# Statistics for Particle Physics Lecture day 1





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# Outline

$\rightarrow$ Monday 9-11 :	Introduction
	Probability
	Hypothesis tests
	Machine Learning
Tuesday 9-11 :	Parameter estimation
	Confidence limits
	Systematic uncertainties
	Experimental sensitivity
Tuesday 15:30:	Tutorial on parameter estimation

Almost everything is a subset of the University of London course: http://www.pp.rhul.ac.uk/~cowan/stat\_course.html

# 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.
- Luca Lista, *Statistical Methods for Data Analysis in Particle Physics*, Springer, 2017.
- 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.
- R.L. Workman et al. (Particle Data Group), Prog. Theor. Exp. Phys. 083C01 (2022); pdg.1b1.gov sections on probability, statistics, MC.

# Theory ↔ Statistics ↔ Experiment

Theory (model, hypothesis):

Experiment (observation):

$$F = -G \frac{m_i m_2}{r^2}, \dots$$

+ response of measurement apparatus

= model prediction



## A quick review of probability

Frequentist (*A* = outcome of repeatable observation)

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

Subjective (*A* = hypothesis)

$$P(A) =$$
degree of belief that A is true

Conditional probability:

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

A and B are independent iff:

$$P(A \cap B) = P(A)P(B)$$

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

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#### Bayes' theorem

Use definition of conditional probability and  $P(A \cap B) = P(B \cap A)$ 

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

(Bayes' theorem)

If set of all outcomes  $S = \bigcup_i A_i$ with  $A_i$  disjoint, then law of total probability for P(B) says



$$P(B) = \sum_{i} P(B|A_i)P(A_i)$$

so that Bayes' theorem becomes

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

Bayes' theorem holds regardless of how probability is interpreted (frequency, degree of belief...).

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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: x).

Probability = limiting frequency

Probabilities such as

P (string theory is true), P (0.117 <  $\alpha_s$  < 0.119), P (Biden wins in 2024),

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.

Preferred theories (models, hypotheses, ...) are those that predict a high probability for data "like" the data observed.

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

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!)

# Hypothesis, likelihood

Suppose the entire result of an experiment (set of measurements) is a collection of numbers x.

A (simple) hypothesis is a rule that assigns a probability to each possible data outcome:

 $P(\mathbf{x}|H)$  = the likelihood of H

Often we deal with a family of hypotheses labeled by one or more undetermined parameters (a composite hypothesis):

$$P(\mathbf{x}|oldsymbol{ heta}) = L(oldsymbol{ heta})$$
 = the "likelihood function"

Note:

1) For the likelihood we treat the data x as fixed.

2) The likelihood function  $L(\theta)$  is not a pdf for  $\theta$ .

# Frequentist hypothesis tests

Suppose a measurement produces data x; consider a hypothesis  $H_0$  we want to test and alternative  $H_1$ 

 $H_0$ ,  $H_1$  specify probability for  $\mathbf{x}$ :  $P(\mathbf{x}|H_0)$ ,  $P(\mathbf{x}|H_1)$ 

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.,

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

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  $H_0$ .



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# Definition of a test (2)

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

Use the alternative hypothesis  $H_1$  to motivate where to place the critical region.

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



#### Classification viewed as a statistical test

Suppose events come in two possible types:

s (signal) and b (background)

For each event, test hypothesis that it is background, i.e.,  $H_0 = b$ .

Carry out test on many events, each is either of type s or b, i.e., here the hypothesis is the "true class label", which varies randomly from event to event, so we can assign to it a frequentist probability.

Select events for which where  $H_0$  is rejected as "candidate events of type s". Equivalent Particle Physics terminology:

background efficiency 
$$arepsilon_{
m b} = \int_W f({f x}|H_0) \, d{f x} = lpha$$

 $\varepsilon_{\mathbf{s}} = \int_{W} f(\mathbf{x}|H_1) \, d\mathbf{x} = 1 - \beta = \text{power}$ 

signal efficiency

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# Example of a test for classification



For each event in a mixture of signal (s) and background (b) test

 $H_0$ : event is of type b

using a critical region W of the form:  $W = \{x : x \le x_c\}$ , where  $x_c$  is a constant that we choose to give a test with the desired size  $\alpha$ .

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# Classification example (2)

Suppose we want  $\alpha = 10^{-4}$ . Require:

$$\alpha = P(x \le x_{c}|b) = \int_{0}^{x_{c}} f(x|b) \, dx = \frac{4x^{4}}{4} \Big|_{0}^{x_{c}} = x_{c}^{4}$$

and therefore  $x_{\rm c} = \alpha^{1/4} = 0.1$ 

For this test (i.e. this critical region W), the power with respect to the signal hypothesis (s) is

$$M = P(x \le x_{\rm c}|{\rm s}) = \int_0^{x_{\rm c}} f(x|{\rm s}) \, dx = 2x_{\rm c} - x_{\rm c}^2 = 0.19$$

Note: the optimal size and power is a separate question that will depend on goals of the subsequent analysis.

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## Classification example (3)

Suppose that the prior probabilities for an event to be of type s or b are:

 $\pi_{\rm s} = 0.001$  $\pi_{\rm b} = 0.999$ 

The "purity" of the selected signal sample (events where b hypothesis rejected) is found using Bayes' theorem:

$$P(\mathbf{s}|x \le x_{\mathbf{c}}) = \frac{P(x \le x_{\mathbf{c}}|\mathbf{s})\pi_{\mathbf{s}}}{P(x \le x_{\mathbf{c}}|\mathbf{s})\pi_{\mathbf{s}} + P(x \le x_{\mathbf{c}}|\mathbf{b})\pi_{\mathbf{b}}}$$

= 0.655

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# Testing significance / goodness-of-fit

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

We observe a single point in this space:  $x_{obs}$ .

 $X_i$ 

How can we quantify the level of compatibility between the data and the predictions of *H*?

Decide what part of the data space represents equal or less compatibility with H than does the point  $x_{obs}$ . (Not unique!)



# *p*-values

Express level of compatibility between data and hypothesis (sometimes 'goodness-of-fit') by giving the *p*-value for *H*:

 $p = P(\mathbf{x} \in \omega_{\leq}(\mathbf{x}_{obs})|H)$ 

- probability, under assumption of H, to observe data
   with equal or lesser compatibility with H relative to the
   data we got.
- probability, under assumption of H, to observe data as discrepant with H as the data we got or more so.

Basic idea: if there is only a very small probability to find data with even worse (or equal) compatibility, then *H* is "disfavoured by the data".

If the *p*-value is below a user-defined threshold  $\alpha$  (e.g. 0.05) then *H* is rejected (equivalent to hypothesis test of size  $\alpha$  as seen earlier).



The *p*-value of H is not the probability that *H* is true!

In frequentist statistics we don't talk about P(H) (unless H represents a repeatable observation).

If we do define P(H), e.g., in Bayesian statistics as a degree of belief, then we need to use Bayes' theorem to obtain

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

where  $\pi(H)$  is the prior probability for *H*.

For now stick with the frequentist approach; result is p-value, regrettably easy to misinterpret as P(H). 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.

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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# 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.



$$p = \int_{Z}^{\infty} \frac{1}{\sqrt{2\pi}} e^{-x^2/2} dx = 1 - \Phi(Z)$$
  
 $Z = \Phi^{-1}(1-p)$ 

in ROOT: p = 1 - TMath::Freq(Z) Z = TMath::NormQuantile(1-p)Z = r

in python (scipy.stats):
p = 1 - norm.cdf(Z) = norm.sf(Z)
Z = norm.ppf(1-p)

Result Z is a "number of sigmas". Note this does not mean that the original data was Gaussian distributed.

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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.

# Particle Physics context for a hypothesis test

#### A simulated SUSY event ("signal"):



# **Background events**



This event from Standard Model ttbar production also has high  $p_{\rm T}$  jets and muons, and some missing transverse energy.

→ can easily mimic a signal event.

# Classification of proton-proton collisions

Proton-proton collisions can be considered to come in two classes: signal (the kind of event we're looking for, y = 1) background (the kind that mimics signal, y = 0)

For each collision (event), we measure a collection of features:

 $x_1 =$  energy of muon $x_4 =$  missing transverse energy $x_2 =$  angle between jets $x_5 =$  invariant mass of muon pair $x_3 =$  total jet energy $x_6 = \dots$ 

The real events don't come with true class labels, but computersimulated events do. So we can have a set of simulated events that consist of a feature vector x and true class label y (0 for background, 1 for signal):

$$(x, y)_1, (x, y)_2, ..., (x, y)_N$$

The simulated events are called "training data".

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# **Distributions of the features**

If we consider only two features  $x = (x_1, x_2)$ , we can display the results in a scatter plot (red: y = 0, blue: y = 1).

For real events, the dots are black (true type is not known).

For each real event test the hypothesis that it is background.

(Related to this: test that a sample of events is *all* background.)

The test's critical region is defined by a "decision boundary" – without knowing the event type, we can classify them by seeing where their measured features lie relative to the boundary.



# Decision function, test statistic



A surface in an *n*-dimensional

space can be described by

Different values of the constant  $t_c$  result in a family of surfaces.

Problem is reduced to finding the best decision function or test statistic t(x).



# Distribution of t(x)

By forming a test statistic t(x), the boundary of the critical region in the *n*-dimensional *x*-space is determined by a single single value  $t_c$ .



# Types of decision boundaries

So what is the optimal boundary for the critical region, i.e., what is the optimal test statistic t(x)?

First find best t(x), later address issue of optimal size of test.

Remember *x*-space can have many dimensions.



#### Test statistic based on likelihood ratio

How can we choose a test's critical region in an 'optimal way', in particular if the data space is multidimensional?

Neyman-Pearson lemma states:

For a test of  $H_0$  of size  $\alpha$ , to get the highest power with respect to the alternative  $H_1$  we need for all x in the critical region W

"likelihood ratio (LR)" 
$$\frac{P(\mathbf{x}|H_1)}{P(\mathbf{x}|H_0)} \ge c_{\alpha}$$

inside W and  $\leq c_{\alpha}$  outside, where  $c_{\alpha}$  is a constant chosen to give a test of the desired size.

Equivalently, optimal scalar test statistic is

$$t(\mathbf{x}) = \frac{P(\mathbf{x}|H_1)}{P(\mathbf{x}|H_0)}$$

N.B. any monotonic function of this is leads to the same test.

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#### Neyman-Pearson doesn't usually help

We usually don't have explicit formulae for the pdfs f(x|s), f(x|b), so for a given x we can't evaluate the likelihood ratio

$$t(\mathbf{x}) = \frac{f(\mathbf{x}|s)}{f(\mathbf{x}|b)}$$

Instead we may have Monte Carlo models for signal and background processes, so we can produce simulated data:

generate 
$$\boldsymbol{x} \sim f(\boldsymbol{x}|\mathbf{s}) \rightarrow \boldsymbol{x}_1, \dots, \boldsymbol{x}_N$$

generate 
$$\mathbf{x} \sim f(\mathbf{x}|\mathbf{b}) \rightarrow \mathbf{x}_1, ..., \mathbf{x}_N$$

This gives samples of "training data" with events of known type.

 Use these to construct a statistic that is as close as possible to the optimal likelihood ratio (→ Machine Learning).

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# Approximate LR from histograms

Want t(x) = f(x|s)/f(x|b) for x here



One possibility is to generate MC data and construct histograms for both signal and background.

Use (normalized) histogram values to approximate LR:

$$t(x) \approx \frac{N(x|s)}{N(x|b)}$$

Can work well for single variable.

# Approximate LR from 2D-histograms Suppose problem has 2 variables. Try using 2-D histograms:



Approximate pdfs using N(x,y|s), N(x,y|b) in corresponding cells. But if we want M bins for each variable, then in n-dimensions we have  $M^n$  cells; can't generate enough training data to populate.  $\rightarrow$  Histogram method usually not usable for n > 1 dimension.

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#### Strategies for multivariate analysis

Neyman-Pearson lemma gives optimal answer, but cannot be used directly, because we usually don't have  $f(\mathbf{x}|\mathbf{s}), f(\mathbf{x}|\mathbf{b})$ .

Histogram method with M bins for n variables requires that we estimate  $M^n$  parameters (the values of the pdfs in each cell), so this is rarely practical.

A compromise solution is to assume a certain functional form for the test statistic t(x) with fewer parameters; determine them (using MC) to give best separation between signal and background.

Alternatively, try to estimate the probability densities f(x|s) and f(x|b) (with something better than histograms) and use the estimated pdfs to construct an approximate likelihood ratio.

# Multivariate methods (Machine Learning)

#### Many new (and some old) methods:

Fisher discriminant (Deep) Neural Networks Kernel density methods Support Vector Machines Decision trees Boosting

Bagging

#### Linear test statistic

 $\boldsymbol{n}$ 

Suppose there are *n* input variables:  $\mathbf{x} = (x_1, ..., x_n)$ .

Consider a linear function: 
$$y(\mathbf{x}) = \sum_{i=1}^{n} w_i x_i$$

For a given choice of the coefficients  $w = (w_1, ..., w_n)$  we will get pdfs f(y|s) and f(y|b):


#### Linear test statistic

Fisher: to get large difference between means and small widths for f(y|s) and f(y|b), maximize the difference squared of the expectation values divided by the sum of the variances:

$$J(\mathbf{w}) = \frac{(E[y|s] - E[y|b])^2}{V[y|s] + V[y|b]}$$

Setting  $\partial J / \partial w_i = 0$  gives:

$$\mathbf{w} \propto W^{-1}(\boldsymbol{\mu}_{\rm b} - \boldsymbol{\mu}_{\rm s})$$
$$W_{ij} = \operatorname{cov}[x_i, x_j | {\rm s}] + \operatorname{cov}[x_i, x_j | {\rm b}]$$
$$\mu_{i, {\rm s}} = E[x_i | s], \qquad \mu_{i, {\rm b}} = E[x_i | b]$$

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#### The Fisher discriminant

The resulting coefficients  $w_i$  define a Fisher discriminant.

Coefficients defined up to multiplicative constant; can also add arbitrary offset, i.e., usually define test statistic as

$$y(\mathbf{x}) = w_0 + \sum_{i=1}^n w_i x_i$$

Boundaries of the test's critical region are surfaces of constant y(x), here linear (hyperplanes):



### Nonlinear decision boundaries

From the scatter plot below it's clear that some nonlinear boundary would be better than a linear one:



And to have a nonlinear boundary, the decision function t(x) must be nonlinear in x.

#### **Neural Networks**

A simple nonlinear decision function can be constructed as

$$t(\mathbf{x}) = h\left(w_0 + \sum_{i=1}^n w_i x_i\right)$$

where *h* is called the "activation function". For this one can use, e.g., a logistic sigmoid function,



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### **Single Layer Perceptron**

In this form, the decision function is called a Single Layer Perceptron – the simplest example of a Neural Network.



But the surface described by  $t(\mathbf{x}) = t_c$  is the same as by

$$h^{-1}(t(\mathbf{x})) = w_0 + \sum_{i=1}^n w_i x_i = h^{-1}(t_c)$$

So here we still have a linear decision boundary.

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### **Multilayer Perceptron**

The Single Layer Perceptron can be generalized by defining first a set of functions  $\varphi_i(\mathbf{x})$ , with i = 1, ..., m:

$$\varphi_i(\mathbf{x}) = h\left(w_{i0}^{(1)} + \sum_{j=1}^n w_{ij}^{(1)} x_j\right)$$
  $i = 1, \dots, m$ 

The  $\varphi_i(\mathbf{x})$  are then treated as if they were the input variables, in a perceptron, i.e., the decision function (output node) is

$$t(\mathbf{x}) = h\left(w_{10}^{(2)} + \sum_{j=1}^{n} w_{1j}^{(2)}\varphi_j(\mathbf{x})\right)$$

# Multilayer Perceptron (2)



Each line in the graph represents one of the weights  $w_{ij}^{(k)}$ , which must be adjusted using the training data.

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#### **Training a Neural Network**

To train the network (i.e., determine the best values for the weights), define a loss function, e.g.,

$$E(\mathbf{w}) = \frac{1}{2} \sum_{i=1}^{N} |t(\mathbf{x}_i) - y_i|^2$$

where w represents the set of all weights, the sum is over the set of training events, and  $y_i$  is the (numeric) true class label of each event (0 or 1).

The optimal values of the weights are found by minimizing E(w) with respect to the weights (non-trivial algorithms: backpropagation, stochastic gradient descent,...).

The desired result for an event with feature vector x is: if the event is of type 0, want  $t(x) \sim 0$ , if the event is of type 1, want  $t(x) \sim 1$ .

## Distribution of neural net output

Degree of separation between classes now much better than with linear decision function:



# Neural network example from LEP II

Signal:  $e^+e^- \rightarrow W^+W^-$  (often 4 well separated hadron jets) Background:  $e^+e^- \rightarrow qqgg$  (4 less well separated hadron jets)



← input variables based on jet
 structure, event shape, ...
 none by itself gives much separation.

#### Neural network output:



(Garrido, Juste and Martinez, ALEPH 96-144)

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# **Deep Neural Networks**

The multilayer perceptron can be generalized to have an arbitrary number of hidden layers, with an arbitrary number of nodes in each (= "network architecture").

A "deep" network has several (or many) hidden layers:



# Other types of classifiers

#### We have seen only two types of classifiers:

Linear (Fisher discriminant) Neural Network

There are many others:

. . .

Support Vector Machine Boosted Decision Tree *K*-Nearest Neighbour

The field is rapidly developing with advances, e.g., that allow one to use feature vectors of very high dimension, such as the pixels of an image.

 $\rightarrow$  face/handwriting recognition, driverless cars...

#### Extra slides

# Some distributions

Distribution/pdf	Example use in Particle Physics
Binomial	Branching ratio
Multinomial	Histogram with fixed $N$
Poisson	Number of events found
Uniform	Monte Carlo method
Exponential	Decay time
Gaussian	Measurement error
Chi-square	Goodness-of-fit
Cauchy	Mass of resonance
Landau	Ionization energy loss
Beta	Prior pdf for efficiency
Gamma	Sum of exponential variables
Student's t	Resolution function with adjustable tails

### **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

$$\overline{n!(N-n)!}$$

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

## 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:

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

# Binomial distribution (3)

Binomial distribution for several values of the parameters:



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

# **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  $\vec{n} = (n_1, \dots, n_m)$ 

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

## 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.

## **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  $\nu = \sigma \int L dt$ .







### **Uniform distribution**



Notation: *x* follows a uniform distribution between  $\alpha$  and  $\beta$  write as:  $x \sim U[\alpha, \beta]$ 

# Uniform distribution (2)

Very often used with  $\alpha = 0$ ,  $\beta = 1$  (e.g., Monte Carlo method).

For any r.v. x with pdf f(x), cumulative distribution F(x), the function y = F(x) is uniform in [0,1]:

### **Exponential distribution**

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



# **Exponential distribution (2)**

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):  $f(t - t_0 | t \ge t_0) = f(t)$ 

#### Question for discussion:

A cosmic ray muon is created 30 km high in the atmosphere, travels to sea level and is stopped in a block of scintillator, giving a start signal at  $t_0$ . At a time t it decays to an electron giving a stop signal. What is distribution of the difference between stop and start times, i.e., the pdf of  $t - t_0$  given  $t > t_0$ ?

#### Gaussian (normal) distribution

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



х

#### Standardized random variables

If a random variable y has pdf f(y) with mean  $\mu$  and std. dev.  $\sigma$ , then the *standardized* variable

$$x = rac{y-\mu}{\sigma}$$
 has the pdf  $g(x) = f(y(x)) \left| rac{dy}{dx} \right| = \sigma f(\mu + \sigma x)$ 

has mean of zero and standard deviation of 1.

Often work with the *standard* Gaussian distribution ( $\mu = 0. \sigma = 1$ ) using notation:

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

Then e.g.  $y = \mu + \sigma x$  follows

$$f(y) = \frac{1}{\sigma}\varphi\left(\frac{y-\mu}{\sigma}\right) = \frac{1}{\sqrt{2\pi}\sigma}e^{-(y-\mu)^2/2\sigma^2}$$

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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,

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

Marginal pdf of each  $x_i$  is Gaussian with mean  $\mu_i$ , standard deviation  $\sigma_i = \sqrt{V_{ii}}$ .

#### **Two-dimensional Gaussian distribution**

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

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



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

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

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.

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#### Cauchy (Breit-Wigner) distribution

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

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

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

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

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

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

$$x_0 = \text{full width at half maximum}$$

Example: mass of resonance particle, e.g.  $\rho$ ,  $K^*$ ,  $\phi^0$ , ...  $\Gamma$  = 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**

 $E[x] = \frac{\alpha}{\alpha + \beta}$ 

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

$$V[x] = \frac{\alpha\beta}{(\alpha + \beta)^2(\alpha + \beta + 1)}$$

Often used to represent pdf of continuous r.v. nonzero only between finite limits, e.g.,  $y = a_0 + a_1 x$ ,  $a_0 \le y \le a_0 + a_1$ 



#### Gamma distribution

$$f(x; \alpha, \beta) = \frac{1}{\Gamma(\alpha)\beta^{\alpha}} x^{\alpha-1} e^{-x/\beta}$$
$$E[x] = \alpha\beta$$
$$V[x] = \alpha\beta^{2}$$

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

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



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.
## Proof of Neyman-Pearson Lemma

Consider a critical region W and suppose the LR satisfies the criterion of the Neyman-Pearson lemma:

 $P(\mathbf{x}|H_1)/P(\mathbf{x}|H_0) \ge c_{\alpha} \text{ for all } \mathbf{x} \text{ in } W,$  $P(\mathbf{x}|H_1)/P(\mathbf{x}|H_0) \le c_{\alpha} \text{ for all } \mathbf{x} \text{ not in } W.$ 

Try to change this into a different critical region W' retaining the same size  $\alpha$ , i.e.,

$$P(\mathbf{x} \in W'|H_0) = P(\mathbf{x} \in W|H_0) = \alpha$$

To do so add a part  $\delta W_+$ , but to keep the size  $\alpha$ , we need to remove a part  $\delta W_-$ , i.e.,

$$W \to W' = W + \delta W_+ - \delta W_-$$

$$P(\mathbf{x} \in \delta W_+ | H_0) = P(\mathbf{x} \in \delta W_- | H_0)$$





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# Proof of Neyman-Pearson Lemma (2)

But we are supposing the LR is higher for all x in  $\delta W_{-}$  removed than for the x in  $\delta W_{+}$  added, and therefore

$$P(\mathbf{x} \in \delta W_+ | H_1) \le P(\mathbf{x} \in \delta W_+ | H_0) c_\alpha$$

$$\frac{\delta W_{+}}{\delta W_{-}}$$

 $P(\mathbf{x} \in \delta W_{-} | H_{1}) \ge P(\mathbf{x} \in \delta W_{-} | H_{0}) c_{\alpha}$ 

The right-hand sides are equal and therefore

 $P(\mathbf{x} \in \delta W_+ | H_1) \le P(\mathbf{x} \in \delta W_- | H_1)$ 

# Proof of Neyman-Pearson Lemma (3)

### We have

$$W \cup W' = W \cup \delta W_+ = W' \cup \delta W_-$$

Note W and  $\delta W_+$  are disjoint, and W' and  $\delta W_-$  are disjoint, so by Kolmogorov's 3<sup>rd</sup> axiom,



$$P(\mathbf{x} \in W') + P(\mathbf{x} \in \delta W_{-}) = P(\mathbf{x} \in W) + P(\mathbf{x} \in \delta W_{+})$$

#### Therefore

$$P(\mathbf{x} \in W'|H_1) = P(\mathbf{x} \in W|H_1) + P(\mathbf{x} \in \delta W_+|H_1) - P(\mathbf{x} \in \delta W_-|H_1)$$

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## Proof of Neyman-Pearson Lemma (4)

And therefore

$$P(\mathbf{x} \in W'|H_1) \le P(\mathbf{x} \in W|H_1)$$

i.e. the deformed critical region W' cannot have higher power than the original one that satisfied the LR criterion of the Neyman-Pearson lemma.