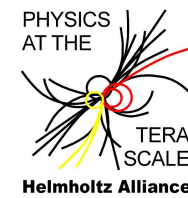




An Introduction to BAT

BAT Workshop Bologna 2011

February 24th - 25th 2011



Kevin Kröninger
University of Göttingen

for the

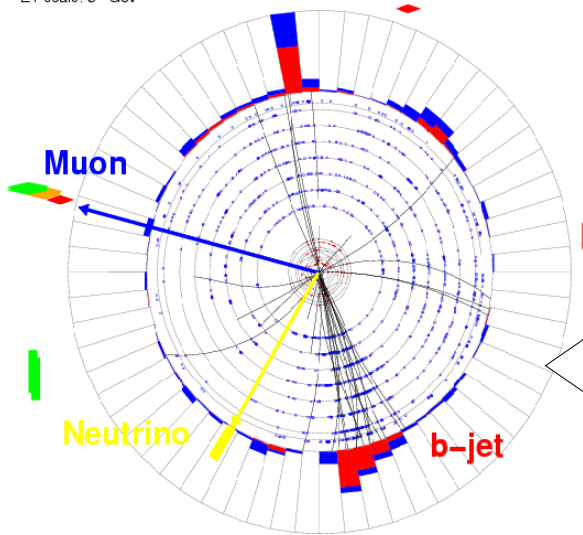


Frederik Beaujean, Allen Caldwell,
Daniel Kollar, Kevin Kröninger,
Shabnaz Pashapour, Arnulf Quadt

Motivation ◦ BAT overview ◦ MCMC ◦ A working example ◦ this course ◦ summary

Rur 190059 Evt49300403 Sat Mar 6 11:15:43 2004

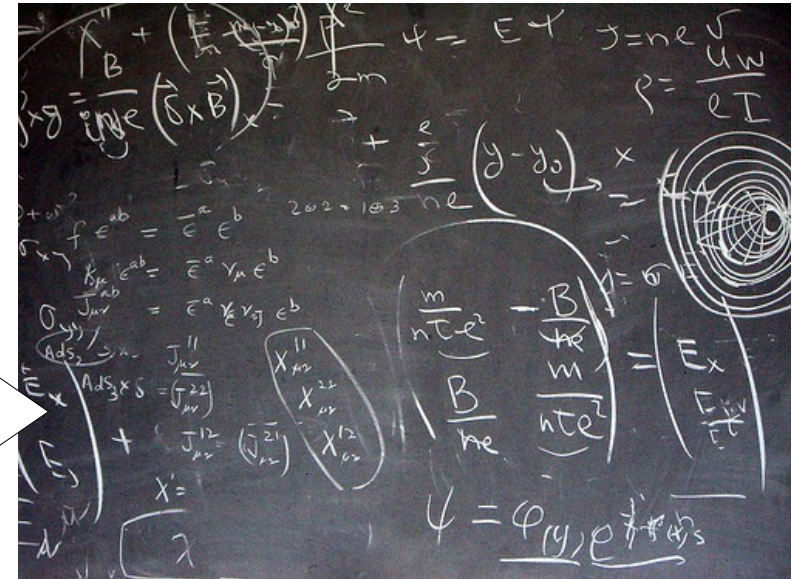
ET scale: 3 GeV



Experiment



Data analysis



Theory

Questions in data analysis:

- What does the data tell us about our model?
- Which model is favored by the data?
- Is the model compatible with the data?

- Parameter estimation
- Model comparison
- Goodness-of-fit test

Need methods and tools to extract information:

$$p(\vec{\lambda} | \vec{D}) = \frac{p(\vec{D} | \vec{\lambda}) p_0(\vec{\lambda})}{\int p(\vec{D} | \vec{\lambda}) p_0(\vec{\lambda}) d\vec{\lambda}}$$



Requirements

- Allow to phrase arbitrary models and data sets
- Interface to HEP software
- Estimate parameters (point estimates)
- Find probability densities (interval estimates)
- Propagate uncertainties
- Compare models
- Test validity of model against the data

Implementation:

- C++ library based on ROOT.
- Models are implemented as base classes and need to be defined by the user, or
- A set of of pre-defined models can be used.
- A set of algorithms can used to perform the actual analysis

Requirements

- Allow to phrase arbitrary models and data sets
- Interface to HEP software
- Estimate parameters (point estimates)
- Find probability densities (interval estimates)
- Propagate uncertainties
- Compare models
- Test validity of model against the data

Implementation:

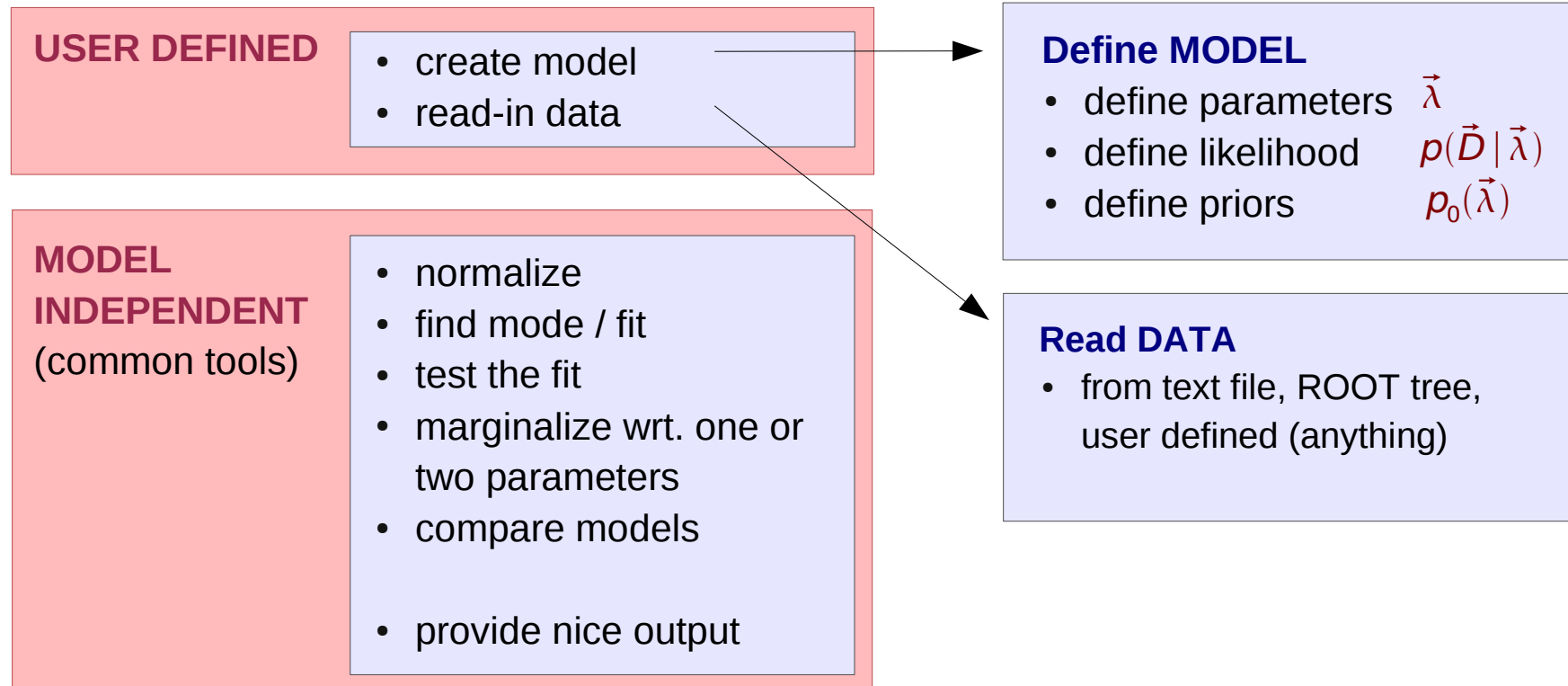
- Minimization can be done via a **Minuit** interface or via **Simulated Annealing**.
- Marginalization and uncertainty estimation can be done via **Markov Chain Monte Carlo (MCMC)**.
- Propagation of uncertainties (without Gaussian assumptions) can also be done via MCMC

Requirements

- Allow to phrase arbitrary models and data sets
- Interface to HEP software
- Estimate parameters (point estimates)
- Find probability densities (interval estimates)
- Propagate uncertainties
- Compare models
- Test validity of model against the data

Implementation:

- Direct comparison of model probabilities (Bayes factors)
- Integration methods from **Cuba** library linked
- Possibilities to do p-value tests



$$p(\vec{\lambda} | \vec{D}) = \frac{p(\vec{D} | \vec{\lambda}) p_0(\vec{\lambda})}{\int p(\vec{D} | \vec{\lambda}) p_0(\vec{\lambda}) d\vec{\lambda}}$$

Tools:

- **Point estimates:**

- Minuit
- Simulated Annealing
- **MCMC**
- simple Monte Carlo

- **Marginalization:**

- **MCMC**
- simple Monte Carlo

- **Integration:**

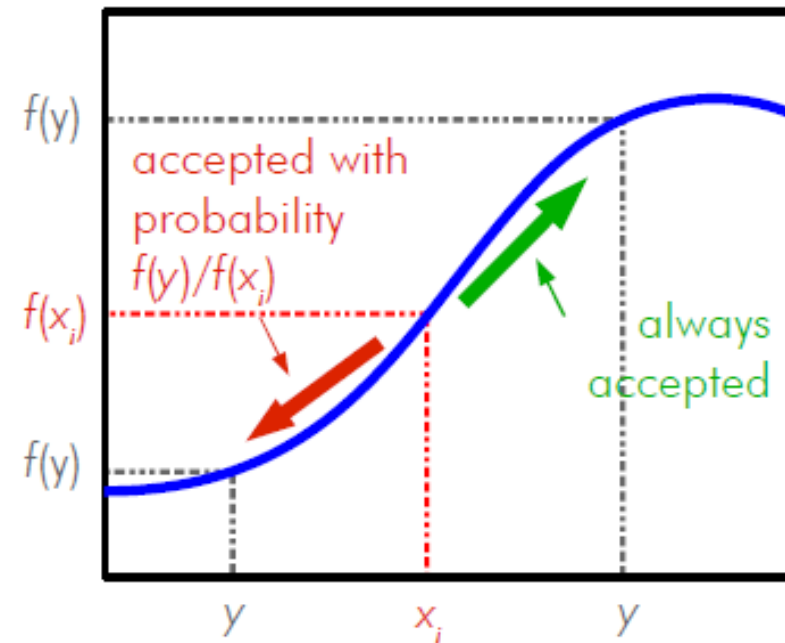
- sampled mean
- importance sampling
- CUBA (Vega, Suave, Divonne, Cuhre)

- **Sampling:**

- simple Monte Carlo
- **MCMC**

How does MCMC work?

- Output of Bayesian analyses are posterior probability densities, i.e., functions of an arbitrary number of parameters (dimensions).
- Sampling large dimensional functions is difficult.
- Idea: use random walk heading towards region of larger values (probabilities)
- **Metropolis algorithm:**

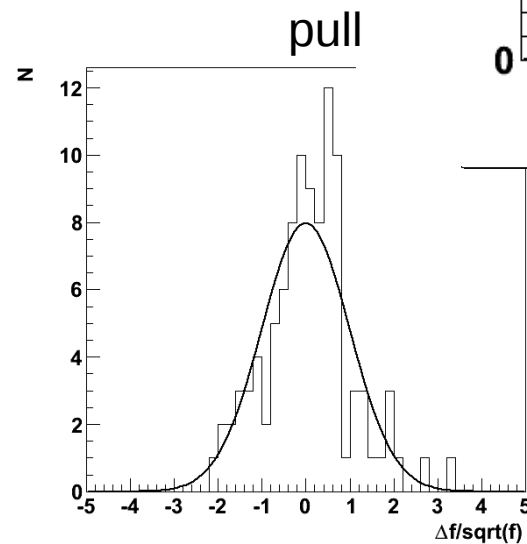
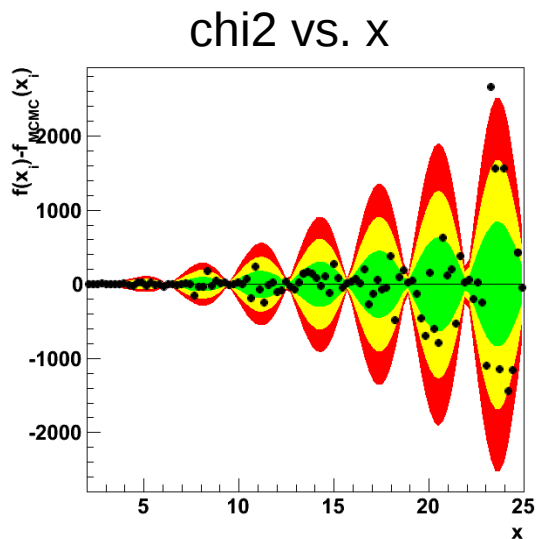
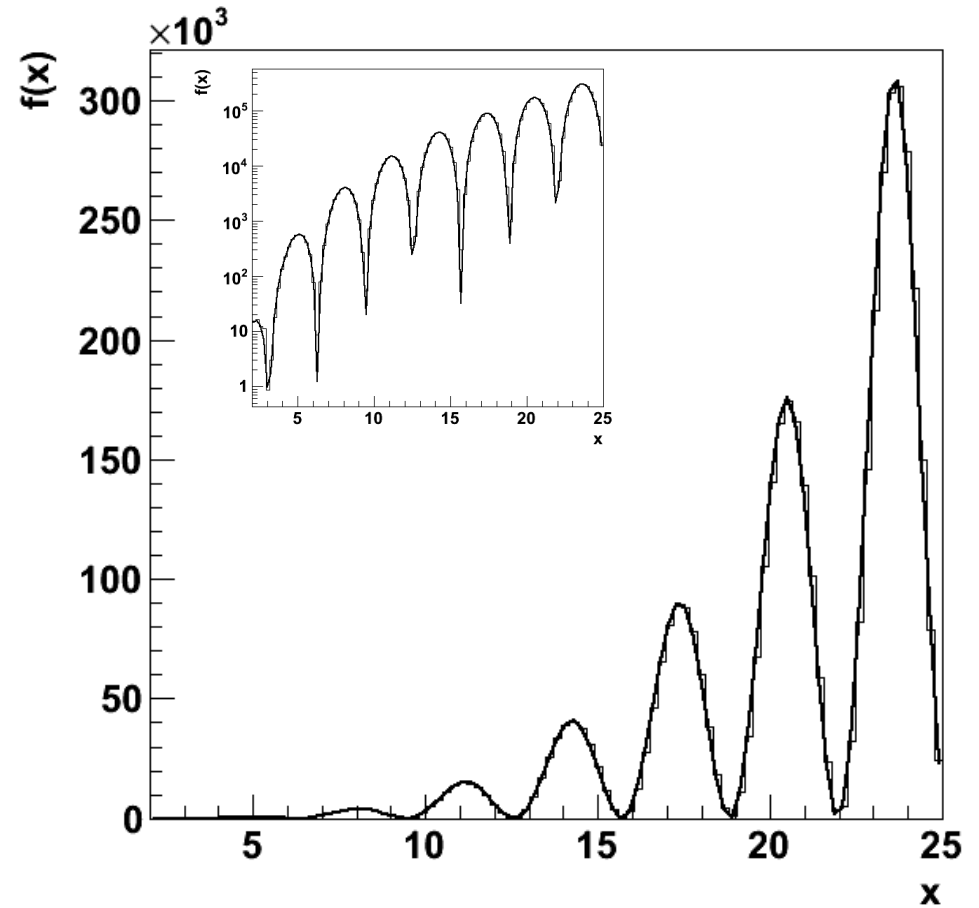


- Start at some randomly chosen x_i
- Randomly generate y around x_i
- If $f(y) \geq f(x_i)$, set $x_{i+1} = y$
- If $f(y) < f(x_i)$, set $x_{i+1} = y$ with probability $p = \frac{f(y)}{f(x_i)}$
- If y not accepted, stay where you are, i.e., set $x_{i+1} = x_i$
- Start over

N. Metropolis et al., J. Chem. Phys. 21 (1953) 1087.

Does it work for difficult functions?

- Test MCMC on a function:
 $f(x) = x^4 \cdot \sin(x^2)$
- Compare MCMC distribution to analytic function
- Several minima/maxima are no problem.
- Different orders of magnitude are no problem.



For more examples, see our test suite on the BAT web page.

How does MCMC help in Bayesian inference?

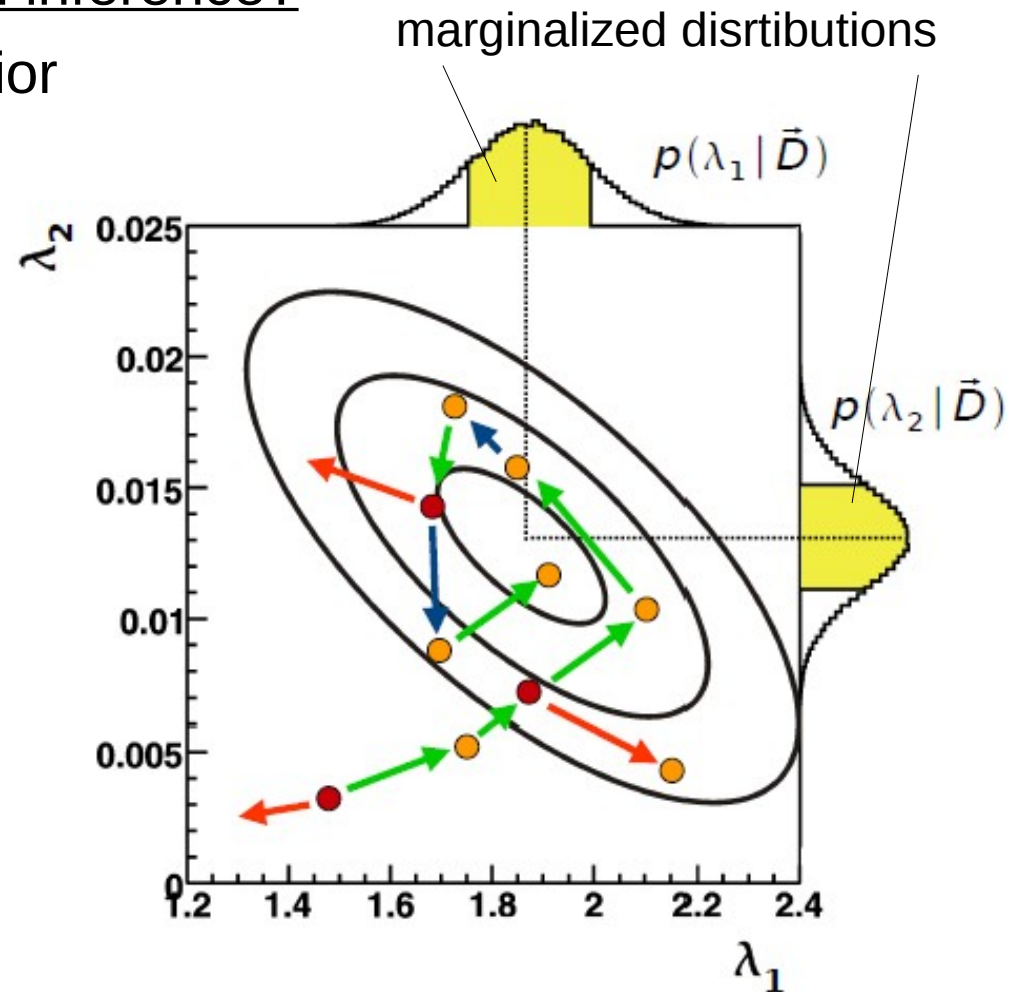
- Use MCMC to sample the posterior probability, i.e.

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

- Marginalization of posterior:

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

- Fill a histogram with just one coordinate while sampling
- Error propagation: calculate any function of the parameters while sampling
- Point estimate: find mode while sampling



Metropolis is ~3 lines of code, fairly easy, but ...

Technical details:

- How are the new points generated?
- How many points can I afford to throw away?
- How many iterations do we need?
- How correlated are the points?

Proposal function

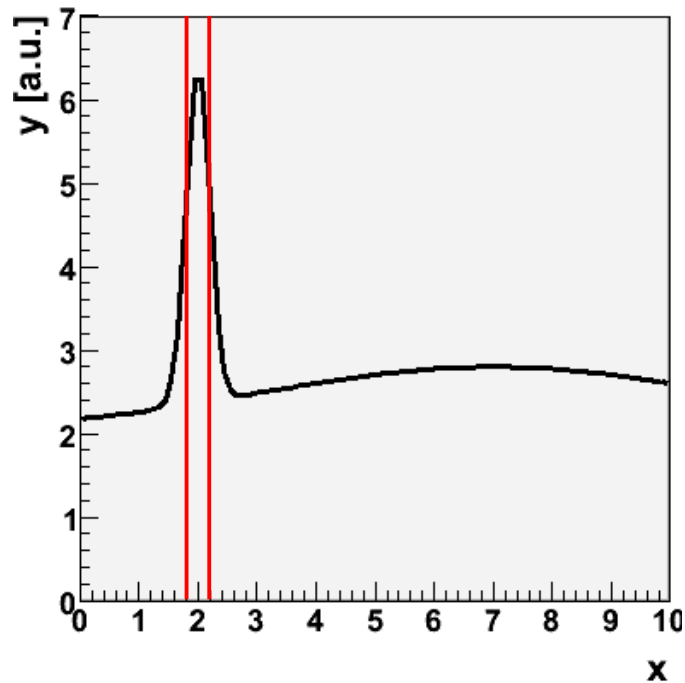
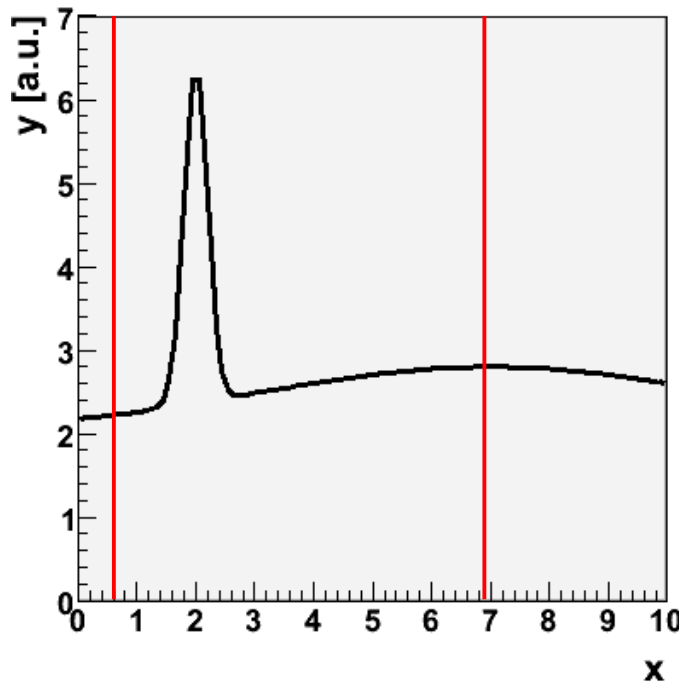
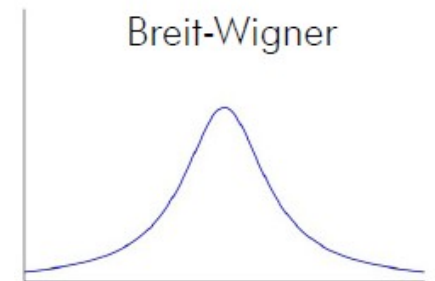
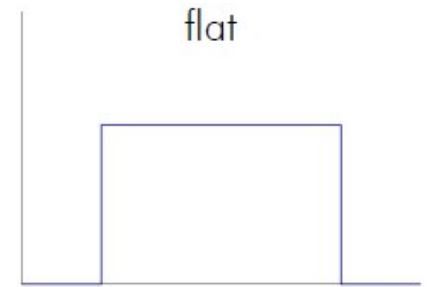
Efficiency

Convergence criterion

Auto-correlation/lag

How are the new points generated?

- **Proposal function:** probability density for taking one step during the random walk
- Should be independent of the underlying distribution, i.e., the same everywhere
- Shape is important (default: Breit-Wigner)
- Width defines efficiency = fraction of accepted points



- Small width = large efficiency
- Large width = small efficiency
- Trade off: efficiency ~25%

How many iterations do we need?

- MCMC distribution should converge asymptotically to underlying function.
- In practice: need to stop the chain at some point. Need criteria.
- Two strategies:
 - Single chain convergence
 - Multi-chain convergence
- **Single chain convergence:**
 - Could monitor auto-correlation
 - Very CPU-time intensive
 - Could be done offline
- **Multi-chain convergence:**
 - Test convergence of multiple chains to each other
 - Use Gelman&Rubin criterion

Gelman & Rubin convergence:

- Calculate average variance of all chains

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

- Estimate variance of target distribution

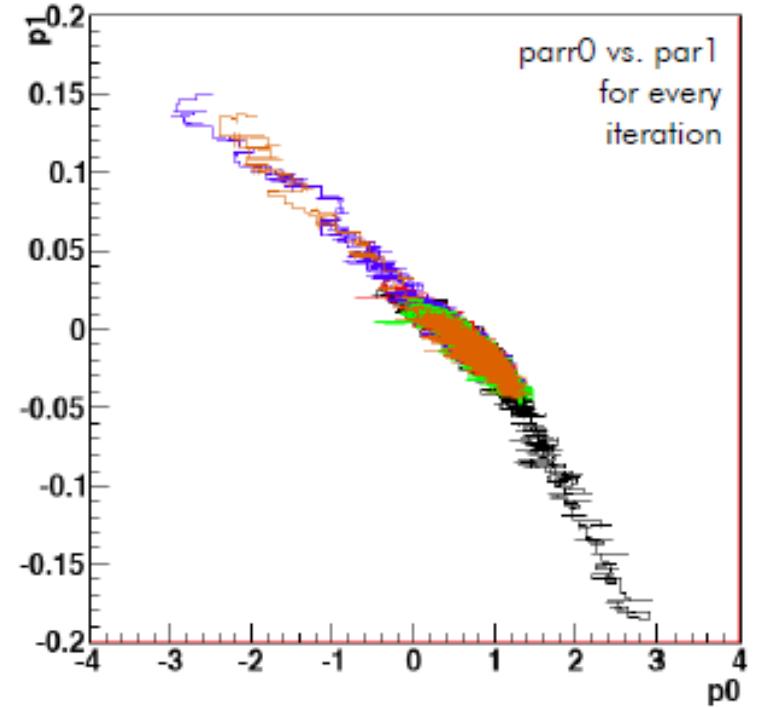
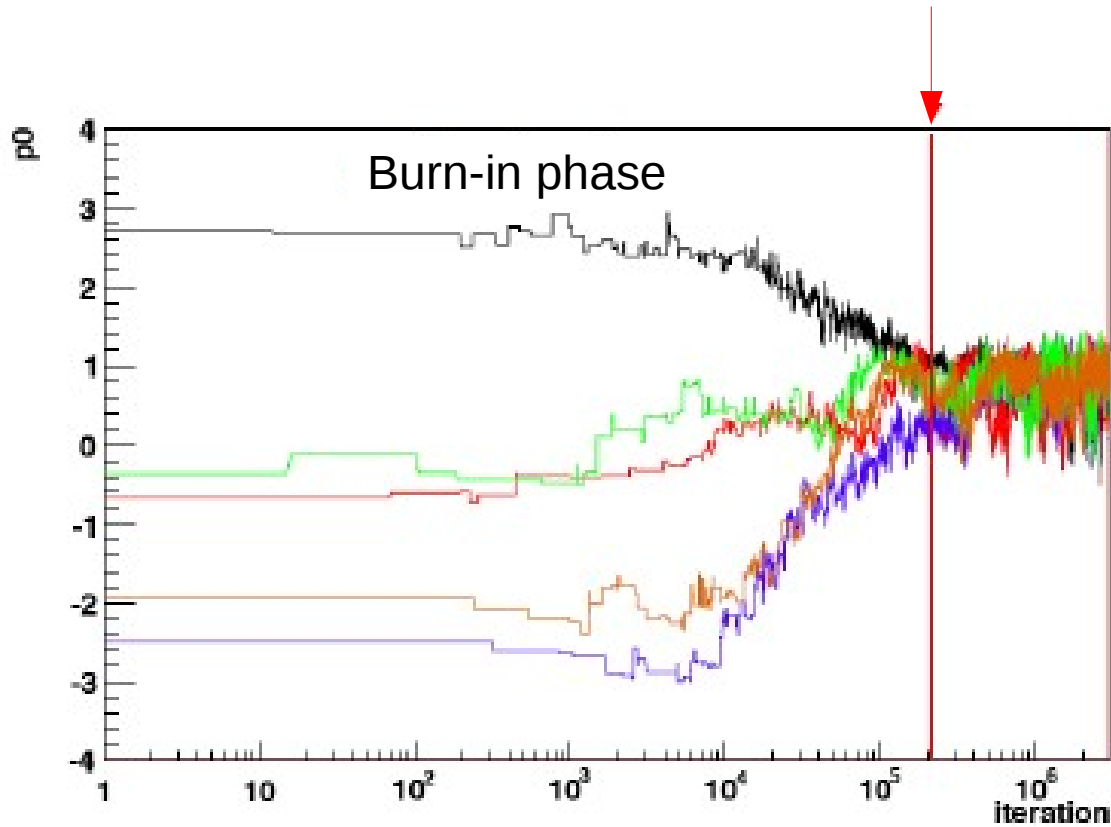
$$\hat{V} = \left(1 - \frac{1}{n}\right) W + \frac{1}{m-1} \sum_{j=1}^m (\bar{x}_j - \bar{x})^2$$

- Calculate ratio and compare with stopping criterion (relaxed version):

$$r = \sqrt{\frac{\hat{V}}{W}} < 1.x \quad (x = 0.1 \text{ default})$$

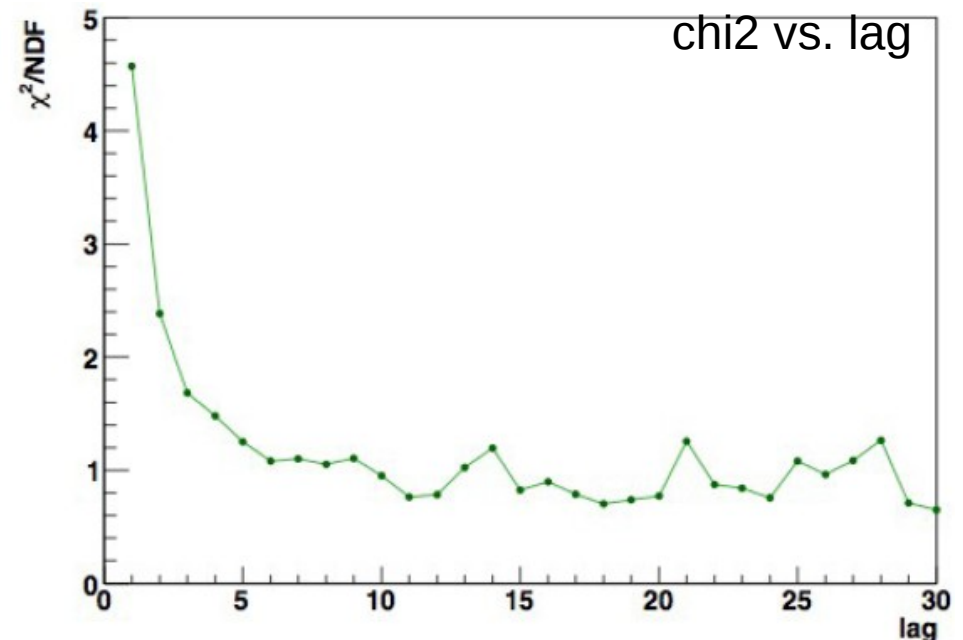
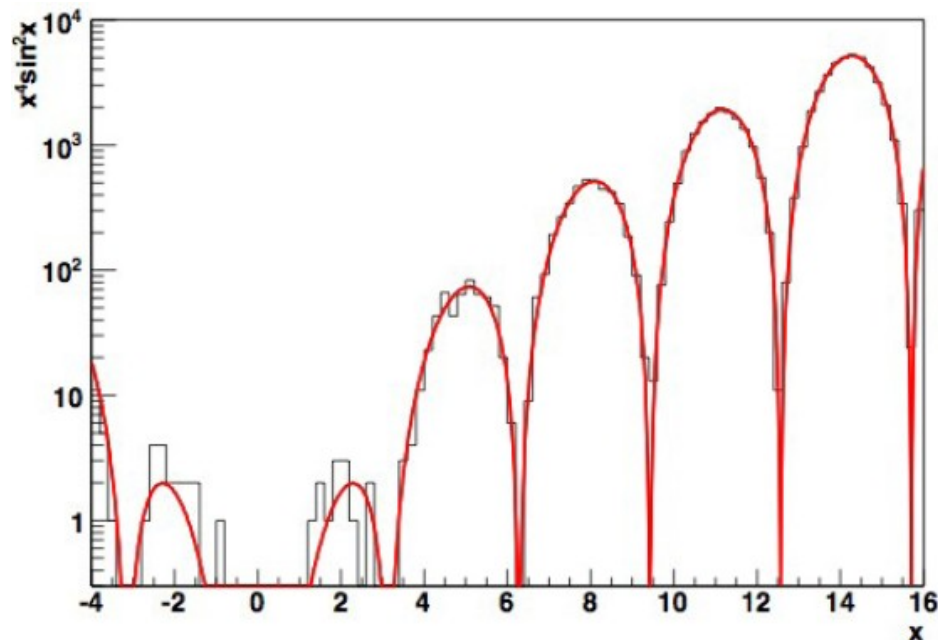
Gelman&Rubin, StatSci 7, 1992

Convergence a la Gelman & Rubin



How correlated are the points?

- True Monte Carlo and random walk create sets of points without (auto-correlation) while MCMC algorithm can cause **auto-correlation**, e.g., when rejecting a point (since the old one is taken again)
- Size of the correlation depends on the underlying posterior and the proposal function
- Can thin the MCMC sample by introducing a lag, i.e., take only every n^{th} point to calculate the marginalized distributions
- Cost: need to run a factor of n longer to get the same stat. precision



What exactly is being done in BAT?

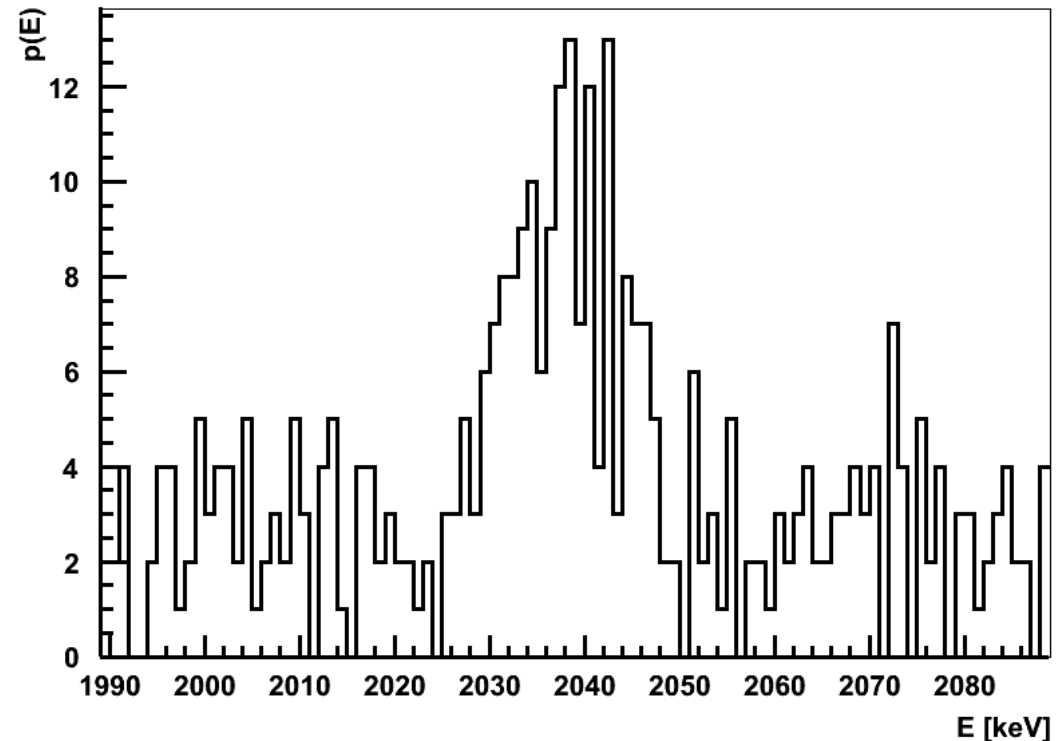
- **Step 1: Starting points**
 - Random within parameter space (default)
 - Center or user defined
- **Step 2: Burn-in**
 - Use multiple chains (default: 5)
 - Run until convergence is reached and chains are efficient
 - Or run until the maximum number of iterations is reached
 - Chains are efficient if the efficiency is between 15% and 50%
 - Run in sequences to adjust the width of the proposal functions:
 - If efficiency $> 50\%$: increase the width
 - If efficiency $< 15\%$: decrease the width
- **Step 3: Main run**
 - Use width obtained from efficiency optimization and convergence
 - Store information (next slide)

What is done in each step?

- **Marginalization:**
 - Fill 1-D and 2-D histograms
 - Large number: $N \cdot (N+1)/2$, e.g., for $N=50$ there are 1275 histograms
 - Individual histograms can be switched on/off
- **Optimization:**
 - Search for maximum of posterior
 - Not precise, but helpful as starting point for other algorithms
- **Error propagation:**
 - Calculate arbitrary (user-defined) functions from parameters
- **Misc:**
 - Write points to ROOT tree for offline analysis
 - Perform any user-defined analysis, histogram filling, etc.

Phrasing the problem:

- Estimate signal strength of Gaussian signal on top of flat background
- Data generated with the following settings:
 - **Gaussian signal:**
 - position $\mu = 2039$ keV
 - width $\sigma = 5$ keV
 - strength $\langle S \rangle = 100$
 - **Flat background:**
 - strength $\langle B \rangle = 3/\text{keV}$
- Number of events per bin fluctuate with Poisson distribution



Statistical model:

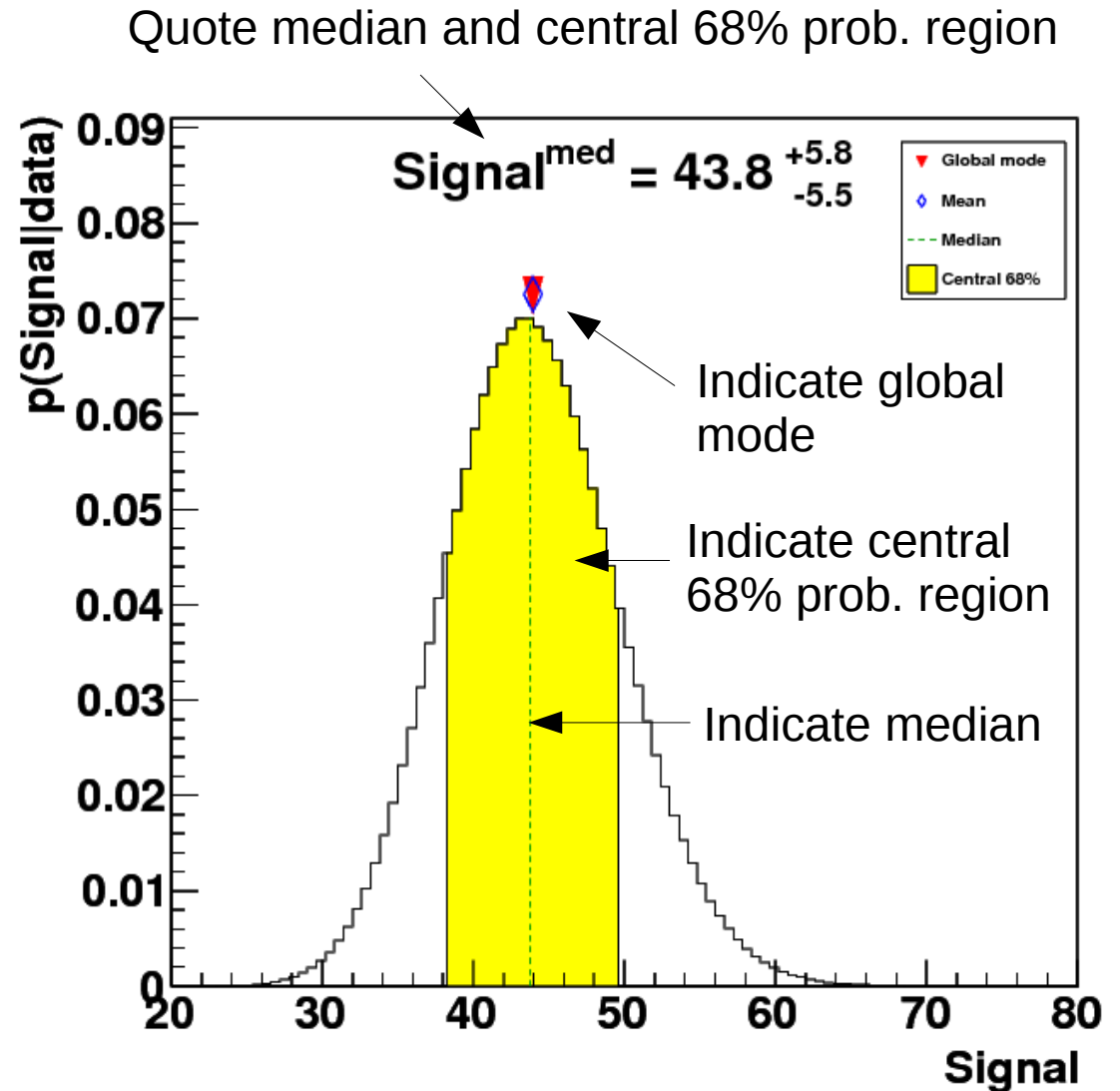
- Gaussian signal on top of flat background
- 4 (+2) fit parameters: Gauss (3) and flat (1) (+2 nuisance parameters for efficiency)
- **Prior knowledge:**
 - Background: 300 +/- 173 in 100 keV (e.g., from sideband analysis)
 - Signal strength: exponentially decreasing (e.g., theoretical intuition)
 - Signal position: flat (e.g., no idea about the mass of a resonance)
 - Signal width: 5 +/- 1 keV (detector resolution)
 - Signal and background efficiency fixed to 1 (in this example)
- **Statistical model:**
 - Bin data
 - Assume independent Poisson fluctuations in each bin

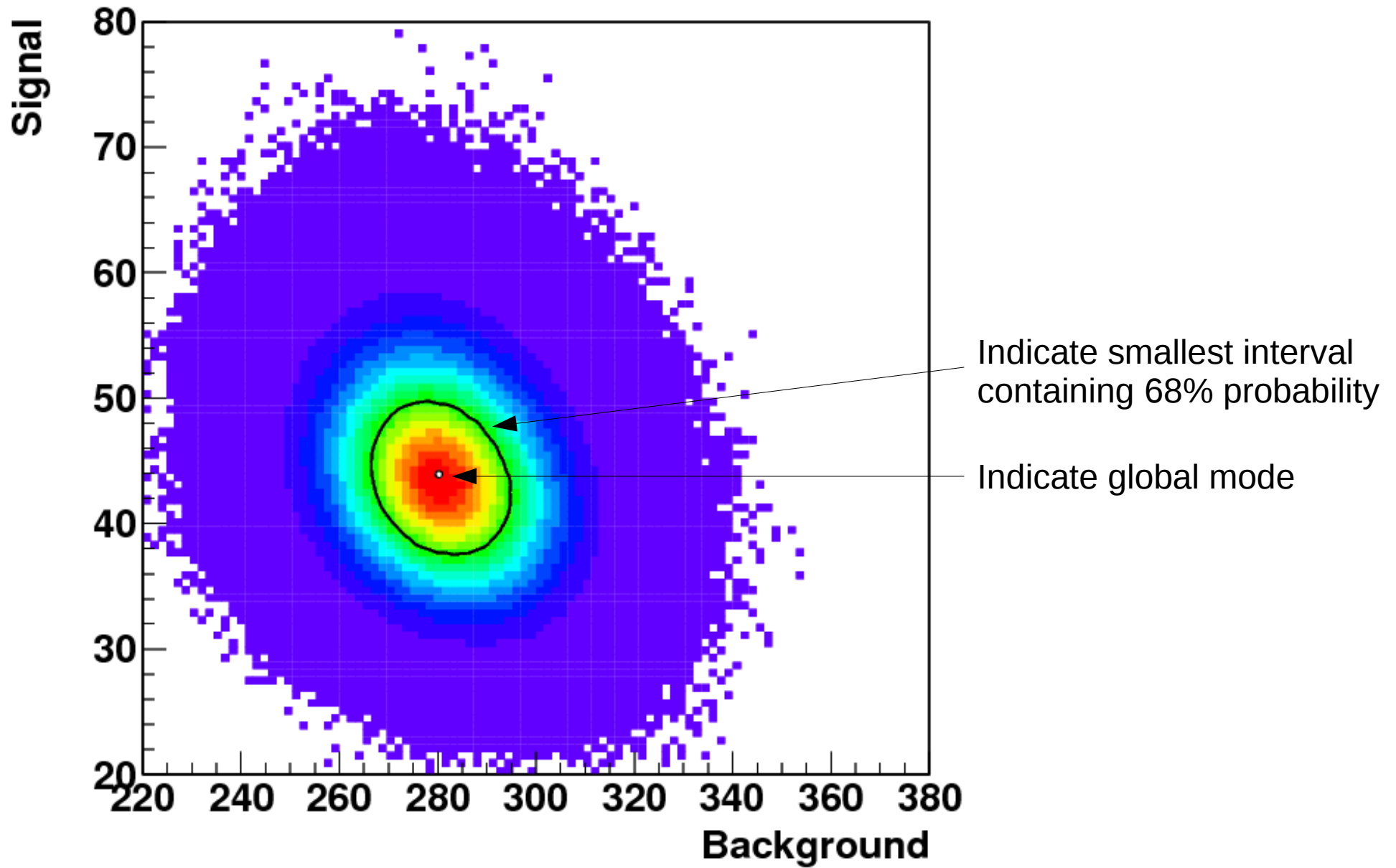
$$p(D|S, \mu, \sigma, B) = \prod_{i=1}^{N_{bins}} \frac{\lambda_i^{n_i}}{n_i!} e^{-\lambda_i}$$

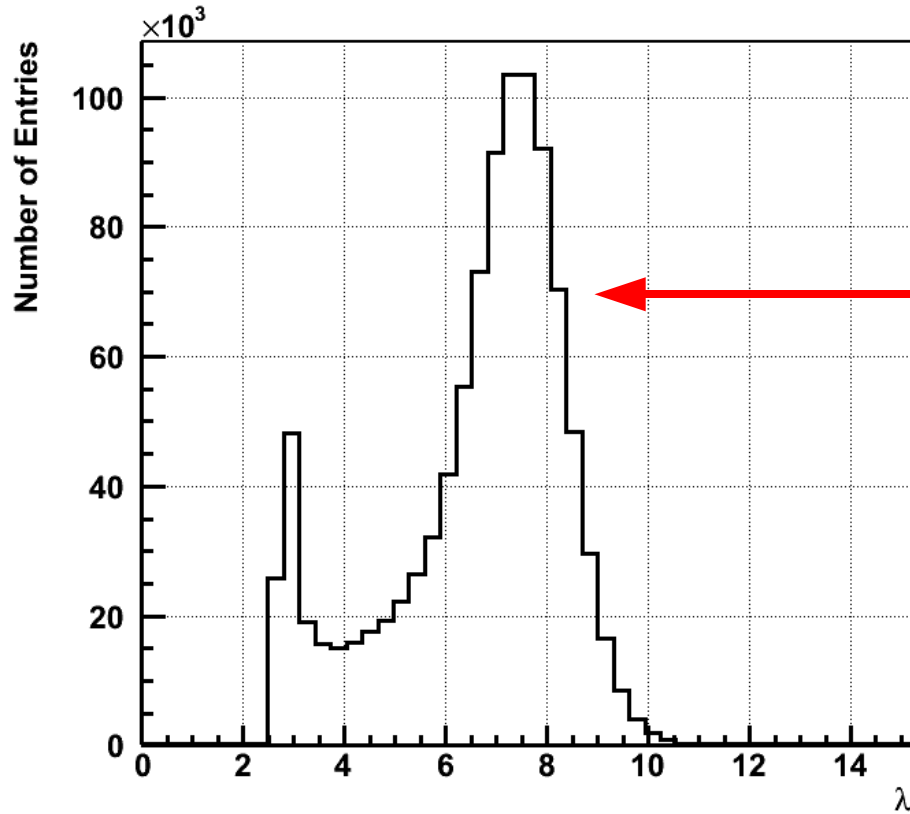
$$\lambda_i = \int_{\Delta_i} \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}} dx + \frac{B}{\Delta_i}$$

Marginalized distributions:

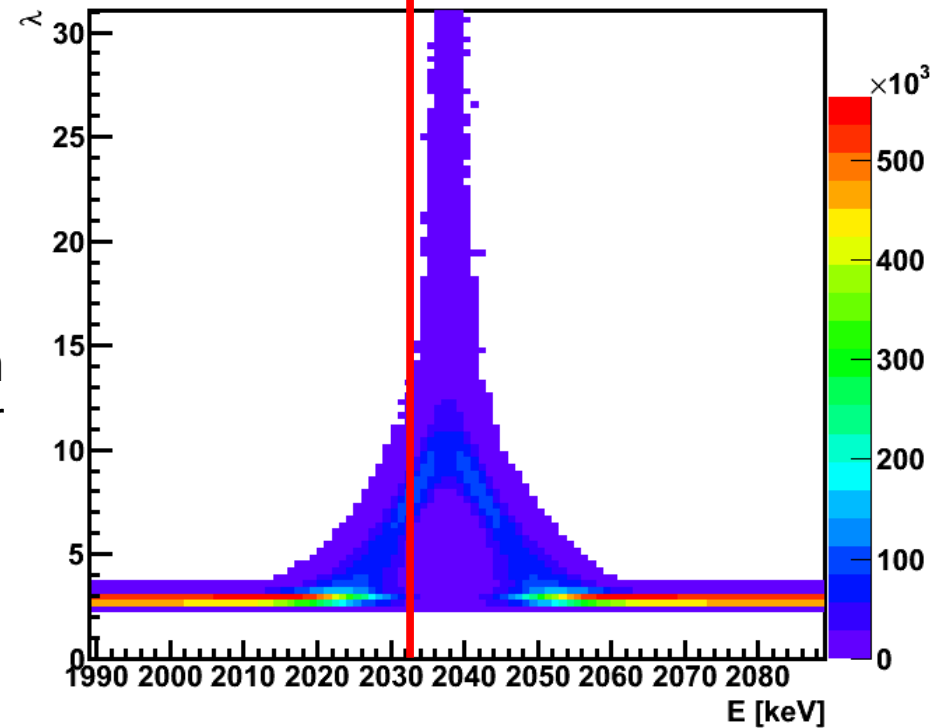
- Project posterior onto one parameter axis, i.e., integrate over all other parameters
- Global mode and mode of marginalized distribution do not have to coincide
- Full (correlated) information in Markov Chain
- **Default output:**
 - Mean +/- std. deviation
 - Median and central int.
 - Mode and smallest int.
- All 1-D and 2-D distributions are written out during main run





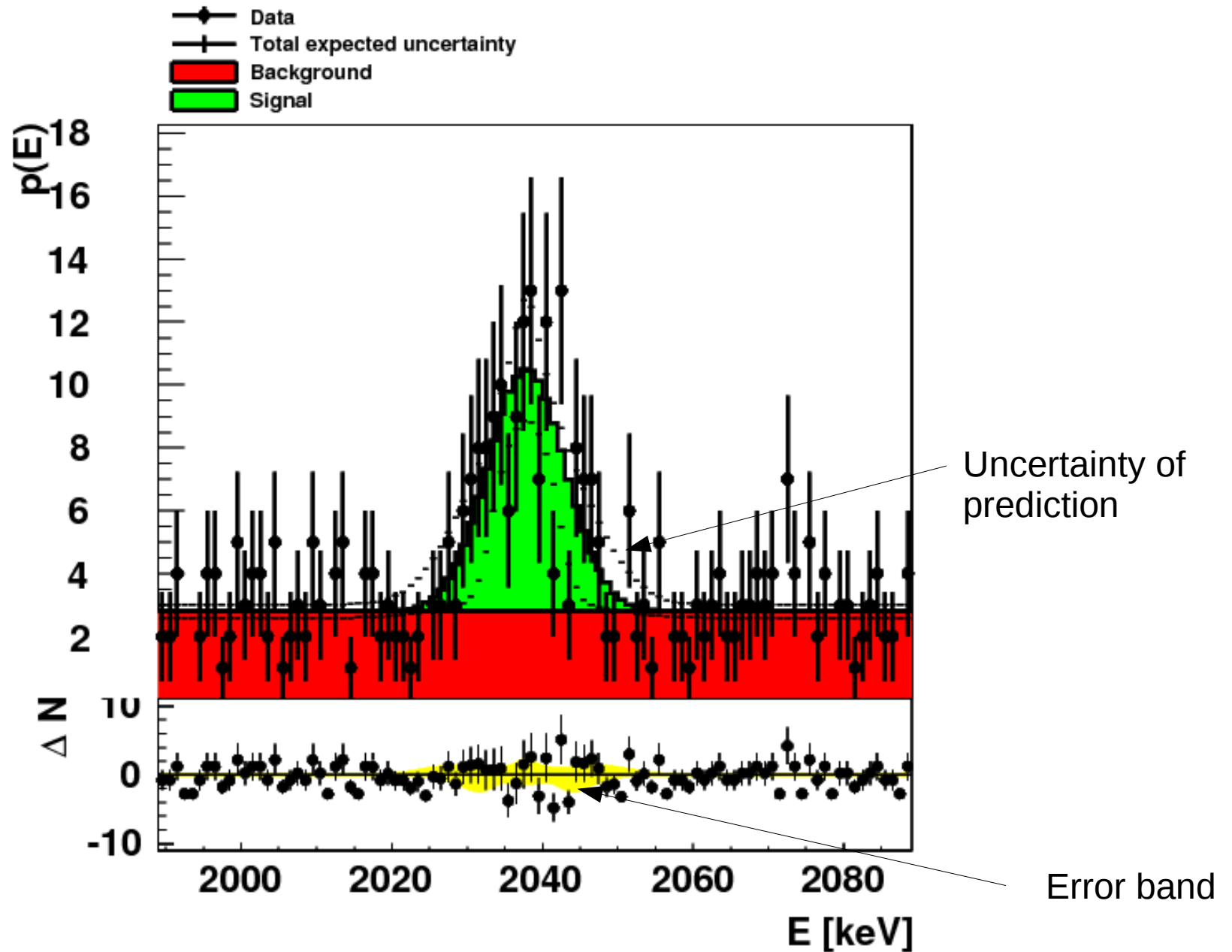


Posterior probability for the number of expected events with energy $E=2032$ keV



Sum of all possible fit functions weighted with posterior: calculate fit function at energy E for all parameter values

Use as error band



Results of the marginalization

=====

List of parameters and properties of the marginalized distributions:

(0) Parameter "Background":

Mean +- sqrt(V): 280.8 +- 13.16
Median +- central 68% interval: 280.7 + 13.2 - 13.02
(Marginalized) mode: 280
5% quantile: 259.2
10% quantile: 263.9
16% quantile: 267.7
84% quantile: 294.4
90% quantile: 297.7
95% quantile: 302.6

Smallest interval(s) containing 68% and local modes:
(266.4, 295.2) (local mode at 280 with rel. height 1; rel. area 0.6978)

(2) Parameter "Signal":

Mean +- sqrt(V): 43.94 +- 5.724
Median +- central 68% interval: 43.78 + 5.849 - 5.532
(Marginalized) mode: 43.7
5% quantile: 34.8
10% quantile: 36.71
16% quantile: 38.25
84% quantile: 49.88
90% quantile: 51.38
95% quantile: 53.62

Smallest interval(s) containing 68% and local modes:
(38, 50) (local mode at 43.7 with rel. height 1; rel. area 0.6821)

(4) Parameter "Signal mass":

Mean +- sqrt(V): 2038 +- 0.7871
Median +- central 68% interval: 2038 + 0.7806 - 0.7781
(Marginalized) mode: 2038
5% quantile: 2037
10% quantile: 2037
16% quantile: 2037
84% quantile: 2039
90% quantile: 2039
95% quantile: 2039

Smallest interval(s) containing 68% and local modes:
(2037, 2039) (local mode at 2038 with rel. height 1; rel. area 0.693)

...

Results of the optimization

=====

Optimization algorithm used: Metropolis MCMC

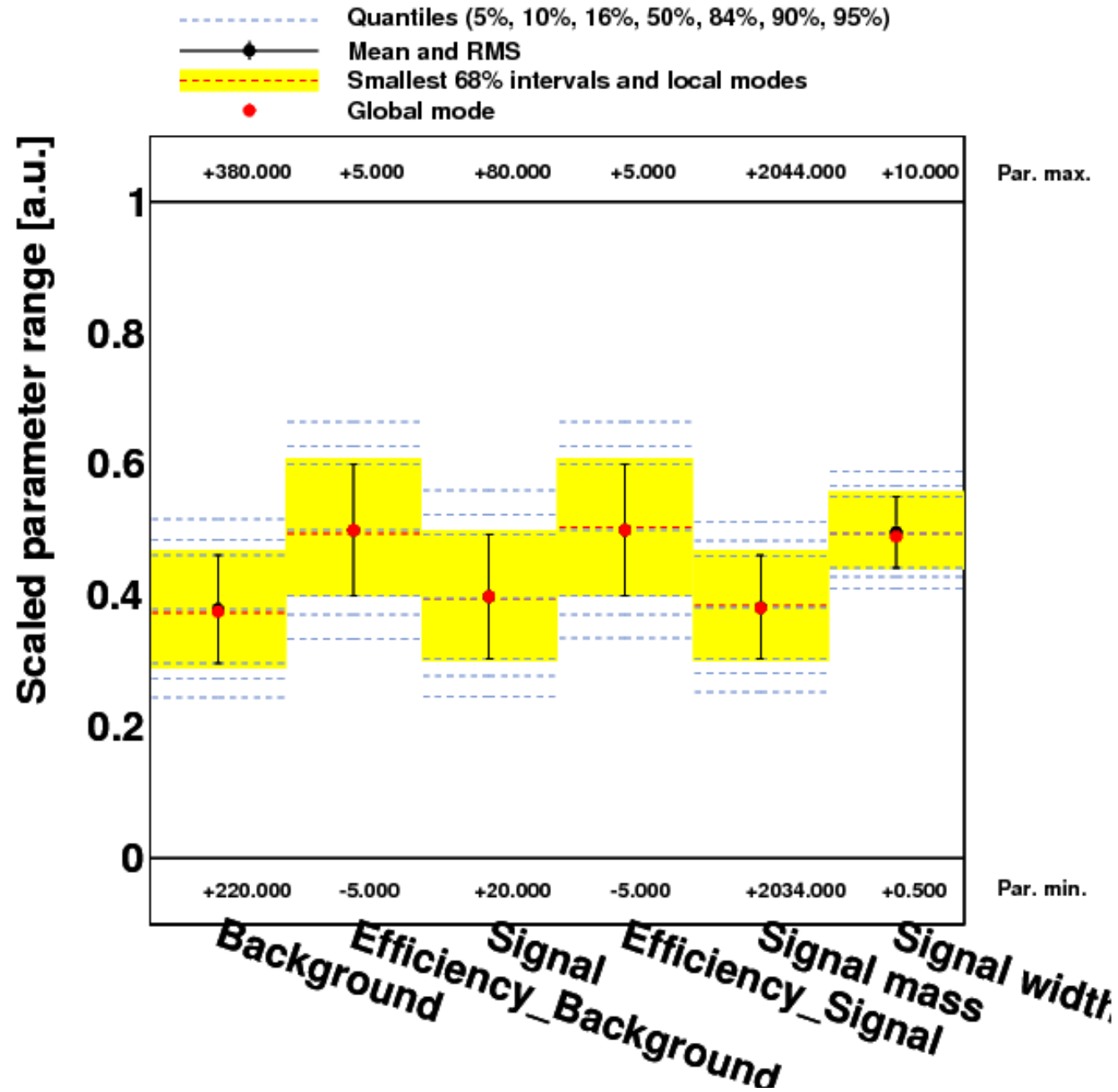
List of parameters and global mode:

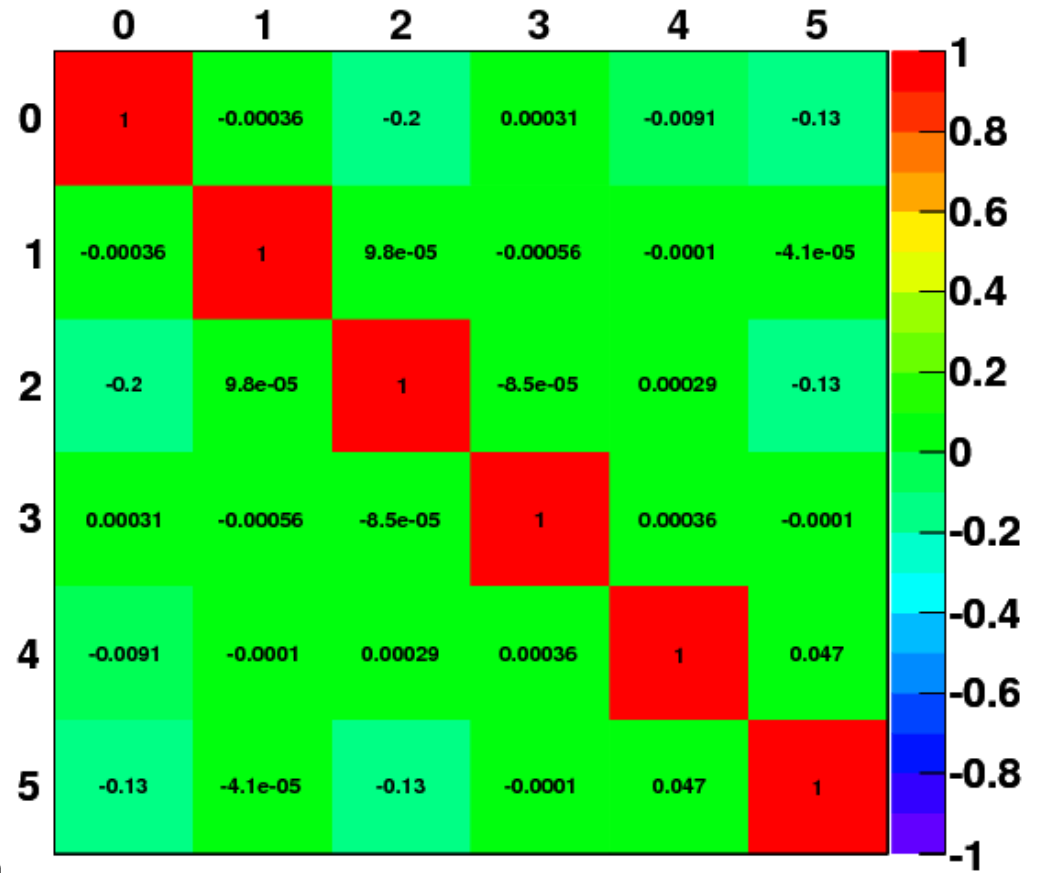
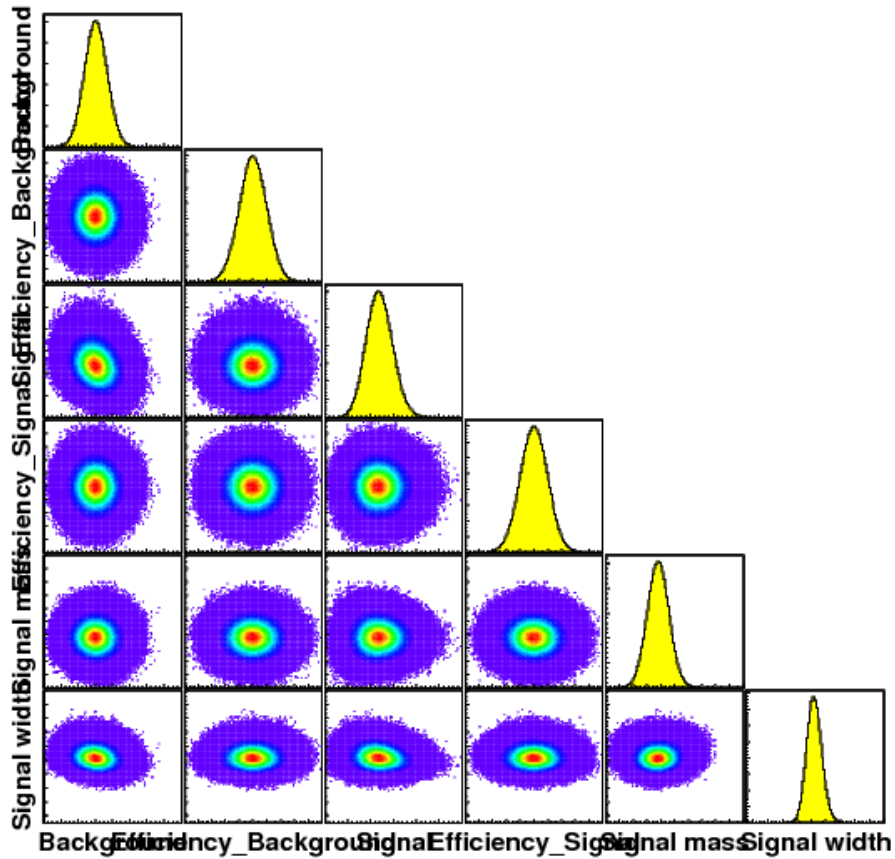
(0) Parameter "Background": 280.2 +- 13.08
(2) Parameter "Signal": 43.94 +- 5.674
(4) Parameter "Signal mass": 2038 +- 0.7652
(5) Parameter "Signal width": 5.159 +- 0.5012

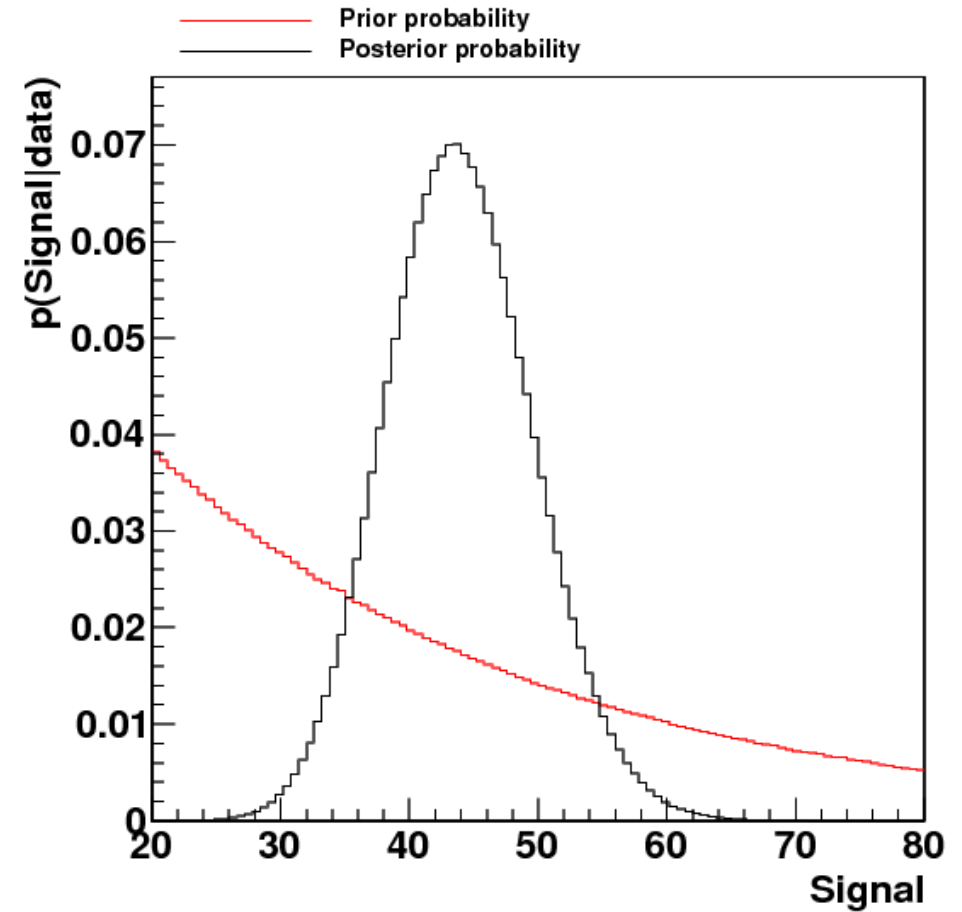
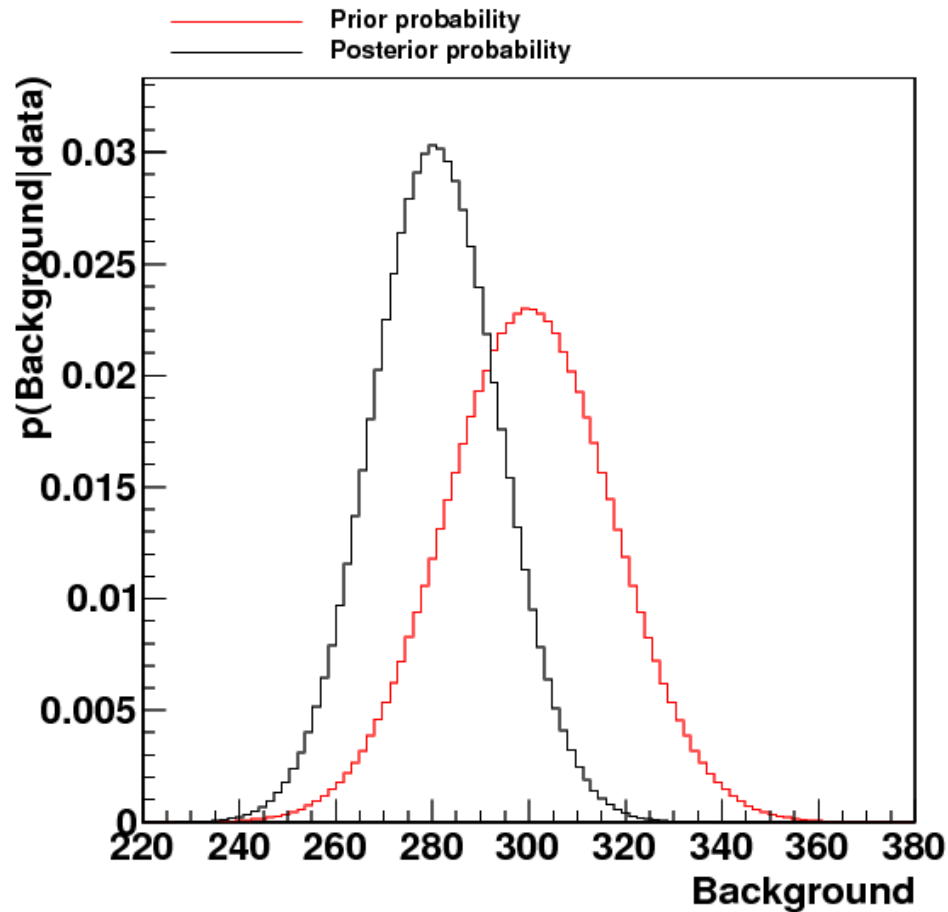
Status of the MCMC

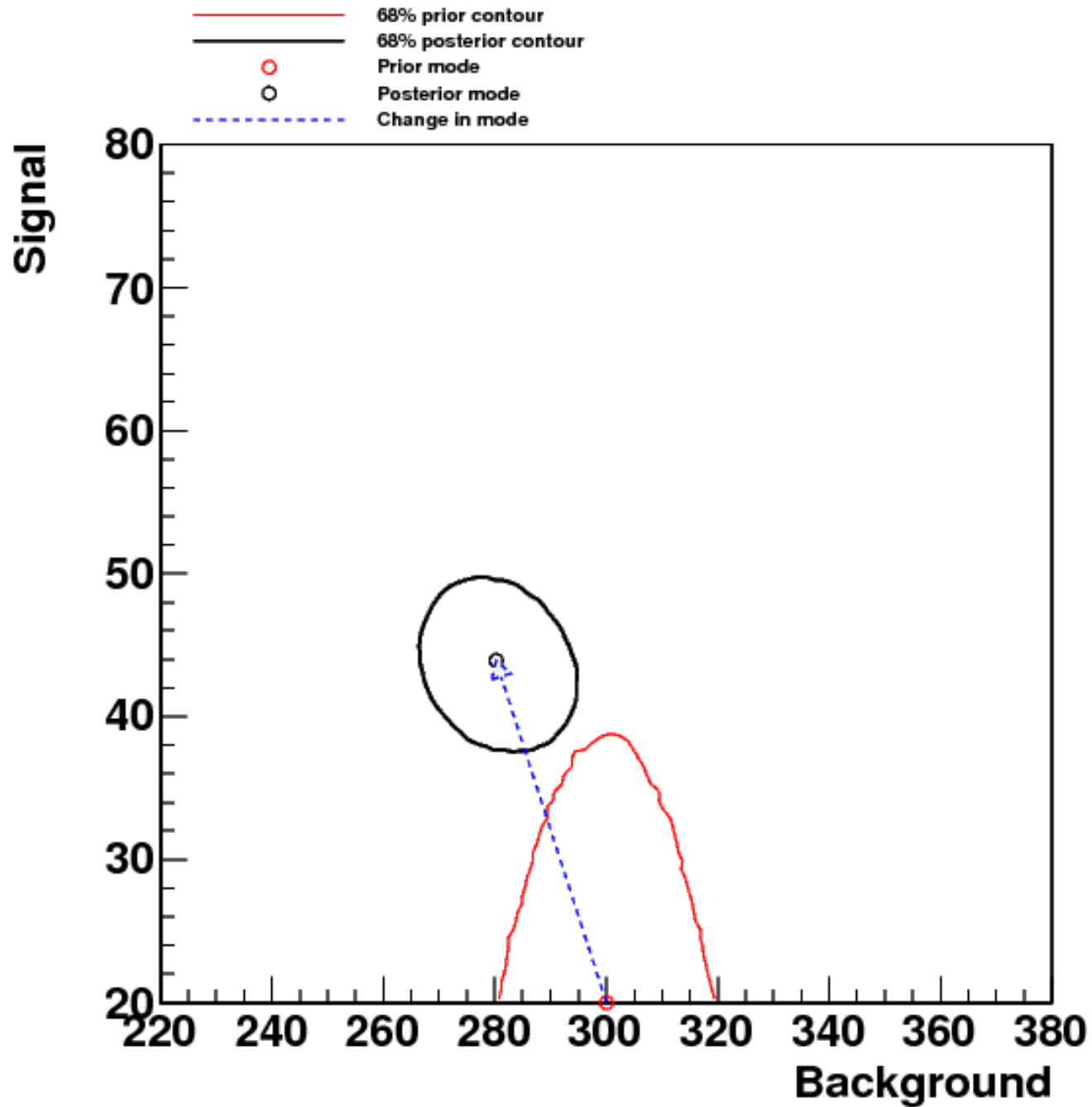
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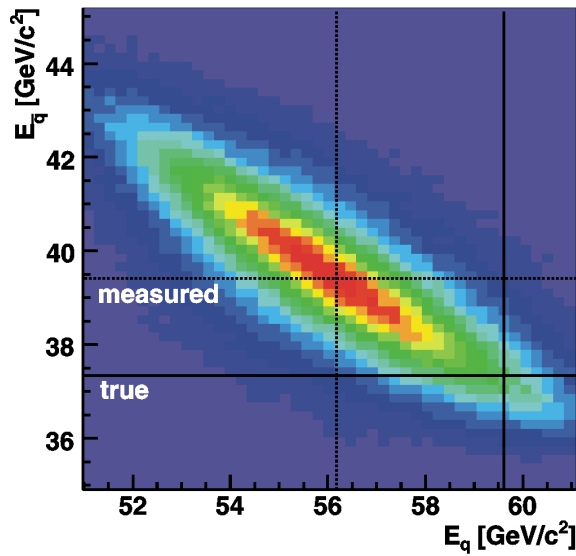
Convergence reached: yes
Number of iterations until convergence: 24000
Number of chains: 10
Number of iterations per chain: 10000000
Average efficiencies:
(0) Parameter "Background": 20.03%
(2) Parameter "Signal": 17.35%
(4) Parameter "Signal mass": 24.52%
(5) Parameter "Signal width": 19.56%



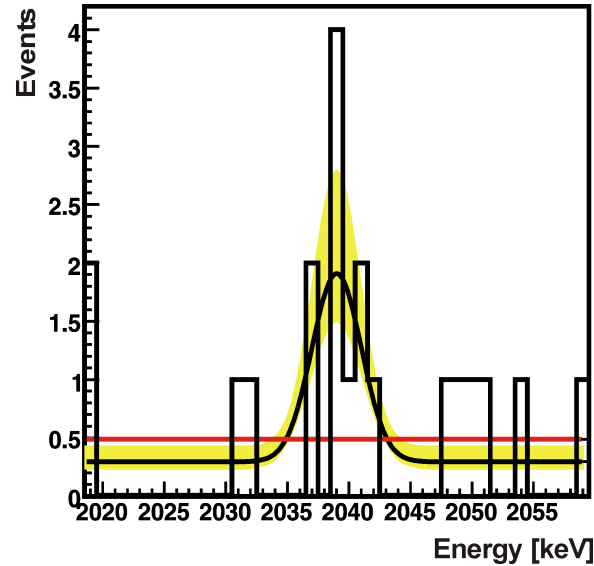




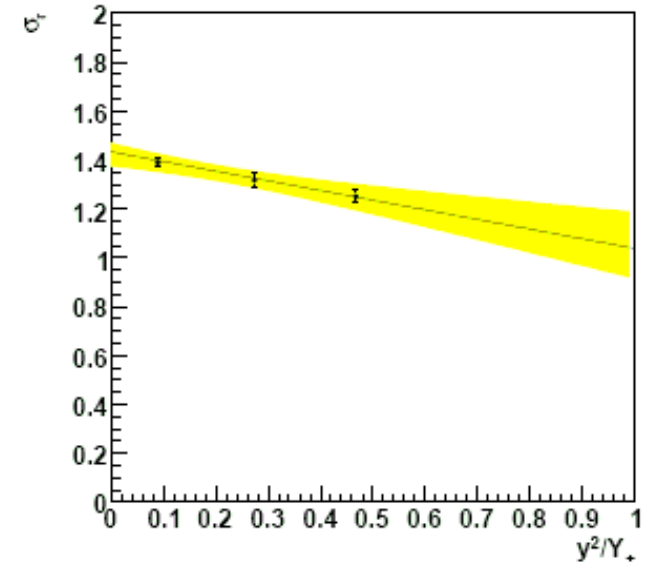




ATLAS:
Example of kinematic fitting in top quark decays



GERDA:
Fitting signal on top of a background



ZEUS:
Extraction of the longitudinal structure function

Contact:

- Web page: <http://www.mppmu.mpg.de/bat/>
- Contact: bat@mppmu.mpg.de
- Paper on BAT:
A. Caldwell, D. Kollar, K. Kröninger, BAT - The Bayesian Analysis Toolkit
Comp. Phys. Comm. 180 (2009) 2197-2209 [arXiv:0808.2552].



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Latest version: **0.4.2** (development)

Urgency: **medium**

Release date: **19.11.2010**

Source code: [BAT-0.4.2.tar.gz](#) (636kB)

[installation instructions](#) | [reference guide](#) | [changelog](#) | [performance testing](#)

Release notes

This version contains several improvements and updates and a few bugfixes. The most important changes are summarized below.

ROOT

- Required version of ROOT is now 5.22 or later.



Bayesian Analysis Toolkit

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BAT Tutorials

The tutorials are intended for the latest version of BAT (unless stated otherwise). However, after a new release they may need some adjustment to work. We try to do the necessary adjustments shortly after the release.

Title	Category	Level
Measuring a decay rate	counting experiment	basic
Estimating trigger efficiencies	fitting	basic
Charged current cross-section analysis	limit setting	basic
Signal search in the presence of background	hypothesis testing, template fitting	intermediate
Combination of cross-sections	combination	intermediate

Tutorials:

- Quite a few on the web
- Our program here:
 - Counting experiment
 - Charged-current cross-section analysis
 - Using BAT for searches

Summary:

- Bayesian inference requires some computational effort (e.g., nuisance parameters)
- Markov Chain Monte Carlo is the key tool to solve these issues
- **BAT is a tool to combine Bayesian inference with MCMC**
- Toolbox with more algorithms (integration, optimization, etc.)
- C++ library, modular, easy to use
- Informative output with predefined plots, numbers, etc.
- Did not talk about:
 - Hypothesis testing and goodness-of-fit
 - p-values
 - Bayes factors, information criteria
 - ...
- Upgrade of BAT ongoing, more to come
- **Participation and feedback are always welcome**