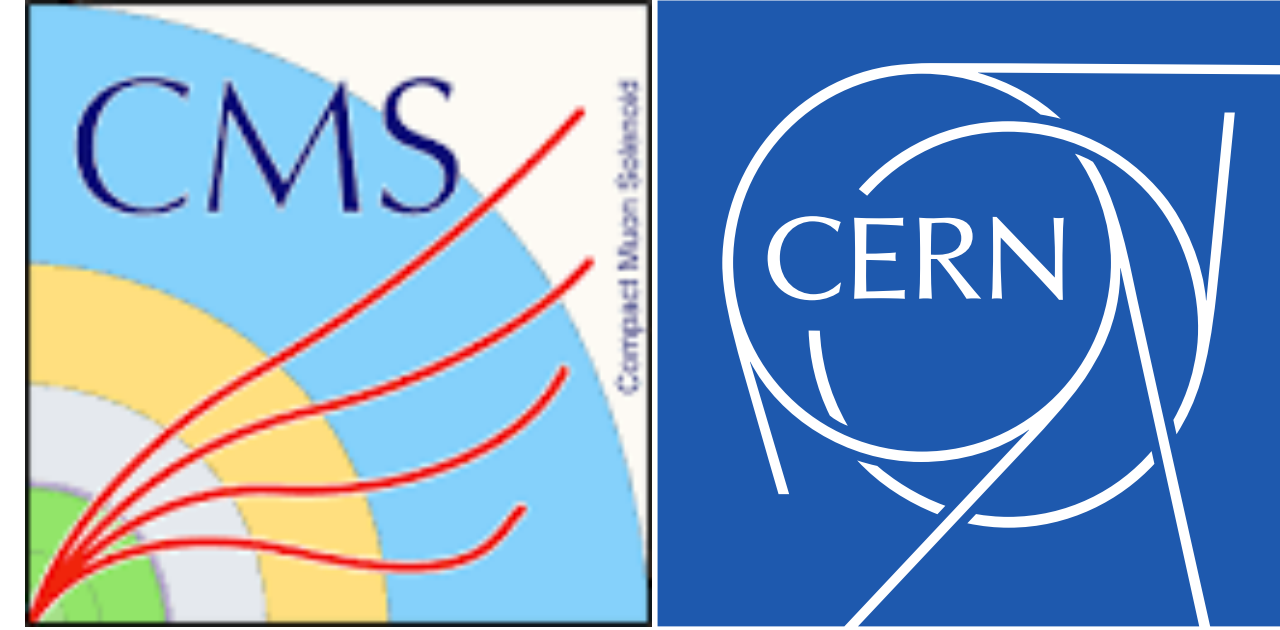




UNIVERSITY
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DATA ANALYSIS

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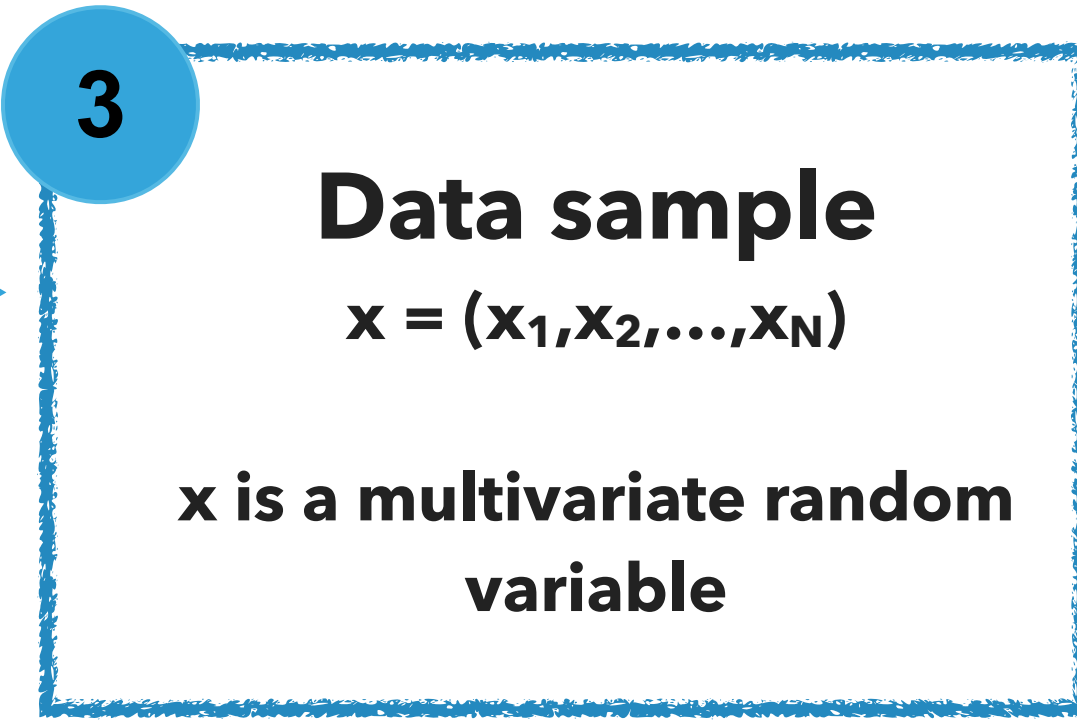
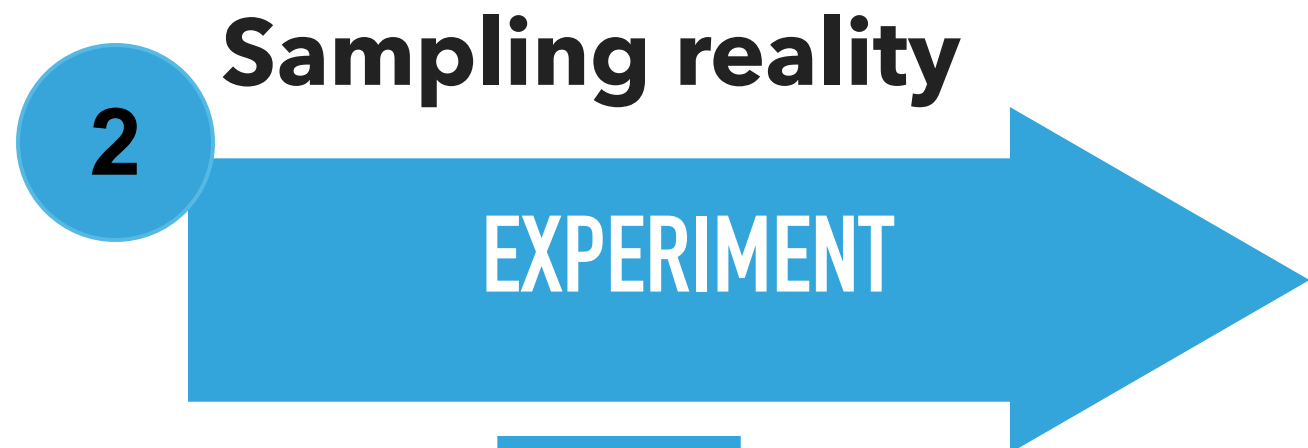
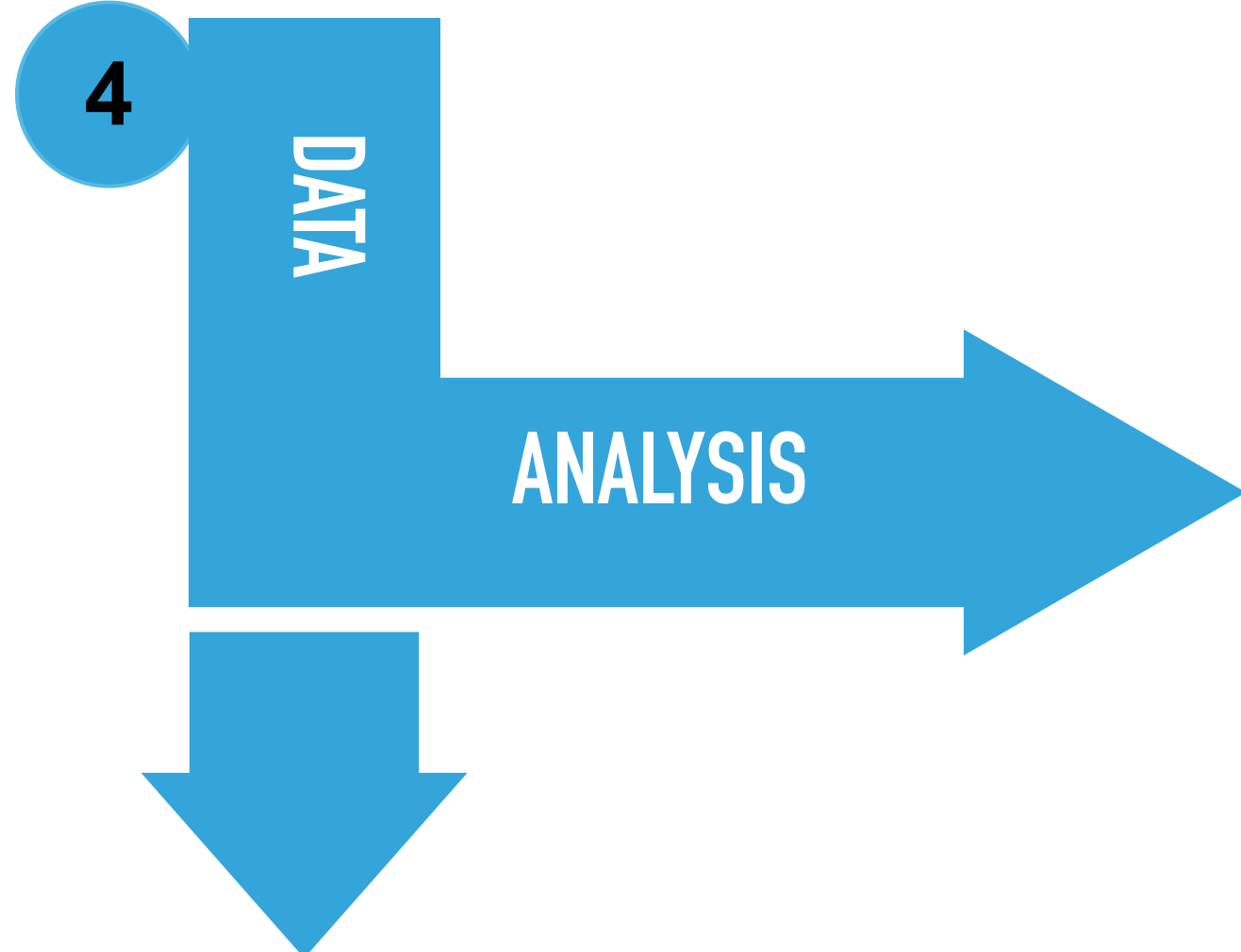
CERN School of Computing 2023, Tartu, Estonia

LECTURES OUTLINE

- 1) Introduction to Data Analysis
- 2) Probability density functions and Monte Carlo methods
- 3) Parameter estimation and Confidence intervals
- 4) Hypothesis testing and p-value

PARAMETER ESTIMATION AND CONFIDENCE INTERVALS

GENERAL PICTURE REMINDER



1

$$ie(W_\mu^- W_\nu^+ - W_\mu^+ W_\nu^-)|^2 -$$

Physical phenomena
$$- W_\nu^+ A_\mu) + ig' c_w (W_\mu^+ Z_\nu -$$

$$- \partial_\nu Z_\mu + ig' c_w (W_\mu^- W_\nu^+ - W$$

Described by PDFs,
depending on unknown parameters
with true values
 $\theta^{\text{true}} = (m_H^{\text{true}}, \Gamma_H^{\text{true}}, \dots, \sigma^{\text{true}})$

- 5 **Results**
- parameter estimates
 - confidence limits
 - hypothesis tests

- The parameters of a PDF are constants that characterise its shape:

$$f(x; \theta) = \frac{1}{\theta} e^{-\frac{x}{\theta}}$$

- where x is measured data, and θ are parameters that we are trying to estimate (measure)
- Suppose we have a sample of observed values $\vec{x} = (x_1, x_1, \dots, x_n)$
- Our goal is to find some function of the data to estimate the parameter(s)
 - we write the **parameter estimator** with a hat $\hat{\theta}(\vec{x})$
 - we usually call the procedure of estimating parameter(s): **parameter fitting**

EXAMPLE - PARAMETER ESTIMATION

- Task: find the average height of all students in a university on the basis of an (honestly selected) sample of N students
- Some possible ways of getting the result:
 - 1) Add up all the heights and divide by N
 - 2) Add up the first 10 heights and divide by 10. Ignore the rest
 - 3) Add up all the heights and divide by $N-1$
 - 4) Throw away the data and give the answer as 1.8 m
 - 5) Multiply all the heights and take the N -th root
 - 6) Choose the most popular height (the mode)
 - 7) Add up the tallest and shortest height and divide by 2
 - 8) Add up the second, fourth, etc. and divide by $N/2$ for N even or by $(N-1)/2$ for N odd

● Consistent

- Estimate converges to the true value as amount of data increases

$$\hat{\theta} \xrightarrow{\text{more data}} \theta^{true}$$

● Unbiased

- Bias is the difference between expected value of the estimator and the true value of the parameter

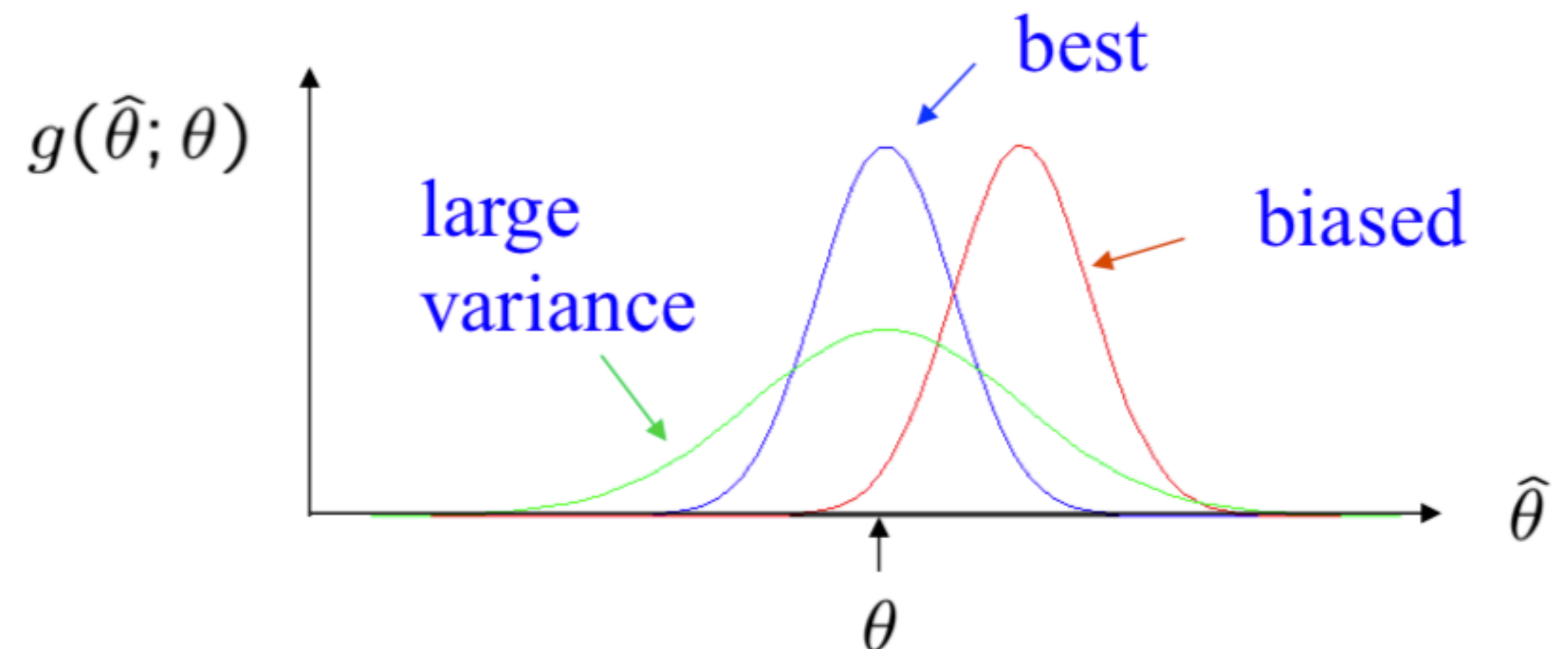
$$b = E(\hat{\theta}) - \theta^{true} = 0$$

● Efficient

- Its variance is small

● Robust

- Insensitive to departures from assumptions in the PDF



BONUS PROBLEM - 3

Quarks produced in high energy collisions will hadronize and form “jets” of particles. We call jets coming from the hadronization of b quarks “b-jets”. Algorithms to identify b-jets, referred to as b-tagging, will tag jets with a high probability to be b-jets. Their performance is characterized by two numbers:

1. The efficiency to tag real b-jets: $\varepsilon_b = P(\text{tag} \mid \text{b jet})$
2. The mistag rate to tag light flavour jets: $\varepsilon_{\text{mistag}} = P(\text{tag} \mid \text{light flavour jet})$

In an event with n_b true b-jets and n_{light} true light jets what is the probability to find n_{tag} tagged jets given ε_b and $\varepsilon_{\text{mistag}}$?

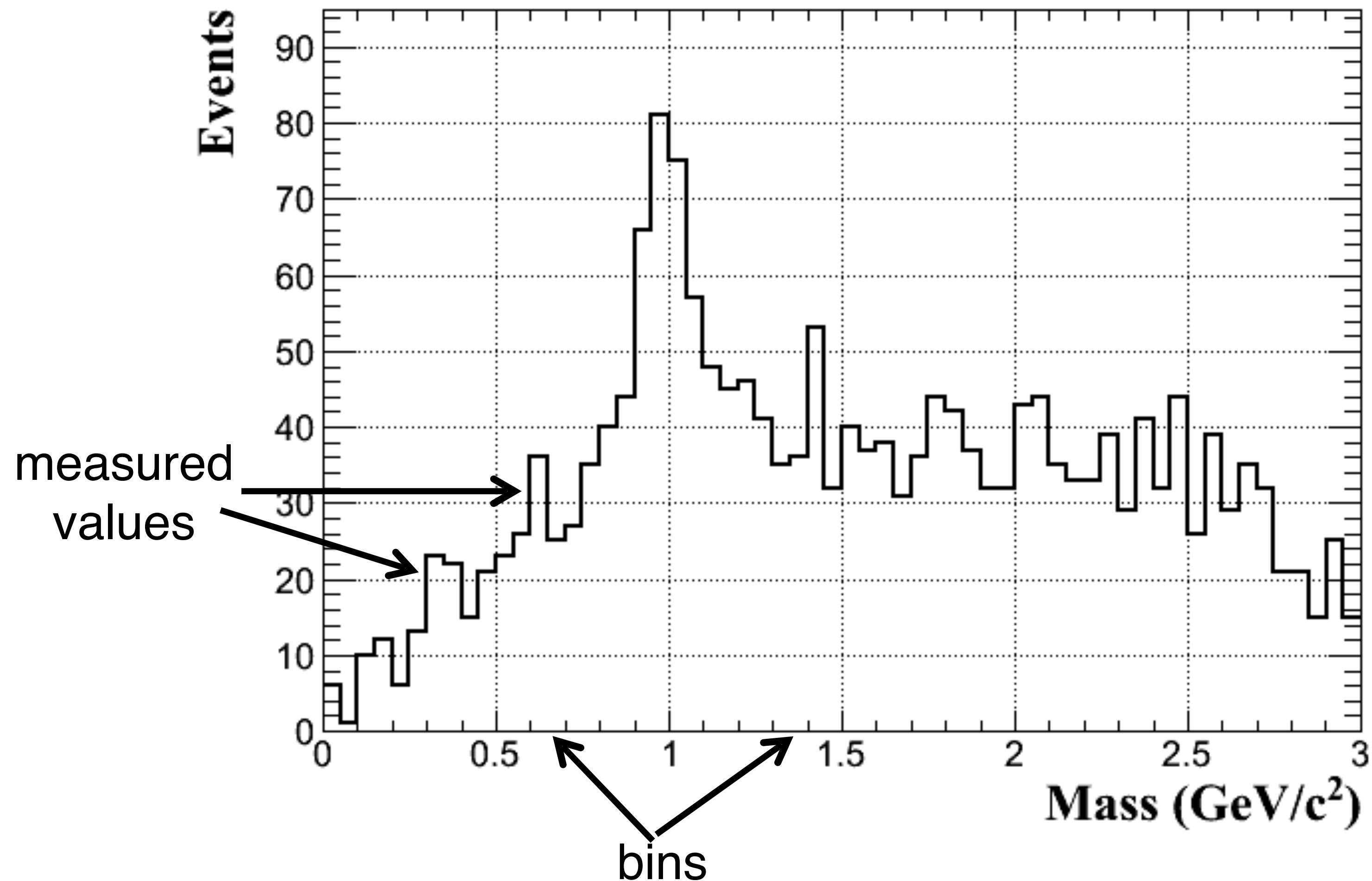
As example consider the process in which the Higgs boson is produced together with a top anti-top pairs, with the H decaying into a pair of b-jets, one of the top quarks decaying hadronically and the other semileptonically: $ttH \rightarrow blvl + bqq' + bb$ (4b – jets + 2 light jets)

What is the probability to tag 2, 3, 4, 5 or 6 jets if $\varepsilon_b = 68\%$ and $\varepsilon_{\text{mistag}} = 1\%$

Hint! Let binomial distribution and python help you solve this one!

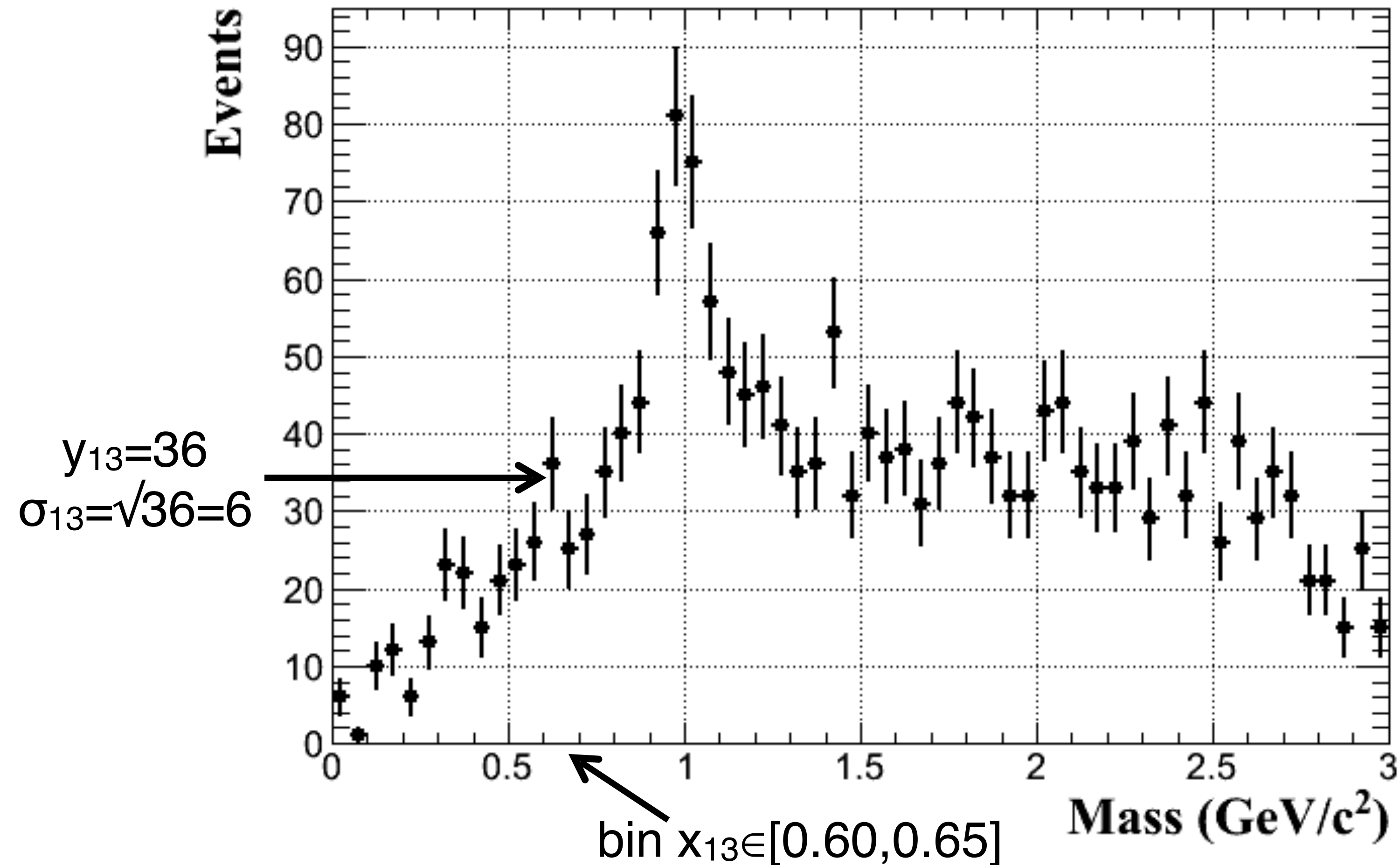
EXAMPLE IN HEP - HISTOGRAM FITTING

- In counting experiments we usually represent data in histograms
- In the following example we will study a particle mass histogram

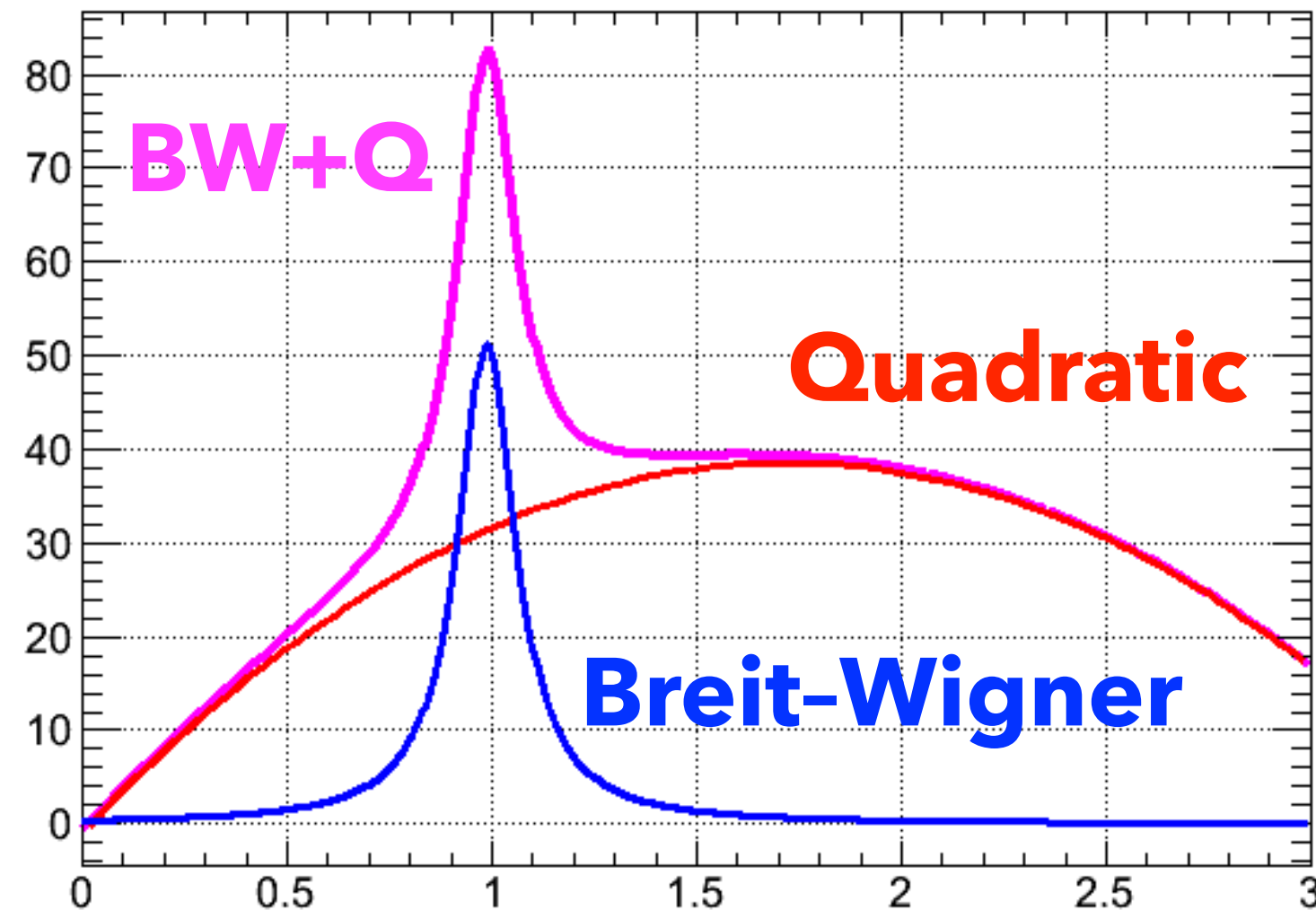


EXAMPLE IN HEP - HISTOGRAM FITTING

- Measured values have statistical uncertainties so it is better to represent them with points and error bars
 - each bin has a Poisson uncertainty



- Therefore we have
 - a set of precisely known values $\mathbf{x} = (x_1, \dots, x_N)$ - **histograms bins**
 - At each x_i
 - a measured value y_i - **number of events in a given bin**
 - a corresponding **error on measured value** σ_i
- We are missing a theoretical PDF $f(x_i; \theta^{true})$ with true parameters θ^{true} so we can calculate **parameter estimator** $\hat{\theta}$

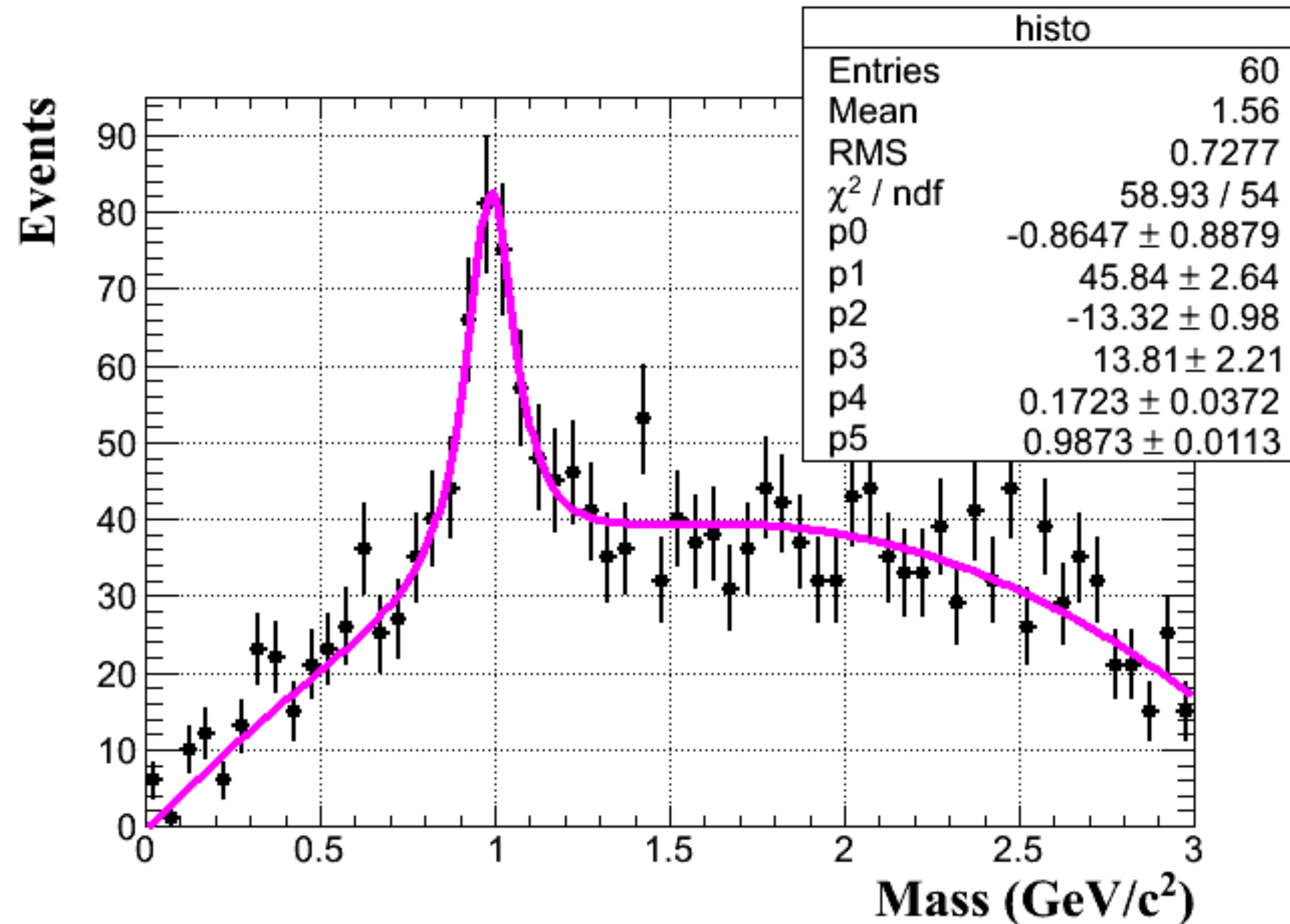


$$BW(x; D, \Gamma, M) \approx \frac{D\Gamma}{(x^2 - M^2)^2 + 0.25\Gamma^2}$$

$$Q(x; A, B, C) = A + Bx + Cx^2$$

EXAMPLE IN HEP - HISTOGRAM FITTING

$$f(x_i, \theta^{true}) = f(x_i; D, \Gamma, M, A, B, C) = BW(x_i; D, \Gamma, M) + Q(x_i; A, B, C)$$

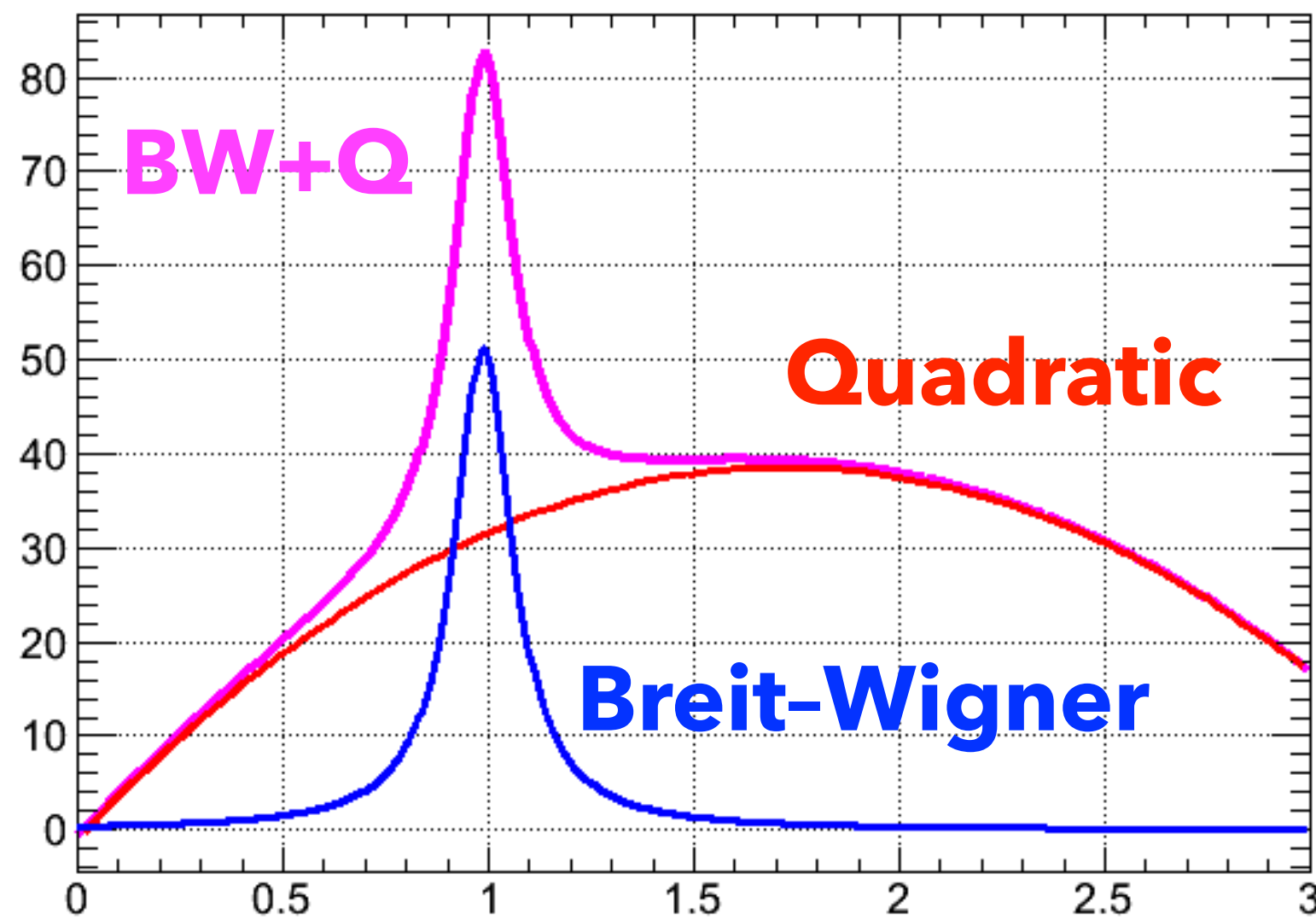


EXAMPLE IN HEP - HISTOGRAM FITTING

1

Physical phenomena

Described by a theory



2

Sampling reality

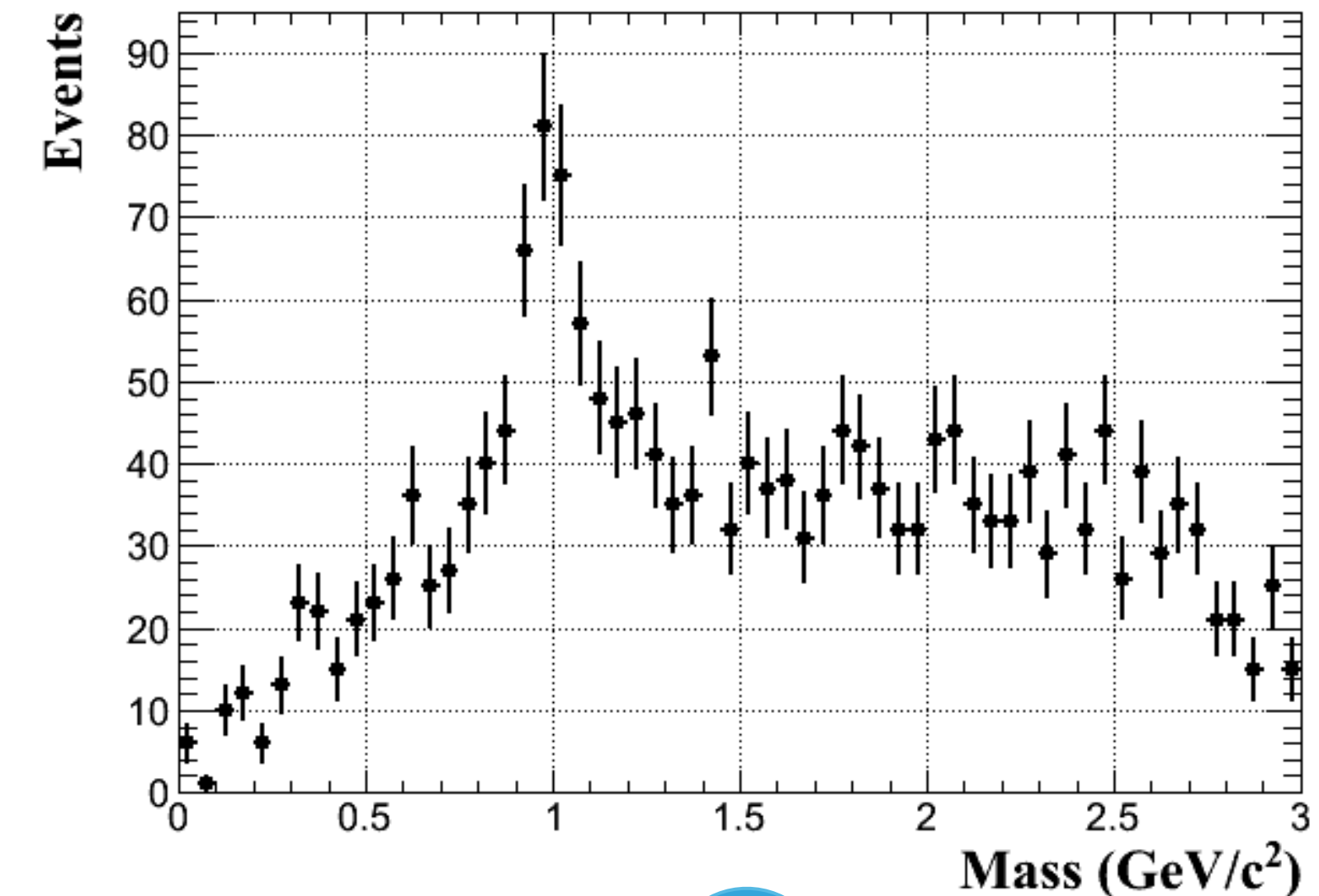
EXPERIMENT

3

Data sample

$x = (x_1, x_2, \dots, x_N)$

x is a multivariate random variable



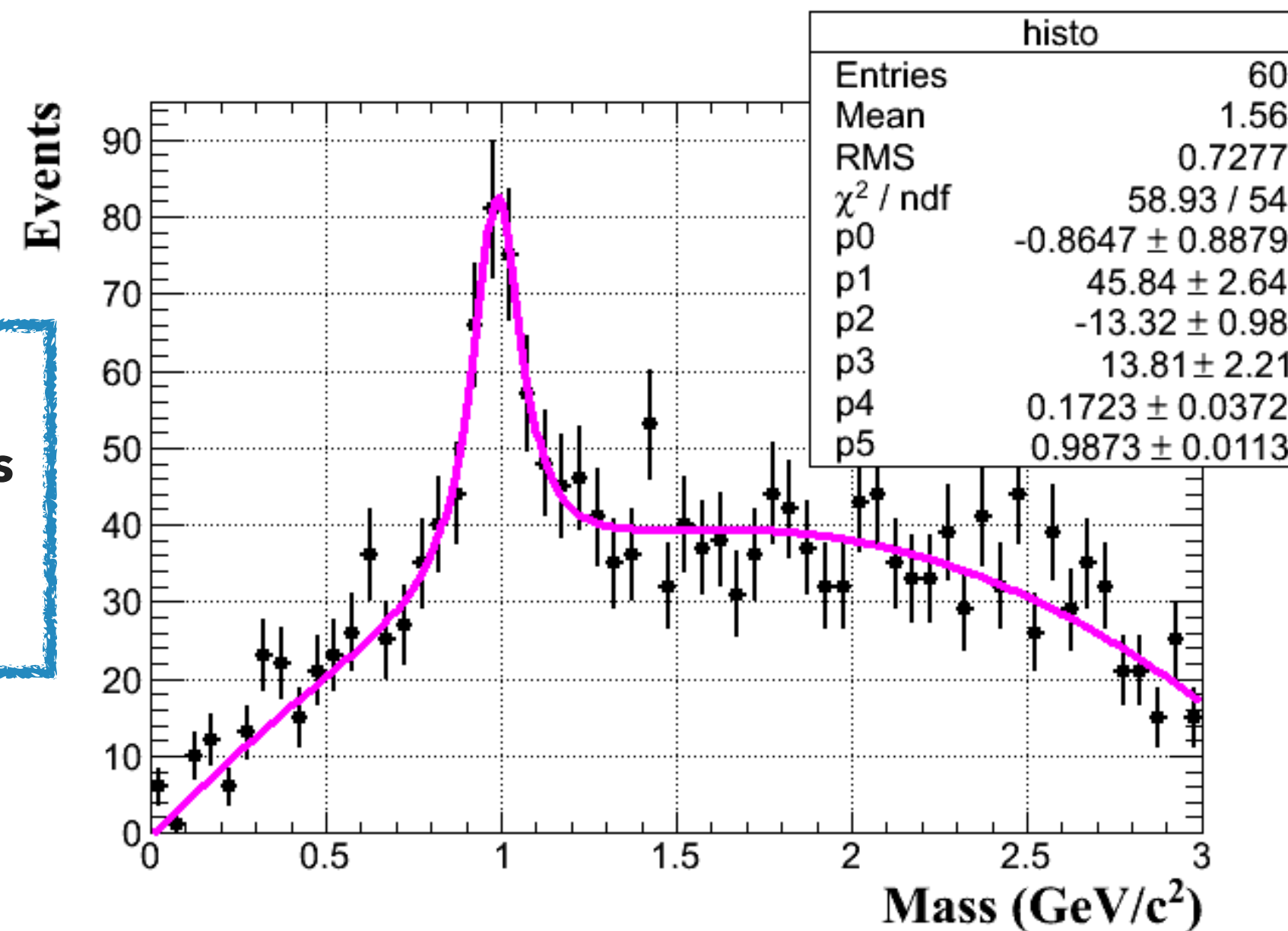
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DATA ANALYSIS

5

Results

- parameter estimates
- confidence limits
- hypothesis tests



-
- Be careful: **statistic** is not **statistics**!
 - Any new random variable (f.g. T), defined as a function of a measured sample x is called a statistic $T = T(x_1, x_2, \dots, x_N)$
 - For example, the sample mean $\bar{x} = \frac{1}{N} \sum x_i$ is a statistic!
 - A statistic used to estimate a parameter is called an **estimator**
 - For instance, the **sample mean** is a statistic and an estimator for the **population mean**, which is an unknown parameter
 - **Estimator** is a function of the data
 - **Estimate**, a value of estimator, is our “best” guess for the true value of parameter
 - Some other example of statistics (plural of statistic!): sample median, variance, standard deviation, t-statistic, chi-square statistic, kurtosis, skewness, ...

THE MAXIMUM LIKELIHOOD METHOD

- Gives consistent and asymptotically unbiased estimators
- Widely used in practice

THE LEAST SQUARES (CHI-SQUARE) METHOD

- Gives consistent estimator
- Linear Chi-Square estimator is unbiased
- Frequently used in histogram fitting

- Assume that observations (events) are independent
 - With the PDF depending on parameters $\theta: f(x_i; \theta)$
- The **probability that all N events will happen** is a product of all single events probabilities:
 - $P(x; \theta) = P(x_1; \theta)P(x_2; \theta) \cdots P(x_N; \theta) = \prod P(x_i; \theta)$
- When the variable **x is replaced by the observed** data x^{OBS} , then P is no longer a PDF
- It is usual to denote it by L and called $L(x^{\text{OBS}}; \theta)$ **the likelihood function**
 - Which is now a function of θ only $L(\theta) = P(x^{\text{OBS}}; \theta)$
- Often in the literature, it's convenient to keep X as a variable and continue to use notation $L(X; \theta)$

- The probability that all N independent events will happen is given by the likelihood function $L(x; \theta) = \prod f(x_i; \theta)$
- The principle of maximum likelihood (ML) says: **The maximum likelihood estimator $\hat{\theta}$ is the value of θ for which the likelihood is a maximum!**
- In words of R. J. Barlow: “You determine the value of θ that makes the probability of the actual results obtained, $\{x_1, \dots, x_N\}$, as large as it can possibly be.”
- In practice it's easier to maximize the **log-likelihood function**
 $\ln L(x; \theta) = \sum \ln f(x_i; \theta)$
- For p parameters we get a set of p **likelihood equations**: $\frac{\partial \ln L(x; \theta)}{\partial \theta_j} = 0$
- It is often more convenient to **minimise $-\ln L$ or $-2\ln L$**

- ML estimator is **consistent**
- ML estimate is approximately **unbiased** and **efficient** for large samples
 - Usually biased for small samples
- ML estimate is **invariant**
 - A transformation of parameter won't change the answer
 - Keep in mind that invariance comes at the cost of a bias!
- Extra care to be taken when the best value of parameters are near imposed limits
- **ML estimate is not the most likely value of parameter; it is the estimate that makes your data the most likely!**
- ML method can be used in the Bayesian approach where both θ and x are random variables
- We want to know the conditional PDF for θ given the data x :
$$p(\theta | x) = \frac{L(x | \theta)\pi(\theta)}{\int L(x | \theta')\pi(\theta')d\theta'}$$

- Likelihood function (L) is constructed by replacing the variable x by the observed data in a product of single events probabilities
- Maximising (minimising) the $\ln L$ ($-2 \ln L$) function gives the parameter estimate $\hat{\theta}_{ML}$
- $\hat{\theta}_{ML}$ does not mean that the estimate is the “most likely” value of θ , it is the value that makes your data most likely
- ML estimate is unbiased and efficient for large samples, be careful if you want to use it for small samples
- ML can be used to fit binned data
- ML can be extended to deal with the case where the number of expected events is not a fixed number but a random number

- Suppose you have a set of precisely known (without error) values $x(x_1, \dots, x_N)$ with a corresponding set of measured values $y(y_1, \dots, y_N)$ with corresponding uncertainties $\sigma(\sigma_1, \dots, \sigma_N)$
 - For example x_i histogram mass bins with y_i events with Poissonian uncertainty σ_i
- Suppose you also know a function $f(x; \theta)$ which predicts the value of y_i for any x_i . It depends on an unknown parameter θ , which you are trying to determine.
 - In our example function $f(x; \theta)$ would be theoretical prediction for number of events at a given mass
- To find best estimate of θ we minimise the suitably weighted sum of squared differences between measured and predicted values, the so called “**least squares**” or “**chi-square**”:

$$\chi^2(\theta) = \sum_{i=1}^N \frac{(y_i - f(x_i; \theta))^2}{\sigma_i^2}$$

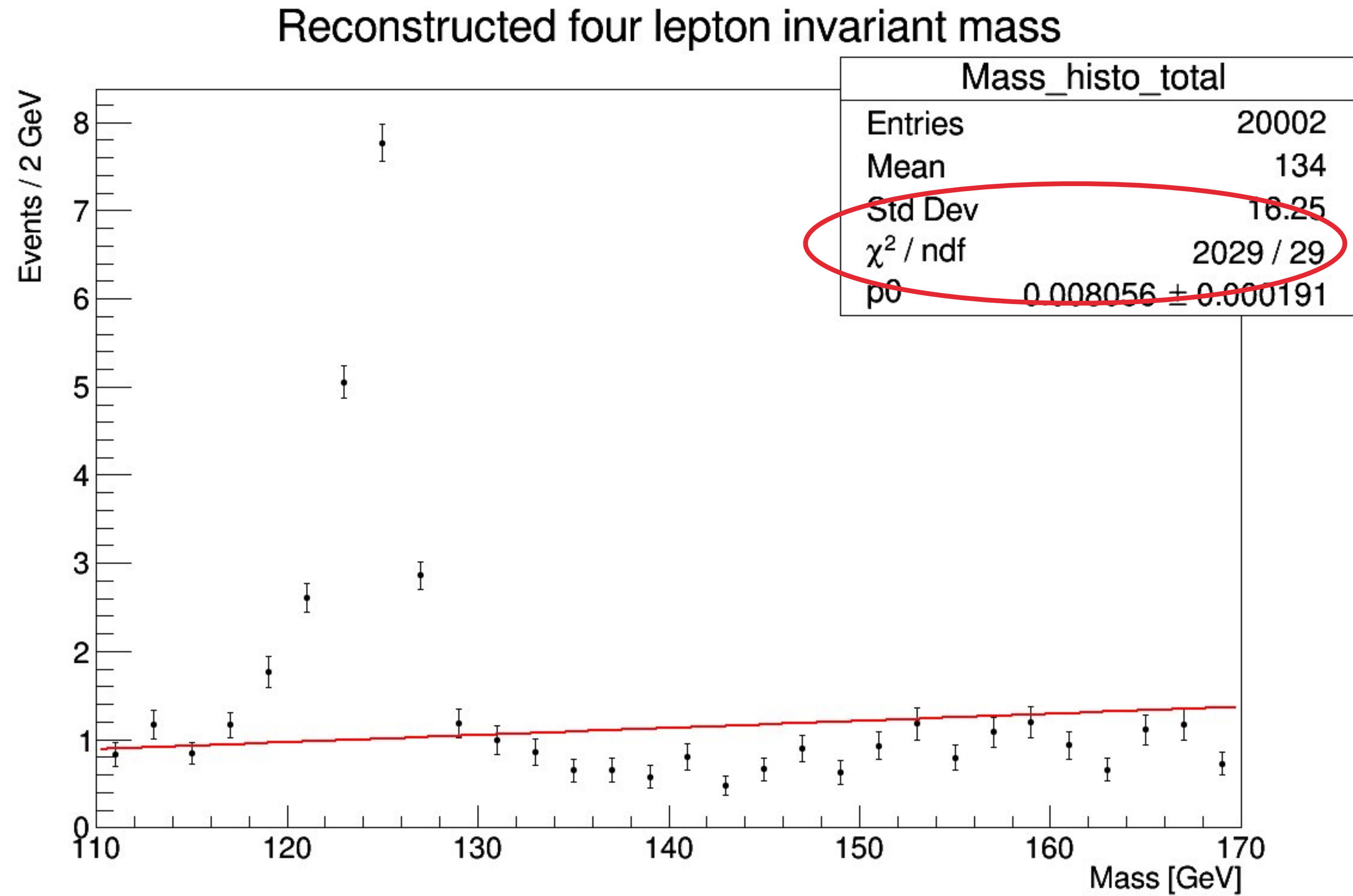
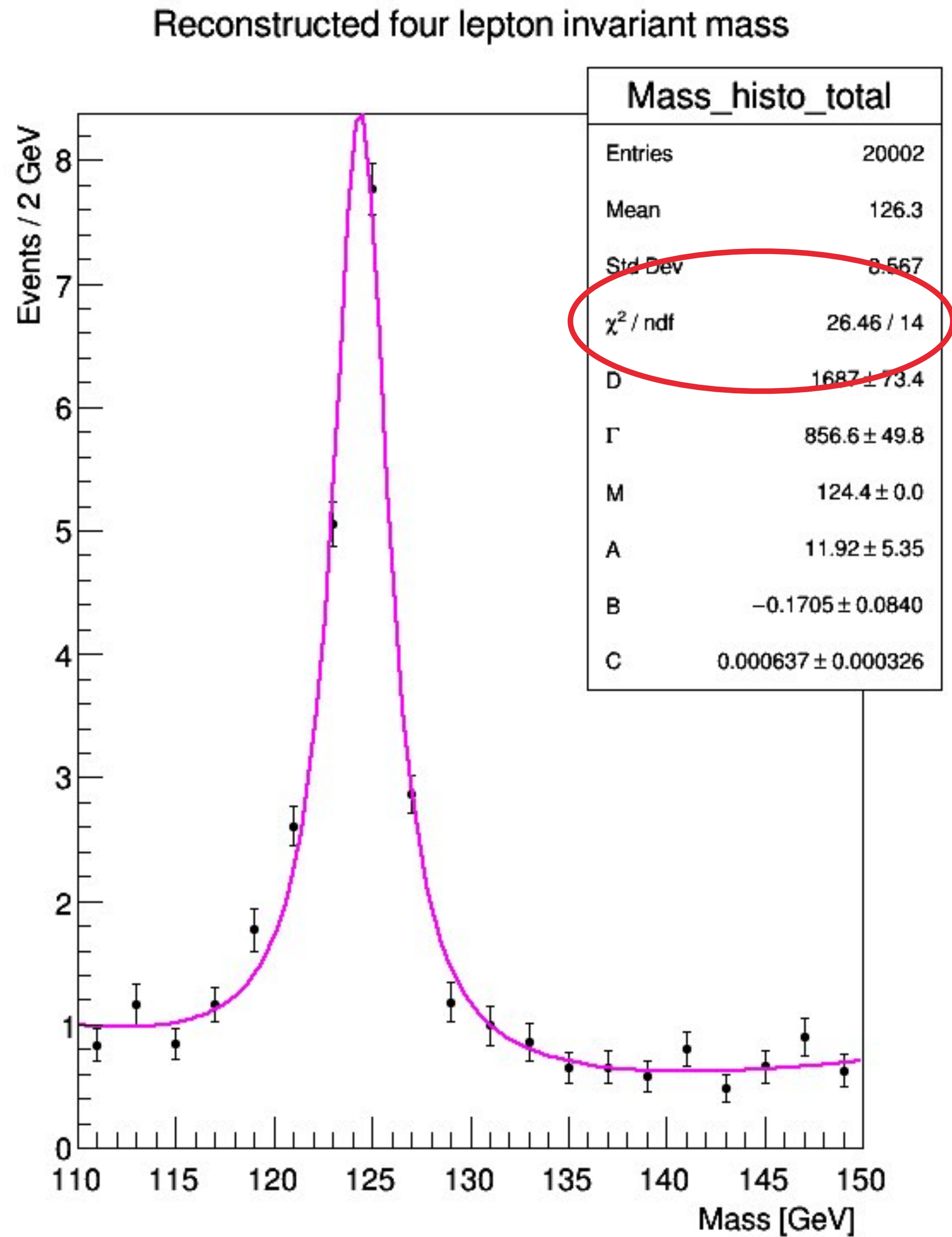
● Estimator is found by finding the value which minimises χ^2 : $\frac{\partial \chi^2}{\partial \theta} = 0$

● The quantity $\chi^2 = \sum_{i=1}^N \frac{(y_i^{data} - y_i^{ideal})^2}{(expected\ error)^2}$ gives information about the fit quality

small χ^2	large χ^2
good fit	bad fit (bad model)
overestimated errors	underestimated errors

● Since $\langle \chi^2 \rangle = N$, easy way to estimate the fit quality is to check if $\frac{\chi^2}{N.D.O.F} \approx 1$, N.D.O.F is calculated as (N - free parameters)

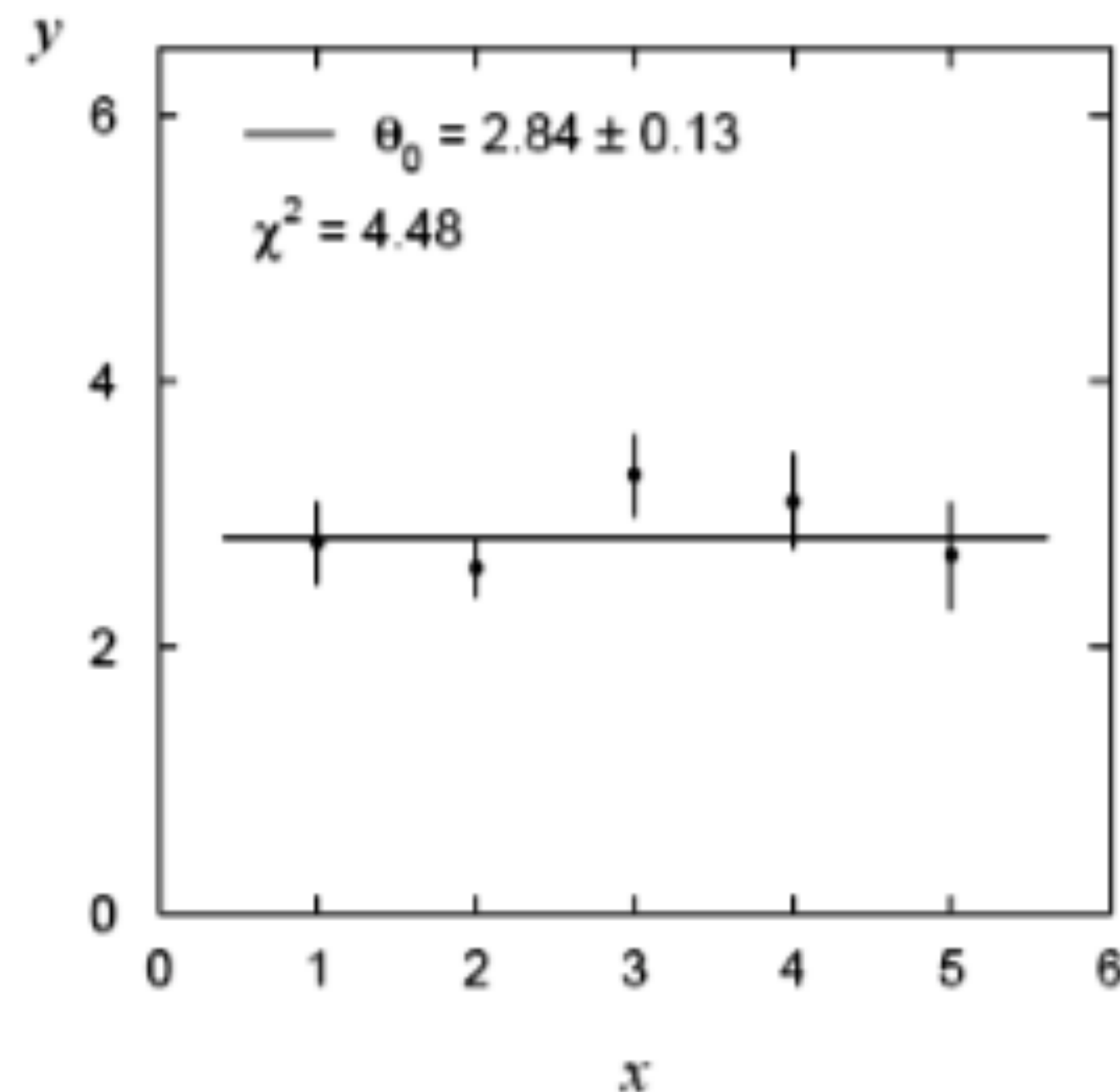
CHI-SQUARE FIT TEST - EXAMPLE



- LS has particularly desirable properties if $f(x; \theta)$ is a linear function of θ :

$$f(x; \theta) = \sum_{j=1}^m a_j(x)\theta_j, \text{ where } a_j(x) \text{ are linearly independent functions of } x$$

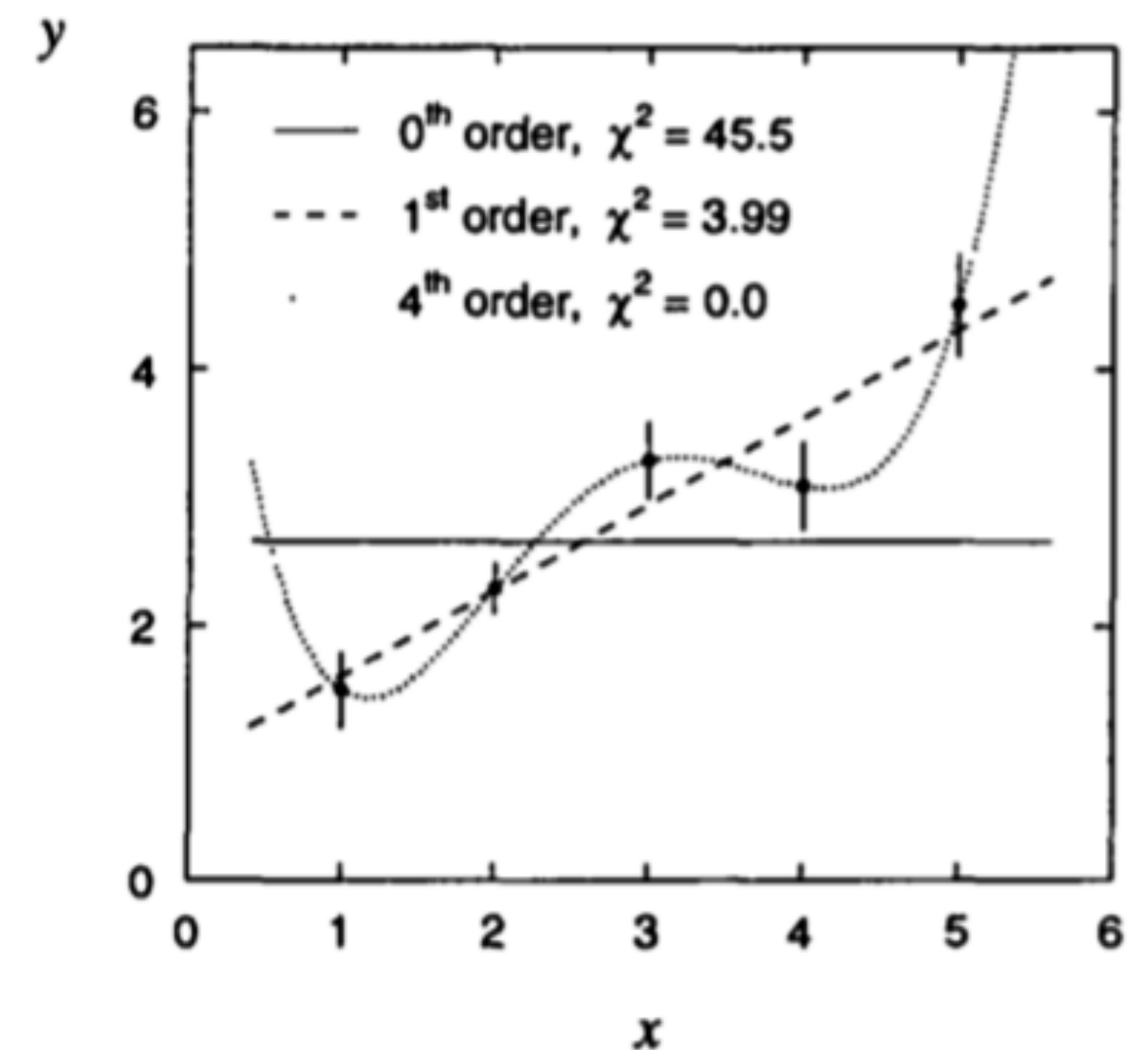
- estimators and their variances can be found analytically
- the estimators have zero bias and minimum variance



- Assume we measure 5 values of a quantity y , measured with errors σ_y at different values of x

- For the fit function we try polynomial of order m :
$$f(x; \theta) = \sum_{j=0}^m x^j \theta_j$$

- 0-th order: the weighted average
- 1-st order: a very good description
- 4-th order: equal number of parameters as points
- For Gaussian distributed y LS = ML!

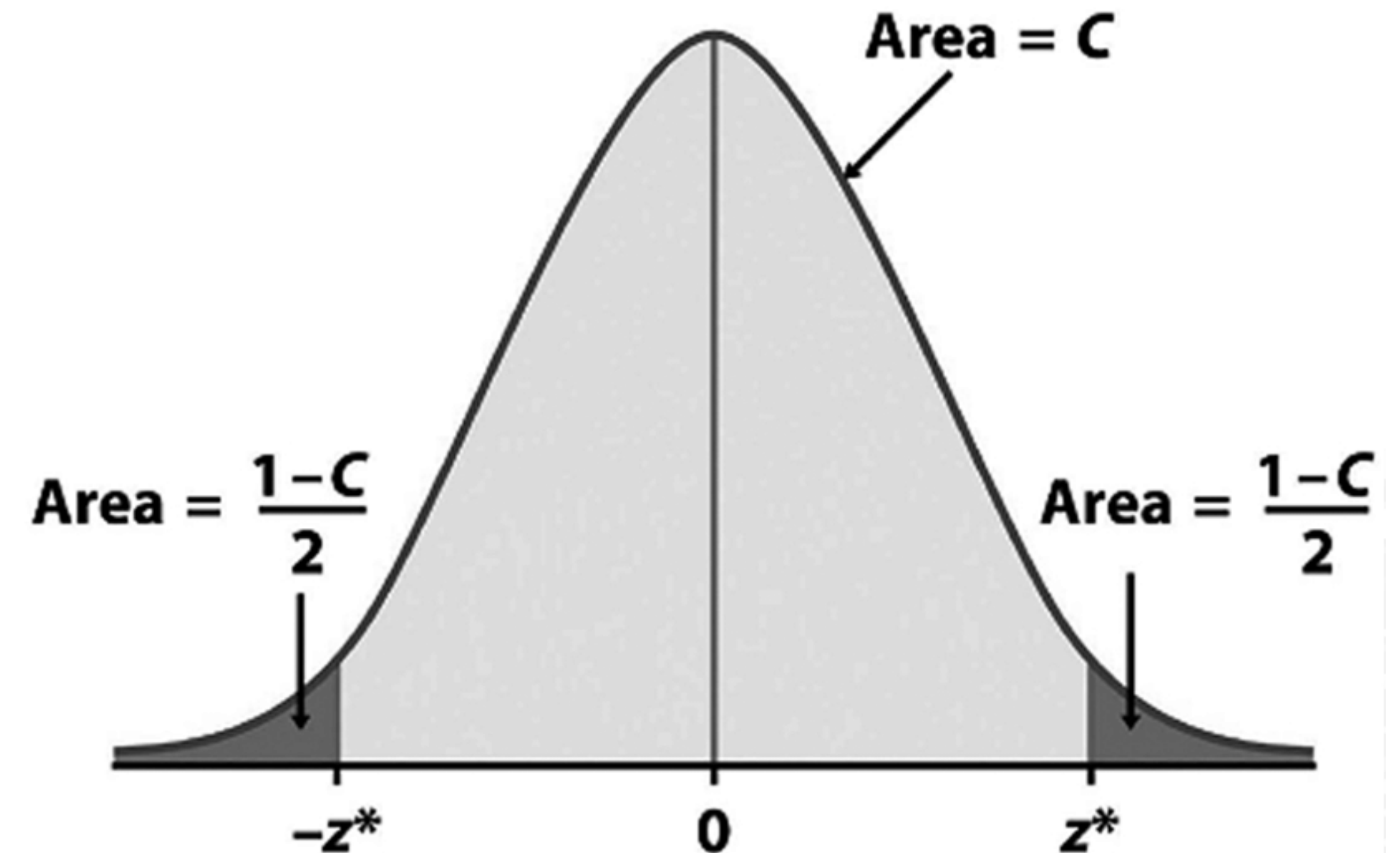


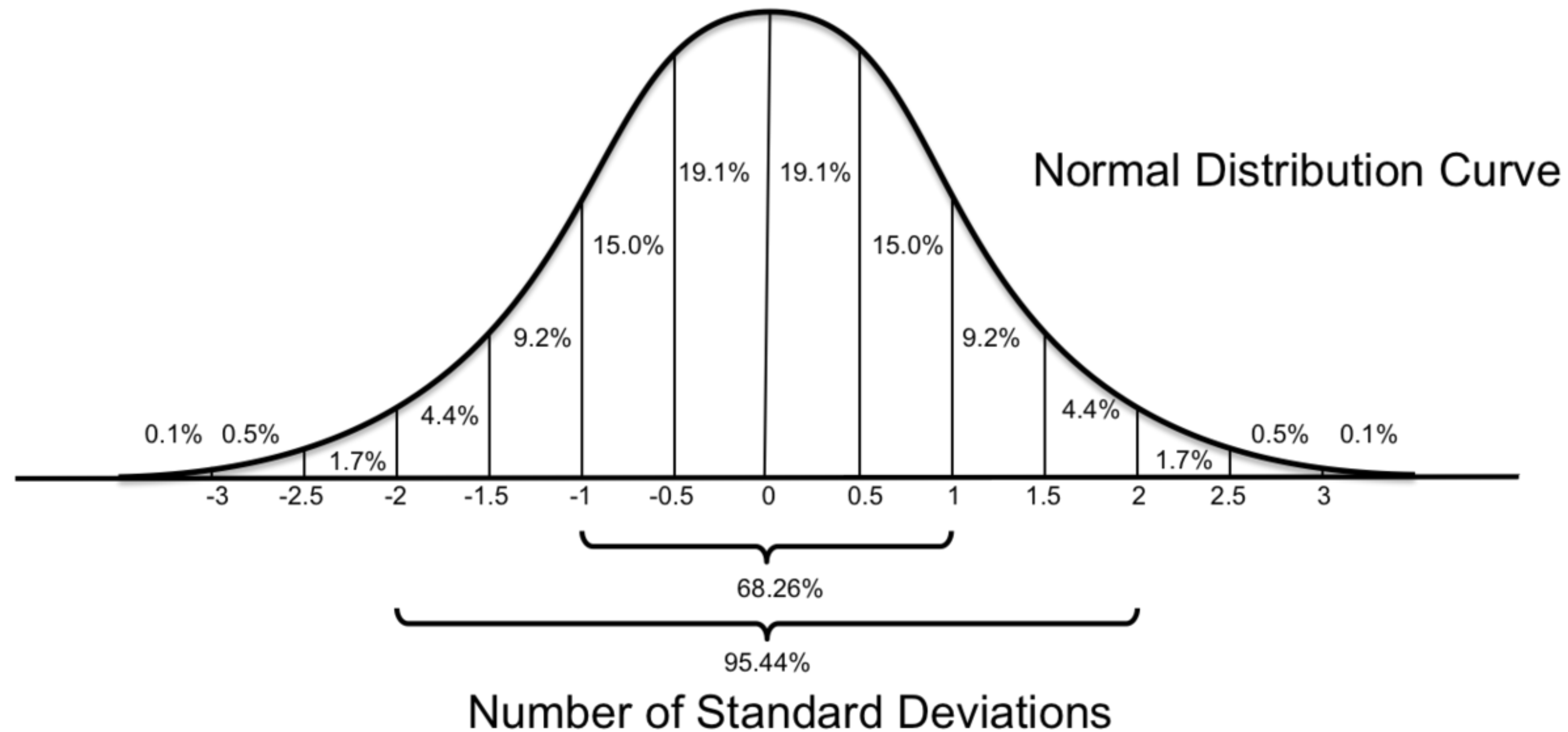
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- In addition to a “point estimate” of a parameter we should report an interval reflecting its statistical uncertainty.
 - Desirable properties of such an interval:
 - communicate objectively the result of the experiment
 - have a given probability of containing the true parameter
 - provide information needed to draw conclusions about the parameter
 - communicate incorporated prior beliefs and relevant assumptions
 - Often use \pm the estimated standard deviation (σ) of the estimator
 - In some cases, however, this is not adequate:
 - estimate near a physical boundary
 - if the PDF is not Gaussian

- Let some measured quantity be distributed according to some PDF $f(x; \theta)$, we can determine the probability that x lies within some interval, with some confidence C :

$$P(x_- < x < x_+) = \int_{x_-}^{x_+} f(x; \theta) dx = C$$

- We say that x lies in the interval $[x_-, x_+]$ with confidence C





● If $f(x; \theta)$ is a Gaussian distribution with mean μ and variance σ^2 :

● $x_{\pm} = \mu \pm 1 \cdot \sigma \quad C = 68 \%$

● $x_{\pm} = \mu \pm 2 \cdot \sigma \quad C = 95.4 \%$

● $x_{\pm} = \mu \pm 1.64 \cdot \sigma \quad C = 90 \%$

● $x_{\pm} = \mu \pm 1.96 \cdot \sigma \quad C = 95 \%$

$$P(x_- < x < x_+) = \int_{x_-}^{x_+} f(x; \theta) dx = C$$

● There are 3 conventional ways to choose an interval around the centre:

1) **Symmetric interval:** x_- and x_+ equidistant from the mean

2) **Shortest interval:** minimizes $(x_+ - x_-)$

3) **Central interval:** $\int_{-\infty}^{x_-} f(x; \theta) dx = \int_{x_+}^{+\infty} f(x; \theta) dx = \frac{1 - C}{2}$

● For the Gaussian, and any symmetric distributions, 3 definitions are equivalent

- So far we have considered only two-tailed intervals, but sometimes one-tailed limits are also useful

- for example in the case of measuring a parameter near a physical boundary

- **Upper limit:** x lies below x_+ at confidence level C :
$$\int_{-\infty}^{x_+} f(x; \theta) dx = C$$

- **Lower limit:** x lies above x_- at confidence level C :
$$\int_{x_-}^{+\infty} f(x; \theta) dx = C$$

- In a measurement two things involved:
 - True physical parameters: θ^{true}
 - Measurement of the physical parameter (parameter estimation): $\hat{\theta}$
- Given the measurement $\hat{\theta} \pm \sigma_{\theta}$ what can we say about θ^{true} ?
- Can we say that θ^{true} lies within $\hat{\theta} \pm \sigma_{\theta}$ with 68% probability?
 - **NO!!!**
 - θ^{true} is **not a random variable!** It lies in the measured interval or it does not!
- We can say that if we repeat the experiment many times with the same sample size, construct the interval according to the same prescription each time, in 68% of the experiments $\hat{\theta} \pm \sigma_{\theta}$ interval will cover θ^{true} .

- There are two ways to obtain confidence intervals for the parameter estimated by the Maximum Likelihood method

- **Analytical way:**

- If we assume the **Gaussian approximation** we can estimate the confidence interval by matrix inversion:

$$\text{cov}^{-1}(\theta_i, \theta_j) = \frac{\partial^2 \ln L}{\partial \theta_i \partial \theta_j} \bigg|_{\theta = \hat{\theta}}$$

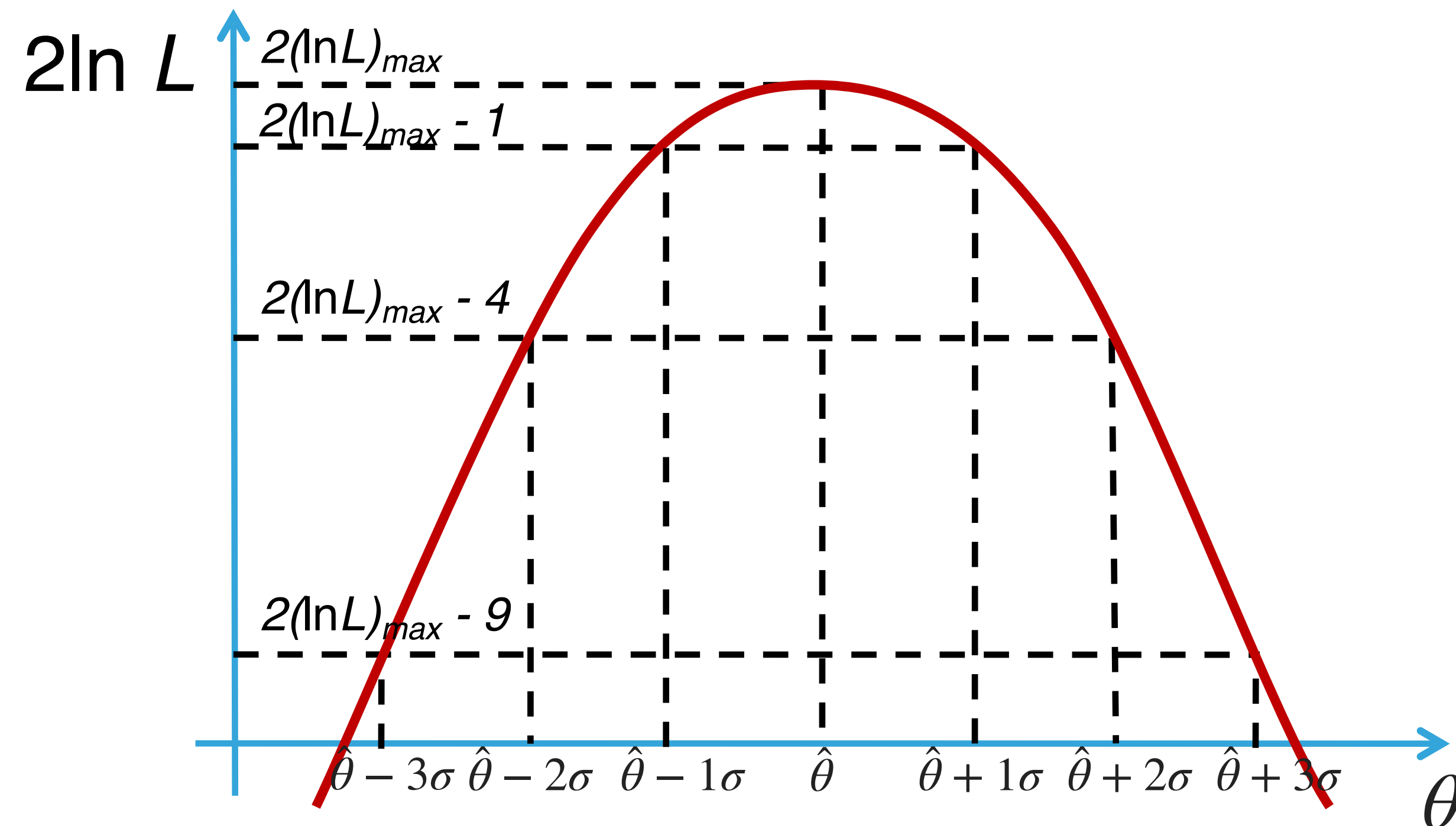
- If the likelihood function is non-Gaussian and in the limit of small number of events this approximation will give symmetrical interval while that might not be the case
- Possible to solve by hand only for very simple PDF cases, otherwise numerical solution needed
 - Matrix inversion done with HESSE/MINUIT algorithm in ROOT

- **From the Log-Likelihood curve**

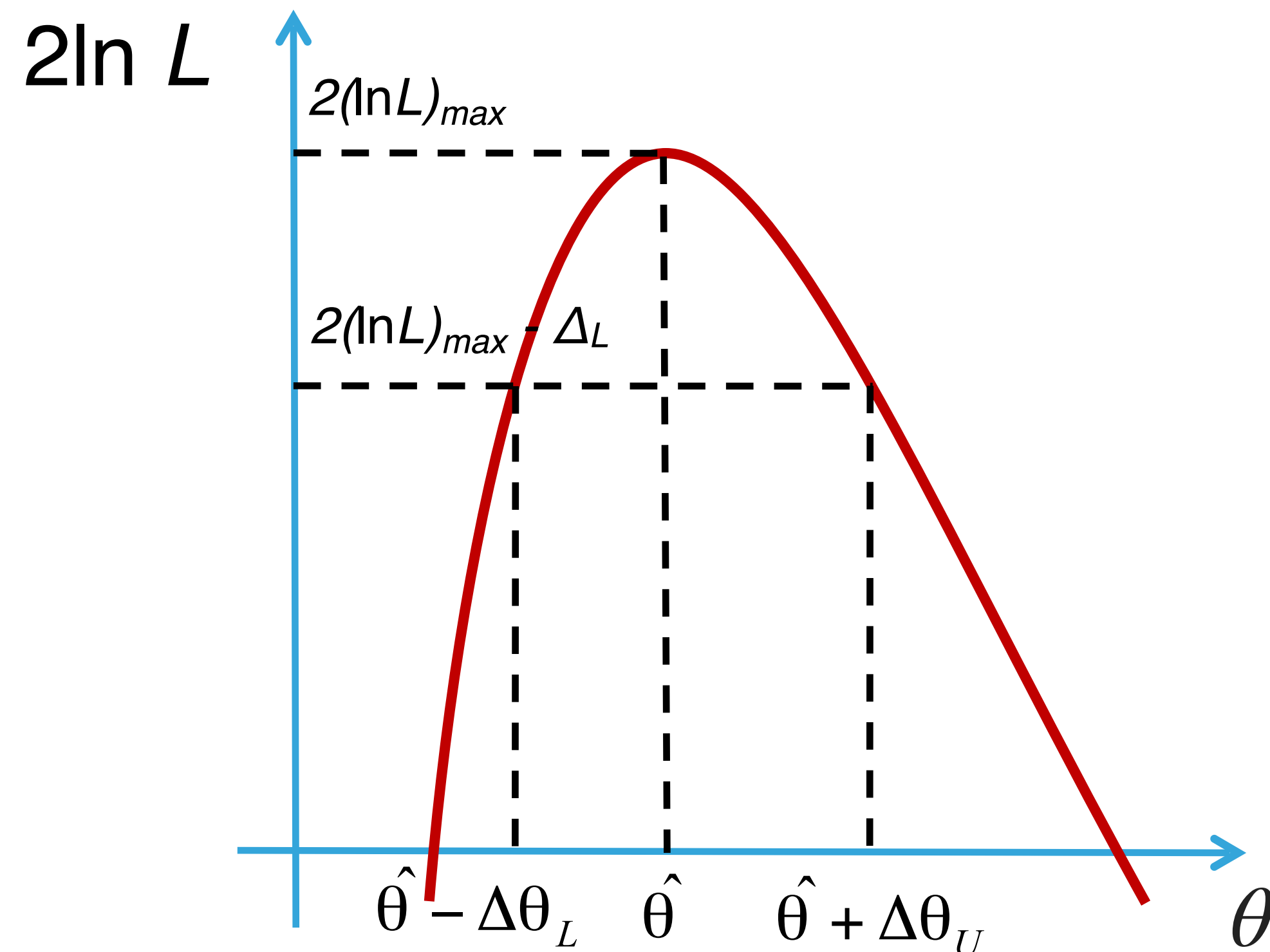
- Extract $\sigma_{\hat{\theta}}$ from log-likelihood scan using:

$$\ln L(\hat{\theta} \pm N \cdot \sigma_{\hat{\theta}}) = \ln L_{max} - \frac{N^2}{2}$$

- This is the same as looking for $2\ln L_{max} - N^2$



- The Log-Likelihood function can be asymmetric
 - for smaller samples, very non-Gaussian PDFs, non-linear problems,...
- The confidence interval is still extracted from the Log-Likelihood curve using the same prescription
 - This leads to asymmetrical confidence interval that should be used when quoting the final result

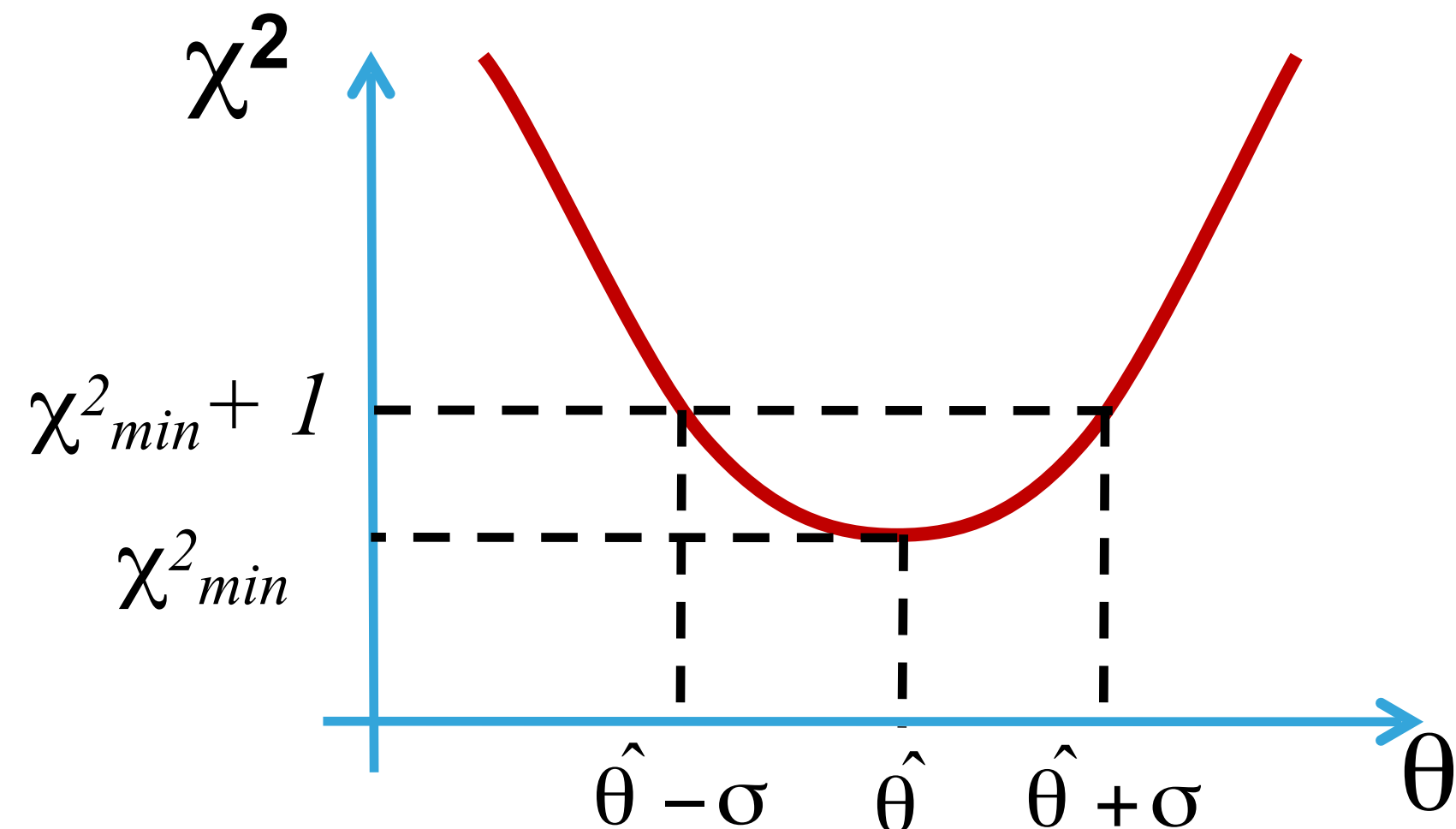


CL	Δ_L
68.27	1
95.45	4
99.73	9

- The confidence intervals for the Least Squares (Chi-Square) method are obtained in the identical way as for the Maximum likelihood method
- **Analytical way of matrix inversion:**
 - Solving analytically (or numerically):

$$cov^{-1}(\theta_i, \theta_j) = \frac{1}{2} \frac{\partial^2 \chi^2}{\partial \theta_i \partial \theta_j} \Bigg|_{\theta=\hat{\theta}}$$

● **From the Chi-Square curve**

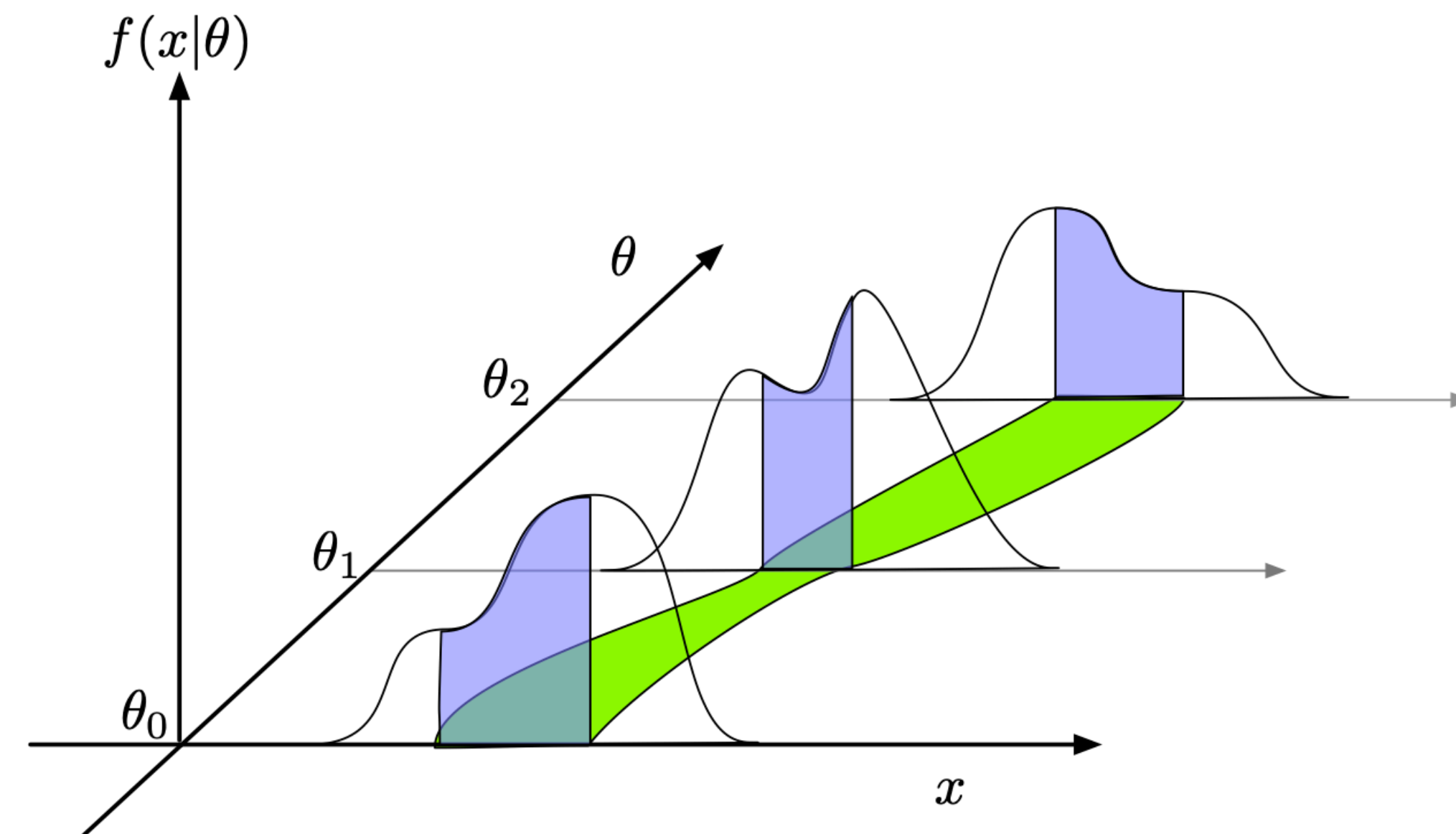


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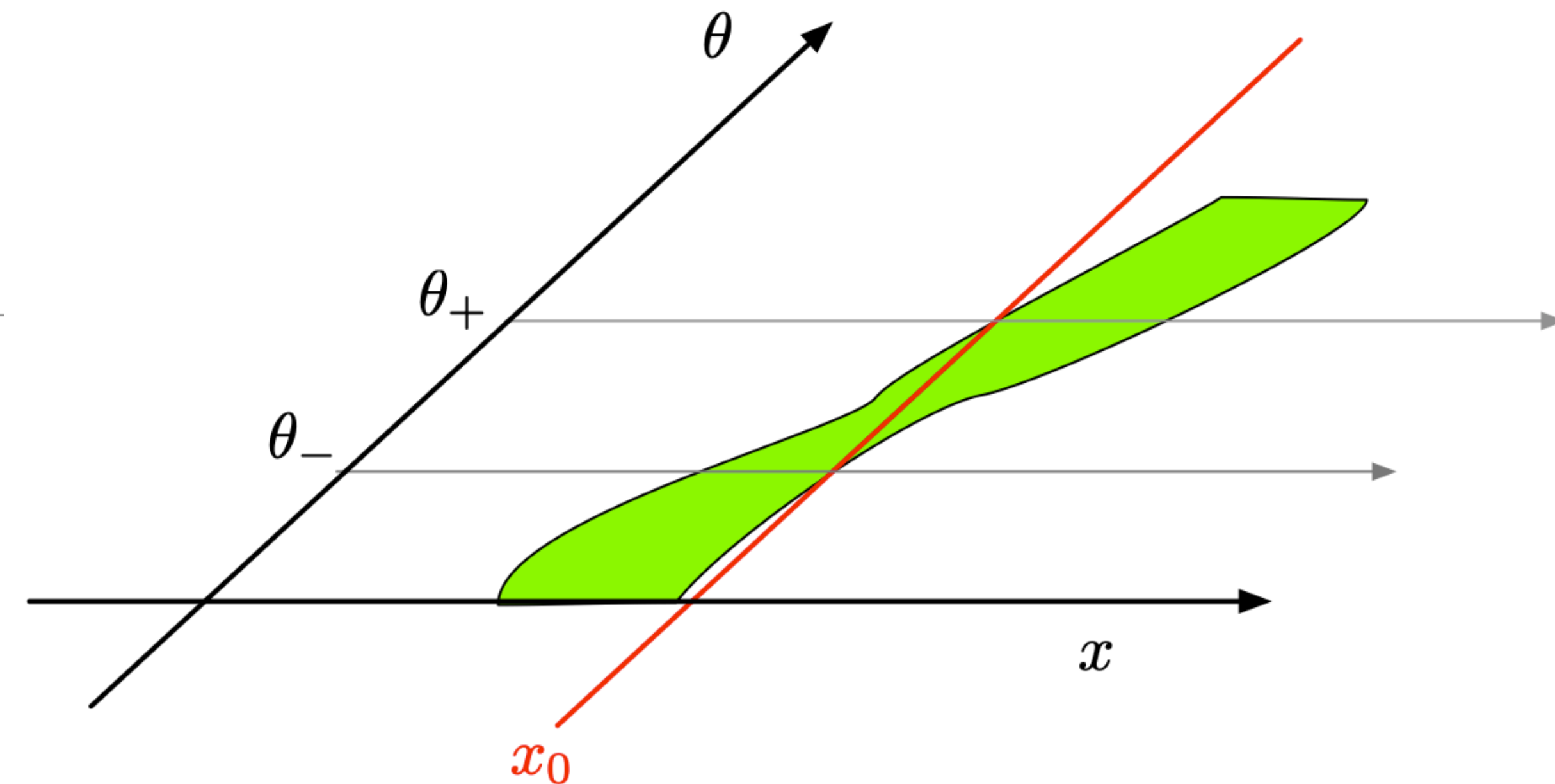
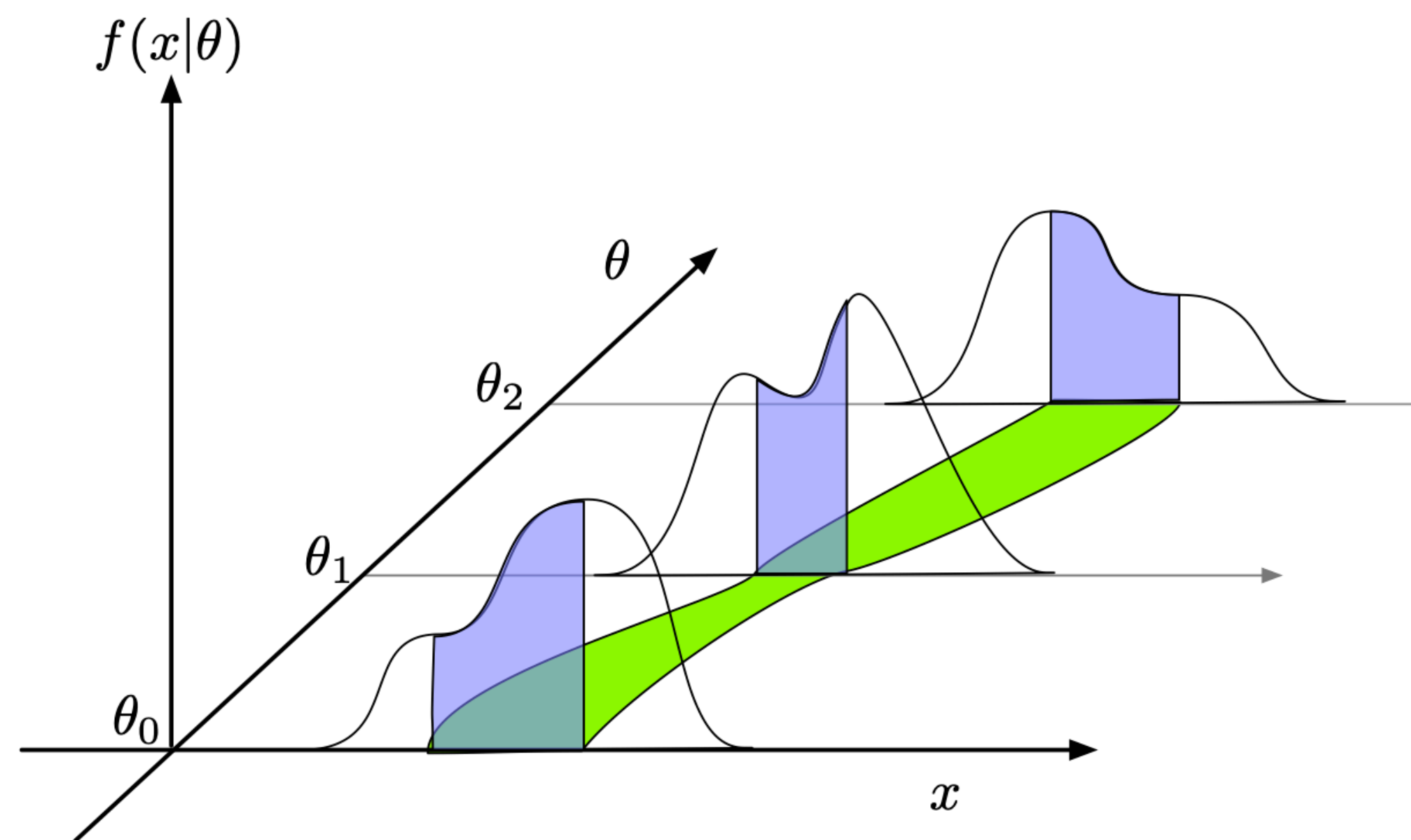
- Using frequentist approach Neyman defines confidence interval of the unknown parameter θ :

$$P(x_1 < x < x_2; \theta) = \int_{x_1}^{x_2} f(x; \theta) dx = CL$$

- x is the measurement and CL is predefined confidence level
- Union of $[x_1, x_2]$ segments for all values of the parameter θ is known as the **confidence belt**
- All of these steps are performed **before measuring the data**



- Now we perform the measurement to obtain x_0
- the points θ where the belt intersects x_0 are part of the **confidence interval** $[\theta_-, \theta_+]$ for this measurement
- For every point θ , if it were true, the data would fall in its acceptance region with probability CL, so the interval $[\theta_-, \theta_+]$ covers the true value with probability CL



- Still a frequentist approach!