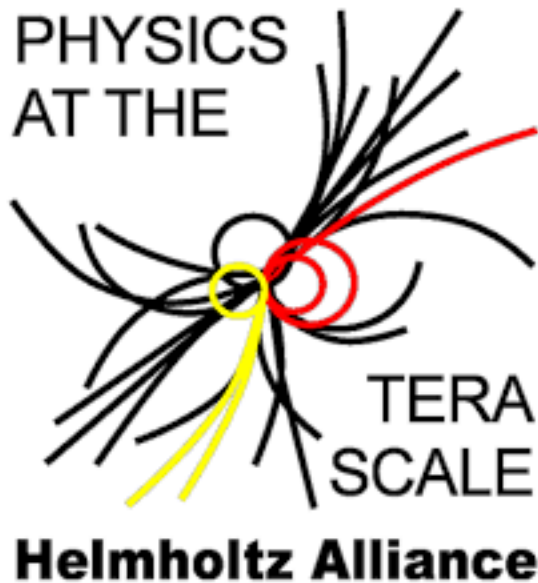


# Statistics for Particle Physics

## Lecture 2



Terascale Statistics School

<https://indico.desy.de/event/46667/>

DESY, Hamburg  
24-28 Feb 2025



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# Outline

## Lectures/tutorials from me:

Monday 15:30 Hypothesis testing

Monday 16:30 Hypothesis testing tutorial

Tuesday 10:30 Parameter estimation tutorial

→ Tuesday 14:00 Setting limits

Wednesday 9:00 Bayesian parameter estimation

Wednesday 11:00 Errors on errors

# Confidence intervals by inverting a test

In addition to a ‘point estimate’ of a parameter we should report an interval reflecting its statistical uncertainty.

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

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

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

Now invert the test to define a confidence interval as:

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

# Relation between confidence interval and $p$ -value

Equivalently we can consider a significance test for each hypothesized value of  $\theta$ , resulting in a  $p$ -value,  $p_\theta$ .

If  $p_\theta \leq \alpha$ , then we reject  $\theta$ .

The confidence interval at  $CL = 1 - \alpha$  consists of those values of  $\theta$  that are not rejected.

E.g. an upper limit on  $\theta$  is the greatest value for which  $p_\theta > \alpha$ .

In practice find by setting  $p_\theta = \alpha$  and solve for  $\theta$ .

For a multidimensional parameter space  $\theta = (\theta_1, \dots, \theta_M)$  use same idea – result is a confidence “region” with boundary determined by  $p_\theta = \alpha$ .

# Coverage probability of confidence interval

If the true value of  $\theta$  is rejected, then it's not in the confidence interval. The probability for this is by construction (equality for continuous data):

$$P(\text{reject } \theta | \theta) \leq \alpha = \text{type-I error rate}$$

Therefore, the probability for the interval to contain or “cover”  $\theta$  is

$$P(\text{conf. interval “covers” } \theta | \theta) \geq 1 - \alpha$$

This assumes that the set of  $\theta$  values considered includes the true value, i.e., it assumes the composite hypothesis  $P(x|H, \theta)$ .

# Frequentist upper limit on Poisson parameter

Consider again the case of observing  $n \sim \text{Poisson}(s + b)$ .

Suppose  $b = 4.5$ ,  $n_{\text{obs}} = 5$ . Find upper limit on  $s$  at 95% CL.

Relevant alternative is  $s = 0$  (critical region at low  $n$ )

$p$ -value of hypothesized  $s$  is  $P(n \leq n_{\text{obs}}; s, b)$

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

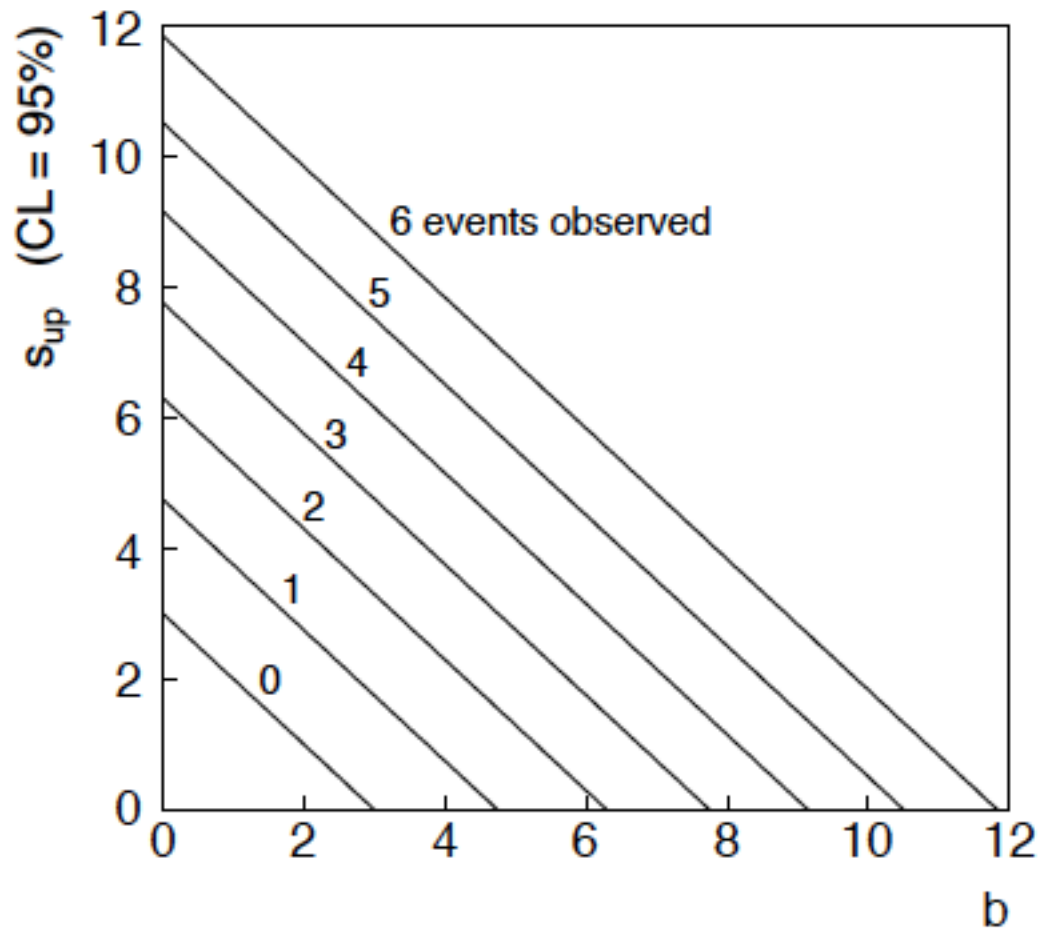
$$\alpha = P(n \leq n_{\text{obs}}; s_{\text{up}}, b) = \sum_{n=0}^{n_{\text{obs}}} \frac{(s_{\text{up}} + b)^n}{n!} e^{-(s_{\text{up}} + b)}$$

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

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

$n \sim \text{Poisson}(s+b)$ : frequentist upper limit on  $s$

For low fluctuation of  $n$ , formula can give negative result for  $s_{\text{up}}$ ; i.e. confidence interval is empty; all values of  $s \geq 0$  have  $p_s \leq \alpha$ .



# Limits near a boundary of the parameter space

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

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

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

Physicist:

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

Statistician:

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

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



# Expected limit for $s = 0$

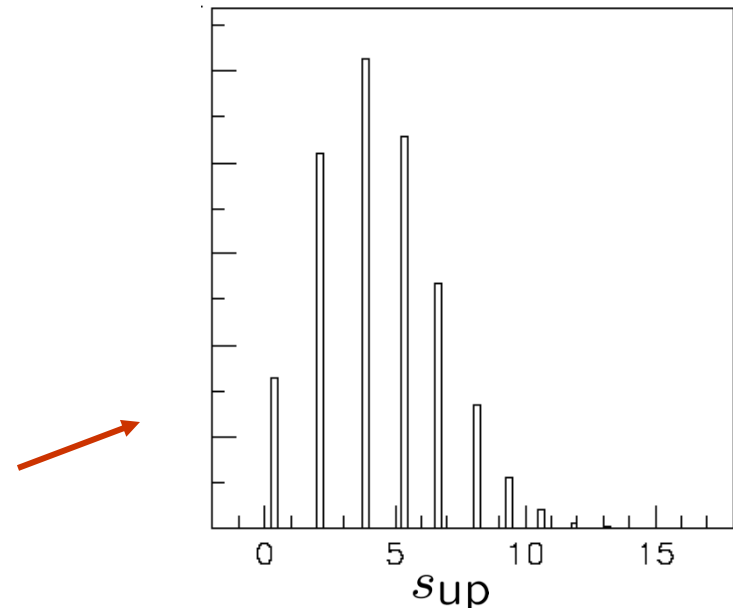
Physicist: I should have used  $\text{CL} = 0.95$  — then  $s_{\text{up}} = 0.496$

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

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

Look at the mean limit for the no-signal hypothesis ( $s = 0$ ) (sensitivity).

Distribution of 95% CL limits  
with  $b = 2.5$ ,  $s = 0$ .  
Mean upper limit = 4.44



# The Bayesian approach to limits

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

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

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

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

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

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

# Bayesian prior for Poisson parameter

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

Could try to reflect ‘prior ignorance’ with e.g.

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

Not normalized; can be OK provided  $p(n|s)$  dies off quickly for large  $s$ .

Not invariant under change of parameter — if we had used instead a flat prior for a nonlinear function of  $s$ , then this would imply a non-flat prior for  $s$ .

Doesn’t really reflect a reasonable degree of belief, but often used as a point of reference; or viewed as a recipe for producing an interval whose frequentist properties can be studied (e.g., coverage probability, which will depend on true  $s$ ).

# Bayesian upper limit with flat prior for $s$

Put Poisson likelihood and flat prior into Bayes' theorem:

$$p(s|n) \propto p(n|s)\pi(s) = \frac{(s+b)^n}{n!} e^{-(s+b)} \times 1, \quad s \geq 0$$

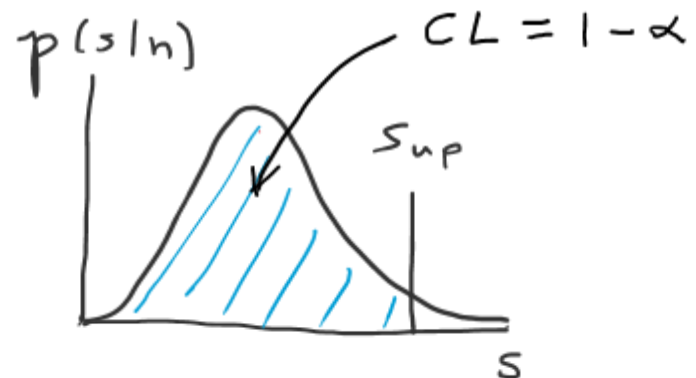
Normalize to unit area:

$$p(s|n) = \frac{(s+b)^n e^{-(s+b)}}{\Gamma(b, n+1)}$$

upper incomplete  
gamma function

Upper limit  $s_{\text{up}}$  determined by

$$1 - \alpha = \int_0^{s_{\text{up}}} p(s|n) ds$$



# Bayesian interval with flat prior for $s$

Solve to find limit  $s_{\text{up}}$ :

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

where

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

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

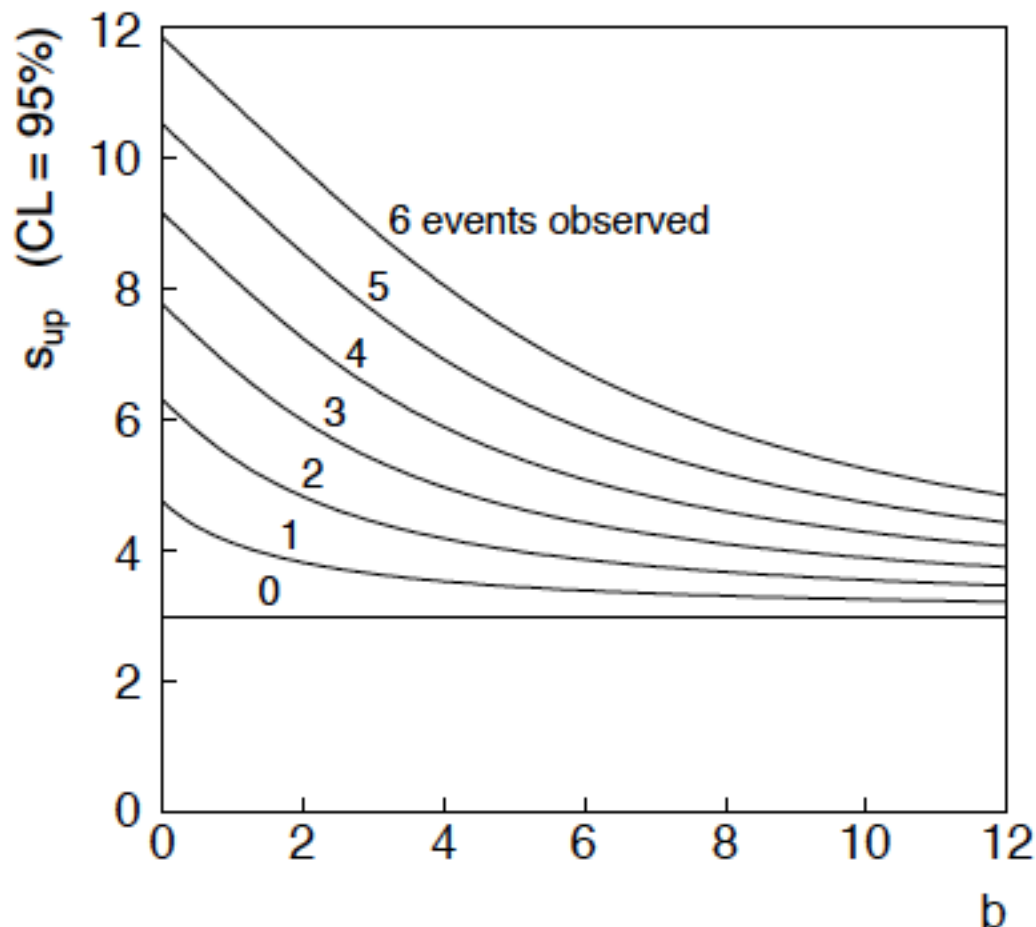
# Bayesian interval with flat prior for $s$

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

For  $b = 0$ , Bayesian and frequentist upper limits come out equal.

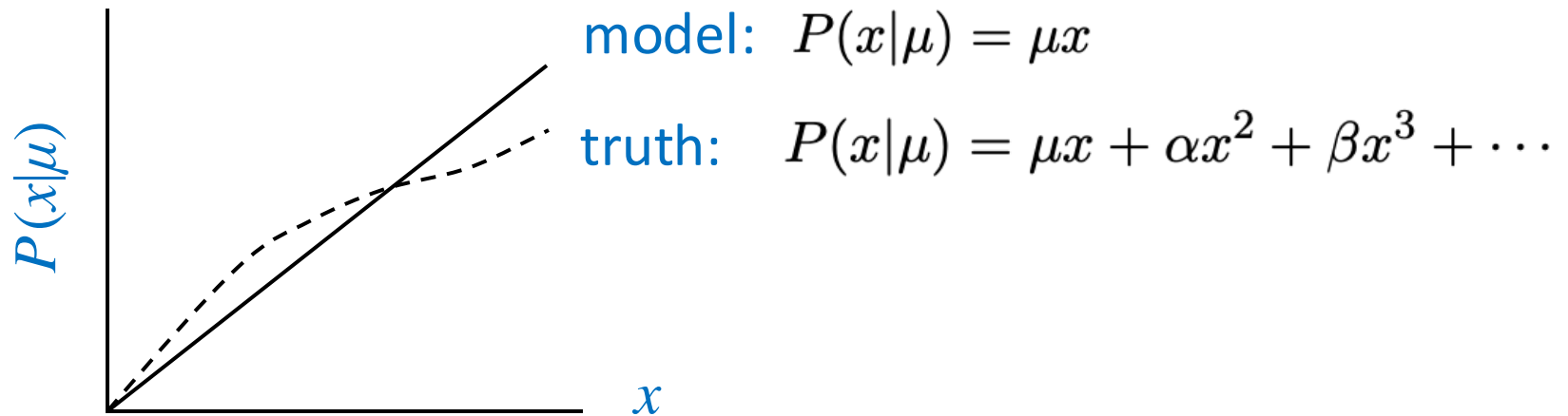
Never goes negative.

Doesn't depend on  $b$  if  $n = 0$ .



# Systematic uncertainties and nuisance parameters

In general, our model of the data is not perfect:



Can improve model by including additional adjustable parameters.

$$P(x|\mu) \rightarrow P(x|\mu, \boldsymbol{\theta})$$

Nuisance parameter  $\leftrightarrow$  systematic uncertainty. Some point in the parameter space of the enlarged model should be “true”.

Presence of nuisance parameter decreases sensitivity of analysis to the parameter of interest (e.g., increases variance of estimate).

# Profile Likelihood

Suppose we have a likelihood  $L(\boldsymbol{\mu}, \boldsymbol{\theta}) = P(\mathbf{x}|\boldsymbol{\mu}, \boldsymbol{\theta})$  with  $N$  parameters of interest  $\boldsymbol{\mu} = (\mu_1, \dots, \mu_N)$  and  $M$  nuisance parameters  $\boldsymbol{\theta} = (\theta_1, \dots, \theta_M)$ . The “profiled” (or “constrained”) values of  $\boldsymbol{\theta}$  are:

$$\hat{\hat{\boldsymbol{\theta}}}(\boldsymbol{\mu}) = \operatorname{argmax}_{\boldsymbol{\theta}} L(\boldsymbol{\mu}, \boldsymbol{\theta})$$

and the profile likelihood is:  $L_p(\boldsymbol{\mu}) = L(\boldsymbol{\mu}, \hat{\hat{\boldsymbol{\theta}}})$

The profile likelihood depends only on the parameters of interest; the nuisance parameters are replaced by their profiled values.

The profile likelihood can be used to obtain confidence intervals/regions for the parameters of interest in the same way as one would for all of the parameters from the full likelihood.



# Profile Likelihood Ratio – Wilks theorem

Goal is to test/reject regions of  $\mu$  space (param. of interest).

Rejecting a point  $\mu$  should mean  $p_\mu \leq \alpha$  for all possible values of the nuisance parameters  $\theta$ .

Test  $\mu$  using the “profile likelihood ratio”:  $\lambda(\mu) = \frac{L(\mu, \hat{\theta})}{L(\hat{\mu}, \hat{\theta})}$

Let  $t_\mu = -2\ln\lambda(\mu)$ . Wilks’ theorem says in large-sample limit:

$$t_\mu \sim \text{chi-square}(N)$$

where the number of degrees of freedom is the number of parameters of interest (components of  $\mu$ ). So  $p$ -value for  $\mu$  is

$$p_\mu = \int_{t_{\mu,\text{obs}}}^{\infty} f(t_\mu | \mu, \theta) dt_\mu = 1 - F_{\chi_N^2}(t_{\mu,\text{obs}})$$

## Profile Likelihood Ratio – Wilks theorem (2)

Set  $p_{\mu} = \alpha$  and solve for  $t_{\mu}$  (here means observed value)

$$t_{\mu} = F_{\chi_N^2}^{-1}(1 - \alpha) \quad \text{which is also} \quad t_{\mu} = -2 \ln \frac{L_p(\mu)}{L(\hat{\mu}, \hat{\theta})}$$

Equate and solve for  $\ln L_p(\mu)$  gives the recipe to get confidence regions/intervals for the parameters of interest at  $CL = 1 - \alpha$ :

$$\ln L_p(\mu) = \ln L_{\max} - \frac{1}{2} F_{\chi_N^2}^{-1}(1 - \alpha)$$

where the number of degrees of freedom  $N$  for the chi-square quantile is equal to the number of parameters of interest.

If the large-sample limit is not justified, then use e.g. Monte Carlo to get distribution of  $t_{\mu}$ .

# Prototype search analysis

Search for signal in a region of phase space; result is histogram of some variable  $x$  giving numbers:

$$\mathbf{n} = (n_1, \dots, n_N)$$

Assume the  $n_i$  are Poisson distributed with expectation values

$$E[n_i] = \mu s_i + b_i$$

strength parameter

where

$$s_i = s_{\text{tot}} \int_{\text{bin } i} f_s(x; \boldsymbol{\theta}_s) dx, \quad b_i = b_{\text{tot}} \int_{\text{bin } i} f_b(x; \boldsymbol{\theta}_b) dx.$$

signal

background

## Prototype analysis (II)

Often also have a subsidiary measurement that constrains some of the background and/or shape parameters:

$$\mathbf{m} = (m_1, \dots, m_M)$$

Assume the  $m_i$  are Poisson distributed with expectation values

$$E[m_i] = u_i(\boldsymbol{\theta})$$

 nuisance parameters ( $\theta_s, \theta_b, b_{\text{tot}}$ )

Likelihood function is

$$L(\mu, \boldsymbol{\theta}) = \prod_{j=1}^N \frac{(\mu s_j + b_j)^{n_j}}{n_j!} e^{-(\mu s_j + b_j)} \prod_{k=1}^M \frac{u_k^{m_k}}{m_k!} e^{-u_k}$$

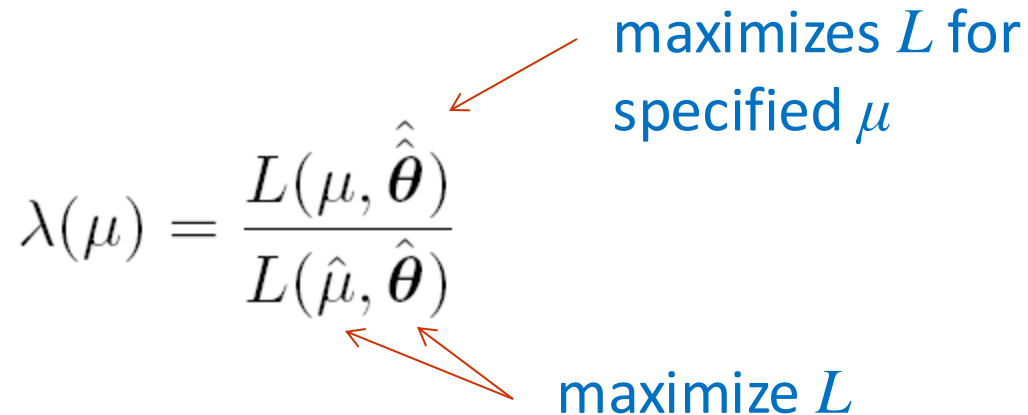
# The profile likelihood ratio

Base significance test on the profile likelihood ratio:

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

maximizes  $L$  for specified  $\mu$

maximize  $L$

The diagram shows the formula for the profile likelihood ratio, λ(μ) = L(μ, θ̂̂) / L(μ̂, θ̂). Two red arrows point from the text 'maximizes L for specified μ' to the θ̂̂ in the numerator. Two red arrows point from the text 'maximize L' to the μ̂ and θ̂ in the denominator.

Define critical region of test of  $\mu$  by the region of data space that gives the lowest values of  $\lambda(\mu)$ .

Important advantage of profile LR is that its distribution becomes independent of nuisance parameters in large sample limit.

# Test statistic for discovery

Suppose relevant alternative to background-only ( $\mu = 0$ ) is  $\mu \geq 0$ .

So take critical region for test of  $\mu = 0$  corresponding to high  $q_0$  and  $\hat{m} > 0$  (data characteristic for  $\mu \geq 0$ ).

That is, to test background-only hypothesis define statistic

$$q_0 = \begin{cases} -2 \ln \lambda(0) & \hat{\mu} \geq 0 \\ 0 & \hat{\mu} < 0 \end{cases}$$

i.e. here only large (positive) observed signal strength is evidence against the background-only hypothesis.

Note that even though here physically  $\mu \geq 0$ , we allow  $\hat{m}$  to be negative. In large sample limit its distribution becomes Gaussian, and this will allow us to write down simple expressions for distributions of our test statistics.

## Distribution of $q_0$ in large-sample limit

Assuming approximations valid in the large sample (asymptotic) limit, we can write down the full distribution of  $q_0$  as

$$f(q_0|\mu') = \left(1 - \Phi\left(\frac{\mu'}{\sigma}\right)\right) \delta(q_0) + \frac{1}{2} \frac{1}{\sqrt{2\pi}} \frac{1}{\sqrt{q_0}} \exp\left[-\frac{1}{2} \left(\sqrt{q_0} - \frac{\mu'}{\sigma}\right)^2\right]$$

The special case  $\mu' = 0$  is a “half chi-square” distribution:

$$f(q_0|0) = \frac{1}{2} \delta(q_0) + \frac{1}{2} \frac{1}{\sqrt{2\pi}} \frac{1}{\sqrt{q_0}} e^{-q_0/2}$$

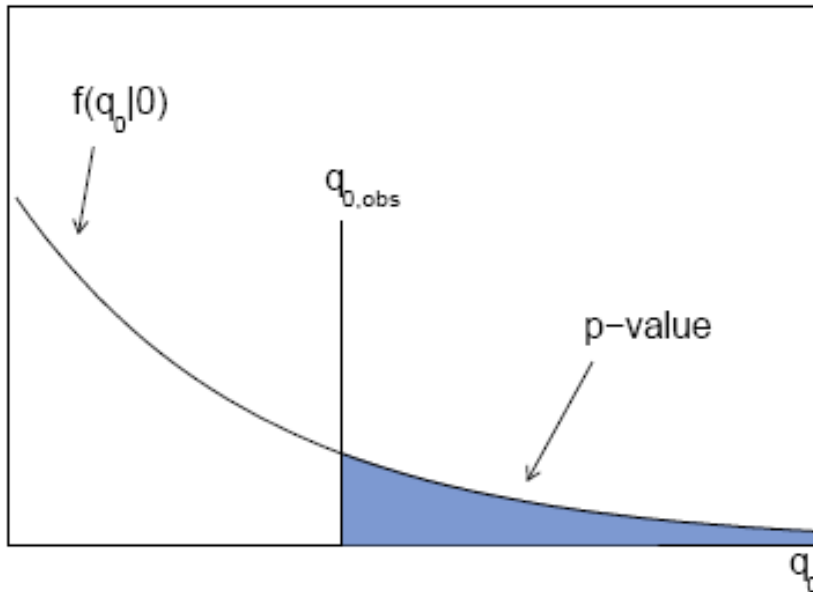
In large sample limit,  $f(q_0|0)$  independent of nuisance parameters;  $f(q_0|\mu')$  depends on nuisance parameters through  $\sigma$ .

# $p$ -value for discovery

Large  $q_0$  means increasing incompatibility between the data and hypothesis, therefore  $p$ -value for an observed  $q_{0,\text{obs}}$  is

$$p_0 = \int_{q_{0,\text{obs}}}^{\infty} f(q_0|0) dq_0$$

use e.g. asymptotic formula



From  $p$ -value get equivalent significance,

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



# Cumulative distribution of $q_0$ , significance

From the pdf, the cumulative distribution of  $q_0$  is found to be

$$F(q_0|\mu') = \Phi\left(\sqrt{q_0} - \frac{\mu'}{\sigma}\right)$$

The special case  $\mu' = 0$  is

$$F(q_0|0) = \Phi(\sqrt{q_0})$$

The  $p$ -value of the  $\mu = 0$  hypothesis is

$$p_0 = 1 - F(q_0|0)$$

Therefore the discovery significance  $Z$  is simply

$$Z = \Phi^{-1}(1 - p_0) = \sqrt{q_0}$$

# Monte Carlo test of asymptotic formula

$$n \sim \text{Poisson}(\mu s + b)$$

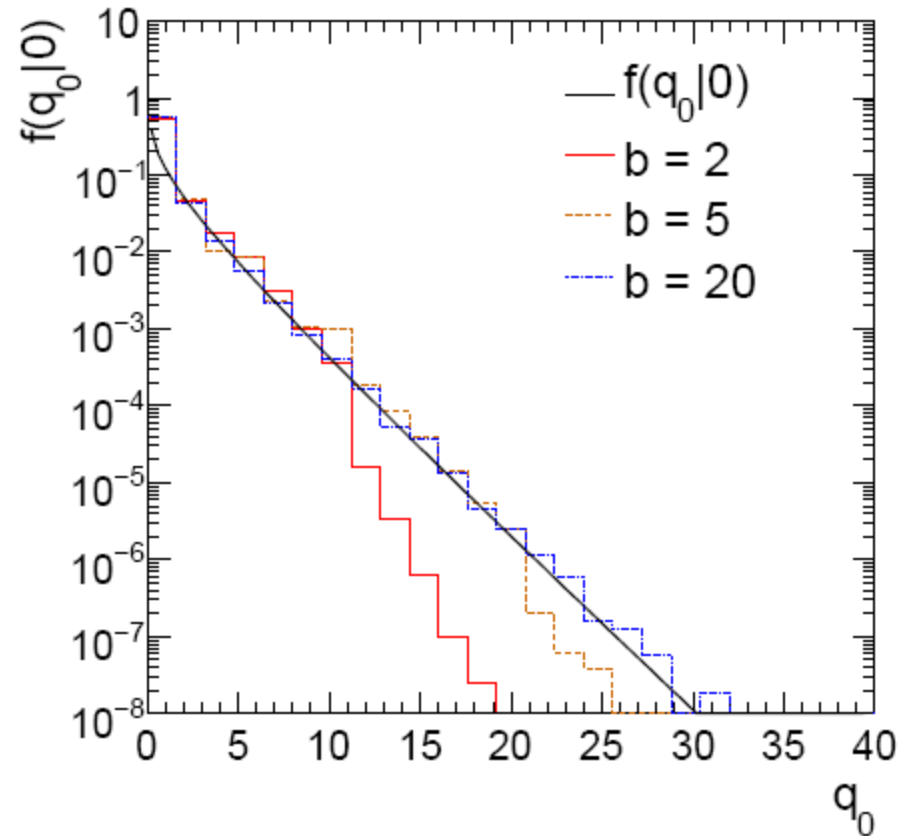
$$m \sim \text{Poisson}(\tau b)$$

$\mu$  = param. of interest

$b$  = nuisance parameter

Here take  $s$  known,  $\tau = 1$ .

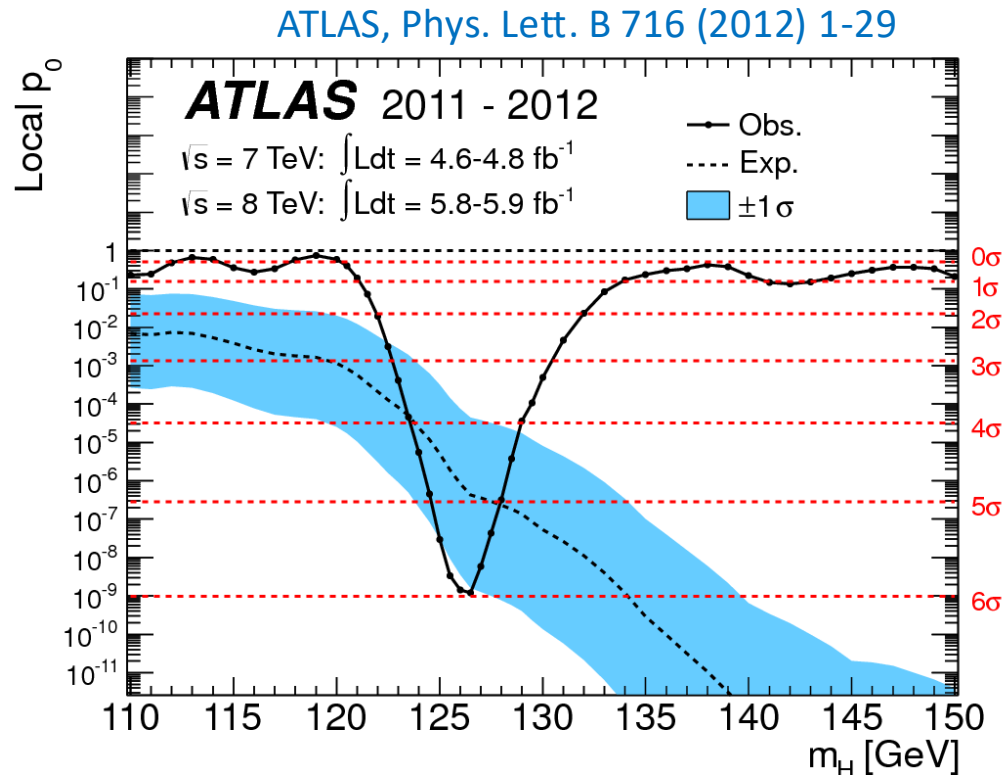
Asymptotic formula is good approximation to  $5\sigma$  level ( $q_0 = 25$ ) already for  $b \sim 20$ .



# How to read the $p_0$ plot

The “local”  $p_0$  means the  $p$ -value of the background-only hypothesis obtained from the test of  $\mu = 0$  at each individual  $m_H$ , without any correct for the Look-Elsewhere Effect.

The “Expected” (dashed) curve gives the median  $p_0$  under assumption of the SM Higgs ( $\mu = 1$ ) at each  $m_H$ .



The blue band gives the width of the distribution ( $\pm 1\sigma$ ) of significances under assumption of the SM Higgs.

# Test statistic for upper limits

For purposes of setting an upper limit on  $\mu$  use

$$q_\mu = \begin{cases} -2 \ln \lambda(\mu) & \hat{\mu} \leq \mu \\ 0 & \hat{\mu} > \mu \end{cases} \quad \text{where} \quad \lambda(\mu) = \frac{L(\mu, \hat{\boldsymbol{\theta}})}{L(\hat{\mu}, \hat{\boldsymbol{\theta}})}$$

I.e. when setting an upper limit, an upwards fluctuation of the data is not taken to mean incompatibility with the hypothesized  $\mu$  :

From observed  $q_\mu$  find  $p$ -value:  $p_\mu = \int_{q_{\mu, \text{obs}}}^{\infty} f(q_\mu | \mu) dq_\mu$

Large sample approximation:

$$p_\mu = 1 - \Phi(\sqrt{q_\mu})$$

To find upper limit at  $\text{CL} = 1 - \alpha$ , set  $p_\mu = \alpha$  and solve for  $\mu$ .

# Monte Carlo test of asymptotic formulae

Consider again  $n \sim \text{Poisson}(\mu s + b)$ ,  $m \sim \text{Poisson}(\tau b)$

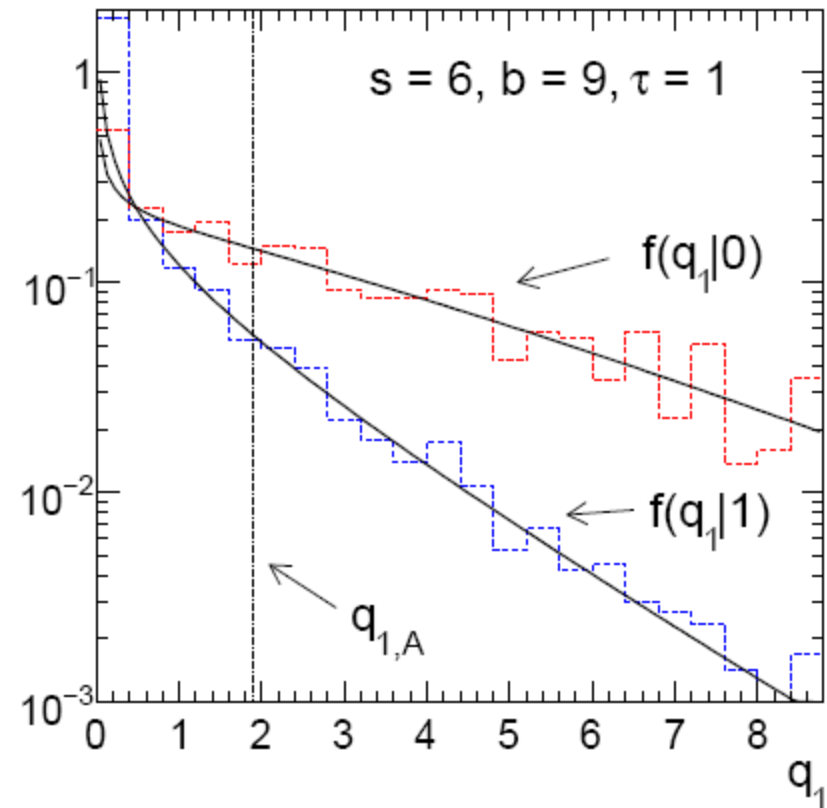
Use  $q_\mu$  to find  $p$ -value of hypothesized  $\mu$  values.

E.g.  $f(q_1|1)$  for  $p$ -value of  $\mu = 1$ .

Typically interested in 95% CL, i.e.,  
 $p$ -value threshold = 0.05, i.e.,  
 $q_1 = 2.69$  or  $Z_1 = \sqrt{q_1} = 1.64$ .

Median[ $q_1 | 0$ ] gives “exclusion sensitivity”.

Here asymptotic formulae good  
 for  $s = 6$ ,  $b = 9$ .

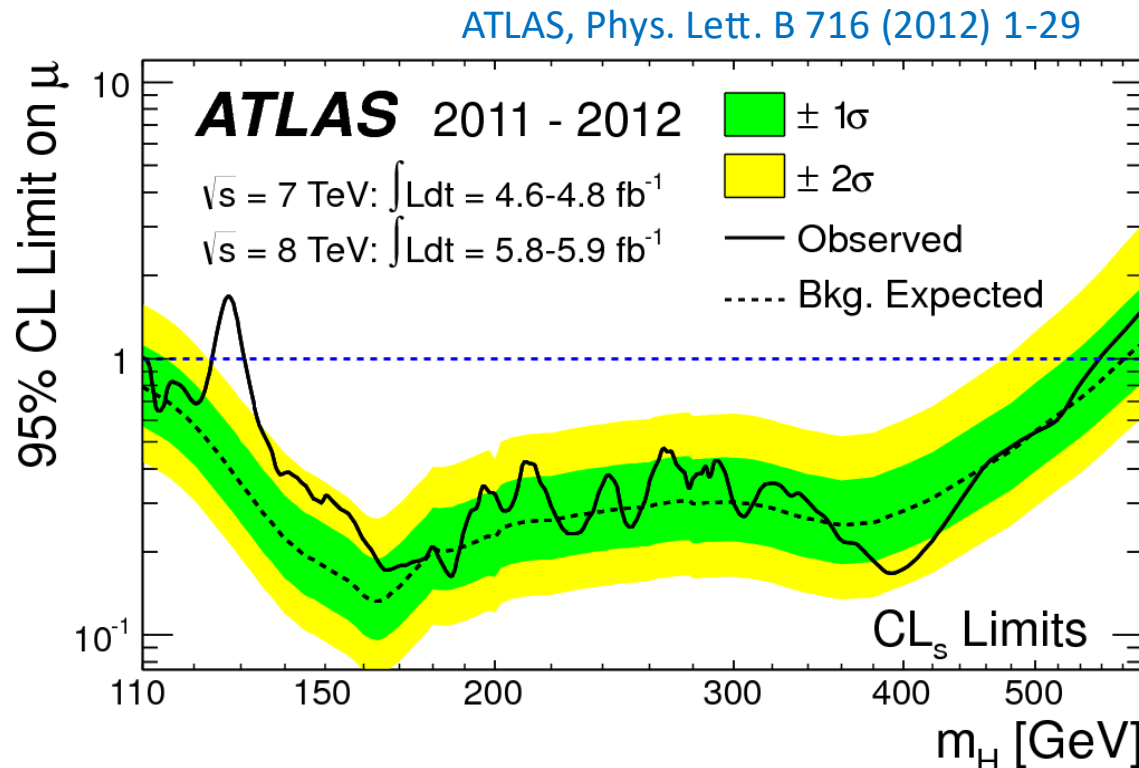


# How to read the green and yellow limit plots

For every value of  $m_H$ , find the upper limit on  $\mu$ .

Also for each  $m_H$ , determine the distribution of upper limits  $\mu_{\text{up}}$  one would obtain under the hypothesis of  $\mu = 0$ .

The dashed curve is the median  $\mu_{\text{up}}$ , and the green (yellow) bands give the  $\pm 1\sigma$  ( $2\sigma$ ) regions of this distribution.



# Extra slides

# Approximate confidence intervals/regions from the likelihood function

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

$$\lambda(\theta) = \frac{L(\theta)}{L(\hat{\theta})} \quad 0 \leq \lambda(\theta) \leq 1$$

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

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

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

$p$ -value of  $\theta$  therefore

$$p_\theta = \int_{t_{\theta, \text{obs}}}^{\infty} f(t_\theta | \theta) dt_\theta$$

need pdf



# Confidence region from Wilks' theorem

Wilks' theorem says (in large-sample limit and provided certain conditions hold...)

$$f(t_\theta|\theta) \sim \chi_n^2$$

chi-square dist. with # d.o.f. =  
# of components in  $\theta = (\theta_1, \dots, \theta_n)$ .

Assuming this holds, the  $p$ -value is

$$p_\theta = 1 - F_{\chi_n^2}(t_\theta) \quad \leftarrow \text{set equal to } \alpha$$

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

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

Recall also

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

# Confidence region from Wilks' theorem (cont.)

i.e., boundary of confidence region in  $\theta$  space is where

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

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

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

and so the 68.3% confidence level interval is determined by

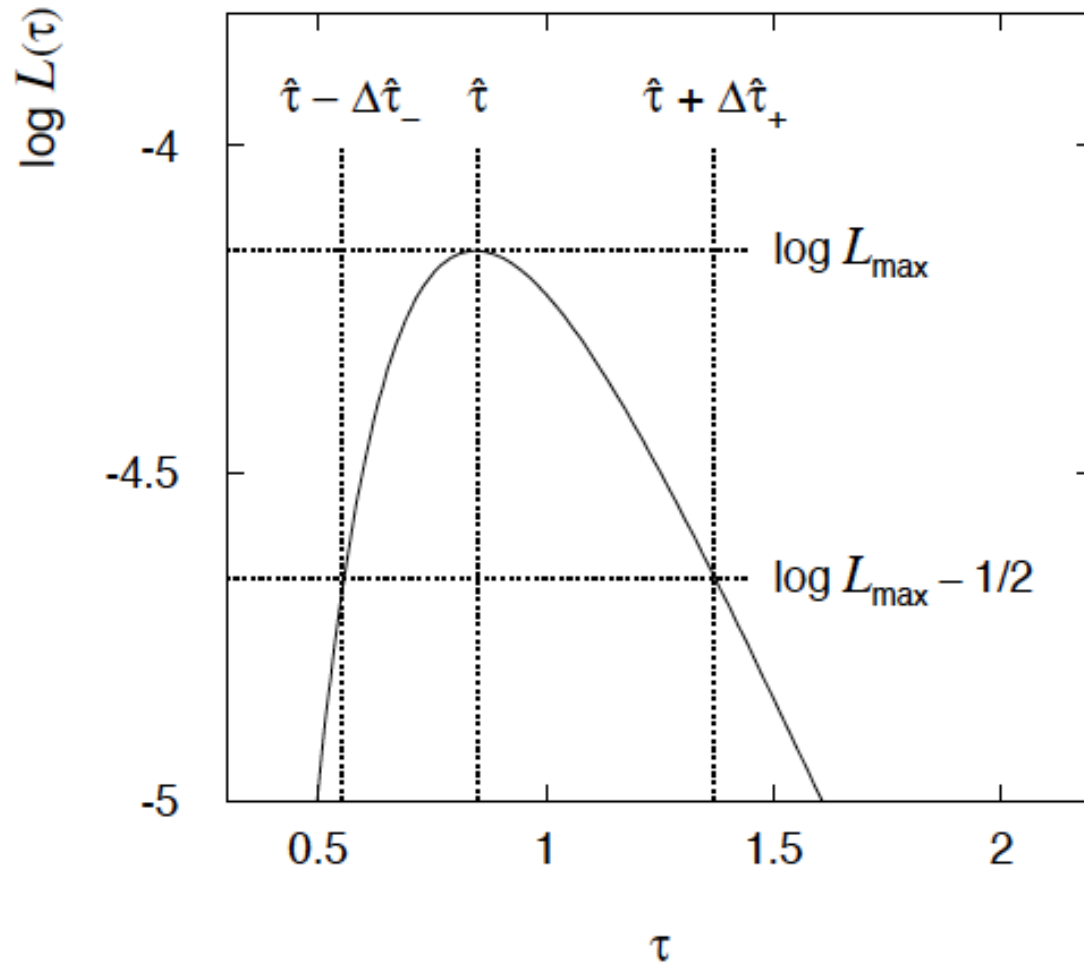
$$\ln L(\theta) = \ln L(\hat{\theta}) - \frac{1}{2}$$

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

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

# Example of interval from $\ln L(\theta)$

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



Our exponential example, now with only  $n = 5$  events.

Can report ML estimate with approx. confidence interval from  $\ln L_{\max} - 1/2$  as “asymmetric error bar”:

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

# Multiparameter case

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

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

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

# Multiparameter case (cont.)

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

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

# Expected significance (sensitivity)

To characterize sensitivity to discovery, give expected (mean or median)  $Z$  under assumption of a given  $s$ .

For large  $s + b$ ,  $n \rightarrow x \sim \text{Gaussian}(\mu, \sigma)$ ,  $\mu = s + b$ ,  $\sigma = \sqrt{s + b}$ .

For observed value  $x_{\text{obs}}$ ,  $p$ -value of  $s = 0$  is  $\text{Prob}(x > x_{\text{obs}} \mid s = 0), :$

$$p_0 = 1 - \Phi\left(\frac{x_{\text{obs}} - b}{\sqrt{b}}\right)$$

Significance for rejecting  $s = 0$  is therefore

$$Z_0 = \Phi^{-1}(1 - p_0) = \frac{x_{\text{obs}} - b}{\sqrt{b}}$$

Expected (median) significance assuming signal rate  $s$  is

$$\text{median}[Z_0 \mid s + b] = \frac{s}{\sqrt{b}}$$

# Better approximation for significance

Poisson likelihood for parameter  $s$  is

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

For now  
no nuisance  
params.

To test for discovery use profile likelihood ratio:

$$q_0 = \begin{cases} -2 \ln \lambda(0) & \hat{s} \geq 0, \\ 0 & \hat{s} < 0. \end{cases} \quad \lambda(s) = \frac{L(s, \hat{\hat{\theta}}(s))}{L(\hat{s}, \hat{\theta})}$$

So the likelihood ratio statistic for testing  $s = 0$  is

$$q_0 = -2 \ln \frac{L(0)}{L(\hat{s})} = 2 \left( n \ln \frac{n}{b} + b - n \right) \quad \text{for } n > b, \quad 0 \text{ otherwise}$$

# Approximate Poisson significance (continued)

For sufficiently large  $s + b$ , (use Wilks' theorem),

$$Z = \sqrt{2 \left( n \ln \frac{n}{b} + b - n \right)} \quad \text{for } n > b \text{ and } Z = 0 \text{ otherwise.}$$

To find  $\text{median}[Z|s]$ , let  $n \rightarrow s + b$  (i.e., the Asimov data set):

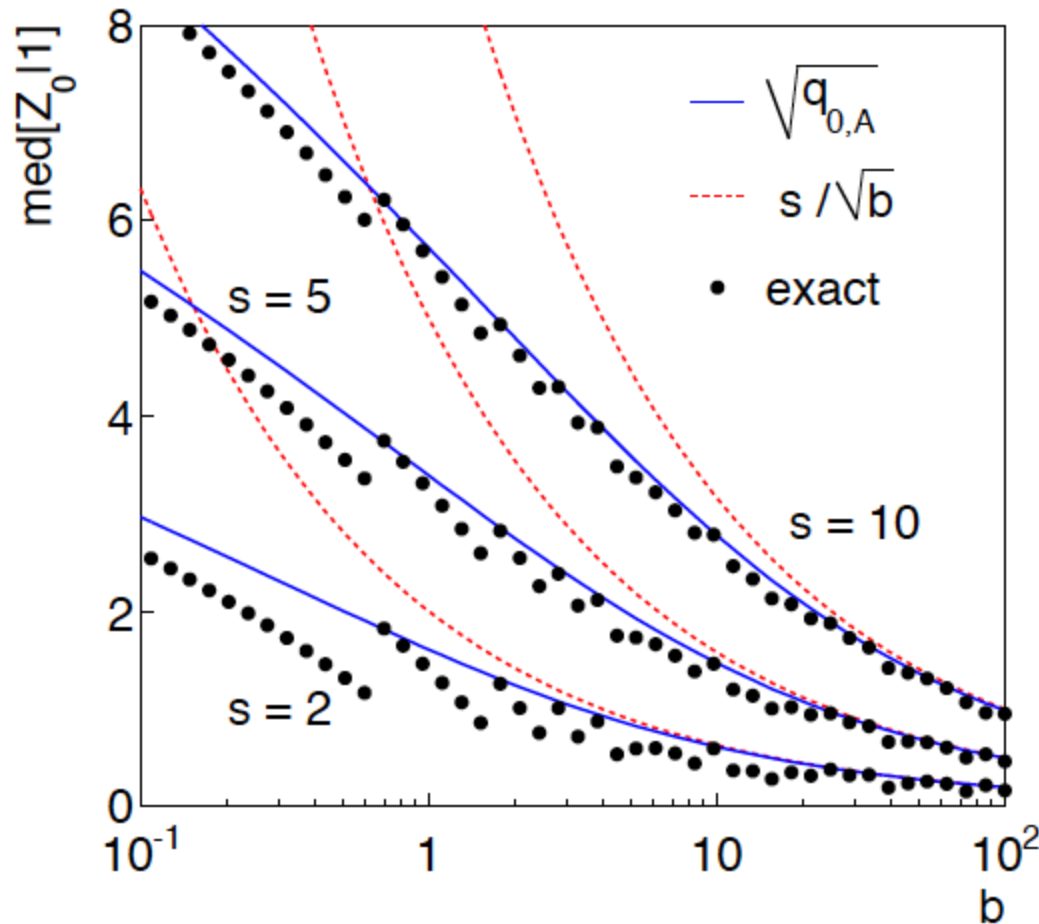
$$Z_A = \sqrt{2 \left( (s + b) \ln \left( 1 + \frac{s}{b} \right) - s \right)}$$

This reduces to  $s/\sqrt{b}$  for  $s \ll b$ .



$n \sim \text{Poisson}(s+b)$ , median significance,  
assuming  $s$ , of the hypothesis  $s = 0$

CCGV, EPJC 71 (2011) 1554, arXiv:1007.1727



“Exact” values from MC,  
jumps due to discrete data.

Asimov  $\sqrt{q_{0,A}}$  good approx.  
for broad range of  $s, b$ .

$s/\sqrt{b}$  only good for  $s \ll b$ .

# Extending $s/\sqrt{b}$ to case where $b$ uncertain

The intuitive explanation of  $s/\sqrt{b}$  is that it compares the signal,  $s$ , to the standard deviation of  $n$  assuming no signal,  $\sqrt{b}$ .

Now suppose the value of  $b$  is uncertain, characterized by a standard deviation  $\sigma_b$ .

A reasonable guess is to replace  $\sqrt{b}$  by the quadratic sum of  $\sqrt{b}$  and  $\sigma_b$ , i.e.,

$$\text{med}[Z|s] = \frac{s}{\sqrt{b + \sigma_b^2}}$$

This has been used to optimize some analyses e.g. where  $\sigma_b$  cannot be neglected.

# Profile likelihood with $b$ uncertain

This is the well studied “on/off” problem: Cranmer 2005; Cousins, Linnemann, and Tucker 2008; Li and Ma 1983,...

Measure two Poisson distributed values:

$n \sim \text{Poisson}(s+b)$  (primary or “search” measurement)

$m \sim \text{Poisson}(\tau b)$  (control measurement,  $\tau$  known)

The likelihood function is

$$L(s, b) = \frac{(s+b)^n}{n!} e^{-(s+b)} \frac{(\tau b)^m}{m!} e^{-\tau b}$$

Use this to construct profile likelihood ratio ( $b$  is nuisance parameter):

$$\lambda(0) = \frac{L(0, \hat{\hat{b}}(0))}{L(\hat{s}, \hat{b})}$$

# Ingredients for profile likelihood ratio

To construct profile likelihood ratio from this need estimators:

$$\hat{s} = n - m/\tau ,$$

$$\hat{b} = m/\tau ,$$

$$\hat{\hat{b}}(s) = \frac{n + m - (1 + \tau)s + \sqrt{(n + m - (1 + \tau)s)^2 + 4(1 + \tau)sm}}{2(1 + \tau)} .$$

and in particular to test for discovery ( $s = 0$ ),

$$\hat{\hat{b}}(0) = \frac{n + m}{1 + \tau}$$

# Asymptotic significance

Use profile likelihood ratio for  $q_0$ , and then from this get discovery significance using asymptotic approximation (Wilks' theorem):

$$Z = \sqrt{q_0};$$
$$= \left[ -2 \left( n \ln \left[ \frac{n+m}{(1+\tau)n} \right] + m \ln \left[ \frac{\tau(n+m)}{(1+\tau)m} \right] \right) \right]^{1/2}$$

for  $n > \hat{b}$  and  $Z = 0$  otherwise.

Essentially same as in:

Robert D. Cousins, James T. Linnemann and Jordan Tucker, NIM A 595 (2008) 480–501; arXiv:physics/0702156.

Tipei Li and Yuqian Ma, Astrophysical Journal 272 (1983) 317–324.

# Asimov approximation for median significance

To get median discovery significance, replace  $n$ ,  $m$  by their expectation values assuming background-plus-signal model:

$$n \rightarrow s + b$$

$$m \rightarrow \tau b$$

$$Z_A = \left[ -2 \left( (s + b) \ln \left[ \frac{s + (1 + \tau)b}{(1 + \tau)(s + b)} \right] + \tau b \ln \left[ 1 + \frac{s}{(1 + \tau)b} \right] \right) \right]^{1/2}$$

Or use the variance of  $\hat{b} = m/\tau$ ,  $V[\hat{b}] \equiv \sigma_b^2 = \frac{b}{\tau}$ , to eliminate  $\tau$ :

$$Z_A = \left[ 2 \left( (s + b) \ln \left[ \frac{(s + b)(b + \sigma_b^2)}{b^2 + (s + b)\sigma_b^2} \right] - \frac{b^2}{\sigma_b^2} \ln \left[ 1 + \frac{\sigma_b^2 s}{b(b + \sigma_b^2)} \right] \right) \right]^{1/2}$$

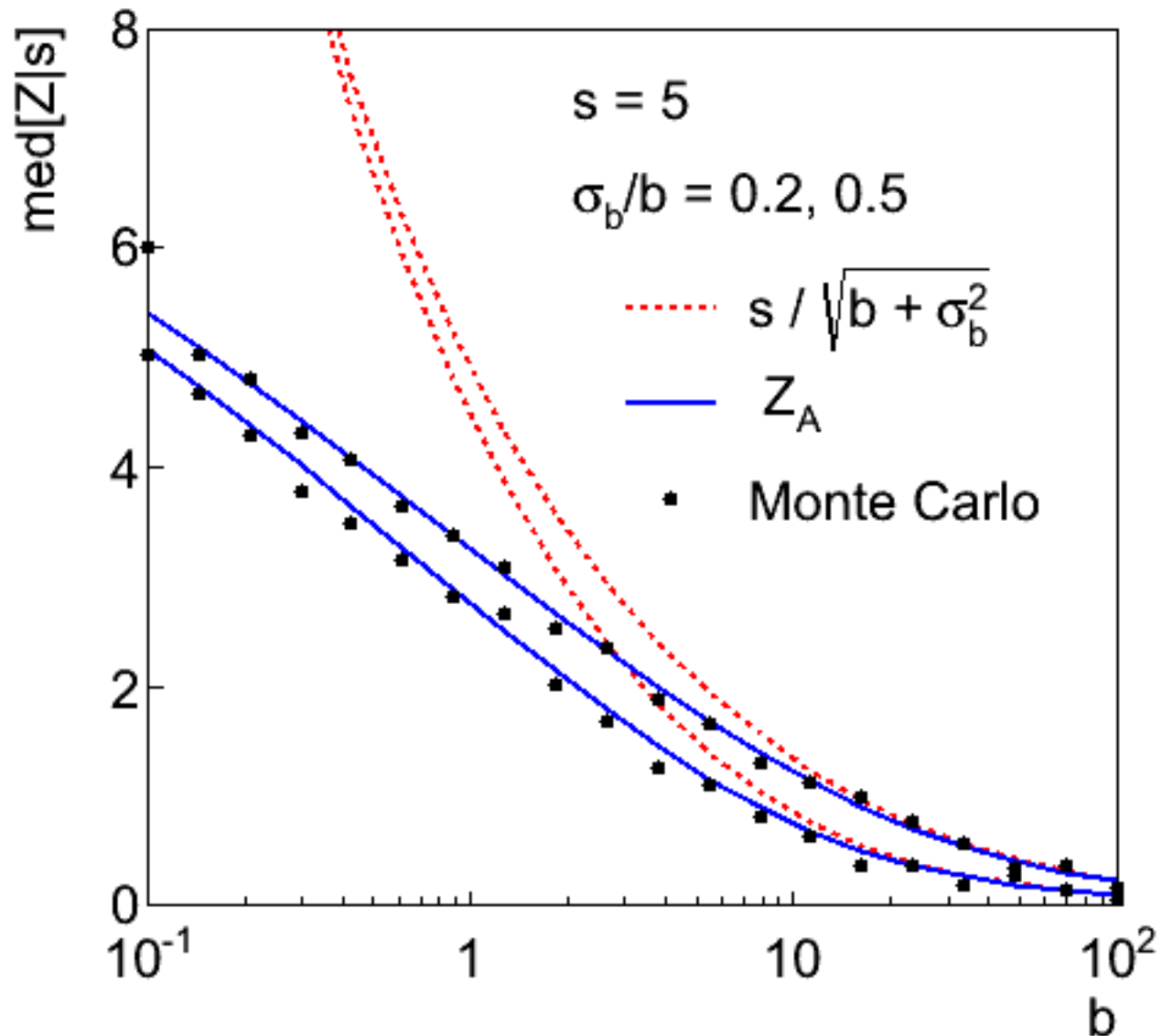
# Limiting cases

Expanding the Asimov formula in powers of  $s/b$  and  $\sigma_b^2/b$  ( $= 1/\tau$ ) gives

$$Z_A = \frac{s}{\sqrt{b + \sigma_b^2}} \left( 1 + \mathcal{O}(s/b) + \mathcal{O}(\sigma_b^2/b) \right)$$

So the “intuitive” formula can be justified as a limiting case of the significance from the profile likelihood ratio test evaluated with the Asimov data set.

## Testing the formulae: $s = 5$





# Using sensitivity to optimize a cut

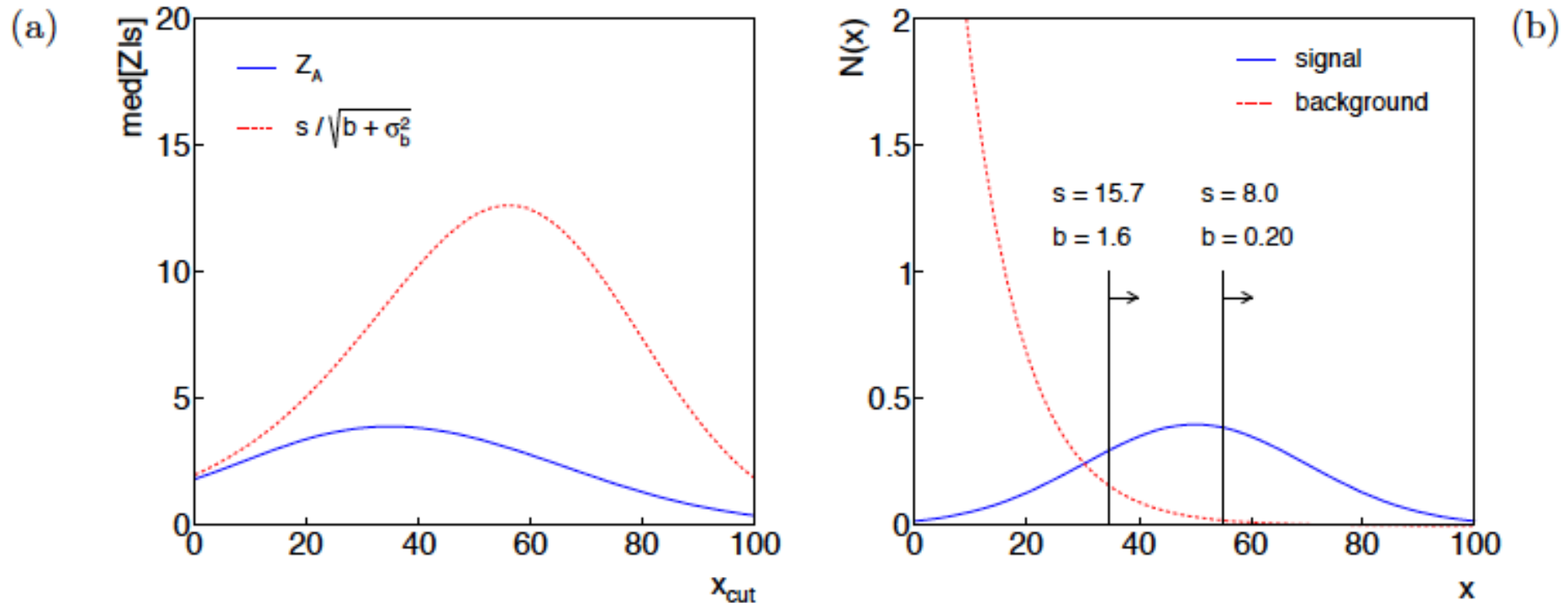


Figure 1: (a) The expected significance as a function of the cut value  $x_{\text{cut}}$ ; (b) the distributions of signal and background with the optimal cut value indicated.

# Summary on discovery sensitivity

Simple formula for expected discovery significance based on profile likelihood ratio test and Asimov approximation:

$$Z_A = \left[ 2 \left( (s + b) \ln \left[ \frac{(s + b)(b + \sigma_b^2)}{b^2 + (s + b)\sigma_b^2} \right] - \frac{b^2}{\sigma_b^2} \ln \left[ 1 + \frac{\sigma_b^2 s}{b(b + \sigma_b^2)} \right] \right) \right]^{1/2}$$

For large  $b$ , all formulae OK.

For small  $b$ ,  $s/\sqrt{b}$  and  $s/\sqrt{(b+\sigma_b^2)}$  overestimate the significance.

Could be important in optimization of searches with low background.

Formula maybe also OK if model is not simple on/off experiment, e.g., several background control measurements (check this).