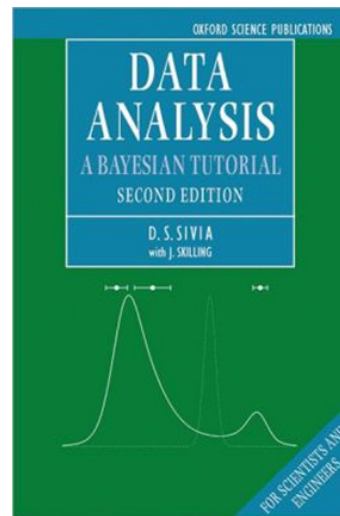


FABADA: a Fitting Algorithm for Bayesian Analysis of DAta



Luis Carlos Pardo Soto
Grup de Caracterització de Materials (GCM)



- The ubiquitous χ^2
- Advantages of Bayesian analysis
- Some examples
 - ✓ Quantitative guide to the eye
 - ✓ Analysis of QENS data
 - ✓ Intramolecular structure determination
- Robustness and simulated annealing
- Summary and conclusions



Why are Nonlinear Fits to Data so Challenging?

Mark K. Transtrum,^{*} Benjamin B. Machta,[†] and James P. Sethna[‡]

Laboratory of Atomic and Solid State Physics, Cornell University, Ithaca, New York 14853, USA

(Received 22 September 2009; published 10 February 2010)

In other words

- Why do they get stuck?
- Why do software thinks that it has finished to fit?
- How sure am I about the results?

Why need a tool to efficiently explore the χ^2 landscape?

Bayes theorem



Likelihood

Probability that our data
describes the hypothesis

Prior

Our forehand knowledge

$$P(H | D) = \frac{P(D | H) \cdot P(H)}{P(D)}$$

Posterior

Probability that the hypothesis is true
given the experimental data

T. Bayes.

Evidence

Normalization factor

In our case is very simple...

Bayes theorem



Likelihood

Probability that our data
describes the hypothesis

Maximum ignorance Prior

$$P(H | D) \propto \frac{P(D | H) \cdot P(H)}{P(D)}$$

Posterior

Probability that the hypothesis is true
given the experimental data

T. Bayes.

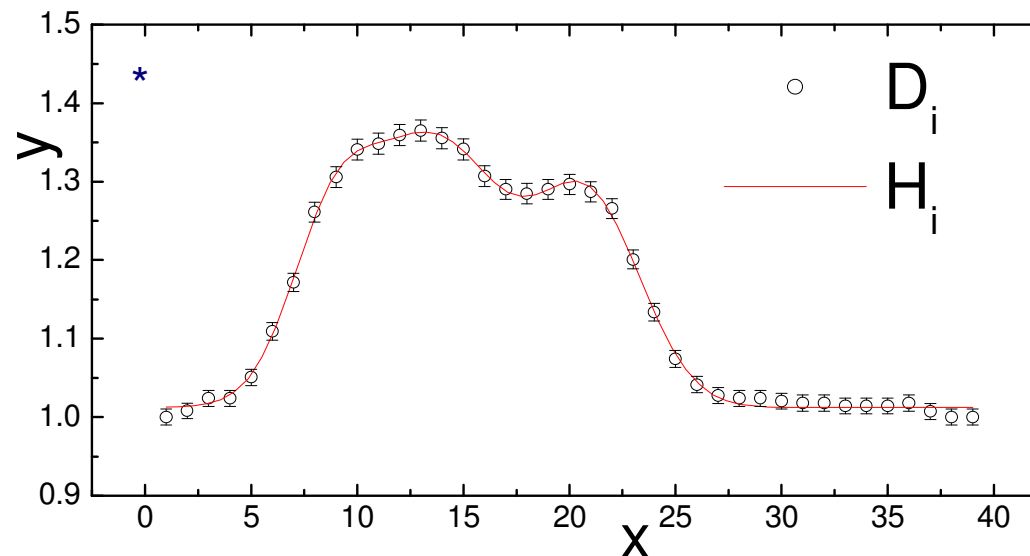
We only care about proportionality

Bayes theorem



$$P(H \mid D) \propto P(D \mid H) \equiv L$$

D_i Data ($i=1,n$) $H_i\{P_l\}$ Hypothesis ($i=1,n$) using a parameter set $\{P_l\}$ ($l=1,m$)



$$P(H_i\{P_l\} \mid D_i) \propto P(D_i \mid H_i\{P_l\})$$

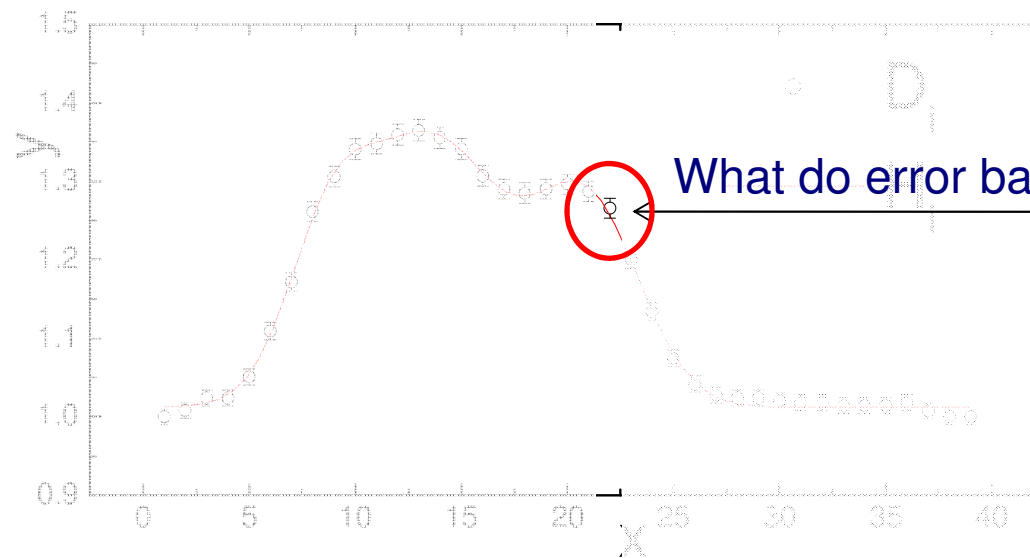
* Figure adapted from „Le petit prince“ A. Saint Exupery (1943)

Bayes theorem



$$P(H | D) \propto P(D | H)$$

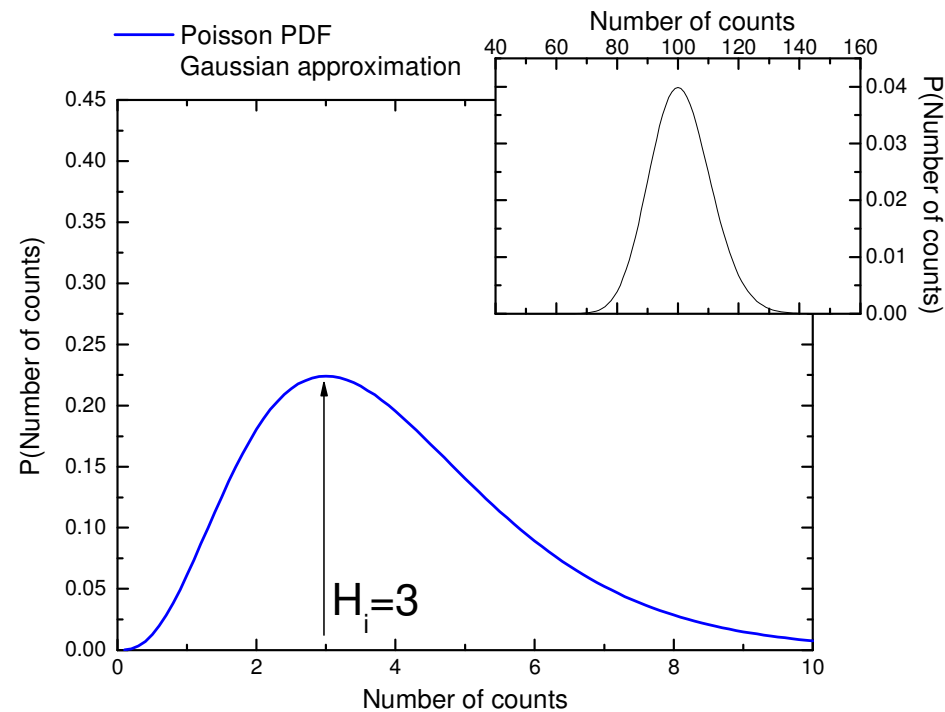
D_i Data ($i=1,n$) $H_i, \{P_l\}$ Hypothesis ($i=1,n$) using a parameter set $\{P_l\}$ ($l=1,m$)



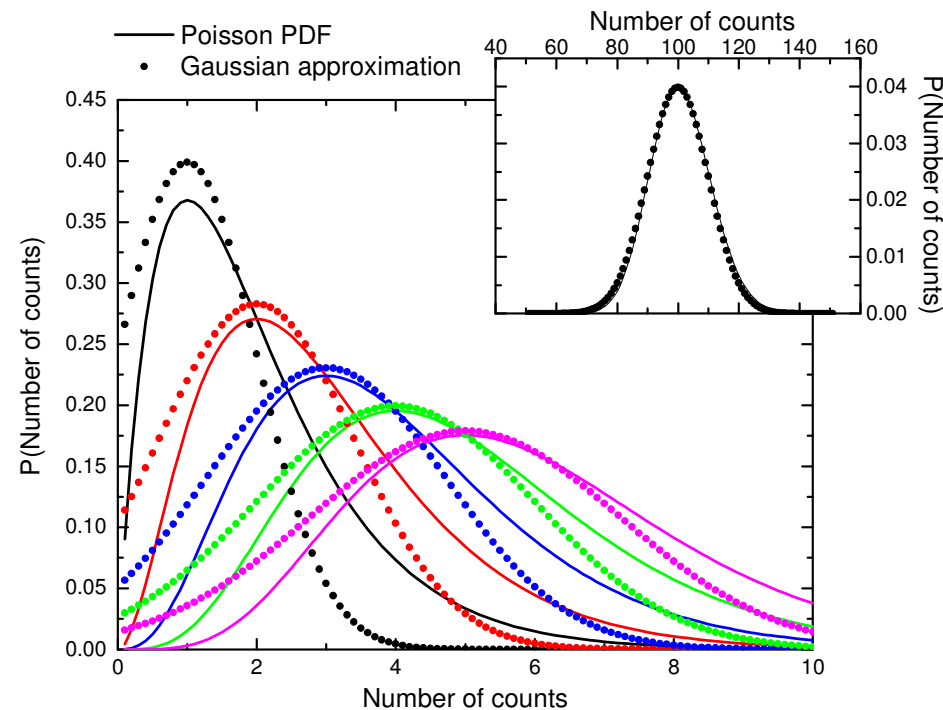
$$P(H_i, \{P_l\} | D_i) \propto P(D_i | H_i, \{P_l\})$$

In a counting experiment, if the expected value is H_i measured values will follow a Poisson statistics

$$P(D_{i=k} | H_{i=k}) \propto H_i^{D_{i=k}} e^{-H_{i=k}}$$



In a counting experiment, if the expected value is H_i measured values will follow a Poisson statistics $P(D_{i=k} | H_{i=k}) \propto H_i^{D_{i=k}} e^{-H_{i=k}}$



If the number of counts is high enough
poisson statistics = normal distribution
with $\sigma_i = \sqrt{D_i}$

$$P(D_{i=k} | H_{i=k}) \propto \exp - \frac{(H_{i=k} - D_{i=k})^2}{2\sigma_{i=k}^2}$$

The error is the square root of the variance $\epsilon_i = \sigma_i$

Bayes theorem $P(H_i \{P_l\} | D_i) \propto P(D_i | H_i \{P_l\})$



We now consider all the points $i=1,2,\dots, n$

$$\begin{aligned} L = P(D_i | H_i \{P_l\}) &\propto \prod_{i=1}^n \exp - \frac{(H_i - D_i)^2}{2\sigma_i^2} \\ &= \exp \sum_{i=1}^n - \frac{(H_i - D_i)^2}{2\sigma_i^2} = \exp - \frac{\chi^2}{2} \end{aligned}$$

Therefore χ^2 is related to the likelihood:

$$\chi^2 \propto -2 \cdot \ln L$$

**So, we got the exact meaning on probability bases of χ^2
Let's use it!**

Similar to Montecarlo simulations, (to compare let's assume that all errors σ_i are equal)

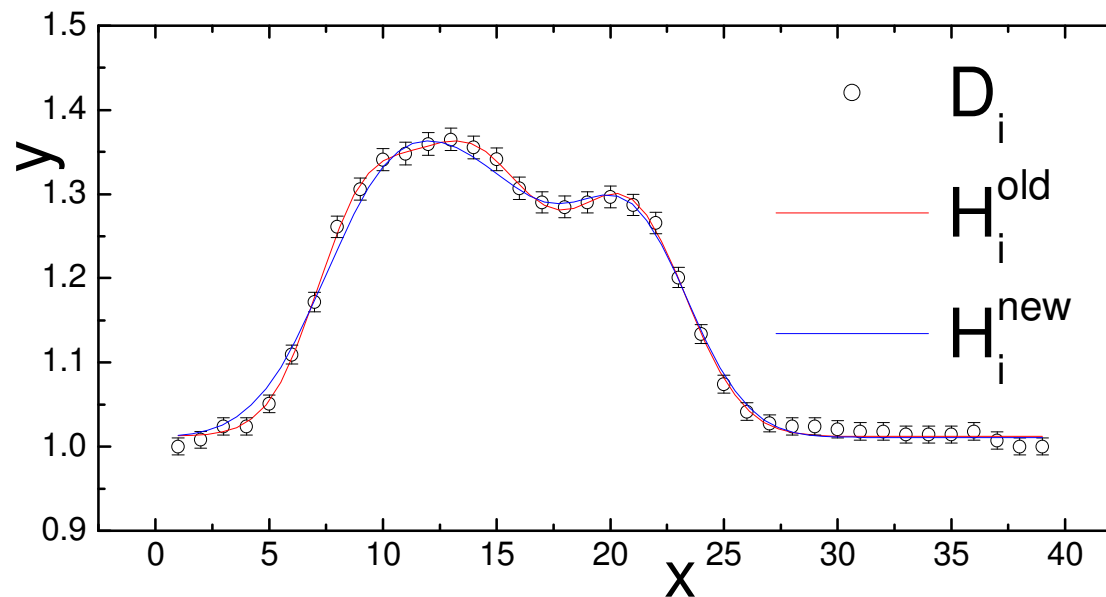
$$\frac{P(H_i\{P_l^{new}\} | D_i)}{P(H_i\{P_l^{old}\} | D_i)} = \exp - \frac{\sum_{i=1}^n (H_i^{new} - D_i)^2 - \sum_{i=1}^n (H_i^{old} - D_i)^2}{2\sigma^2} = \exp - \frac{(\chi_{new}^2 - \chi_{old}^2)}{2}$$

This term makes the fitting better

$$E \leftrightarrow \sum_{i=1}^n (H_i^{new} - D_i)^2$$

This term allows parameters that make the fitting worst

$$T \leftrightarrow 2\sigma^2$$

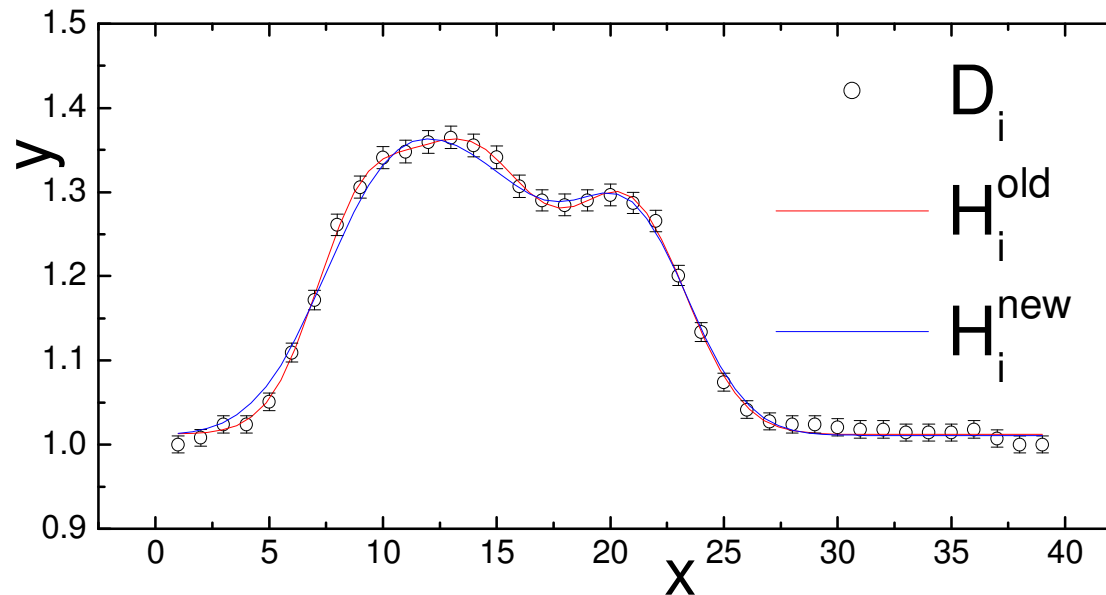


What does FABADA do?

$$\frac{P(H_i\{P_l^{new}\} | D_i)}{P(H_i\{P_l^{old}\} | D_i)} = \exp - \frac{\sum_{i=1}^n (H_i^{new} - D_i)^2 - \sum_{i=1}^n (H_i^{old} - D_i)^2}{2\sigma^2} = \exp - \frac{(\chi_{new}^2 - \chi_{old}^2)}{2}$$

step	a1	c1	sigma1	chisq
100	9.806162E-01	5.169688E+01	5.290837E+00	4.267798E+03
200	9.999395E-01	5.180603E+01	5.411015E+00	4.265669E+03
300	1.015084E+00	5.172740E+01	5.411015E+00	4.267409E+03
400	9.944127E-01	5.171415E+01	5.395535E+00	4.265770E+03
500	1.005000E+00	5.180460E+01	5.440930E+00	4.266081E+03

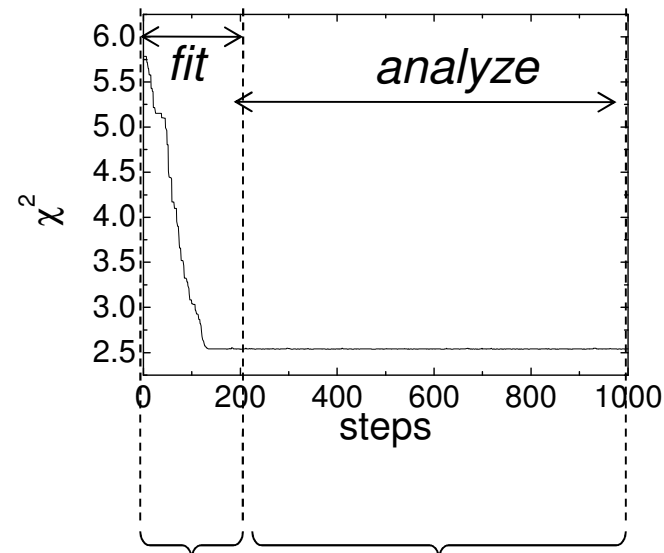
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It “simply” generates a Markov Chain with parameters and χ^2 supported by the data

- The ubiquitous χ^2
- **Advantages of Bayesian analysis**
- Some examples
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Three main advantages

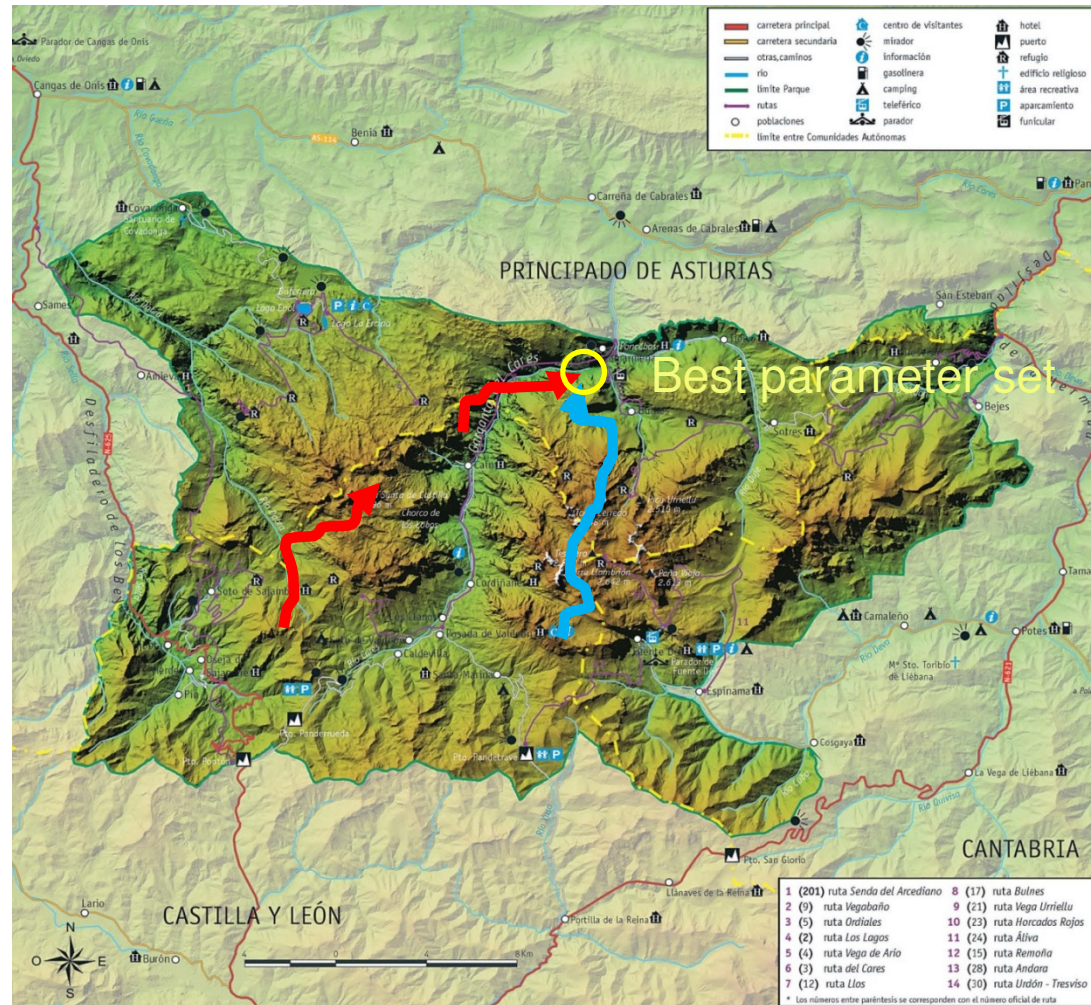


➤ The fitting process

➤ Parameter estimation
➤ Model selection

Fitting in $\chi^2\{P_1\}$ landscape

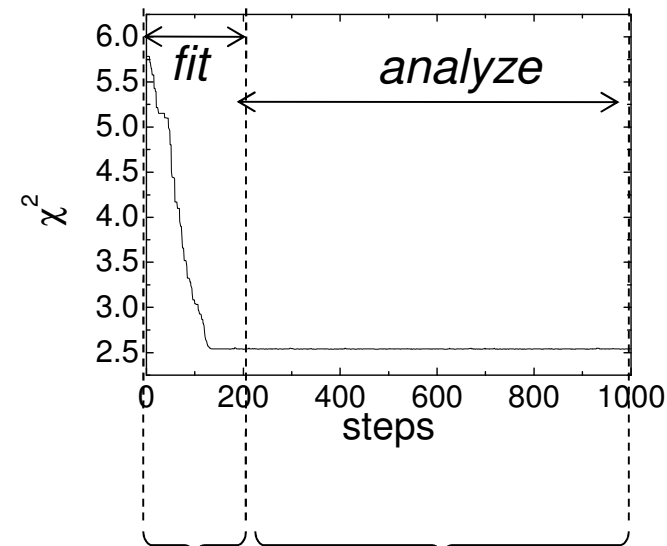
Classical fitting



Bayesian fitting

It does not get stuck!!!

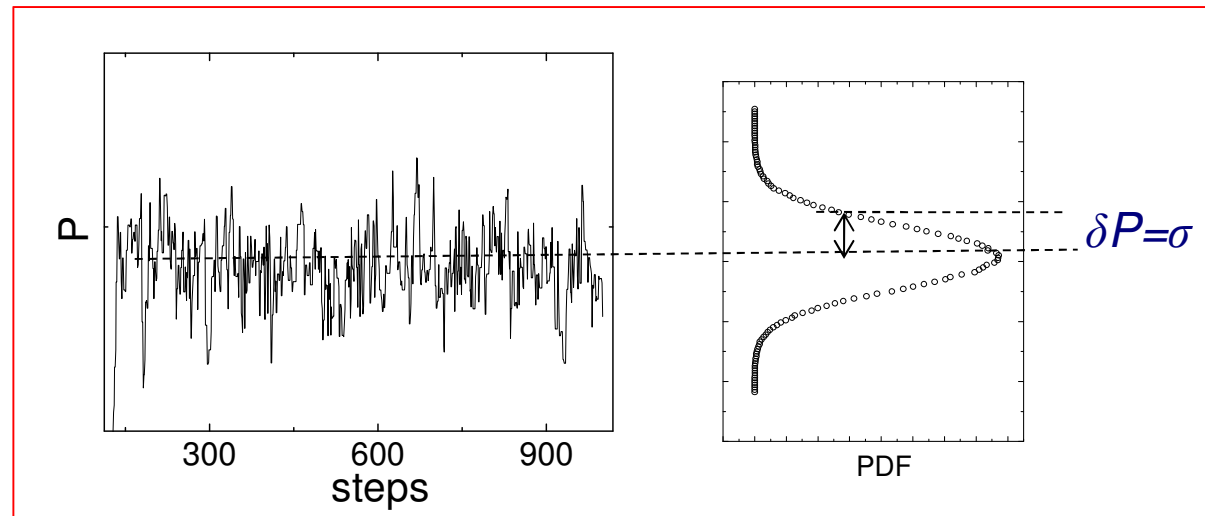
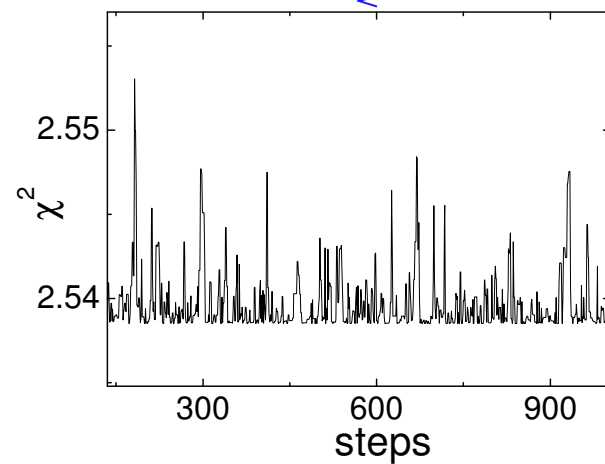
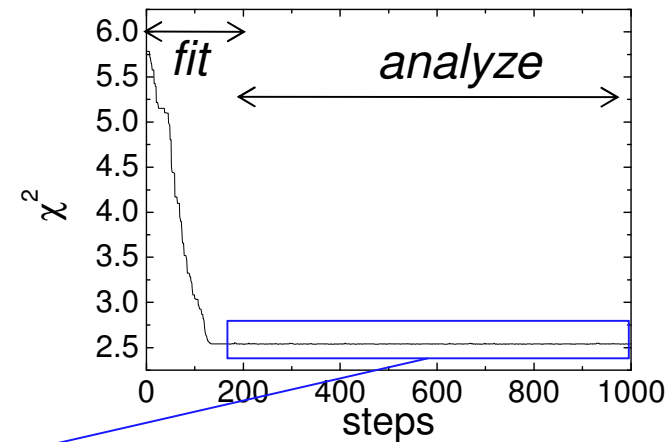
Three main advantages



➤ The fitting process

➤ Parameter estimation
➤ Model selection

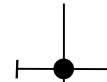
Parameter estimation



*Parameter estimation:
Parameters are obtained as PDF's not as $P \pm \delta P$*

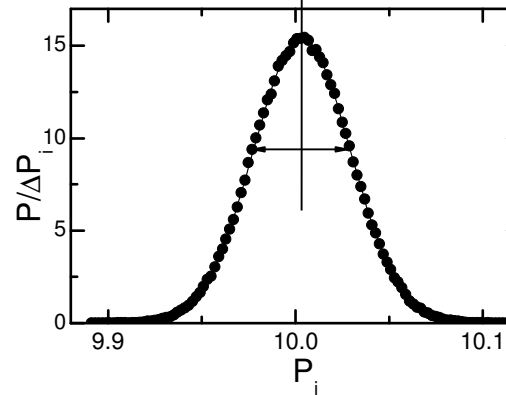
Parameter determination

“Classic”
(frequentist)



$$P \pm \delta P$$

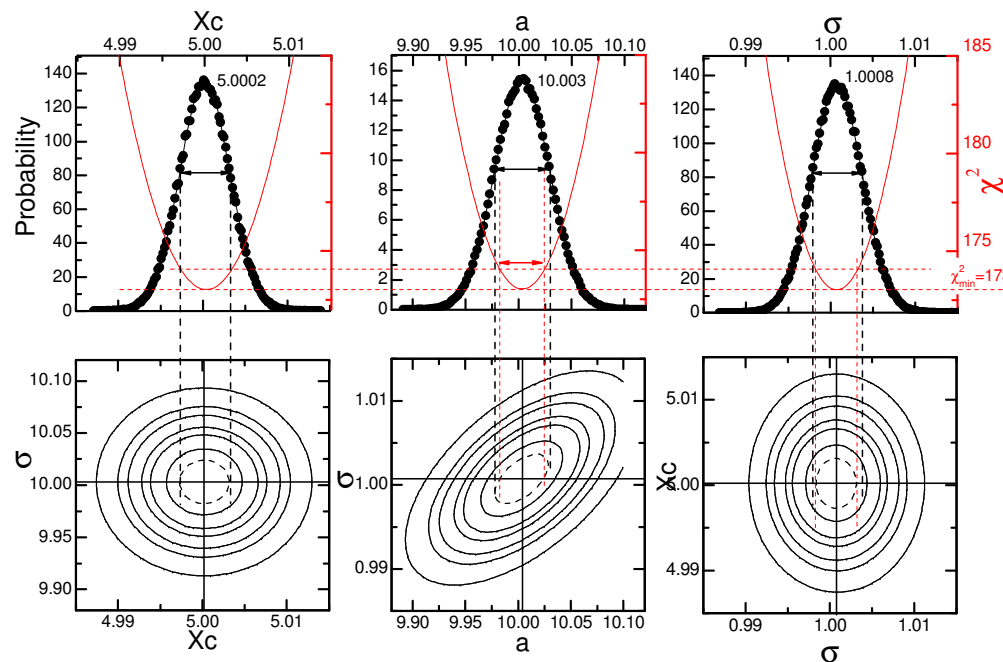
Bayesian



*Probability Density Function
(PDF)*

Parameter determination

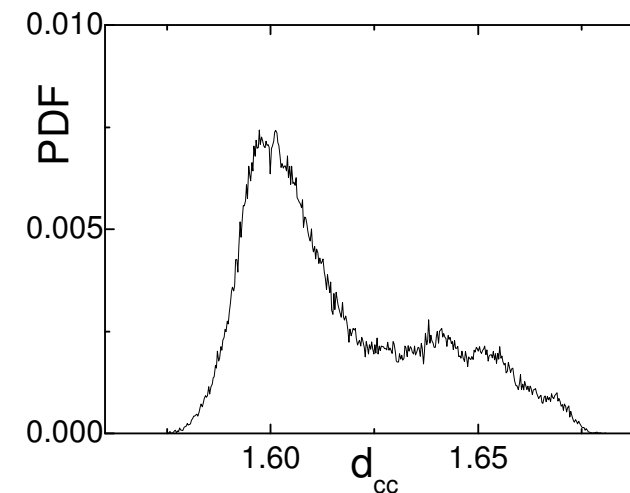
Correlation between parameters
are automatically had into account



Fit of a "simple" Gaussian

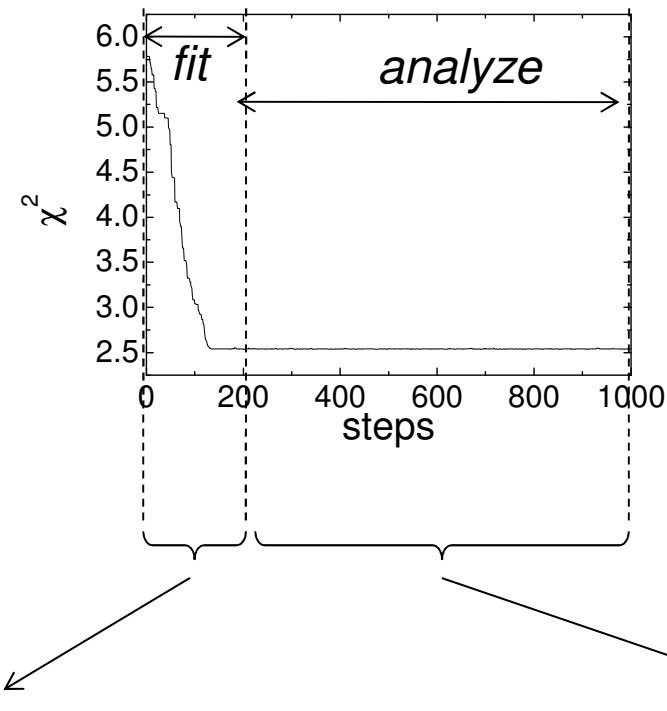
$$f(x) = a \cdot \exp - \frac{(x - x_c)^2}{2\sigma^2}$$

No supposition on the
minimum geometry is made



It might be as terrible as this one...

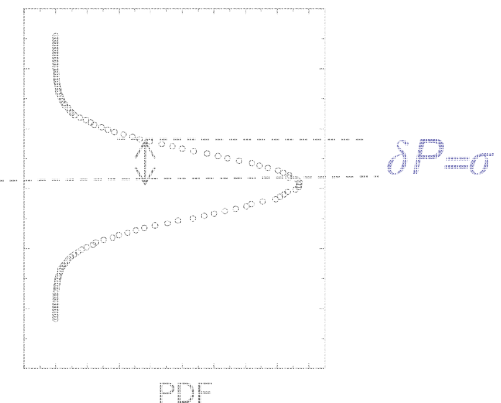
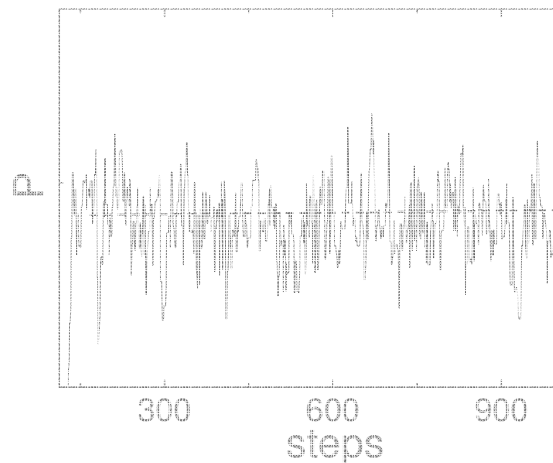
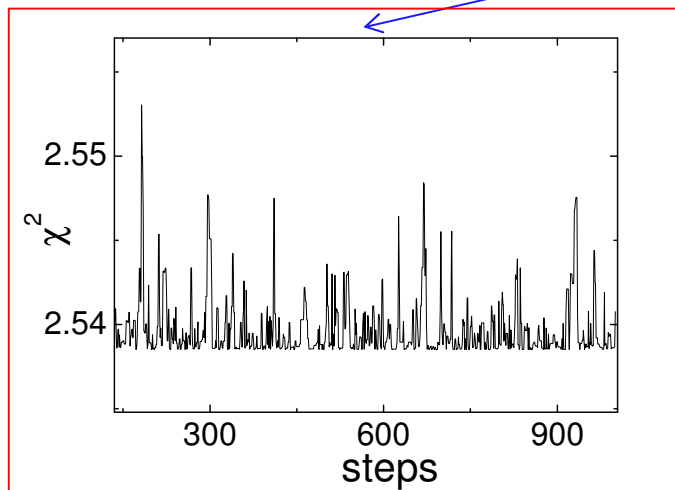
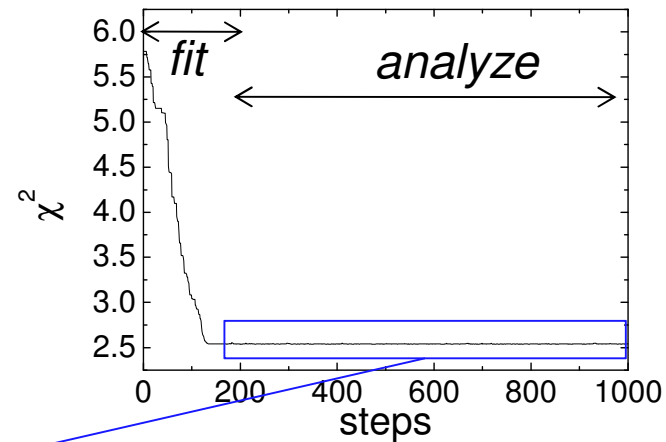
Three main advantages



➤ The fitting process

➤ Parameter estimation

➤ Model selection



*Model selection:
all possible combinations of parameters are investigated
and their χ^2 calculated*

Model Selection

Usual methods

- The "guide to the eye" method
- The reduced χ^2 method

$$\chi_{red}^2 = \frac{\chi^2}{n - m}$$

n : is the number of points

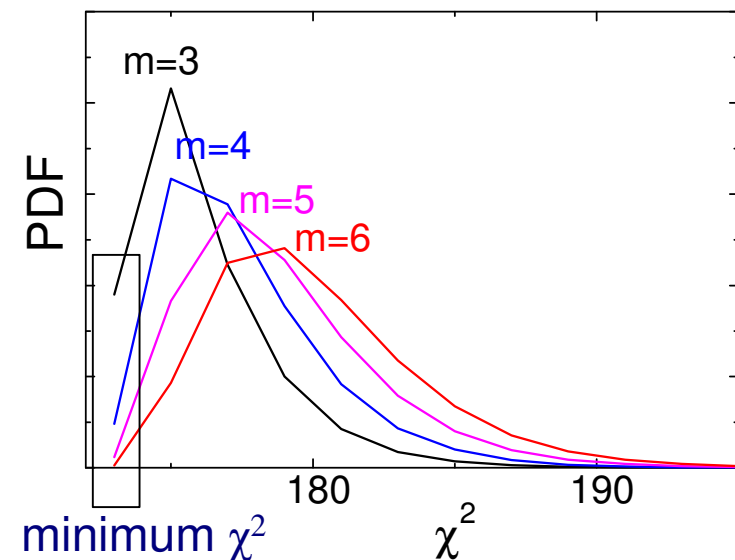
m : is the number of parameters

This works only if:

- ✓ There is no correlation between parameters
- ✓ The PDF in **all** parameters is gaussian
- ✓ The minimum is not multimodal

Bayesian method

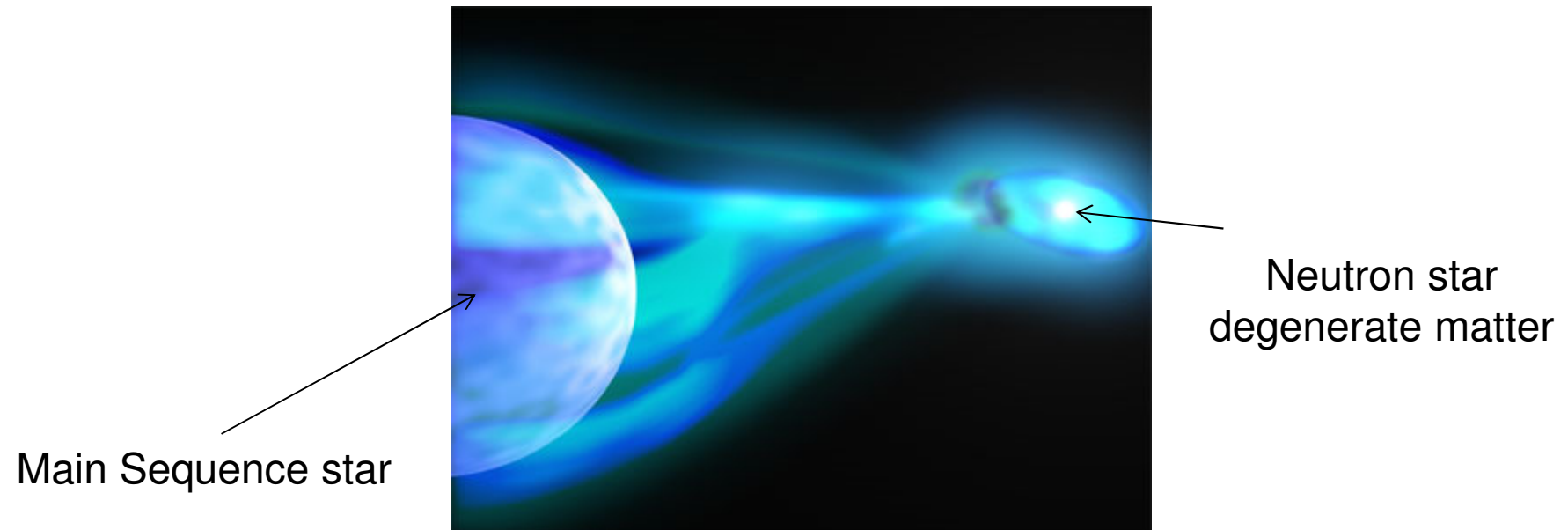
- Directly compares the PDF related to χ^2



an increasing number of parameters
broadens the χ^2 PDF

- The ubiquitous χ^2
- Advantages of Bayesian analysis
- **Some examples**
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A far away example...

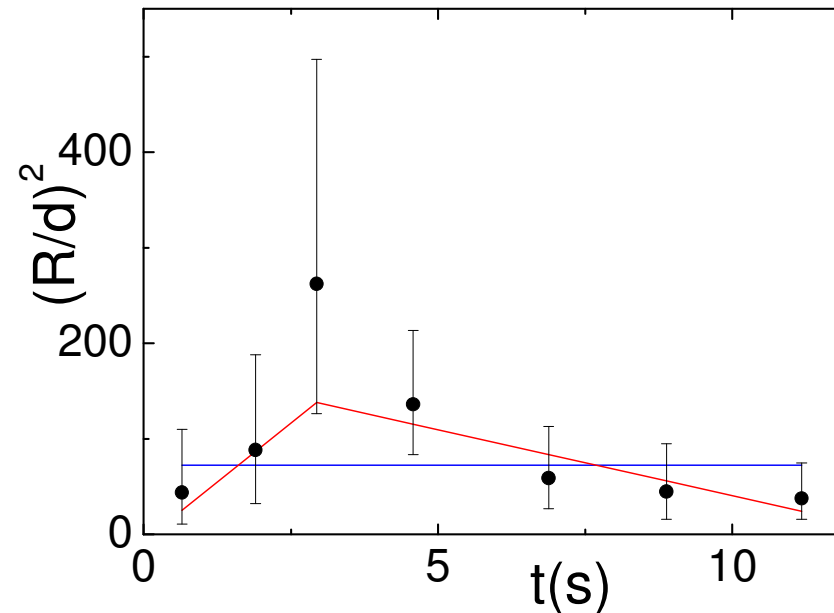


Two models: does the neutron star envelop expand?

YES: radiation pressure=" escape velocity
you know the mass! (you have a paper!)

NO: you don't know the mass! (*Game over!*)

A far away example...



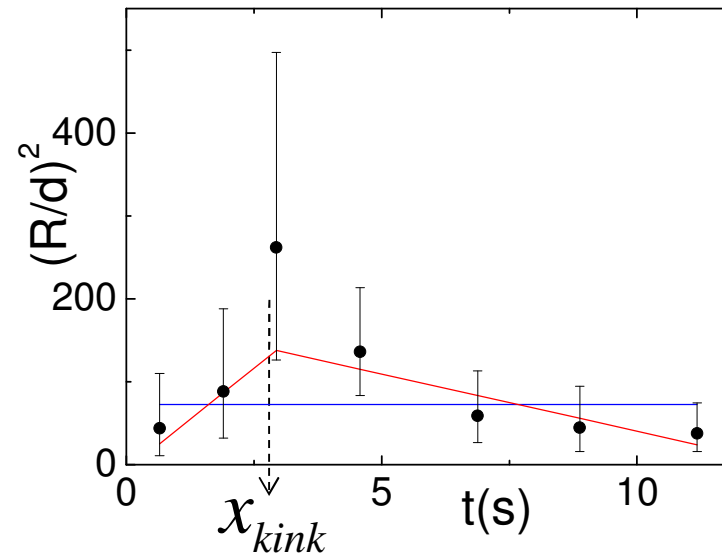
Two models: does the neutron star envelop expand?

YES: Eddington luminosity: radiation pressure=" escape velocity
you know the mass! (you have a paper!)

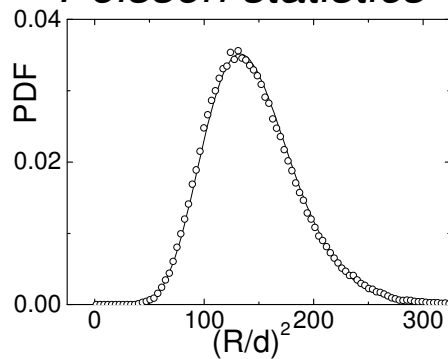
NO: you don't know the mass! (*Game over!*)

A Bayesian Dream...

(Levenberg-Marquart does not work)



Poisson statistics



$$\chi^2 = -2 \sum_{i=1}^n \left[\left(D_i \ln \frac{H_i}{D_i} - H_i + D_i \right) \right]$$

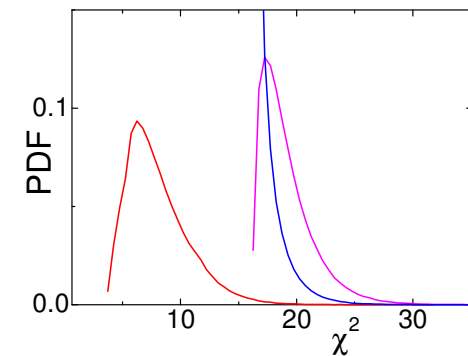
Non differentiable model

$$y_1 = a_1 + b_1 x$$

$$a_2 = a_1 + (m_1 - m_2) x_{kink}$$

$$y_2 = a_2 + b_2 x$$

Model Selection

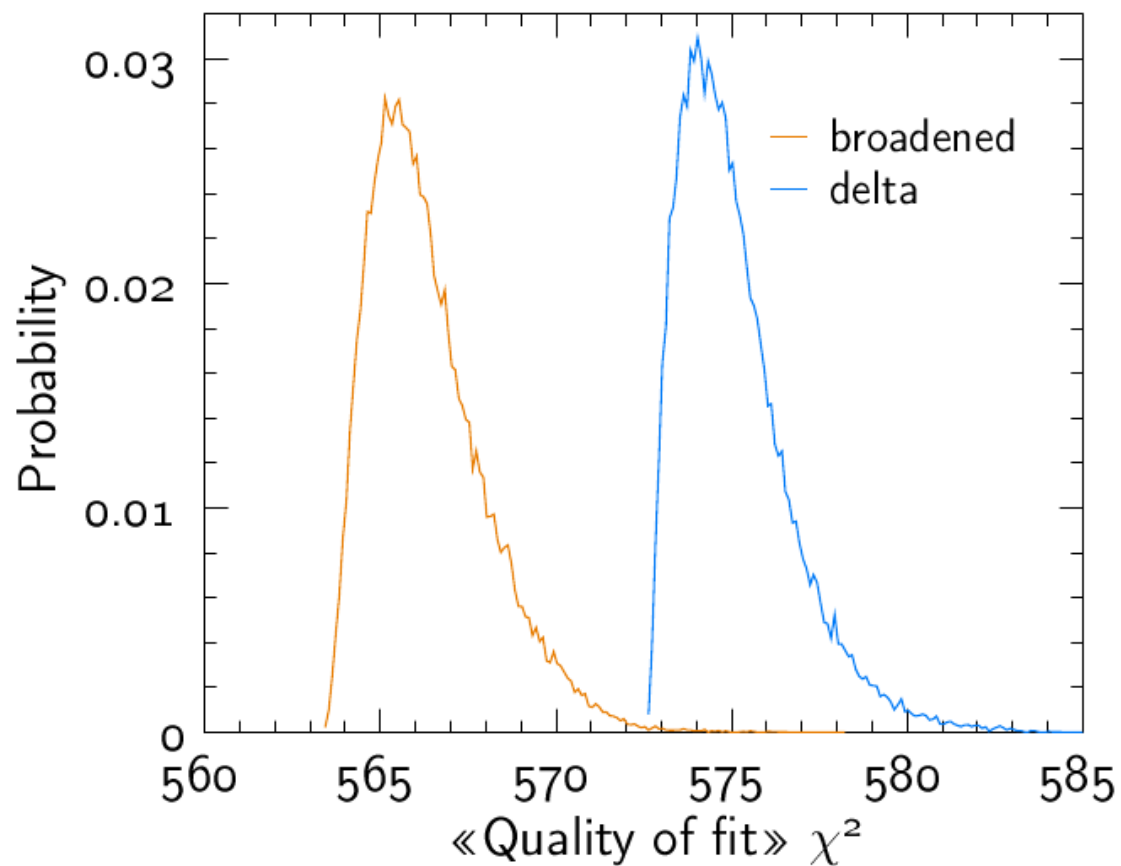
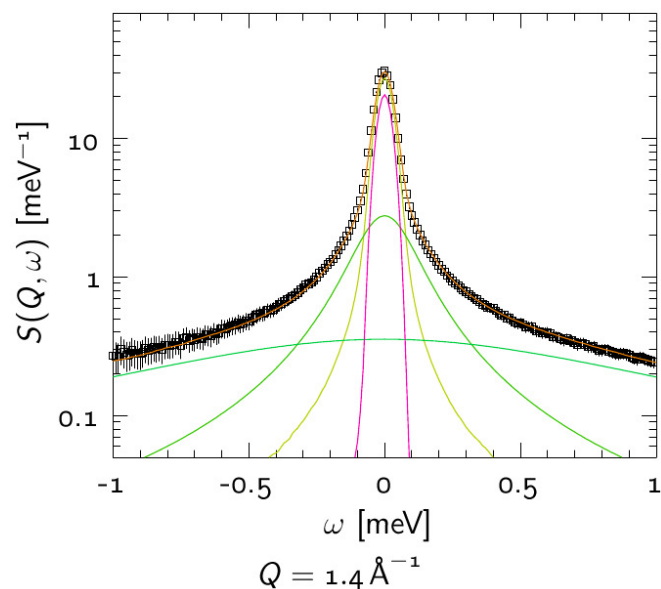


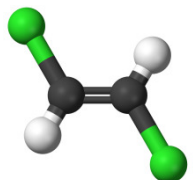
YES!!
you know the mass

- The ubiquitous χ^2
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Motion of phospholipids in the membrane

Is there a broadening? Delta model versus Broadened model

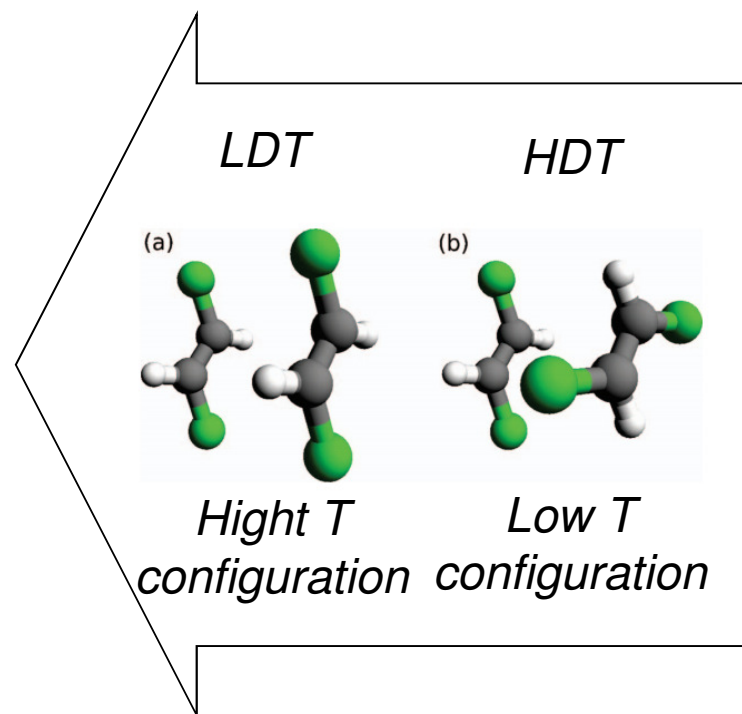
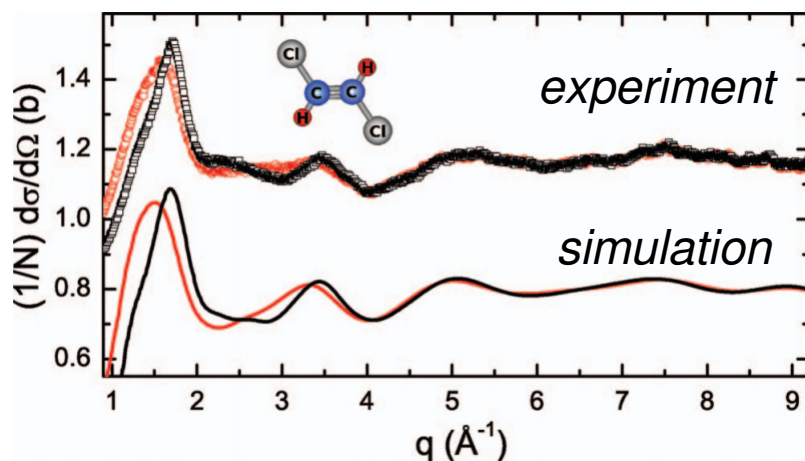




Is there a liquid-liquid phase transition in trans-dichloroethylene?

- Reported changes in density
- Reported changes in dynamics (NMR, infrared spc, viscosity)

There is a change in the structure



Is there any change in the dynamics?

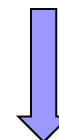
M. Rovira-Esteva et al. *J. Chem. Phys.* 136, 124514 (2012)

M. Rovira-Esteva et al. *Phys. Rev. B* 81(9) 092202 (2010)

Is there any change in the dynamics?

Dynamics transition in trans-dicloroethylene

We analyze TOF spectra with a diffusion+rotation model

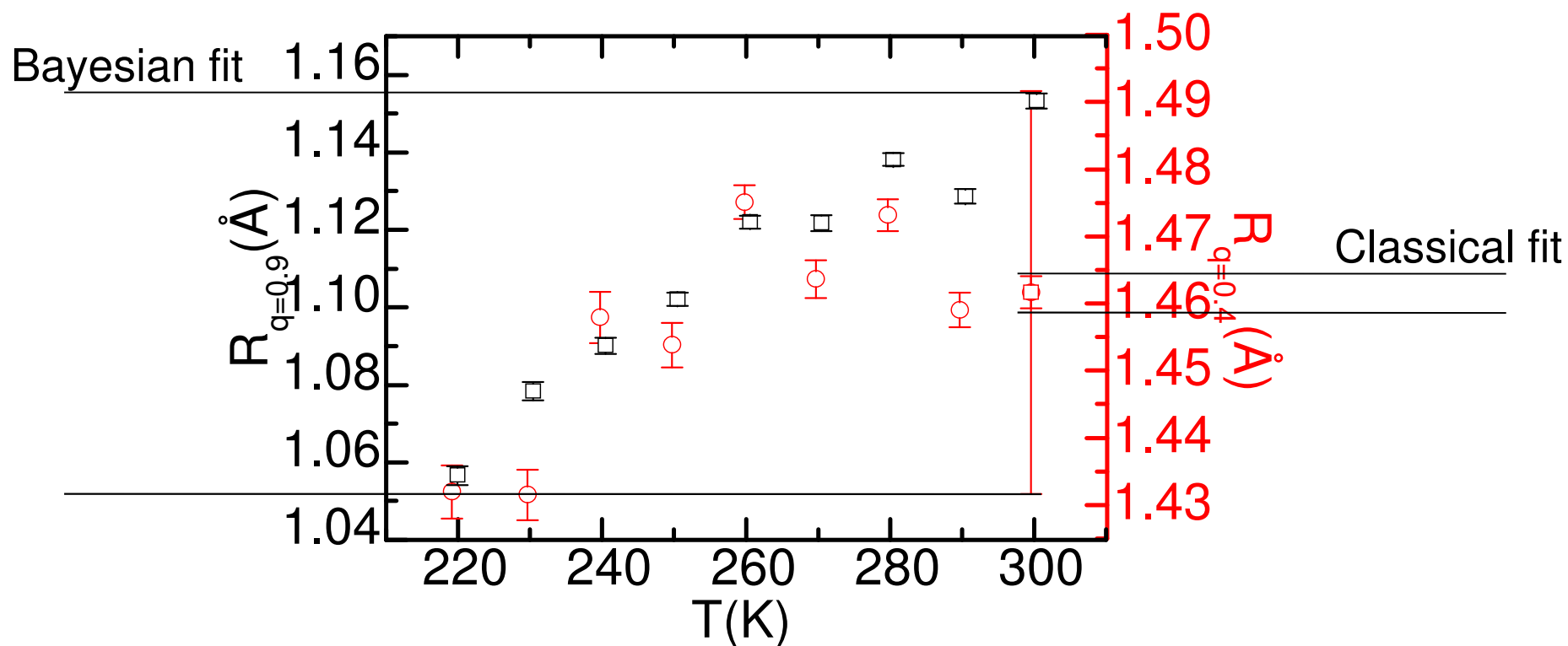


$$S_{rot}(q, \omega) = A_0(qR) \cdot \delta(\omega) + \sum_l A_l(qR) \cdot L_l(\omega, \gamma = l(l+1)D_r)$$

$$S(q, \omega) = S_{diff}(q, \omega) \otimes S_{rot}(q, \omega) \otimes R(q, \omega)$$

$$S_{diff}(q, \omega) = L(\omega, \gamma = Dq^2)$$

We analyze just one q value

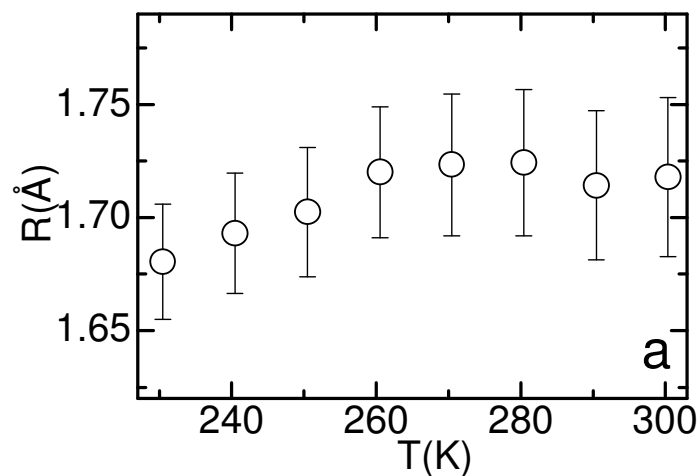


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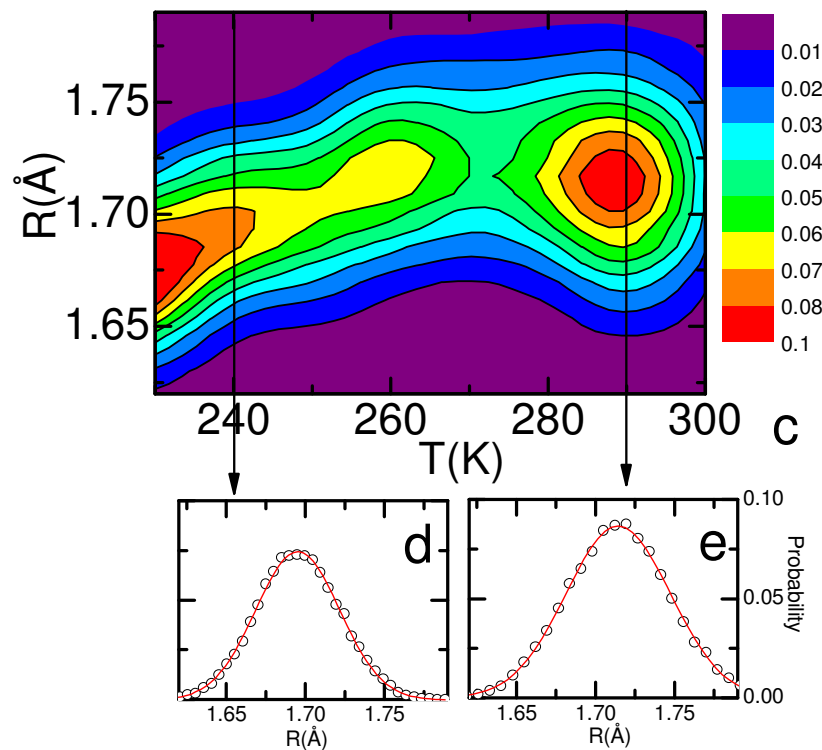
M. Rovira-Esteva et al. Phys. Rev. B 81(9) 092202 (2010)

All q values at once!!
i.e. the whole scattering law

Classical representation

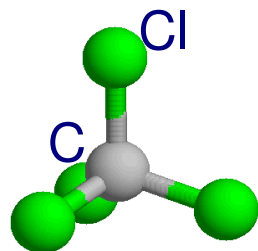


PDF representation

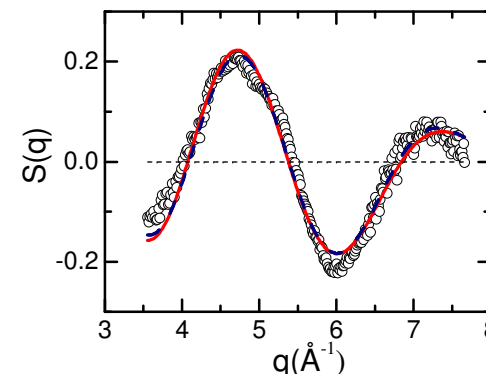


- The ubiquitous χ^2
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A simple case: CCl_4

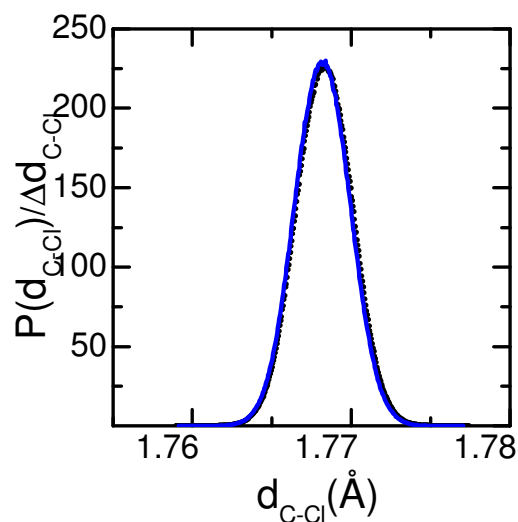


Molecular structure
fit of the high q region of $s(q)$

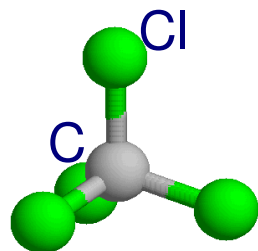


$$S(q) = h \cdot \sum_{i,j}^m b_i b_j \cdot \frac{\sin(qr_{ij})}{qr_{ij}} \cdot e^{-\frac{u_{ij}^2 q^2}{2}}$$

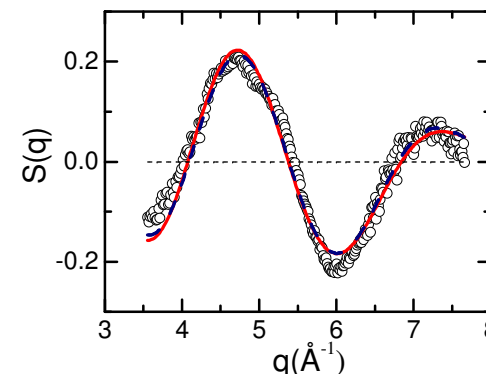
d_{CCl} distance is well defined



A simple case: CCl_4

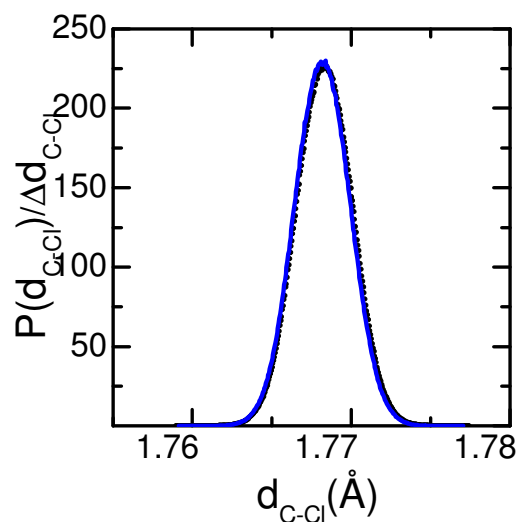


Molecular structure
fit of the high q region of $s(q)$

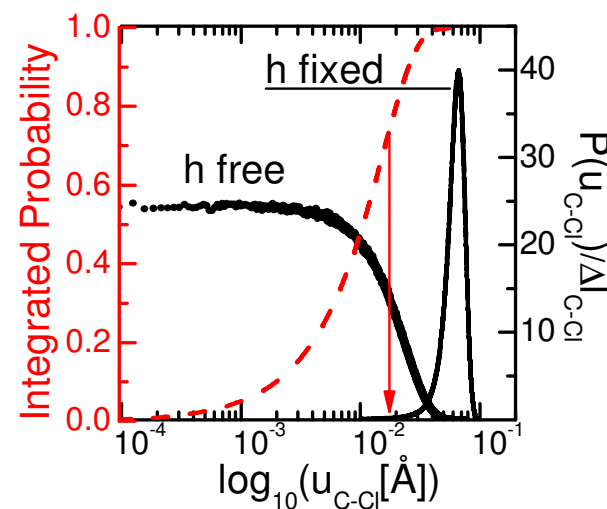


$$S(q) = h \cdot \sum_{i,j}^m b_i b_j \cdot \frac{\sin(qr_{ij})}{qr_{ij}} \cdot e^{-u_{ij}^2 q^2 / 2}$$

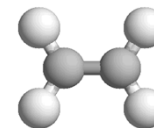
$d_{\text{C-Cl}}$ distance is well defined



$U_{\text{C-Cl}}$ depends on the scaling factor h



A more complicated test case: C_2D_4



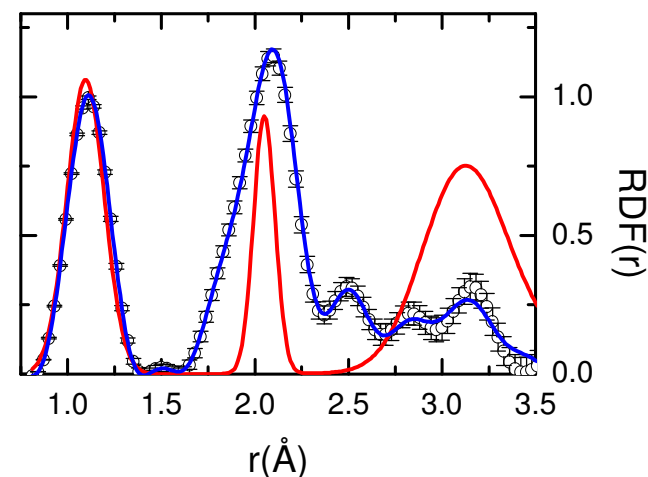
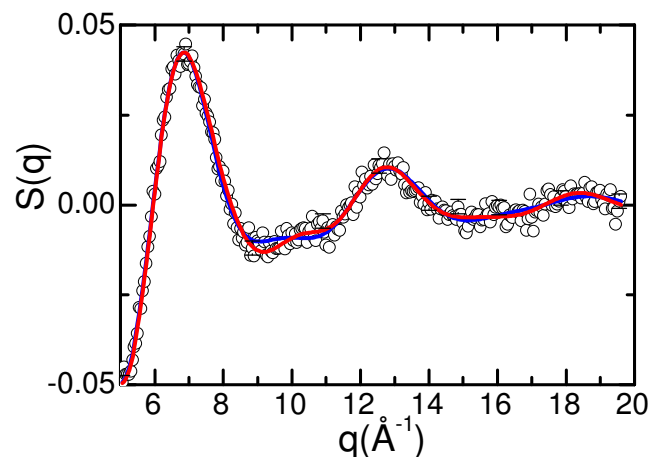
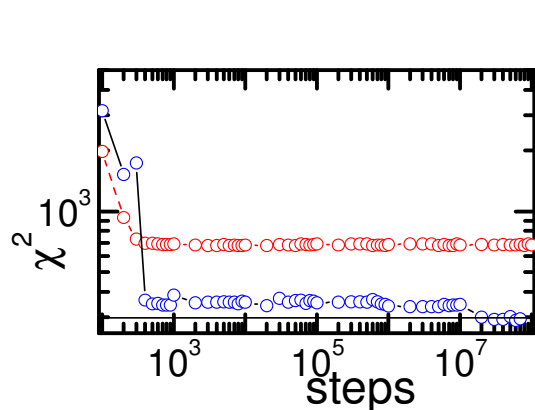
fit of the high q region of $s(q)$

fit of the small r region of $G(r)$

$$S(q) = h \cdot \sum_{i,j}^m b_i b_j \cdot \frac{\sin(qr_{ij})}{qr_{ij}} \cdot e^{-\frac{u_{ij}^2 q^2}{2}} \longleftrightarrow G(r) \approx h \cdot \sum_{i,j}^m b_i b_j \cdot \exp\left(-\frac{(r-r_{ij})^2}{2u_{ij}^2}\right)$$

Fit using only $s(q)$

Fit using both $s(q)$ and $g(r)$

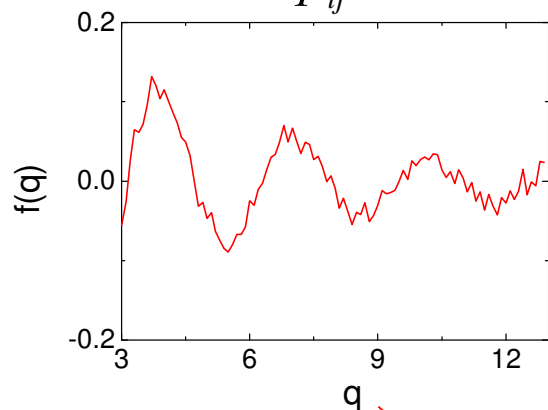


Why is it better fitting both at the same time?

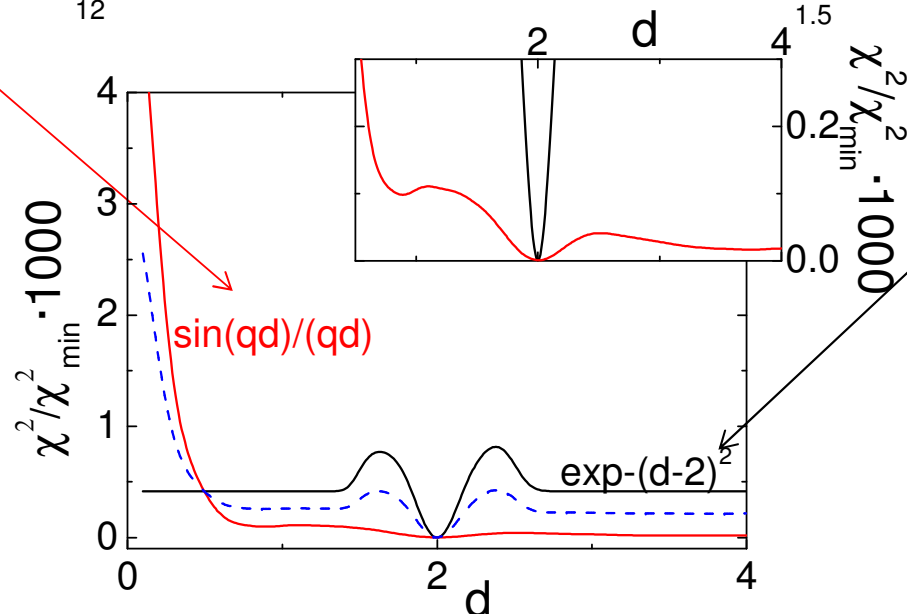
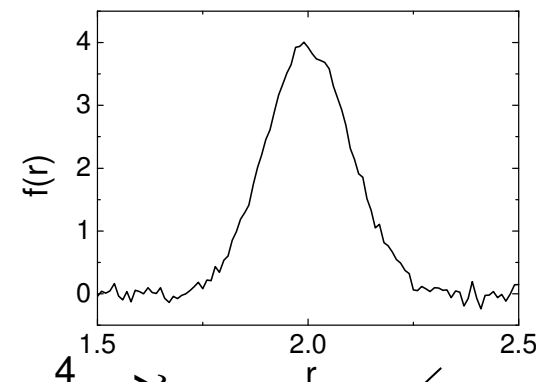
Why is it better fitting both at the same time?

The simplest case (with $U_{ij}=0.1$, $r_{ij}=2$):

$$S(q) = \frac{\sin(qr_{ij})}{qr_{ij}} \cdot e^{-\frac{u_{ij}^2 q^2}{2}}$$



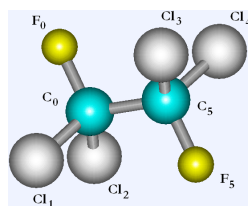
$$RDF(r) = \exp\left(-\frac{(r-r_{ij})^2}{2u_{ij}^2}\right)$$



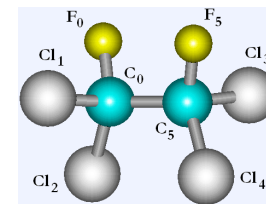
χ^2 landscapes are quite different!!

A really complicated case: $C_2Cl_4F_2$

It has two conformers

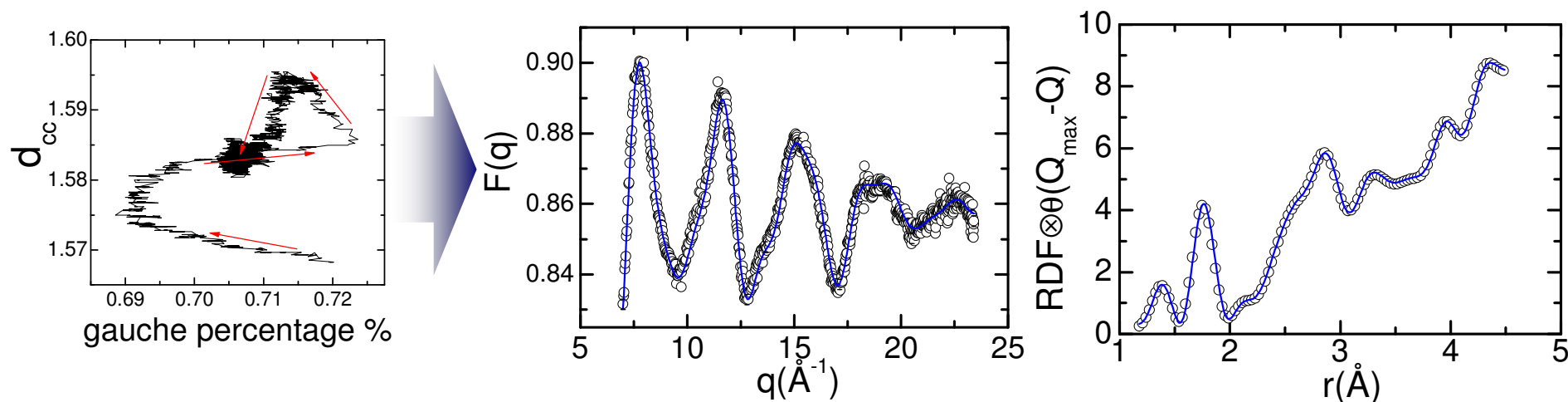


trans



gauche

We have to fit **37 parameters**, and they are not independent:
for example a change in d_{cc} implies changes in the whole molecule



We simply let the program to wander through the parameter space...

- The ubiquitous χ^2
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 - ✓ Analysis of QENS data
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- Robustness and simulated annealing
- Summary and conclusions

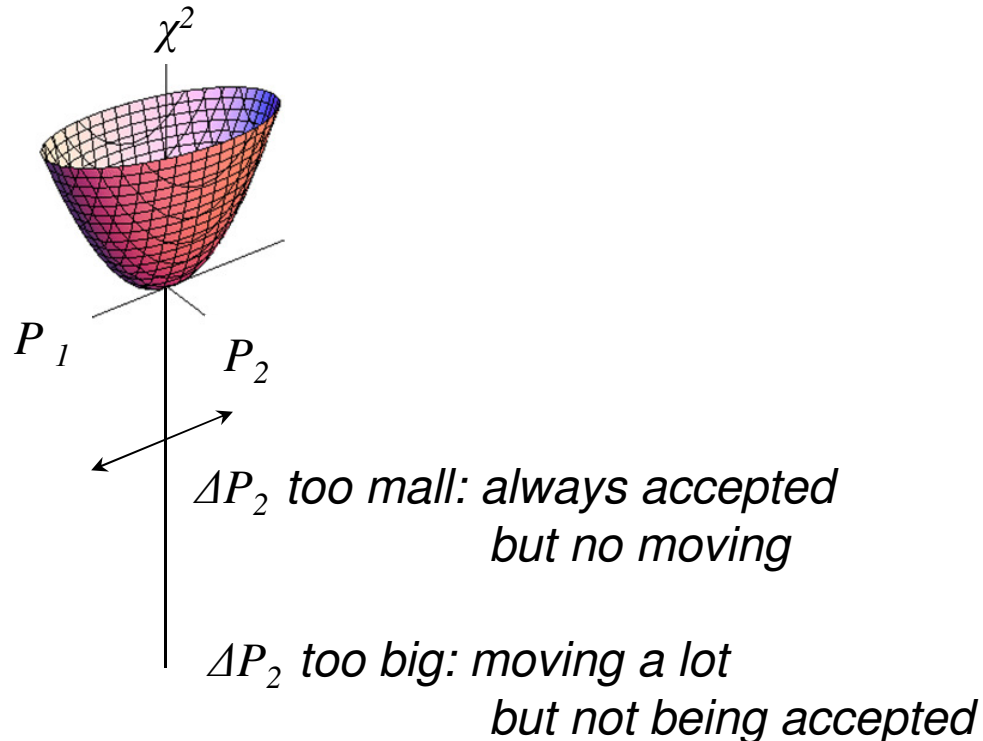
Adaptive Markov Chain Montecarlo

$$\Delta P_{new} = \Delta P_{old} \frac{R_i}{R_{i,desired}}$$

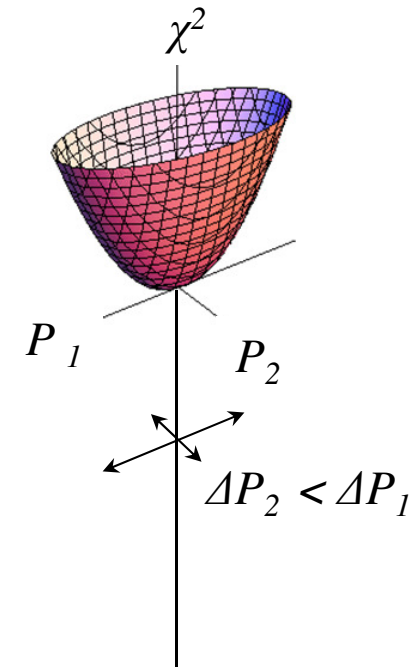
step	a1	c1	sigma1	chisq
100	9.806162E-01	5.169688E+01	5.290837E+00	4.267798E+03
200	9.999395E-01	5.180603E+01	5.411015E+00	4.265669E+03

ΔP is optimized!

To optimize de number of accepted moves

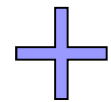


To explore the χ^2 landscape in all directions



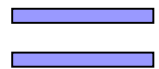
Adaptive Markov Chain Montecarlo

$$\Delta P_{new} = \Delta P_{old} \frac{R_i}{R_{i,desired}}$$

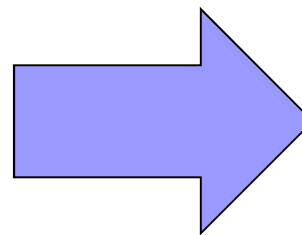
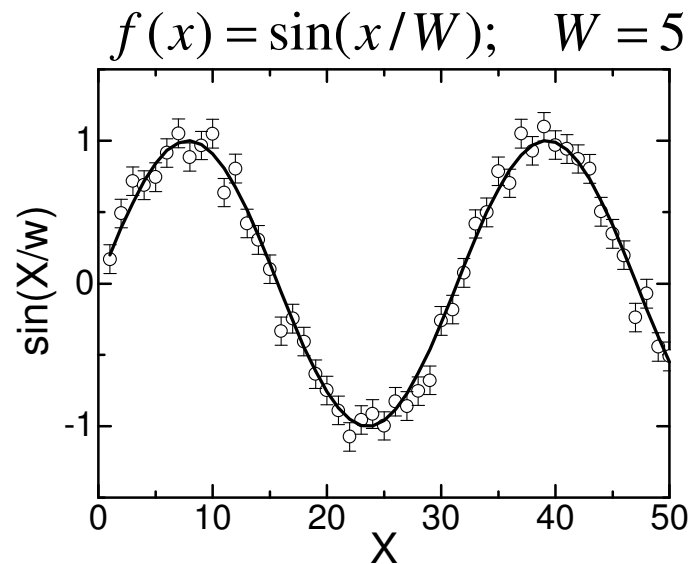


Simulated annealing...

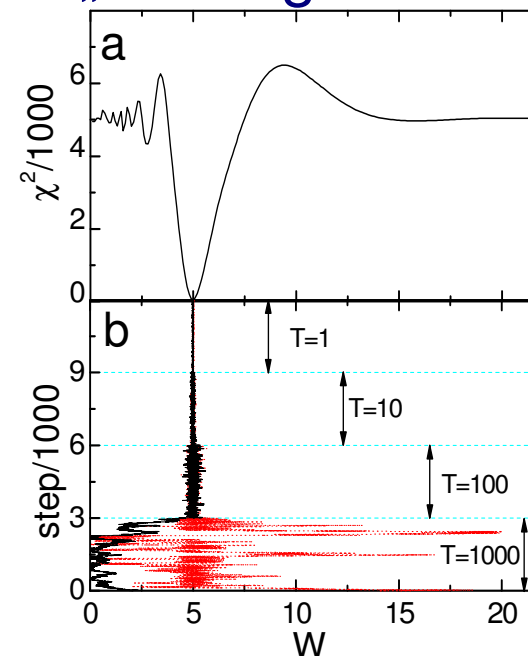
$$\frac{P(H_i\{P_l^{new}\} | D_i)}{P(H_i\{P_l^{old}\} | D_i)} = \exp - \frac{(\chi_{new}^2 - \chi_{old}^2)}{2T}$$

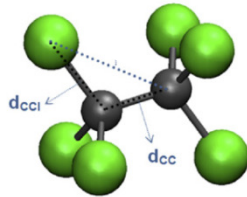
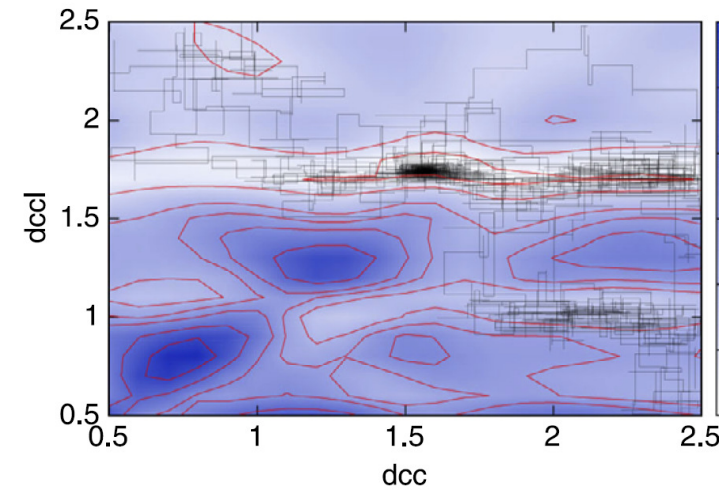


a test case

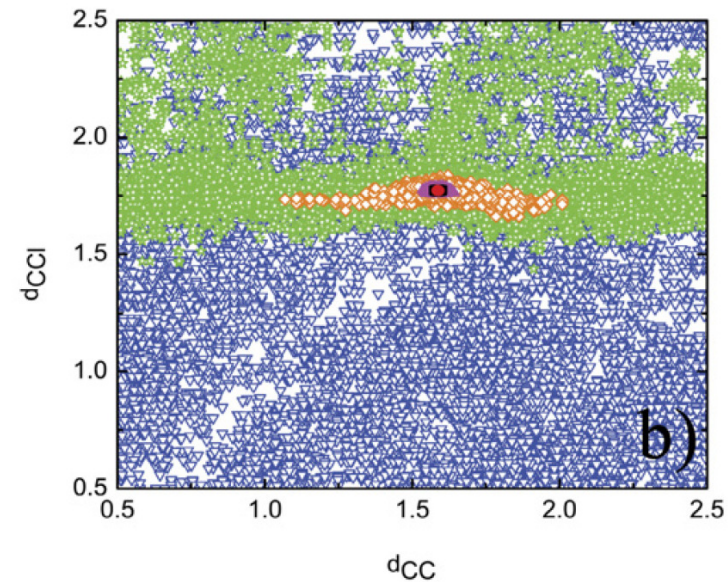
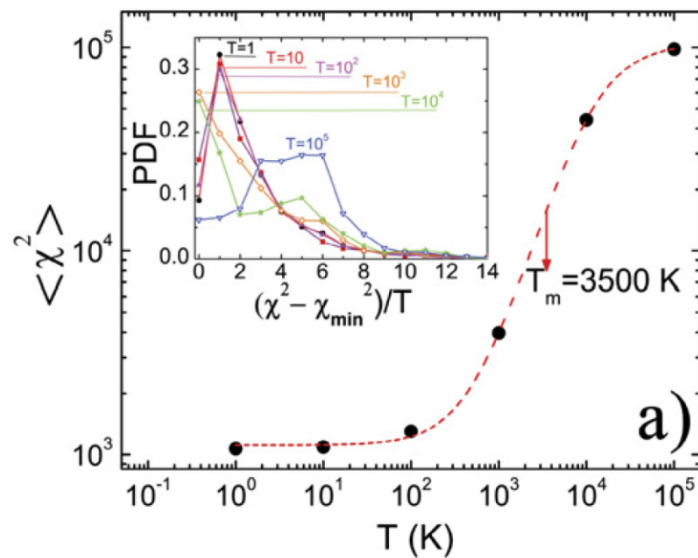


„Cooling the fit“



The structure of C_2Cl_6  $\chi^2(d_{cc}, d_{cci})$ landscape

„Melting“ the fit



Just a word about priors:

From some paper (that I trust) I know $P \pm \delta P$

Can I include this information in the fit?

$$P(H | D) = \frac{P(D | H) \cdot P(H)}{P(D)}$$

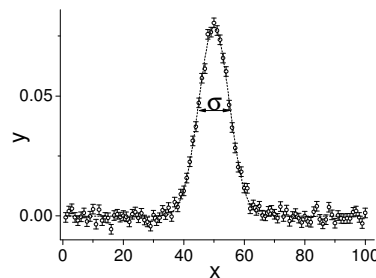
$$P(H | D) \propto \exp - \frac{\chi^2}{2} \cdot \exp - \frac{(P - P_i)^2}{2\delta P^2}$$

$$\chi_{+prior}^2 = \sum_{i=1}^n \frac{(H_i - D_i)^2}{\sigma_i^2} + \sum_{i=1}^{n_p} \frac{(P - P_i)^2}{\delta P^2}$$

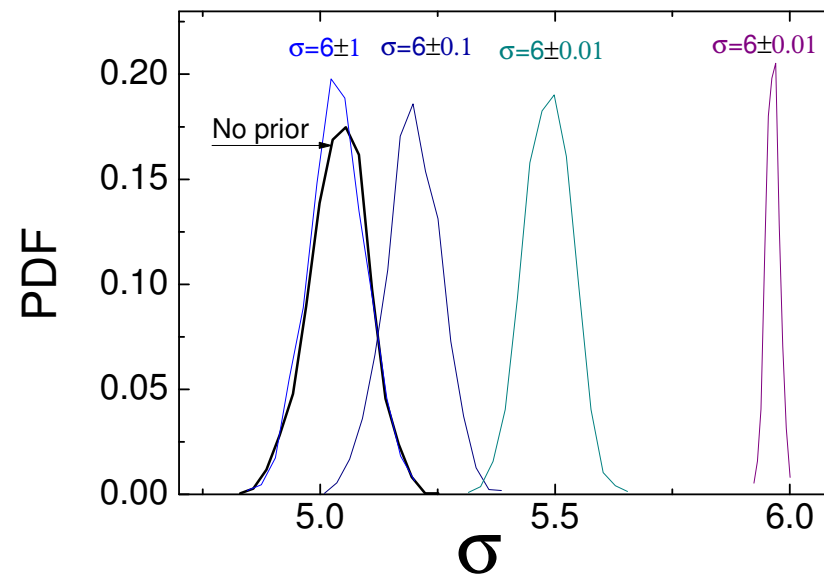
Just a word about priors:

From some paper (that I trust) I know $P \pm \delta P$

Can I include this information in the fit?



Fitting a gaussian with $\sigma=5$, and $P=6 \pm \delta P$



- The ubiquitous χ^2
- Advantages of Bayesian analysis
- Some examples
 - ✓ Quantitative guide to the eye
 - ✓ Analysis of QENS data
 - ✓ Intramolecular structure determination
- Robustness and simulated annealing
- **Summary and conclusions**



Classical fit

Bayesian “Fit”

Fitting process

Only downhill changes of χ^2 are allowed

Uphill changes of χ^2 are allowed

Parameter determination

Parameters and their error $P_i \pm \varepsilon_i$

Probability Density Function for P_i

Correlation between parameters
are not automatically taken into account

Correlation between parameters
are taken into account

Multiple minima are invisible

Multiple minima are visible

Model selection

A χ^2 PDF is assumed

No assumption is made on χ^2 PDF

When is it **not** worth to work with Bayesian analysis

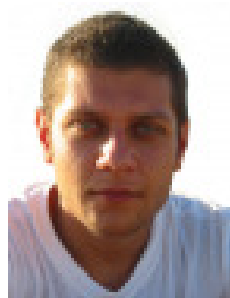
- When you have a simple function, with few parameters
- When parameters can be initialized close to the solution
- When model selection “done by eye” is evident (being m equal!!!)

When is it worth to work with Bayesian analysis

- When your function has too many parameters
- When fit gets stuck every now and then
- When model selection is not evident
- When you have different number of parameters for each model



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Andres Henao
(GCM / UPC)



Gabriel Cuello
(ILL / D4)

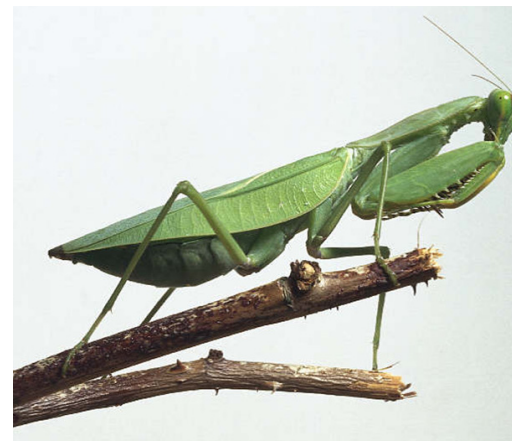
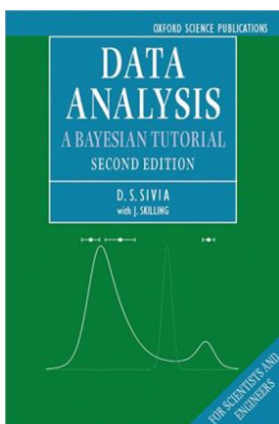


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(GAA-UPC)

Thank you for your attention



To read more about the examples

Astrophysics

G. Sala et al. *The Astrophysical Journal* (2012)

Diffraction

M. Rovira-Esteve et al. *Phys. Rev. B* 84, 064202 (2011)

QENS

M. Rovira-Esteve et al. *Phys. Rev. B* 81(9) 092202 (2010)

S. Busch et al. *J. Am. Chem. Soc.* 132(10) 3232 (2010)

Dielectric spectroscopy

J. C. Martínez et al. *J. Phys. Chem. B* 114 6099 (2010)

To read more about FABADA and download it

Aristizabal A. H. et al. *Journal of Physics: condensed matter. Special issue: Reverse Monte Carlo 2012*

L.C. Pardo et al. *Phys. Rev. E.* 84, 046711 (2011)

L.C Pardo et al. *J. Phys.: Conf. Ser.* 325, 012006 (2011)

L. C. Pardo et al. *arXiv:0907.3711v3 [physics.data-an]*

Download the program, and see these slides:

<http://gcm.upc.edu/members/luis-carlos/bayesiano>