The Flavor Structure of the Nucleon Sea
Institute for Nuclear Theory
October 2, 2017

# New approaches to global PDF analysis



with Nobuo Sato



Connecticut / JLab

#### **Outline**

- Motivation why the need for a new paradigm?
- Bayesian approach to fitting
  - → single-fit (Hessian) vs. Monte Carlo approaches
  - → shortcomings of Hessian (Gaussian) approach
- Incompatible data sets
  - "tolerance" factors
    (uncertainties should not depend on # of parameters!)
- Monte Carlo methods
  - → iterative MC, nested sampling, ...
- Generalization to non-Gaussian likelihoods
  - → disjoint probabilities, empirical Bayes, ...
- Outlook

- With limited number of observables and finite statistics, need a robust analysis framework to extract meaningful parton information from experiment
- Over the first  $\sim 2-3$  decades of global PDF analysis efforts,  $\chi^2$  minimization (single-fit) analysis (with Hessian error propagation) has generally been sufficient to map out global characteristics of partonic structure
  - $\rightarrow$  e.g. shapes of quark PDFs from DIS, where data are plentiful
- A major challenge has been to characterize PDF uncertainties
   in a statistically meaningful way in the presence of tensions among data sets

- Previous attempts sought to address tensions in data sets by introducing
  - → "tolerance" factors (artificially inflating PDF errors)
  - "neural net" parametrization (instead of polynomial parametrization), together with MC techniques
- However, to address the problem in a more statistically rigorous way, one requires going beyond the standard  $\chi^2$  minimization paradigm
  - → utilize modern techniques based on Bayesian statistics!

- In the near future, standard  $\chi^2$  minimization techniques will be unsuitable even in the absence of tensions e.g. for
  - → simultaneous analysis of collinear distributions (unpolarized & polarized PDFs, fragmentation functions)

→ "JAM17": Jake Ethier (Tuesday)

 $\rightarrow$  new types of observables — TMDs or GPDs — that will involve  $>\mathcal{O}(10^5)$  data points, with  $\mathcal{O}(10^3)$  parameters

■ Typically PDF parametrizations are nonlinear functions of the PDF parameters, e.g.

$$xf(x,\mu) = Nx^{\alpha}(1-x)^{\beta} P(x)$$

where P is a polynomial e.g.  $P(x) = 1 + \epsilon \sqrt{x} + \eta x$ , or Chebyshev, neural net, ...

- $\rightarrow$  have multiple local minima present in the  $\chi^2$  function
- Robust parameter estimation that thoroughly scans over a realistic parameter space, including multiple local minima, is only possible using MC methods!
- Need more reliable algorithms "PDFs beyond the LHC"!

Analysis of data requires estimating expectation values E and variances V of "observables"  $\mathcal O$  (= PDFs, FFs) which are functions of parameters  $\vec a$ 

$$E[\mathcal{O}] = \int d^n a \, \mathcal{P}(\vec{a}|\text{data}) \, \mathcal{O}(\vec{a})$$
$$V[\mathcal{O}] = \int d^n a \, \mathcal{P}(\vec{a}|\text{data}) \, \left[\mathcal{O}(\vec{a}) - E[\mathcal{O}]\right]^2$$

"Bayesian master formulas"

lacktriangle Using Bayes' theorem, probability distribution  ${\cal P}$  given by

$$\mathcal{P}(\vec{a}|\mathrm{data}) = \frac{1}{Z} \mathcal{L}(\mathrm{data}|\vec{a}) \pi(\vec{a})$$

in terms of the <u>likelihood function</u>  $\mathcal{L}$ 

#### Likelihood function

$$\mathcal{L}(\text{data}|\vec{a}) = \exp\left(-\frac{1}{2}\chi^2(\vec{a})\right)$$

is a Gaussian form in the data, with  $\chi^2$  function

$$\chi^2(\vec{a}) = \sum_i \left( \frac{\text{data}_i - \text{theory}_i(\vec{a})}{\delta(\text{data})} \right)^2$$

with priors  $\pi(\vec{a})$  and "evidence" Z

$$Z = \int d^n a \, \mathcal{L}(\text{data}|\vec{a}) \, \pi(\vec{a})$$

 $\rightarrow$  Z tests if e.g. an n-parameter fit is statistically different from (n+1)-parameter fit

■ Two methods generally used for computing Bayesian master formulas:

 $\frac{\text{Maximum Likelihood}}{(\chi^2 \text{ minimization})}$ 

**Monte Carlo** 

■ Two methods generally used for computing Bayesian master formulas:

# Maximum Likelihood ( $\chi^2$ minimization)

 $\longrightarrow$  maximize probability distribution  ${\cal P}$  by minimizing  $\chi^2$  for a set of best-fit parameters  $\vec{a}_0$ 

$$E\left[\vec{a}\right] = \vec{a}_0$$

■ Two methods generally used for computing Bayesian master formulas:

# Maximum Likelihood ( $\chi^2$ minimization)

 $\longrightarrow$  maximize probability distribution  ${\cal P}$  by minimizing  $\chi^2$  for a set of best-fit parameters  $\vec{a}_0$ 

$$E\left[\vec{a}\right] = \vec{a}_0$$

 $\longrightarrow$  if  ${\mathcal O}$  is  $\approx$  linear in the parameters, and if probability is symmetric in all parameters

$$E\left[\mathcal{O}(\vec{a})\right] \approx \mathcal{O}(\vec{a}_0)$$

■ Two methods generally used for computing Bayesian master formulas:

#### Maximum Likelihood

 $(\chi^2 \text{ minimization})$ 

 $\rightarrow$  variance computed by expanding  $\mathcal{O}(\vec{a})$  about  $\vec{a}_0$  e.g. in 1 dimension have "master formula"

$$V[\mathcal{O}] \approx \frac{1}{4} \Big[ \mathcal{O}(a + \delta a) - \mathcal{O}(a - \delta a) \Big]^2$$

where

$$\delta a^2 = V[a]$$

■ Two methods generally used for computing Bayesian master formulas:

# $\frac{\text{Maximum Likelihood}}{(\chi^2 \text{ minimization})}$

→ generalization to multiple dimensions via Hessian approach:

find set of (orthogonal) contours in parameter space around  $\vec{a}_0$  such that  $\mathcal{L}$  along each contour is parametrized by statistically independent parameters — directions of contours given by eigenvectors  $\hat{e}_k$  of Hessian matrix H, with elements

$$H_{ij} = \frac{1}{2} \left. \frac{\partial^2 \chi^2(\vec{a})}{\partial a_i \partial a_j} \right|_{\vec{a} = \vec{a}_0}$$

and contours parametrized as  $\Delta a^{(k)}=a^{(k)}-a_0=t_k\frac{\hat{e}_k}{\sqrt{v_k}}$  , with  $v_k$  eigenvectors of H

Two methods generally used for computing Bayesian master formulas:

# $\frac{\text{Maximum Likelihood}}{(\chi^2 \text{ minimization})}$

ightharpoonup basic assumption:  $\mathcal P$  factorizes along each eigendirection

$$\mathcal{P}(\Delta a) \approx \prod_{k} \mathcal{P}_k(t_k)$$

where

$$\mathcal{P}_k(t_k) = \mathcal{N}_k \exp\left[-\frac{1}{2}\chi^2\left(a_0 + t_k \frac{\hat{e}_k}{\sqrt{v_k}}\right)\right]$$

<u>note</u>: in quadratic approximation for  $\chi^2$ , this becomes a normal distribution

■ Two methods generally used for computing Bayesian master formulas:

# Maximum Likelihood $(\chi^2 \text{ minimization})$

 $\rightarrow$  uncertainties on  $\mathcal{O}$  along each eigendirection (assuming linear approximation)

$$(\Delta \mathcal{O}_k)^2 \approx \frac{1}{4} \left[ \mathcal{O}\left(a_0 + T_k \frac{\hat{e}_k}{\sqrt{v_k}}\right) - \mathcal{O}\left(a_0 - T_k \frac{\hat{e}_k}{\sqrt{v_k}}\right) \right]^2$$

where  $T_k$  is finite step size in  $t_k$ , with total variance

$$V[\mathcal{O}] = \sum_{k} (\Delta \mathcal{O}_k)^2$$

■ Two methods generally used for computing Bayesian master formulas:

#### Monte Carlo

- $\longrightarrow$  in practice, generally one has  $E[\mathcal{O}(\vec{a})] \neq \mathcal{O}(E[\vec{a}])$  so the maximal likelihood method will sometimes fail
- $\rightarrow$  Monte Carlo approach samples parameter space and assigns weights  $w_k$  to each set of parameters  $a_k$
- -> expectation value and variance are then weighted averages

$$E[\mathcal{O}(\vec{a})] = \sum_{k} w_k \, \mathcal{O}(\vec{a}_k), \qquad V[\mathcal{O}(\vec{a})] = \sum_{k} w_k \, \left(\mathcal{O}(\vec{a}_k) - E[\mathcal{O}]\right)^2$$

■ Two methods generally used for computing Bayesian master formulas:

# $\frac{\text{Maximum Likelihood}}{(\chi^2 \text{ minimization})}$

- fast
- assumes Gaussianity
- no guarantee that global minimum has been found
- errors only characterize local geometry of  $\chi^2$  function

#### Monte Carlo

- Slow
- does not rely on Gaussian assumptions
- includes all possible solutions
- accurate

- Incompatible data sets can arise because of errors in determining central values, or underestimation of systematic experimental uncertainties
  - --> requires some sort of modification to standard statistics
- lacksquare Often one modifies the master formula by introducing a "tolerance" factor T

$$V[\mathcal{O}] \rightarrow T^2 V[\mathcal{O}]$$

e.g. for one dimension

$$V[\mathcal{O}] = \frac{T^2}{4} \left[ \mathcal{O}(a + \delta a) - \mathcal{O}(a - \delta a) \right]^2$$

effectively modifies the likelihood function

lacktriangle Simple example: consider observable m, and two measurements

$$(m_1, \delta m_1), (m_2, \delta m_2)$$

 $\rightarrow$  compute exactly the  $\chi^2$  function

$$\chi^2 = \left(\frac{m - m_1}{\delta m_1}\right)^2 + \left(\frac{m - m_2}{\delta m_2}\right)^2$$

and, from Bayesian master formula, the mean value

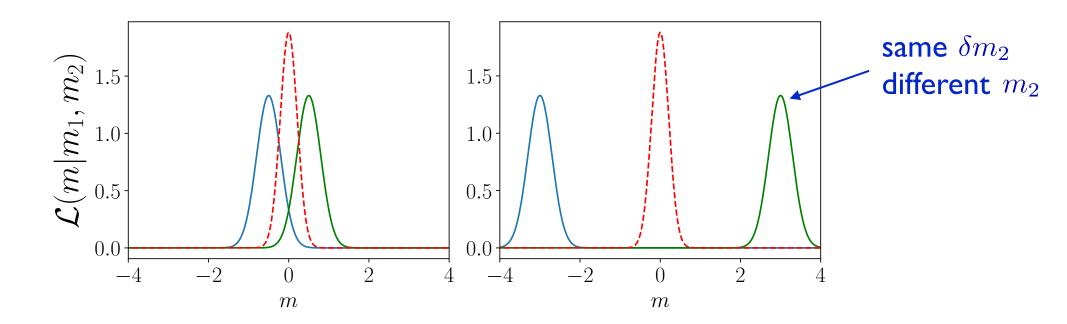
$$E[m] = \frac{m_1 \delta m_2^2 + m_2 \delta m_1^2}{\delta m_1^2 + \delta m_2^2}$$

and variance

$$V[m] = H^{-1} = \frac{\delta m_1^2 \, \delta m_2^2}{\delta m_1^2 + \delta m_2^2}$$

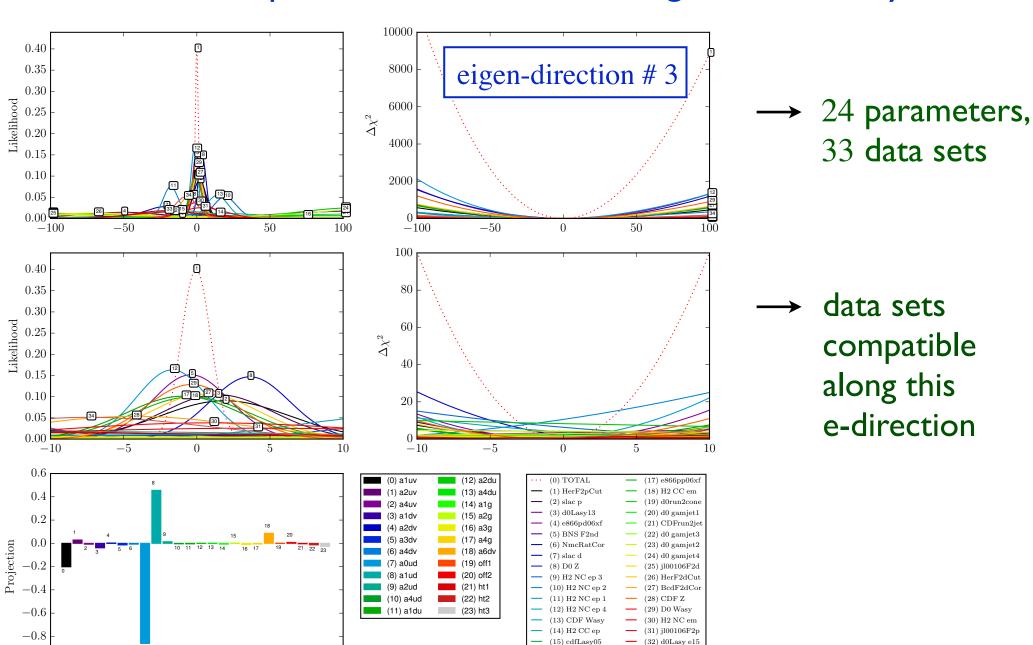
does not depend on  $m_1 - m_2$ !

■ Simple example: consider observable m, and two measurements  $(m_1, \delta m_1), (m_2, \delta m_2)$ 



- total uncertainty remains independent of degree of (in)compatibility of data
- Gaussian likelihood gives unrealistic representation of true uncertainty

■ Realistic example: recent CJ (CTEQ-JLab) global PDF analysis

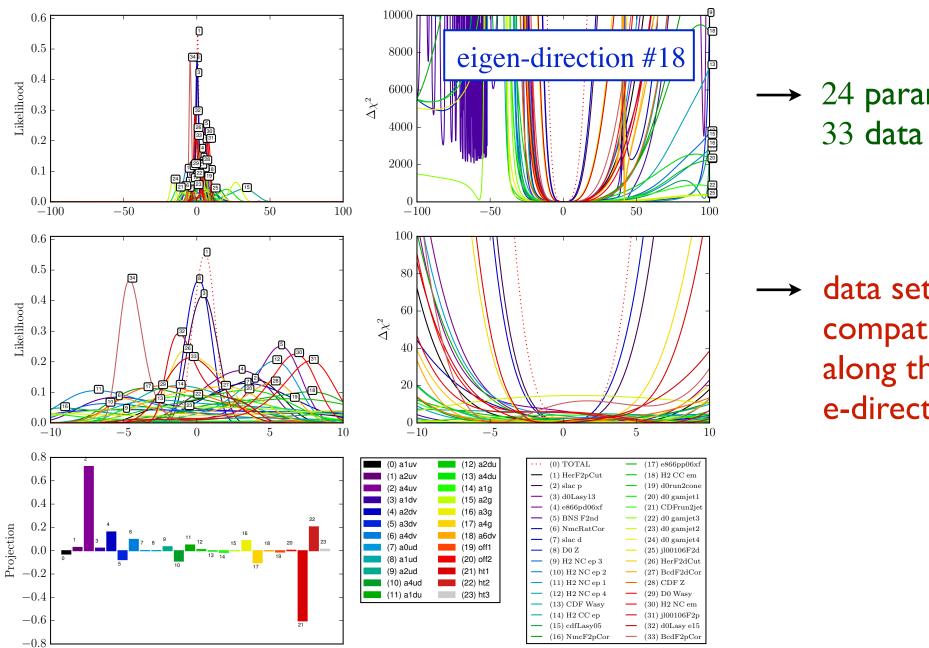


(16) NmcF2pCor

-1.0

— (33) BcdF2pCor

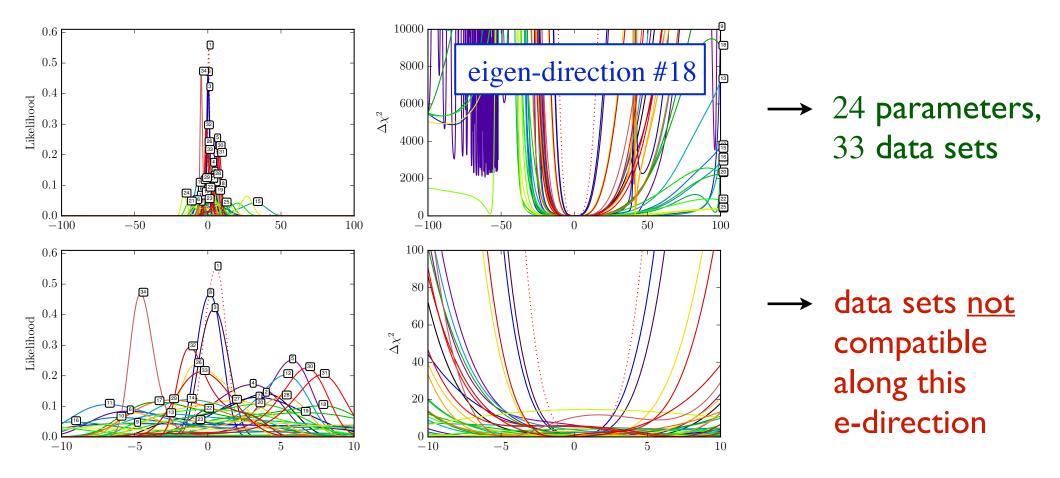
■ Realistic example: recent CJ (CTEQ-JLab) global PDF analysis



 $\rightarrow$  24 parameters, 33 data sets

data sets not compatible along this e-direction

■ Realistic example: recent CJ (CTEQ-JLab) global PDF analysis



standard Gaussian likelihood incapable of accounting for underestimated individual errors (leading to incompatible data sets)
 not designed for such scenarios!

- Two ways in which tolerance factors usually implemented
  - → CTEQ "tolerance criteria"
     (variations adopted by other groups, e.g., MMHT, CJ)

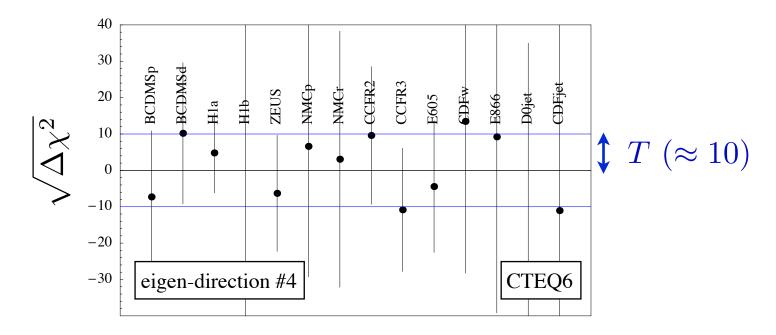
Pumplin, Stump, Huston, Lai, Nadolsky, Tung JHEP 07 (2002) 012

 $\rightarrow$  scaling of  $\Delta\chi^2$  with number of parameters (or number of degrees of freedom)

e.g. Brodsky, Gardner PRL (Comment) **116**, 019101 (2016)

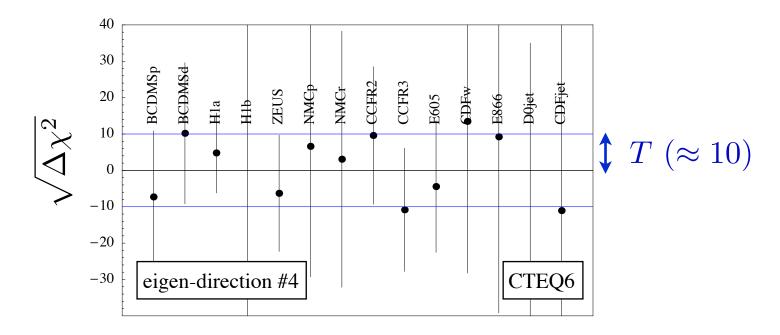
JDHLM assess their PDF errors using a tolerance criteria of  $\Delta\chi^2=1$  at  $1\sigma$ ; however, the actual value of  $\Delta\chi^2$  to be employed depends on the number of parameters to be simultaneously determined in the fit. This is illustrated in Table 38.2 of Ref. [15] and is used broadly, noting, e.g., Refs. [16–19]. Ref. [7] employs the CT10 PDF analysis [20], so that it contains 25 parameters, plus one for intrinsic charm. Figure 38.2 of Ref. [15] then shows that  $\Delta\chi^2\approx 29$  at  $1\sigma$  (68% CL), whereas  $\Delta\chi^2\approx 36$  at 90% CL. Ref. [7] uses the criterion  $\Delta\chi^2>100$ , determined on empirical grounds, to indicate a poor fit. JDHLM employs the framework of Ref. [21] which contains 25 parameters for the PDFs and 12 for the higher-twist contributions, so that a much larger tolerance than  $\Delta\chi^2=1$  is warranted.

#### 1 CTEQ tolerance criteria



- ullet for each experiment, find minimum  $\chi^2$  along given e-direction
- $\circ$  from  $\chi^2$  distribution determine 90% CL for each experiment
- o along each side of e-direction, determine maximum range  $d_k^\pm$  allowed by the most constraining experiment
- ullet T computed by averaging over all  $\,d_k^\pm\,$  (typically  $T\sim 5-10$ )

CTEQ tolerance criteria



- This approach is not consistent with Gaussian likelihood
  - → no clear Bayesian interpretation of uncertainties (ultimately, a prescription...)

- Scaling of  $\Delta \chi^2$  with # of parameters: " $\Delta \chi^2$  paradox"
- Simple example: two parameters  $\theta_i$  (i = 1, 2) with mean values  $\mu_i$  and standard deviation  $\sigma_i$ 
  - → joint probability distribution

$$\mathcal{P}(\theta_1, \theta_2) = \prod_{i=1,2} \frac{1}{\sqrt{2\pi\sigma_i^2}} \exp\left[-\frac{1}{2} \left(\frac{\theta_i - \mu_i}{\sigma_i}\right)^2\right]$$

 $\rightarrow$  change variables  $\theta_i \rightarrow t_i = (\theta_i - \mu_i)/\sigma_i$  and use polar coordinates  $r^2 = t_1^2 + t_2^2, \ \phi = \tan^{-1}(t_2/t_1)$ 

$$d\theta_1 d\theta_2 \mathcal{P}(\theta_1, \theta_2) = \frac{d\phi}{2\pi} r dr \exp\left[-\frac{1}{2}r^2\right]$$

- Scaling of  $\Delta \chi^2$  with # of parameters: " $\Delta \chi^2$  paradox"
  - → confidence volume

$$CV \equiv \int d\theta_1 d\theta_2 \, \mathcal{P}(\theta_1, \theta_2) = \int_0^R dr \, r \, \exp\left[-\frac{1}{2}r^2\right]$$
$$= 68\% \text{ for } R = 2.279$$

 $\longrightarrow$  note that  $R^2=t_1^2+t_2^2\equiv\chi^2$ , so that confidence region for parameters  $\max{[t_i]}=R$ 

 $\rightarrow$  implies that  $\theta_i = \mu_i \pm \sigma_i R$ , which contradicts original premise that  $\theta_i = \mu_i \pm \sigma_i$ !

- Scaling of  $\Delta \chi^2$  with # of parameters: " $\Delta \chi^2$  paradox"
  - → to resolve paradox, use Bayesian master formulas

$$E[\theta_i] = \int_0^{2\pi} \frac{d\phi}{2\pi} \int_0^{\infty} dr \, \mathcal{P}(r,\phi) \, \theta_i$$
$$= \int_0^{2\pi} \frac{d\phi}{2\pi} \int_0^{\infty} dr \, r \, e^{-r^2/2} \left(\mu_i + t_i \, \sigma_i\right) = \mu_i \quad \checkmark$$

- Scaling of  $\Delta \chi^2$  with # of parameters: " $\Delta \chi^2$  paradox"
  - → to resolve paradox, use Bayesian master formulas

$$E[\theta_i] = \int_0^{2\pi} \frac{d\phi}{2\pi} \int_0^{\infty} dr \, \mathcal{P}(r,\phi) \, \theta_i$$

$$= \int_0^{2\pi} \frac{d\phi}{2\pi} \int_0^{\infty} dr \, r \, e^{-r^2/2} \left(\mu_i + t_i \, \sigma_i\right) = \mu_i \quad \checkmark$$

$$V[\theta_{i}] = \int_{0}^{2\pi} \frac{d\phi}{2\pi} \int_{0}^{\infty} dr \, \mathcal{P}(r,\phi) \, (\theta_{i} - \mu_{i})^{2}$$

$$= \int_{0}^{2\pi} \frac{d\phi}{2\pi} \int_{0}^{\infty} dr \, r \, e^{-r^{2}/2} \, (t_{i} \, \sigma_{i})^{2} = \sigma_{i}^{2} \quad \checkmark$$

■ Scaling of  $\Delta \chi^2$  with # of parameters: " $\Delta \chi^2$  paradox"

 $\rightarrow$  no paradox if use  $\Delta \chi^2 = 1$  for <u>any number</u> of parameters to characterize the  $1\sigma$  CL

 $\rightarrow$  only consistent tolerance for Gaussian likelihood is T=1

#### To summarize standard maximum likelihood method...

- Gradient search (in parameter space) depends how "good" the starting point is
  - → for ~30 parameters trying different starting points is impractical, if do not have some information about shape
- Common to free parameters initially, then <u>freeze</u> those not sensitive to data ( $\chi^2$  flat locally)
  - $\rightarrow$  introduces bias, does not guarantee that flat  $\chi^2$  globally
- Cannot guarantee solution is <u>unique</u>
- lacktriangle Error propagation characterized by quadratic  $\chi^2$  near minimum
  - $\rightarrow$  no guarantee this is quadratic globally (e.g. Student t-distribution?)
- Introduction of tolerance modifies Gaussian statistics

# Monte Carlo methods

#### Monte Carlo

- Designed to faithfully compute Bayesian master formulas
- Do not assume a <u>single minimum</u>, include all possible solutions (with appropriate weightings)
- <u>Do not</u> assume likelihood is <u>Gaussian</u> in parameters
- Allows likelihood analysis to be extended to <u>address tensions</u> among data sets via Bayesian inference
- More computationally demanding compared with Hessian method

#### Monte Carlo

lacktriangle First group to use MC for global PDF analysis was NNPDF, using neural network to parametrize P(x) in

$$f(x) = N x^{\alpha} (1 - x)^{\beta} P(x)$$

- $\alpha, \beta$  are fitted "preprocessing coefficients"
- Iterative Monte Carlo (IMC), developed by JAM Collaboration, variant of NNPDF, tailored to non-neutral net parametrizations

→ J. Ethier

Markov Chain MC (MCMC) / Hybid MC (HMC)
 recent "proof of principle" analysis, ideas from lattice QCD

Gbedo, Mangin-Brinet, PRD **96**, 014015 (2017)

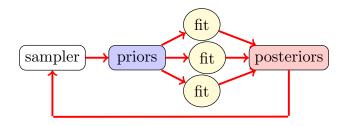
Nested sampling (NS) — computes integrals in Bayesian master formulas (for E, V, Z) explicitly

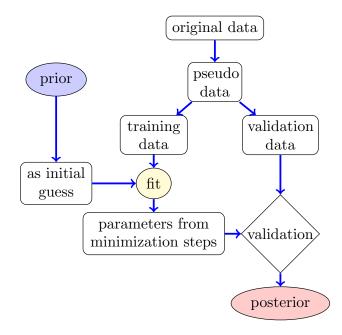
Skilling (2004)

## Iterative Monte Carlo (IMC)

 Use traditional functional form for input distribution shape, but sample significantly larger parameter space than possible in single-fit analyses

### Iterative Monte Carlo (IMC)

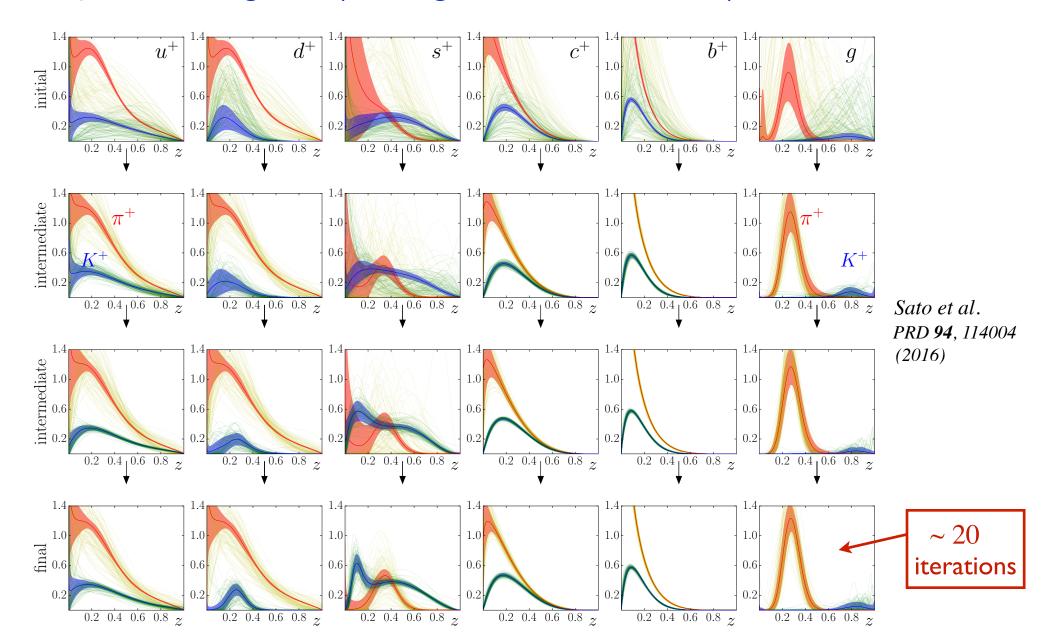




- → no assumptions for exponents
- cross-validation to avoid overfitting
- iterate until convergence criteria satisfied

## Iterative Monte Carlo (IMC)

### $\blacksquare$ e.g. of convergence (for fragmentation functions) in IMC

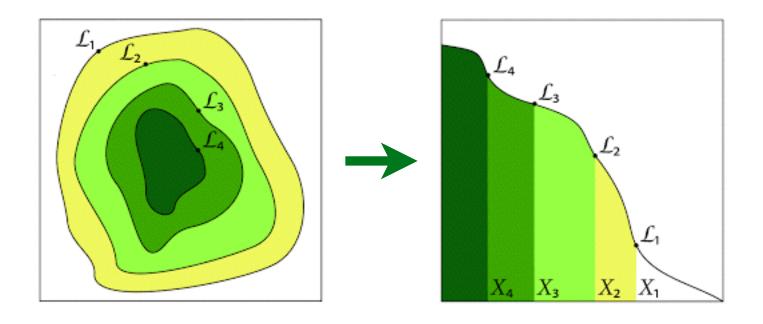


## **Nested Sampling**

lacktriangle Basic idea: transform n-dimensional integral to 1-D integral

$$Z = \int d^n a \, \mathcal{L}(\text{data}|\vec{a}) \, \pi(\vec{a}) = \int_0^1 dX \, \mathcal{L}(X)$$

where prior volume  $dX = \pi(\vec{a}) d^n a$ 



such that 
$$0 < \cdots < X_2 < X_1 < X_0 = 1$$

Feroz et al. arXiv:1306.2144 [astro-ph]

## **Nested Sampling**

Approximate evidence by a weighted sum

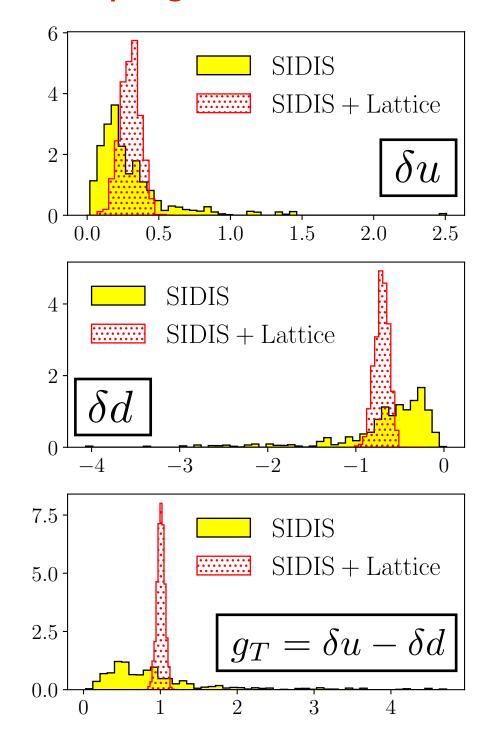
$$Z \approx \sum_i \mathcal{L}_i \, w_i$$
 with weights  $w_i = \frac{1}{2}(X_{i-1} - X_{i+1})$ 

- Algorithm:
  - $\rightarrow$  randomly select samples from full prior s.t. initial volume  $X_0 = 1$
  - $\rightarrow$  for each iteration, remove point with lowest  $\mathcal{L}_i$ , replacing it with point from prior with constraint that its  $\mathcal{L} > \mathcal{L}_i$
  - repeat until entire prior volume has been traversed
    - can be parallelized
  - performs better than VEGAS for large dimensions
  - o increasingly used in fields outside of (nuclear) analysis

## **Nested Sampling**

 Recent application in global analysis of transversity TMD PDF

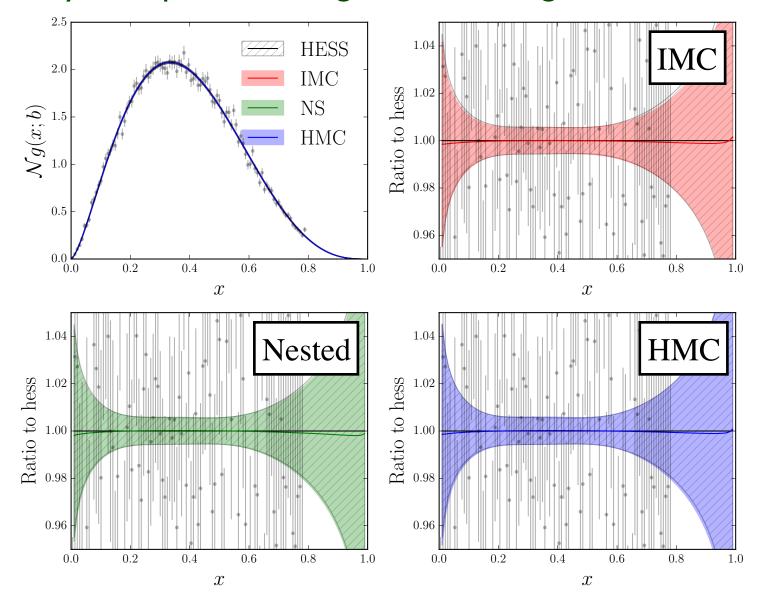
→ H.-W. Lin



Lin, WM, Prokudin, Sato, Shows (2017)

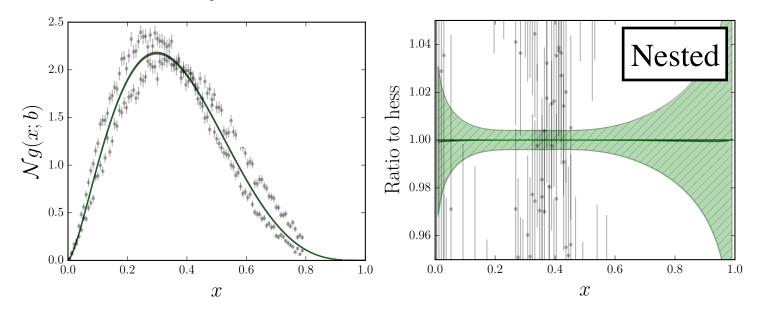
## MC Error Analysis

- Assuming a single minimum, a Hessian or MC analysis *must* give same results, if using same likelihood function
  - analysis of pseudodata, generated using Gaussian distribution



## MC Error Analysis

- Assuming a single minimum, a Hessian or MC analysis must give same results, if using same likelihood function
  - → also for discrepant data



almost identical uncertainty bands for Hessian and for MC!

## MC Error Analysis

- Assuming a single minimum, a Hessian or MC analysis must give same results, if using same likelihood function
- Approaches that use Hessian + tolerance factor not consistent with Gaussian likelihood function
- NNPDF group claim that within their neural net MC methodology, no need for a tolerance factor, since uncertainties similar to other groups who use Hessian + tolerance
  - $\rightarrow$  how can this be?
- Assuming sufficient observables to determine PDFs, then PDF uncertainties cannot depend on parametrization!

# Non-Gaussian likelihood

## Incompatible data sets

- Rigorous (Bayesian) way to address incompatible data sets is to use generalization of Gaussian likelihood
  - joint vs. disjoint distributions
  - empirical Bayes
  - hierarchical Bayes
  - o others, used in different fields

### Disjoint distributions

Instead of using total likelihood that is a <u>product</u> ("and") of individual likelihoods, e.g. for simple example of two measurements

$$\mathcal{L}(m_1 m_2 | m; \delta m_1 \delta m_2) = \mathcal{L}(m_1 | m; \delta m_1) \times \mathcal{L}(m_2 | m; \delta m_2)$$

use instead <u>sum</u> ("or") of individual likelihoods

$$\mathcal{L}(m_1 m_2 | m; \delta m_1 \delta m_2) = \frac{1}{2} \left[ \mathcal{L}(m_1 | m; \delta m_1) + \mathcal{L}(m_2 | m; \delta m_2) \right]$$

→ gives rather different expectation value and variance

$$E[m] = \frac{1}{2}(m_1 + m_2)$$
 
$$V[m] = \frac{1}{2}(\delta m_1^2 + \delta m_2^2) + \left(\frac{m_1 - m_2}{2}\right)^2$$
 depends on separation!

## Disjoint distributions

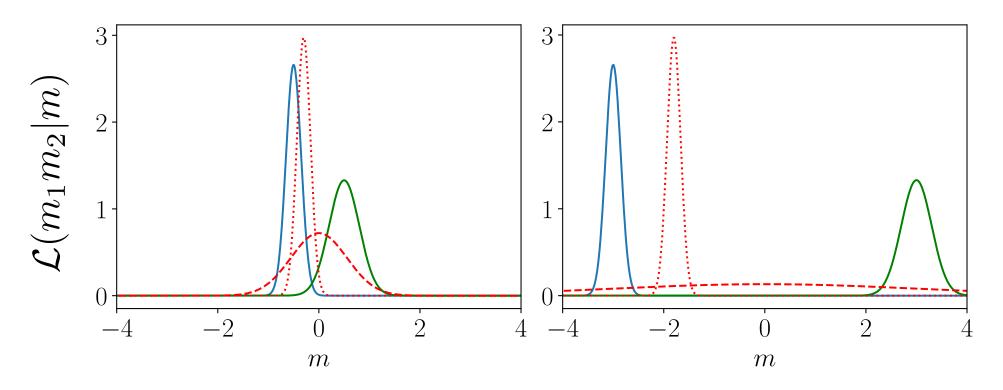
Symmetric uncertainties  $\delta m_1 = \delta m_2$ 

disjoint: 
$$V[m] = \frac{1}{2}(\delta m_1^2 + \delta m_2^2) + \left(\frac{m_1 - m_2}{2}\right)^2$$

joint: 
$$V[m] = \frac{\delta m_1^2 \ \delta m_2^2}{\delta m_1^2 + \delta m_2^2}$$

## Disjoint distributions

**Asymmetric uncertainties**  $\delta m_1 \neq \delta m_2$ 



→ disjoint likelihood gives broader overall uncertainty, overlapping individual (discrepant) data

## **Empirical Bayes**

- Shortcoming of conventional Bayesian still <u>assume</u> <u>prior</u> distribution follows specific form (e.g. Gaussian)
- Extend approach to more fully represent prior uncertainties,
   with final uncertainties that do not depend on initial choices
- In generalized approach, data uncertainties modified by distortion parameters, whose probability distributions given in terms of "hyperparameters" (or "nuisance parameters")
- Hyperparameters determined from data
  - → give posteriors for both PDF and hyperparameters

### **Empirical Bayes**

Standard mean and variance that characterize data

$$\theta = \mu + \sigma \longrightarrow f(\mu) + g(\sigma)$$

where  $f(\mu),g(\sigma)$  are unknown functions that account for faulty measurements

Simple choice is

$$(\mu, \sigma) \rightarrow (\zeta_1 \mu + \zeta_2, \zeta_3 \sigma)$$

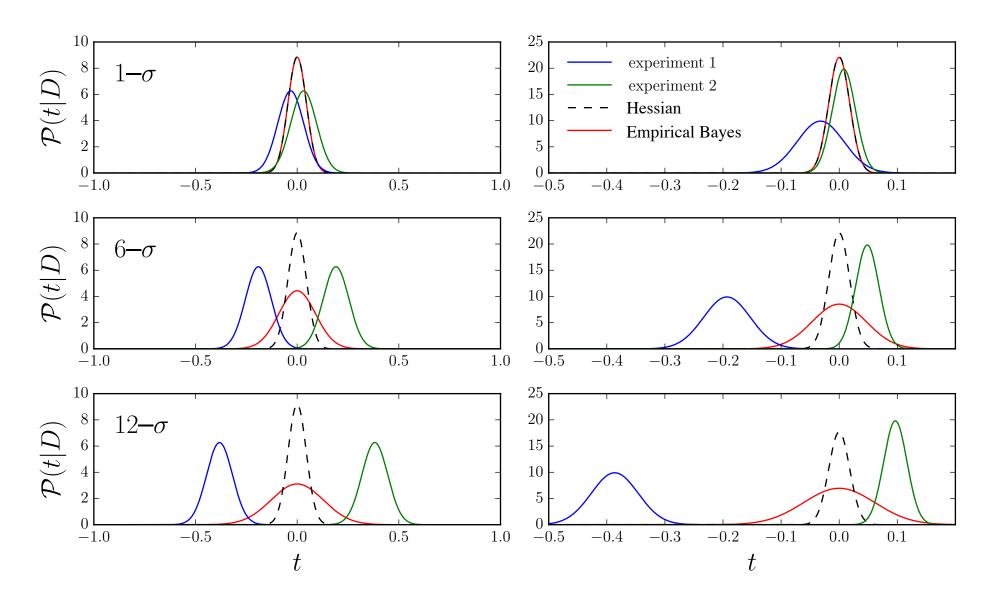
where  $\zeta_{1,2,3}$  are <u>distortion parameters</u>, with prob. dists. described by hyperparameters  $\phi_{1,2,3}$ 

Likelihood function is then

$$\mathcal{L}(\text{data}|\vec{a},\zeta_{1,2,3}) \sim \exp\left[-\frac{1}{2}\sum_{i}\left(\frac{d_{1}-f(\mu_{i}(\vec{a},\zeta_{1,2}))}{g(\sigma,\zeta_{3})}\right)^{2}\right]\pi_{1}(\zeta_{1}|\phi_{1})\pi_{2}(\zeta_{1}|\phi_{2})\pi_{3}(\zeta_{1}|\phi_{3})$$

## **Empirical Bayes**

# Simple example of EB for symmetric & asymmetric errors



### Outlook

- New paradigm needed in global QCD analysis
  - <u>simultaneous</u> determination of collinear distributions
     (also TMDs) using <u>Monte Carlo</u> sampling of parameter space
- Treatment of discrepant data sets needs serious attention
  - Bayesian perspective has clear merits
- Necessary to benchmark MC extractions (not just NNPDF)
- Near-term future: "universal" QCD analysis of all observables sensitive to collinear (unpolarized & polarized) PDFs and FFs
- Longer-term: apply MC technology to global QCD analysis of transverse momentum dependent (TMD) PDFs and FFs