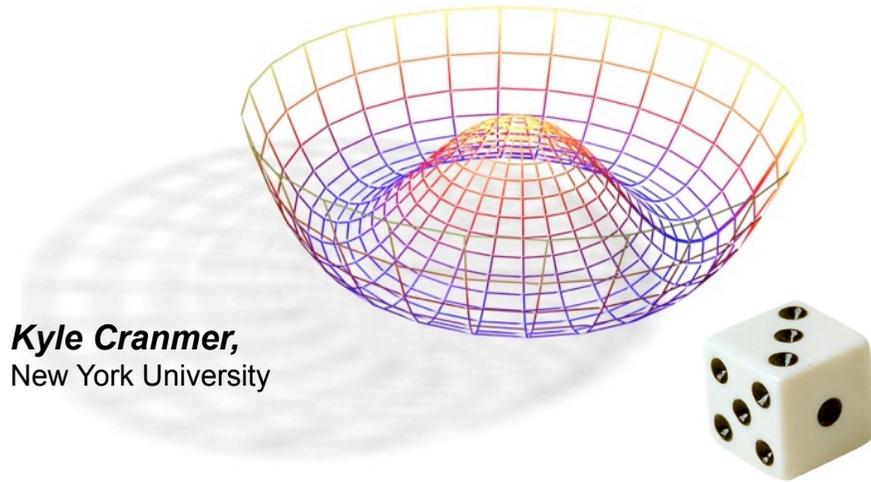


Practical Statistics for Particle Physics



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Lecture 2

Outline

Lecture 1: Building a probability model

- preliminaries, the marked Poisson process
- incorporating systematics via nuisance parameters
- constraint terms
- examples of different “narratives” / search strategies

Lecture 2: Hypothesis testing

- simple models, Neyman-Pearson lemma, and likelihood ratio
- composite models and the profile likelihood ratio
- review of ingredients for a hypothesis test

Lecture 3: Limits & Confidence Intervals

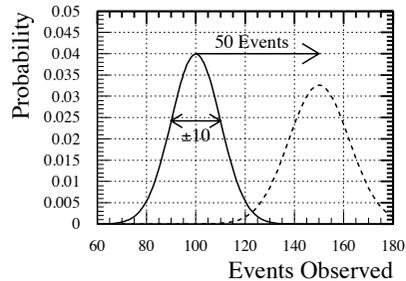
- the meaning of confidence intervals as inverted hypothesis tests
- asymptotic properties of likelihood ratios
- Bayesian approach

Hypothesis Testing

Hypothesis testing

One of the most common uses of statistics in particle physics is Hypothesis Testing (e.g. for discovery of a new particle)

- assume one has pdf for data under two hypotheses:
 - Null-Hypothesis, H_0 : eg. background-only
 - Alternate-Hypothesis H_1 : eg. signal-plus-background
- one makes a measurement and then needs to decide whether to **reject** or **accept** H_0



Hypothesis testing

Before we can make much progress with statistics, we need to decide what it is that we want to do.

- first let us define a few terms:

- Rate of Type I error α
- Rate of Type II β
- Power = $1 - \beta$

		Actual condition	
		Guilty	Not guilty
Decision	Verdict of 'guilty'	True Positive	False Positive (i.e. guilt reported unfairly) Type I error
	Verdict of 'not guilty'	False Negative (i.e. guilt not detected) Type II error	True Negative

Treat the two hypotheses asymmetrically

- the Null is special.
 - Fix rate of Type I error, call it “the size of the test”

Now one can state “a well-defined goal”

- Maximize power for a fixed rate of Type I error

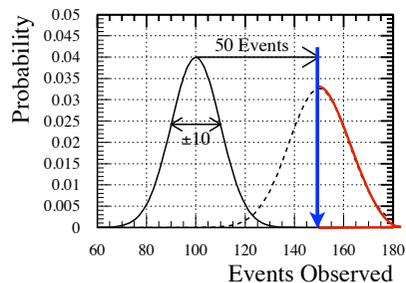
Hypothesis testing

The idea of a “ 5σ ” discovery criteria for particle physics is really a conventional way to specify the size of the test

- usually 5σ corresponds to $\alpha = 2.87 \cdot 10^{-7}$
 - eg. a very small chance we reject the standard model

In the simple case of number counting it is obvious what region is sensitive to the presence of a new signal

- but in higher dimensions it is not so easy



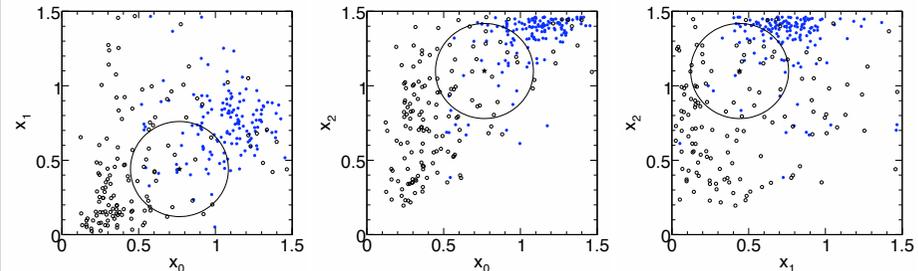
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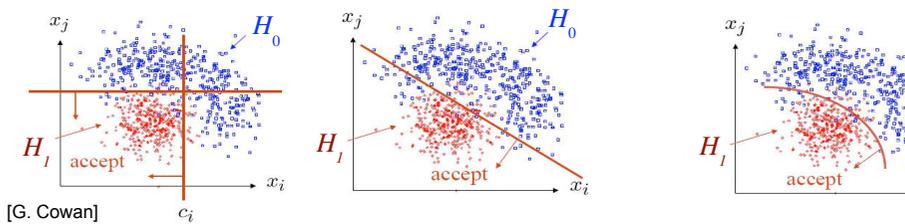


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[G. Cowan]

In 1928-1938 Neyman & Pearson developed a theory in which one must consider competing Hypotheses:

- the Null Hypothesis H_0 (background only)
- the Alternate Hypothesis H_1 (signal-plus-background)

Given some probability that we wrongly reject the Null Hypothesis

$$\alpha = P(x \notin W | H_0)$$

(Convention: if data falls in W then we accept H_0)

Find the region W such that we minimize the probability of wrongly accepting the H_0 (when H_1 is true)

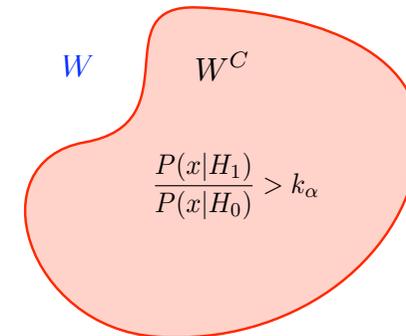
$$\beta = P(x \in W | H_1)$$

The region W that minimizes the probability of wrongly accepting H_0 is just a contour of the Likelihood Ratio

$$\frac{P(x|H_1)}{P(x|H_0)} > k_\alpha$$

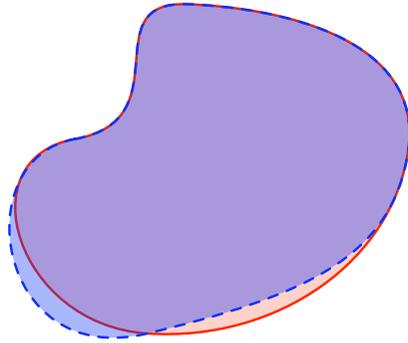
Any other region of the same size will have less power

The likelihood ratio is an example of a **Test Statistic**, eg. a real-valued function that summarizes the data in a way relevant to the hypotheses that are being tested



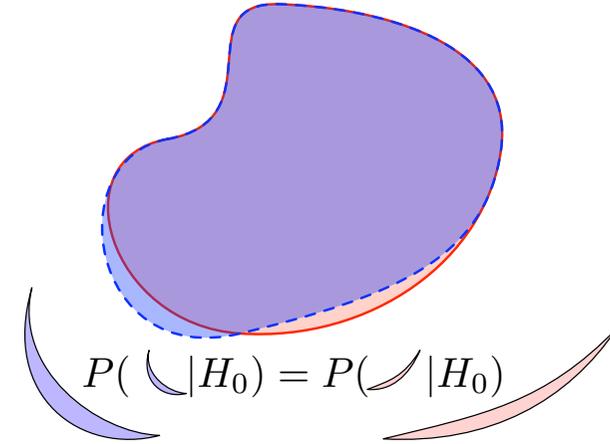
Consider the contour of the likelihood ratio that has size a given size (eg. probability under H_0 is $1-\alpha$)

A short proof of Neyman-Pearson



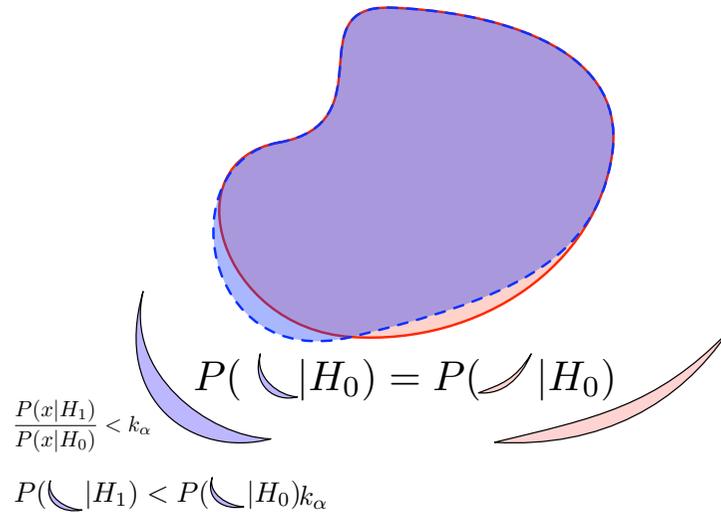
Now consider a variation on the contour that has the same size

A short proof of Neyman-Pearson



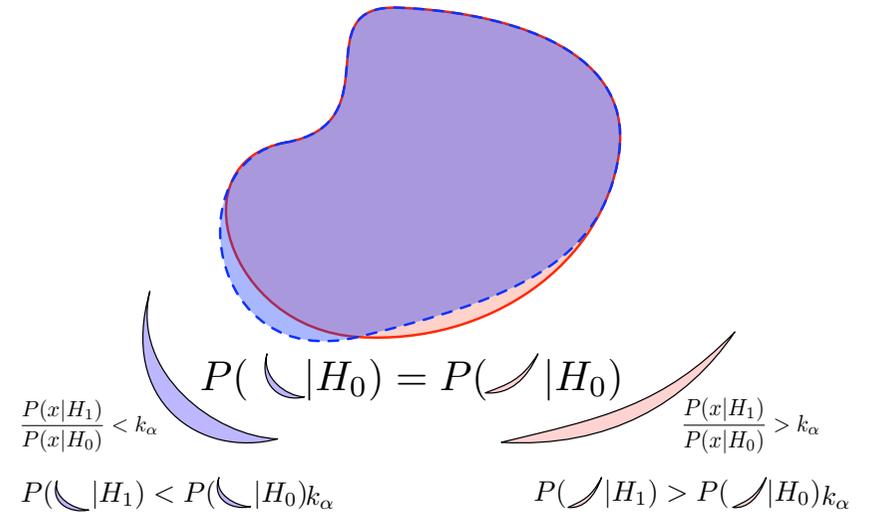
Now consider a variation on the contour that has the same size (eg. same probability under H_0)

A short proof of Neyman-Pearson



Because the new area is outside the contour of the likelihood ratio, we have an inequality

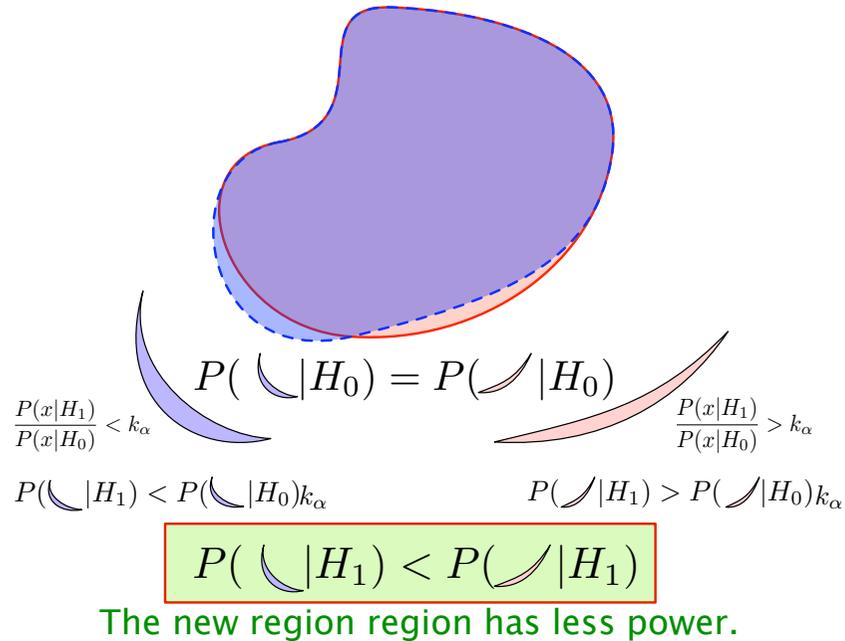
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And for the region we lost, we also have an inequality

Together they give...

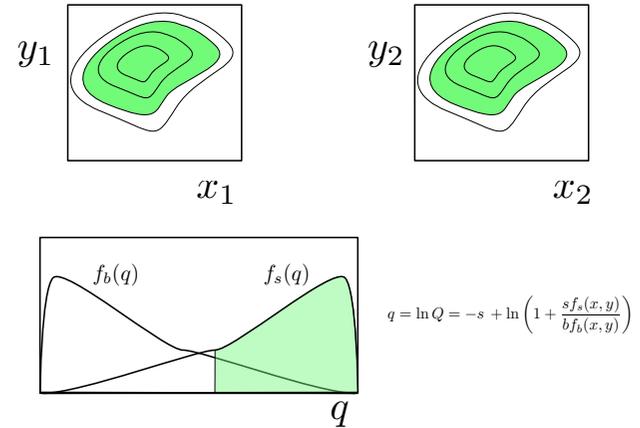
A short proof of Neyman-Pearson



2 discriminating variables

Often one uses the output of a neural network or multivariate algorithm in place of a true likelihood ratio.

- That's fine, but what do you do with it?
- If you have a fixed cut for all events, this is what you are doing:

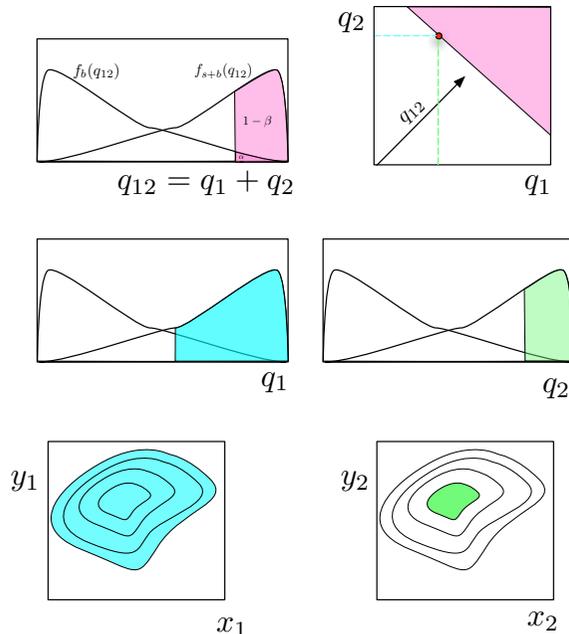


Experiments vs. Events

Ideally, you want to cut on the likelihood ratio for your experiment

- equivalent to a sum of log likelihood ratios

Easy to see that includes experiments where one event had a high LR and the other one was relatively small

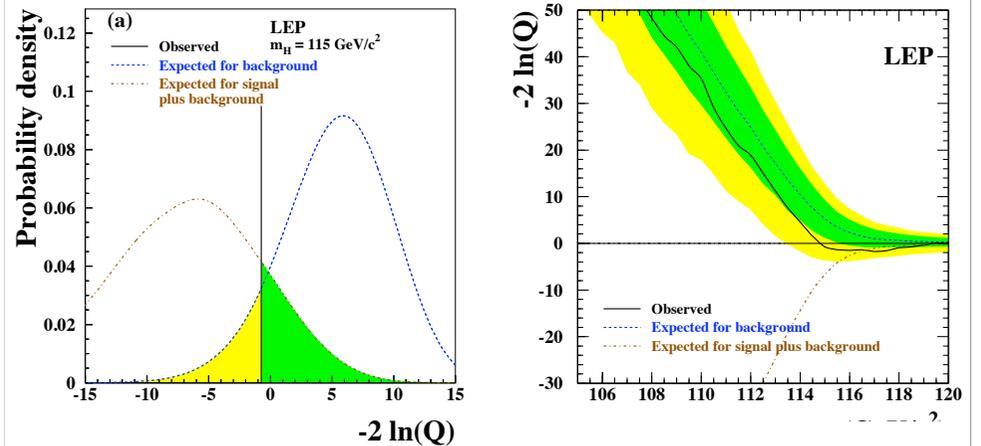


LEP Higgs

A simple likelihood ratio with no free parameters

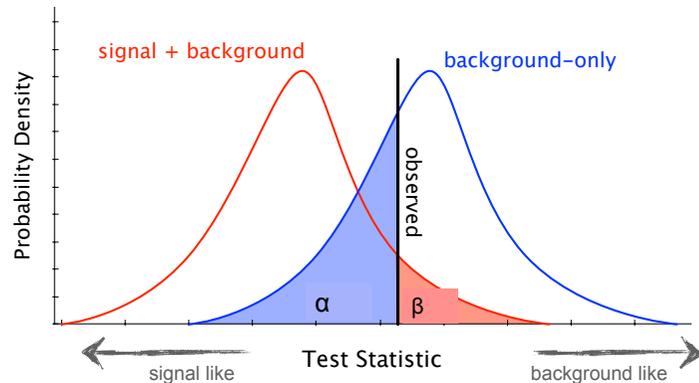
$$Q = \frac{L(x|H_1)}{L(x|H_0)} = \frac{\prod_i^{N_{chan}} Pois(n_i | s_i + b_i) \prod_j^{n_i} \frac{s_i f_s(x_{ij}) + b_i f_b(x_{ij})}{s_i + b_i}}{\prod_i^{N_{chan}} Pois(n_i | b_i) \prod_j^{n_i} f_b(x_{ij})}$$

$$q = \ln Q = -s_{tot} + \sum_i^{N_{chan}} \sum_j^{n_i} \ln \left(1 + \frac{s_i f_s(x_{ij})}{b_i f_b(x_{ij})} \right)$$



The Test Statistic and its distribution

Consider this schematic diagram



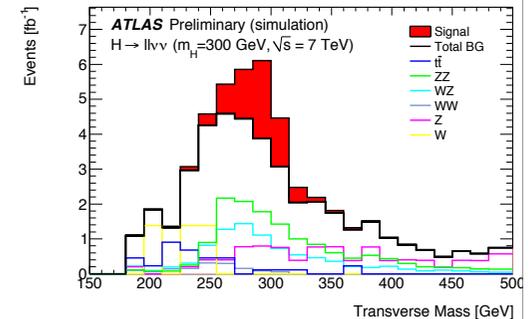
The “**test statistic**” is a single number that quantifies the entire experiment, it could just be number of events observed, but often its more sophisticated, like a likelihood ratio. What test statistic do we choose?

And how do we build the **distribution**? Usually “toy Monte Carlo”, but what about the uncertainties... what do we do with the nuisance parameters?

The Marked Poisson model

Recall our marked Poisson model

- **observables:** n events each with some value of discriminating variable m
- **auxiliary measurements:** a_i
- **parameters:** α



$$P(\mathbf{m}, \mathbf{a} | \alpha) = \text{Pois}(n | s(\alpha) + b(\alpha)) \prod_j \frac{s(\alpha) f_s(m_j | \alpha) + b(\alpha) f_b(m_j | \alpha)}{s(\alpha) + b(\alpha)} \times \prod_{i \in \text{syst}} G(a_i | \alpha_i, \sigma_i)$$

Useful to separate parameters into $\alpha = (\mu, \nu)$

- **parameters of interest μ :** cross sections, masses, coupling constants, ...
- **nuisance parameters ν :** reconstruction efficiencies, energy scales, ...
- note: not all of the nuisance parameters need to have constraint terms

Our number counting example

From our general model

$$P(\mathbf{m}, \mathbf{a} | \alpha) = \text{Pois}(n | s(\alpha) + b(\alpha)) \prod_j \frac{s(\alpha) f_s(m_j | \alpha) + b(\alpha) f_b(m_j | \alpha)}{s(\alpha) + b(\alpha)} \times \prod_{i \in \text{syst}} G(a_i | \alpha_i, \sigma_i)$$

Consider a simple number counting model with $s(\alpha) \rightarrow s$, $b(\alpha) \rightarrow b$, and replace the constraint $G(a | \alpha, \sigma) \rightarrow \text{Pois}(n_{\text{off}} | \tau b)$ with τ known.

$$P(n_{\text{on}}, n_{\text{off}} | s, b) = \text{Pois}(n_{\text{on}} | s + b) \text{Pois}(n_{\text{off}} | \tau b).$$

We could simply use n_{on} as our test statistic, but to calculate the p-value we need to know distribution of n_{on} .

$$p = \sum_{n_{\text{on}}=n_{\text{obs}}}^{\infty} \text{Pois}(n_{\text{on}} | s + b) \times \underbrace{\text{Pois}(n_{\text{off}} | \tau b)}_{\text{constant}}$$

Observations:

- The distribution of n_{on} explicitly depends on both s and b .
- The distribution of n_{off} is independent of s
- If τb is very different from n_{off} , then the data are not consistent with the model parameters. However, the p-value derived from n_{on} is not small.

With nuisance parameters: Hybrid Solutions

Goal of Bayesian-frequentist hybrid solutions is to provide a frequentist treatment of the main measurement, while eliminating nuisance parameters (deal with systematics) with an intuitive Bayesian technique.

$$P(n_{\text{on}} | s) = \int db \text{Pois}(n_{\text{on}} | s + b) \pi(b), \quad p = \sum_{n_{\text{on}}=n_{\text{obs}}}^{\infty} P(n_{\text{on}} | s)$$

Tracing back the origin of $\pi(b)$

- clearly state prior $\eta(b)$; identify control samples (sidebands) and use:

$$\pi(b) = P(b | n_{\text{off}}) = \frac{P(n_{\text{off}} | b) \eta(b)}{\int db P(n_{\text{off}} | b) \eta(b)}.$$

In a purely Frequentist approach we must need a test statistic that depends on both n_{on} and n_{off} and we must consider both random (eg. when generating toy Monte Carlo)

$$P(n_{\text{on}}, n_{\text{off}} | s, b) = \text{Pois}(n_{\text{on}} | s + b) \text{Pois}(n_{\text{off}} | \tau b).$$

Does it matter?

This on/off problem has been studied extensively.

- instead of arguing about the merits of various methods, just go and check their rate of Type I error
- Results indicated large discrepancy in “claimed” significance and “true” significance for various methods
- eg. 5σ is really $\sim 4\sigma$ for some points

So, yes, it does matter.

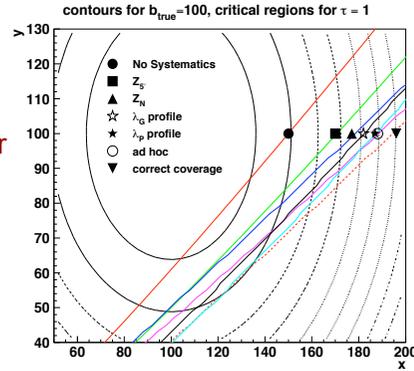


Figure 7. A comparison of the various methods critical boundary $x_{crit}(y)$ (see text). The concentric ovals represent contours of L_G from Eq. 15.

$$P(n_{on}, n_{off}|s, b) = \text{Pois}(n_{on}|s + b) \text{Pois}(n_{off}|\tau b).$$

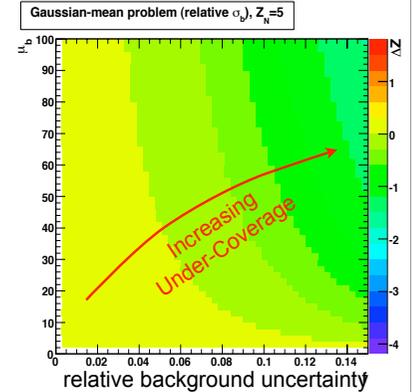
http://www.physics.ox.ac.uk/phystat05/proceedings/files/Cranmer_LHCStatisticalChallenges.ps

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Follow-up work by Bob Cousins & Jordan Tucker, [physics/0702156]

$$P(n_{on}, n_{off}|s, b) = \text{Pois}(n_{on}|s + b) \text{Pois}(n_{off}|\tau b).$$

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The Profile Likelihood Ratio

Consider our general model with a single parameter of interest μ

- let $\mu=0$ be no signal, $\mu=1$ nominal signal

In the LEP approach the likelihood ratio is equivalent to:

$$Q_{LEP} = \frac{P(\mathbf{m}|\mu = 1, \nu)}{P(\mathbf{m}|\mu = 0, \nu)}$$

- but this variable is sensitive to uncertainty on ν and makes no use of auxiliary measurements \mathbf{a}

Alternatively, one can define **profile likelihood ratio**

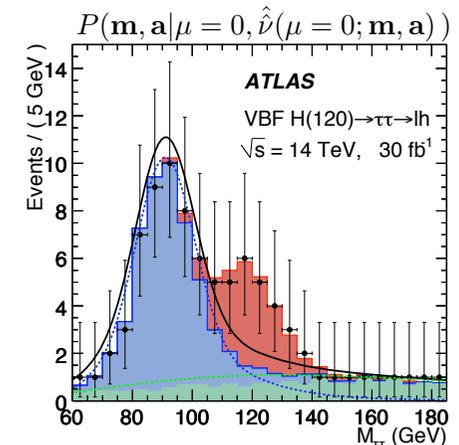
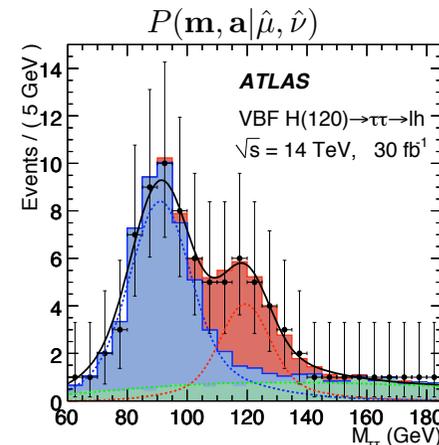
$$\lambda(\mu) = \frac{P(\mathbf{m}, \mathbf{a}|\mu, \hat{\nu}(\mu; \mathbf{m}, \mathbf{a}))}{P(\mathbf{m}, \mathbf{a}|\hat{\mu}, \hat{\nu})}$$

- where $\hat{\nu}(\mu; \mathbf{m}, \mathbf{a})$ is best fit with μ fixed (the constrained maximum likelihood estimator, depends on data)
- and $\hat{\nu}$ and $\hat{\mu}$ are best fit with both left floating (unconstrained)
- Tevatron used $Q_{Tev} = \lambda(\mu=1)/\lambda(\mu=0)$ as generalization of Q_{LEP}

An example

Essentially, you need to fit your model to the data twice:
once with everything floating, and once with signal fixed to 0

$$\lambda(\mu = 0) = \frac{P(\mathbf{m}, \mathbf{a}|\mu = 0, \hat{\nu}(\mu = 0; \mathbf{m}, \mathbf{a}))}{P(\mathbf{m}, \mathbf{a}|\hat{\mu}, \hat{\nu})}$$



After a close look at the profile likelihood ratio

$$\lambda(\mu) = \frac{P(\mathbf{m}, \mathbf{a} | \mu, \hat{\nu}(\mu; \mathbf{m}, \mathbf{a}))}{P(\mathbf{m}, \mathbf{a} | \hat{\mu}, \hat{\nu})}$$

one can see the function is independent of true values of ν

- ▶ though its distribution might depend indirectly

Wilks's theorem states that under certain conditions the distribution of $-2 \ln \lambda(\mu = \mu_0)$ given that the true value of μ is μ_0 converges to a chi-square distribution

- ▶ more on this tomorrow, but the important points are:
- ▶ "asymptotic distribution" is known and it is independent of ν
 - more complicated if parameters have boundaries (eg. $\mu \geq 0$)

Thus, we can calculate the p-value for the background-only hypothesis without having to generate Toy Monte Carlo!

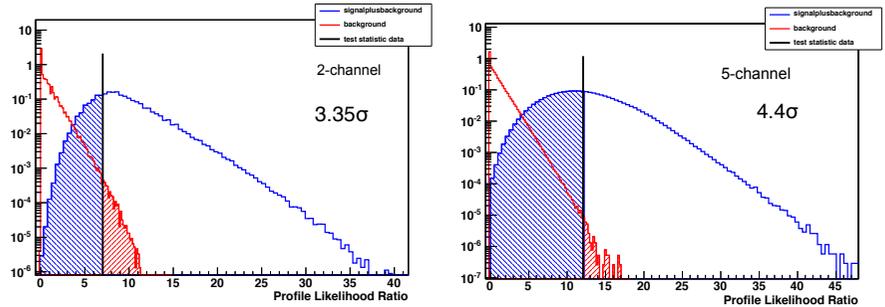
Explicitly build distribution by generating "toys" / pseudo experiments assuming a specific value of μ and ν .

- ▶ randomize both main measurement \mathbf{m} and auxiliary measurements \mathbf{a}
- ▶ fit the model twice for the numerator and denominator of profile likelihood ratio
- ▶ evaluate $-2 \ln \lambda(\mu)$ and add to histogram

Choice of μ is straight forward: typically $\mu=0$ and $\mu=1$, but choice of ν is less clear

- ▶ more on this tomorrow

This can be very time consuming. Plots below use millions of toy pseudo-experiments on a model with ~ 50 parameters.



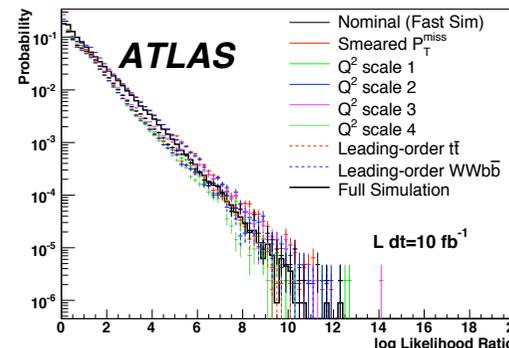
To describe a statistical method, you should clearly specify

- choice of a test statistic
 - simple likelihood ratio (LEP) $Q_{LEP} = L_{s+b}(\mu = 1) / L_b(\mu = 0)$
 - ratio of profiled likelihoods (Tevatron) $Q_{TEV} = L_{s+b}(\mu = 1, \hat{\nu}) / L_b(\mu = 0, \hat{\nu}')$
 - profile likelihood ratio (LHC) $\lambda(\mu) = L_{s+b}(\mu, \hat{\nu}) / L_{s+b}(\hat{\mu}, \hat{\nu})$
- how you build the distribution of the test statistic
 - toy MC randomizing nuisance parameters according to $\pi(\nu)$
 - aka Bayes-frequentist hybrid, prior-predictive, Cousins-Highland
 - toy MC with nuisance parameters fixed (Neyman Construction)
 - assuming asymptotic distribution (Wilks and Wald, more tomorrow)
- what condition you use for limit or discovery
 - more on this tomorrow

So far this looks a bit like magic. How can you claim that you incorporated your systematic just by fitting the best value of your uncertain parameters and making a ratio?

It won't unless the parametrization is sufficiently flexible.

So check by varying the settings of your simulation, and see if the profile likelihood ratio is still distributed as a chi-square



Here it is pretty stable, but it's not perfect (and this is a log plot, so it hides some pretty big discrepancies)

For the distribution to be independent of the nuisance parameters your parametrization must be sufficiently flexible.

A very important point

If we keep pushing this point to the extreme, the physics problem goes beyond what we can handle practically

The p-values are usually predicated on the assumption that the **true distribution** is in the family of functions being considered

- eg. we have sufficiently flexible models of signal & background to incorporate all systematic effects
- but we don't believe we simulate everything perfectly
- ..and when we parametrize our models usually we have further approximated our simulation.
 - nature -> simulation -> parametrization

At some point these approaches are limited by honest systematics uncertainties (not statistical ones). Statistics can only help us so much after this point. Now we must be physicists!