Statistical Data Analysis 2024/25 Lecture Week 9



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



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Course web page via RHUL moodle (PH4515) and also www.pp.rhul.ac.uk/~cowan/stat_course.html

Statistical Data Analysis Lecture 9-1

• Least squares with histogram data

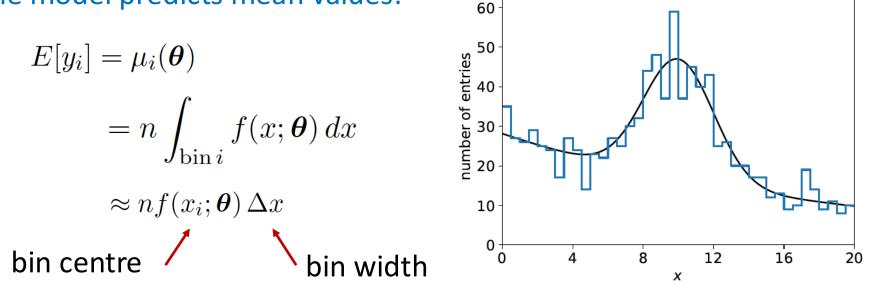
LS with histogram data

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 $x_1, ..., x_n$ 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*: $y = (y_1, ..., y_N)$.

The model predicts mean values:



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LS with histogram data (2)

The usual models:

for fixed sample size n, take $y \sim$ 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 ~ Gauss with $\sigma_i \approx \sqrt{\mu_i}$. ($\rightarrow \sigma_i$ depends on θ).

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 :

 $\sigma_i = \sqrt{\mu_i(\theta)}$ (LS method) $\sigma_i = \sqrt{y_i}$ (Modified LS method)

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

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

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LS with histogram data — normalization

Do not "fit" the normalization, i.e., $n \rightarrow$ free parameter v:

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

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

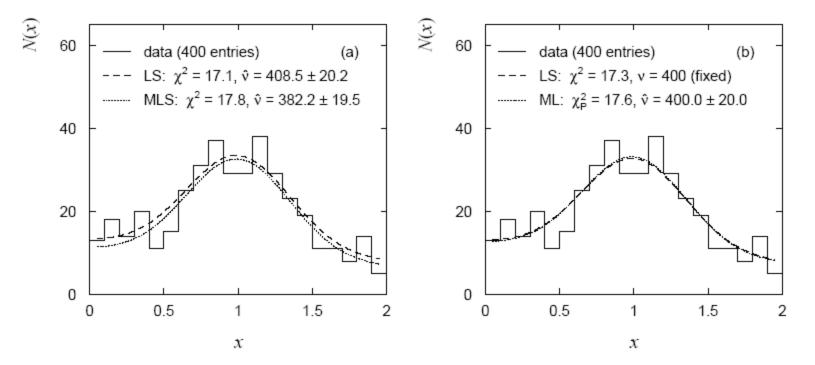
$$\hat{\nu}_{\rm LS} = n + \frac{\chi^2_{\rm min}}{2}$$

$$\hat{\nu}_{\rm MLS} = n - \chi^2_{\rm min}$$

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^2_{\min} around N-m,

 \rightarrow relative error in $\hat{\nu}$ large when N large, n small Either get n directly from data for LS (or better, use ML).

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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 $L(\mu)$ that depends on N parameters $\mu = (\mu_1, ..., \mu_N)$. Define the statistic

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

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

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

Larger t_{μ} means less compatibility between data and μ .

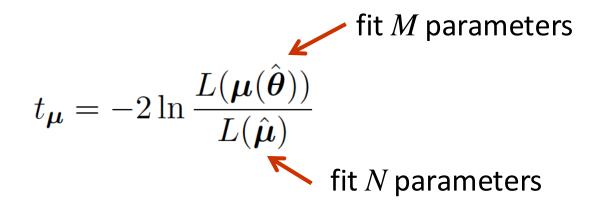
Quantify "goodness of fit" with *p*-value: $p_{\mu} = \int_{t_{\mu,obs}}^{\infty} f(t_{\mu}|\mu) dt_{\mu}$ need this pdf

Likelihood ratio (2)

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

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

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



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

Wilks' Theorem

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

$$t_{\mu} = -2 \ln \frac{L(\mu(\hat{\theta}))}{L(\hat{\mu})}$$
 follows a chi-square distribution for
$$N - M \text{ degrees of freedom.}$$

MLE of $(\mu_1, ..., \mu_N)$

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

Proof boils down to having all estimators ~ Gaussian.

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

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Wilks' Theorem (2)

To find $p_{\theta} = \int_{t_{\mu},\text{obs}}^{\infty} f(t_{\mu}|\mu(\theta)) dt_{\mu}$ e.g. with Monte Carlo we

would need to choose a point in θ space, then $p = \max_{\theta} p_{\theta}$

But if we can use Wilks', the chi-square dist. should hold for all θ .

The chi-square pdf for $-2\ln\lambda$ breaks down:

if the sample size is too small;

if the true value of a parameter is on the boundary of the allowed parameter space;

if the model in the numerator is not a special case of the denominator (models must be "nested");

if variance of estimators of any components of μ too large (e.g., parameter refers to location of a feature not present in the null hypothesis, such as the position of a peak).

Goodness of fit with Gaussian data

Suppose the data are *N* independent Gaussian distributed values:

$$y_i \sim \mathrm{Gauss}(\mu_i, \sigma_i) \;, \qquad i=1,\ldots,N$$
 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: $\hat{\mu}_i = y_i$ $i = 1, \dots, N$

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Likelihood ratio for Gaussian data

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

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

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

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

Here t_{μ} is the same as $\chi^2_{\rm min}$ from an LS fit.

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

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Likelihood ratio for Poisson data

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

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

First (for LR denominator) use saturated model, i.e., treat $v = (v_1, ..., v_N)$ as all adjustable:

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

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

ML estimators: $\hat{
u}_{m{i}}=n_{m{i}}$, $m{i}=1,\ldots,N$

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Goodness of fit with Poisson data (2)

For LR numerator find $v(\theta)$ with *M* fitted parameters $\theta = (\theta_1, ..., \theta_M)$:

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

if $n_i = 0$, skip log term

Wilks' theorem: in large-sample limit $t_{m
u} \sim \chi^2_{N-M}$

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

As before use t_v to get p-value of $v(\theta)$,

independent of heta

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

Goodness of fit with multinomial data

Similar if data $\mathbf{n} = (n_1, ..., n_N)$ 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:

$$n_{\rm tot} = \sum_{i=1}^{N} n_i$$

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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
$$n_{\rm tot}$$
 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 $t_{\nu} \sim \chi^2_{N-M-1}$

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:

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

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

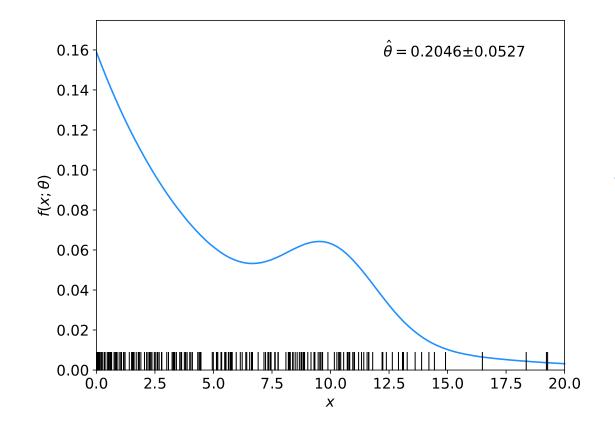
Steve Baker and Robert D. Cousins, *Clarification of the use of the chi-square and likelihood functions in fits to histograms*, NIM **221** (1984) 437.

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Examples of ML/LS fits

Unbinned maximum likelihood (mlFit.py, minimize negLogL)

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

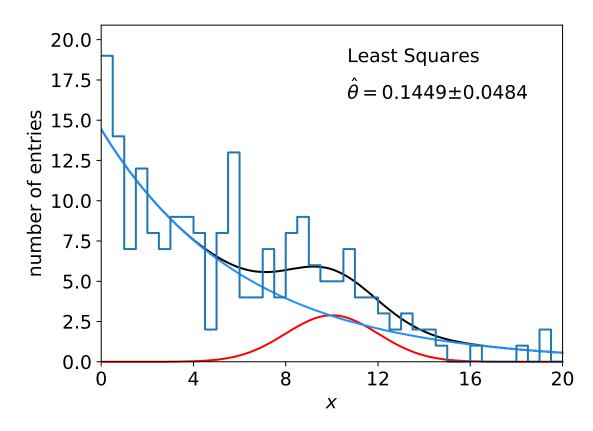


No useful measure of goodness-of-fit from unbinned ML.

Examples of ML/LS fits

Least Squares fit (histFit.py, minimize chi2LS)

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



$$\chi^{2}_{min} = 32.7$$

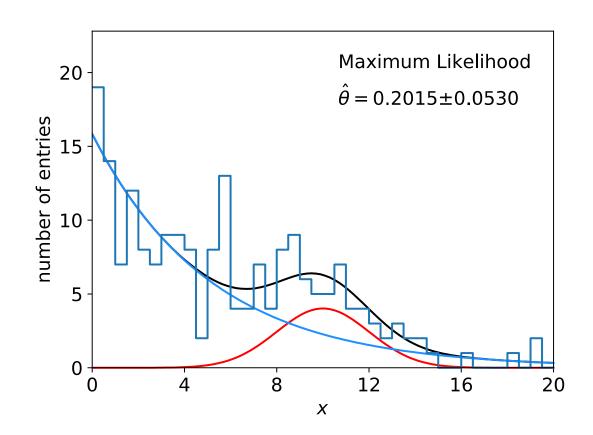
 $n_{dof} = 38$
 $p = 0.71$

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

Examples of ML/LS fits

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

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



$$\chi^2_{\rm min} = 35.3$$

 $n_{\rm dof} = 37$
 $p = 0.55$

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 $P(\text{data in critical region} | \theta) \le \alpha$ 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(conf. interval "covers" $\theta | \theta \ge 1 \square \alpha$

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

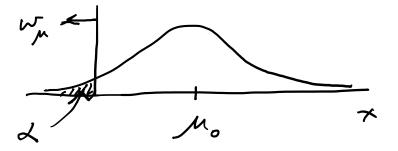
Example: upper limit on mean of Gaussian

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 $H_1: \mu < \mu_0$

 \rightarrow Put w_{μ} at region of x-space characteristic of low μ (i.e. at low x)

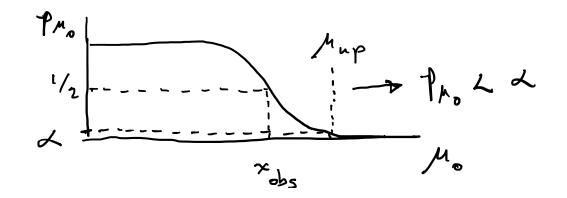


Equivalently, take the *p*-value to be

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

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



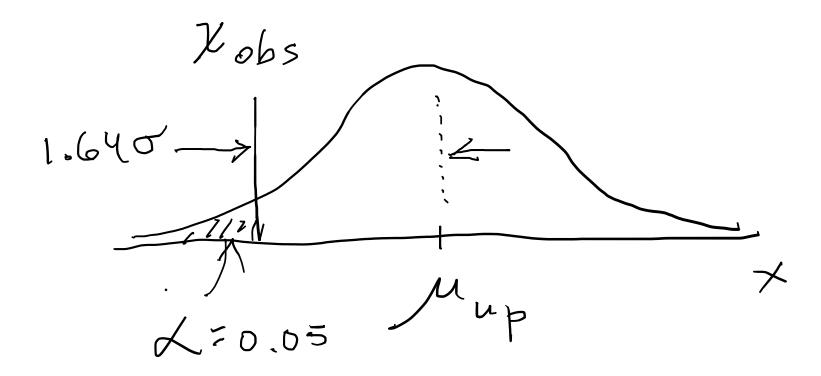
$$\mu_0 \to \mu_{\rm up} = x_{\rm obs} - \sigma \Phi^{-1}(\alpha) = x_{\rm obs} + \sigma \Phi^{-1}(1 - \alpha)$$

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)

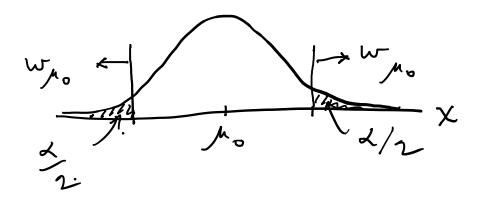
 μ_{up} = the hypothetical value of μ such that there is only a probability α to find $x < x_{obs}$.



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 $[\mu_{lo}, \mu_{up}]$:

$$\mu_{\rm lo} = x_{\rm obs} - \sigma \Phi^{-1} \left(1 - \frac{\alpha}{2} \right) \qquad \text{E.g. for } \alpha = 0.05$$
$$\mu_{\rm up} = x_{\rm obs} + \sigma \Phi^{-1} \left(1 - \frac{\alpha}{2} \right) \qquad \Phi^{-1} \left(1 - \frac{\alpha}{2} \right) = 1.96 \approx 2$$

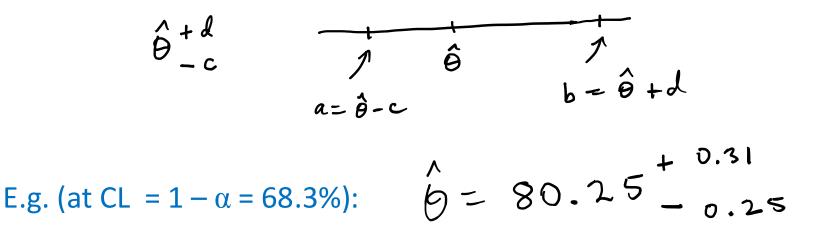
Note upper edge of two-sided interval is higher (i.e. not as tight of a limit) than obtained from the one-sided test.

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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.,



Does this mean P(80.00 < θ < 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):

$$P(alx) L \Theta L b(x)) = 1 - \alpha$$

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Example with binomial parameter

Suppose $m \sim \text{Binomial}(N,\theta)$ with N trials (known) and success probability per trial θ (unknown). We observe a single value m.

The likelihood function is

$$L(\theta) = P(m|N,\theta) = \frac{N!}{m!(N-m)!}\theta^m (1-\theta)^{N-m}$$

so the log-likelihood is $\ln L(\theta) = m \ln \theta + (N-m) \ln(1-\theta) + C$

Set its derivative to zero

$$\frac{\partial \ln L}{\partial \theta} = \frac{m}{\theta} - \frac{N-m}{\theta} = 0$$

to find the MLE $\hat{\theta} = \frac{m}{N}$.

Since $V[m] = N\theta(1-\theta) \rightarrow \sigma_{\hat{\theta}} = \frac{1}{N}\sqrt{\theta(1-\theta)} \rightarrow \hat{\sigma}_{\hat{\theta}} = \frac{1}{N}\sqrt{\frac{m}{N}\left(1-\frac{m}{N}\right)}$

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Limits on binomial parameter

To give the MLE and a 68.3% central confidence interval, it is often sufficient to report $\hat{\theta} \pm \sigma_{\hat{\theta}}$.

Suppose we find $m_{\rm obs}$ and we want to know an upper limit on θ .

To quantify how big θ could be, find upper limit at CL = $1-\alpha$ = 95%.

$$p_{\theta} = P(m \le m_{\text{obs}} | \theta) = \sum_{m=0}^{m_{\text{obs}}} \frac{N!}{m!(N-m)!} \theta^m (1-\theta)^{N-m}$$

Set $p_{\theta} = \alpha$ and solve for $\theta \to \theta_{up}$.

Can be done in closed form; see PDG Eq. (40.83):

$$\theta_{\rm up} = \frac{(m+1)F_F^{-1}[1-\alpha;2(m+1),2(N-m)]}{(N-m)+(m+1)F_F^{-1}[1-\alpha;2(m+1),2(N-m)]}$$

usually just solve with computer

where F is the Fisher-Snedecor distribution .

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Upper limit for θ for $m_{\rm obs} = 0$

Suppose we find $m_{\rm obs} = 0$.

 $\hat{ heta}=0$ makes sense

 $\hat{\sigma}_{\hat{ heta}} = 0$ not incorrect but does not provide a useful interval

For the *p*-value (for upper limit) we find

$$p_{\theta} = \sum_{m=0}^{0} \frac{N!}{0!(N-0)!} \theta^{0} (1-\theta)^{N-0} = (1-\theta)^{N}$$

Set $p_{\theta} = \alpha$ and solving for θ gives the upper limit $\theta_{up} = 1 - \alpha^{1/N}$

For example, $N = 20, \alpha = 0.05, \rightarrow \theta_{up} = 0.14$ at 95% CL.

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) $\theta = (\theta_1, ..., \theta_N)$ using the ratio

$$\lambda(\theta) = \frac{L(\theta)}{L(\hat{\theta})} \qquad \qquad 0 \le \lambda(\theta) \le 1$$

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

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Confidence region from Wilks' theorem

Wilks' theorem says (in large-sample limit and provided certain conditions hold...)

 $f(t_{\theta}|\theta) \sim \chi_N^2$ chi-square dist. with # d.o.f. = # of components in $\theta = (\theta_1, ..., \theta_N)$.

Assuming this holds, the *p*-value is

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

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

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

Recall also

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

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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

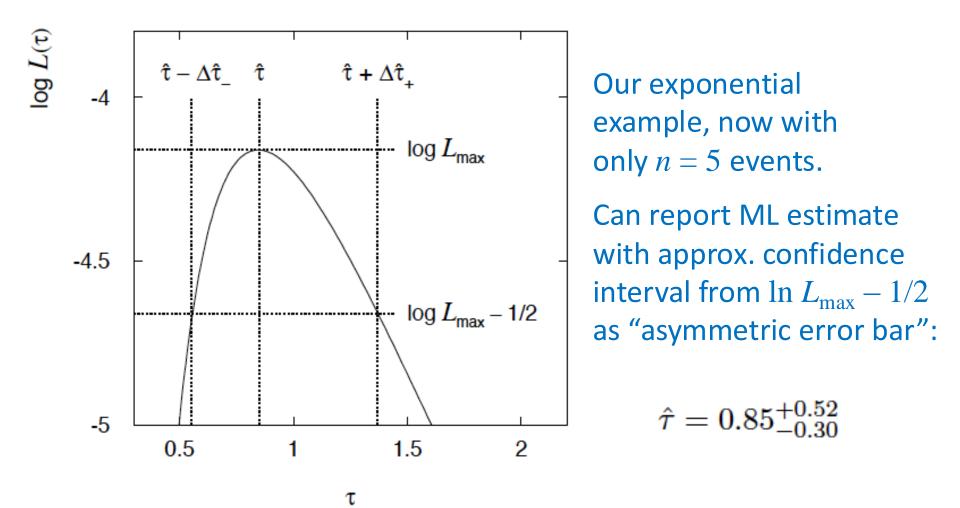
$$\ln L(\theta) = \ln L(\hat{\theta}) - \frac{1}{2}$$

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 conf. interval (in large sample limit).

Example of interval from $\ln L(\theta)$

For N = 1 parameter, CL = 0.683, $Q_{\alpha} = 1$.



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_{lpha}		-				
	n = 1	n = 2	n = 3	n = 4	n = 5	\leftarrow # of par.
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	

Multiparameter case (cont.)

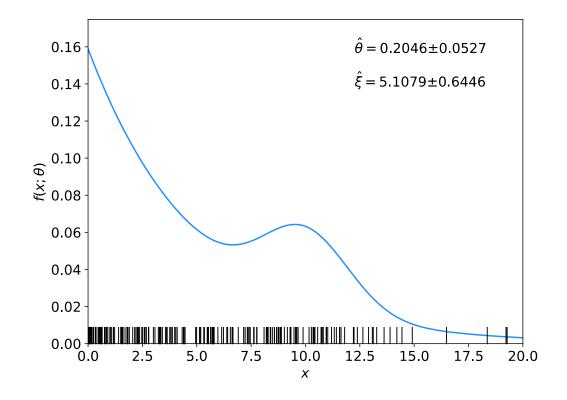
Equivalently, Q_{α} increases with *n* for a given $CL = 1 - \alpha$.

$1 - \alpha$		_				
	n = 1	n = 2	n = 3	n = 4	n = 5	\leftarrow # of par.
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	_

Example: 2 parameter fit:

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

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

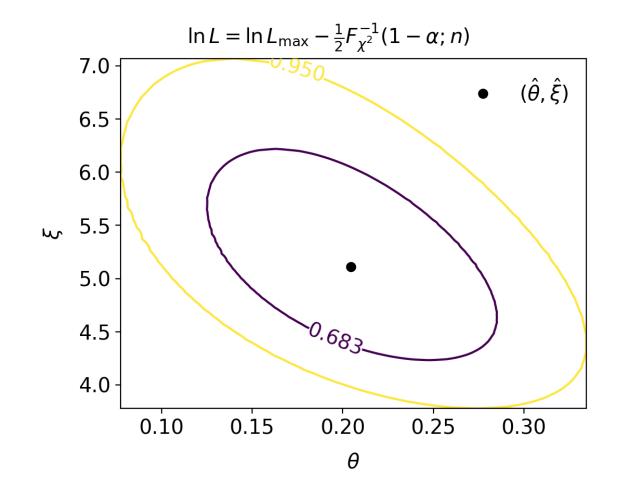


Here fit two parameters: θ and ξ .

Example: 2 parameter fit:

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

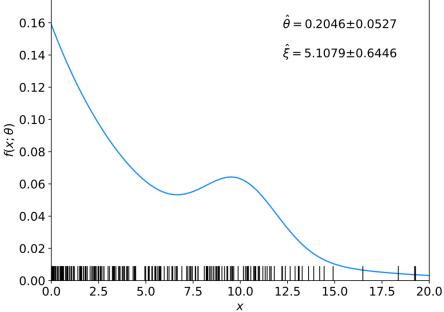
m.draw_mncontour('theta', 'xi', cl=[0.683, 0.95], size=200)



Extra slides

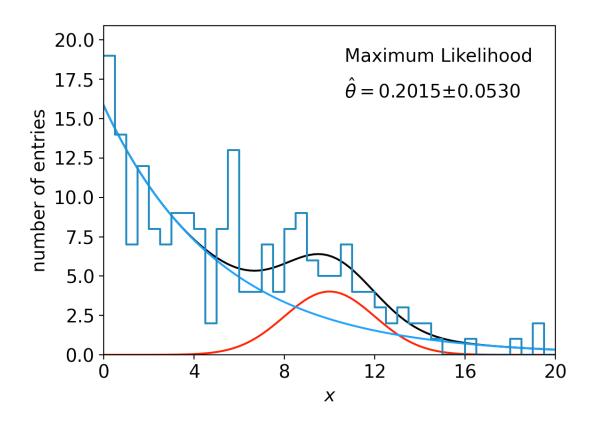
Comments on using iminuit

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):



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:



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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
   \ln L = 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

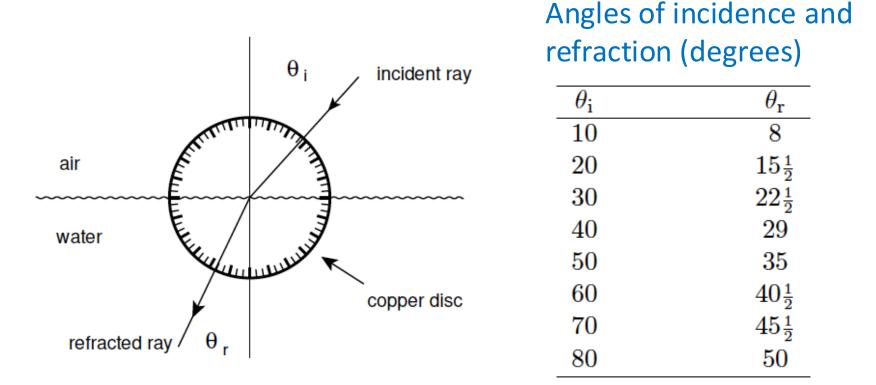
```
# Put data values into a histogram
numBins=40
xHist, bin_edges = np.histogram(xData, bins=numBins, range=(xMin, xMax))
binSize = bin_edges[1] - bin_edges[0]
```

```
# 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)]
chisg = ChiSguared(xHist, bin edges, fitType)
m = Minuit(chisg, parin, name=parname)
m.errors = parstep
m.fixed = parfix
m.limits = parlim
                                     # errors from chi2 = chi2min + 1
m.errordef = 1.0
```

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.:



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

 $heta_{
m r} = lpha heta_{
m i}$,

although it is reported that Ptolemy preferred

$$\theta_{\rm r} = \alpha \theta_{\rm i} - \beta \theta_{\rm i}^2$$
.

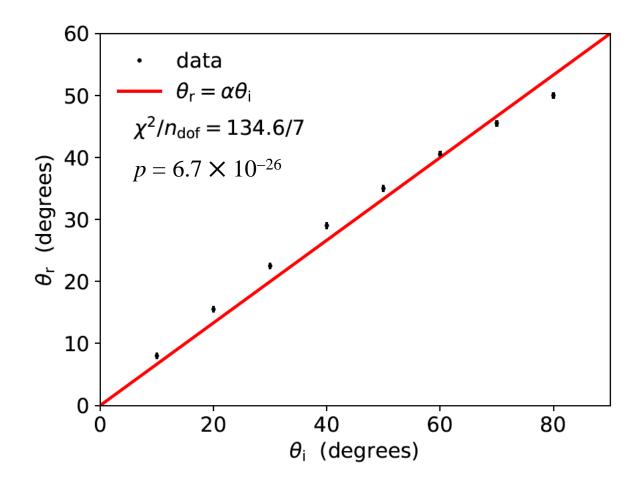
The law of refraction discovered by Ibn Sahl in 984 (and rediscovered by Snell in 1621) is

$$\theta_{\rm r} = \sin^{-1} \left(\frac{\sin \theta_{\rm i}}{r} \right).$$

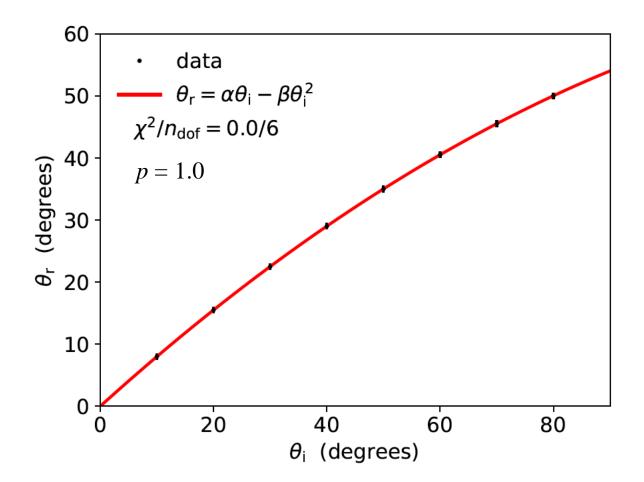
where $r = n_r/n_i$ is the ratio of indices of refraction of the two media.

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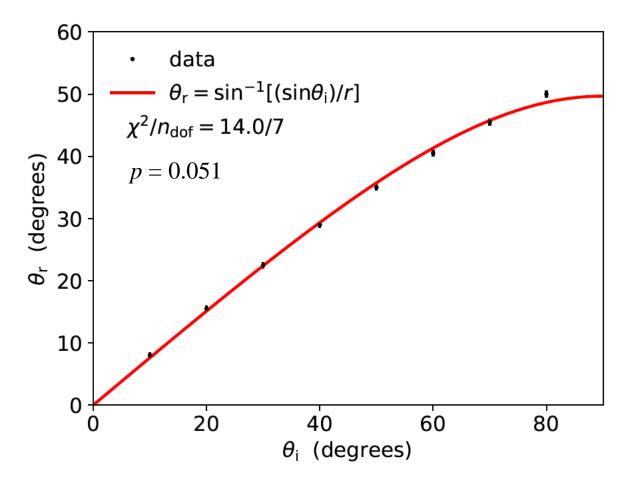
LS fit: $\theta_{\rm r} = \alpha \theta_{\rm i}$



LS fit: $\theta_{\rm r} = \alpha \theta_{\rm i} - \beta \theta_{\rm i}^2$



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.

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Statistical Data Analysis / lecture week 9