

Outline

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Statistical Modeling
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Computing statistical results

Discovery

Confidence intervals

Upper limits

Reparameterization and presentation of results

Expected results

Today

Profiling

Bayesian methods

Look elsewhere effect

Highlights: Discovery

Given a statistical model P(data; μ), define likelihood $L(\mu) = P(data; \mu)$

To estimate a parameter, use the value $\hat{\mathbf{p}}$ that maximizes $L(\mu) \rightarrow$ best-fit value

To decide between hypotheses H_0 and H_1 , use the likelihood ratio $\frac{L(H_0)}{L(H_1)}$

To test for **discovery**, use
$$q_0 = -2\log\frac{L(S=0)}{L(\hat{S})}$$
 $\hat{S} \ge 0$

For large enough datasets (n >~ 5), $Z = \sqrt{q_0}$

For a Gaussian measurement,
$$Z = \frac{\hat{S}}{\sqrt{B}}$$

For a Poisson measurement,
$$Z = \sqrt{2\left[(\hat{S} + B) \log \left(1 + \frac{\hat{S}}{B} \right) - \hat{S} \right]}$$

Highlights: Confidence intervals

 $\mu^{+\delta\mu^{+}}_{-\delta\mu^{-}}$ μ^{*}

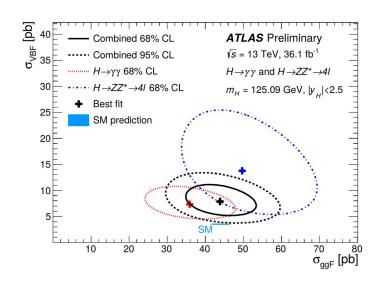
Contain the true value with given probability

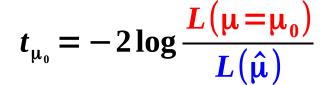
To obtain, compute the log-likelihood ratio as a function of μ_0 .

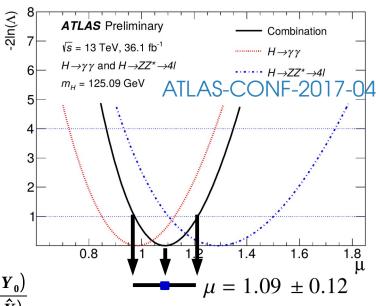
Interval endpoints = μ^{\pm} for which $t_{\mu^{\pm}} = 1$

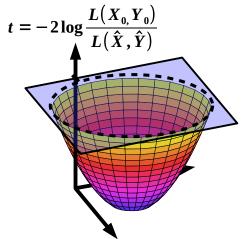
Gaussian case : $\hat{\mu} \pm \sigma$

Works also to obtain **contours in 2D**:







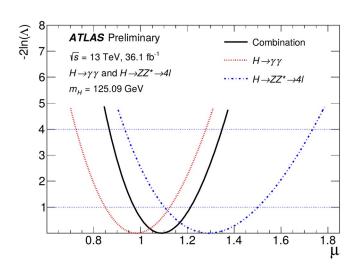


Highlights: Upper Limits

Confidence intervals: use
$$t_{\mu_0} = -2\log\frac{L(\mu = \mu_0)}{L(\hat{\mu})}$$

 \rightarrow Crossings with $t_{\mu 0} = Z^2$ for $\pm Z\sigma$ intervals (in 1D)

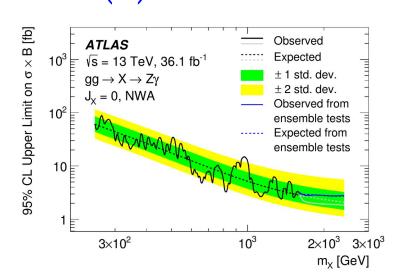
Gaussian regime: $\mu = \hat{\mu} \pm \sigma_{\mu}$ (1 σ interval)



Limits: use LR-based test statistic: $q_{S_0} = -2\log\frac{L(S-S_0)}{L(\hat{S})}$ $S_0 \geq \hat{S}$

→ Use CL_s procedure to avoid negative limits

Poisson regime, n=0: $S_{up} = 3$ events



Outline

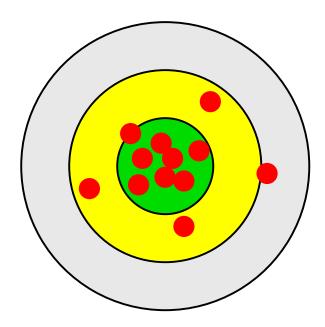
Profiling

Bayesian methods

Look elsewhere effect

Systematic Errors

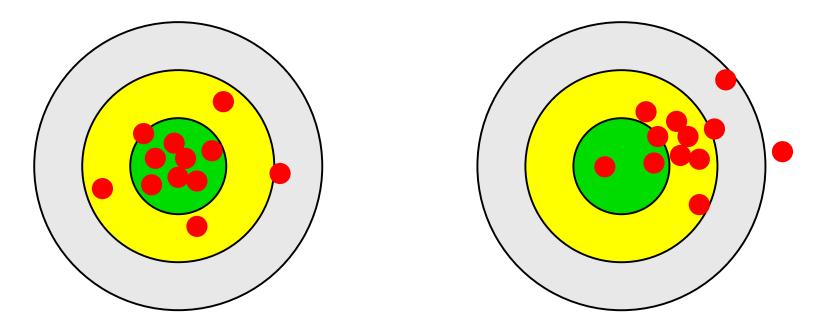
The statistical model (PDF) is a way to express **uncertainty** on the outcome of an experiment. e.g. 2D Gaussian:



These uncertainties are also called **Statistical Uncertainties** – they are the ones encoded in the model.

Systematic Errors

The statistical model (PDF) is a way to express **uncertainty** on the outcome of an experiment. e.g. 2D Gaussian:



These uncertainties are also called **Statistical Uncertainties** – they are the ones encoded in the model.

However the model itself may be wrong: this is a systematic error

- → To account for them, need a set of **Systematic uncertainties**
- → Can often add them "by hand", but how to treat this in a general way?

Systematic Uncertainties

Likelihood typically includes

- Parameters of interest (POIs): S, σ×B, m_w, ...
- Nuisance parameters (NPs): other parameters needed to define the model
 - → Ideally, **constrained by data** like the POI

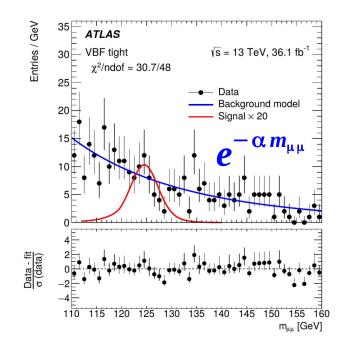
What about systematics?

- = what we don't know about the random processs
- ⇒ Parameterize using additional NPs
- ⇒ Add constraints in the likelihood

$$L(\mu, \theta; \text{data}) = L_{\text{measurement}}(\mu, \theta; \text{data}) C(\theta)$$

$$\downarrow \text{Systematics} \text{Measurement} \text{NP Constraint} \text{term}$$

Phys. Rev. Lett. 119 (2017) 051802



"Systematic uncertainty is, in any statistical inference procedure, the uncertainty due to the incomplete knowledge of the probability distribution of the observables.

G. Punzi, What is systematics?

 $C(\theta)$ represents extra knowledge about the NP

Frequentist Systematics

Prototype: NP measured in a separate *auxiliary* experiment e.g. luminosity measurement

→ Build the combined likelihood of the main+auxiliary measurements

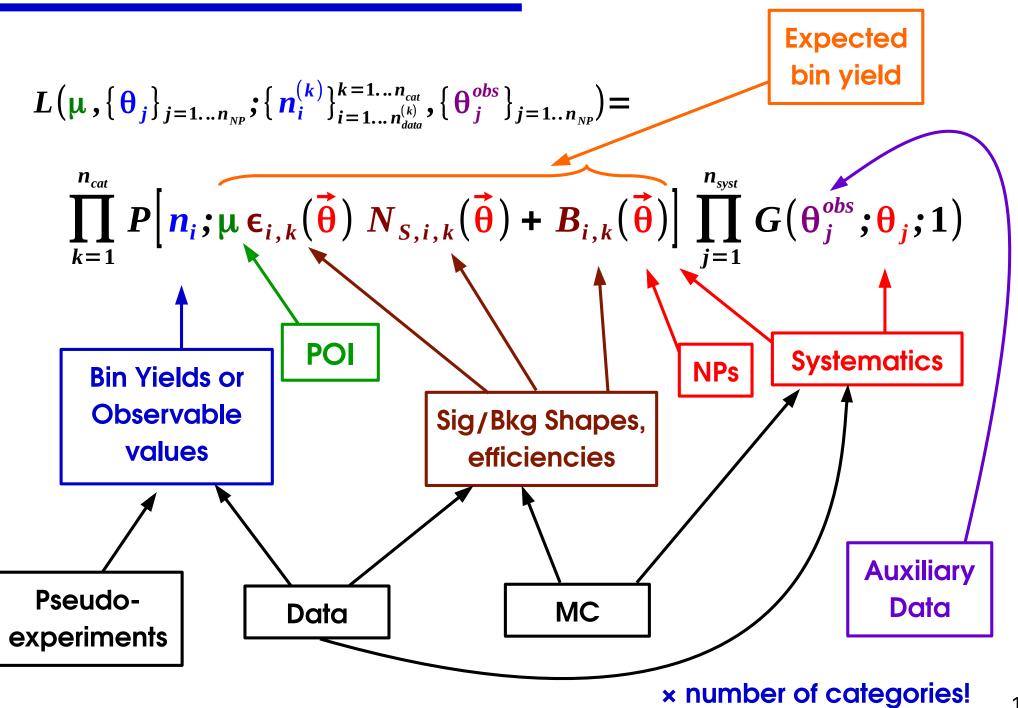
$$L(\mu, \theta; \text{data}) = L_{\text{main}}(\mu, \theta; \text{main data}) \quad L_{\text{aux}}(\theta; \text{aux. data}) \quad \text{independent measurements:} \\ \Rightarrow \text{ just a product}$$

Gaussian form often used by default: $L_{\text{aux}}(\theta; \text{aux. data}) = G(\theta^{\text{obs}}; \theta, \sigma_{\text{syst}})$

In the combined likelihood, systematic NPs are constrained

- → now same as e.g. NPs constrained in sidebands.
- → Often no clear setup for auxiliary measurements
 e.g. theory uncertainties on missing HO terms from scale variations
 - → Implemented in the same way nevertheless ("pseudo-measurement")

Likelihood, the full version (binned case)



Reminder: Wilks' Theorem

Cowan, Cranmer, Gross & Vitells Eur. Phys. J. C71:1554,2011

Consider
$$t_{S_0} = -2 \log \frac{L(S=S_0)}{L(\hat{S})}$$

→ Assume **Gaussian regime** (e.g. large n_{evts}, Central-limit theorem) : then:

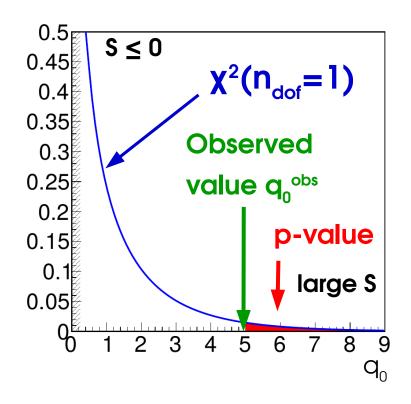
Wilk's Theorem: t_{s0} is distributed as a χ^2

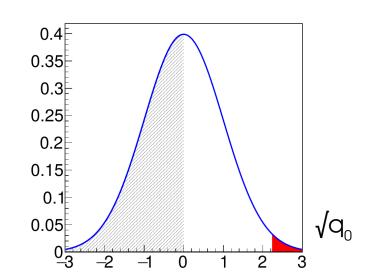
under $H_{SO}(S=S_0)$:

$$f(t_{S_0} | S = S_0) = f_{\chi^2(n_{dof} = 1)}(t_{S_0})$$

→ The significance is:

$$Z = \sqrt{q_0}$$





Profiling

How to deal with nuisance parameters in likelihood ratios?

 \rightarrow Let the data choose \Rightarrow use the best-fit values (*Profiling*)

$$\textbf{Profile Likelihood Ratio} \text{ (PLR)} \\ t_{S_0} = -2\log \frac{L(S=S_0, \hat{\hat{\theta}}(S_0))}{L(\hat{S}, \hat{\theta})} \\ \hat{\theta}(S_0) \text{ best-fit value for } S=S_0 \\ \text{ (conditional MLE)} \\ \hat{\theta} \text{ overall best-fit value} \\ \text{ (unconditional MLE)}$$

Wilks' Theorem: same properties as plain likelihood ratio

$$f(t_{S_0} | S = S_0) = f_{\chi^2(n_{dof} = 1)}(t_{S_0})$$
 also with NPs present

- → Profiling "builds in" the effect of the NPs
- \Rightarrow Can use t_{sn} to compute limits, significance, etc. in the same way as before

Homework 7: Gaussian Profiling

Counting experiment with background uncertainty: n = S + B:

$$\hat{\hat{B}}(S) = B_{\text{obs}} + \frac{\sigma_{\text{bkg}}^2}{\sigma_{\text{stat}}^2 + \sigma_{\text{bkg}}^2} (\hat{S} - S)$$

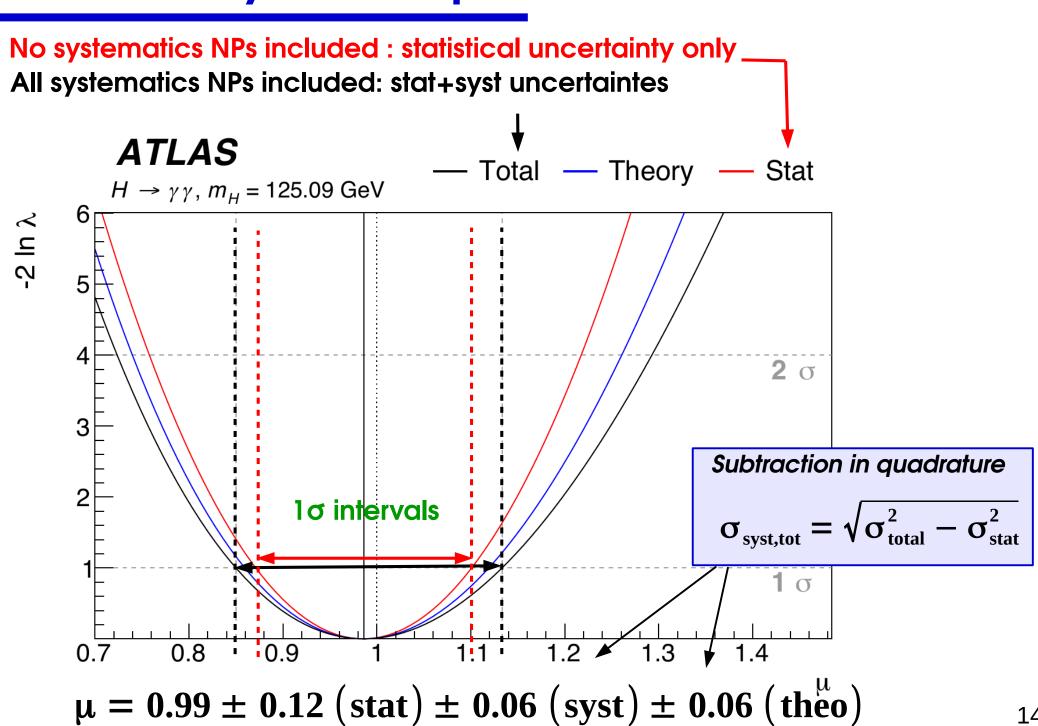
- Recall: Signal region only (fixed B): $t_S = \left(\frac{S n_{\rm obs}}{\sigma_{\rm stat}}\right)^2$ $S = (n_{\rm obs} B) \pm \sigma_{\rm stat}$ \rightarrow Compute the best-fit (MLEs) for S and B \rightarrow Show that the conditional MLE for B is $\hat{B}(S) = B_{\rm obs} + \frac{\sigma_{\rm bkg}^2}{\sigma_{\rm stat}^2 + \sigma_{\rm bkg}^2}(\hat{S} S)$
- → Compute the profile likelihood t_s
- → Compute the 1σ confidence interval on S

$$S = (n_{\text{obs}} - B_{\text{obs}}) \pm \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{bkg}}^2}$$

$$\sigma_{S} = \sqrt{\sigma_{\text{stat}}^{2} + \sigma_{\text{bkg}}^{2}}$$

Stat uncertainty (on n) and systematic (on B) add in quadrature

Uncertainty decomposition



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Pull/Impact plots

Systematics are described by NPs included in the fit. Define **pull** as

$$(\hat{\theta} - \theta_0) / \sigma_{\theta}$$

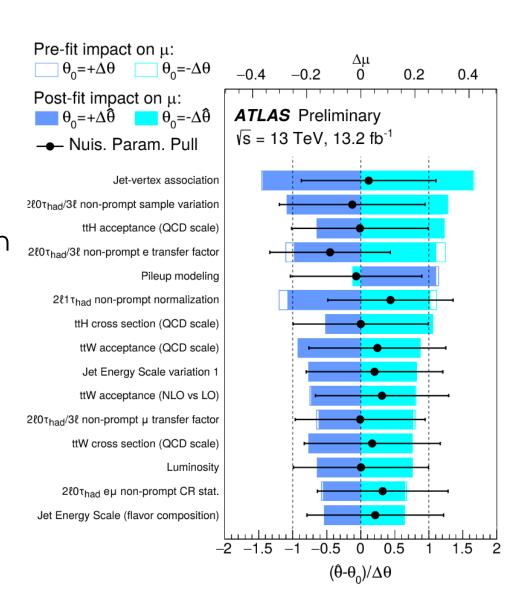
Nominally:

- pull = 0 : i.e. the pre-fit expectation
- pull uncertainty = 1: from the Gaussian

However fit results may be different:

- Central value ≠ 0: some data feature differs from MC expectation
 ⇒ Need investigation if large
- Uncertainty < 1 : effect is constrained by the data ⇒ Needs checking if this legitimate or a modeling issue

 \rightarrow Impact on result of $\pm 1\sigma$ shift of NP allows to gauge which NPs matter most .



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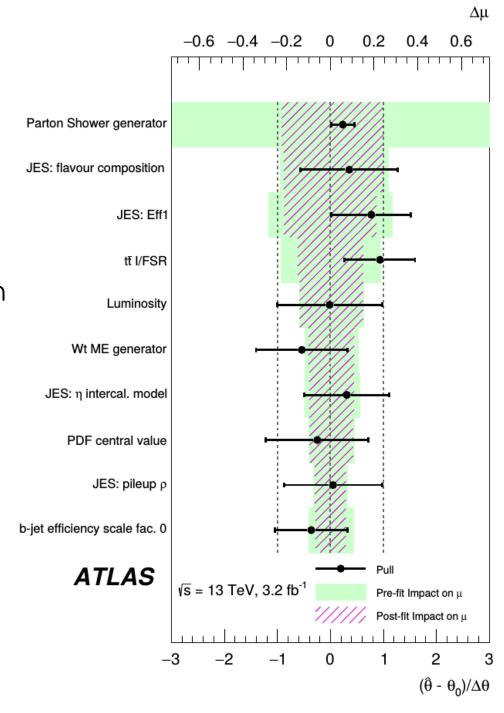
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- \rightarrow Impact on result of $\pm 1\sigma$ shift of NP allows to gauge which NPs matter most .

13 TeV single-t XS (arXiv:1612.07231)



Profiling Takeaways

When testing a hypothesis, use the best-fit values of the nuisance parameters: *Profile Likelihood Ratio*.

$$\frac{L(\mu = \mu_{0}, \hat{\hat{\theta}}_{\mu_{0}})}{L(\hat{\mu}, \hat{\theta})}$$

Allows to include systematics as uncertainties on nuisance parameters.

Profiling systematics includes their effect into the total uncertainty. Gaussian:

$$\sigma_{\text{total}} = \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}$$

Guaranteed to work well as long as everything is Gaussian, but typically also robust against non-Gaussian behavior.

Profiling can have unintended effects – need to carefully check behavior

Outline

Profiling

Bayesian methods

Look elsewhere effect

Bayesian methods

Probability distribution (= likelihood):

- → Same as frequentist case, but treat systematics by marginalization, i.e. integrating over priors, instead of profiling:
 - ightarrow Integrate out θ to get $P(\mu)$: $P(\mu) = \int P(\mu, \theta) C(\theta) d\theta$
 - \rightarrow Use probability distribution P(μ) directly for limits & intervals

e.g. 68% CL ("Credibility Level") interval [A, B] is:
$$\int\limits_A^B P(\mu)\pi(\mu)d\mu = 68\,\%$$

where $\pi(\mu)$ is the prior on μ . Uses **Bayes' Theorem**: $P(\mu \mid n) = P(n \mid \mu) \frac{P(\mu)}{P(n)}$

- No simple way to test for discovery
- Integration over NPs can be CPU-intensive (but can use MCMC methods)

Priors: most analyses use flat priors in the analysis variable(s)

- \Rightarrow **Parameterization-dependent**: if flat in $\sigma \times B$, them not flat in couplings....
- → Can use the Jeffreys' or reference priors, but difficult in practice

Homework 8: Bayesian methods and CL₂

Gaussian counting problem with systematic on background: $n = S + B + \sigma_{syst}\theta$

$$P(n;S,\theta) = G(n;S+B+\sigma_{\text{syst}}\theta,\sigma_{\text{stat}}) G(\theta_{\text{obs}}=0;\theta,1)$$

→ What is the 95% CL upper limit on S, given a measurement n_{obs}?

1. CLs computation:

- Use the result of Homework 7 to compute the PLR for S
- Use the result of Homework 6 to compute the CLs upper limit

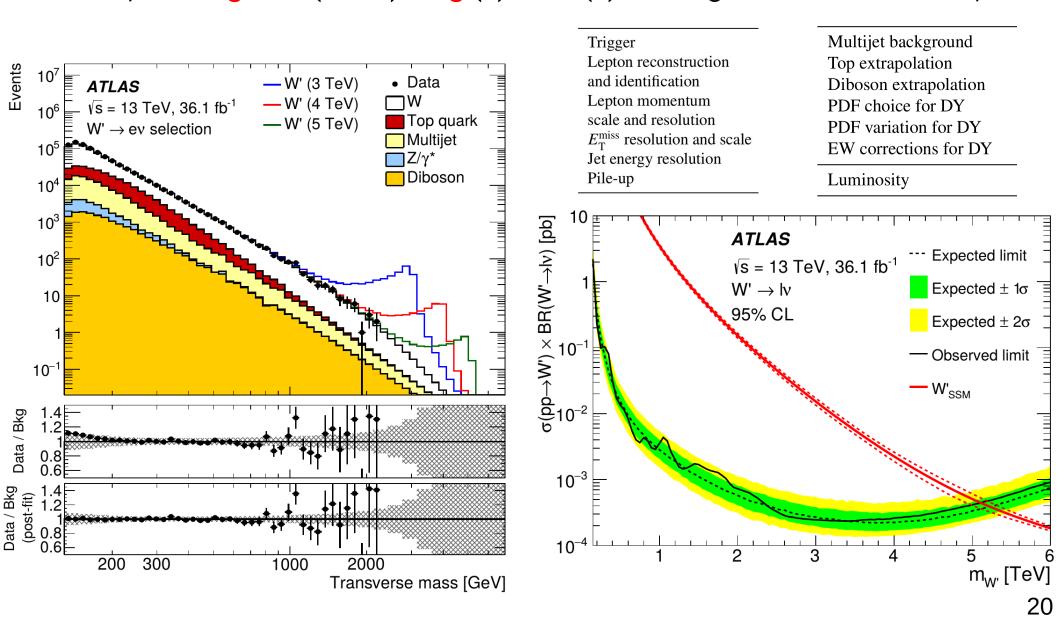
2. Bayesian computation:

- Integrate $P(n; S, \theta)$ over θ to get the marginalized P(n|S)
- S>0.
- Find the 95% CL limit by solving $\int_{S}^{\infty} P(S|n)dS = 5\%$

Solution: In both cases
$$S_{\rm up}^{\rm CL_s} = n - B + \left[\Phi^{-1} \left(1 - 0.05 \ \Phi \left(\frac{n - B}{\sqrt{\sigma_{\rm stat}^2 + \sigma_{\rm syst}^2}} \right) \right) \right] \sqrt{\sigma_{\rm stat}^2 + \sigma_{\rm syst}^2}$$

Example: W'→Iv Search

- POI: W' $\sigma \times B \rightarrow \text{use flat prior over } [0, +\infty[$.
- NPs: syst on signal ε (6 NPs), bkg (6), lumi (1) → integrate over Gaussian priors



Outline

Profiling

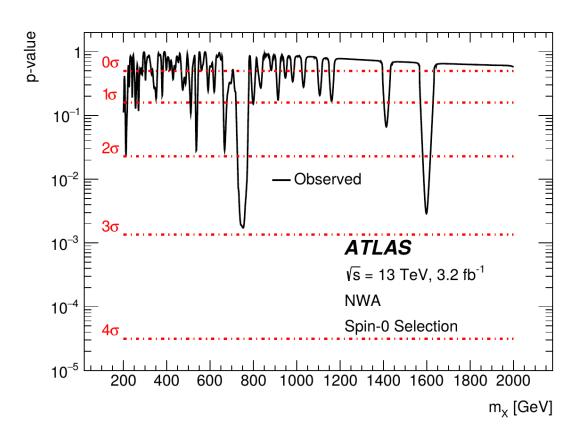
Bayesian methods

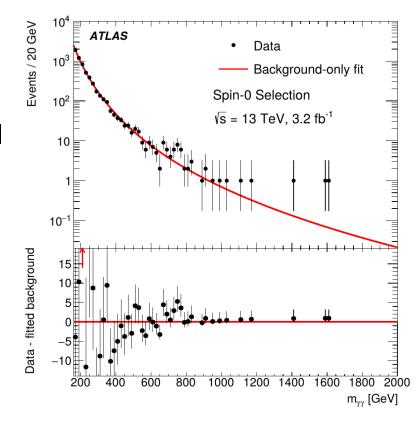
Look elsewhere effect

Look-Elsewhere effect

Sometimes, unknown parameters in signal model e.g. p-values as a function of m_v

- ⇒ Effectively: multiple, simultaneous searches
- → If e.g. small resolution and large scan range, many independent experiments

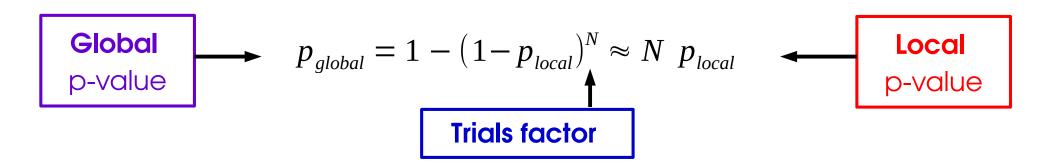




- → More likely to find an excess anywhere in the range, rather than in a predefined location
- **→ Look-elsewhere effect** (LEE)

Global Significance

Probability for a fluctuation *anywhere* in the range \rightarrow **Global** p-value. at a given location \rightarrow **Local** p-value



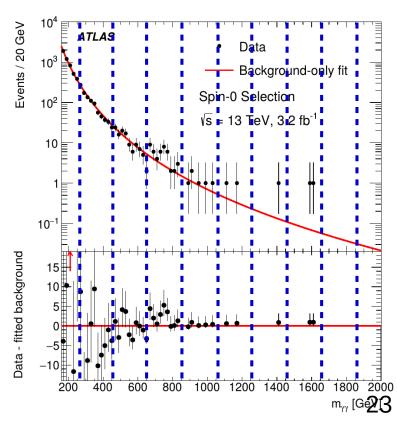
ightarrow $\mathbf{p}_{\mathsf{global}}$ $\mathbf{p}_{\mathsf{local}}$ \Rightarrow $\mathbf{Z}_{\mathsf{global}}$ < $\mathbf{Z}_{\mathsf{local}}$: global fluctuation more likely \Rightarrow less significant

 \triangle

Trials factor: naively = # of independent intervals:

$$N_{\text{trials}} = N_{\text{indep}} = \frac{\text{scan range}}{\text{peak width}}$$

However this is usually **wrong** – more on this later



Global Significance

Probability for a fluctuation *anywhere* in the range \rightarrow **Global** p-value. at a given location \rightarrow **Local** p-value

For searches over a parameter range, the global p-value is the relevant one

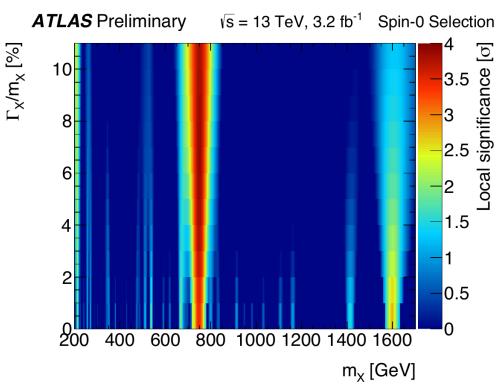
→ Accounts for the actual search procedure: look for an excess anywhere in

the scanned range

→ Depends on the scanned parameter ranges

e.g. $X \rightarrow \gamma \gamma$:

- $200 < m_x < 2000 GeV$
- $0 < \Gamma_x < 10\% \, \text{m}_x$.

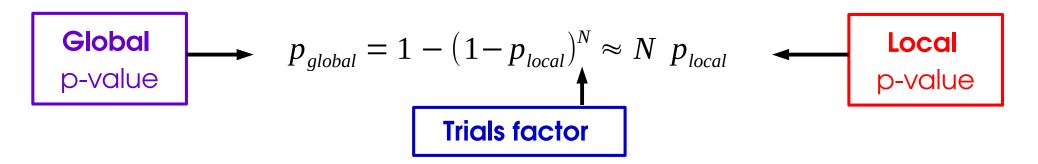


 $\rightarrow p_{local}$ is what comes out of the usual formulas

How to compute p_{global} (or N_{trials})?

Trials Factor

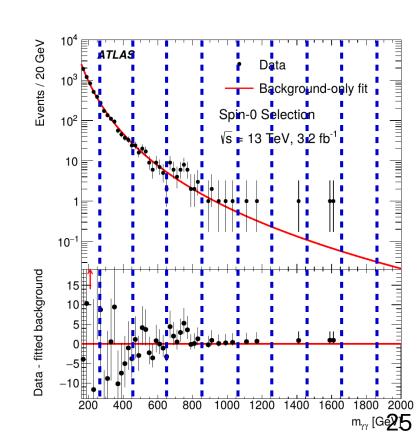
Trials factor N = # of independent searches:



Naively, one could expect

$$N_{\text{trials}} = N_{\text{indep}} = \frac{\text{scan range}}{\text{peak width}}$$

However this is only correct for a discrete Number of experiments (i.e. 10 different regions)

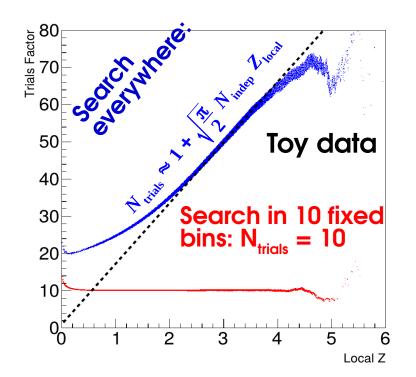


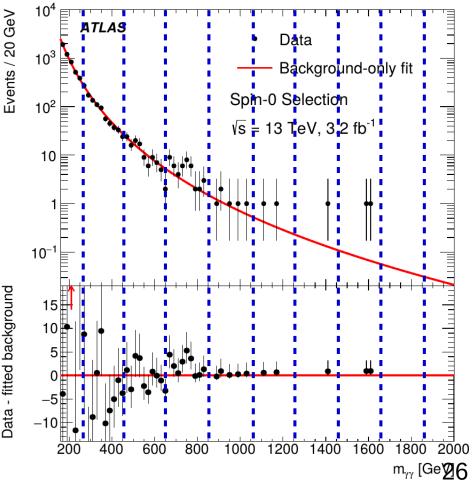
Asymptotic limit: trials factor (1 POI) is

$$m{N_{ ext{trials}}} = 1 + \sqrt{rac{\pi}{2}} \; m{N_{ ext{indep}}} \; m{Z_{ ext{local}}}$$
 and son $m{Z_{ ext{local}}}$!

 \rightarrow Trials factor is **not just N**_{indep}, also depends on Z_{local} !

Why? Slicing range into N_{indep} regions misses peaks sitting on edges between regions true N_{trials} is $> N_{indep}$!





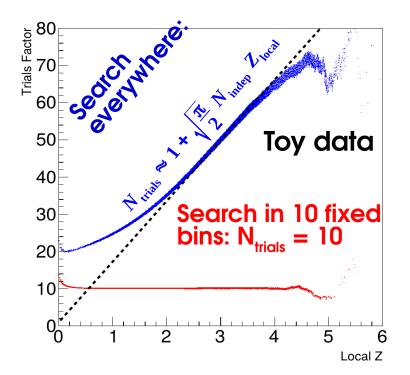
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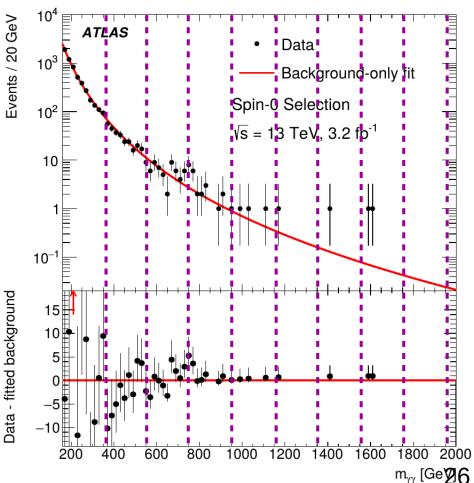
$$N_{\text{trials}} = 1 + \sqrt{\frac{\pi}{2}} N_{\text{indep}} Z_{\text{local}}$$

 \rightarrow Trials factor is **not just N**_{indep}, also depends on Z_{local} !

 $N_{indep} = \frac{scan range}{peak width}$

Why? Slicing range into N_{indep} regions misses peaks sitting on edges between regions true N_{trials} is $> N_{indep}$!





Global Significance from Toys

ATLAS

Data

Background-only fit

Spin-0 Selection

Vs = 13 TeV, 3.2 fb⁻¹

1000 1200 1400 1600

m_{yy} [GeV]

Principle: repeat the analysis in toy data:

- → generate pseudo-dataset
- → perform the search, scanning over parameters as in the data
- → report the largest significance found
- → repeat many times
- \Rightarrow The frequency at which a given Z_0 is found is the global p-value

e.g.
$$X \rightarrow \gamma \gamma$$
 Search: $Z_{local} = 3.9\sigma \ (\Rightarrow p_{local} \sim 5 \ 10^{-5})$,

- \rightarrow However we are scanning 200 < m_x < 2000 GeV and 0 < $\Gamma_{\rm x}$ < 10% m_x!
- → Toys : find such an excess 2% of the time somewhere in the range
- \Rightarrow p_{global} ~ 2 10⁻², $\mathbf{Z}_{global} = 2.1\sigma$ Less exciting, and better indication of true Z!
- **Exact treatment**
- CPU-intensive especially for large Z (need ~O(100)/p_{alobal} toys)

Conclusion

- Significant evolution in the statistical methods used in HEP
- Variety of methods, adapted to various situations and target results
- Allow to
 - model the statistical process with high precision in difficult situations (large systematics, small signals)
 - make optimal use of available information
- Implemented in standard RooFit/RooStat toolkits within the ROOT framework, as well as other tools (BAT)

- Still many open questions and areas that could use improvement
 - → e.g. how to present results with all available information

Homework solutions for Lecture 3

Homework 1: Gaussian Counting

Count number of events n in data

- → assume n large enough so process is Gaussian
- → assume B is known, measure S

$$L(S;n) = e^{-\frac{1}{2}\left(\frac{n-(S+B)}{\sqrt{S+B}}\right)^{2}}$$

Likelihood:

$$L(S;n) = e^{-\frac{1}{2} \left(\frac{n - (S+B)}{\sqrt{S+B}}\right)^{2}}$$
$$\lambda(S;n) = \left(\frac{n - (S+B)}{\sqrt{S+B}}\right)^{2}$$

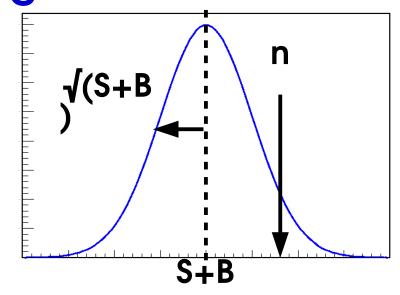
MLE for $S: \hat{S} = n - B$

Test statistic: assume $\hat{S} > 0$,

$$q_0 = -2\log\frac{L(S=0)}{L(\hat{S})} = \lambda(S=0) - \lambda(\hat{S}) = \left|\frac{n-B}{\sqrt{B}}\right|^2 = \left|\frac{\hat{S}}{\sqrt{B}}\right|^2$$

$$Z = \sqrt{q_0} = \frac{\hat{S}}{\sqrt{B}}$$

Known formula! → Strictly speaking only valid in Gaussian regime



Homework 2: Poisson Counting

Same problem but now **not** assuming Gaussian behavior:

$$L(S;n) = e^{-(S+B)}(S+B)^n$$
 $\lambda(S;n) = 2(S+B)-2n\log(S+B)$

MLE: $\hat{S} = n - B$, same as Gaussian

Test statistic (for $\hat{S} > 0$):

$$q_0 = \lambda(S=0) - \lambda(\hat{S}) = -2\hat{S} - 2(\hat{S}+B) \log \frac{B}{\hat{S}+B}$$

Assuming asymptotic distribution for q_0 ,

$$Z = \sqrt{2\left[\left(\hat{S} + B\right)\log\left(1 + \frac{\hat{S}}{B}\right) - \hat{S}\right]}$$

Homework solutions for Lecture 4

Homework 3: Gaussian CL_{s+b}

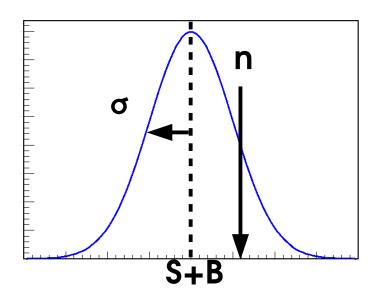
Usual Gaussian counting example with known B:

$$\lambda(S) = \left| \frac{n - (S + B)}{\sigma_S} \right|^2$$

Reminder:

Best fit signal : $\hat{S} = n - B$

Significance: $Z = \hat{S} / \sqrt{B}$



Compute the 95% CL upper limit on S:

$$q_{S_0} = -2\log\frac{L(S=S_0)}{L(\hat{S})} = \lambda(S_0) - \lambda(\hat{S}) = \left(\frac{n - (S_0 + B)}{\sigma_S}\right)^2 = \left(\frac{S_0 - \hat{S}}{\sigma_S}\right)^2 \qquad \stackrel{\text{for }}{S_0 > \hat{S}}$$

so
$$q_{S_0} = 2.70$$
 for $S_0 = \hat{S} + \sqrt{2.70} \sigma_S$

And finally
$$S_{up} = \hat{S} + 1.64 \sigma_S$$
 at 95 % CL

Homework 4: Gaussian CL

Usual Gaussian counting example with known B:

$$\lambda(S) = \left(\frac{n - (S + B)}{\sigma_S}\right)^2$$

Reminder

Best fit signal : $\hat{S} = n - B$

$$S_{up} = \hat{S} + 1.64 \sigma_s$$
 at 95 % CL

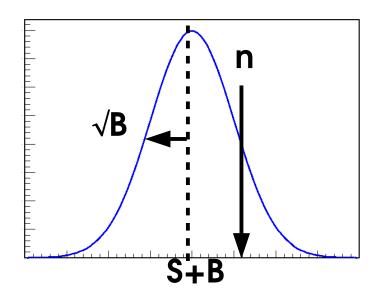
CL_{s+b} limit: $S_{up} = \hat{S} + 1.64 \sigma_S \text{ at } 95 \% \text{ CL}$ CL_s upper limit: still have $q_{S_0} = \left(\frac{S_0 - \hat{S}}{\sigma_S}\right)^2 \text{ (for } S_0 > \hat{S}\text{)}$

so need to solve

$$p_{CL_s} = \frac{p_{S_0}}{1 - p_B} = \frac{1 - \Phi(\sqrt{q_{S_0}})}{1 - \Phi(\sqrt{q_{S_0}} - S_0/\sigma_S)} = 5\%$$

for
$$\hat{S} = 0$$
,

$$S_{up} = \hat{S} + \left[\Phi^{-1}\left(1 - 0.05 \Phi(\hat{S}/\sigma_s)\right)\right] \sigma_s$$
 at 95% CL



$$\hat{S} \sim G(S, \sigma_s)$$
 so

Under $H_0(S = S_0)$:

$$\sqrt{q_{S_0}} \sim G(0,1)$$

$$p_{S_0} = 1 - \Phi(\sqrt{q_{S_0}})$$

Under $H_0(S = 0)$:

$$\sqrt{q_{S_0}} \sim G(S_0/\sigma_S, 1)$$

$$p_B = \Phi(\sqrt{q_{S_0}} - S_0/\sigma_S)$$

Homework 5: Poisson CL_s

Same exercise, for the Poisson case

Exact computation: sum probabilities of cases "at least as extreme as data" (n)

$$p_{S_0}(n) = \sum_{0}^{n} e^{-(S_0 + B)} \frac{(S_0 + B)^k}{k!}$$
 and one should solve $p_{CL_s} = \frac{p_{S_{up}}(n)}{p_0(n)} = 5\%$ for S_{up}

For n = 0:
$$p_{CL_s} = \frac{p_{S_{up}}(0)}{p_0(0)} = e^{-S_{up}} = 5\% \Rightarrow S_{up} = \log(20) = 2.996 \approx 3$$

 \Rightarrow Rule of thumb: when $n_{obs}=0$, the 95% CL_s limit is 3 events (for any B)

Asymptotics: as before,
$$q_{S_0} = \lambda(S_0) - \lambda(\hat{S}) = 2(S_0 + B - n) - 2n \log \frac{S_0 + B}{n}$$

For n = 0,
$$q_{S_0}(n=0) = 2(S_0 + B)$$

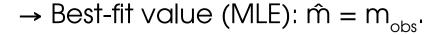
$$p_{CL_s} = \frac{p_{S_0}}{p_0} = \frac{1 - \Phi(\sqrt{q_{S_0}(n=0)})}{1 - \Phi(\sqrt{q_{S_0}(n=0)} - \sqrt{q_{S_0}(n=B)})} = 5\%$$

- \Rightarrow S_{up} ~ 2, exact value depends on B
- \Rightarrow Asymptotics not valid in this case (n=0) need to use exact results, or toys

Homework 6: Gaussian Intervals

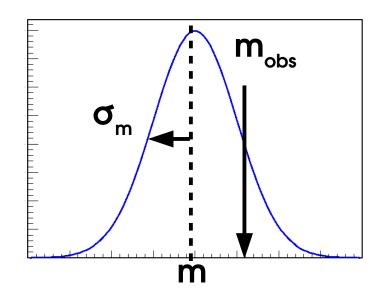
Consider a parameter m (e.g. Higgs boson mass) whose measurement is Gaussian with known width $\sigma_{\rm m}$, and we measure $m_{\rm obs}$:

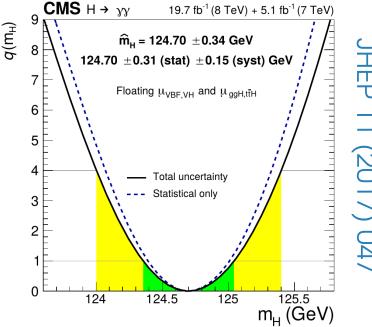
$$\lambda(m; m_{\text{obs}}) = \left(\frac{m - m_{\text{obs}}}{\sigma_m}\right)^2$$



→ Test statistic:
$$t_m = \left(\frac{m - m_{\text{obs}}}{\sigma_m}\right)^2$$

$$\rightarrow$$
 1 σ Interval $m = m_{\rm obs} \pm \sigma_{\rm m}$





Homework solutions for Lecture 5

Homework 7: Gaussian Profiling

Counting experiment with background uncertainty: $\mathbf{n} = \mathbf{S} + \mathbf{\theta}$:

→ Signal region:
$$\mathbf{n} \sim \mathbf{G}(\mathbf{S} + \mathbf{\theta}, \sigma_{\text{stat}})$$

→ Control region: $\mathbf{\theta}^{\text{obs}} \sim \mathbf{G}(\mathbf{\theta}, \sigma_{\text{syst}})$

$$L(S, \theta) = G(n; S + \theta, \sigma_{\text{stat}}) \ G(\theta^{\text{obs}}; \theta, \sigma_{\text{syst}})$$

Then:
$$\lambda(S, \theta) = \left(\frac{n - (S + \theta)}{\sigma_{\text{stat}}}\right)^2 + \left(\frac{\theta^{\text{obs}} - \theta}{\sigma_{\text{syst}}}\right)^2$$

For S = Ŝ, matches MLE as it should

MLEs:
$$\hat{S} = n - \theta^{\text{obs}}$$
 Conditional MLE: $\hat{\hat{\theta}}(S) = \theta^{\text{obs}} + \frac{\sigma_{\text{syst}}^2}{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2} (\hat{S} - S)$

PLR:
$$t_S = -2\log\frac{L(S,\hat{\theta}(S))}{L(\hat{S},\hat{\theta})} = \lambda(S,\hat{\theta}(S)) - \lambda(\hat{S},\hat{\theta}) = \frac{(S-\hat{S})^2}{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}$$

1
$$\sigma$$
 interval $S = \hat{S} \pm \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}$ $\sigma_S = \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}$

Stat uncertainty (on n) and systematic (on θ) add in quadrature

Homework 8: CL_s computation

Gaussian counting with systematic on background: $n = S + B + \sigma_{syst}\theta$

$$L(n;S,\theta) = G(n;S+B+\sigma_{\text{syst}}\theta,\sigma_{\text{stat}}) G(\theta_{\text{obs}}=0;\theta,1)$$

MLE:
$$\hat{S} = n - B$$

Conditional MLE: $\hat{\hat{\theta}}(\mu) = \frac{\sigma_{\text{syst}}}{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2} (n - S - B)$

PLR: $\lambda(\mu) = \left(\frac{S + B - n}{\sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}}\right)^2$

This boils down to the Gaussian case of HW 6, so the CL_s limit is

CL_s:
$$S_{up}^{CL_s} = n - B + \left[\Phi^{-1} \left(1 - 0.05 \Phi \left(\frac{n - B}{\sqrt{\sigma_{stat}^2 + \sigma_{syst}^2}} \right) \right) \right] \sqrt{\sigma_{stat}^2 + \sigma_{syst}^2}$$

Homework 8: Bayesian computation

Gaussian counting with systematic on background: $n = S + B + \sigma_{syst}\theta$

$$P(n \mid S, \theta) = G(n; S+B+\sigma_{\text{syst}}\theta, \sigma_{\text{stat}}) G(\theta \mid 0, 1)$$

Bayesian: $G(\theta)$ is actually a **prior** on $\theta \Rightarrow$ perform integral (**marginalization**)

$$P(n|S) = G(S; n-B, \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2})$$

same effect as profiling!

0.4

0.35

0.3

0.25

0.15

0.1

Need P(S|n) \Rightarrow a prior for S – take flat PDF over S > 0 \Rightarrow Truncate Gaussian at S=0: $P(S \mid n) = P(n \mid S) P(S)$

$$P(S \mid n) = P(n \mid S) P(S)$$

$$P(S \mid n) = G(S; n-B, \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}) \left[\Phi \left(\frac{n-B}{\sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}} \right) \right]^{-1}$$

Bayesian Limit:

$$\int_{S_{up}}^{\infty} P(S \mid n) dS = 5\% = \left[1 - \Phi\left(\frac{S_{up} - (n - B)}{\sqrt{\sigma_{stat}^2 + \sigma_{syst}^2}}\right)\right] \left[\Phi\left(\frac{n - B}{\sqrt{\sigma_{stat}^2 + \sigma_{syst}^2}}\right)\right]^{-1}$$

$$\int_{S_{up}}^{\infty} P(S|n) dS = 5\% = \left[1 - \Phi\left(\frac{S_{up} - (n-B)}{\sqrt{\sigma_{stat}^2 + \sigma_{syst}^2}}\right)\right] \left[\Phi\left(\frac{n-B}{\sqrt{\sigma_{stat}^2 + \sigma_{syst}^2}}\right)\right]^{-1}$$

$$S_{\text{up}}^{\text{Bayes}} = n - B + \left[\Phi^{-1} \left[1 - 0.05 \, \Phi \left(\frac{n - B}{\sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}} \right) \right] \right] \sqrt{\sigma_{\text{stat}}^2 + \sigma_{\text{syst}}^2}$$

same result as CL,!