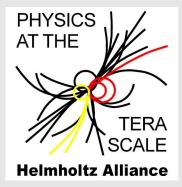




# BAT A Bayesian Analysis Toolkit



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#### Outline:

- Motivation
- Requirements/Specifications/Implementation
- Markov Chain Monte Carlo
- A simple example
- Applications and status





**Motivation** 

**Experimental data**  $\leftrightarrow$  **Data analysis**  $\leftrightarrow$  **Models and parameters** 

What does the data tell us about our model?

Which model is favored by the data?

Is the model in agreement with the data?

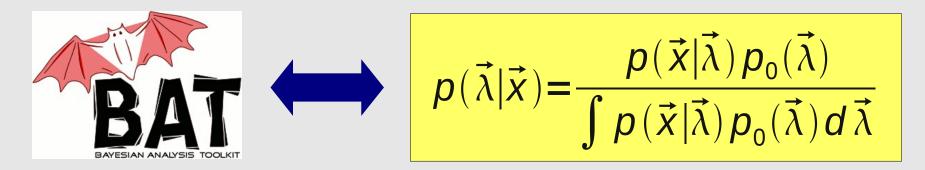
Parameter estimation

**Model comparison** 

**Goodness-of-fit test** 

... from simple line fitting to cosmological model testing!

#### **Need tools to extract information!**







# **Requirements / Specifications**

## Should be able to ...

- (1) phrase arbitrary problems,
- (2) interface to HEP-software,
- (3) estimate parameters,
- (4) extract pdfs for single parameters and correlations; evaluate limits,
- (5) propagate uncertainties,
- (6) compare models,
- (7) check validity of a models.

#### Implementation:

- C++ framework based on ROOT core functionality. BAT comes as library.
- Define base class for models which then need to be specified by the user, or
- run ROOT interactively using pre-defined fitter methods.
- Use modules for numerical tasks (optimization, integration, etc.). Chose from a set of algorithms.





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#### **Implementation:**

- Use minimization methods, e.g. interface to TMinuit. Working on simulated annealing.
- Use Markov Chain Monte Carlo (MCMC) for marginalization and error estimation.
- Use MCMC for uncertainty propagation. (no Gaussian assumptions here)

## **Key element: MCMC**





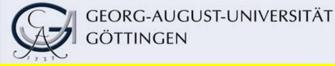
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#### **Implementation:**

- Model comparison using common definitions (e.g. direct comparison of probabilities)
- Model validity is a difficult topic. Use **p-value** to judge. (generalization of  $\chi^2$ -probability).





## Implementation

#### User defined (Pseudo code)

#### class **myModel**:

- Parameters
- Conditional probability
- Prior probabilities

## int main(){

MyModel \* mm;

mm  $\rightarrow$  "read in data"();

mm  $\rightarrow$  "perform analysis"(); }

#### Common tasks

#### Analysis

- Normalize (Integration)
- Maximize (Minimization)
- Marginalize/Sampling
- Goodness-of-fit
- Model comparison
- Error propagation
- Write output file
- Graphical output





# **Tools and algorithms**

#### Integration:

- Simple Monte Carlo algorithms (sampled mean, importance sampling)
- Interface to CUBA (VEGAS, ...)
- Marginalization:
  - MCMC (Metropolis)
- Sampling:
  - Monte Carlo sampling
  - MCMC (Metropolis)
- Propagation of uncertainies:
  - Calculate any function of the parameters during MCMC.

## • Minimization:

- Monte Carlo (hit & miss)
- MCMC (during random walk)
- Interface to Minuit
- Simulated Annealing planned
- Goodness-of-fit:
  - Ensemble tests and p-value

# Key: Markov Chain Monte Carlo (MCMC)





## **MCMC and Bayesian Inference**

#### MCMC in BAT:

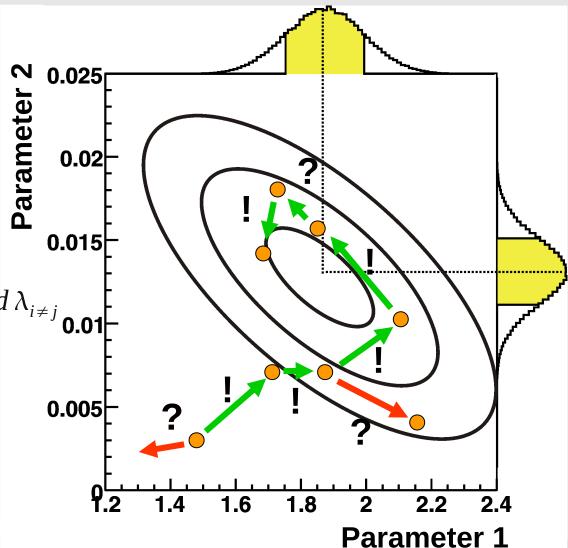
- Sample parameter space using MCMC
- Function:

$$f(\vec{\lambda}) = p(D|\vec{\lambda}) \cdot p_0(\vec{\lambda})$$

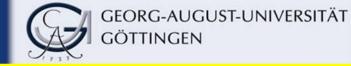
• Marginalize pdf while walking:

$$p(D|\vec{\lambda}) = \int p(D|\vec{\lambda}) \cdot p_0(\vec{\lambda}) d\lambda_{i \neq j}$$

- Calculate any function of parameters while walking (uncertainty propagation)
- Find global maximum
- Step size?





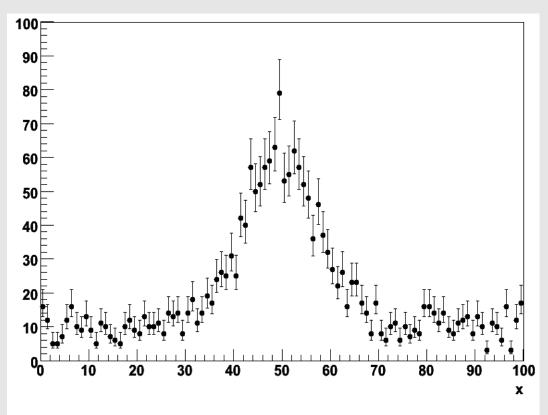


#### Simple spectrum:

- Gaussian signal: position  $\mu = 50$ width  $\sigma = 7.5$ strength  $\langle S \rangle = 1,000$
- Flat background: strength <B> = 10/bin
- Number of events per bin fluctuate with Poisson distribution

$$p(D|S,\mu,\sigma,B) = \prod_{i=1}^{N_{bins}} \frac{\lambda_i^{n_i}}{n_i!} e^{-\lambda_i}$$

$$\lambda_i = S \int_{\Delta_i} \frac{1}{\sqrt{2\pi\sigma}} e^{\frac{-(x-\mu)^2}{2\sigma^2}} dx + \frac{B}{\Delta_i}$$



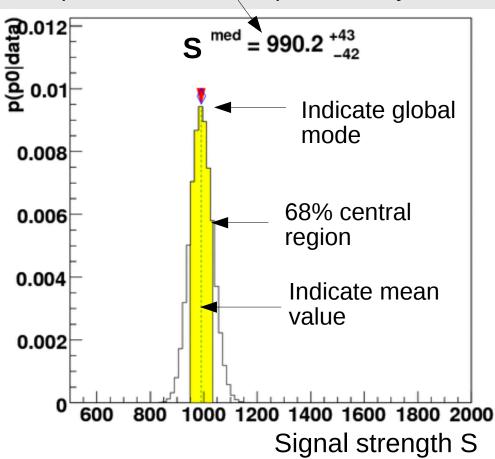




## **Marginalized Distributions 1-D**

#### **Marginalized distributions:**

- Project sampled distribution onto one (or two) parameters. Integrate over all others.
- Mode of marginalized distribution is in general not equal global mode. User's decision.
- Full information in Markov chain.
- Default output:
  - Global mode
  - Mean value
  - Central and smallest 68% interval

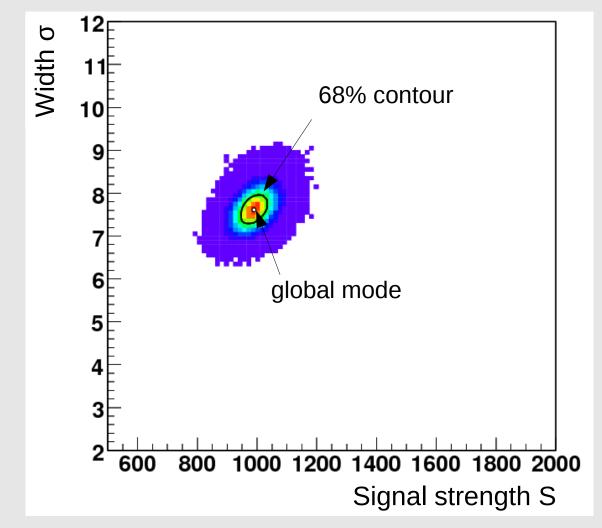






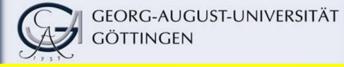


## **Marginalized Distributions 2-D**

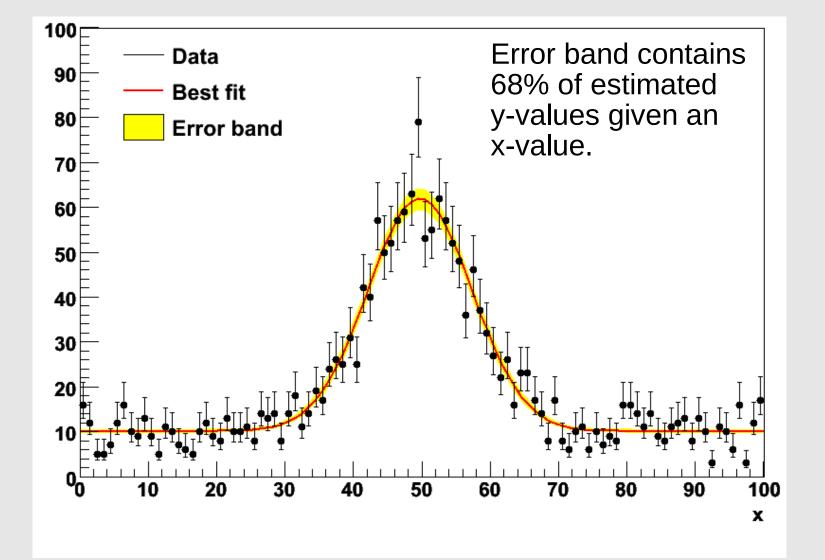


All distributions (1-d and 2-d) are stored during single run



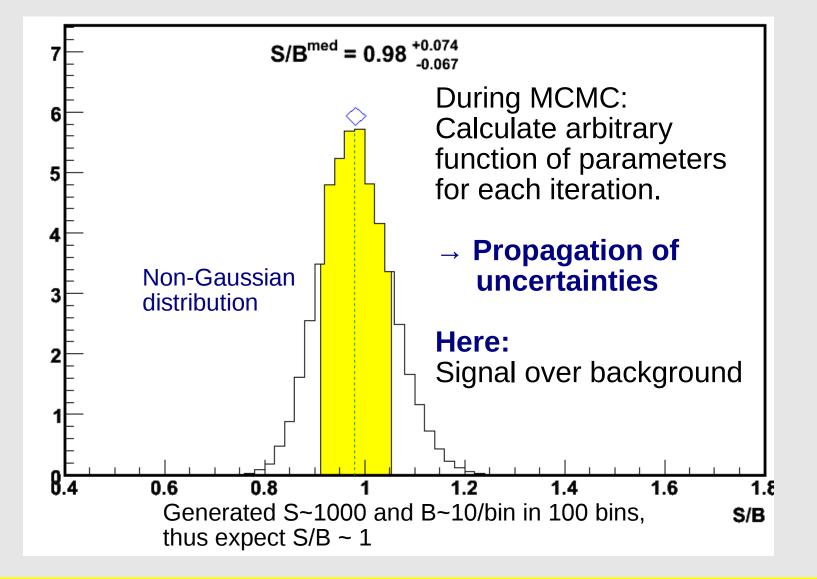


#### **Best Fit and Error Band**





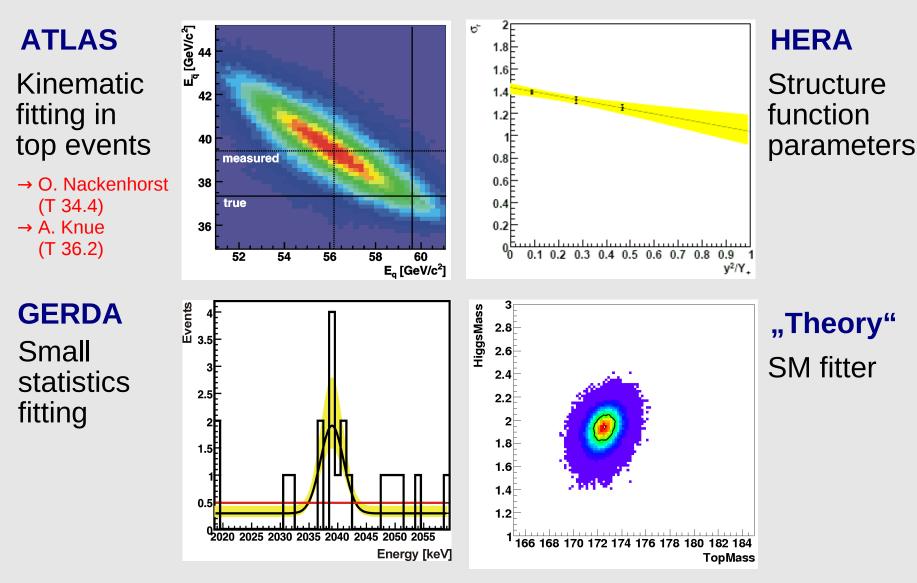








## **Applications**







- Current software release available on web-site: www.mppmu.mpg.de/bat
- Paper on arxiv: http://arxiv.org/abs/0808.2552 (submitted to *Computer Physics Communications*)
- Examples and use-cases available
- Interface to RooStat (part of next BAT release)
- Implementation (of parts) into ROOT planned (with ROOT-team)
- More algorithms are currently being implemented
- Developers and beta-testers welcome!





#### Backup





4. Goto 2.

## **MCMC: Metropolis Algorithm**

# Sampling parameter space can efficiently be done using MCMC

**Idea:** random walk heading towards regions of larger probability

#### Metropolis algorithm:

- 1. Start at random point x<sub>i</sub>.
- Generate proposal point x<sub>p</sub> vicinity of x<sub>i</sub>.
- 3. If  $f(x_p) > f(x_i)$  chose  $x_{i+1} = x_p$ , Else, accept proposal only with probability  $p=f(x_p)/f(x_i)$ .

 $f(x_{p})$   $f(x_{p})$ 

Causes migration towards region of large function values (probability)





## Simple MCMC Examples

## **Test function:**

 $f(x) = x^4 \cdot \sin^2 x$ 

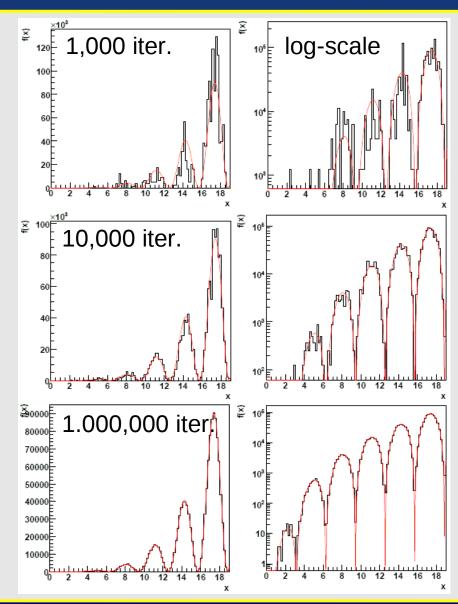
#### Here:

- chains converge quickly towards underlying function.
- Several maxima/minima are no problem. Can sample from complicated functions.

## But:

when is convergence reached?

→ Convergence criteria have to be defined.







#### **Flow of algorithm for MCMC:**

- Run several chains in parallel (default: 5).
- Start a pre-run:
  - forget about initial position
  - sample through parameter space:
    - adjust proposal function
    - monitor convergence criterion
  - stop, if (adjusted and converged) or (max. number of iterations)
  - summarize MCMC-parameters
- Start main run:
  - sample through parameter space:
    - update marginalized distributions (1-D and 2-D)
    - update global mode (if necessary)
    - calculate any function of parameters for uncertainty propagation
  - stop, if maximum number of iterations reached (default: 100,000)





#### **Proposal function:**

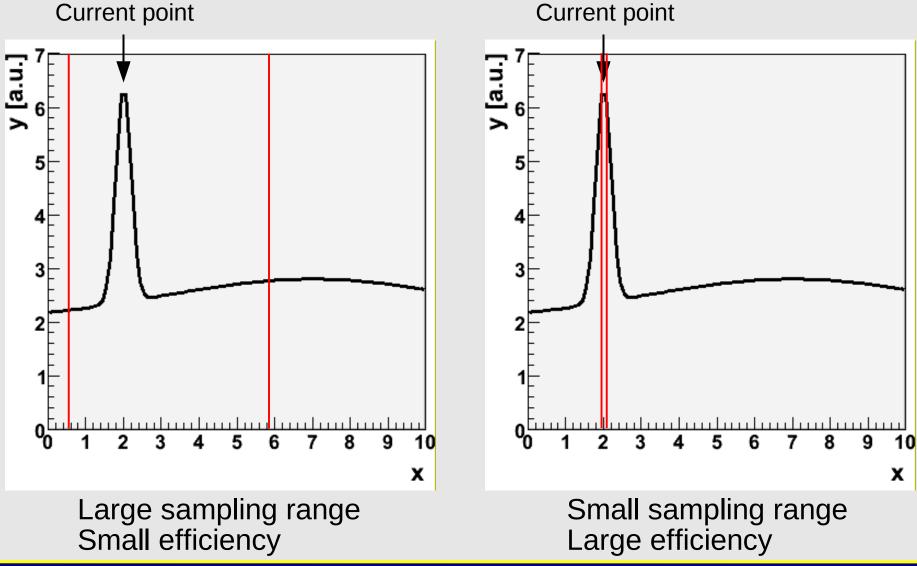
- Proposal function is flat in a box around current point
- Start with width = interval of parameter
- Adjust width during MCMC-prerun.
- Efficiency  $\epsilon$  is defined as the fraction of accepted proposals:
  - $\epsilon \rightarrow 1$ : random sampling (no migration towards maxima)
  - $\epsilon \rightarrow 0$ : no sampling (no walk at all)
  - $\epsilon \sim 0.25$ : (empirical) optimal value (trade-off)
- Calculate efficiency for a consecutive sub-chains. If  $\varepsilon < 0.1$ : reduce width by factor  $\frac{1}{2}$  ("curve" becomes flatter) if  $\varepsilon > 0.5$ : increase width by factor 2 ("jump around" more)
- Do not change efficiency during main run  $\rightarrow$  bias.
- Form of proposal function can be changed.





## **MCMC** in **BAT**: Proposal function

Current point





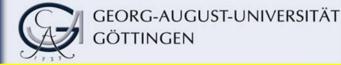




#### **Convergence:**

- How do you know the chain converged towards the underlying distribution?
- Can build criteria which give hints
- Two types of convergence: single chain: compare chain and distributions or chain with itself several chains: compare chains to each other
- Convergence criteria for single chains:
  - could use, e.g., auto-correlation function, etc.
  - can be CPU-intensive
  - can be tested offline if chain is stored to file
  - not yet implemented.





#### **Convergence:**

- Convergence criteria for several chains:
  - can be tested offline if chains are stored to file
  - use approach from Gelman & Rubin, StatSci 7, 1992:
    - compare mean of variances of single chains with variance of target distribution.  $\hat{V}$
    - define **r-value criterion**: <sup>r</sup>

$$W = \frac{1}{m} \frac{1}{n-1} \sum_{j=1}^{m} \sum_{i=1}^{n} (x_i - \bar{x}_j)^2$$
$$\hat{V} = (1 - \frac{1}{n})W + \frac{1}{m-1} \sum_{i=1}^{m} (\bar{x}_j - \bar{x})^2$$

Mean of variances of all chains

Estimate variance of target distribution

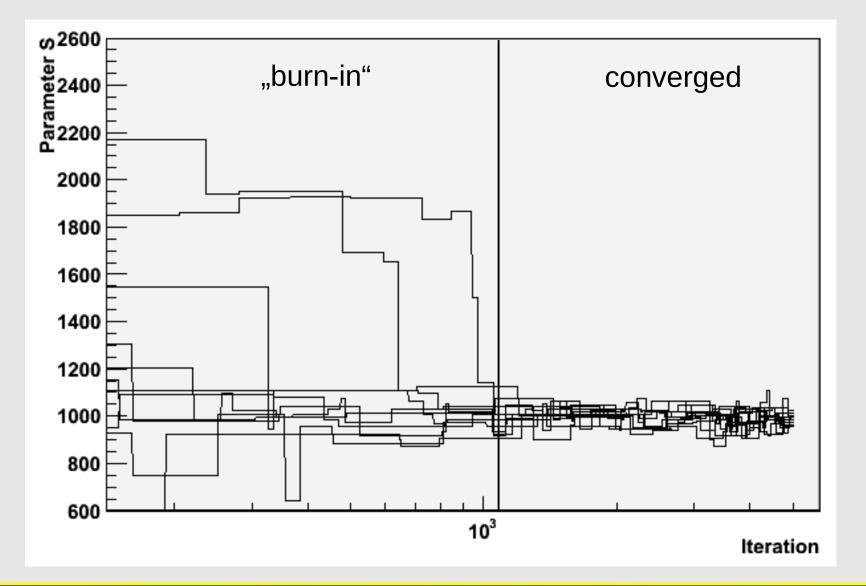
- require: (r-1) < 0.1 for each parameter and log prob.
- General question: Use one long chain or several shorter ones?







#### MCMC in BAT: Convergence Criteria







#### MCMC in BAT: Parameters of Running Chains

## **Option 1: "fast mode":**

- Single chain ran for a fixed number of iterations.
- Short pre-run to forget initial position.
- No convergence test
- No adjustment of proposal function

#### **Option 2: "precision mode":**

- 5 chains ran in parallel
- Pre-run including convergence test and adjustment of proposal function
- Default number of iterations: 100,000

#### In general:

User can set all parameters of running individually:

- Iterations, convergence criteria, pre-run, ...
- Requires some knowledge or intuition on the problem studied.





## Input/Output

#### Input to BAT:

- Possible input formats:
  - User defined (on the fly)
  - ROOT trees
  - ROOT histograms
  - ASCII files
- Data-classes for easy handling of data.
- Depending on model classes, input can be made available by user.

(Poisson)

(Binomial)

(Gauss)

- For fitter classes:
  - TH1D
  - 2 x TH1D
  - TGraphErrors

**Output from BAT:** 

- Log file with processing details
- ASCII summary to screen and to file:
  - Model summary
  - Mean values, modes, etc.
  - Posterior probabilities
- Plots:
  - Marginalized distributions (1-D)
  - Correlation plots (2-D)
  - Fit function and error band
- ROOT-file:
  - Summary data
  - add distributions
  - Markov chain (all points)



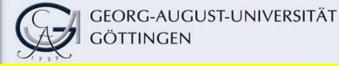


#### Fit a function to spectrum:

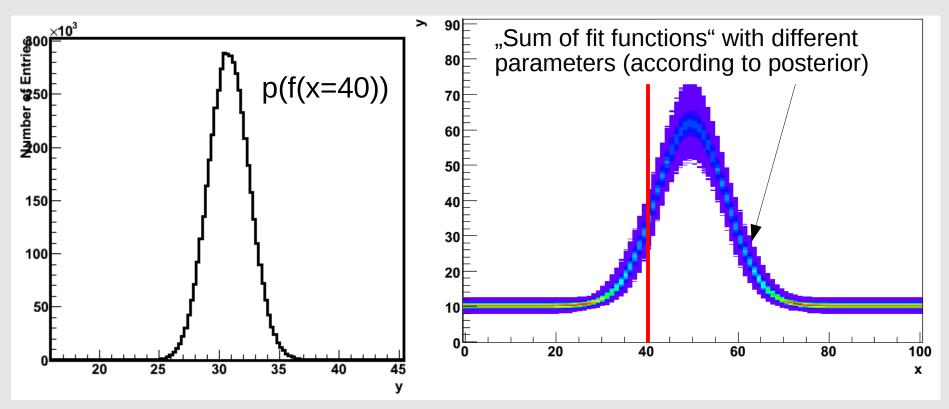
- Fit function *f* is Gaussian (3) + const. (1)
- 4 parameters
- No prior knowledge on the parameters, i.e.,  $p_0$ =const.
- Assume that Poissonian fluctuation per bin are independent, i.e., conditional probability is product of Poisson terms:

$$p(D|S,\mu,\sigma,B) = \prod_{i=1}^{N_{bins}} \frac{\lambda_i^{n_i}}{n_i!} e^{-\lambda_i}$$
$$\lambda_i = \int_{\Delta_i} \frac{1}{\sqrt{2\pi\sigma}} e^{\frac{-(x-\mu)^2}{2\sigma^2}} dx + \frac{B}{\Delta_i}$$





While sampling through parameter space: calculate *f* at every x (at that point in parameter space)



At x = 40: probability (density) for function f to have value y **Use this to define uncertainty band** (e.g., use 68% center interval)





#### How to judge if the model describes the data (at all)?

#### Strategy:

- Find best parameters from original data,  $\vec{\lambda}^*$
- Generate data sets, *x*, using these parameters (ensemble tests)
- Calculate and histogram the probability  $p(x|\vec{\lambda}^*)$
- Compare prob. distribution with prob. of original data, *D*,  $p(x=D|\vec{\lambda}^*)$
- Calculate the *p*-value:
  - probability that  $p(x|\vec{\lambda}^*)$  is equal to or less than  $p(x=D|\vec{\lambda}^*)$
  - Returns value between 0 and 1
  - Large *p*-value means good agreement
- Note: for the Gaussian case, the p-value is equivalent to a X<sup>2</sup>-probability.





#### **Goodness-of-fit**

For a good model, the probability to find a better matching between data and theory (larger likelihood) should be small.

#### Here:

*p*-value = 0.49

