Elements of Statistics
Fundamental Concepts

HDM 2017 and African School of Fundamental Physics and its Applications
Contributing authors : Kadono Marumi, Simon Connell, Ketevi Assamagan

Goal

Describe the fundamental concepts of statistics for HEP

Explore these concepts with Root-Macros for hands-on experience
Using the random number generator … seeing some sampling theory ….
Statistics : quantify knowledge and uncertainty : communicate results

How to interpret the plot below ?
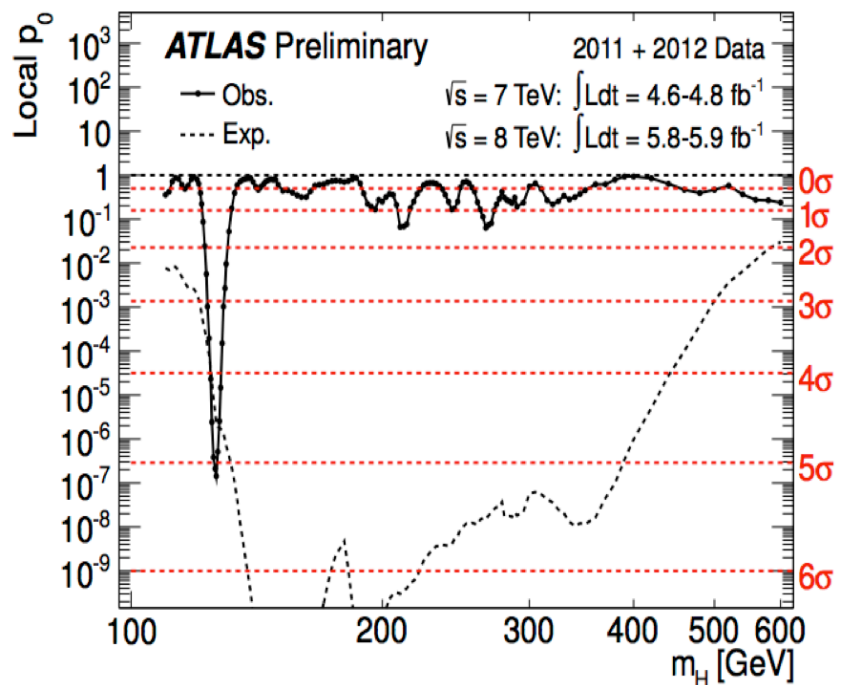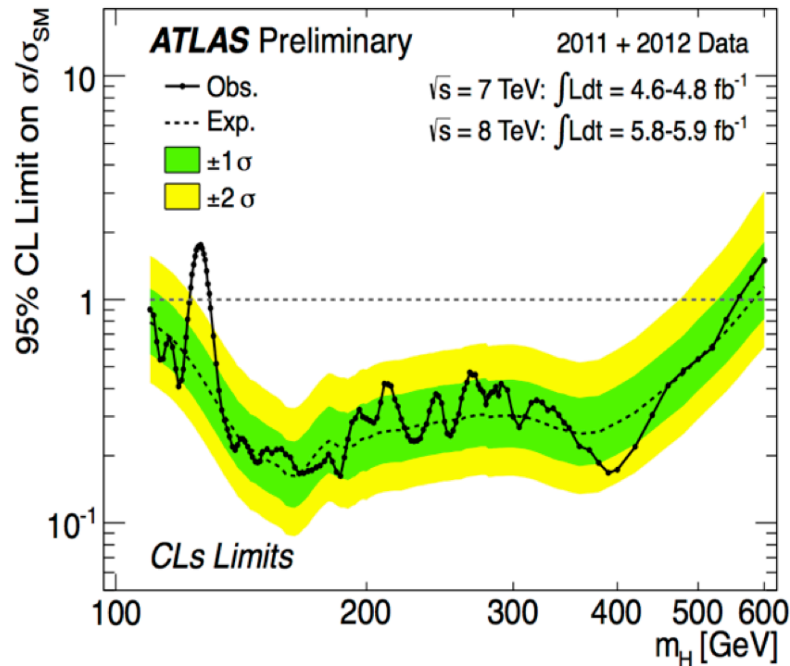
Appreciate there is a lot more for you/us to learn about statistical techniques
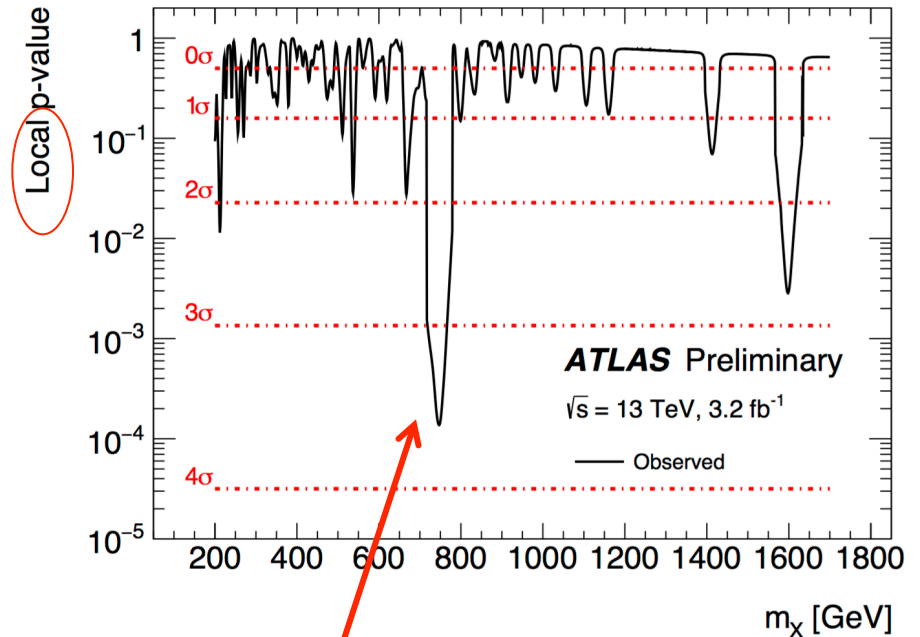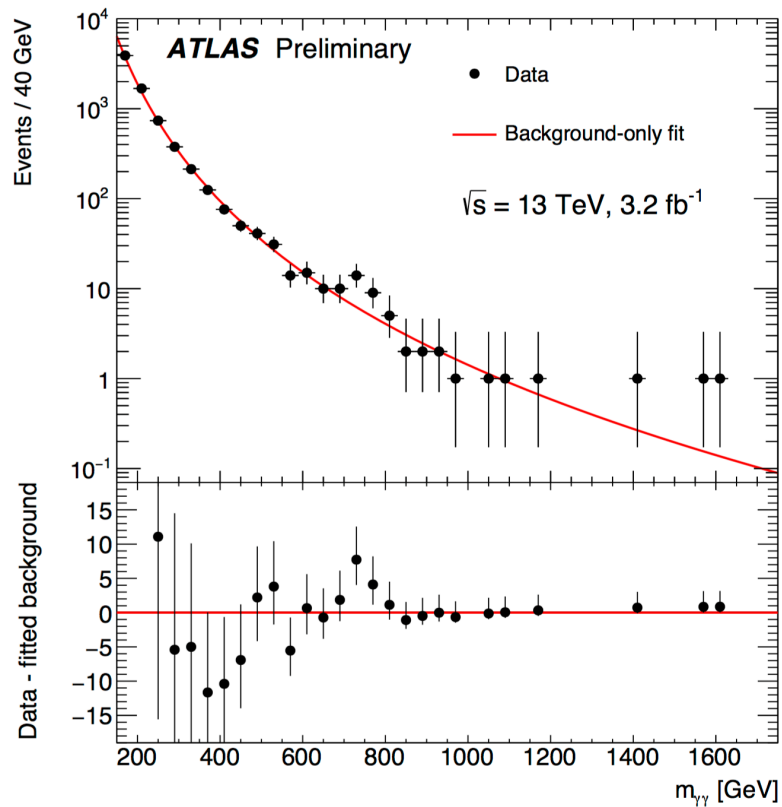
In particular concerning the treatment of systematics

Apply these results to Discovery and Exclusion in HEP

So be patient and take some time to understand the techniques step by step…

Another example

Announced December 2015 : "excess" Peak in the diphoton reconstructed mass spectrum

global p-value 2.0σ

not confirmed with additional
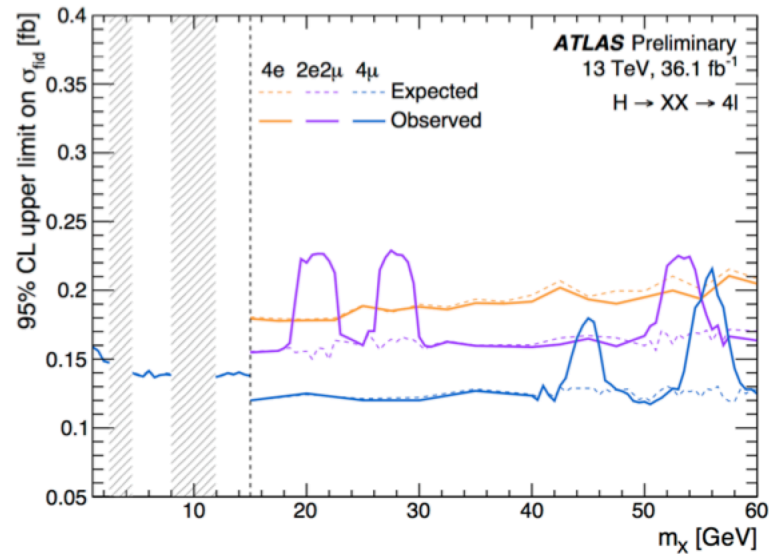data at ICHEP this summer 2016
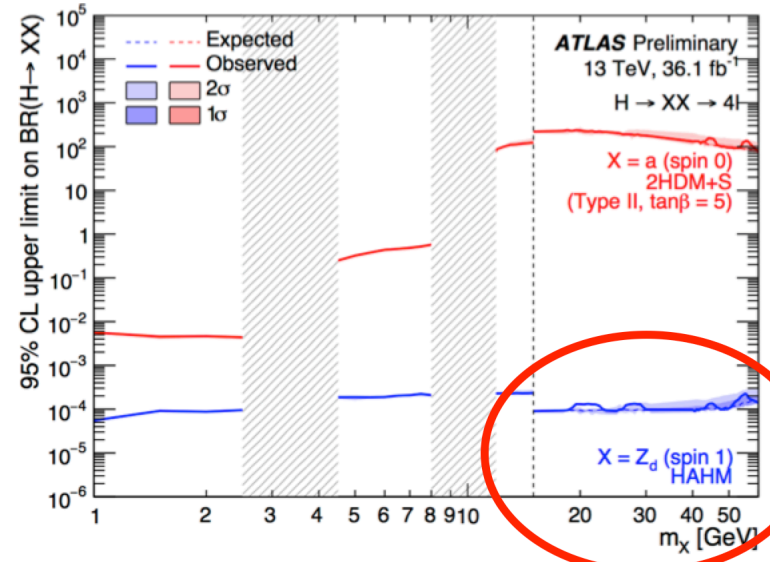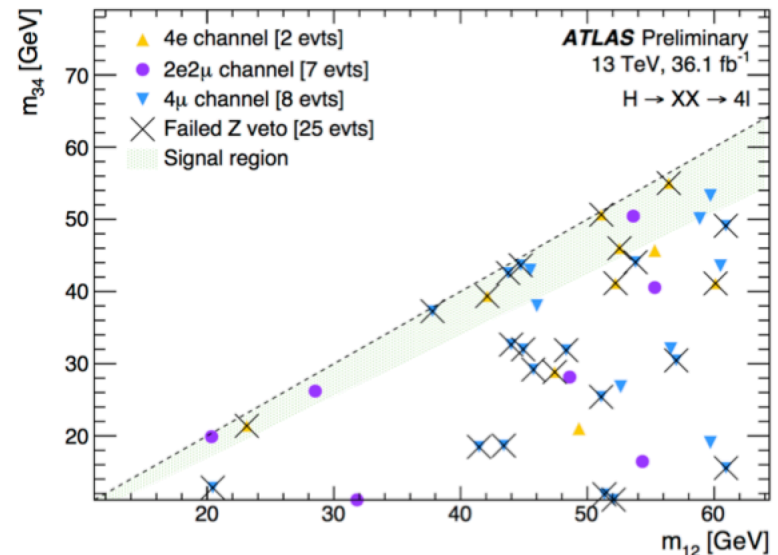
ATLAS CONF Note
ATLAS-CONF-2017-042

Search for Higgs boson decays to Beyond-the-Standard-Model light bosons in four-lepton events with the ATLAS detector at $\sqrt{s} = 13$ TeV

The ATLAS Collaboration

10th July 2017

(a) Fiducial cross sections

(b) Branching Ratios

Further Reading

By physicists, for physicists

G. Cowan, Statistical Data Analysis, Clarendon Press, Oxford, 1998.

R.J.Barlow, A Guide to the Use of Statistical Methods in the Physical Sciences, John Wiley, 1989;

F. James, Statistical Methods in Experimental Physics, 2nd ed., World Scientific, 2006;
  - W.T. Eadie et al., North-Holland, 1971 (1st ed., hard to find);

S.Brandt, Statistical and Computational Methods in Data Analysis, Springer, New York, 1998.

L.Lyons, Statistics for Nuclear and Particle Physics, CUP, 1986.
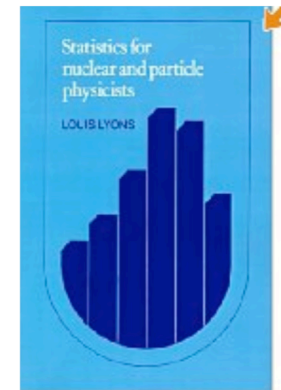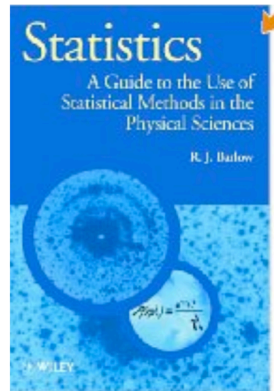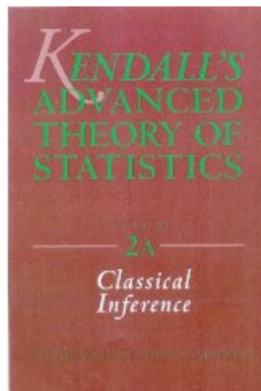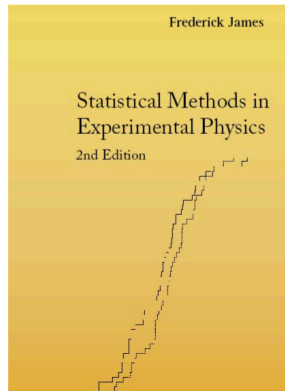
My favorite statistics book by a statistician:

Stuart, Ord, Arnold. "Kendall's Advanced Theory of Statistics" Vol. 2A *Classical Inference & the Linear Model*.
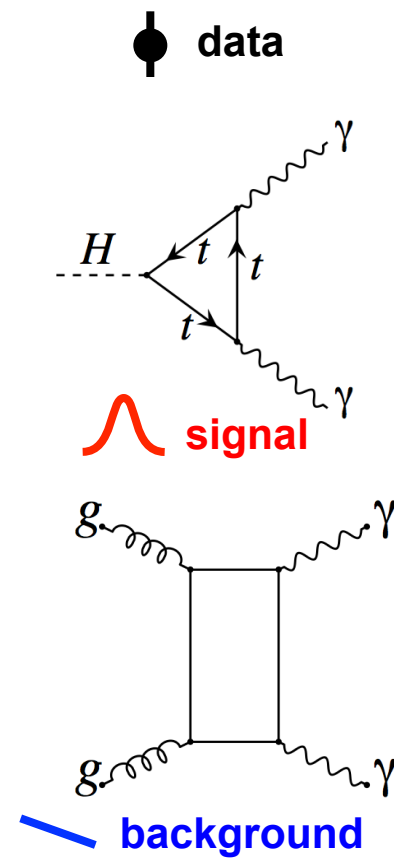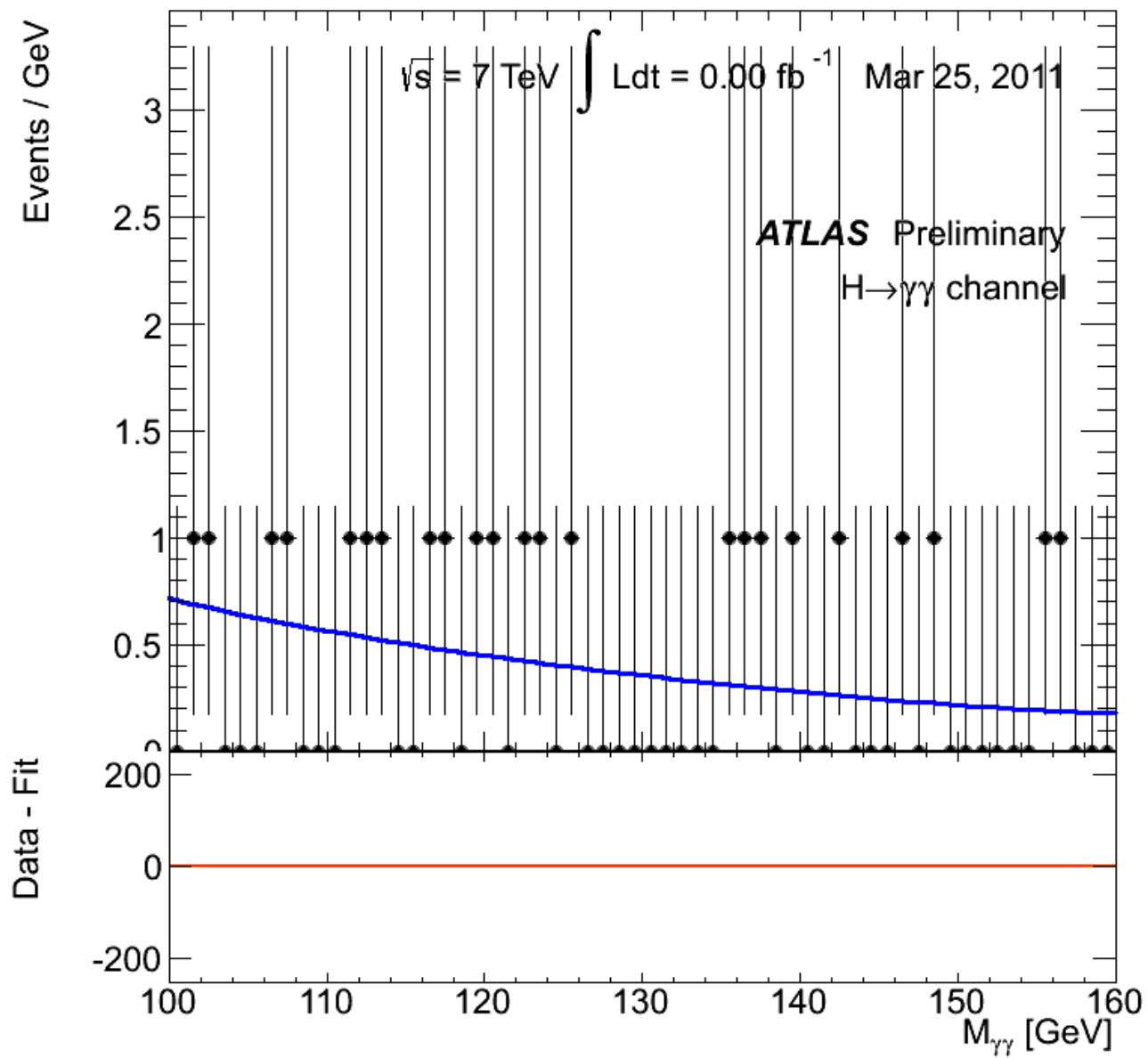
Disclaimer :
What this lecture is not going to be about…

- It will not be a lecture on the fundamental theory of statistics

- Multivariate techniques

- Bayesian confidence intervals

- Goodness of fit theory

- In depth discussion of systematics and their treatment

- Bayesian vs. Frequentist diatribe

Why are Statistics so Important in Particle Physics ?

Because we need to give quantitative statements about processes that have some inherent randomness…

… May this randomness be of measurement nature or quantum …

How did it all start ?

Liber de ludo aleae

To study games of chance !

G. Cardano (1501-1576)

And many others to follow (Pascal, Fermat, etc..)

"La theorie des probabilités n'est, au fond, que le bon sens reduit en calcul"

"The theory of probabilities is at ultimately nothing more than common sense reduced to calculation"

P. S. Laplace (1749-1824)

Example …..

From the very innocuous seeming assumption ….

"There is a random process characterised by a constant average event rate, μ."

… many significant and fundamental results follow – perhaps the prime example of the dramatic yield of results from an assumption in all physics.

radioactive decay

The random deviate represented by the waiting time between such events may be shown to be drawn from the exponential probability density distribution.

$$p_E(t;\mu) = \mu e^{-\mu dt}$$
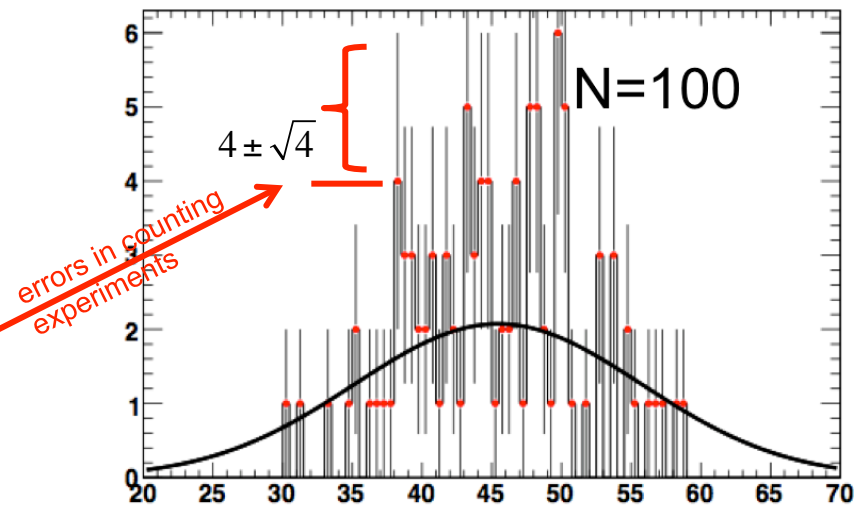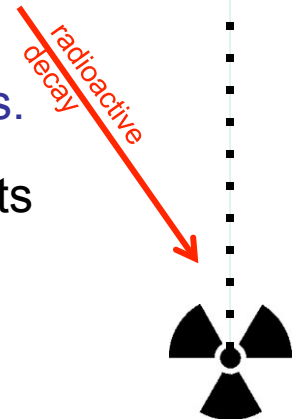
The random deviate represented by the number of such events within a time bin T is drawn from the Binomial Distribution, well approximated by the Poisson Distribution.

$$p_P(n;\bar{n}) = \frac{\bar{n}^n}{n!} e^{-\bar{n}}$$

$4 \pm \sqrt{4}$

N=100

errors in counting experiments

Where the expectation value $\bar{n} = T\mu$

and std. deviation $\sigma = \sqrt{\bar{n}} \approx \sqrt{n}$

ROOT

Data Analysis Framework

Google Custom Search

Download   **Documentation**   **News**   **Support**   **About**   **Development**   **Contribute**

Getting Started   **Reference Guide**   **Forum**   **Gallery**

ROOT is ...

A modular scientific software framework. It provides all the functionalities needed to deal with big data processing, statistical analysis, visualisation and storage. It is mainly written in C++ but integrated with other languages such as Python and R.

Try it in your browser! (Beta)

Download   or **Read More ...**

```
import ROOT

cppFunctionCode = '''
void f() {
    std::cout << "Hi jitted C++ world!" << std::endl;
}
'''

ROOT.gInterpreter.Declare(cppFunctionCode)

ROOT.f() # Hello!
```

Previous Pause Next

Under the Spotlight

03-08-2017 **The ROOT Docker container (alpha version)**
Do you like Docker? Would you like to use ROOT? We provide an *alpha* version of the ROOT Docker container!

08-03-2017 **Development release 6.09/02 is out!**
This is the first ROOT development release of the 6.09 series! It is meant to offer a preview of the many features which will be included in the 6.10 production release.

05-09-2016 **Get the most out of the ROOT tutorials!**
All ROOT tutorials are now available as ROOTBooks which can be statically visualized via NBViewer or interactively explored with SWAN.

06-07-2016 **CERN Summer Students' Course**
The CERN Summer Student program is in full swing and ROOT is part of it.

Other News

16-04-2016 The status of reflection in C++

05-01-2016 Wanted: A tool to 'warn' user of inefficient (for I/O) construct in data model

03-12-2015 ROOT::TSeq::GetSize() or ROOT::seq::size()?

02-09-2015 Wanted: Storage of HEP data via key/value storage solutions

Latest Releases

Release 6.10/08 - 2017-10-16

Release 6.11/02 - 2017-10-06

Release 6.10/06 - 2017-09-19

Release 6.10/04 - 2017-07-28

To run the C++ ROOT Code snippets

- Install ROOT ... See previous page for instructions ... See google if you get stuck

- Install an editor, like notepad++ for Windows, atom for a Mac, gedit for Linux.

- Have access to a shell ... eg command prompt in windows, terminal in a Mac or Linux.

- Do Chapters 2,3,4 in the ROOT User Guide as a self taught tutorial.

- Put the code snippets into a folder, go there in your shell, say

  `root`

  to get to the ROOT environment.

- From here

  `.x <code snipet file name>`

- You can modify things like the number of events etc ... and you can teach yourself both stats and coding by following and modifying the code

What is a Statistical Error ?

Imagine I have a billion white ○ and blue ● golf balls

I decide to throw one million of them into a well with an admixture of 15 out of one hundred blue ones…

I then know PRECISELY the probability that if you pick one at RANDOM, it will be blue…

$$p = 15\%$$

You of course don't know this number and you want to measure it…

All you have is a bucket…

Which contains exactly 300 balls

This is approximately how the well looks like inside…
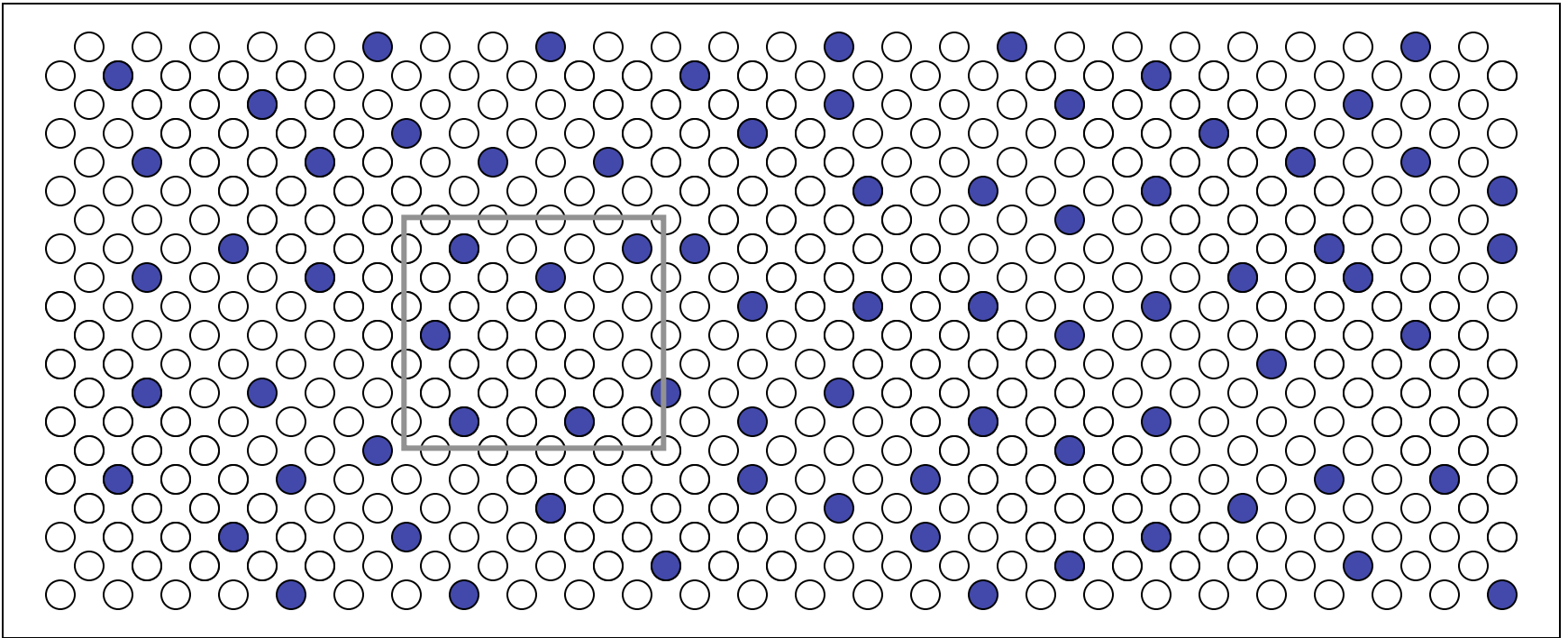
You throw the bucket and pull out the following outcome
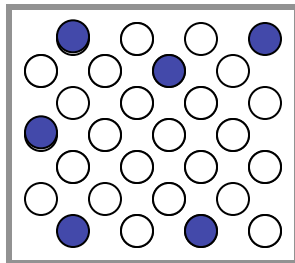
Aha! You have a measurement!

$n = 300$

$k = 36$

The probability is…

$p = 12\%$

… But how precise is it ?

13

Remember you are not supposed to know the true value!

The difference between a measurement and the true value is the Statistical Error

In this case it would be 3% absolute (20% relative), but since you don't know the true value you don't know at all what your statistical error really is !

Of course had you thrown your bucket on a different spot, you would have probably had a different measurement and the statistical error would be different…

What you want to know is your measurement error, or what the average statistical variation of your measurement is…

This can be done provided that you know the law of probability governing the possible outcomes of your experiment …   (and the true value of p, but assume that 12% is a close enough)

You want to know what the probability for an outcome of k golf balls to be blue is.

For one specific outcome the probability is:

$$P = p^k \times (1 - p)^{n-k}$$

What are all possible combination of outcomes of k blue balls out of n?

What are all possible combination of outcomes of k blue balls out of n?

For the first blue ball there are n choices, once this choice is made the second ball has n-1 choices,… the kth ball has (n-k) choices.

In a simple case… n=10 and k=3 this can be seen as:

The first ball has 3 choices The second ball has 2 choices

So the number of combinations is : $n \times (n-1) \times (n-2)$

In the general case : $n \times (n-1) \times (n-2) \times (n-3)... \times (n-k+1) = \dfrac{n!}{(n-k)!}$

Because we do not care about the order in which we have picked the balls

… avoid the double counting!

1	2	3
1	3	2
2	1	3
3	1	2
2	3	1
3	2	1

Each configuration is counted 6 times

This number corresponds in fact to the number of combinations of k blue balls out of k balls and therefore :

$$k \times (k-1) \times (k-2) \times (k-3)...\times 1 = k!$$

Aka the number of re-arrangements of the k blue balls.

In order to account for each combination only once you just need to divide by the number of re-arrangements of the k blue balls.

So the number of combinations of k elements among n is given by :

$$C_n^k = \frac{n!}{k!(n-k)!}$$

The probability to pick k blue balls among n, given a probability P that the a ball is blue is thus :

$$P = C_n^k \times p^k \times (1-p)^{n-k}$$

This is an absolutely fundamental formula in probability and statistics!
It is the so called Binomial Probability!

The Binomial Probability

Binomial coefficients were known since more than a thousand years…

… they were also the foundation of modern probability theory!

B. Pascal (1623-1662)
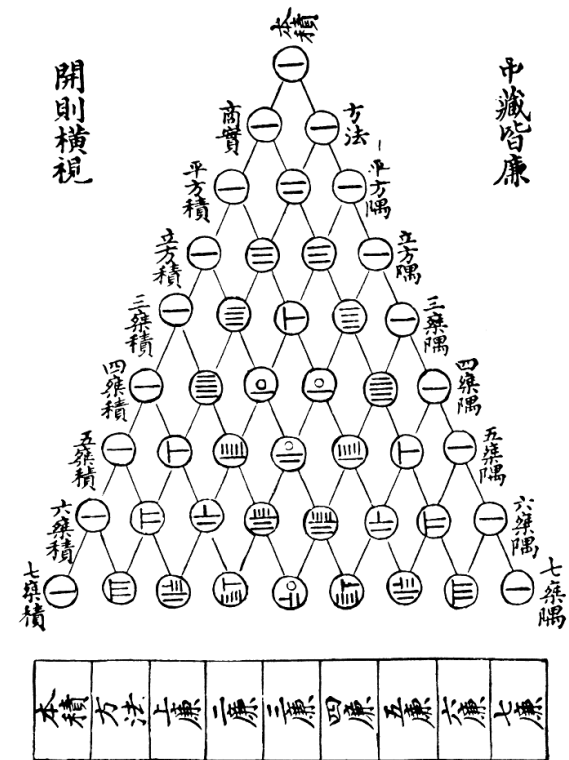
The Pascal Triangle (~1000 AD)

So what is the precision of your measurement ?

A good measure of the precision (not the accuracy) is the *Root Mean Square Deviation* (square root of the variance) of possible outcomes of the measurement.

You will compute it yourself. To do so you need two steps…
(see next slide for the full derivation)

Step 1 : Compute the mean value of the binomial probability

$$\mu = nP$$

Step 2 : Compute the variance of the binomial probability

$$Variance = nP(1 - P)$$

So now you know the variance of your distribution for a given probability P…

In your case : $P = 12\%$ Assuming P is close enough to the true value, the precision is :

$$RMSD = \sqrt{nP(1 - P)} = 5.6$$

The relative precision ~15% is rather poor and the accuracy questionable! (Remember, your statistical error is 45 - 36 = 9, although you are not supposed to know it !)

Step 1 : Compute mean value   Step 2 : Compute variance

Mean of the Binomial Probability

Let us denote by $\mathcal{P}_k^n(p)$ the binomial probability of an outcome k among n with single event probability p, and μ its mean value, then

$$\mu = \frac{\sum_{k=0}^{n} k\, \mathcal{P}_k^n(p)}{\underbrace{\sum_{k=0}^{n} \mathcal{P}_k^n(p)}_{=1}} = \sum_{k=0}^{n} k\, C_n^k\, p^k (1-p)^{n-k}$$

Note that it looks like a derivative – !

Nice trick :   Start from the derivative …

$$\frac{\partial}{\partial p}\left[\sum_{k=0}^{n} C_n^k\, p^k (1-p)^{n-k}\right] = 0 = \frac{\partial [1]}{\partial p}$$

so $$\underbrace{\sum_{k=0}^{n} k\, C_n^k\, p^{k-1}(1-p)^{n-k}}_{\mu/p} - \underbrace{\sum_{k=0}^{n}(n-k)\, C_n^k\, p^k (1-p)^{n-k-1}}_{} = 0$$

$$0 = \frac{\mu}{p} - n\underbrace{\sum_{k=0}^{n} C_n^k\, p^k (1-p)^{n-k-1}}_{1/(1-p)} + \underbrace{\sum_{k=0}^{n} k\, C_n^k\, p^k (1-p)^{n-k-1}}_{\mu/(1-p)}$$

thus

$$\frac{\mu}{p} + \frac{\mu}{1-p} = \frac{n}{1-p} \implies (1-p)\mu + p\mu = np$$

so $\boxed{\mu = np}$ Aha !

Variance of the Binomial Probability

Let us start from the previous formula for the mean value of the binomial probability.

$$\text{Variance} = \sum_{k=0}^{n} k^2\, C_n^k\, p^k (1-p)^{n-k} - \underbrace{\left(\sum k\, C_n^k\, p^k (1-p)^{n-k}\right)}_{\mu^2}$$

given that $\sum_{k=0}^{n} k\, C_n^k\, p^k (1-p)^{n-k} = \mu = np$

then $\frac{\partial}{\partial p}\left[\sum_{k=0}^{n} k\, C_n^k\, p^k (1-p)^{n-k}\right] = n$ thus :

$$\sum_{k=0}^{n} k^2\, C_n^k\, p^{k-1}(1-p)^{n-k} - \sum_{k=0}^{n} k(n-k)\, C_n^k\, p^k (1-p)^{n-k-1} = n$$

$$\frac{1}{p}\sum_{k=0}^{n} k^2\, C_n^k\, p^k (1-p)^{n-k} - \frac{n\mu}{1-p} + \frac{1}{1-p}\sum k^2\, C_n^k\, p^k (1-p)^{n-k} = n$$

so $$(1-p+p)\sum_{k=0}^{n} k^2\, C_n^k\, p^k (1-p)^{n-k} = np(1-p) + \underbrace{np\,\mu}_{\mu^2}$$

when $\mu = np$ so :

$$\underbrace{\sum k^2\, C_n^k\, p^k (1-p)^{n-k} - \mu^2}_{\text{Variance !}} = np(1-p)$$

Thus $\boxed{\text{Variance} = np(1-p)}$

But wait…

Now you are curious to see what happens if you repeat your measurement!

You have noticed that the average binomial probability is the expected value!

Intuitively you will therefore try to repeat and average your measurements…

You will do it 50,000 times and meticulously plot the number of counts. This is what you get :

$N_{throws} = 50000$
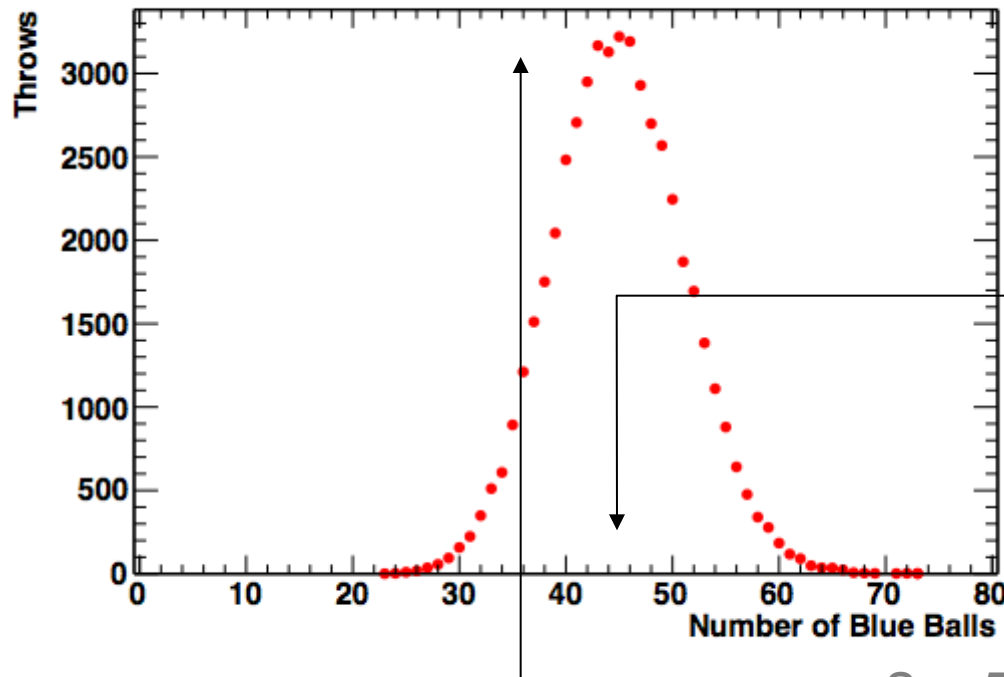
The average number of blue balls in 50,000 throws :

$$\langle Number_{Blue} \rangle = 44.98$$

$$\langle P \rangle = 14.99\%$$

See Binomial.C

Your initial measurement (36) !

Now you decide that your measurement is the average, what is its precision ?

What is the variance of the average ?

Let's start from one straightforward and general property of the Variance for two random variables X and Y :

$$Var\left(aX + bY\right) = \left\langle (aX + bY - \langle aX + bY \rangle)^2 \right\rangle = \left\langle \left[a(X - \langle X \rangle) + b(Y - \langle Y \rangle) \right]^2 \right\rangle$$

$$= a^2 Var(X) + b^2 Var(Y) + 2ab Cov(X,Y)$$

Where the covariance is :
$$Cov(X,Y) = \left\langle (X - \langle X \rangle)(Y - \langle Y \rangle) \right\rangle$$

This formula generalizes to...
$$Var\left(\sum_{i=0}^{n} a_i X_i \right) = \sum_{i=0}^{n} a_i^2 Var(X_i) + \sum_{0 \le i < j \le n} a_i a_j Cov(X_i, X_j)$$

Therefore assuming that each of the bucket throws measurement N_{Blue}^{k} is independent from the previous one, the mean value being a simple sum of the measurements divided by the number of throws :

$$\left\langle Number_{Blue} \right\rangle = \frac{1}{N_{Throws}} \sum_{k=1}^{N_{Throws}} N_{Blue}^{k}$$

The ensemble variance then is :

$$\hat{\sigma}^2 = Var\left(\frac{1}{N_{Throws}} \sum_{k=1}^{N_{Throws}} X_i \right) = \frac{1}{N_{Throws}^2} \sum_{k=1}^{N_{Throws}} Var(X_i) = \frac{1}{N_{Throws}^2} N_{Throws} Var(X_i) = \frac{nP(1-P)}{N_{Throws}}$$

The precision being given by the *Root Mean Square Deviation* :

$$RMSD = \sqrt{\frac{nP(1-P)}{N_{Throws}}} = \frac{RMSD_{Individual}}{\sqrt{N_{Throws}}} = 0.01\%$$

Very interesting behavior : Although you do not know the true value p, you see that the average is converging towards it with increasing precision!

The line here is the true value !

Your initial measurement

See Binomial.C
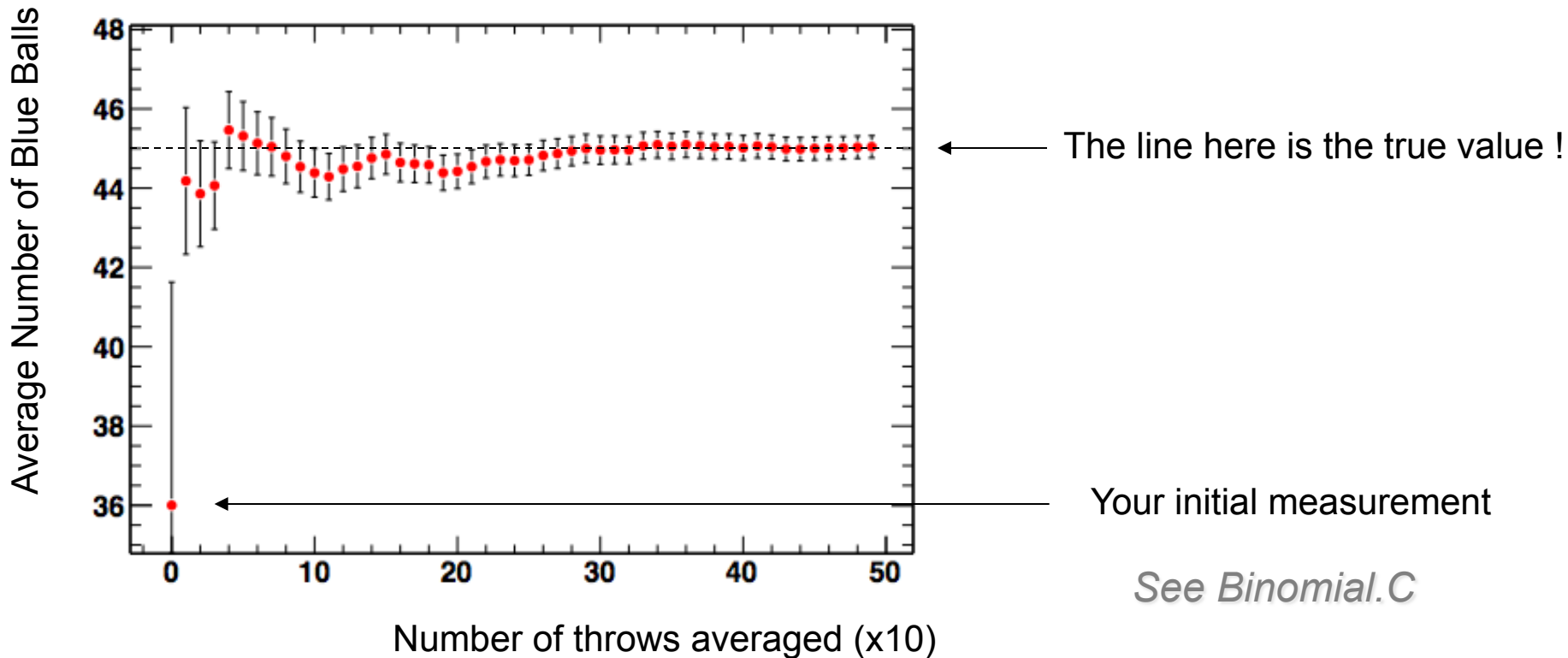
This is an illustration of the *LAW* of *LARGE NUMBERS* ! Extremely important, intuitive but not trivial to demonstrate…

What is the meaning of our first measurement N_{blue} = 36 ?

Now that we know (after 50,000 throws) to a high precision that the probability of a blue ball is very close to 15%.

The frequency of an outcome as low as 12% is ~10% (not so unlikely!)

What difference would it make if you had known true value ?

Frequency at which the measurement is within the precision as estimated from the truth :

$$\left|P_{meas} - p\right| \le \sqrt{np(1-p)} \qquad \Rightarrow 70\% \text{ (of the cases the measurement is within the true statistical RMSD)}$$

Frequency at which the true value is within the precision as estimated from the measurement :

$$\left|P_{meas} - p\right| \le \sqrt{nP_{Meas}(1 - P_{Meas})} \qquad \Rightarrow 67\% \text{ (of the cases the true value is within the measured error)}$$

See Coverage.C

The true value coverage is similar in the two cases, keep these values in mind…

Here all results are derived from a simulation in terms of frequencies…
Computing Binomial probabilities with large numbers of N can be quite difficult !

The Gaussian or Normal Probability

Is there a way to simplify the computation ? Not so trivial to compute 300! directly…

A very nice approximation of the Binomial Probability can be achieved using Stirling's Formula !

$$n! \approx \sqrt{2\pi n}\left(\frac{n}{e}\right)^n \qquad \ln n! \approx n \ln n - n + \tfrac{1}{2}\ln 2\pi n$$
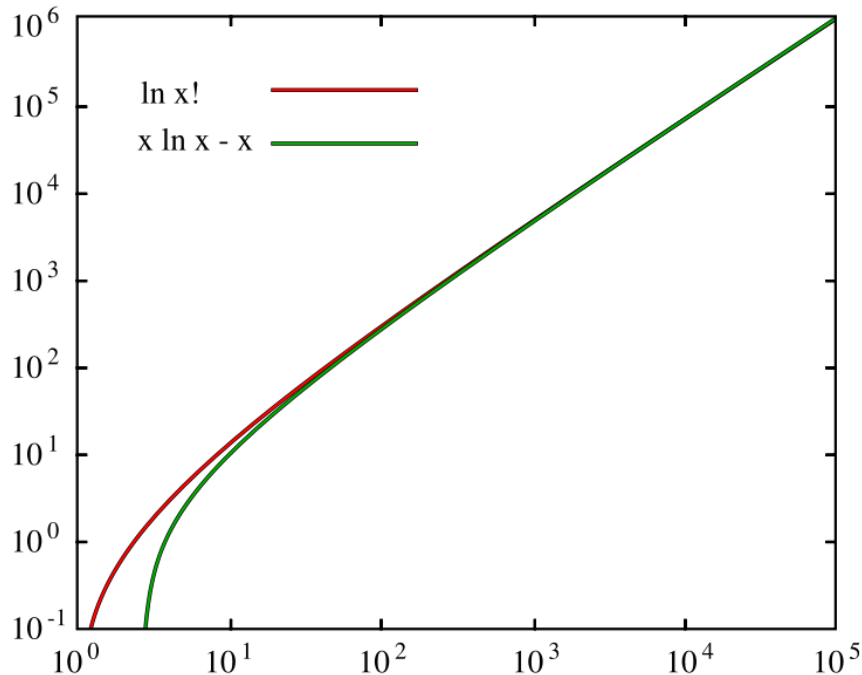
Formula is valid for large values of n…

$$C_n^k p^k (1-p)^{n-k} \approx \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(k-\langle k \rangle)^2}{2\sigma^2}}$$

$$\sigma = \sqrt{np(1-p)}$$

(See derivation in the next slide)

Binomial convergence towards Normal

Derivation of the Normal Probability

... Again start from the binomial probability, but this time use the Stirling formula: $n! \sim \sqrt{2\pi n}\left(\frac{n}{e}\right)^n$

$$\ln(k!) \sim k \ln k - k + \frac{1}{2}\ln(2\pi k)$$

— Very useful !..

then $\ln C_n^k = \ln n! - \ln k! - \ln(n-k)!$

thus
$$\ln\left[C_n^k p^k (1-p)^{n-k}\right] = n \ln n - n + \frac{1}{2}\ln(2\pi n)$$
$$-\left[k \ln k - k + \frac{1}{2}\ln(2\pi k)\right] - \left[(n-k)\ln(n-k) - n+k + \frac{1}{2}\ln(2\pi(n-k))\right]$$
$$+ k \ln p + (n-k)\ln(1-p)$$

Again using a derivative trick
$$\frac{\partial}{\partial k}\left[\ln\left(C_n^k p^k (1-p)^{n-k}\right)\right] = \underbrace{-\ln k + \ln(n-k) + \ln p - \ln(1-p)}_{\ln\left(\frac{p(n-k)}{k(1-p)}\right)}$$
$$\underbrace{-\frac{1}{2k} + \frac{1}{2(n-k)}}_{O\left(\frac{1}{n}\right)}$$

We see that the maximum value will occur at
$$p(n-k) = k(1-p) \quad \text{or} \quad \boxed{k = np} \quad \text{Which is}$$
also the average value.

Then looking at the second derivative:
$$\frac{\partial^2}{\partial k^2}\left[\ln\left(C_n^k p^k (1-p)^{n-k}\right)\right] = -\left(\frac{1}{n-k} + \frac{1}{k}\right) + O\left(\frac{1}{k^2}\right)$$

then taken at maximum value $k = np$:
$$\frac{\partial^2}{\partial k^2}\left[\ln\left(C_n^k p^k (1-p)^{n-k}\right)\right] \sim \frac{-1}{np(1-p)} = -\frac{1}{\sigma^2}$$
where $\sigma = \sqrt{np(1-p)}$

Therefore with a second order Taylor series expansion:
$$\ln\left(C_n^k p^k (1-p)^{n-k}\right) \sim \underbrace{\ln A}_{\text{constant}} + \frac{(k-np)^2}{2}\cdot\left(\frac{-1}{\sigma^2}\right)$$

therefore $\quad C_n^k p^k (1-p)^{n-k} \sim A\, e^{-\frac{(k-np)^2}{2\sigma^2}}$

introducing the notation $\tilde{k} = np$ corresponding to the maximum of the probability - then
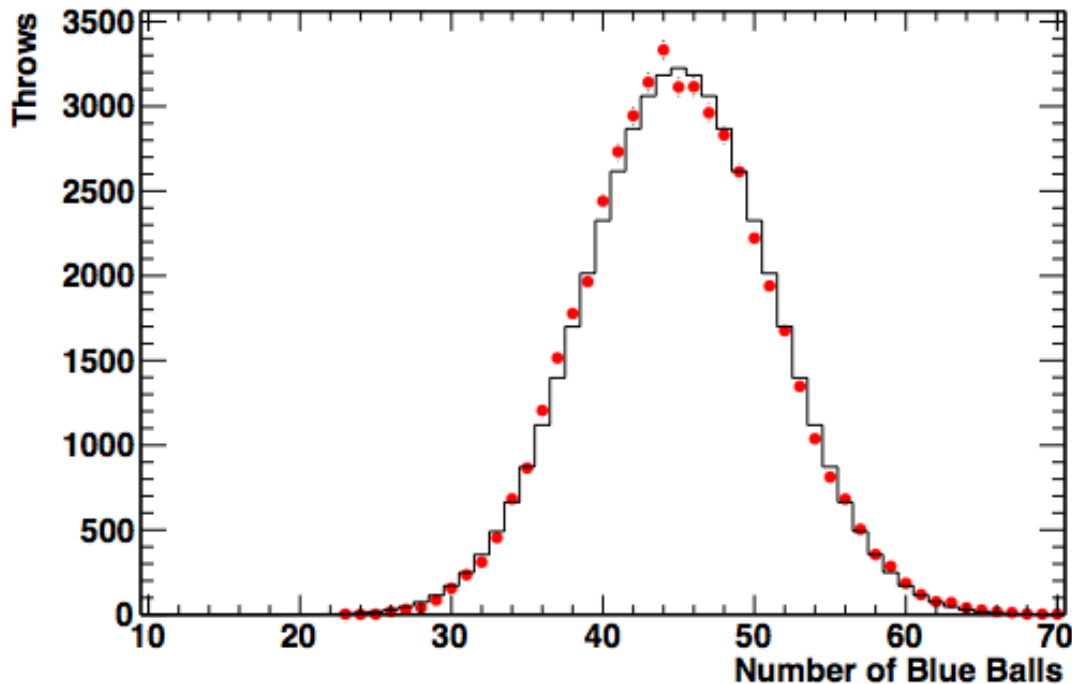$$C_n^k p^k (1-p)^{n-k} \sim A\, e^{-\frac{(k-\tilde{k})^2}{2\sigma^2}}$$

then using the normalization of the probability
$$\int_{-\infty}^{\infty} A\, e^{-\frac{(k-\tilde{k})^2}{2\sigma^2}}\, dk = 1 \quad \text{and the Gauss integral}$$
$$\int_{-\infty}^{+\infty} e^{-\frac{x^2}{2\sigma^2}} = \sqrt{2\pi\sigma^2} \quad \text{so:}$$
$$C_n^k p^k (1-p)^{n-k} = \frac{e^{-\frac{(k-\tilde{k})^2}{2\sigma^2}}}{\sqrt{2\pi\sigma^2}}$$

Validity of the Normal Convergence (Approximation)

Does the approximation apply to our bucket experiment (n=300 and p=15%) ?

See NormalConvergence.C



Not bad (although not perfect) !

C. F. Gauss (1777-1855)

In practice you can use the normal law when approximately n>30 and np>5

What is so "Normal" About the Gaussian?
The Central Limit Theorem…   … at Work !

When averaging various independent random variables (and identically distributed) the distribution of the average converges towards a Gaussian distribution (100 000 events, 100 bins)

See CLT.C

N = 10
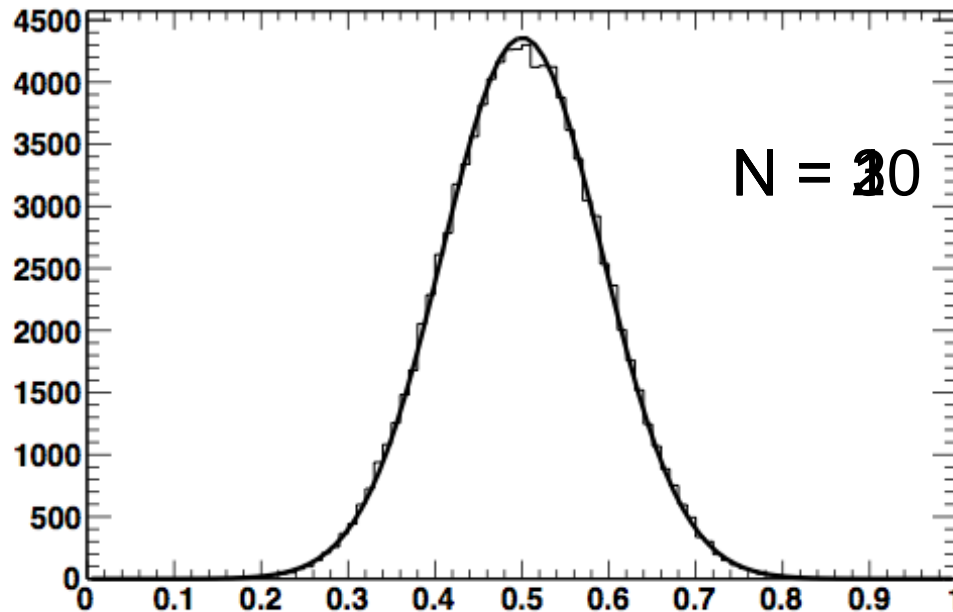
$$\hat{\sigma}^2 = \frac{nP(1-P)}{N_{Throws}}$$

$$RMS = \frac{[0,1]}{\sqrt{12}} \times \frac{1}{\sqrt{10}}$$

At N=10 an excellent agreement with a gaussian distribution is observed

The CLT is one of the main reasons for the great success of the Gaussian law…

On the one hand the CLT is very powerful to describe all those phenomena that result from the superposition of various other phenomena… but on the other hand it is just a limit…
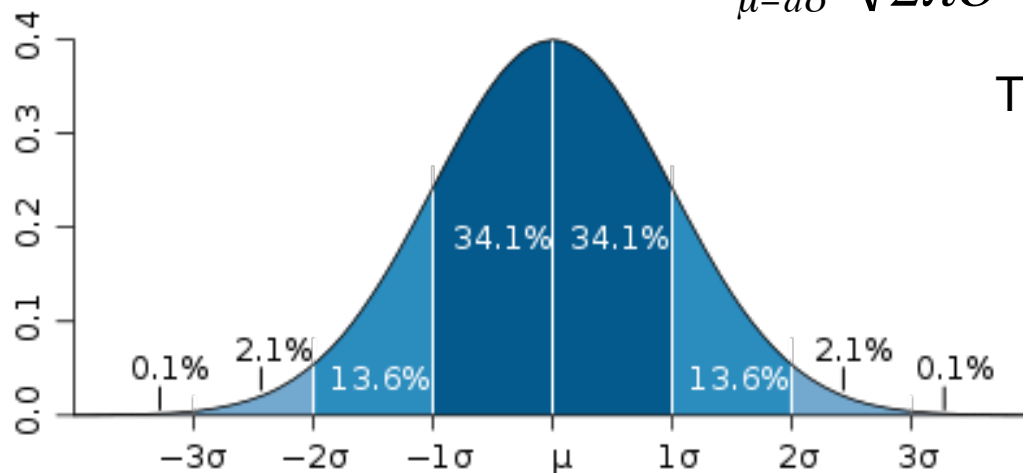
The Notion of Standard Error

Starting from the gaussian PDF :

$$G_{PDF}(x,\mu,\sigma) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

Let's give a first definition of a central confidence interval as the deviation from the central value…

$$P(a\sigma) = \int_{\mu-a\sigma}^{\mu+a\sigma} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}} dx$$

Then for : - a = 1 : P(aσ) = 68.3%

- a = 2 : P(aσ) = 95.4%

- a = 3 : P(aσ) = 99.7 %

See NormalCoverage.C

If you knew the true value of the "error" (σ) then you could say that the in the gaussian limit that the true value has 68.3% probability to be within the 1s, but in many practical examples (such as the well) the true value of the error is not known…

How does the Bucket Experiment Relate to Particle Physics?

The bucket experiment is the measurement of an abundance (blue balls)…

This is precisely what we call in particle physics cross sections…

… except that the bucket contains all collisions collected in an experiment so…

- We try to fill it as much as possible (N is very large and not constant!)

- The processes we are looking for are very rare (p is very small)

The very large N makes it difficult to compute the binomial probability…
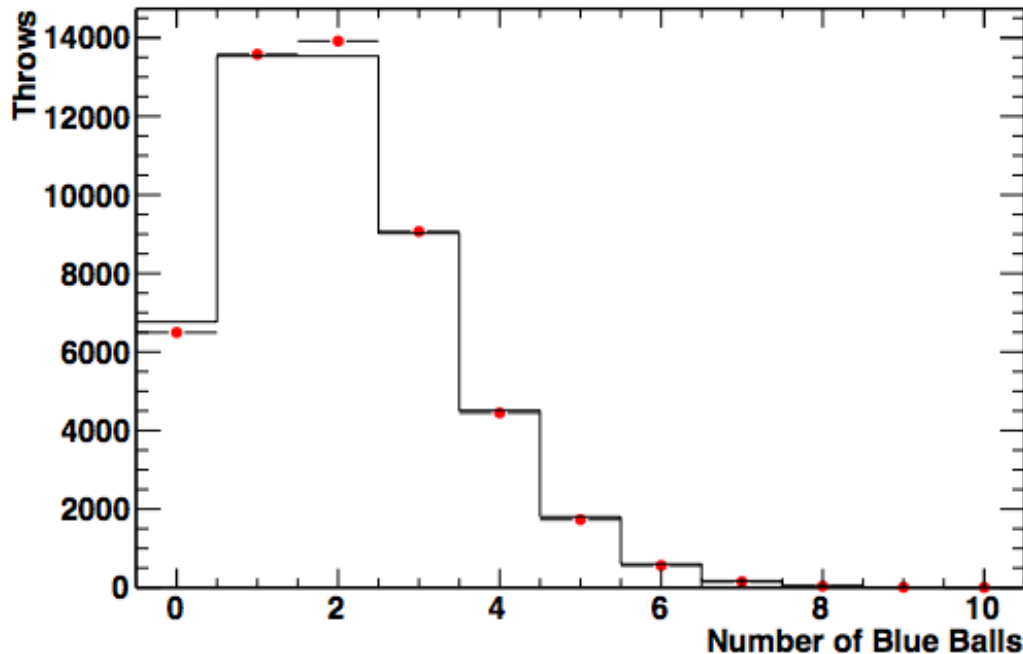
The Poisson Probability

In the large n and small p limit and assuming that $np = \mu$ is finite you can show (see next slide) that …

$$C_n^k p^k (1-p)^{n-k} \approx \frac{(np)^k}{k!} e^{-(np)} = \frac{\mu^k}{k!} e^{-\mu}$$

Much simpler formulation! In practice you can use the normal law when approximately $n>30$ and $np<5$

See PoissonConvergence.C

N=100 and p=26%

S. D. Poisson (1781-1840)

Interesting to note that Poisson developed his theory trying not to solve a game of chance problem but a question of Social Science !

Derivation of the Poisson Probability

... from the binomial probability : $C_n^k \, p^k \, (1-p)^{n-k}$ which can be written in the limit when p is very small and np is finite.

$$C_n^k = \frac{n!}{k!\,(n-k)!} = \frac{1}{k!}\left[n\,(n-1)\ldots(n-k+1)\right] \sim \frac{n^k}{k!}$$

and for p small $(1-p)^{n-k} = 1 - (n-k)p + \frac{(n-k)(n-k-1)}{2!}\,p^2$

$$+ \ldots \simeq 1 - np + \frac{(pn)^2}{2!} + \ldots$$

$$\approx e^{-pn}$$

therefore $C_n^k\,p^k\,(1-p)^{n-k} \approx \frac{n^k}{k!}\,p^k\,e^{-np}$

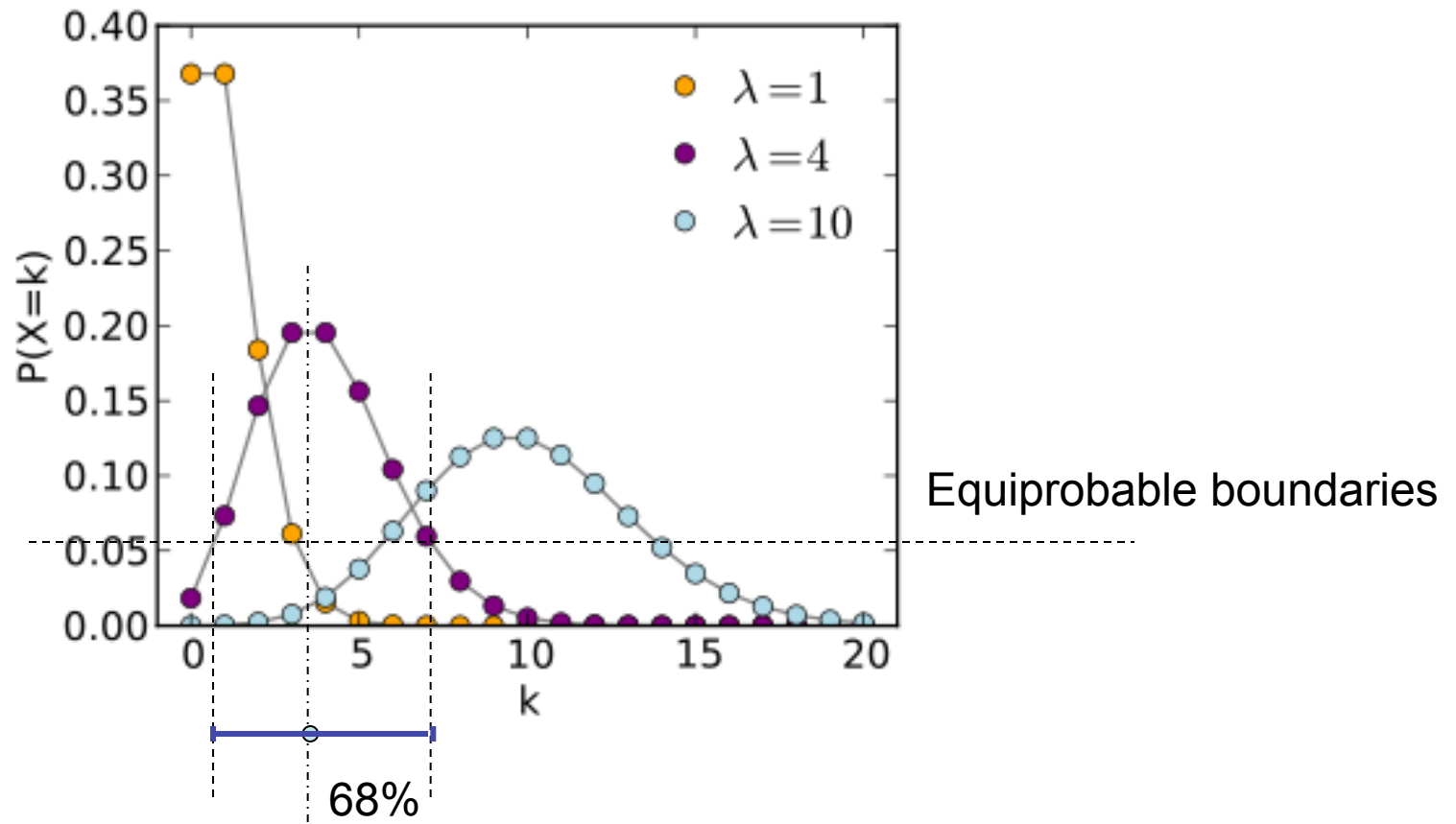$$= \frac{(np)^k}{k!}\,e^{-np}$$

thus denoting $\mu = np$ the poisson probability of an outcome k given an expectation of μ is

$$\boxed{P(k,\mu) = \frac{\mu^k}{k}\,e^{-\mu}}$$

This approximation works essentially when $(n > 50$ and $p < 0.1)$.

Poisson Intervals (or Errors)

Now how will you define a central confidence interval in a non symmetric case ?

Equiprobable boundaries

68%

The integration needs to start from the most probable value downwards…

Here is our first encounter with the necessity of an ordering !

What have we learned ?

…and a few by-products…

1.- Repeating measurements allows to converge towards the true value of an observable more and more precisely …

But never reach it with infinite precision !!!

Even more so accounting for systematics…
(what if the balls do not have an homogeneous distribution ?)

2.- Binomial variance is also useful to compute the so-called binomial error, mostly used for efficiencies :

$$\sigma_\varepsilon = \frac{\sigma_\mu}{N} = \sqrt{\frac{\varepsilon(1-\varepsilon)}{N}} \qquad \mu = np$$

For an efficiency you must consider n fixed !

3.- We came across a very important formula in the previous slides

$$Var\left(\sum_{i=0}^{n} a_i X_i\right) = \sum_{i=0}^{n} a_i^2 Var(X_i) + \sum_{0 \le i < j \le n} a_i a_j Cov(X_i, X_j)$$

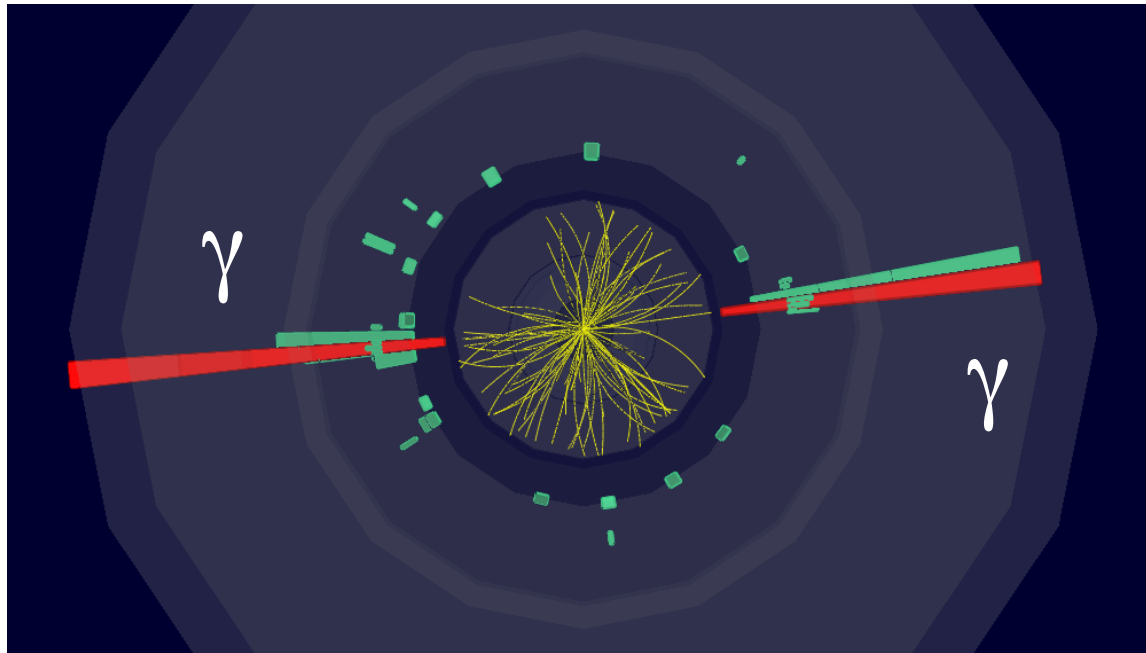That generalizes (with a simple Taylor expansion) to…

$$var(f(x_1,...,x_n)) = \sum_{i=0}^{n} (\frac{\partial f}{\partial x_i})^2 var(x_i) + \sum_{0 \le i < j \le n} \frac{\partial f}{\partial x_i} \frac{\partial f}{\partial x_j} cov(x_i, x_j)$$
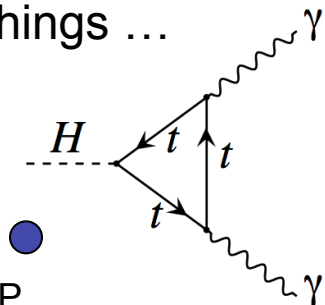
Likelihood

Unfortunately in High Energy physics experiments, events (balls) don't come in single colors (white or blue) … Their properties are not as distinct !
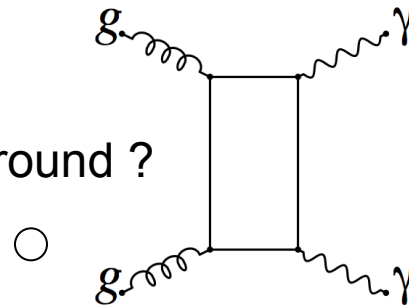
For instance take this simple event :

Could be many things …

Higgs ?

Background ?

Let alone that they can be indistinguishable (quantum interference)

How can we distinguish between the two ?

Very vast question, let's first start with how to measure their properties

(Which is also a very vast question!)

One clear distinctive feature is that the signal is a narrow mass resonance, while the background is a continuum !
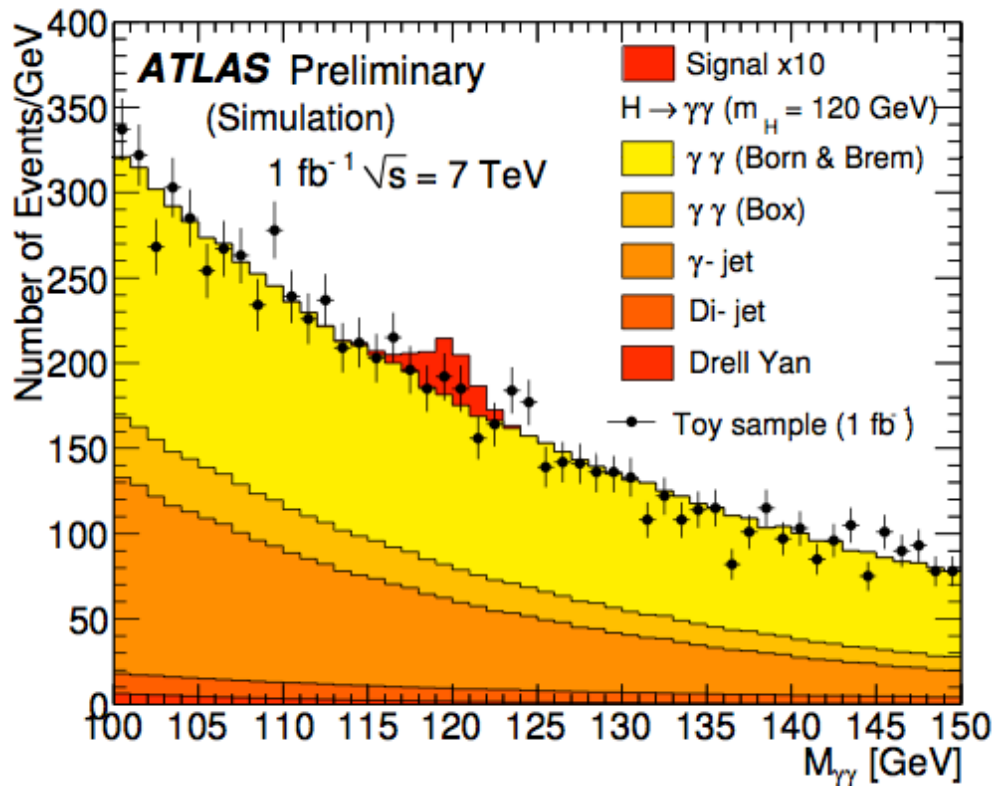
To measure properties in general *(a.k.a. parameter estimation)* among the most commonly used tools is the maximum likelihood fit…

What is a Likelihood ?

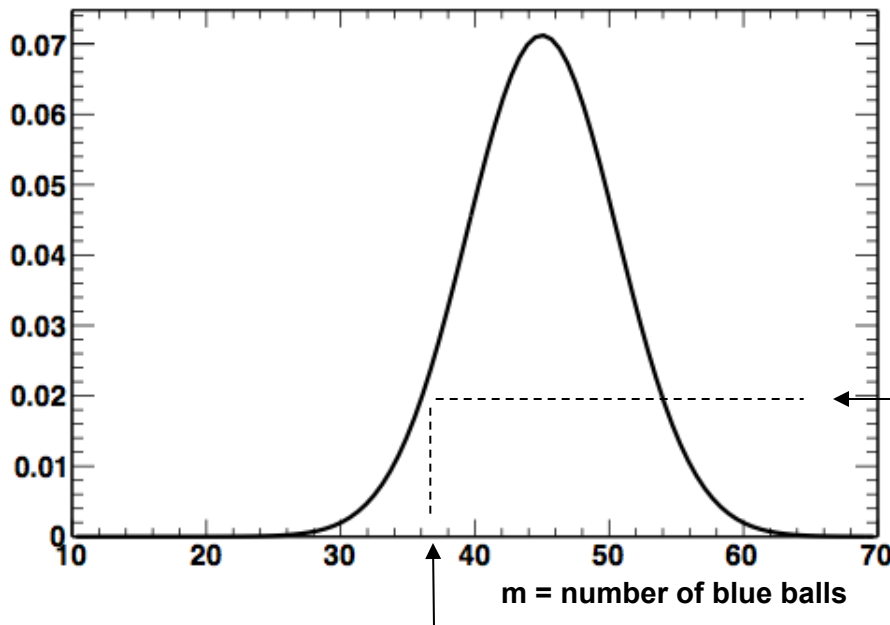A simple way of defining a Likelihood is a Probability Density Function (PDF) which depends on a certain number of parameters…

Simplistic definition is a function with integral equal to 1…

Let's return to the well experiment but under a different angle this time…

(but this applies to any parameter estimate)

Under certain hypothesis :

- Gaussian centered at 45 (p=15%)
- Width equal to error for 1 bucket (300 balls)

$$\sigma = \sqrt{np(1-p)}$$

– $\sigma \sim 6.2$ blue balls)

Here is its probability !

or Likelihood

Not so likely !

Here is your first measurement (36) !

What happens when we throw more buckets ?

$$L(\mu) = \prod_{i=1}^{N} f_\mu(n_i)$$

Then the probability of each bucket can be multiplied!

N=10000

m = number of blue balls

μ = estimate of the mean
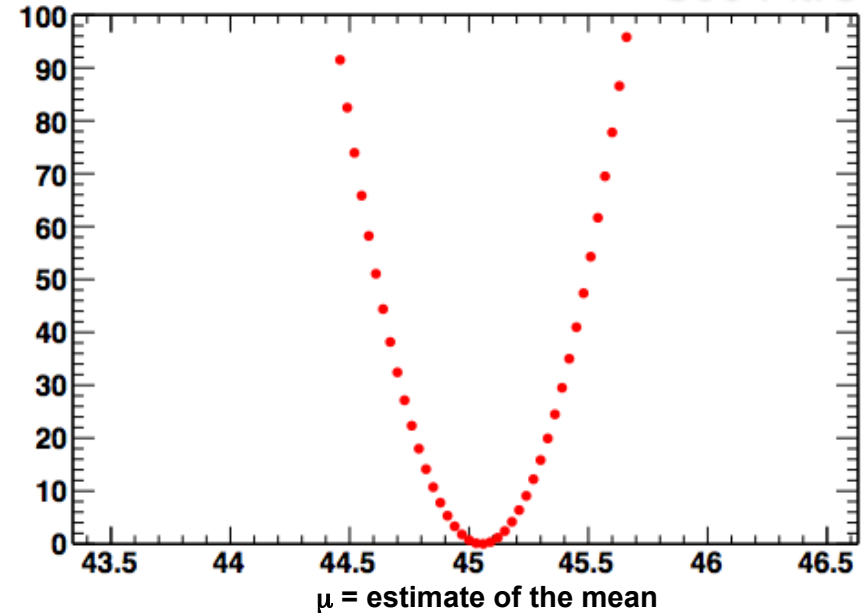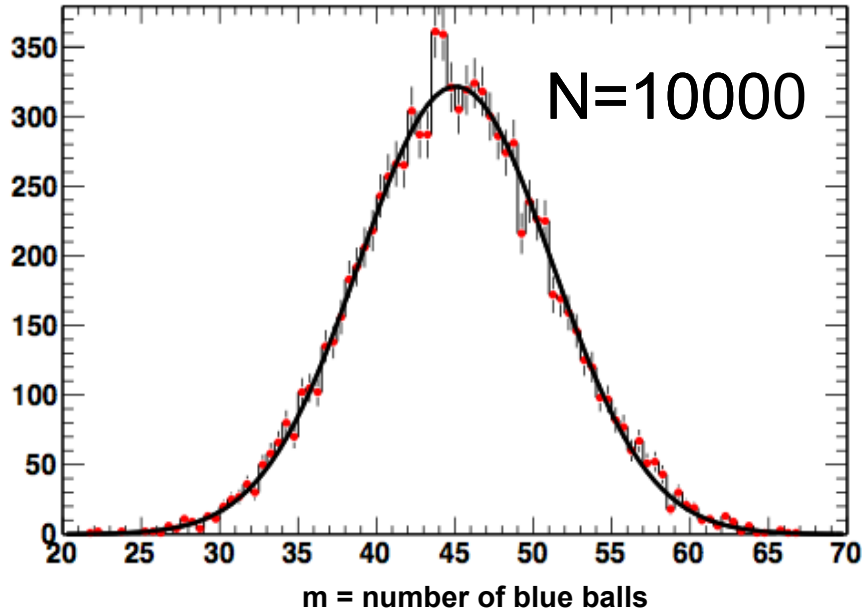
This probability will soon be very very small $(O(0.1))^{100...}$ It is easier to handle its log :

$$\ln(L(\mu)) = \sum_{i=1}^{n} \ln(f_\mu(n_i))$$

Then to estimate a parameter one just has to maximize this function of the parameter μ (or minimize -2ln$L$ you will see why in a slide)…

See how the accuracy translates in the sharpness of the minimum!

In our simple (but not unusual) case we can see that :

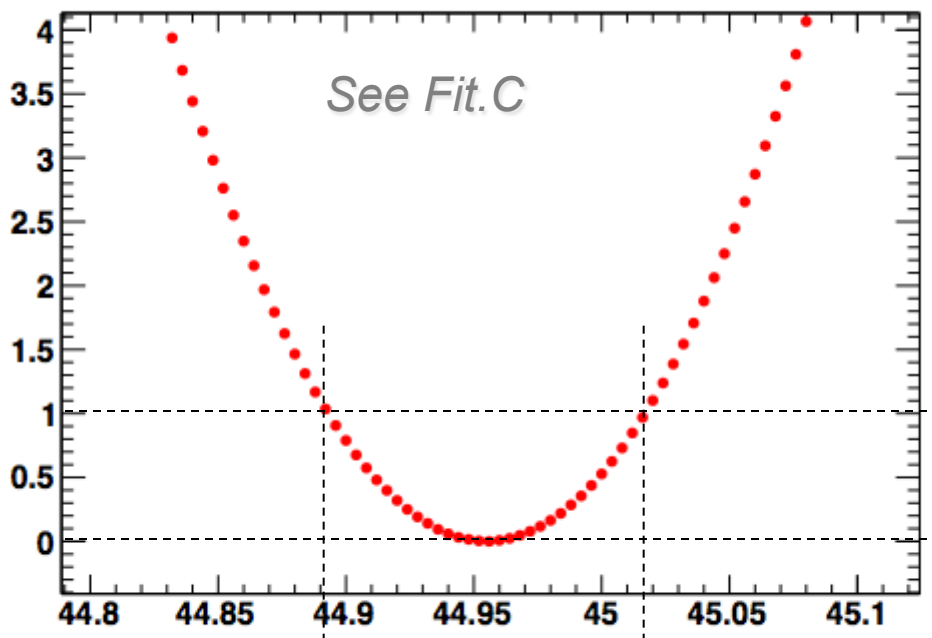$$-2\ln(L(\mu)) = -2\sum_{i=1}^{N}\ln(f_\mu(n_i)) = -2\sum_{i=1}^{N}\ln(\frac{1}{\sqrt{2\pi}\sigma}e^{-\frac{(m_i-\mu)^2}{2\sigma^2}}) = \underbrace{\sum_{i=1}^{N}\frac{(m_i-\mu)^2}{\sigma^2}}_{} + cste$$

This is also called χ^2

There is an exact equivalence between maximizing the Likelihood or minimizing
the χ^2 (Least Squares Method) in the case of a gaussian PDF

See Fit.C

You can also see that the error on
the measured value will be given by
a variation of -2 ln L of one unit :

$$\Delta(-2\ln(L(\mu))) = 1$$

$$\overline{\mu} = 44.95 \pm \underbrace{0.06}_{}$$

Which is precisely $\dfrac{\sigma}{\sqrt{n}}$

"Proof" : Estimation of variance of model parameters (errors)

$$S = \sum_{i=1}^{n} \left[\frac{y_i - f(x_i; a_j)}{\sigma_i} \right]^2 \qquad \underline{V} = \begin{pmatrix} \sigma_1^2 & \mathrm{cov}(1,2) & \mathrm{cov}(1,3) & \dots \\ . & \sigma_2^2 & \mathrm{cov}(2,3) & \dots \\ . & . & \sigma_3^2 & \dots \\ . & . & . & \end{pmatrix} \qquad (\underline{V}^{-1})_{ij} = \frac{1}{2} \frac{\partial^2 S}{\partial a_i \partial a_j}$$

The likelihood function
The mean is estimated by a model
a_j are model parameters

The errors in the parameters are estimated by
the diagonal elements of the covariance matrix,
which for linear least squares is given by the
inverse of the double partial derivative.

$$\sigma^2 = \left| \frac{1}{2} \frac{\partial^2 S}{\partial \theta^2} \right|^{-1}$$ In 1d …. simple case

A robust procedure varies the twice
log likelihood function about the
minimum θ* by 1 to find the root of the
variance in the model parameter.

If we expand S in Taylor series about the minimum

$$S(\theta) = S(\theta^*) + \frac{1}{2} \frac{\partial^2 S}{\partial \theta^2} (\theta - \theta^*)^2$$

$$= S(\theta^*) + \frac{1}{\sigma^2} (\theta - \theta^*)^2 .$$

For higher dimensionality, the tested
model parameter is stepped while the
others are varied to maintain the
minimum condition.

At the point $\theta = \theta^* + \sigma$, we thus find that

$$S(\theta^* + \sigma) = S(\theta^*) + 1 .$$

What have we learned?

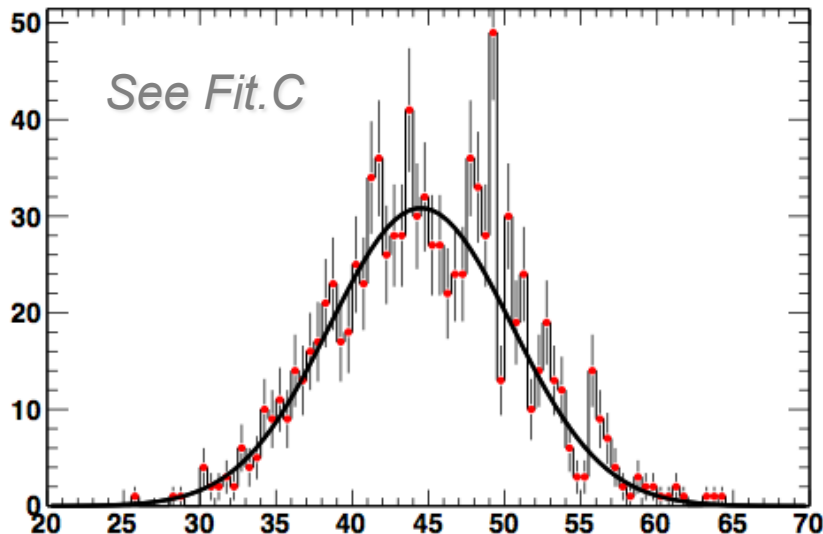How to perform an unbinned likelihood fit :

See Fit.C

For n=1000 the fit yields

$$\overline{\mu} = 44.91 \pm 0.19$$

Using a simple binned fit (as shown here with 100 bins) in the same data yields :

$$\overline{\mu} = 44.81 \pm 0.20$$

LSM between the PDF and the bin value

This can of course be applied to any parameter estimation, as for instance the di-photon reconstructed mass !

More on the χ^2

The χ^2 value is itself a statistic (random variable).
One can repeat the measurement, (collection of the data), and one would get a
different data set, and then calculate a different χ^2_i
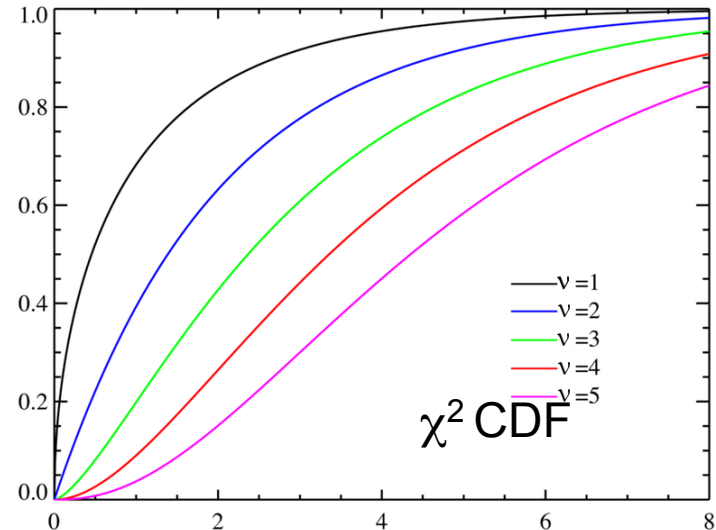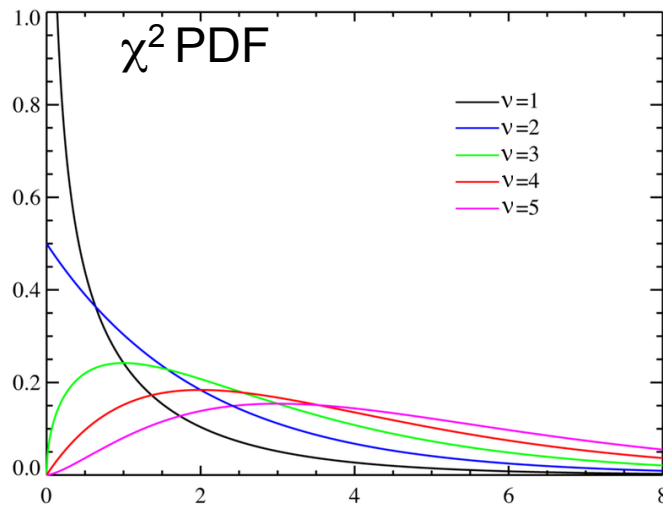This means that the value of χ^2 belongs to a distribution.
As we can write down a closed mathematical expression for the
it follows that the χ^2 distribution is amenable to analysis, and can be calculated as:

$$\chi^2 = -2 \sum_{i=1}^{N} \ln(\frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(m_i - \mu)^2}{2\sigma^2}}) \qquad P(u) = \frac{(u/2)^{(v/2)-1} e^{-u/2}}{2\Gamma(v/2)} \qquad <\chi^2> = v$$

We have used $v = <\chi^2>$ to avoid confusion with the exponent.
$\Gamma(v/2)$ represents the gamma function and v the degrees of freedom (see later).

χ^2 PDF — with curves $v=1$, $v=2$, $v=3$, $v=4$, $v=5$

χ^2 CDF — with curves $v=1$, $v=2$, $v=3$, $v=4$, $v=5$

Max Likelihood Estimation, fitted data, or modeling the distribution of each event

$$-2\ln(L(\mu)) = \sum_{i=1}^{N} \frac{(m_i - \mu)^2}{\sigma^2} + cste$$

See Fit.C

N=10000

MLE$_{obs}$

m = number of blue balls

MLE$_{exp}$

μ = estimate of the mean
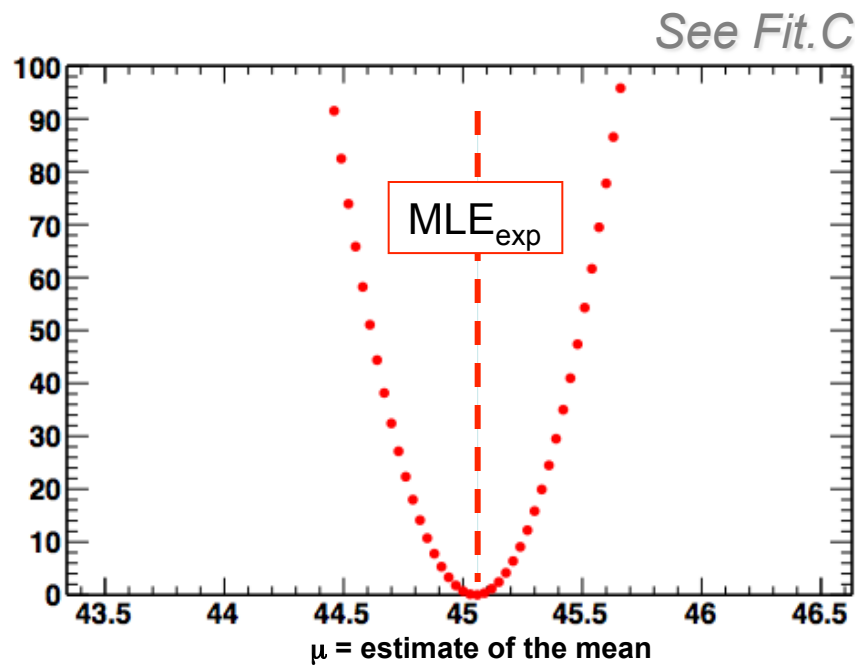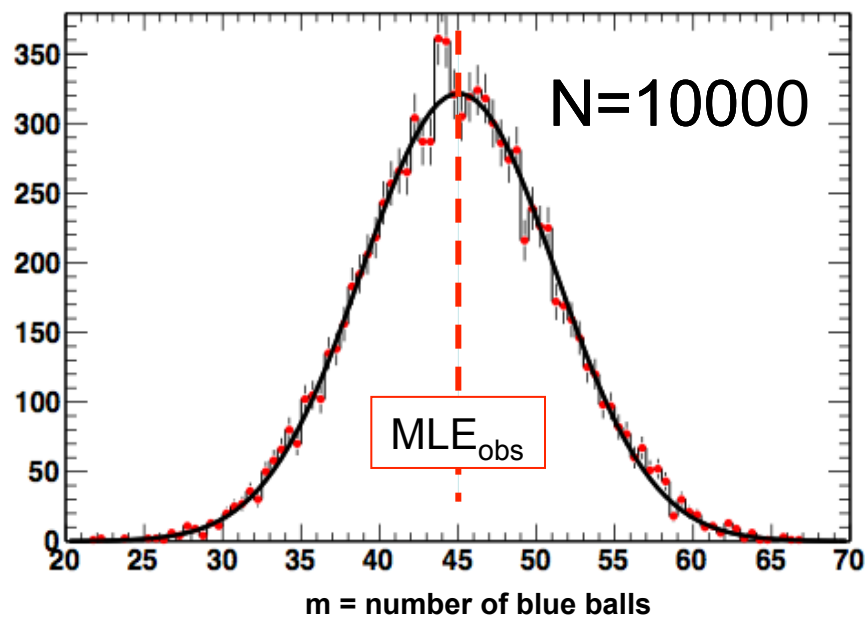
Hypothesis Testing

How to set limits or claim discovery ?

Hypothesis Testing in HEP Boils Down to One Question :
Is there a Signal ?

Exclusion, Observation or Discovery ?

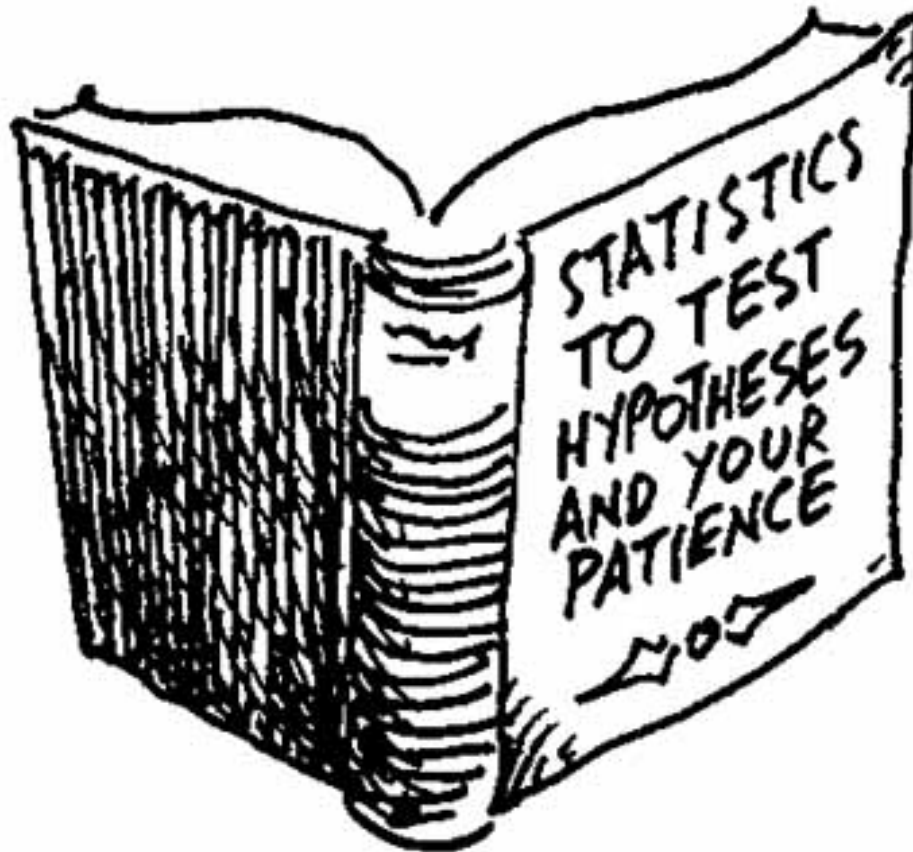The goal here is to assess quantitatively the compatibility of our observation with two hypotheses :

No-Signal (H_0) and presence of Signal (H_1)…

We need to be able to have estimate whether an experiment is more Signal-like or Background-Like.
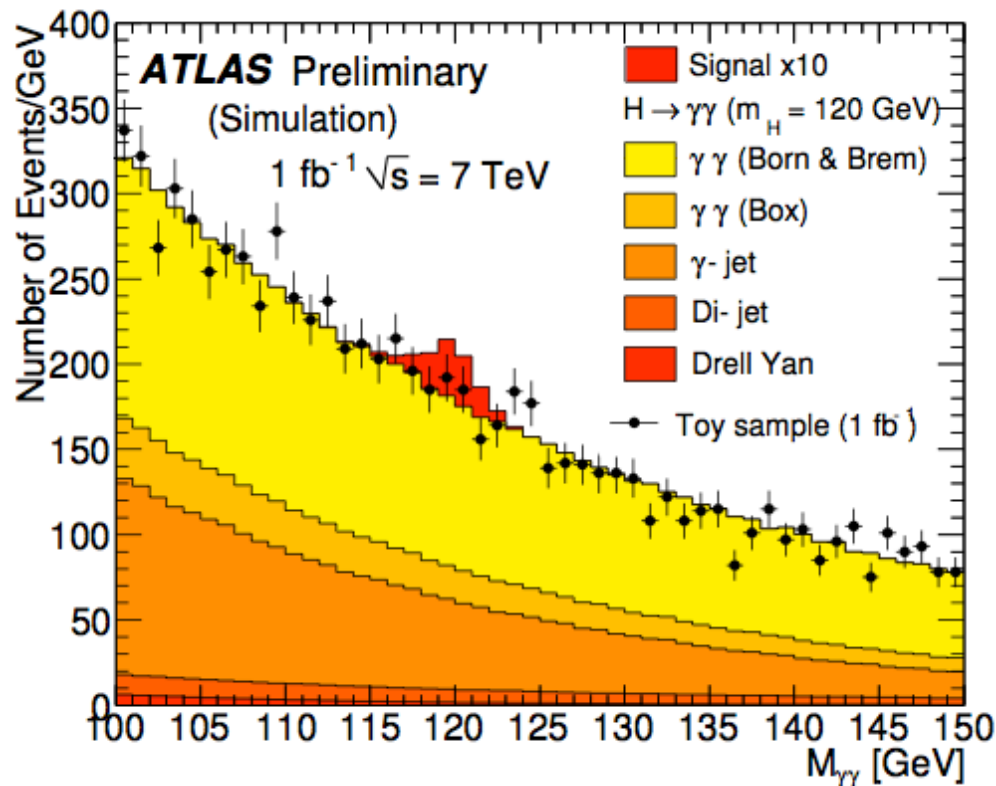
Let's again take the example of the

$gg \rightarrow H \rightarrow \gamma\gamma$ analysis at LHC
(in ATLAS)

Conceptual Intro :
The F-Test as a test statistic

data point (x_i, y_i)

Model dependent on parameters (now a Gaussian)

We can think of the χ^2 defined above equivalently as

$$\chi^2 = \sum_{i=1}^{N} \frac{(m_i - \mu)^2}{\sigma^2} + cste \approx \sum_{i=1}^{N} \frac{[y_i - y(x_i; A, \mu, s)]^2}{\sigma_i^2} + cste_2$$

Error in data point, assumed Gaussian but Poisson in this example.

So ... consider the case where the test statistic is defined as

$$F = \frac{\chi^2(H_1 | x) / v_1}{\chi^2(H_0 | x) / v_2} = \frac{\frac{1}{v_1} \sum \frac{(f_1(x_i; h, \widehat{\theta}) - y_i)^2}{\sigma^2}}{\frac{1}{v_2} \sum \frac{(f_2(x_i; \widehat{\theta}) - y_i)^2}{\sigma^2}}$$

With reference to the High Energy Physics example, in H_1, h is the height of a (Gaussian) peak (of assumed known width) on a smooth background characterised by a function of parameters θ, and in H_0, there is only the smooth background.
The ratio of two χ^2 distributions will be well defined because the χ^2 is well defined.
The ratio is the F statistic, which itself belongs to a distribution.

$$Q(F | v_1, v_2) = I_{\frac{v_2}{v_2 + v_2 F}} \left(\frac{v_2}{2}, \frac{v_1}{2} \right)$$

Where I is the incomplete beta function.
Note : We are asking if the two distributions (with and without the peak) are different.

v_1 and v_2 are the degrees of freedom for H_1 and H_0 respectively.
H_1 is described by $f(x,h,\theta)$ which has n data points and m free parameters.
Then, $v_1 = n - m$. H_0 will have two less degrees of freedom than H_1.
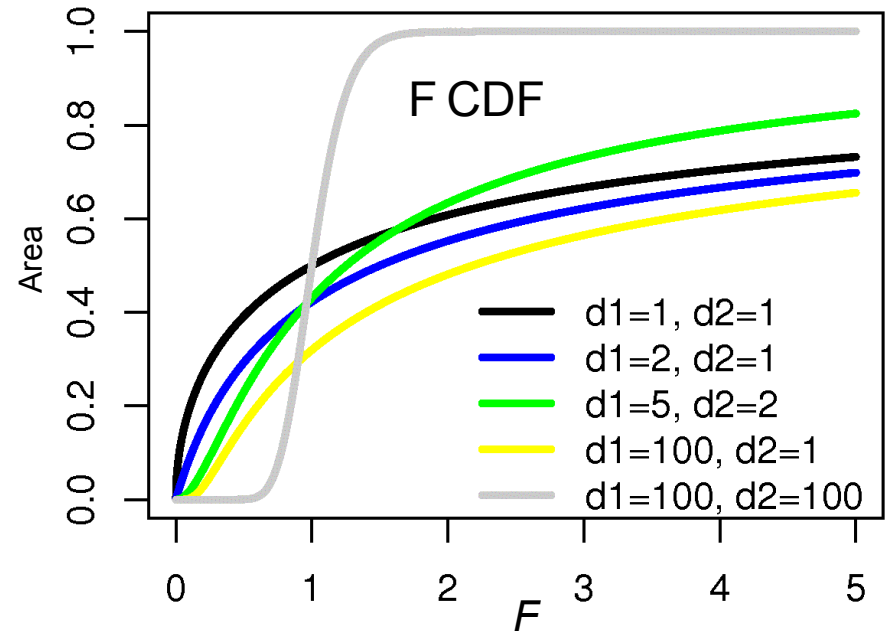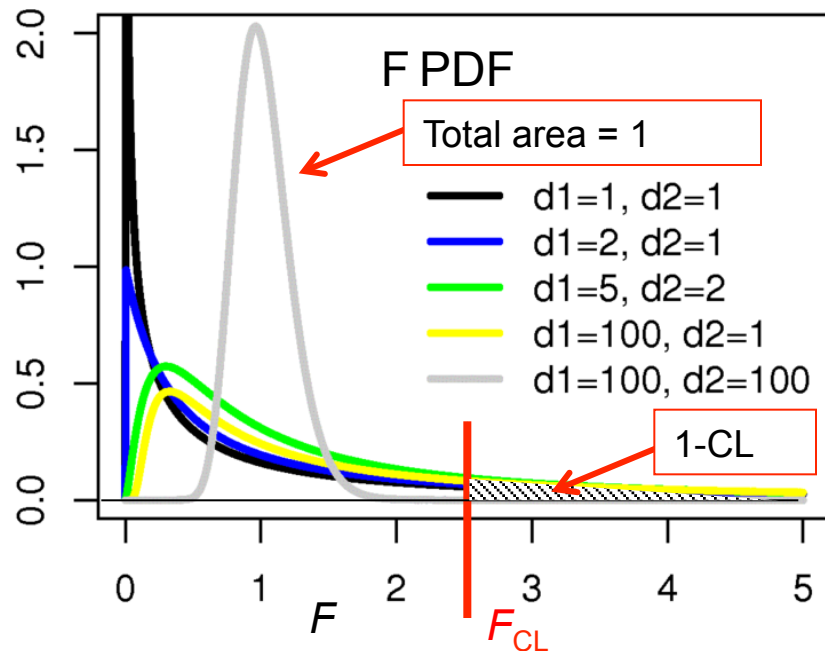
A confidence limit for the rejection (acceptance) of H_0, the null hypothesis, that there is no peak, corresponds to discovery (exclusion).
In the analysis, the confidence limit is pre-set at a desired CL%, and the F distribution is integrated to the the F-value of F_{CL}. Based on the cumulate F distribution to the point F_{CL}, we are CL% certain that a measured F-value larger than F_{CL} is not statistically acceptable as being consistent with H_0.

This analysis is didactic and illustrative, but it suffers from several drawbacks. It does not respect the "*look elsewhere*" effect, it assumes a normal distribution for the data, it cannot easily take into account the full systematics of the measurement, amongst other issues.

The "*look elsewhere*" effect considers that we do not know where the peak should be. The estimated probability of the peak must be diluted by the number of ways that it could have been manifested (~ factor of the measurement interval divided by the peak width) – the wider the search, the more chance for the bg to fluctuate in a peaky way.
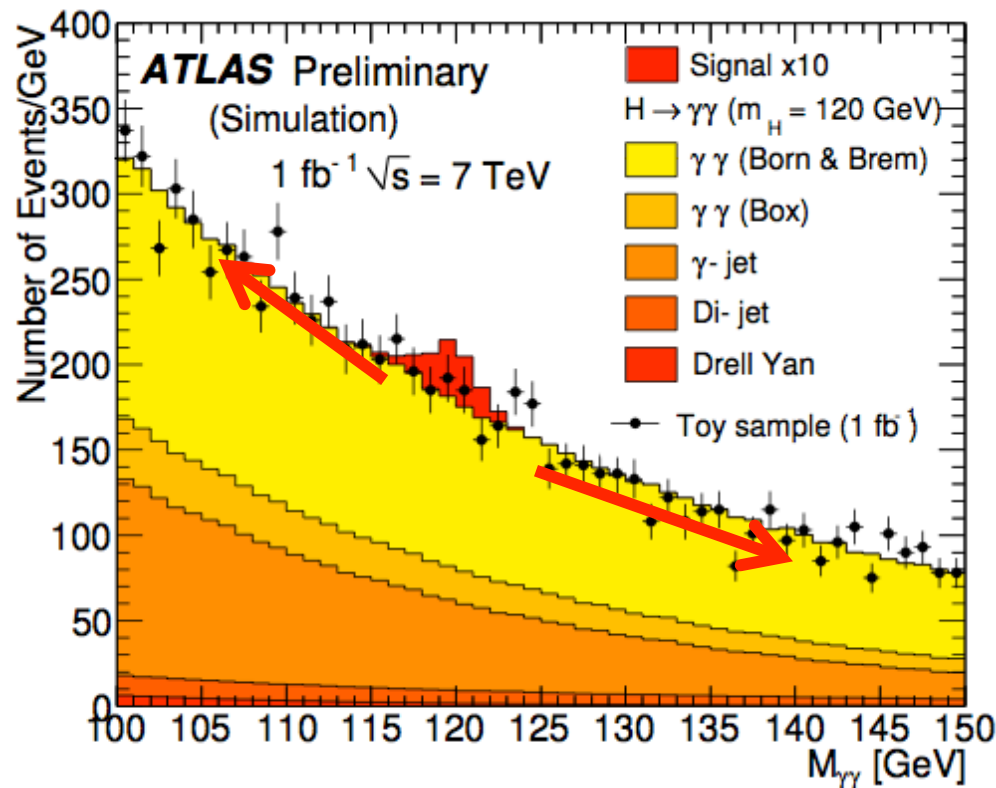
To remove dependence on the previous Gaussian assumption, an improvement is to develop toy Monte Carlo pseudo experiments to get the PDFs for the test statistic.
Test for a peak + bg, H_1, against the bg only, H_0, for a range of invariant masses.

→ Constructing the test statistic

$$E = \frac{P(H_1 \mid x)}{P(H_0 \mid x)}$$

The Neyman-Pearson Lemma

The underlying concept in ordering experiments is really to quantify the compatibility of the observation with the signal hypothesis (H1) …

The problem of testing Hypotheses was studied in the 30's by Jerzy Neyman and Egon Pearson…

They have shown that the ratio of likelihoods of an observation under the two hypotheses is the most powerful tool (or test-statistic or order parameter) to
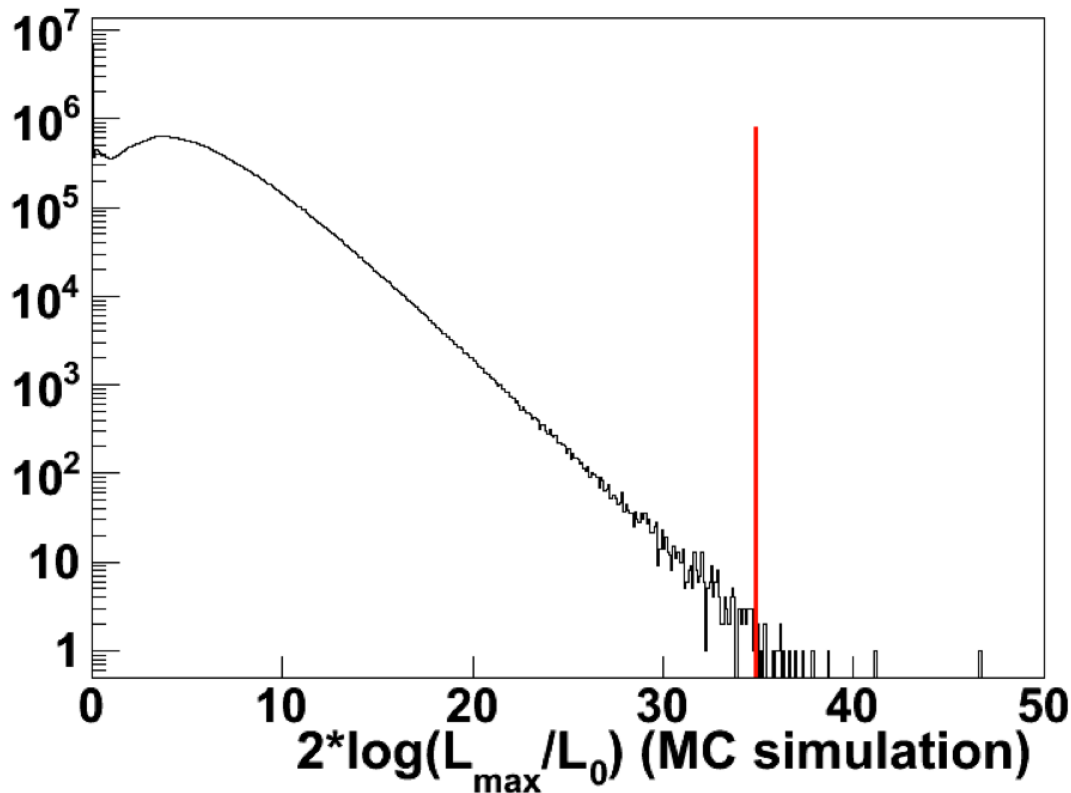
$$E = \frac{P(H_1 \mid x)}{P(H_0 \mid x)}$$

The "χ^2" statistic for H_1 and H_0 can be calculated using the synthetic data.
The toy MC pseudo experiment can be repeated many times, billions of times, and the PDF's of the "χ^2" statistic for H_1 and H_0 can be numerically assembled.

The same can be done for the statistic

$$E = \frac{P(H_1 \mid x)}{P(H_0 \mid x)}$$

Likelihood ratio

2*log(L_{max}/L_0) (MC simulation)

The "*look elsewhere*" effect will be accommodated if the peak position is a free parameter, and it could then range freely in the position where the statistical fluctuations allow it to be found most favorably. Other effects (width variations, systematics are able to be included in developing the PDF's.

The process of setting a CL% and determining a p-value from the CDF can now follow based on these distributions.

The Profile Likelihood

A very useful tool to compute limits, observation or discovery sensitivties and treat systematics is the Profile Likelihood … based on toy MC pseudo experiments.

Let's again take the example of the H→γγ analysis at LHC *(in ATLAS)*
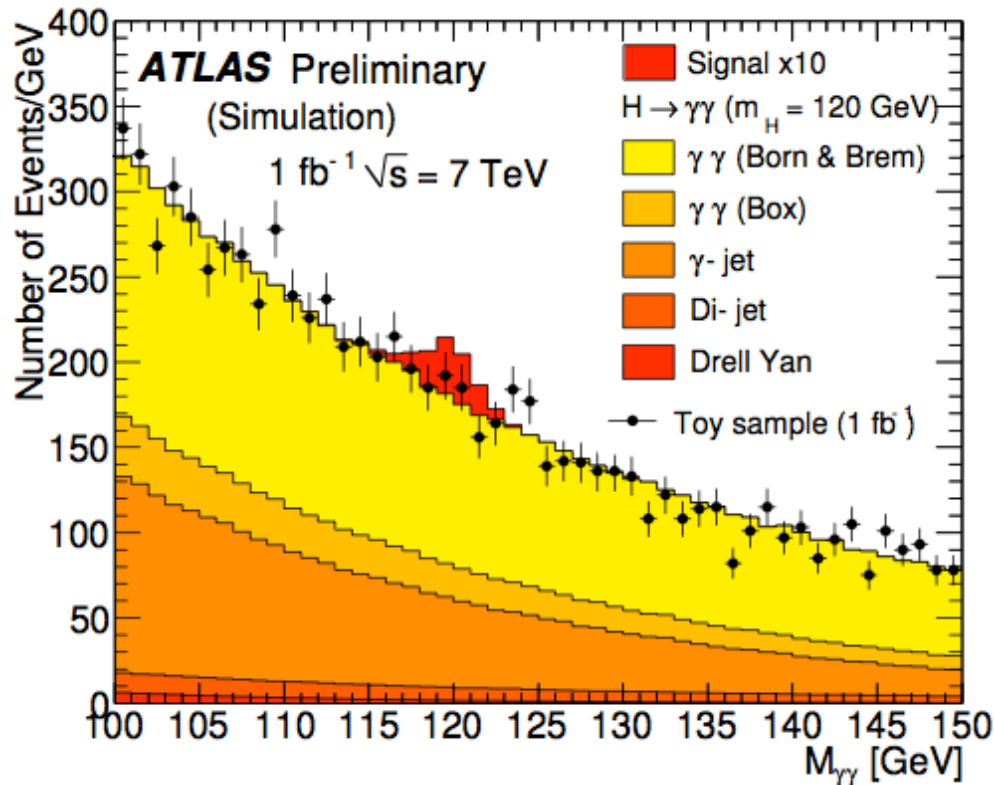
We have a simple model for the background :

$$b(m, \theta) = \theta_1 e^{-\theta_2 m}$$

Relies only on two parameters

Assume a very simple model for the signal :

$$s(m, \mu) = \mu s_0 \times Gauss(m)$$

The Gaussian is centered at 120 GeV/c² and a width of 1.4 GeV/c²

The Profile Likelihood

The overall fit model is very simple :

$$L(\mu, \theta \mid data) = \prod_{i \in data} (s(m_i, \mu) + b(m_i, \theta))$$

This model relies essentially only on two types of parameters :

- The signal strength parameter (μ)　　　It is essentially the signal normalization

- The nuisance parameters (θ)　　Background description in the *"side bands"*
θ Includes effects of ~ luminosity, identification efficiencies, jet energy scale, etc

$$\lambda(\mu) = \frac{L(\mu, \hat{\hat{\theta}}(\mu) \mid data)}{L(\hat{\mu}, \hat{\theta} \mid data)}$$

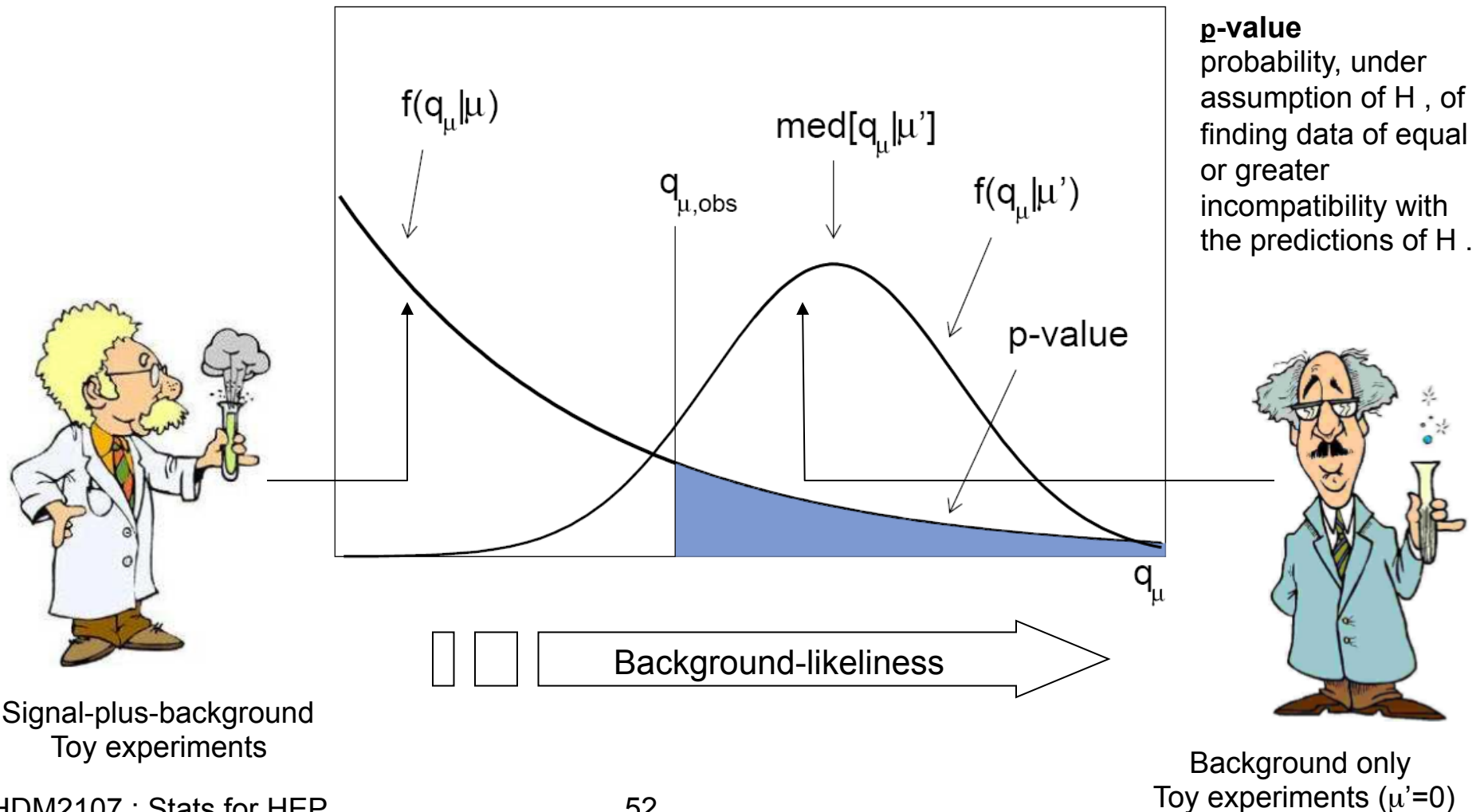⟵　Test of a given signal hypothesis μ

⟵　Best fit of the data

Prescription similar to the Feldman Cousins

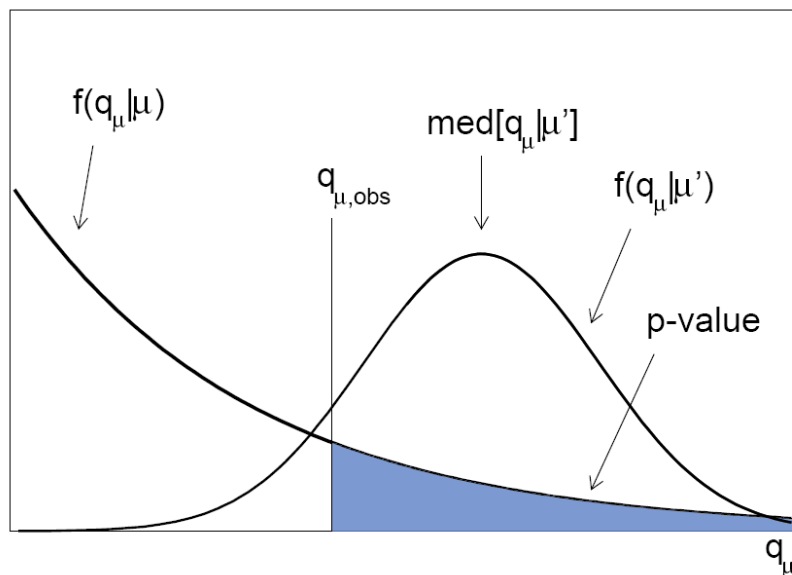Usually work with the estimator :　$q_\mu = -2\ln(\lambda(\mu))$　　Because …

Wilks' Theorem

Under the H_μ Signal hypothesis the PL is distributed in the asymptotic limit as a χ^2 with 1 d.o.f. !   (v.i.z a well know analytical function)

To estimate the overall statistical behavior, toy MC full experiments are simulated and fitted !

$f(q_\mu|\mu)$

$med[q_\mu|\mu']$

$q_{\mu,obs}$

$f(q_\mu|\mu')$

p-value

p-value
probability, under assumption of H , of finding data of equal or greater incompatibility with the predictions of H .

q_μ

Background-likeliness

Signal-plus-background
Toy experiments

Background only
Toy experiments ($\mu'=0$)

95% CL Limits

The observed 95% CL upper limit on μ is obtained by varying μ until the p value :

$$1 - CL_{s+b} = p = \int_{q_{obs}}^{+\infty} f(q_\mu \mid \mu)dq_\mu = 5\%$$

Analytically simple

This means in other words that if there is a signal with strength μ, the false exclusion probability is 5%.

The 95% CL exclusion sensitivity is obtained by varying μ until the p value :

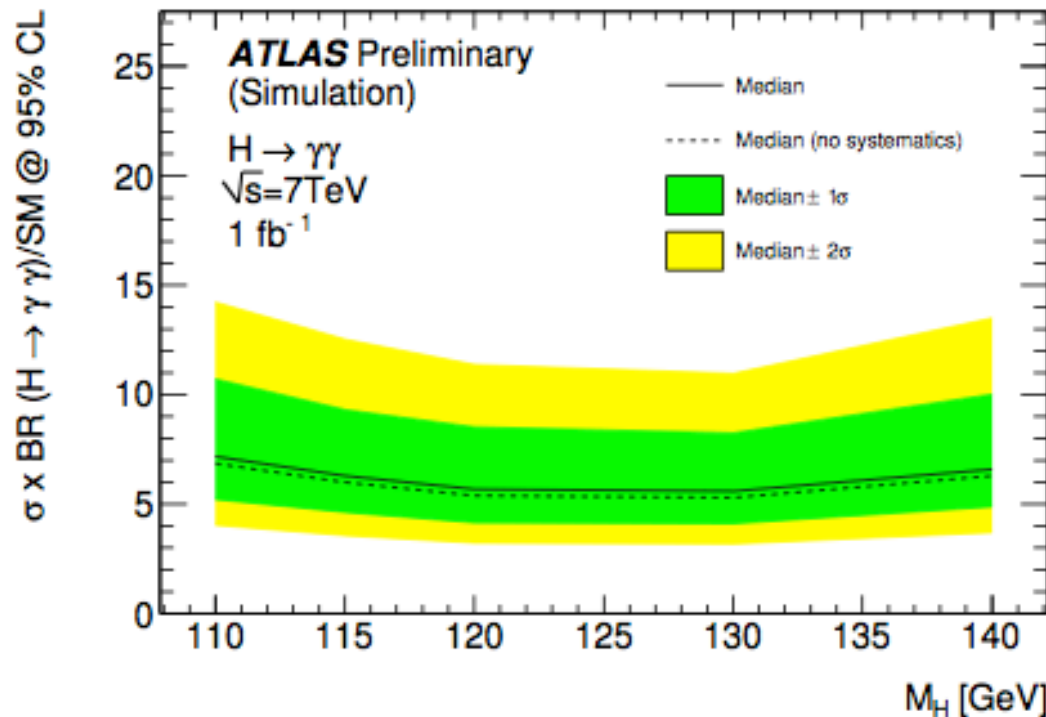$$p = \int_{med(q_\mu \mid 0)}^{+\infty} f(q_\mu \mid \mu)dq_\mu = 5\%$$

Background only experiments

Exclusion Results

Performing this analysis for several mass hypotheses and using CL_{s+b} the exclusion has the same problem as the simple Poisson exclusion with background…

No-Signal (H_0) and presence of Signal (H_1)…

i.e. a signal of 0 can be excluded with a fluctuation of the background

We thus apply the (conservative) "modified frequentist" approach that requires :

$$CL_s = CL_{s+b} / CL_b = 5\%$$ where $$CL_b = \int_{q_{obs}}^{+\infty} f(q_\mu \mid 0) dq_\mu$$
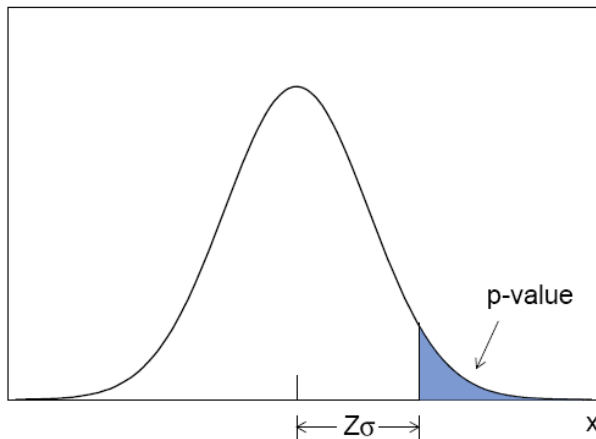
Observation and Discovery

The method is essentially the same, only the estimator changes…we now use q_0

In this case the $f(q_0|0)$ will be distributed as a χ^2 with 1 d.o.f. (Wilks' theorem)

$$p = \int\limits_{q_{obs}}^{+\infty} f(q_0 \mid 0) dq_0$$

- To claim evidence of an observation (3 σ) : the conventional p-value required is 1.35×10^{-3}
- To claim a discovery (5 σ) : the conventional p-value required is 2.87×10^{-7}

Corresponds to the "one sided" convention

This means in other words that in absence of signal, the false discovery probability is p.

« a probability of 1 in 10 000 000 is *almost* impossible to estimate »

R. P. Feynman

(What do you care what other people think?)

p-value .... significance

For **discovery**, reject the background hypothesis with a significance of at least $Z = 5$.

This corresponds to $p = 2.87 \times 10^{-7}$

For **exclusion**, a threshold p-value of 0.05 (95% confidence level) is used, which corresponds to $Z = 1.64$.

For predicting the **sensitivity** to discovery of a given signal process H_1 one assumes H_1 and calculates the expectation value, Z obtained testing for H_0.
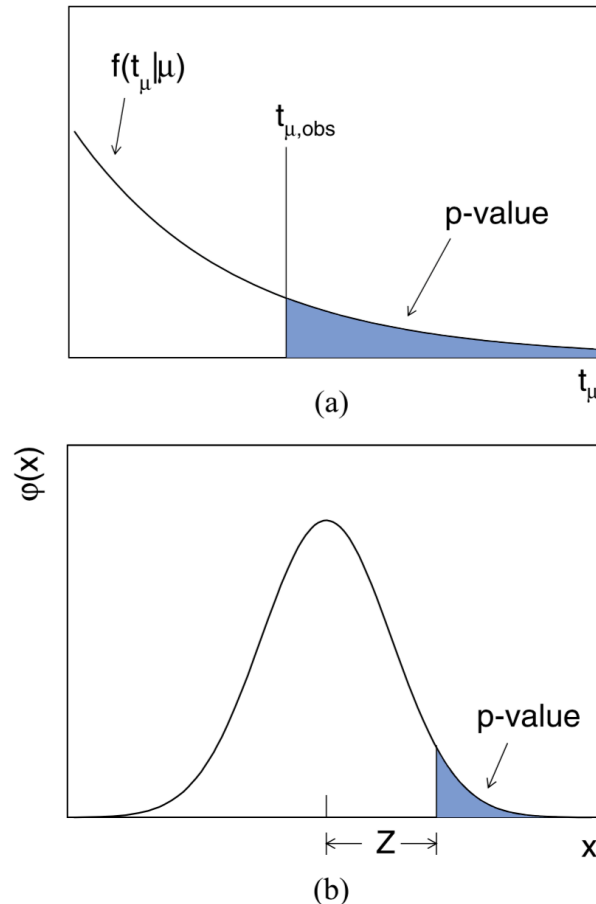
Fig. 1 (a) Illustration of the relation between the p-value obtained from an observed value of the test statistic t_μ. (b) The standard normal distribution $\varphi(x) = (1/\sqrt{2\pi}) \exp(-x^2/2)$ showing the relation between the significance Z and the p-value

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Conclusion

We went through an overview of the fundamental concepts of statistics for HEP

If possible take some time to play with the Root-Macros for hands-on experience

You should now be able to understand the following plot !
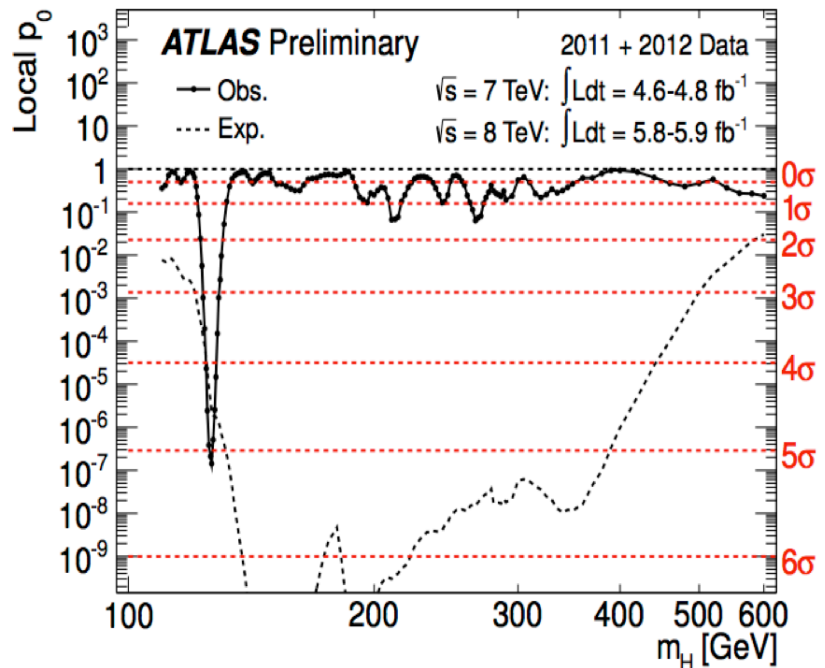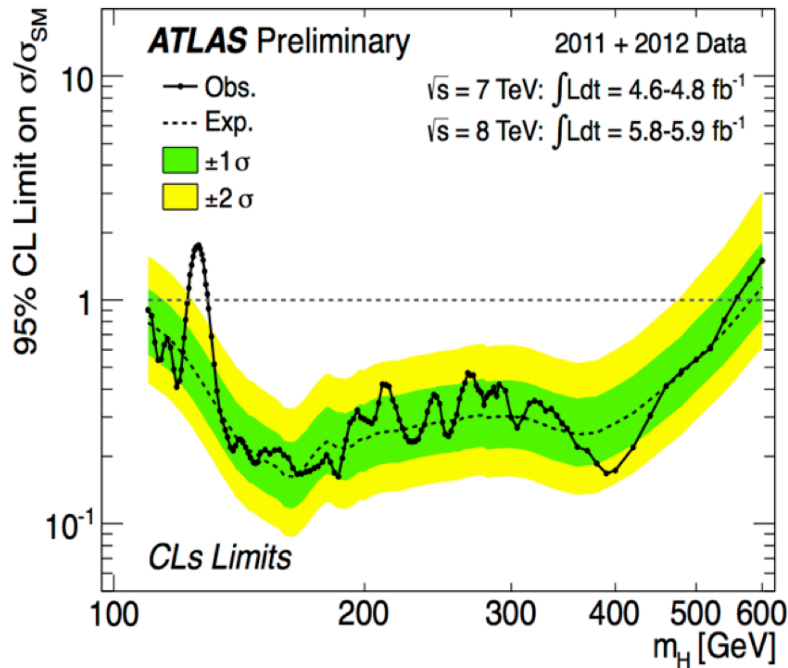
There is a lot more for you/us to learn about statistical techniques

In particular concerning the treatment of systematics

So be patient and take some time to understand the techniques step by step…

… and follow Laplace's advice about statistics !

Additional

PG Positions
- Some funding available : (SA-CERN and Nat. Res. Foundation)
- Sometimes University based funding.
- SA-CERN (Jean Cleymans : jean.cleymans@gmail.com)
- SA-CERN web-site (google)
- Simon Connell shconnell@uj.ac.za

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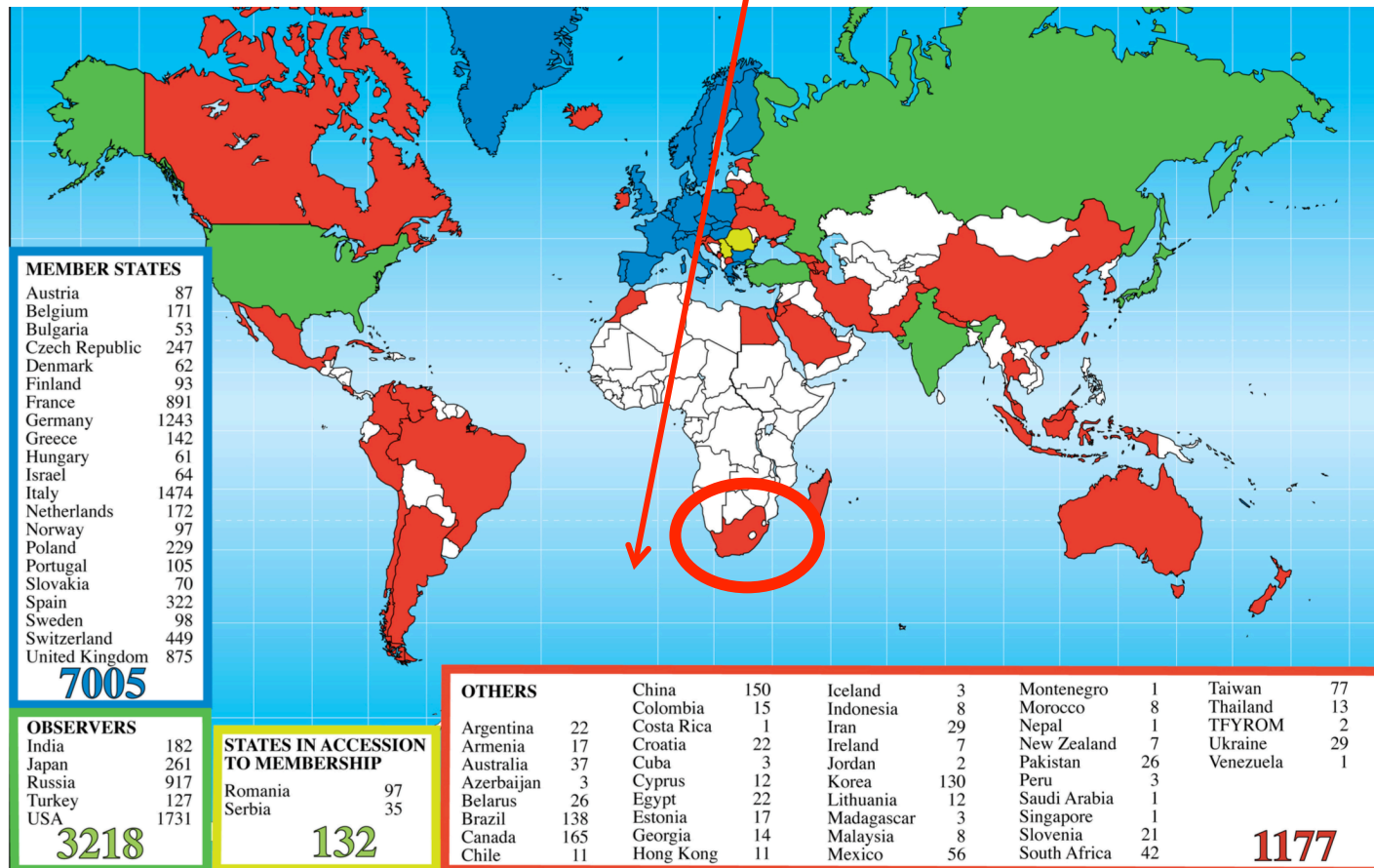
Figure 2. Countries of origin of CERN's visiting scientists (Users) by location of their home institute (January 2015)

Counting statistics for random events

The waiting time distribution between random events is exponentially distributed.

Events occur with a mean rate μ. The probability for an event in time dt is $\mu.dt$.

Consider probability function $F_{no}(t)$ for no event in time t. Then
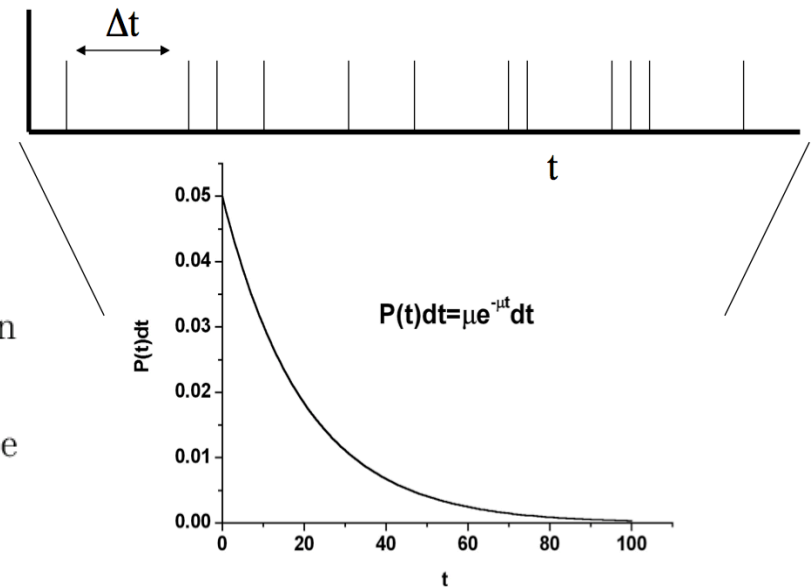
$$F_{no}(t + dt) = F_{no}(t)(1 - \mu dt)$$

solving this equation we get

$$F_{no}(t) = Ce^{-\mu t},$$

C=1 by normalisation

The probability density function $P(t)$ of the random event not occurring for a time t, and then occurring in the next dt.

$$P(t)dt = F_{no}(t)\,\mu dt = \mu\,e^{-\mu t}\,dt$$

Thus, the waiting times between random events are exponentially distributed.

Counting statistics for random events

The number of random events per bin is Poisson distributed

If there are on average $\overline{n} = \mu\Delta t$ events in the tome interval Δt, then the probability of an event in the subinterval will be

$$p = \overline{n}/N \ll 1.$$

The number of events n in the interval Δt then follows a Binomial distribution

$$P(n) = \binom{N}{n} p^n (1-p)^{(N-n)}.$$

Decreasing the size of the subintervals, ie, $N \longrightarrow \infty$ and $p \longrightarrow 0$ but $\overline{n} = Np$ remains finite, we find that the Binomial distribution may be replaced by its approximation for small p, the Poisson distribution.

$$P(n) = \frac{\overline{n}^n e^{-\overline{n}}}{n!}$$

Consequently, the standard deviation of the number of events per bin is

$$\sigma = \sqrt{\overline{n}}$$

So that the relative error in counting random events diminishes as $\sim \frac{1}{\sqrt{n}}$.

n events

1 event

dt

$\Delta t = Ndt$