prior numerically same as one-sided frequentist case ('coincidence').

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Bayesian interval with flat prior for s

For b > 0 Bayesian limit is everywhere greater than the (one sided) frequentist upper limit.

Never goes negative. Doesn't depend on *b* if n = 0.



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#### Priors from formal rules

Because of difficulties in encoding a vague degree of belief in a prior, one often attempts to derive the prior from formal rules, e.g., to satisfy certain invariance principles or to provide maximum information gain for a certain set of measurements.

> Often called "objective priors" Form basis of Objective Bayesian Statistics

The priors do not reflect a degree of belief (but might represent possible extreme cases).

In Objective Bayesian analysis, can use the intervals in a frequentist way, i.e., regard Bayes' theorem as a recipe to produce an interval with certain coverage properties.

#### Priors from formal rules (cont.)

#### For a review of priors obtained by formal rules see, e.g.,

Robert E. Kass and Larry Wasserman, *The Selection of Prior Distributions by Formal Rules*, J. Am. Stat. Assoc., Vol. 91, No. 435, pp. 1343-1370 (1996).

Formal priors have not been widely used in HEP, but there is recent interest in this direction, especially the reference priors of Bernardo and Berger; see e.g.

L. Demortier, S. Jain and H. Prosper, *Reference priors for high energy physics*, Phys. Rev. D 82 (2010) 034002, arXiv:1002.1111.

D. Casadei, *Reference analysis of the signal + background model in counting experiments*, JINST 7 (2012) 01012; arXiv:1108.4270.

### Approximate confidence intervals/regions from the likelihood function

Suppose we test parameter value(s)  $\theta = (\theta_1, ..., \theta_n)$  using the ratio

$$\lambda(\theta) = \frac{L(\theta)}{L(\hat{\theta})} \qquad \qquad 0 \le \lambda(\theta) \le 1$$

Lower  $\lambda(\theta)$  means worse agreement between data and hypothesized  $\theta$ . Equivalently, usually define

$$t_{\theta} = -2\ln\lambda(\theta)$$

so higher  $t_{\theta}$  means worse agreement between  $\theta$  and the data.

*p*-value of 
$$\theta$$
 therefore  $p_{\theta} = \int_{t_{\theta,\text{obs}}}^{\infty} f(t_{\theta}|\theta) dt_{\theta}$   
need pdf

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Confidence region from Wilks' theorem Wilks' theorem says (in large-sample limit and providing certain conditions hold...)

 $f(t_{\theta}|\theta) \sim \chi_n^2 \qquad \text{chi-square dist. with $\#$ d.o.f. =} \\ \# \text{ of components in $\theta = (\theta_1, ..., \theta_n)$.}$ 

Assuming this holds, the *p*-value is

$$p_{\theta} = 1 - F_{\chi_n^2}(t_{\theta})$$

To find boundary of confidence region set  $p_{\theta} = \alpha$  and solve for  $t_{\theta}$ :

$$t_{\theta} = -2\ln\frac{L(\theta)}{L(\hat{\theta})} = F_{\chi_n^2}^{-1}(1-\alpha)$$

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Confidence region from Wilks' theorem (cont.) i.e., boundary of confidence region in  $\theta$  space is where

$$\ln L(\theta) = \ln L(\hat{\theta}) - \frac{1}{2}F_{\chi_n^2}^{-1}(1-\alpha)$$

For example, for  $1 - \alpha = 68.3\%$  and n = 1 parameter,

$$F_{\chi_1^2}^{-1}(0.683) = 1$$

and so the 68.3% confidence level interval is determined by

$$\ln L(\theta) = \ln L(\hat{\theta}) - \frac{1}{2}$$

Same as recipe for finding the estimator's standard deviation, i.e.,

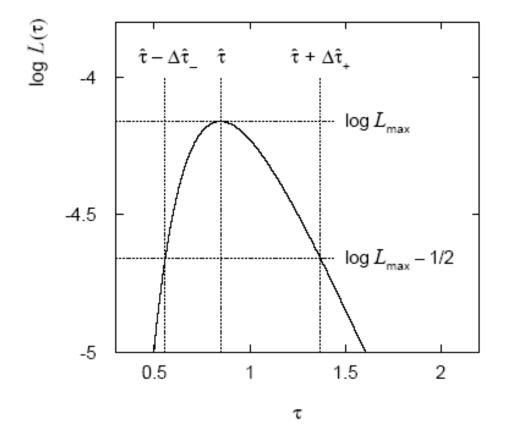
 $[\hat{\theta} - \sigma_{\hat{\theta}}, \hat{\theta} + \sigma_{\hat{\theta}}]$  is a 68.3% CL confidence interval.

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Example of interval from  $\ln L$ 

For n = 1 parameter, CL = 0.683,  $Q_{\alpha} = 1$ .

Exponential example, now with only 5 events:



Parameter estimate and approximate 68.3% CL confidence interval:

 $\hat{\tau} = 0.85^{+0.52}_{-0.30}$ 

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#### Multiparameter case

For increasing number of parameters,  $CL = 1 - \alpha$  decreases for confidence region determined by a given

$$Q_{\alpha} = F_{\chi_n^2}^{-1}(1-\alpha)$$

$Q_{lpha}$	1-lpha						
	n = 1	n = 2	n = 3	n = 4	n = 5		
1.0	0.683	0.393	0.199	0.090	0.037		
2.0	0.843	0.632	0.428	0.264	0.151		
4.0	0.954	0.865	0.739	0.594	0.451		
9.0	0.997	0.989	0.971	0.939	0.891		

#### Multiparameter case (cont.)

Equivalently,  $Q_{\alpha}$  increases with *n* for a given  $CL = 1 - \alpha$ .

$1 - \alpha$	$Q_{lpha}$						
	n = 1	n = 2	n = 3	n = 4	n = 5		
0.683	1.00	2.30	3.53	4.72	5.89		
0.90	2.71	4.61	6.25	7.78	9.24		
0.95	3.84	5.99	7.82	9.49	11.1		
0.99	6.63	9.21	11.3	13.3	15.1		

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#### Some distributions

Distribution/pdf **Binomial** Multinomial Poisson Uniform Exponential Gaussian Chi-square Cauchy Landau Beta Gamma Student's t

Example use in HEP **Branching** ratio Histogram with fixed NNumber of events found Monte Carlo method Decay time Measurement error Goodness-of-fit Mass of resonance **Ionization energy loss** Prior pdf for efficiency Sum of exponential variables Resolution function with adjustable tails

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#### **Binomial distribution**

Consider *N* independent experiments (Bernoulli trials): outcome of each is 'success' or 'failure', probability of success on any given trial is *p*.

Define discrete r.v. n = number of successes ( $0 \le n \le N$ ).

Probability of a specific outcome (in order), e.g. 'ssfsf' is  $pp(1-p)p(1-p) = p^n(1-p)^{N-n}$  N!

But order not important; there are

 $\frac{1}{n!(N-n)!}$ 

ways (permutations) to get *n* successes in *N* trials, total probability for *n* is sum of probabilities for each permutation.

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#### Binomial distribution (2)

The binomial distribution is therefore

$$f(n; N, p) = \frac{N!}{n!(N-n)!}p^n(1-p)^{N-n}$$
random parameters
variable

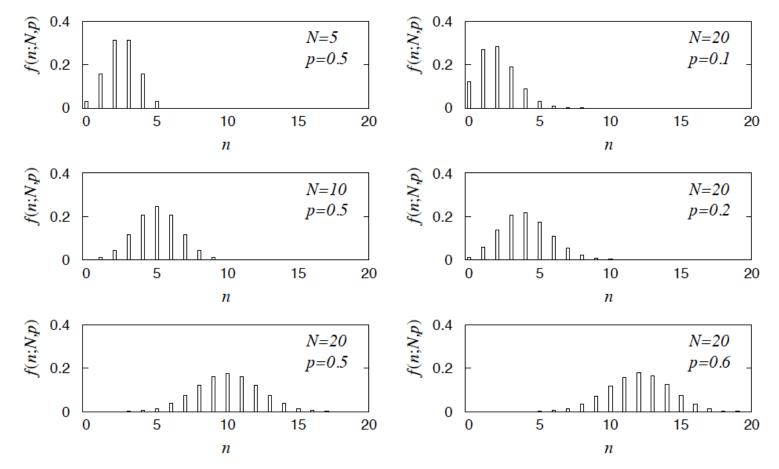
For the expectation value and variance we find:

$$E[n] = \sum_{n=0}^{N} nf(n; N, p) = Np$$
$$V[n] = E[n^{2}] - (E[n])^{2} = Np(1 - p)$$

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### Binomial distribution (3)

Binomial distribution for several values of the parameters:



Example: observe *N* decays of  $W^{\pm}$ , the number *n* of which are  $W \rightarrow \mu \nu$  is a binomial r.v., *p* = branching ratio.

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#### Multinomial distribution

Like binomial but now *m* outcomes instead of two, probabilities are

$$\vec{p} = (p_1, \dots, p_m)$$
, with  $\sum_{i=1}^m p_i = 1$ .

For N trials we want the probability to obtain:

 $n_1$  of outcome 1,  $n_2$  of outcome 2,  $\vdots$  $n_m$  of outcome *m*.

This is the multinomial distribution for  $\vec{n} = (n_1, \ldots, n_m)$ 

$$f(\vec{n}; N, \vec{p}) = \frac{N!}{n_1! n_2! \cdots n_m!} p_1^{n_1} p_2^{n_2} \cdots p_m^{n_m}$$

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#### Multinomial distribution (2)

Now consider outcome *i* as 'success', all others as 'failure'.

 $\rightarrow$  all  $n_i$  individually binomial with parameters  $N, p_i$ 

$$E[n_i] = Np_i, \quad V[n_i] = Np_i(1-p_i) \quad \text{for all } i$$

One can also find the covariance to be

$$V_{ij} = Np_i(\delta_{ij} - p_j)$$

Example:  $\vec{n} = (n_1, \dots, n_m)$  represents a histogram with *m* bins, *N* total entries, all entries independent.

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## Poisson distribution

Consider binomial *n* in the limit

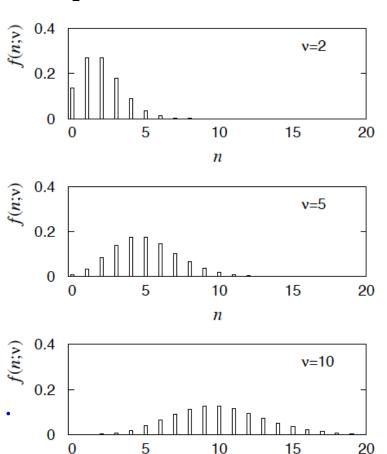
 $N \to \infty, \qquad p \to 0, \qquad E[n] = Np \to \nu.$ 

 $\rightarrow$  *n* follows the Poisson distribution:

$$f(n;\nu) = \frac{\nu^n}{n!}e^{-\nu} \quad (n \ge 0)$$

$$E[n] = \nu, \quad V[n] = \nu.$$

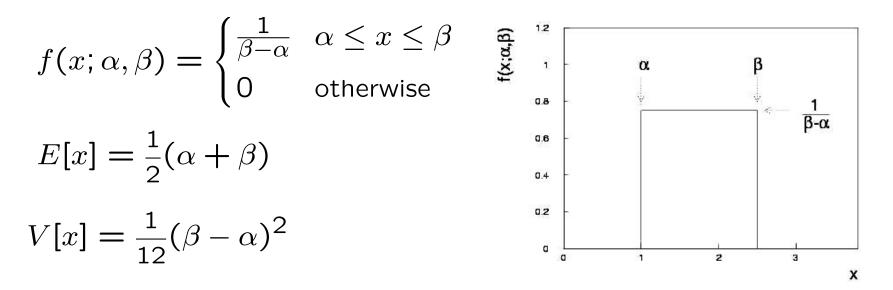
Example: number of scattering events *n* with cross section  $\sigma$  found for a fixed integrated luminosity, with  $\nu = \sigma \int L dt$ .



n

## Uniform distribution

Consider a continuous r.v. *x* with  $-\infty < x < \infty$ . Uniform pdf is:



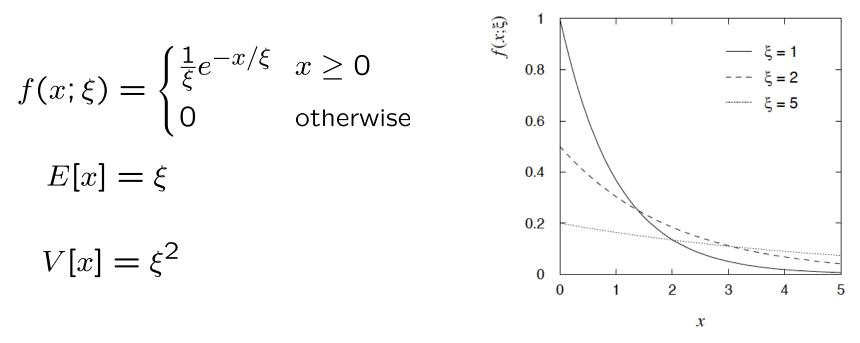
N.B. For any r.v. *x* with cumulative distribution F(x), y = F(x) is uniform in [0,1].

Example: for  $\pi^0 \to \gamma \gamma$ ,  $E_{\gamma}$  is uniform in  $[E_{\min}, E_{\max}]$ , with  $E_{\min} = \frac{1}{2} E_{\pi} (1 - \beta)$ ,  $E_{\max} = \frac{1}{2} E_{\pi} (1 + \beta)$ 

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## Exponential distribution

The exponential pdf for the continuous r.v. x is defined by:



Example: proper decay time *t* of an unstable particle

 $f(t;\tau) = \frac{1}{\tau}e^{-t/\tau}$  ( $\tau$  = mean lifetime)

Lack of memory (unique to exponential):  $f(t - t_0 | t \ge t_0) = f(t)$ 

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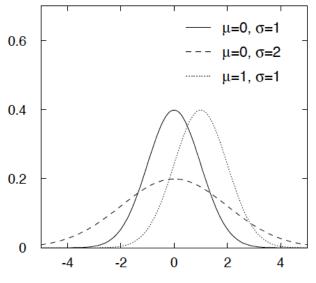
## Gaussian distribution

The Gaussian (normal) pdf for a continuous r.v. *x* is defined by:

$$f(x; \mu, \sigma) = \frac{1}{\sqrt{2\pi\sigma}} e^{-(x-\mu)^2/2\sigma^2}$$

$$E[x] = \mu$$
(N.B. often  $\mu, \sigma^2$  denote mean, variance of any  

$$V[x] = \sigma^2$$
r.v., not only Gaussian.)



х

Special case:  $\mu = 0$ ,  $\sigma^2 = 1$  ('standard Gaussian'):

$$\varphi(x) = \frac{1}{\sqrt{2\pi}} e^{-x^2/2} , \quad \Phi(x) = \int_{-\infty}^x \varphi(x') \, dx'$$

If  $y \sim$  Gaussian with  $\mu$ ,  $\sigma^2$ , then  $x = (y - \mu) / \sigma$  follows  $\varphi(x)$ .

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## Gaussian pdf and the Central Limit Theorem

The Gaussian pdf is so useful because almost any random variable that is a sum of a large number of small contributions follows it. This follows from the Central Limit Theorem:

For *n* independent r.v.s  $x_i$  with finite variances  $\sigma_i^2$ , otherwise arbitrary pdfs, consider the sum

$$y = \sum_{i=1}^{n} x_i$$

In the limit  $n \to \infty$ , y is a Gaussian r.v. with

$$E[y] = \sum_{i=1}^{n} \mu_i \qquad V[y] = \sum_{i=1}^{n} \sigma_i^2$$

Measurement errors are often the sum of many contributions, so frequently measured values can be treated as Gaussian r.v.s.

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## Central Limit Theorem (2)

The CLT can be proved using characteristic functions (Fourier transforms), see, e.g., SDA Chapter 10.

For finite *n*, the theorem is approximately valid to the extent that the fluctuation of the sum is not dominated by one (or few) terms.



Beware of measurement errors with non-Gaussian tails.

Good example: velocity component  $v_x$  of air molecules.

OK example: total deflection due to multiple Coulomb scattering. (Rare large angle deflections give non-Gaussian tail.)

Bad example: energy loss of charged particle traversing thin gas layer. (Rare collisions make up large fraction of energy loss, cf. Landau pdf.)

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## Multivariate Gaussian distribution

Multivariate Gaussian pdf for the vector  $\vec{x} = (x_1, \dots, x_n)$ :

$$f(\vec{x};\vec{\mu},V) = \frac{1}{(2\pi)^{n/2}|V|^{1/2}} \exp\left[-\frac{1}{2}(\vec{x}-\vec{\mu})^T V^{-1}(\vec{x}-\vec{\mu})\right]$$

 $\vec{x}, \vec{\mu}$  are column vectors,  $\vec{x}^T, \vec{\mu}^T$  are transpose (row) vectors,

$$E[x_i] = \mu_i, , \quad \operatorname{cov}[x_i, x_j] = V_{ij} .$$

For n = 2 this is  $f(x_1, x_2; \mu_1, \mu_2, \sigma_1, \sigma_2, \rho) = \frac{1}{2\pi\sigma_1\sigma_2\sqrt{1-\rho^2}}$   $\times \exp\left\{-\frac{1}{2(1-\rho^2)} \left[ \left(\frac{x_1 - \mu_1}{\sigma_1}\right)^2 + \left(\frac{x_2 - \mu_2}{\sigma_2}\right)^2 - 2\rho \left(\frac{x_1 - \mu_1}{\sigma_1}\right) \left(\frac{x_2 - \mu_2}{\sigma_2}\right) \right] \right\}$ 

where  $\rho = \operatorname{cov}[x_1, x_2]/(\sigma_1 \sigma_2)$  is the correlation coefficient.

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## Chi-square $(\chi^2)$ distribution

The chi-square pdf for the continuous r.v.  $z \ (z \ge 0)$  is defined by

$$f(z;n) = \frac{1}{2^{n/2} \Gamma(n/2)} z^{n/2-1} e^{-z/2} \left\{ \begin{array}{c} 0.5 \\ 0.4 \\ 0.3 \\ 0.3 \\ 0.1 \\ 0 \end{array} \right\}_{\substack{n=1, 2, \dots = n \text{ number of 'degrees of freedom' (dof)}} \\ E[z] = n, \quad V[z] = 2n. \end{array} \right\}_{\substack{n=1, 2, \dots = n \text{ number of 'degrees of freedom' (dof)}}$$

For independent Gaussian  $x_i$ , i = 1, ..., n, means  $\mu_i$ , variances  $\sigma_i^2$ ,

$$z = \sum_{i=1}^{n} \frac{(x_i - \mu_i)^2}{\sigma_i^2} \quad \text{follows } \chi^2 \text{ pdf with } n \text{ dof.}$$

Example: goodness-of-fit test variable especially in conjunction with method of least squares.

G. Cowan

## Cauchy (Breit-Wigner) distribution

The Breit-Wigner pdf for the continuous r.v. x is defined by

$$f(x; \Gamma, x_0) = \frac{1}{\pi} \frac{\Gamma/2}{\Gamma^2/4 + (x - x_0)^2}$$
  

$$(\Gamma = 2, x_0 = 0 \text{ is the Cauchy pdf.})$$
  

$$E[x] \text{ not well defined, } V[x] \to \infty.$$
  

$$x_0 = \text{ mode (most probable value)}$$
  

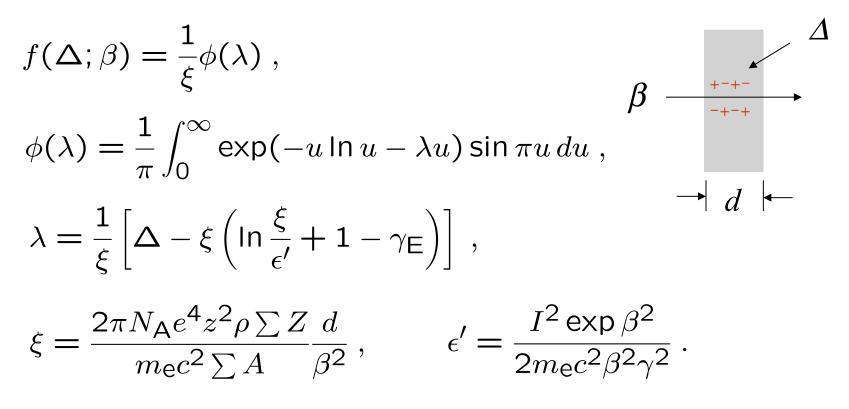
$$\Gamma = \text{ full width at half maximum}$$

Example: mass of resonance particle, e.g.  $\rho$ , K<sup>\*</sup>,  $\phi^0$ , ...  $\Gamma$  = decay rate (inverse of mean lifetime)

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## Landau distribution

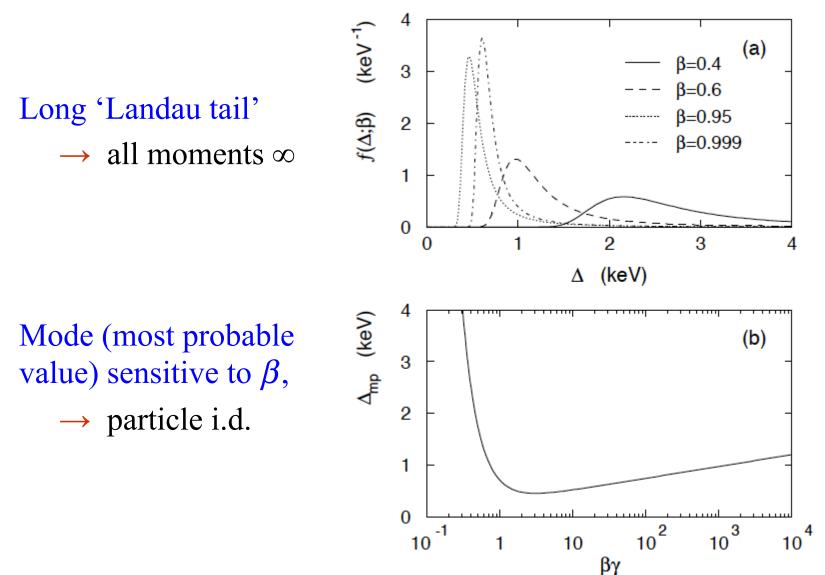
For a charged particle with  $\beta = v/c$  traversing a layer of matter of thickness *d*, the energy loss  $\Delta$  follows the Landau pdf:



L. Landau, J. Phys. USSR **8** (1944) 201; see also W. Allison and J. Cobb, Ann. Rev. Nucl. Part. Sci. **30** (1980) 253.

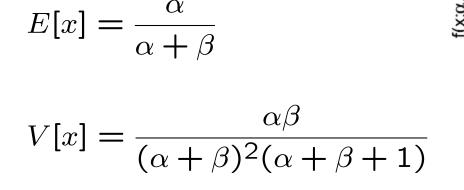
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## Landau distribution (2)

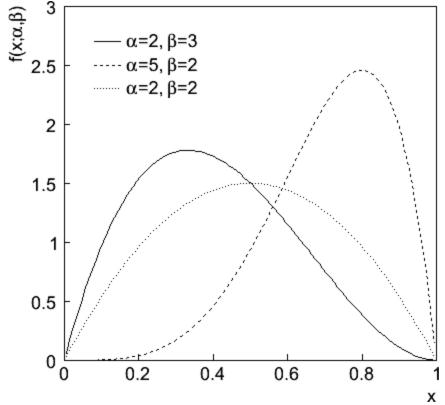


## Beta distribution

$$f(x;\alpha,\beta) = \frac{\Gamma(\alpha+\beta)}{\Gamma(\alpha)\Gamma(\beta)} x^{\alpha-1} (1-x)^{\beta-1}$$



Often used to represent pdf of continuous r.v. nonzero only between finite limits.



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## Gamma distribution

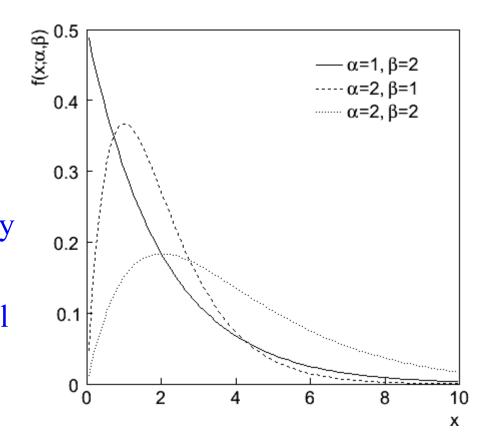
$$f(x; \alpha, \beta) = \frac{1}{\Gamma(\alpha)\beta^{\alpha}} x^{\alpha-1} e^{-x/\beta}$$

$$V[x] = \alpha \beta^2$$

 $E[r] = \alpha \beta$ 

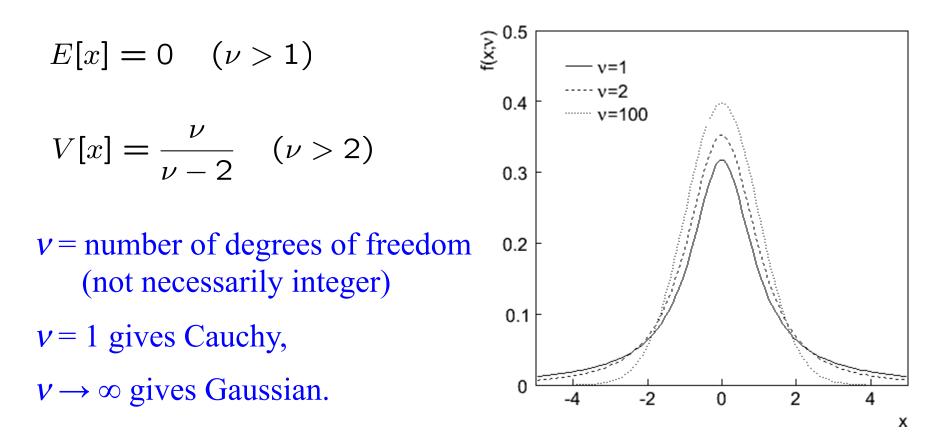
Often used to represent pdf of continuous r.v. nonzero only in  $[0,\infty]$ .

Also e.g. sum of *n* exponential r.v.s or time until *n*th event in Poisson process ~ Gamma



## Student's t distribution

$$f(x;\nu) = \frac{\Gamma\left(\frac{\nu+1}{2}\right)}{\sqrt{\nu\pi}\,\Gamma(\nu/2)} \left(1 + \frac{x^2}{\nu}\right)^{-\left(\frac{\nu+1}{2}\right)}$$



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## Student's *t* distribution (2)

If 
$$x \sim \text{Gaussian}$$
 with  $\mu = 0$ ,  $\sigma^2 = 1$ , and

$$z \sim \chi^2$$
 with *n* degrees of freedom, then

 $t = x / (z/n)^{1/2}$  follows Student's t with v = n.

This arises in problems where one forms the ratio of a sample mean to the sample standard deviation of Gaussian r.v.s.

The Student's *t* provides a bell-shaped pdf with adjustable tails, ranging from those of a Gaussian, which fall off very quickly,  $(v \rightarrow \infty)$ , but in fact already very Gauss-like for v = two dozen), to the very long-tailed Cauchy (v = 1).

Developed in 1908 by William Gosset, who worked under the pseudonym "Student" for the Guinness Brewery.

## The Monte Carlo method

What it is: a numerical technique for calculating probabilities and related quantities using sequences of random numbers.

The usual steps:

- (1) Generate sequence  $r_1, r_2, ..., r_m$  uniform in [0, 1].
- g(r) f(r) f(r) r 0 1
- Use this to produce another sequence x<sub>1</sub>, x<sub>2</sub>, ..., x<sub>n</sub> distributed according to some pdf f(x) in which we're interested (x can be a vector).
- (3) Use the *x* values to estimate some property of f(x), e.g., fraction of *x* values with a < x < b gives  $\int_a^b f(x) dx$ .

 $\rightarrow$  MC calculation = integration (at least formally)

MC generated values = 'simulated data'

 $\rightarrow$  use for testing statistical procedures

## Random number generators

- Goal: generate uniformly distributed values in [0, 1]. Toss coin for e.g. 32 bit number... (too tiring).
  - $\rightarrow$  'random number generator'
  - = computer algorithm to generate  $r_1, r_2, ..., r_n$ .

Example: multiplicative linear congruential generator (MLCG)

 $n_{i+1} = (a \ n_i) \mod m$ , where  $n_i = \text{integer}$  a = multiplier m = modulus $n_0 = \text{seed (initial value)}$ 

N.B. mod = modulus (remainder), e.g. 27 mod 5 = 2. This rule produces a sequence of numbers  $n_0, n_1, ...$ 

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## Random number generators (2)

### The sequence is (unfortunately) periodic!

Example (see Brandt Ch 4):  $a = 3, m = 7, n_0 = 1$ 

$$n_1 = (3 \cdot 1) \mod 7 = 3$$

$$n_2 = (3 \cdot 3) \mod 7 = 2$$

$$n_3 = (3 \cdot 2) \mod 7 = 6$$

$$n_4 = (3 \cdot 6) \mod 7 = 4$$

$$n_5 = (3 \cdot 4) \mod 7 = 5$$

$$n_6 = (3 \cdot 5) \mod 7 = 1 \quad \leftarrow \text{ sequence repeats}$$

Choose *a*, *m* to obtain long period (maximum = m - 1); *m* usually close to the largest integer that can represented in the computer.

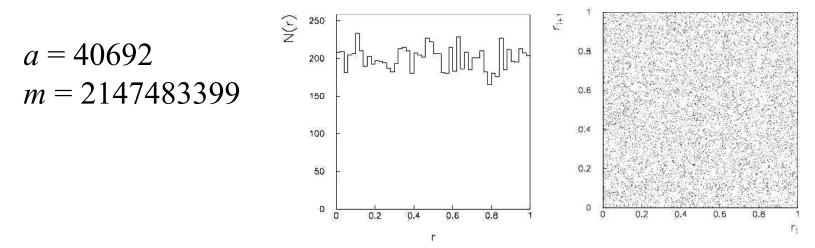
Only use a subset of a single period of the sequence.

## Random number generators (3)

 $r_i = n_i/m$  are in [0, 1] but are they 'random'?

Choose *a*, *m* so that the  $r_i$  pass various tests of randomness: uniform distribution in [0, 1],

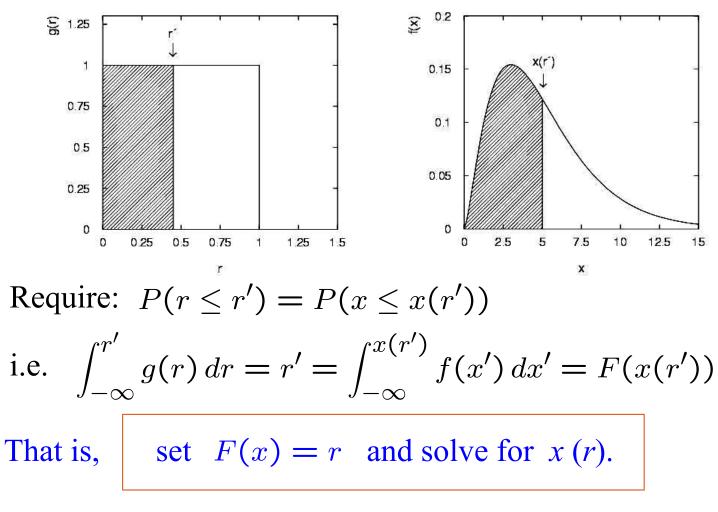
all values independent (no correlations between pairs), e.g. L'Ecuyer, Commun. ACM **31** (1988) 742 suggests



Far better generators available, e.g. **TRandom3**, based on Mersenne twister algorithm, period =  $2^{19937} - 1$  (a "Mersenne prime"). See F. James, Comp. Phys. Comm. 60 (1990) 111; Brandt Ch. 4 G. Cowan iSTEP 2015, Jinan / Statistics for Particle Physics / Lecture 1

## The transformation method

Given  $r_1, r_2, ..., r_n$  uniform in [0, 1], find  $x_1, x_2, ..., x_n$  that follow f(x) by finding a suitable transformation x(r).



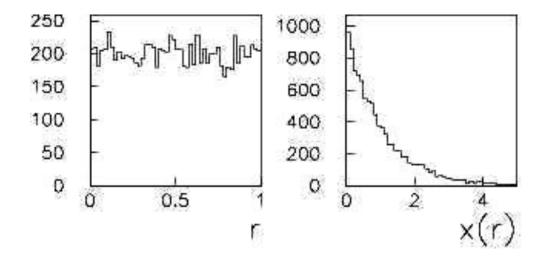
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Example of the transformation method

Exponential pdf: 
$$f(x;\xi) = \frac{1}{\xi}e^{-x/\xi}$$
  $(x \ge 0)$ 

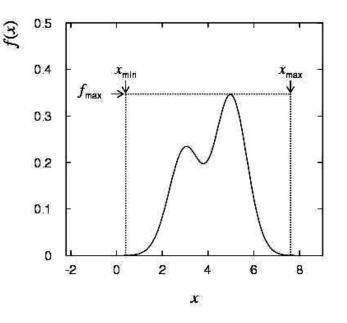
Set 
$$\int_0^x \frac{1}{\xi} e^{-x'/\xi} dx' = r$$
 and solve for  $x(r)$ .

$$\rightarrow x(r) = -\xi \ln(1-r) \quad (x(r) = -\xi \ln r \text{ works too.})$$



## The acceptance-rejection method

Enclose the pdf in a box:



(1) Generate a random number x, uniform in  $[x_{\min}, x_{\max}]$ , i.e.  $x = x_{\min} + r_1(x_{\max} - x_{\min})$ ,  $r_1$  is uniform in [0,1].

(2) Generate a 2nd independent random number u uniformly distributed between 0 and  $f_{\text{max}}$ , i.e.  $u = r_2 f_{\text{max}}$ .

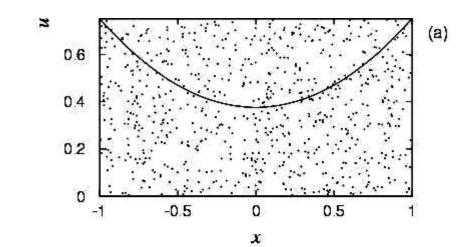
(3) If u < f(x), then accept x. If not, reject x and repeat.

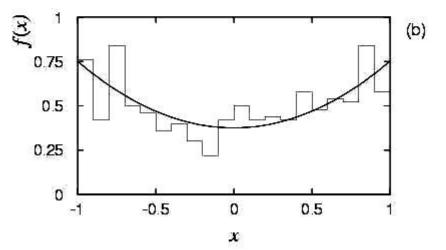
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## Example with acceptance-rejection method

$$f(x) = \frac{3}{8}(1+x^2)$$
  
(-1 \le x \le 1)

If dot below curve, use *x* value in histogram.





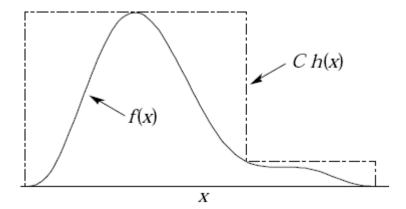
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# Improving efficiency of the acceptance-rejection method

The fraction of accepted points is equal to the fraction of the box's area under the curve.

For very peaked distributions, this may be very low and thus the algorithm may be slow.

Improve by enclosing the pdf f(x) in a curve C h(x) that conforms to f(x) more closely, where h(x) is a pdf from which we can generate random values and C is a constant.



Generate points uniformly over C h(x).

If point is below f(x), accept x.

## Monte Carlo event generators

Simple example:  $e^+e^- \rightarrow \mu^+\mu^-$ 

Generate  $\cos\theta$  and  $\phi$ :

$$e^+$$
  $e^ e^-$ 

$$f(\cos\theta; A_{\text{FB}}) \propto \left(1 + \frac{8}{3}A_{\text{FB}}\cos\theta + \cos^2\theta\right),$$
$$g(\phi) = \frac{1}{2\pi} \quad (0 \le \phi \le 2\pi)$$

Less simple: 'event generators' for a variety of reactions:  $e^+e^- \rightarrow m^+m^-$ , hadrons, ...  $pp \rightarrow$  hadrons, D-Y, SUSY,...

e.g. PYTHIA, HERWIG, ISAJET...

Output = 'events', i.e., for each event we get a list of generated particles and their momentum vectors, types, etc.

>	~~										>	<	
	Event listing (summary)										Δ		
	T	particle/jet	KS	KF	orig	P_X	P_9	p_z	E		m		
	1	ip+i ip+i	21 21	2212 2212	0 0	0.000 0.000		7000,000 -7000,000			0,938 0,938		
ľ	3		21 21 21 21	21 -2 21	1	0,863		1739,862	1739,		0.000		
	4	!ubar!	21	-2	23	-0,621	-0,163	-777,415	777.	.415	0,000		
	5	!9! !9!	21	21	3	-2,427	5,486	1487,857	1487.	X~			
	6	! <u>9</u> !	21	21 1000021	4	-62,910	63,357	-463,274	471.	207		4	047
	7	!~9!	21	1000021	0	314,363	544,843	498,897	979.	397	pi+	1 1	211 22 21 111 111 21 21 21 211 321 -211 311 -311 321 310 321 -211 310 321 -211
	8	!~g!	21	1000021	0	-379,700	-476,000 112,247	525,686	980.	398 399	gamma	1	24
	9 10	!~chi_1-! !sbar!	211	-1000024 -3	7	130,058 259,400	112,247	129,860 83,100	263. 330.	400	gamma (pi0)	11	111
		isbari Ic!	21 21	-5		-79,403	242,409	283,026	381.	401	(pi0)	11	111
	12	!~chi_20!	- 21	1000023	7 8	-326,241	-80,971	113.712	385.	402	(pi0)	11	11
		161	21	1000023 5 -5	8	-51,841	-294.077	389,853	491	403	gamma	1	2
	14	!bbar!	21	5 -5	Ř	-0,597	-99,577	21,299	491. 101.	404	gamma	1 1 1 1 1 1	22
		!"chi_10!	21	1000022	8	103,352	81.316	83,457	175. 65. 22. 181.	405	pi-	1	-211
	16	Isl	21	3	. ě	5,451	38.374	52,302	65	406	pi+	1	211
		[cbar]	21	3 -4	9	20,839	-7,250	-5,938	22.	407	Κ+	1	321
	18	!~chi_10!	21	1000022	12	-136,266	-72,961	53,246	181.	408	pi-	1	-211
	- 19		21	14	12	-78,263	-24,757	21,719	84.	409	(pi0)	11	111
	- 20	!nu_mubar!	21	-14	12	-107,801	16,901	38,226	115.	410	(pi0)	11	111
										411	(Kbar0)	11 1 1	-311
	21	gamma	1	22	- 4	2,636	1,357	0,125	2. 262.	412	Pi-	1	-211
	22	(~chi_1-)		-1000024	9	129,643	112,440	129,820	262.	413	K+	1	321
	-23	(~chi_20)		1000023	12	-322,330	-80,817	113,191	382.	414	(pi0)	11	111
	-24	~chi_10	1	1000022	15	97,944	77,819	80,917	169.	415	(K_S0)	11	510
	25	~chi_10		1000022	18	-136,266	-72,961	53,246	181.	416		11 1 1	521
	26	nu_mu	1	14	19 20	-78,263	-24,757	21,719	84.		pi-	1	-Z11
	27	nu_mubar	1 11	-14 2224	20	-107,801	16,901	38,226 -2734,287	115. 2734.	410	nbar0 (pi0)	11	-211/
	28	(Delta++)	11	2224	Z	0,222	0,012	-2754,287	27.54.		pi+	1	211
:										420	(pi0)	11	111
										422	n0	1	2113
					•					423	pi-	1	-211
					•					424	gamma	1 1 1 1	- 22
					•					425	gamma	1	22
										426	pi+	1	211
										427	(pi0)	11	111
DVTUIA Monto Corlo								428	pi-	1	11: 21: 21: 21: 21: 22: 21: 21: 21: 11: 21: 11: 21: 11: 21: 2		
PYTHIA Monte Carlo								429	(pi0)	11	111		
$pp \rightarrow gluino-gluino$								gamma	1	- 23			
		$nn \rightarrow$	σ	luin	0-	ماييناه	10			431	gamma	1	22
		ЧЧ 🧹	Ĕ	, um		Siuli	U			8			
									-				

## A simulated event

								-OX
	1	211	209	0,006	0,398	-308,296	308,297	0,140
	1 1 1 11	22	211	0,407		1695,458	1695,458	0.000
	1	22	211	0,113	-0,029	-314,822	314,822	0.000
	11	111	212	0,021	0,122	-103,709	103,709	0,135
	11	111	212	0,084	-0,068	-94,276	94.276	0.135
	11	111	212	0,267	-0,052	-144,673	144,674	0,135
	1	22	215	-1,581	2.473	3,306	4.421	0.000
	1	22	215	-1,494	2,143	3,051	4,016	0.000
	1	-211	216	0,007	0,738	4,015	4,085	0,140
	1 1 1 1 1 1 1 1 1	211	216	-0,024	0,293	0,486	0,585	0,140
	1	321	218	4,382	-1,412	-1,799	4,968	0,494
	1	-211	218	1.183	-0,894	-0,176	1,500	0,140
		111	218	0,955	-0,459	-0,590	1,221	0,135
	11	111	218	0,955 2,349	-1,105	-1.181	2.855	0.135
0)	11	-311	219	1,441 2,232 1,380	-1.105 -0.247	-0,472	1.615 2.285 1.644	0,498
	1	-211 321	219	2,232	-0,400 -0,652	-0.249 -0.361	2,285	0.140 0.494
	1 1 11	321	220	1,380	-0,652	-0,361	1.644	0,494
	11	111	220	1.078 1.841	-0,265	0,175	1.132	0,135
)	11	310	222	1.841	0,111	0,894	2,109	0,498
	1 1 1	321	223	0,307	0,107	0,252	0,642	0,494
	1	-211	223	0,266	0.316	-0,201	0,480	0,140
	1	-2112	226	1,335	1,641	2,078	3,111	0,940
	11	111	226	0.899 0.217	1,046	1,311	1,908	0,135
	1	211	227	0,217	1,407	1,356	1,971	0,140
	11	111	227	1,207	2.336 5.324	1,311 1,356 2,767 5,702	3,820 8,592	0,135
	1	2112	228	3,475	5,324	5,702	8,592	0,940
	1	-211	228 229	1,856	2,606	2,808	4,259	0,140
	1	22	229	-0,012	0,247	0,421	0,489	0,000
	1 1 1 1 1	22	229	0,025	0.034	0,009	0,043	0,000
	1	211	230	2,718	5,229	6,403	8,703	0,140
	11	111	230	4,109	6,747	7,597	10,961	0,135
	1	-211	231	0,551 0,645	1,233	1,945	2,372	0,140
	11	111	231	0,645	1,141	0,922	1,608 1,724	0,135
	1 1	22	232	-0,383	1,169	1,208	1,724	0.000
	1	22	232	-0,201	0,070	0,060	0,221	0,000

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## Monte Carlo detector simulation

Takes as input the particle list and momenta from generator.

Simulates detector response:

multiple Coulomb scattering (generate scattering angle), particle decays (generate lifetime), ionization energy loss (generate  $\Delta$ ), electromagnetic, hadronic showers, production of signals, electronics response, ...

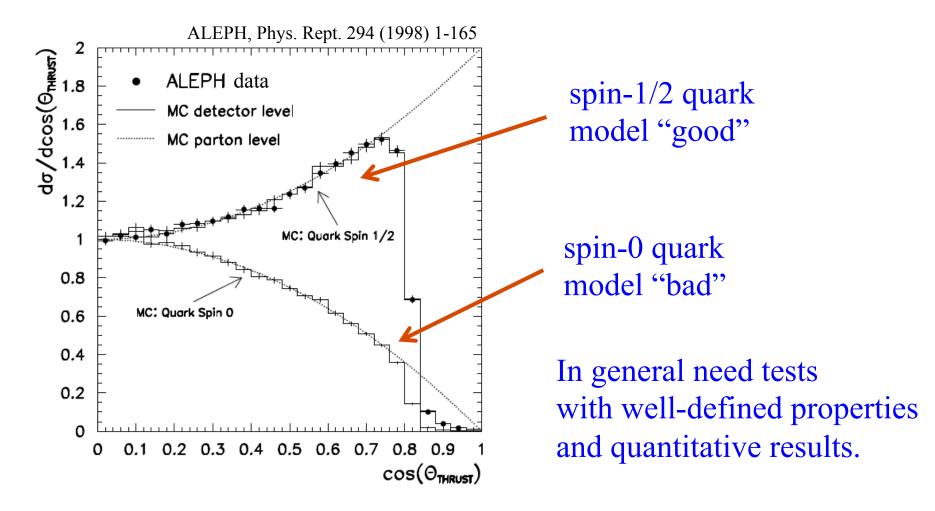
Output = simulated raw data  $\rightarrow$  input to reconstruction software: track finding, fitting, etc.

Predict what you should see at 'detector level' given a certain hypothesis for 'generator level'. Compare with the real data. Estimate 'efficiencies' = #events found / # events generated. Programming package: GEANT

G. Cowan

## Data analysis in particle physics: testing hypotheses

Test the extent to which a given model agrees with the data:



## Choosing a critical region

To construct a test of a hypothesis  $H_0$ , we can ask what are the relevant alternatives for which one would like to have a high power.

Maximize power wrt  $H_1$  = maximize probability to reject  $H_0$  if  $H_1$  is true.

Often such a test has a high power not only with respect to a specific point alternative but for a class of alternatives. E.g., using a measurement  $x \sim \text{Gauss}(\mu, \sigma)$  we may test

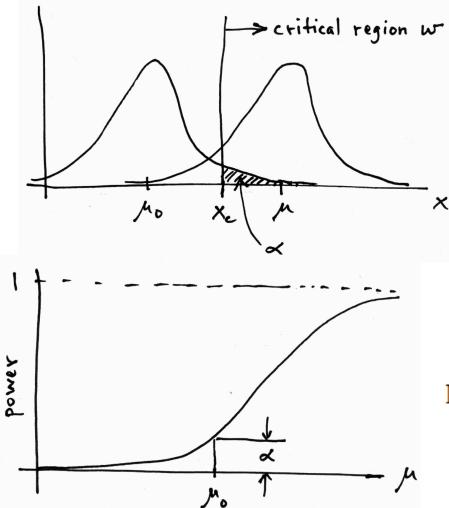
 $H_0: \mu = \mu_0$  versus the composite alternative  $H_1: \mu > \mu_0$ 

We get the highest power with respect to any  $\mu > \mu_0$  by taking the critical region  $x \ge x_c$  where the cut-off  $x_c$  is determined by the significance level such that

$$\alpha = P(x \ge x_c | \mu_0).$$

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Test of 
$$\mu = \mu_0$$
 vs.  $\mu > \mu_0$  with  $x \sim \text{Gauss}(\mu, \sigma)$ 



Standard Gaussian cumulative distribution

$$\alpha = 1 - \Phi\left(\frac{x_{\rm c} - \mu_0}{\sigma}\right)$$

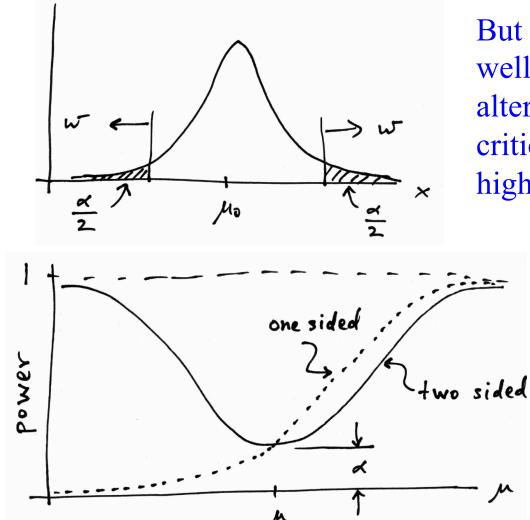
$$x_{\rm c} = \mu_0 + \sigma \Phi^{-1} (1 - \alpha)$$

Standard Gaussian quantile

$$power = 1 - \beta = P(x > x_c | \mu) =$$

$$1 - \Phi\left(\frac{\mu_0 - \mu}{\sigma} + \Phi^{-1}(1 - \alpha)\right)$$

## Choice of critical region based on power (3)



But we might consider  $\mu < \mu_0$  as well as  $\mu > \mu_0$  to be viable alternatives, and choose the critical region to contain both high and low *x* (a two-sided test).

> New critical region now gives reasonable power for  $\mu < \mu_0$ , but less power for  $\mu > \mu_0$  than the original one-sided test.

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No such thing as a model-independent test In general we cannot find a single critical region that gives the maximum power for all possible alternatives (no "Uniformly Most Powerful" test).

In HEP we often try to construct a test of

 $H_0$ : Standard Model (or "background only", etc.)

such that we have a well specified "false discovery rate",

 $\alpha$  = Probability to reject  $H_0$  if it is true,

and high power with respect to some interesting alternative,

 $H_1$ : SUSY, Z', etc.

But there is no such thing as a "model independent" test. Any statistical test will inevitably have high power with respect to some alternatives and less power with respect to others.

G. Cowan

## Rejecting a hypothesis

Note that rejecting  $H_0$  is not necessarily equivalent to the statement that we believe it is false and  $H_1$  true. In frequentist statistics only associate probability with outcomes of repeatable observations (the data).

In Bayesian statistics, probability of the hypothesis (degree of belief) would be found using Bayes' theorem:

$$P(H|x) = \frac{P(x|H)\pi(H)}{\int P(x|H)\pi(H) \, dH}$$

which depends on the prior probability  $\pi(H)$ .

What makes a frequentist test useful is that we can compute the probability to accept/reject a hypothesis assuming that it is true, or assuming some alternative is true.

G. Cowan