

Statistical Data Analysis 2022/23

Lecture Week 9



London Postgraduate Lectures on Particle Physics
University of London MSc/MSci course PH4515



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Statistical Data Analysis

Lecture 9-1

- Least squares with histogram data

LS with histogram data

The fit function in an LS fit is not a pdf, but it could be proportional to one, e.g., when we fit the “envelope” of a histogram.

Suppose for example, we have an i.i.d. data sample of n values x_1, \dots, x_n sampled from a pdf $f(x; \theta)$. Goal is to estimate θ .

Instead of using all n values, put them in a histogram with N bins, i.e., $y_i =$ number of entries in bin i : $\mathbf{y} = (y_1, \dots, y_N)$.

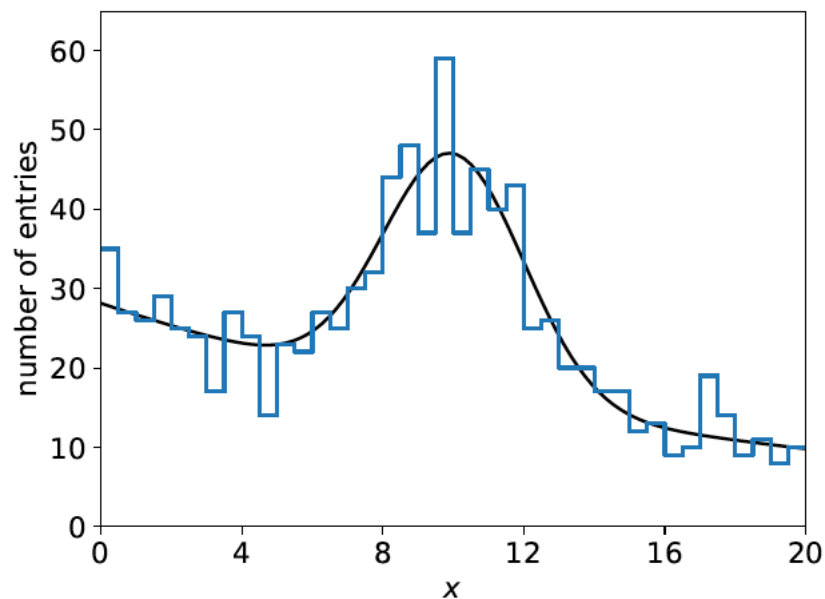
The model predicts mean values:

$$\begin{aligned} E[y_i] &= \mu_i(\theta) \\ &= n \int_{\text{bin } i} f(x; \theta) dx \\ &\approx n f(x_i; \theta) \Delta x \end{aligned}$$

bin centre



bin width



LS with histogram data (2)

The usual models:

for fixed sample size n , take $\mathbf{y} \sim$ multinomial,
if n not fixed, $y_i \sim \text{Poisson}(\mu_i)$

Suppose that the expected number of entries in each μ_i are all $\gg 1$
and probability to be in any individual bin $p_i \ll 1$, one can show

$\rightarrow y_i$ indep. and \sim Gauss with $\sigma_i \approx \sqrt{\mu_i}$. ($\rightarrow \sigma_i$ depends on $\boldsymbol{\theta}$).

The (log-) likelihood functions are then

$$L(\boldsymbol{\theta}) = \prod_{i=1}^N \frac{1}{\sqrt{2\pi}\sigma_i(\boldsymbol{\theta})} e^{-(y_i - \mu_i(\boldsymbol{\theta}))^2 / 2\sigma_i^2(\boldsymbol{\theta})}$$

$$\ln L(\boldsymbol{\theta}) = -\frac{1}{2} \sum_{i=1}^N \frac{(y_i - \mu_i(\boldsymbol{\theta}))^2}{\sigma_i(\boldsymbol{\theta})^2} - \sum_{i=1}^N \ln \sigma_i(\boldsymbol{\theta}) + C$$

LS with histogram data (3)

Still define the least-squares estimators to minimize

$$\chi^2(\boldsymbol{\theta}) = \sum_{i=1}^N \frac{(y_i - \mu_i(\boldsymbol{\theta}))^2}{\sigma_i(\boldsymbol{\theta})^2}$$

No longer equivalent to maximum likelihood (equal for $\mu_i \gg 1$).

Two possibilities for σ_i :

$$\sigma_i = \sqrt{\mu_i(\boldsymbol{\theta})} \quad (\text{LS method})$$

$$\sigma_i = \sqrt{y_i} \quad (\text{Modified LS method})$$

Modified LS can be easier computationally but not defined if any $y_i = 0$.

For either method, $\chi^2_{\min} \sim$ chi-square pdf for $\mu_i \gg 1$, but this breaks down for when the μ_i are not large.

LS with histogram data — normalization

Do **not** “fit” the normalization, i.e., $n \rightarrow$ free parameter ν :

$$\mu_i(\boldsymbol{\theta}, \nu) = \nu \int_{\text{bin } i} f(x; \boldsymbol{\theta}) dx$$

If you do this, one finds the LS estimator for ν is not n , but rather

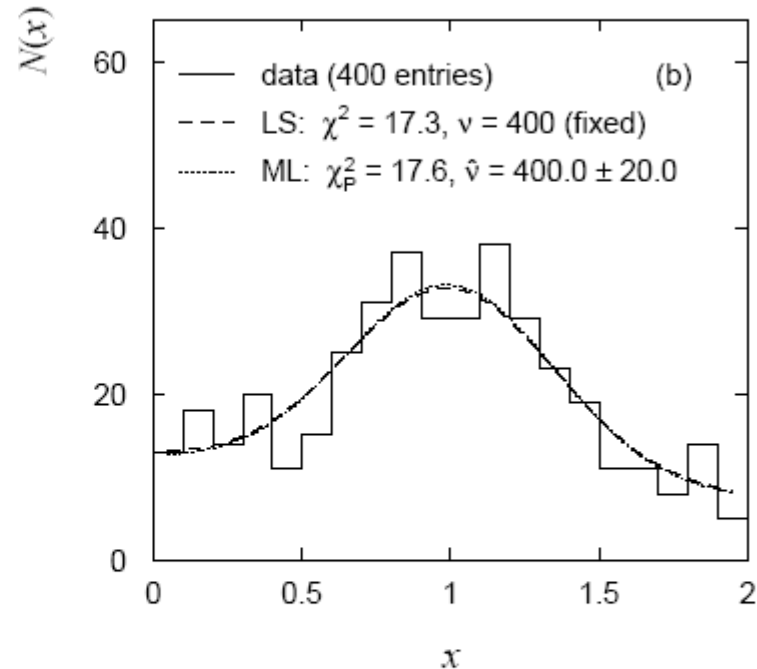
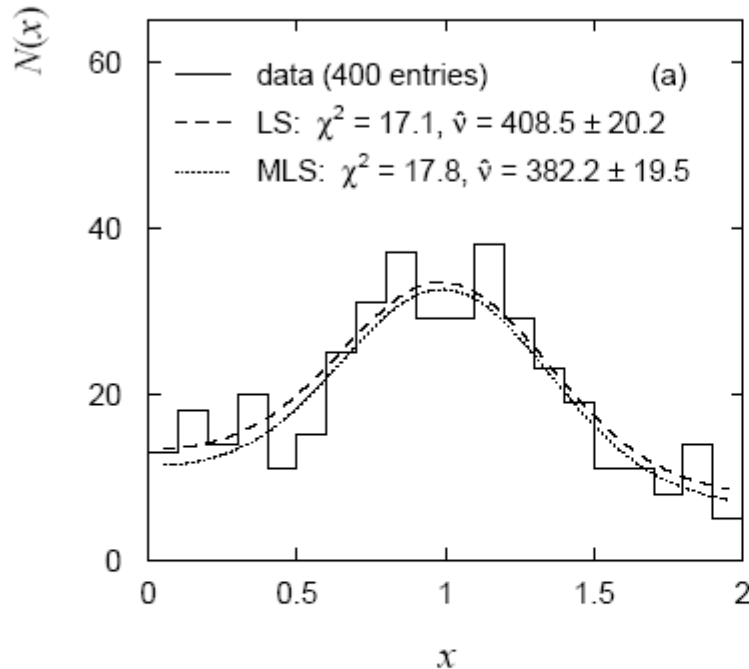
$$\hat{\nu}_{\text{LS}} = n + \frac{\chi_{\text{min}}^2}{2}$$

$$\hat{\nu}_{\text{MLS}} = n - \chi_{\text{min}}^2$$

Software may include adjustable normalization parameter as default; better to use known n .

LS normalization example

Example with $n = 400$ entries, $N = 20$ bins:



Expect χ_{\min}^2 around $N - m$,

→ relative error in \hat{v} large when N large, n small

Either get n directly from data for LS (or better, use ML).

Statistical Data Analysis

Lecture 9-2

- Goodness-of-fit from the likelihood ratio
- Wilks' theorem
- MLE and goodness-of-fit all in one

Goodness of fit from the likelihood ratio


Suppose we model data using a likelihood $L(\boldsymbol{\mu})$ that depends on N parameters $\boldsymbol{\mu} = (\mu_1, \dots, \mu_N)$. Define the statistic

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

where $\hat{\boldsymbol{\mu}}$ is the ML estimator for $\boldsymbol{\mu}$. Value of $t_{\boldsymbol{\mu}}$ reflects agreement between hypothesized $\boldsymbol{\mu}$ and the data.

Good agreement means $\boldsymbol{\mu} \approx \hat{\boldsymbol{\mu}}$, so $t_{\boldsymbol{\mu}}$ is small;

Larger $t_{\boldsymbol{\mu}}$ means less compatibility between data and $\boldsymbol{\mu}$.

Quantify “goodness of fit” with p -value: $p_{\boldsymbol{\mu}} = \int_{t_{\boldsymbol{\mu}, \text{obs}}}^{\infty} f(t_{\boldsymbol{\mu}} | \boldsymbol{\mu}) dt_{\boldsymbol{\mu}}$
need this pdf 

Likelihood ratio (2)

Now suppose the parameters $\boldsymbol{\mu} = (\mu_1, \dots, \mu_N)$ can be determined by another set of parameters $\boldsymbol{\theta} = (\theta_1, \dots, \theta_M)$, with $M < N$.

E.g., curve fit with $\mu_i = E[y_i] = \mu(x_i; \boldsymbol{\theta})$, $i = 1, \dots, N$, $\boldsymbol{\theta} = (\theta_1, \dots, \theta_M)$.

Want to test hypothesis that the true model is somewhere in the subspace $\boldsymbol{\mu} = \boldsymbol{\mu}(\boldsymbol{\theta})$ versus the alternative of the full parameter space $\boldsymbol{\mu}$. Generalize the LR test statistic to be

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

fit M parameters

fit N parameters

To get p -value, need pdf $f(t_{\boldsymbol{\mu}} | \boldsymbol{\mu}(\boldsymbol{\theta}))$.

Wilks' Theorem

Wilks' Theorem: if the hypothesized $\mu_i(\boldsymbol{\theta})$, $i = 1, \dots, N$, are true for some choice of the parameters $\boldsymbol{\theta} = (\theta_1, \dots, \theta_M)$, then in the large sample limit (and provided regularity conditions are satisfied)

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

MLE of $(\theta_1, \dots, \theta_M)$

follows a chi-square distribution for $N - M$ degrees of freedom.

MLE of (μ_1, \dots, μ_N)

The regularity conditions include: the model in the numerator of the likelihood ratio is “nested” within the one in the denominator, i.e., $\boldsymbol{\mu}(\boldsymbol{\theta})$ is a special case of $\boldsymbol{\mu} = (\mu_1, \dots, \mu_N)$.

Proof boils down to having all estimators \sim Gaussian.

S.S. Wilks, *The large-sample distribution of the likelihood ratio for testing composite hypotheses*, Ann. Math. Statist. **9** (1938) 60-2.

Goodness of fit with Gaussian data

Suppose the data are N independent Gaussian distributed values:

$$y_i \sim \text{Gauss}(\mu_i, \sigma_i), \quad i = 1, \dots, N$$

want to estimate

known

N measurements and N parameters (= “saturated model”)

Likelihood:
$$L(\boldsymbol{\mu}) = \prod_{i=1}^N \frac{1}{\sqrt{2\pi}\sigma_i} e^{-(y_i - \mu_i)^2 / 2\sigma_i^2}$$

Log-likelihood:
$$\ln L(\boldsymbol{\mu}) = -\frac{1}{2} \sum_{i=1}^N \frac{(y_i - \mu_i)^2}{\sigma_i^2} + C$$

ML estimators:
$$\hat{\mu}_i = y_i \quad i = 1, \dots, N$$


Likelihood ratio for Gaussian data

Now suppose $\boldsymbol{\mu} = \boldsymbol{\mu}(\boldsymbol{\theta})$, e.g., in an LS fit with $\mu_i(\boldsymbol{\theta}) = \mu(x_i; \boldsymbol{\theta})$.

The goodness-of-fit statistic for the test of the hypothesis $\boldsymbol{\mu}(\boldsymbol{\theta})$ becomes

$$t_{\boldsymbol{\mu}} = -2 \ln \frac{L(\boldsymbol{\mu}(\hat{\boldsymbol{\theta}}))}{L(\hat{\boldsymbol{\mu}})} = \sum_{i=1}^N \frac{(y_i - \mu_i(\hat{\boldsymbol{\theta}}))^2}{\sigma_i^2} \sim \chi_{N-M}^2$$

chi-square pdf for $N-M$
degrees of freedom



Here $t_{\boldsymbol{\mu}}$ is the same as χ_{\min}^2 from an LS fit.

So Wilks' theorem formally states the property that we claimed for the minimized chi-squared from an LS fit with N measurements and M fitted parameters.

Likelihood ratio for Poisson data

Suppose the data are a set of values $\mathbf{n} = (n_1, \dots, n_N)$, e.g., the numbers of events in a histogram with N bins.

Assume $n_i \sim \text{Poisson}(\nu_i)$, $i = 1, \dots, N$, all independent.

First (for LR denominator) treat $\boldsymbol{\nu} = (\nu_1, \dots, \nu_N)$ as all adjustable:

Likelihood:
$$L(\boldsymbol{\nu}) = \prod_{i=1}^N \frac{\nu_i^{n_i}}{n_i!} e^{-\nu_i}$$

Log-likelihood:
$$\ln L(\boldsymbol{\nu}) = \sum_{i=1}^N [n_i \ln \nu_i - \nu_i] + C$$

ML estimators:
$$\hat{\nu}_i = n_i, \quad i = 1, \dots, N$$

Goodness of fit with Poisson data (2)

For LR numerator find $\nu(\theta)$ with M fitted parameters $\theta = (\theta_1, \dots, \theta_M)$:

$$t_\nu = -2 \ln \frac{L(\nu(\hat{\theta}))}{L(\hat{\nu})} = -2 \sum_{i=1}^N \left[n_i \ln \frac{\nu_i(\hat{\theta})}{n_i} - \nu_i(\hat{\theta}) + n_i \right]$$

if $n_i = 0$, skip \ln term

Wilks' theorem: in large-sample limit $t_\nu \sim \chi_{N-M}^2$

Exact in large sample limit; in practice good approximation for surprisingly small n_i (\sim several).

As before use t_ν to get p -value of $\nu(\theta)$,

$$p_\nu = \int_{t_{\nu, \text{obs}}}^{\infty} f(t_\nu | \nu(\theta)) dt_\nu = 1 - F_{\chi^2}(t_{\nu, \text{obs}}; N - M)$$

independent of θ

Goodness of fit with multinomial data

Similar if data $\mathbf{n} = (n_1, \dots, n_N)$ follow multinomial distribution:

$$P(\mathbf{n}|\mathbf{p}, n_{\text{tot}}) = \frac{n_{\text{tot}}!}{n_1!n_2!\dots n_N!} p_1^{n_1} p_2^{n_2} \dots p_N^{n_N}$$

E.g. histogram with N bins but fix: $n_{\text{tot}} = \sum_{i=1}^N n_i$

Log-likelihood: $\ln L(\boldsymbol{\nu}) = \sum_{i=1}^N n_i \ln \frac{\nu_i}{n_{\text{tot}}} + C$ ($\nu_i = p_i n_{\text{tot}}$)

ML estimators: $\hat{\nu}_i = n_i$ (Only $N-1$ independent; one is n_{tot} minus sum of rest.)

Goodness of fit with multinomial data (2)

The likelihood ratio statistics become:

$$t_{\nu} = -2 \ln \frac{L(\nu(\hat{\theta}))}{L(\hat{\nu})} = -2 \sum_{i=1}^N n_i \ln \frac{\nu_i(\hat{\theta})}{n_i}$$

if $n_i = 0$, skip term

Wilks: in large sample limit $t_{\nu} \sim \chi_{N-M-1}^2$

One less degree of freedom than in Poisson case because effectively only $N-1$ parameters fitted in denominator of LR.

Estimators and g.o.f. all at once

Evaluate numerators with θ (not its estimator); if any $n_i = 0$, omit the corresponding log terms:

$$\chi_{\text{P}}^2(\theta) = -2 \sum_{i=1}^N \left[n_i \ln \frac{\nu_i(\theta)}{n_i} - \nu_i(\theta) + n_i \right] \quad (\text{Poisson})$$

$$\chi_{\text{M}}^2(\theta) = -2 \sum_{i=1}^N n_i \ln \frac{\nu_i(\theta)}{n_i} \quad (\text{Multinomial})$$

These are equal to the corresponding $-2 \ln L(\theta)$ plus terms not depending on θ , so minimizing them gives the usual ML estimators for θ .

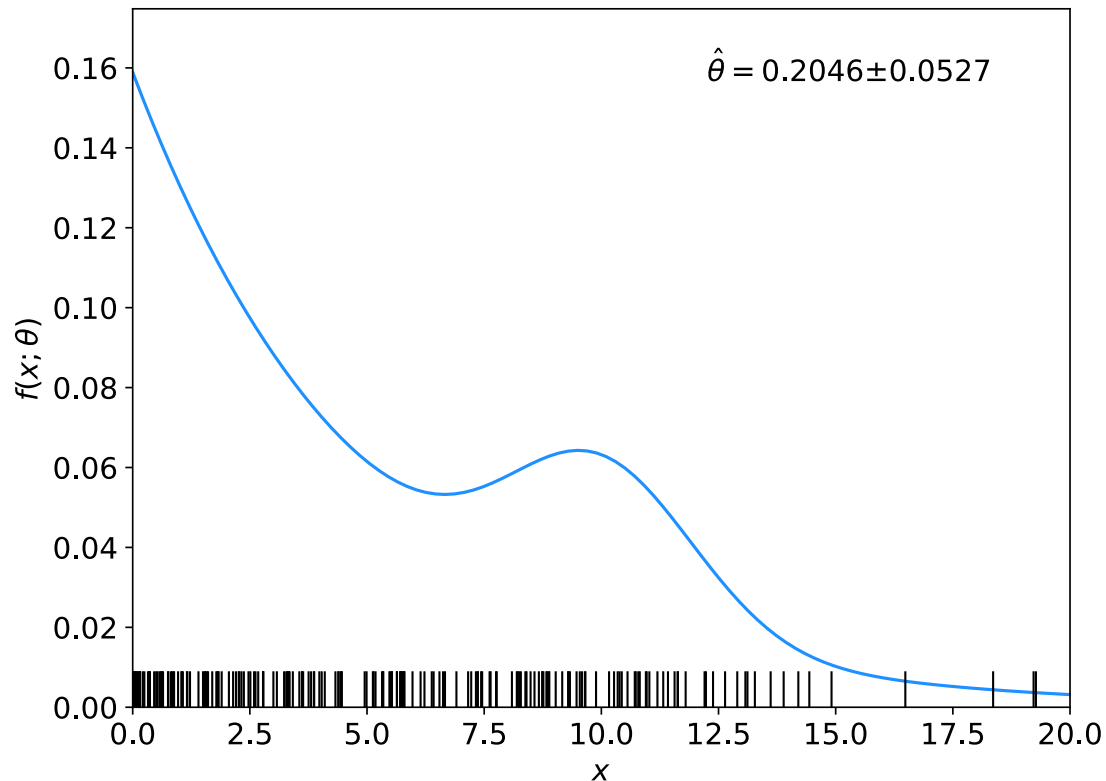
The minimized value gives the statistic t_{ν} , so we get goodness-of-fit for free.

Steve Baker and Robert D. Cousins, *Clarification of the use of the chi-square and likelihood functions in fits to histograms*, NIM **221** (1984) 437.

Examples of ML/LS fits

Unbinned maximum likelihood (mlFit.py, minimize negLogL)

$$\ln L(\boldsymbol{\theta}) = \sum_{i=1}^n \ln f(x_i; \boldsymbol{\theta})$$

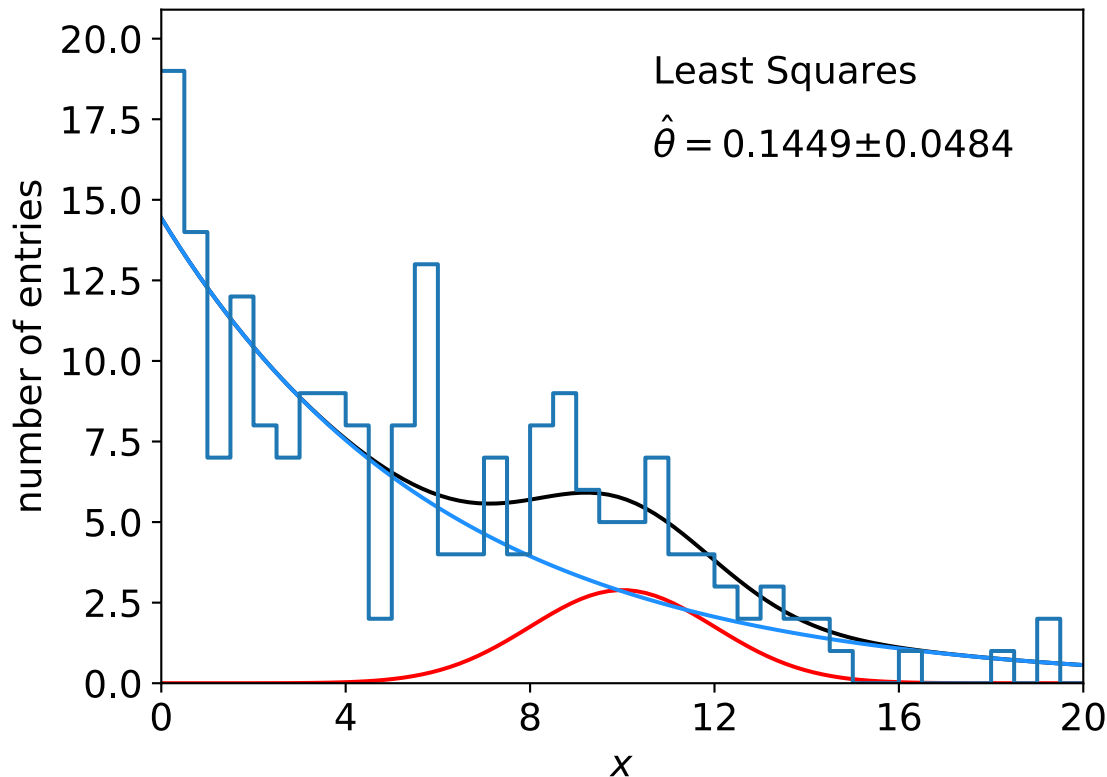


No useful measure
of goodness-of-fit
from unbinned ML.

Examples of ML/LS fits

Least Squares fit (histFit.py, minimize chi2LS)

$$\chi^2(\boldsymbol{\theta}) = \sum_{i=1}^N \frac{(y_i - \mu_i(\boldsymbol{\theta}))^2}{\mu_i(\boldsymbol{\theta})}$$



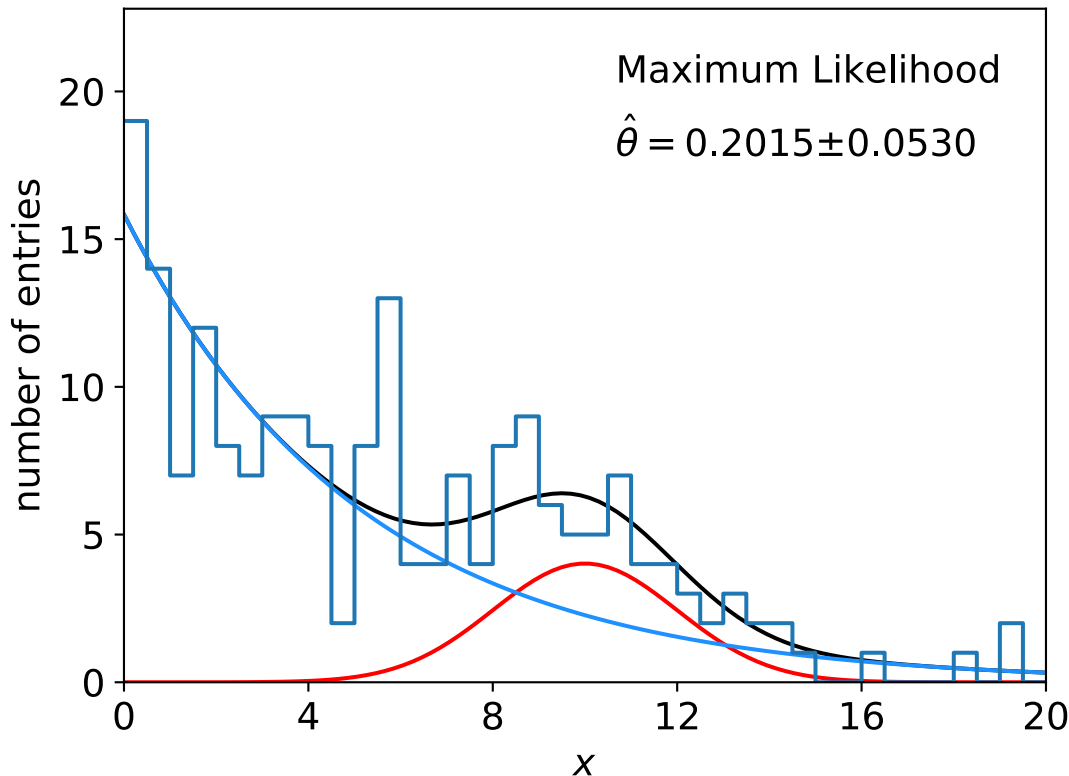
$$\chi^2_{\min} = 32.7$$
$$n_{\text{dof}} = 38$$
$$p = 0.71$$

Many bins with few entries, LS not expected to be reliable.

Examples of ML/LS fits

Multinomial maximum likelihood fit (histFit.py, minimize chi2M)

$$\chi_M^2(\boldsymbol{\theta}) = -2 \sum_{i=1}^N n_i \ln \frac{\nu_i(\boldsymbol{\theta})}{n_i}$$



$$\chi_{\min}^2 = 35.3$$
$$n_{\text{dof}} = 37$$
$$p = 0.55$$

Essentially same result
as unbinned ML.

Statistical Data Analysis

Lecture 9-3

- Interval estimation
- Confidence interval from inverting a test
- Example: limits on mean of Gaussian

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 θ can be found by defining a test of the hypothesized value θ (do this for all θ):

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

If data observed in the critical region, reject the value θ .

Now invert the test to define a confidence interval as:

set of θ values that are not rejected in a test of size α (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 θ , resulting in a p -value, p_θ .

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

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

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

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

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 θ 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” θ is

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

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

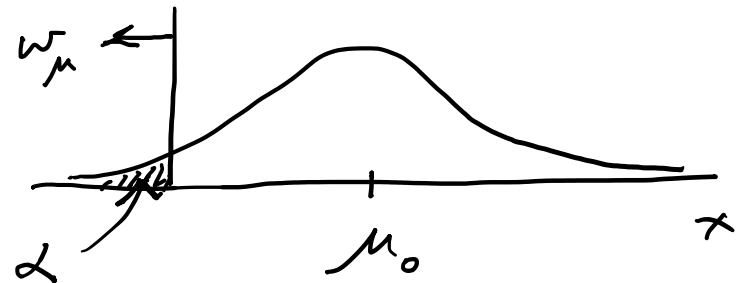
Example: upper limit on mean of Gaussian

When we test the parameter, we should take the critical region to maximize the power with respect to the relevant alternative(s).

Example: $x \sim \text{Gauss}(\mu, \sigma)$ (take σ known)

Test $H_0 : \mu = \mu_0$ versus the alternative $H_1 : \mu < \mu_0$

→ Put w_μ at region of x -space characteristic of low μ (i.e. at low x)

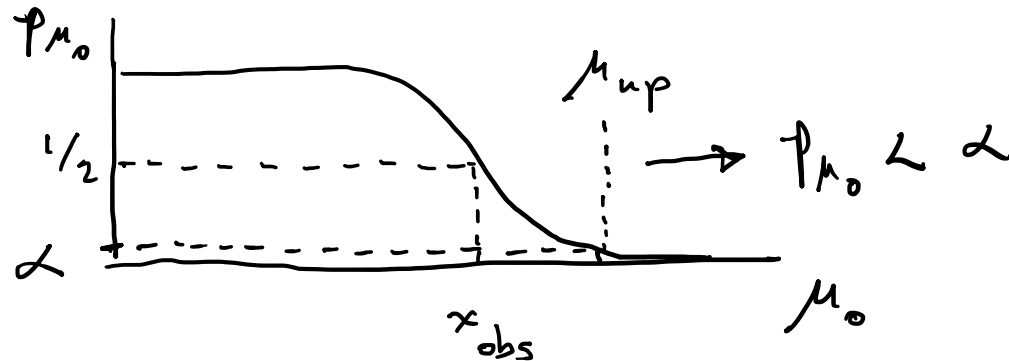


Equivalently, take the p -value to be

$$p_{\mu_0} = P(x \leq x_{\text{obs}} | \mu_0) = \int_{-\infty}^{x_{\text{obs}}} \frac{1}{\sqrt{2\pi}\sigma} e^{-(x-\mu_0)^2/2\sigma^2} dx = \Phi\left(\frac{x_{\text{obs}} - \mu_0}{\sigma}\right)$$

Upper limit on Gaussian mean (2)

To find confidence interval, repeat for all μ_0 , i.e., set $p_{\mu_0} = \alpha$ and solve for μ_0 to find the interval's boundary



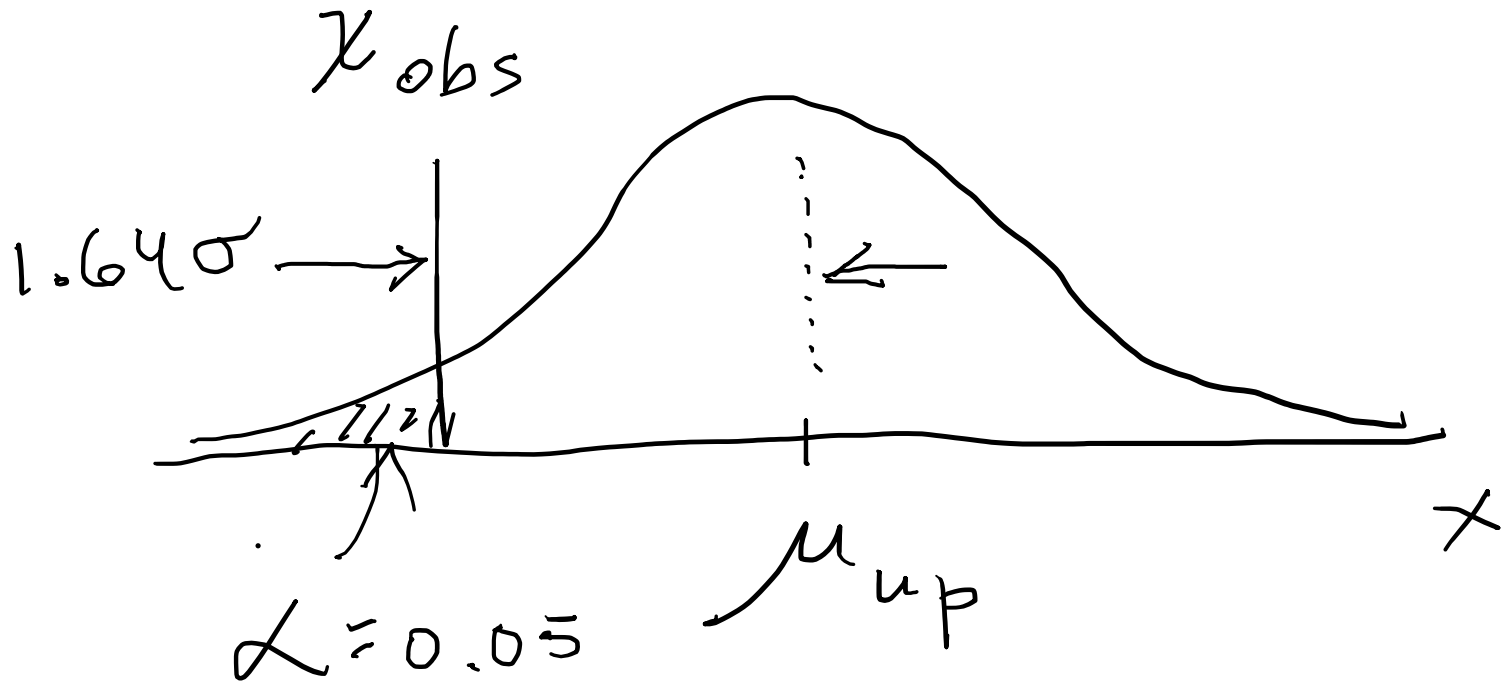
$$\mu_0 \rightarrow \mu_{\text{up}} = x_{\text{obs}} - \sigma \Phi^{-1}(\alpha) = x_{\text{obs}} + \sigma \Phi^{-1}(1 - \alpha)$$

This is an upper limit on μ , i.e., higher μ have even lower p -value and are in even worse agreement with the data.

Usually use $\Phi^{-1}(\alpha) = -\Phi^{-1}(1-\alpha)$ so as to express the upper limit as x_{obs} plus a positive quantity. E.g. for $\alpha = 0.05$, $\Phi^{-1}(1-0.05) = 1.64$.

Upper limit on Gaussian mean (3)

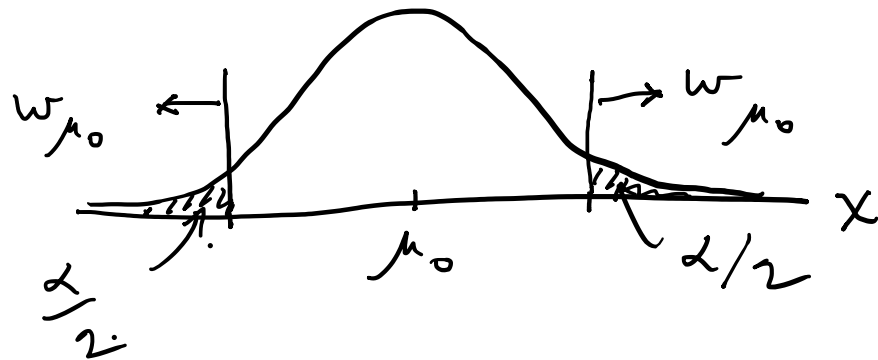
μ_{up} = the hypothetical value of μ such that there is only a probability α to find $x < x_{\text{obs}}$.



1- vs. 2-sided intervals

Now test: $H_0 : \mu = \mu_0$ versus the alternative $H_1 : \mu \neq \mu_0$

I.e. we consider the alternative to μ_0 to include higher and lower values, so take critical region on both sides:



Result is a “central” confidence interval $[\mu_{lo}, \mu_{up}]$:

$$\mu_{lo} = x_{obs} - \sigma \Phi^{-1} \left(1 - \frac{\alpha}{2} \right)$$

E.g. for $\alpha = 0.05$

$$\mu_{up} = x_{obs} + \sigma \Phi^{-1} \left(1 - \frac{\alpha}{2} \right)$$

$$\Phi^{-1} \left(1 - \frac{\alpha}{2} \right) = 1.96 \approx 2$$

Note upper edge of two-sided interval is higher (i.e. not as tight of a limit) than obtained from the one-sided test.

On the meaning of a confidence interval

Often we report the confidence interval $[a, b]$ together with the point estimate as an “asymmetric error bar”, e.g.,

$$\begin{array}{c}
 \hat{\theta} + d \\
 \hat{\theta} \\
 -c
 \end{array}
 \quad
 \begin{array}{c}
 \text{---|---|---} \\
 \uparrow \quad \hat{\theta} \quad \uparrow \\
 a = \hat{\theta} - c \quad b = \hat{\theta} + d
 \end{array}$$

E.g. (at $CL = 1 - \alpha = 68.3\%$):

$$\hat{\theta} = 80.25 \begin{array}{l} + 0.31 \\ - 0.25 \end{array}$$

Does this mean $P(80.00 < \theta < 80.56) = 68.3\%$? No, not for a frequentist confidence interval. The parameter θ does not fluctuate upon repetition of the measurement; the endpoints of the interval do, i.e., the endpoints of the interval fluctuate (they are functions of data):

$$P(a(x) < \theta < b(x)) = 1 - \alpha$$

Statistical Data Analysis

Lecture 9-4

- Confidence intervals from the likelihood function

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 θ . Equivalently, usually define

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

so higher t_θ means worse agreement between θ and the data.

p -value of θ 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...)

$$f(t_{\boldsymbol{\theta}}|\boldsymbol{\theta}) \sim \chi_N^2$$

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

Assuming this holds, the p -value is

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

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

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

Recall also

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

Confidence region from Wilks' theorem (cont.)

i.e., boundary of confidence region in θ space is where

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

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

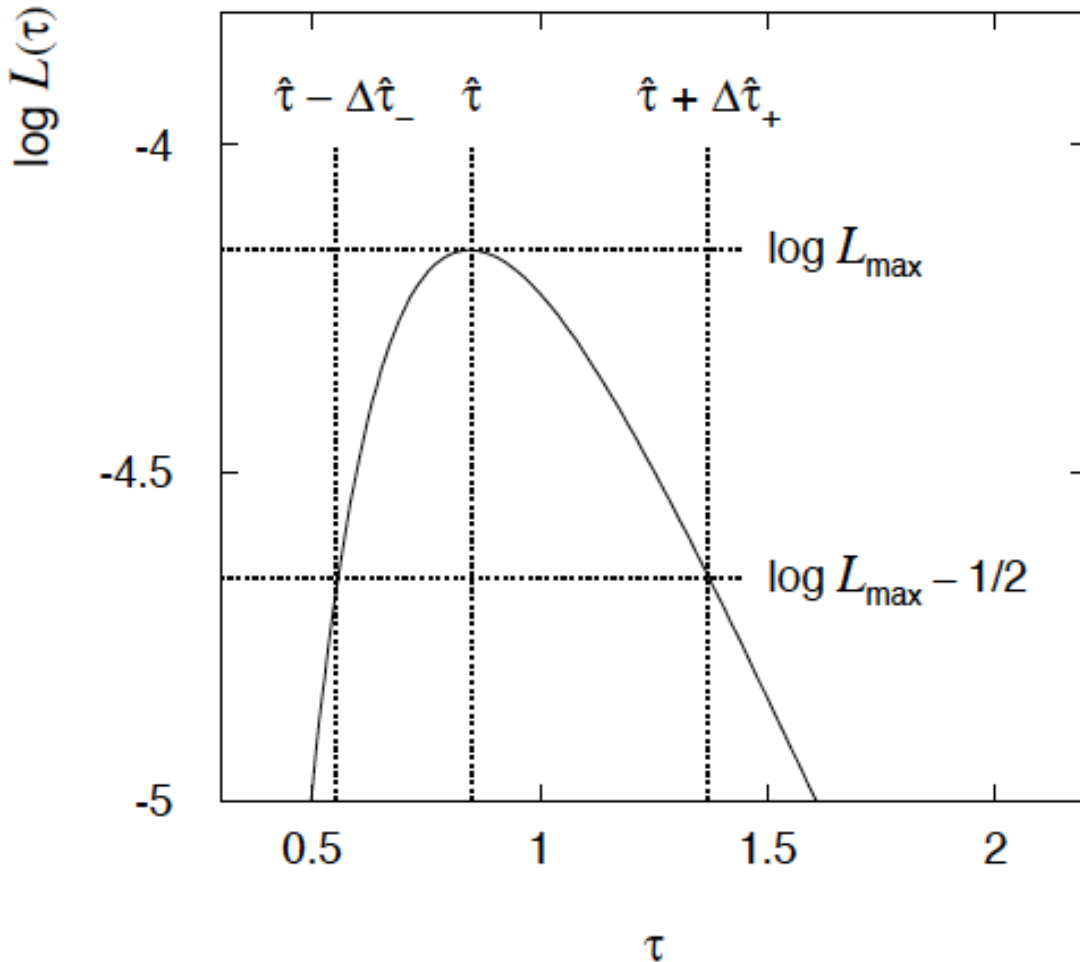
$$\ln L(\theta) = \ln L(\hat{\theta}) - \frac{1}{2}$$

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, $CL = 0.683$, $Q_\alpha = 1$.



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_α	$1 - \alpha$				
	$n = 1$	$n = 2$	$n = 3$	$n = 4$	$n = 5$
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

← # of par.

Multiparameter case (cont.)

Equivalently, Q_α increases with n for a given $CL = 1 - \alpha$.

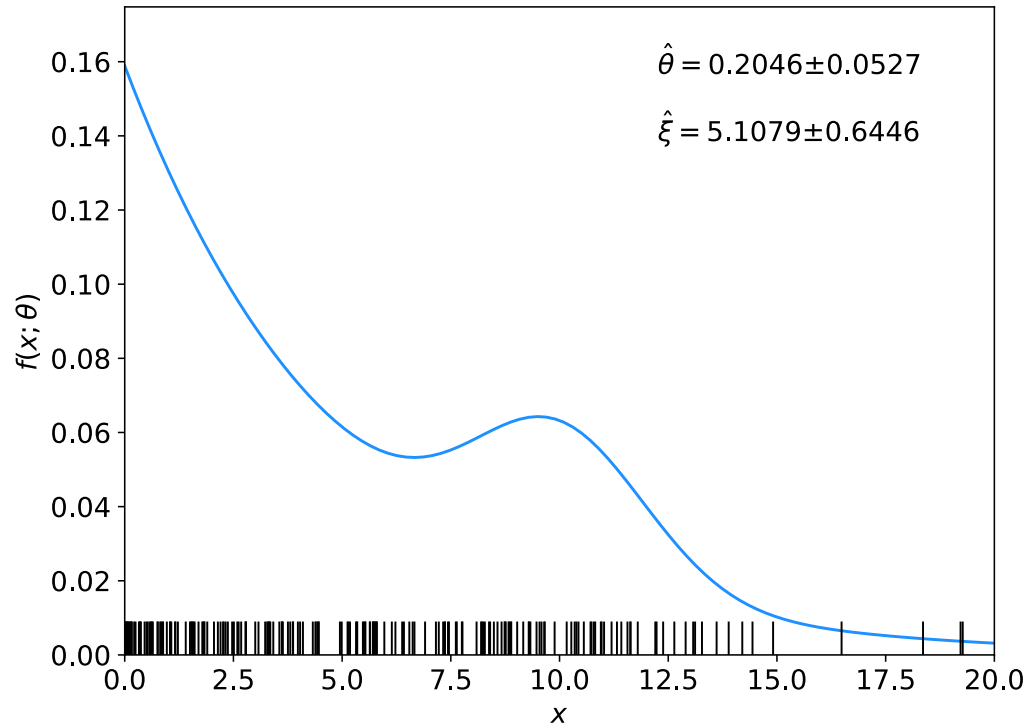
$1 - \alpha$	\bar{Q}_α				
	$n = 1$	$n = 2$	$n = 3$	$n = 4$	$n = 5$
0.683	1.00	2.30	3.53	4.72	5.89
0.90	2.71	4.61	6.25	7.78	9.24
0.95	3.84	5.99	7.82	9.49	11.1
0.99	6.63	9.21	11.3	13.3	15.1

← # of par.

Example: 2 parameter fit:

Example from problem sheet 8, i.i.d. sample of size 200

$$x \sim f(x; \theta, \xi) = \theta \frac{1}{\sqrt{2\pi}\sigma} e^{-(x-\mu)^2/2\sigma^2} + (1 - \theta) \frac{1}{\xi} e^{-x/\xi}$$

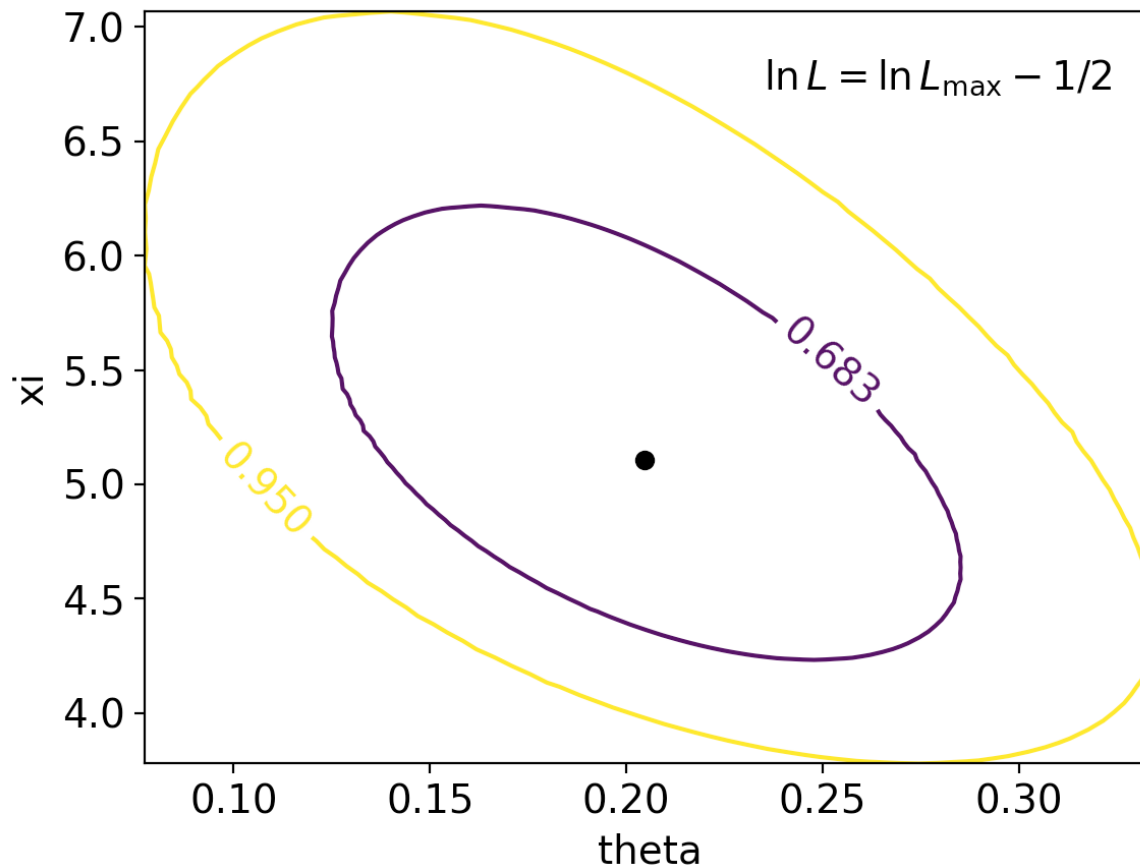


Here fit two parameters:
 θ and ξ .

Example: 2 parameter fit:

In iminuit v2, user can set $CL = 1 - \alpha$

```
m.draw_mncontour('theta', 'xi', cl=[0.683, 0.95], size=200)
```

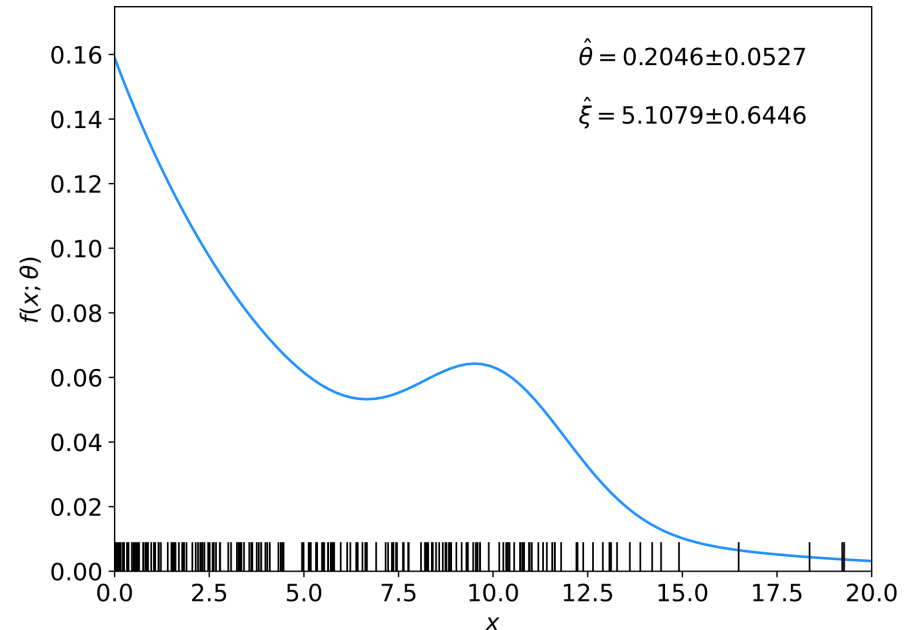


Extra slides

Comments on using iminuit

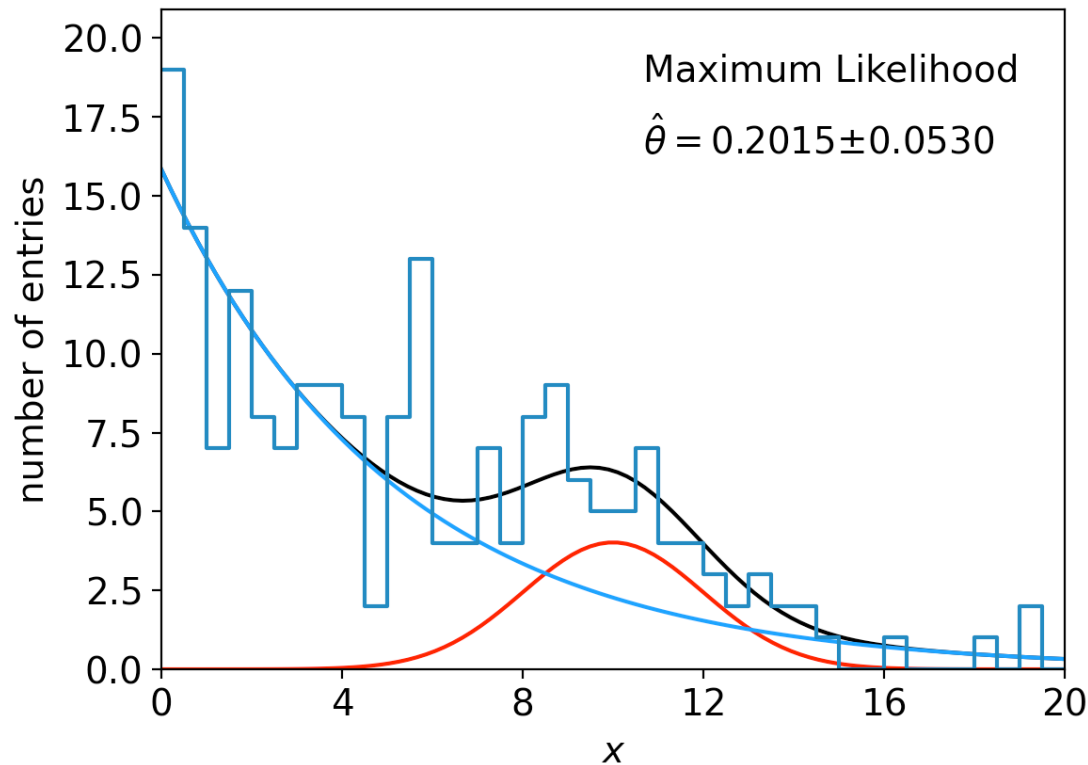
In our earlier iminuit example mlFit.py, the only argument of the log-likelihood function was the parameter array, and the data array xData entered as global (usually not a good idea):

```
def negLogL(par):  
    pdf = f(xData, par)  
    return -np.sum(np.log(pdf))  
  
    ⋮  
  
m = Minuit(negLogL, par, name=parname)
```



InL in a class, binned data,...

Sometimes it is convenient to have the function being minimized as a method of a class. An example of this is shown in the program `histFit.py`, which does the same fit as in `mlFit.py` but with a histogram of the data:



Commentary on histFit.py

The global data can be avoided if we make the objective function a method of a class:

```
class ChiSquared:                                # function to be minimized

    def __init__(self, xHist, bin_edges, fitType):
        self.setData(xHist, bin_edges)
        self.fitType = fitType

    def setData(self, xHist, bin_edges):
        numVal = np.sum(xHist)
        numBins = len(xHist)
        binSize = bin_edges[1] - bin_edges[0]
        self.data = xHist, bin_edges, numVal, numBins, binSize

    def chi2LS(self, par):                        # least squares
        xHist, bin_edges, numVal, numBins, binSize = self.data
        xMid = bin_edges[:numBins] + 0.5*binSize
        binProb = f(xMid, par)*binSize
        nu = numVal*binProb
        sigma = np.sqrt(nu)
        z = (xHist - nu)/sigma
        return np.sum(z**2)
```

class ChiSquared (continued)

```
def chi2M(self, par):          # multinomial maximum likelihood
    xHist, bin_edges, numVal, numBins, binSize = self.data
    xMid = bin_edges[:numBins] + 0.5*binSize
    binProb = f(xMid, par)*binSize
    nu = numVal*binProb
    lnL = 0.
    for i in range(len(xHist)):
        if xHist[i] > 0.:
            lnL += xHist[i]*np.log(nu[i]/xHist[i])
    return -2.*lnL

def __call__(self, par):
    if self.fitType == 'LS':
        return self.chi2LS(par)
    elif self.fitType == 'M':
        return self.chi2M(par)
    else:
        print("fitType not defined")
        return -1
```

Using the ChiSquared class

```
# Put data values into a histogram
numBins=40
xHist, bin_edges = np.histogram(xData, bins=numBins, range=(xMin, xMax))
binSize = bin_edges[1] - bin_edges[0]

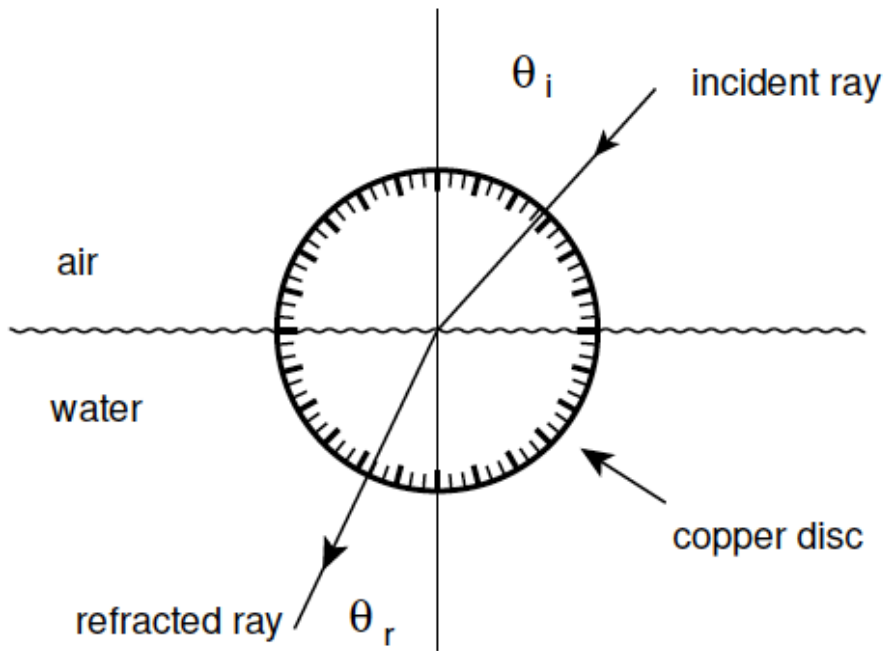
# Initialize Minuit and set up fit:
parin = np.array([theta, mu, sigma, xi]) # initial values (here = true)
parname = ['theta', 'mu', 'sigma', 'xi']
parstep = np.array([0.1, 1., 1., 1.]) # initial setp sizes
parfix = [False, True, True, False] # change to fix/free param.
parlim = [(0.,1), (None, None), (0., None), (0., None)]
chisq = ChiSquared(xHist, bin_edges, fitType)
m = Minuit(chisq, parin, name=parname)
m.errors = parstep
m.fixed = parfix
m.limits = parlim
m.errordef = 1.0 # errors from chi2 = chi2min + 1
```

For full program see

<https://www.pp.rhul.ac.uk/~cowan/stat/exercises/fitting/python/>

LS example: refraction data from Ptolemy

Astronomer Claudius Ptolemy obtained data on refraction of light by water in around 140 A.D.:



Angles of incidence and refraction (degrees)

θ_i	θ_r
10	8
20	$15\frac{1}{2}$
30	$22\frac{1}{2}$
40	29
50	35
60	$40\frac{1}{2}$
70	$45\frac{1}{2}$
80	50

Suppose the angle of incidence is set with negligible error, and the measured angle of refraction has a standard deviation of $\frac{1}{2}^\circ$.

Laws of refraction

A commonly used law of refraction was

$$\theta_r = \alpha\theta_i ,$$

although it is reported that Ptolemy preferred

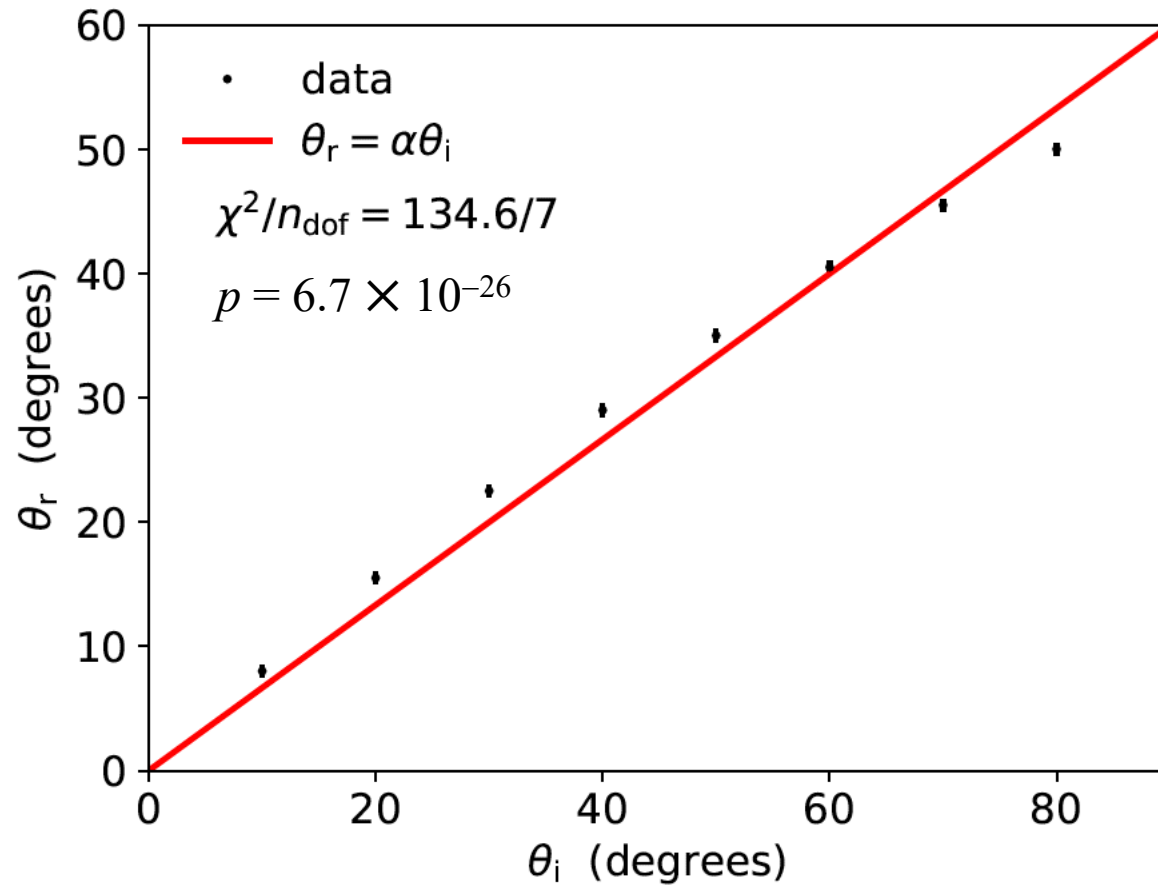
$$\theta_r = \alpha\theta_i - \beta\theta_i^2 .$$

The law of refraction discovered by Ibn Sahl in 984 (and rediscovered by Snell in 1621) is

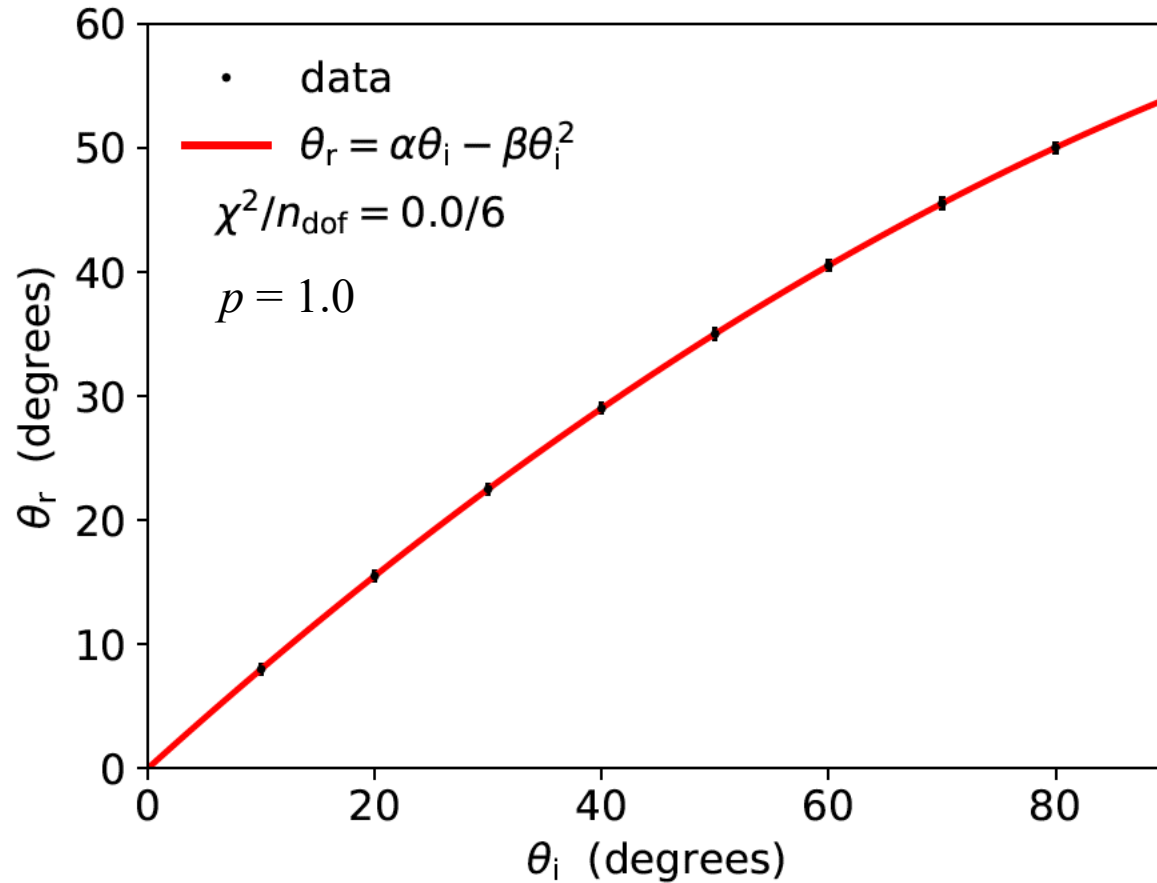
$$\theta_r = \sin^{-1} \left(\frac{\sin \theta_i}{r} \right) .$$

where $r = n_r/n_i$ is the ratio of indices of refraction of the two media.

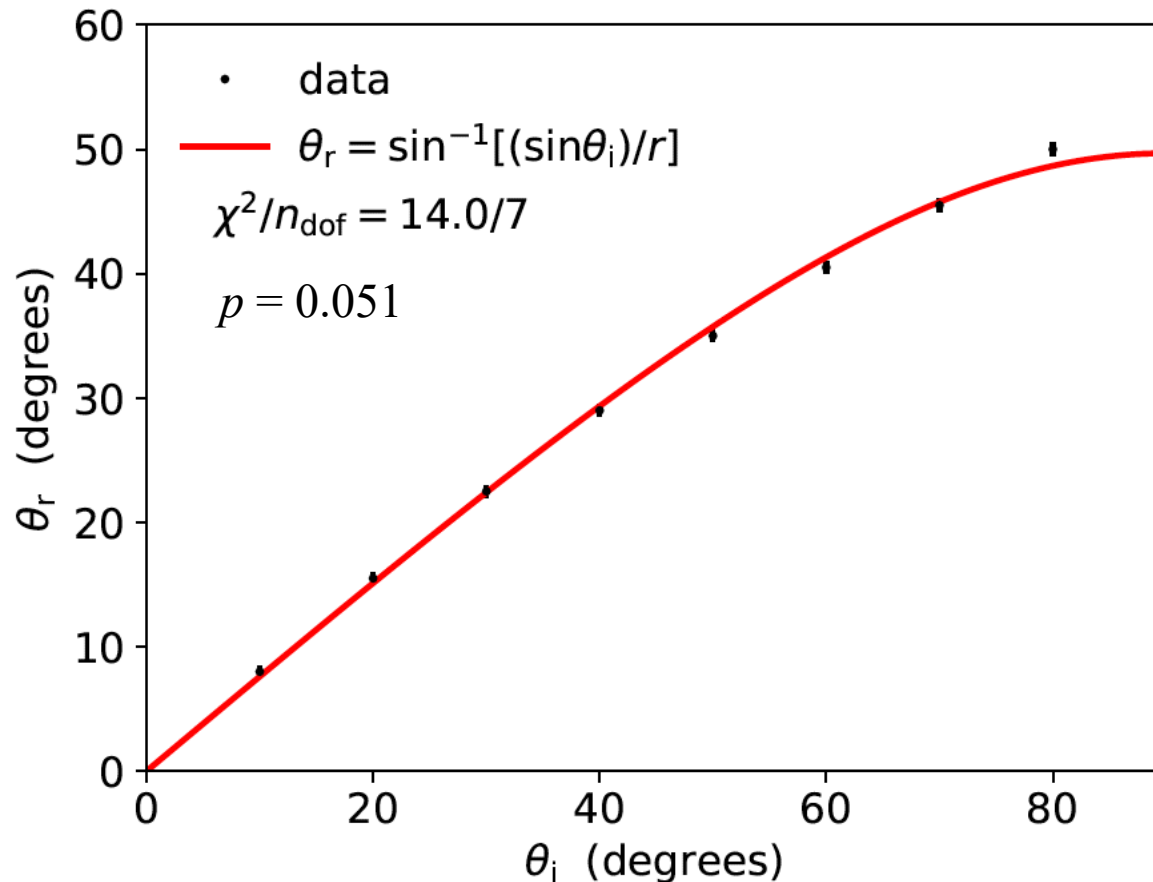
LS fit: $\theta_r = \alpha\theta_i$



LS fit: $\theta_r = \alpha\theta_i - \beta\theta_i^2$



LS fit: Snell's Law



Fitted index of refraction of water $r = 1.3116 \pm 0.0056$ found not quite compatible with currently known value 1.330.