

# Data analysis (in lattice QCD)

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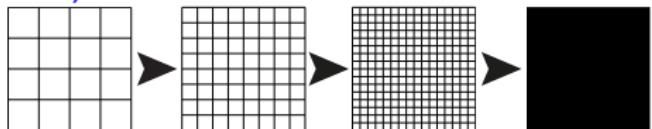
Lattice Practices 2024, The Cyprus Institute



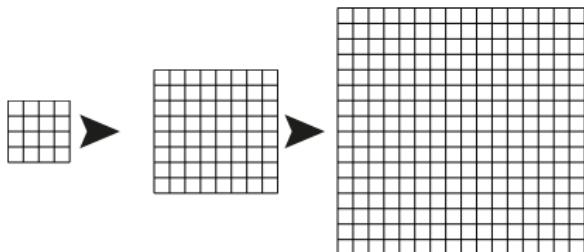
# Lattice

Lattice QCD=QCD when

- Cutoff removed (continuum limit)



- Infinite volume limit taken



- At physical hadron masses (Especially  $\pi$ )
  - Numerically challenging to reach light quark masses

Statistical error from stochastic estimate of the path integral

# Basic task

- Goal of phenomenological lattice QCD:
  - Compute expectation values of physical observables (masses, matrix elements,...)
  - Get reliable total errors of physical predictions
  - Use a minimum amount of computer time to obtain them
- Data analysis should:
  - provide results with reliable total errors
  - show how to efficiently improve the results

It's not about the final number, it's all about reliable errors

# Errors

Errors fall into 2 broad categories:

- Statistical errors:

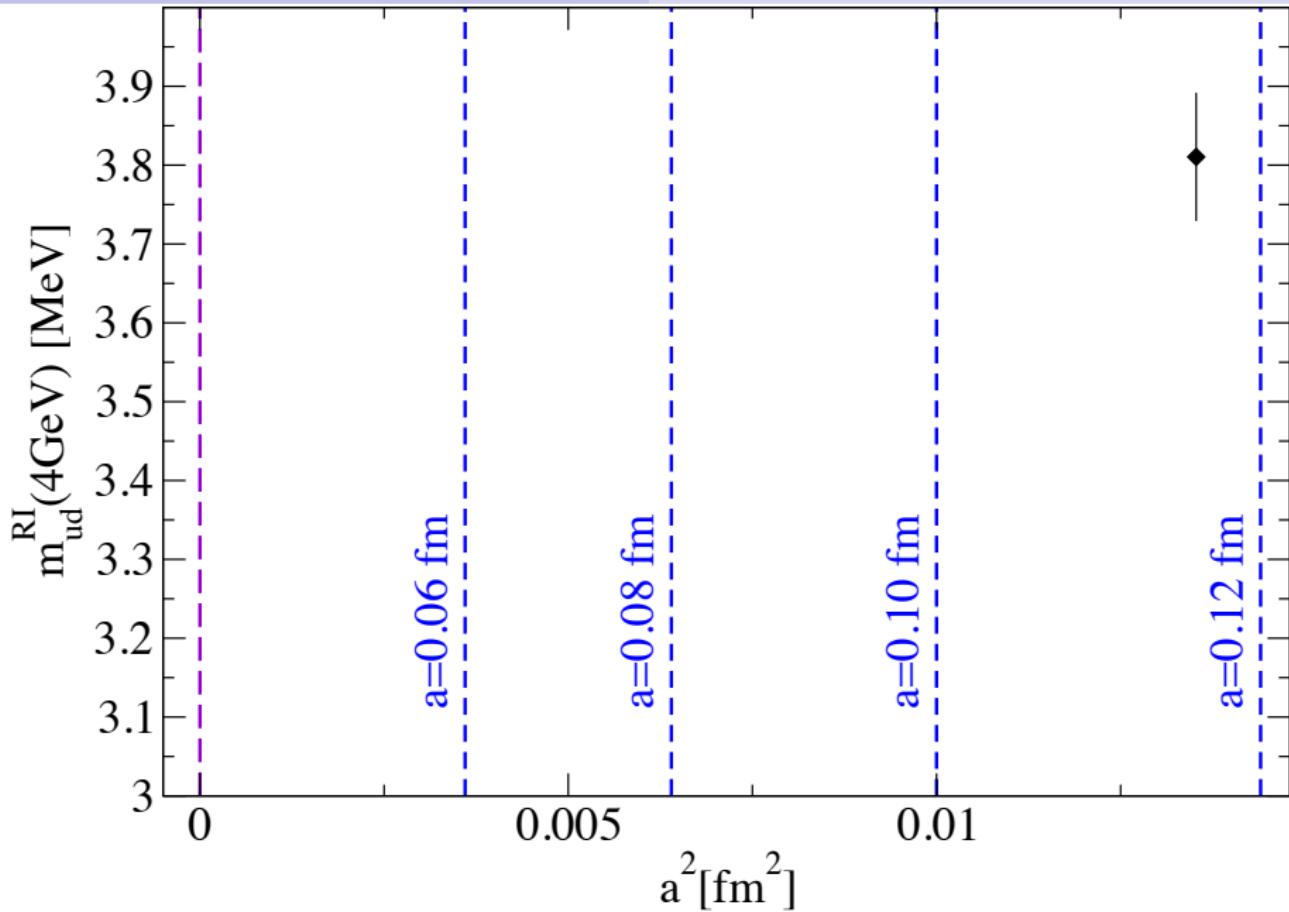
- Origin: stochastic evaluation of the path integral
- Can be treated by standard methods (e.g. bootstrap)

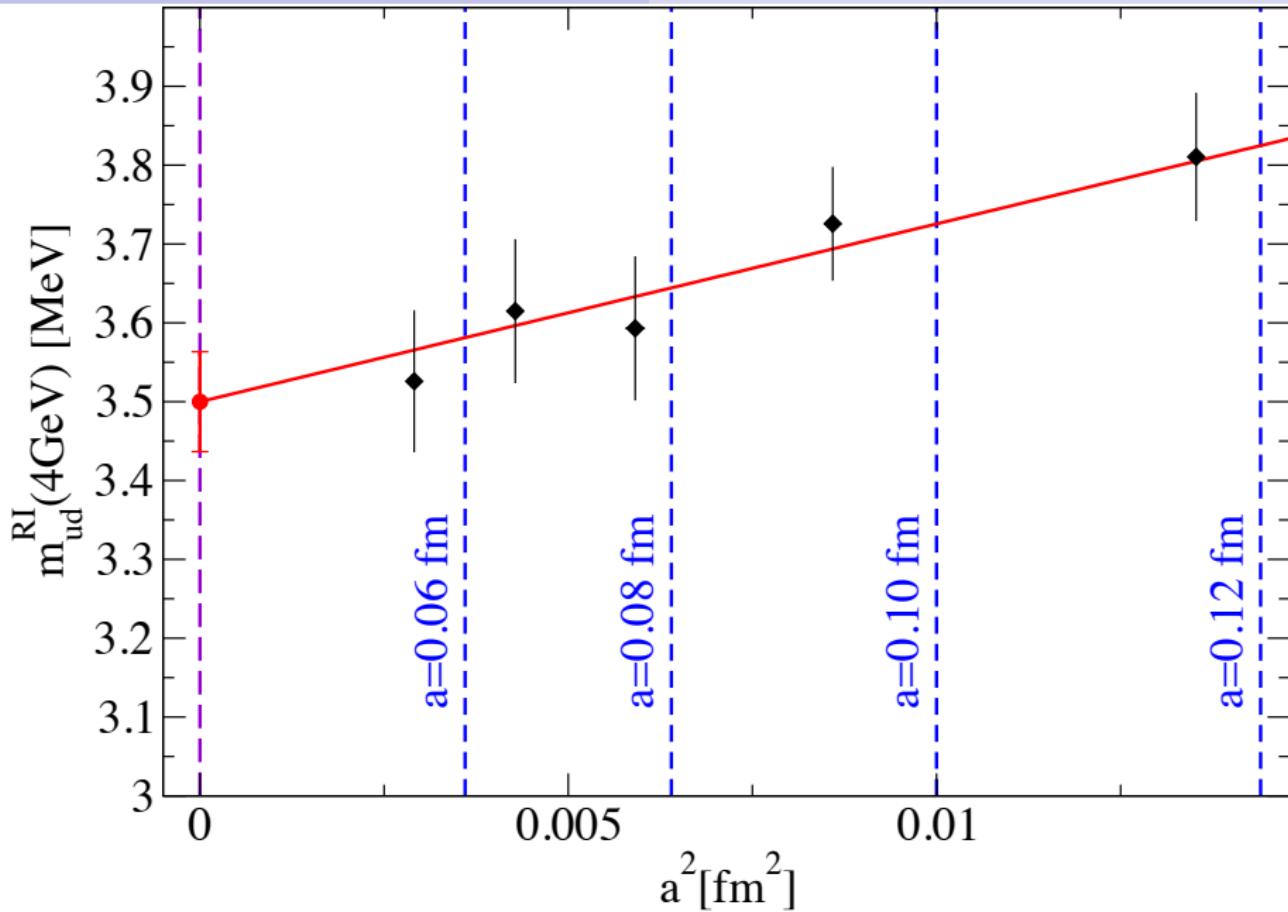
- Systematic errors:

- Origin: our lack of knowledge
- Can not be computed, only estimated

Keep good balance between the two!

- All systematics needs to be included for a correct result!





# What we will practice

In this course, we will:

- Generate fake propagators
  - Everyone deals with a separate set
  - We know the solution
- Extract ground state mass (exercise 1)
- Extrapolate an observable to the “physical point” (exercise 2)
- Tutorial: focus on practical aspects

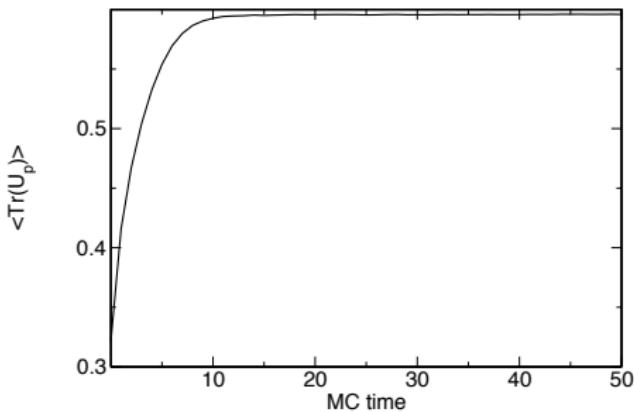
# Autocorrelation

Lattice data are typically Markov chains:

- Each ensemble is based on the previous one
- Need independent ensembles in equilibrium distribution

Two problems:

- Thermalization
  - Affects only beginning
  - Cut initial configs
- Autocorrelation
  - Reduces number of independent configs
  - Different per observable



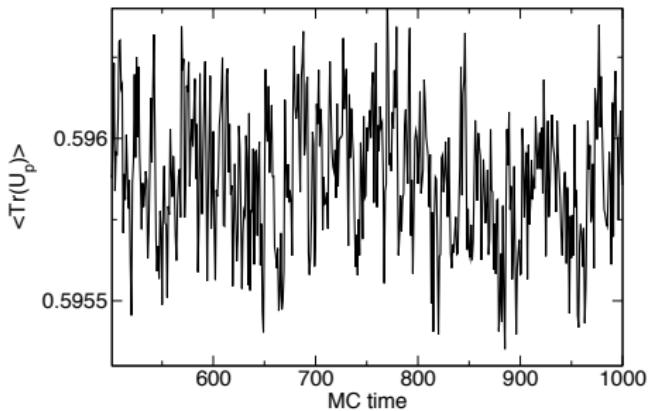
# Autocorrelation

Lattice data are typically Markov chains:

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  - Reduces number of independent configs
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# Autocorrelation - definitions

Given a time series  $a_t$ , the autocorrelation is the correlation of the time series with itself at a lag  $T$

$$R(T) = \frac{\langle (a_t - \langle a_t \rangle)(a_{T+t} - \langle a_{T+t} \rangle) \rangle}{\langle a_t \rangle \langle a_{T+t} \rangle}$$

In a stationary random process

$$R(T) \sim e^{-T/\tau}$$

with the autocorrelation time  $\tau$

# Autocorrelation - effects

We usually compute the integrated autocorrelation time

$$\tau_{\text{int}} = \sum_{T=1}^N R(T) \sim \int_0^\infty dT e^{-T/\tau} = \tau$$

Autocorrelation reduces the effective number of measurements

$$\sigma_{\langle a \rangle}^2 \approx \frac{\sigma_a^2}{N}$$

Minimize autocorrelation: blocking the data

$$a_x = \frac{1}{B} \sum_{b=0}^{B-1} a_{bx+b}$$

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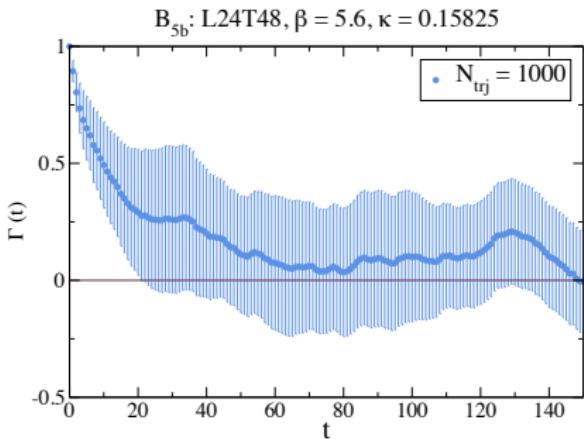
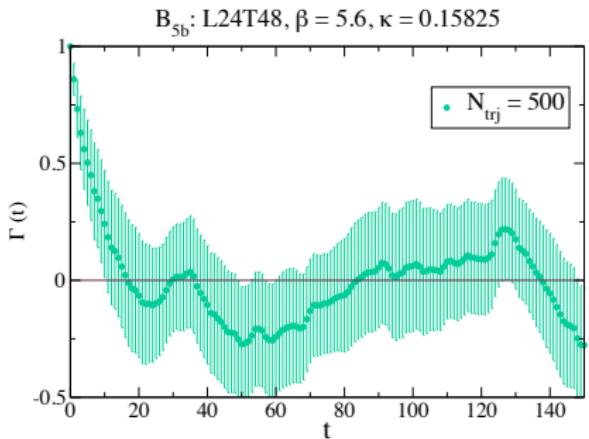
Autocorrelation reduces the effective number of measurements

$$\sigma_{\langle a \rangle}^2 \approx \frac{\sigma_a^2}{N} (1 + 2\tau_{\text{int}})$$

Minimize autocorrelation: blocking the data

$$a_x = \frac{1}{B} \sum_{b=0}^{B-1} a_{bx+b}$$

## Autocorrelation

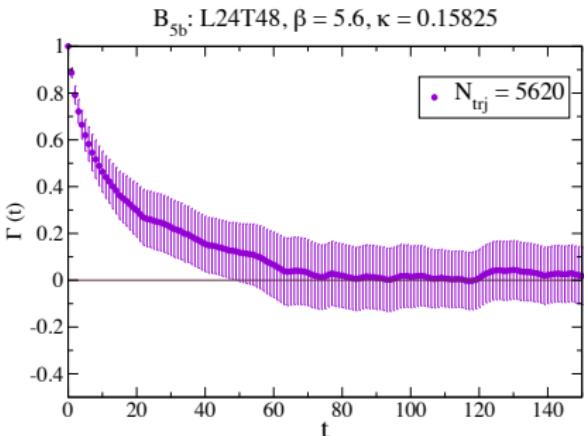


Difficult to compute  $\tau_{\text{int}}$  accurately

- Time series long enough
- Observable dependent
- Global observables slower

Example: plaquette in DDHMC

(Chowdhury et. al (2012))



# Autocorrelation - packages

There is a standard package you can feed your time series to:

U. Wolff, Monte Carlo errors with less errors,

Comput.Phys.Commun. 156:143-153,2004;

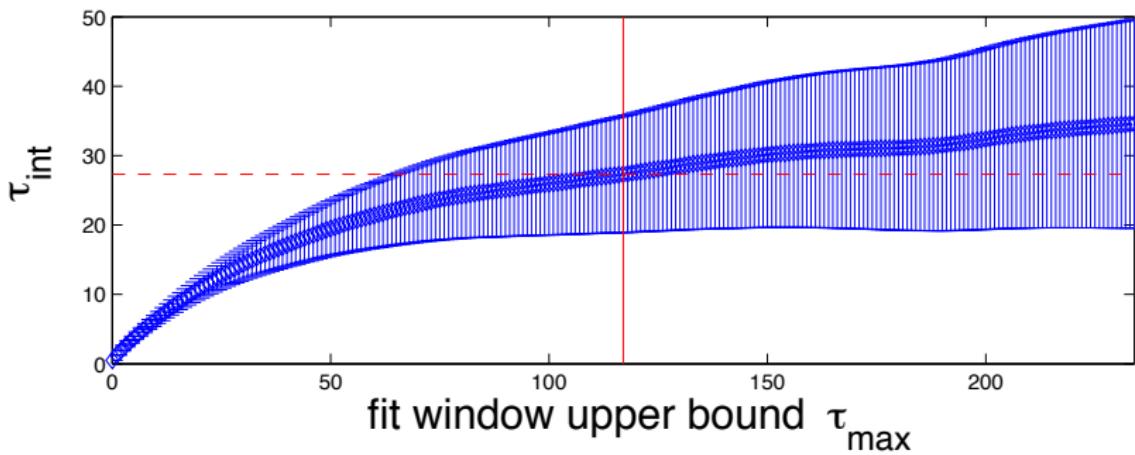
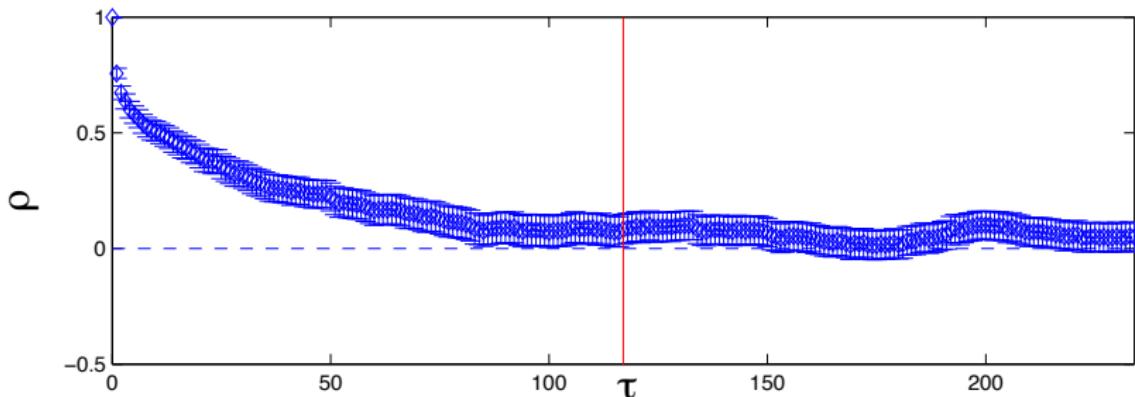
Erratum-ibid.176:383,2007

hep-lat/0306017

MATLAB code can be found at:

<http://www.physik.hu-berlin.de/com/ALPHAssoft/>

### Autocorrelation



# Ground state extraction

- Euclidean correlation function

$$c_t = \langle 0 | \mathcal{O}^\dagger(t) \mathcal{O}(0) | 0 \rangle$$

- Insert 1 =  $|i\rangle\langle i|$

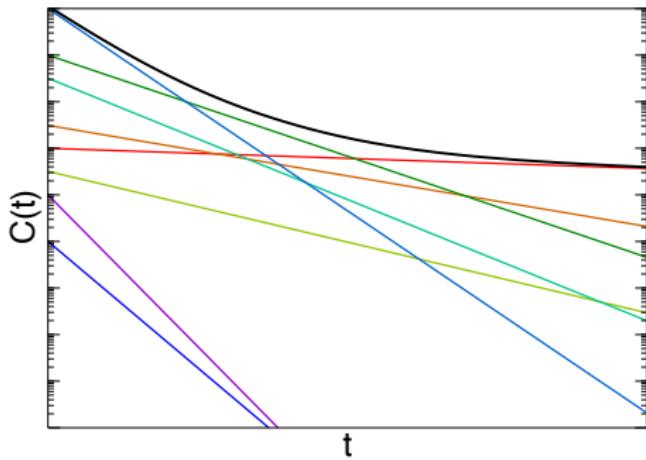
$$\sum_i \langle 0 | e^{Ht} \mathcal{O}^\dagger(0) e^{-Ht} | i \rangle \langle i | \mathcal{O}(0) | 0 \rangle$$

- Eigenbasis  $|i\rangle$  of  $H$

$$\sum_i |\langle 0 | \mathcal{O}(0) | i \rangle|^2 e^{-(E_i - E_0)t}$$

For  $t \rightarrow \infty$ :

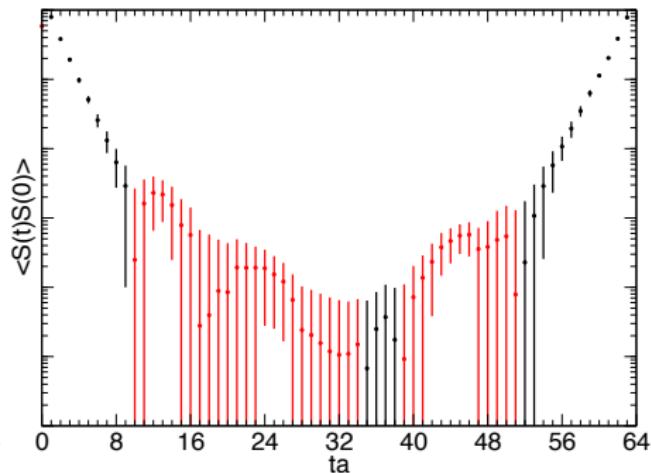
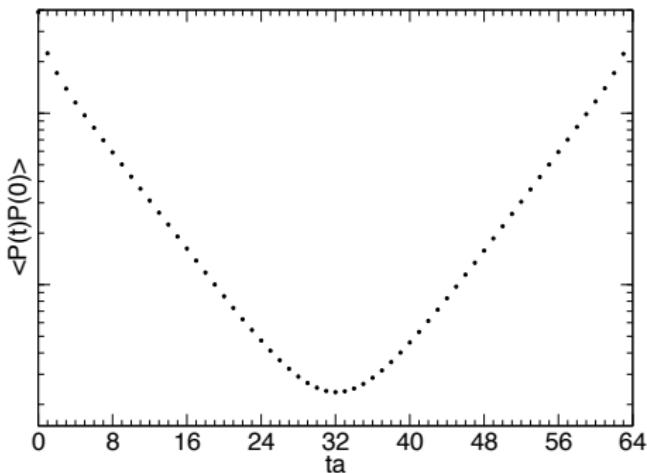
- Lightest state coupling to  $\mathcal{O}$  dominates:  $c_t \propto e^{-M \cdot t}$
- $M_{t+\frac{1}{2}} = \log[c_t/c_{t+1}]$ , prefactor  $\rightarrow$  matrix element



# Signals from propagators

There are several complications

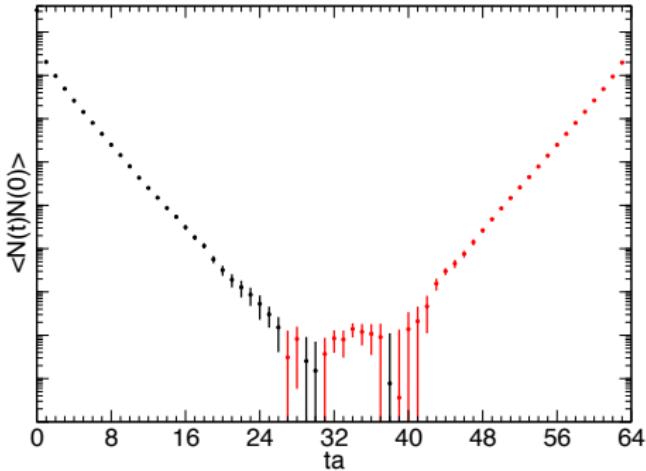
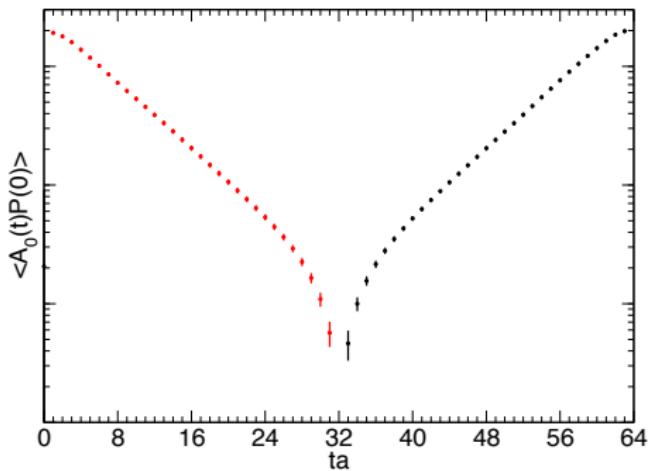
- Ground state coupling may be small
- Signal decays exponentially, noise not always
- There are backward (periodic BC) or border (open/fixed BC) contributions



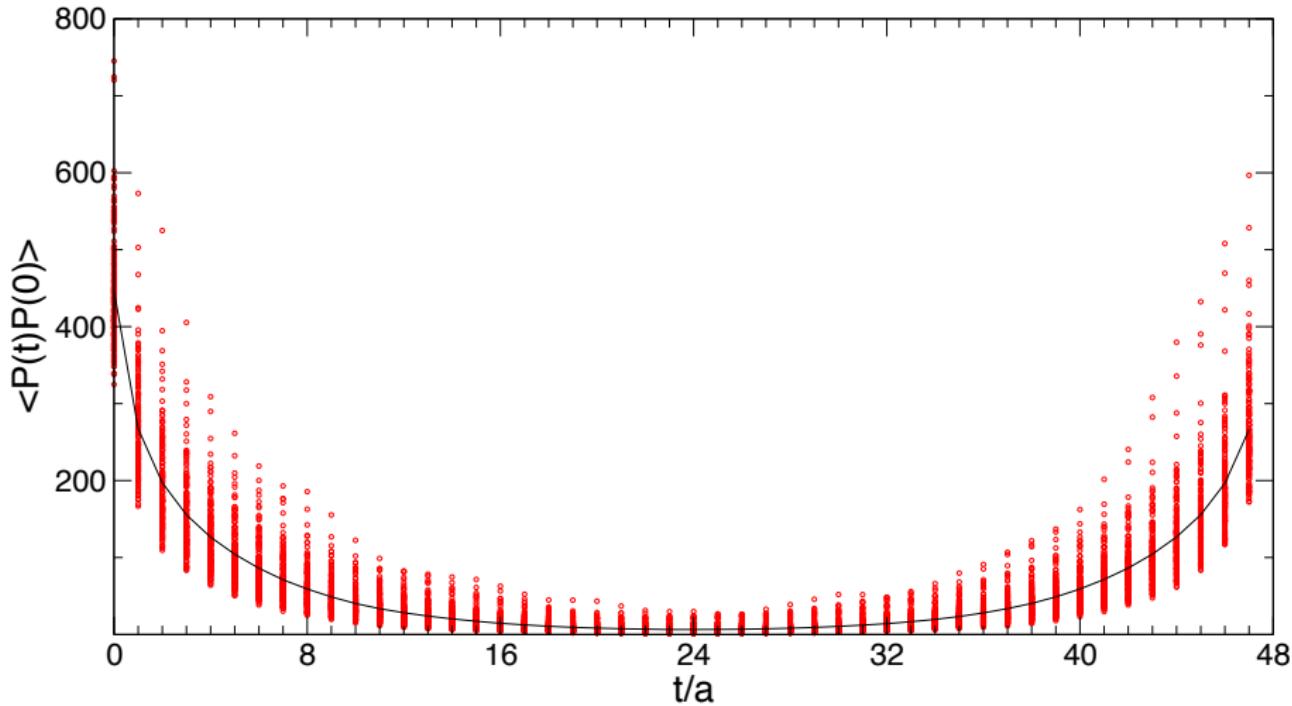
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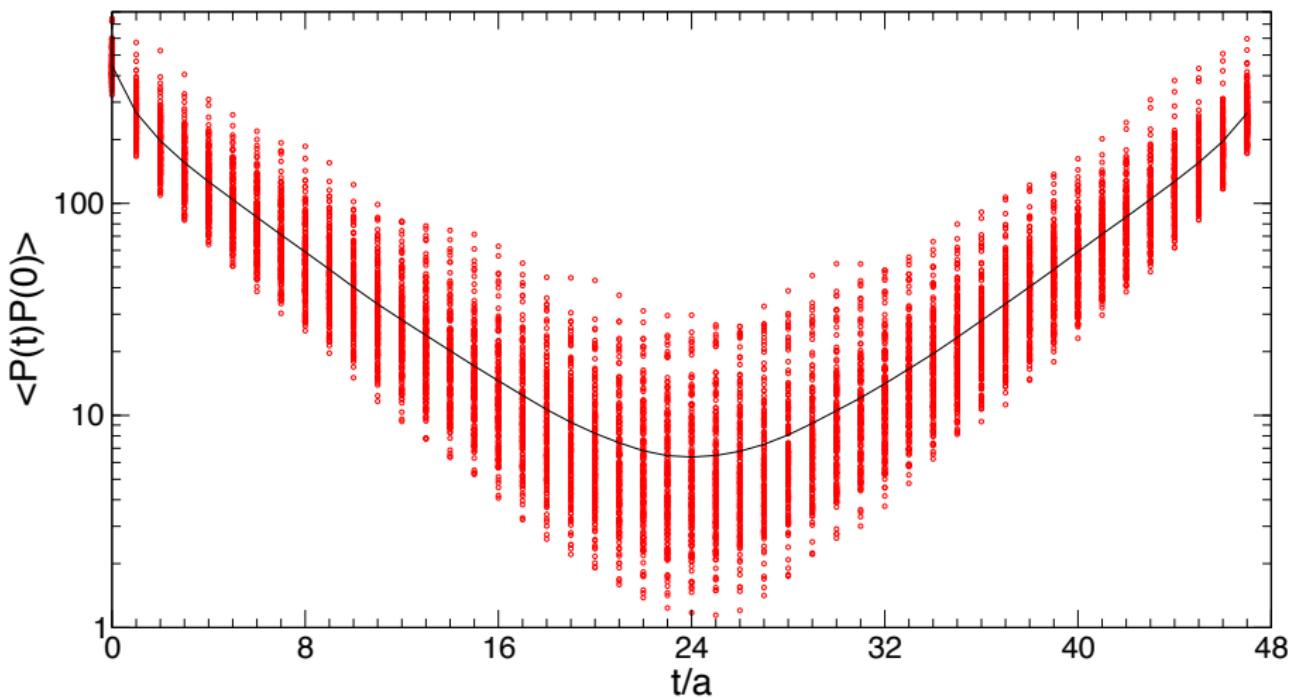
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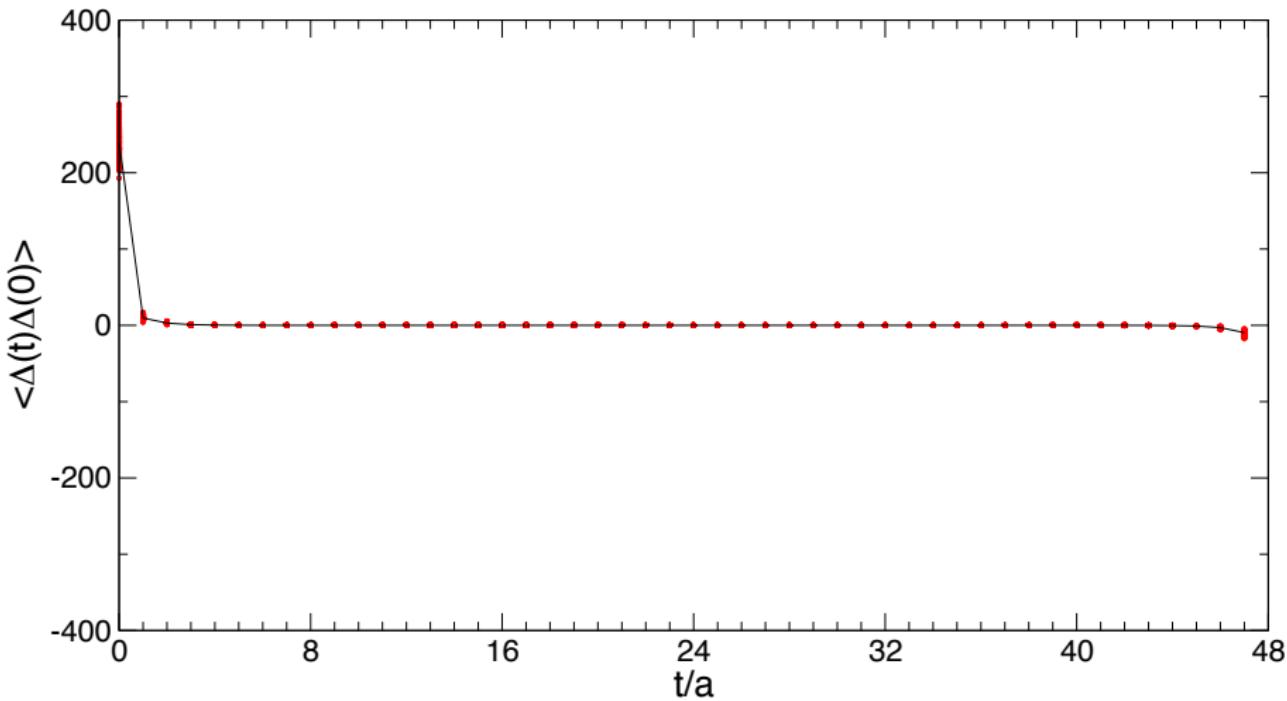
# Nice propagator example



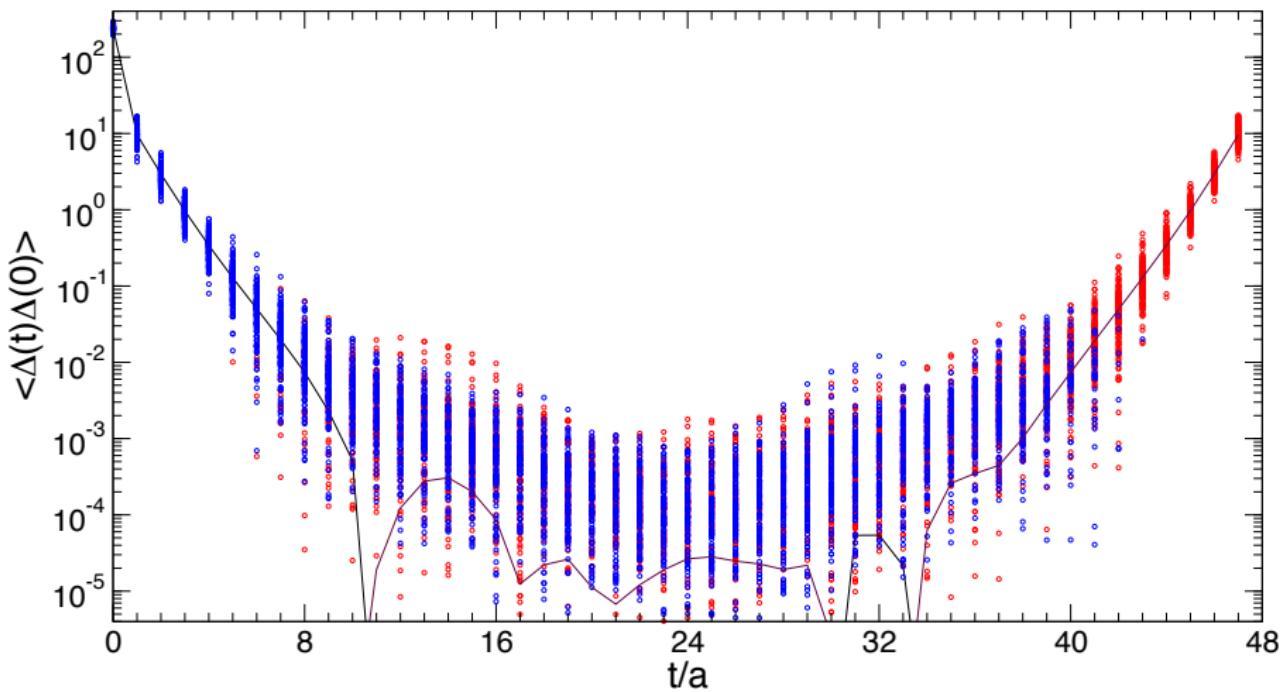
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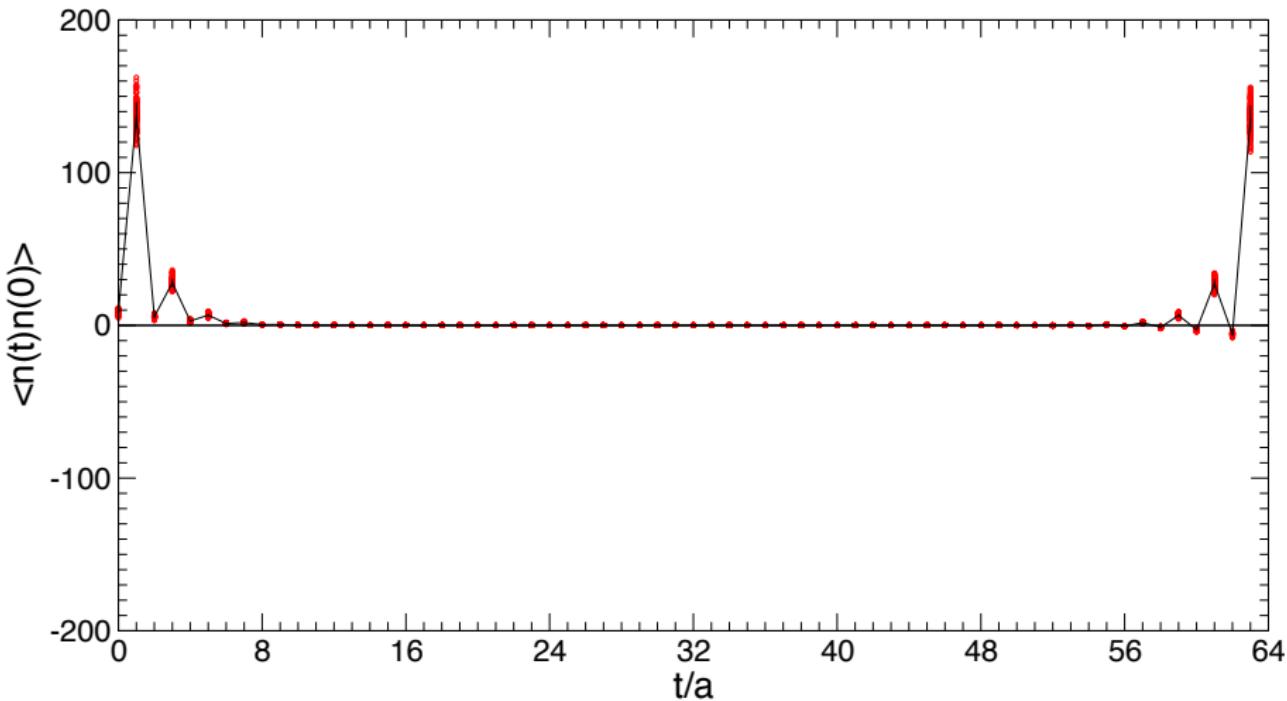
# Not so nice propagator example



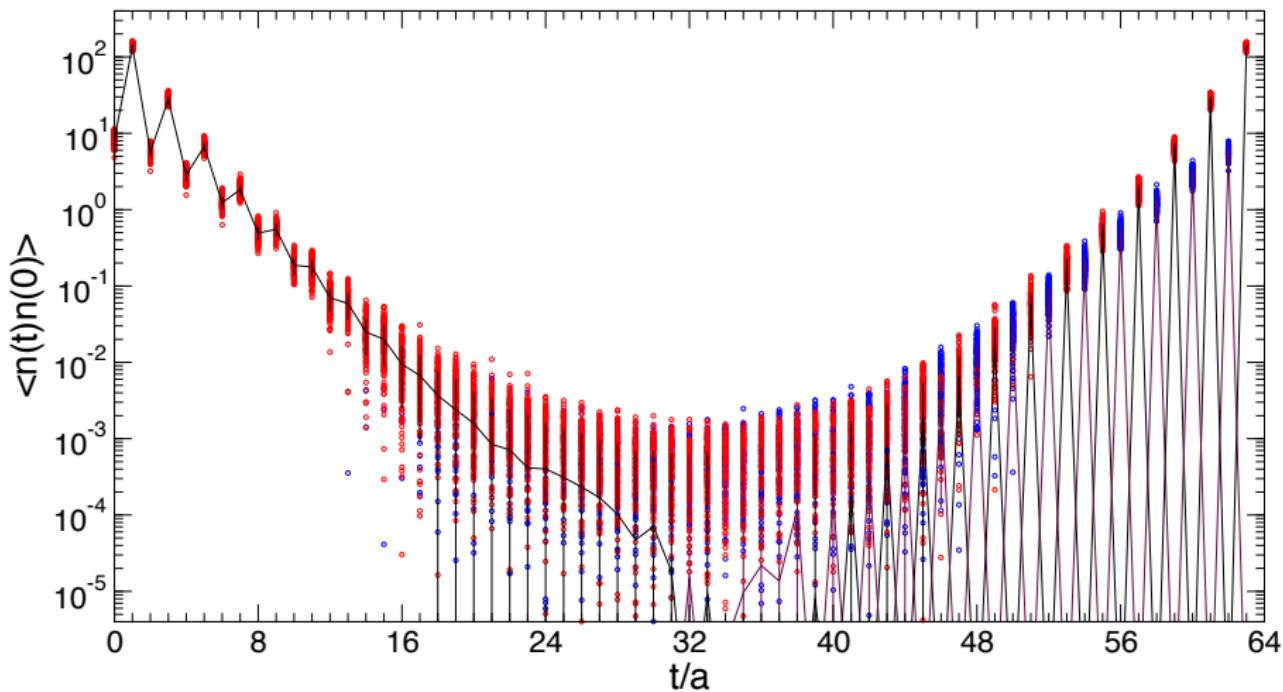
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# Not so nice propagator example



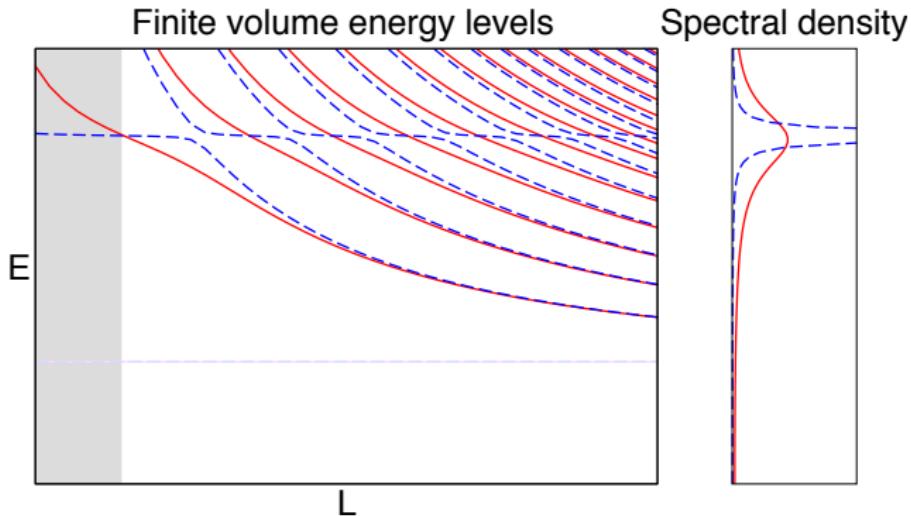
# Not so nice propagator example



# Excited state dominance

Small coupling of ground state is not an academic problem

- Occurs especially in resonant channels
- Ground state needs virtual  $q\bar{q}$  production
- Different operators couple very differently



# Propagator forms

Single state, propagating forward:

$$c_f(t) = c_f^0 e^{-mt}$$

The backward contribution:

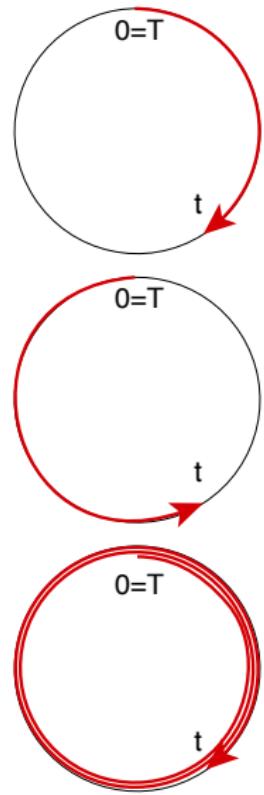
$$c_b(t) = c_b^0 e^{-m(T-t)}$$

Include contributions warping around the lattice (tiny):

$$c_f(t) = c_f^0 \left( e^{-mt} + e^{-m(T+t)} + \dots \right)$$

$$= c_f^0 e^{-mt} \times \sum_{n=0}^{\infty} e^{-nmT}$$

$$= c_f^0 e^{-mt} \frac{1}{1 - e^{-mT}}$$



# Propagator forms

For T (P) symmetric ( $c^0 = c_f^0 = c_b^0$ )  
 resp. antisymmetric ( $c^0 = c_f^0 = -c_b^0$ ):

$$\begin{aligned} c_t &= \frac{c^0}{1 - e^{-mT}} \left( e^{-mt} - e^{-m(T-t)} \right) \\ &= \frac{c^0}{1 - e^{-mT}} e^{-m\frac{T}{2}} \times \begin{cases} \cosh(m(\frac{T}{2} - t)) \\ \sinh(m(\frac{T}{2} - t)) \end{cases} \end{aligned}$$

Effective mass  $M_{t+\frac{1}{2}}$  from numerical solution of:

$$\frac{c_{t+1}}{c_t} = \frac{\cosh(M_{t+\frac{1}{2}}(\frac{T}{2} - t - 1))}{\cosh(M_{t+\frac{1}{2}}(\frac{T}{2} - t))}$$

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# Effective masses

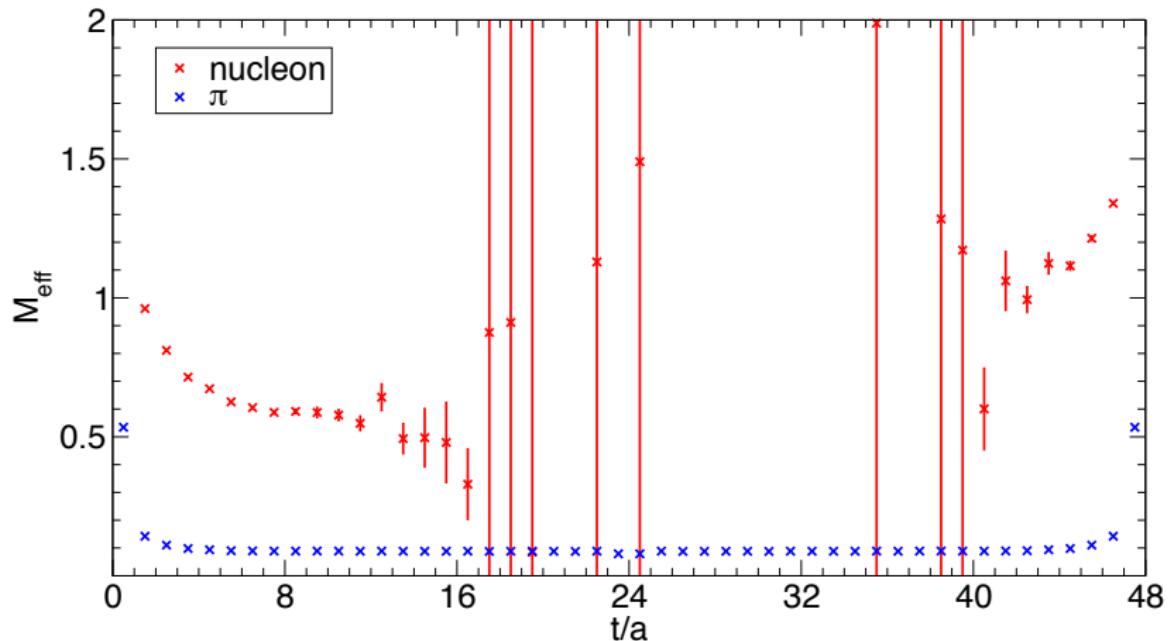
To identify asymptotic region, define an effective mass:

$$M_{\text{eff}}(t) = \frac{1}{l} \ln \frac{C(t - l/2)}{C(t + l/2)}$$

- Lag  $l$ : typically 1 or 2
- Modified versions including backward contributions
- Analytical 3-point expression for symmetric case:

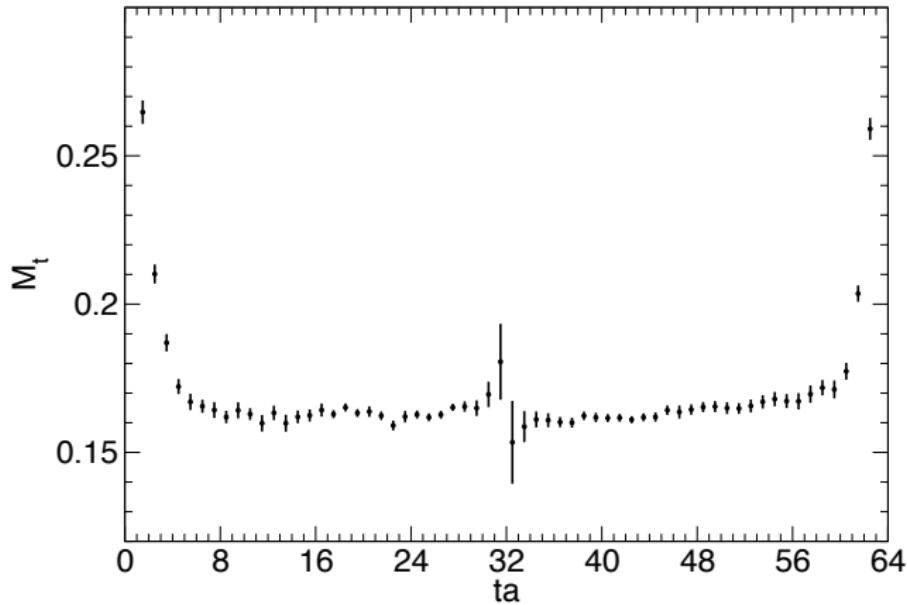
$$M_{\text{eff}}(t) = \text{acosh} \frac{c_{t+1} + c_{t-1}}{2c_t}$$

# Mass plateaus



Identify plausible fit ranges from mass plateaux

# Mass plateaus



Analytical 3-point expression (we will use this):

$$M_{t+\frac{1}{2}} = \text{acosh} \frac{c_{t+1} + c_{t-1}}{2c_t}$$

# Mass fit

After identifying plateau range, we fit the propagators with

$$p_t = \frac{c^0}{1 - e^{-mt}} \left( e^{-mt} \pm e^{-m(T-t)} \right)$$

where  $m$  and  $c^0$  are fit parameters

Maximum likelihood fit assuming normal error distribution:

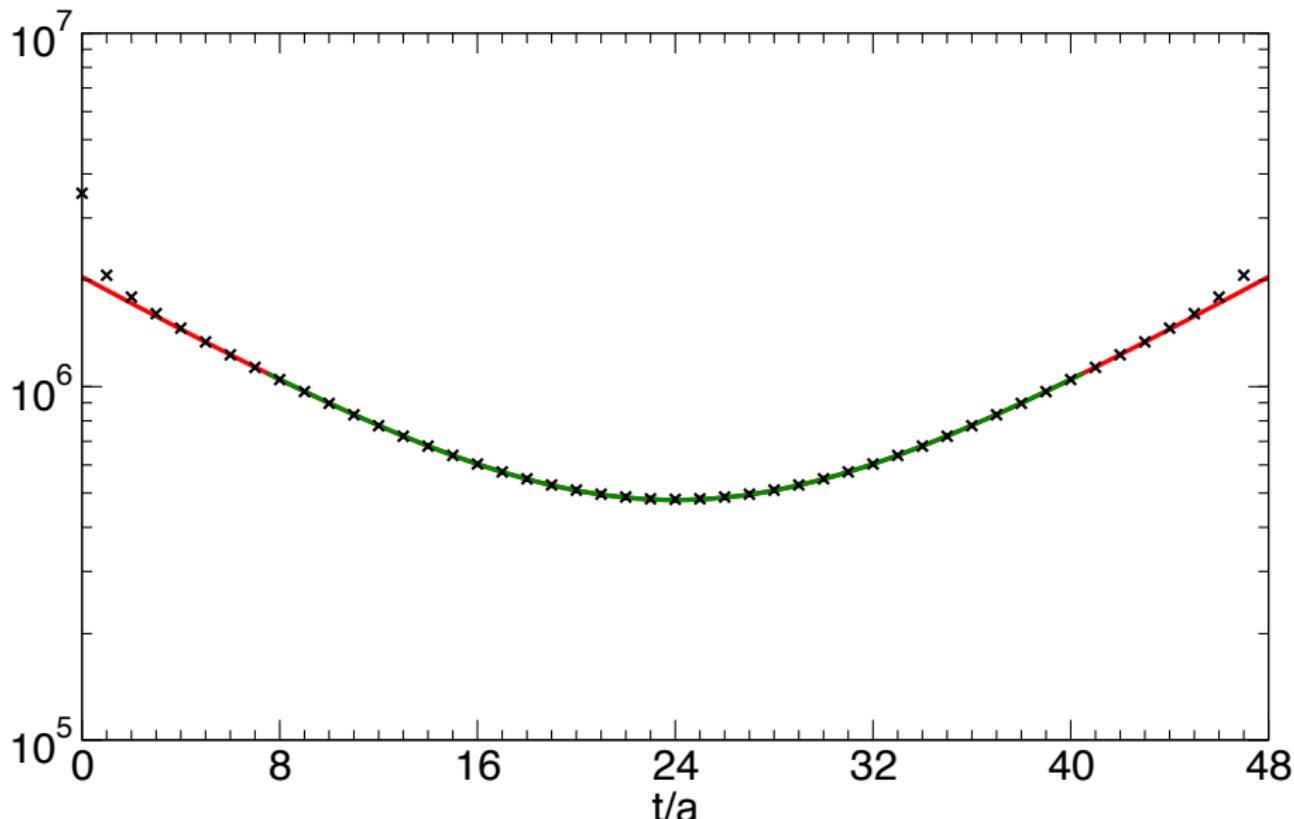
$$\chi^2 = (\mathbf{c} - \mathbf{p})_s (\Sigma^{-1})_{st} (\mathbf{c} - \mathbf{p})_t \rightarrow \min$$

Data points  $\mathbf{c}$ , fit function  $\mathbf{p}$  and covariance matrix  $\Sigma$

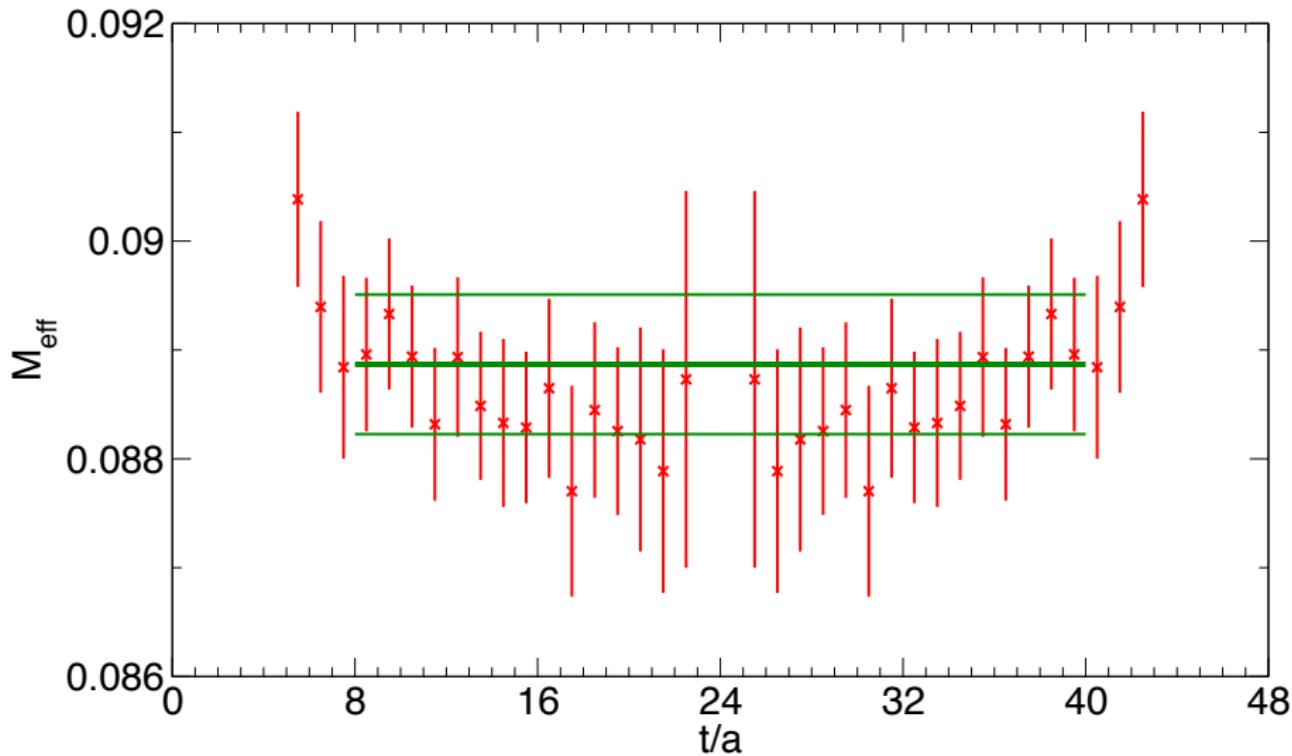
$$\Sigma_{st} = \langle (c_s - \langle c_s \rangle)(c_t - \langle c_t \rangle) \rangle$$

Usual variance in diagonal elements  $\Sigma_{tt} = \sigma(c_t)^2$

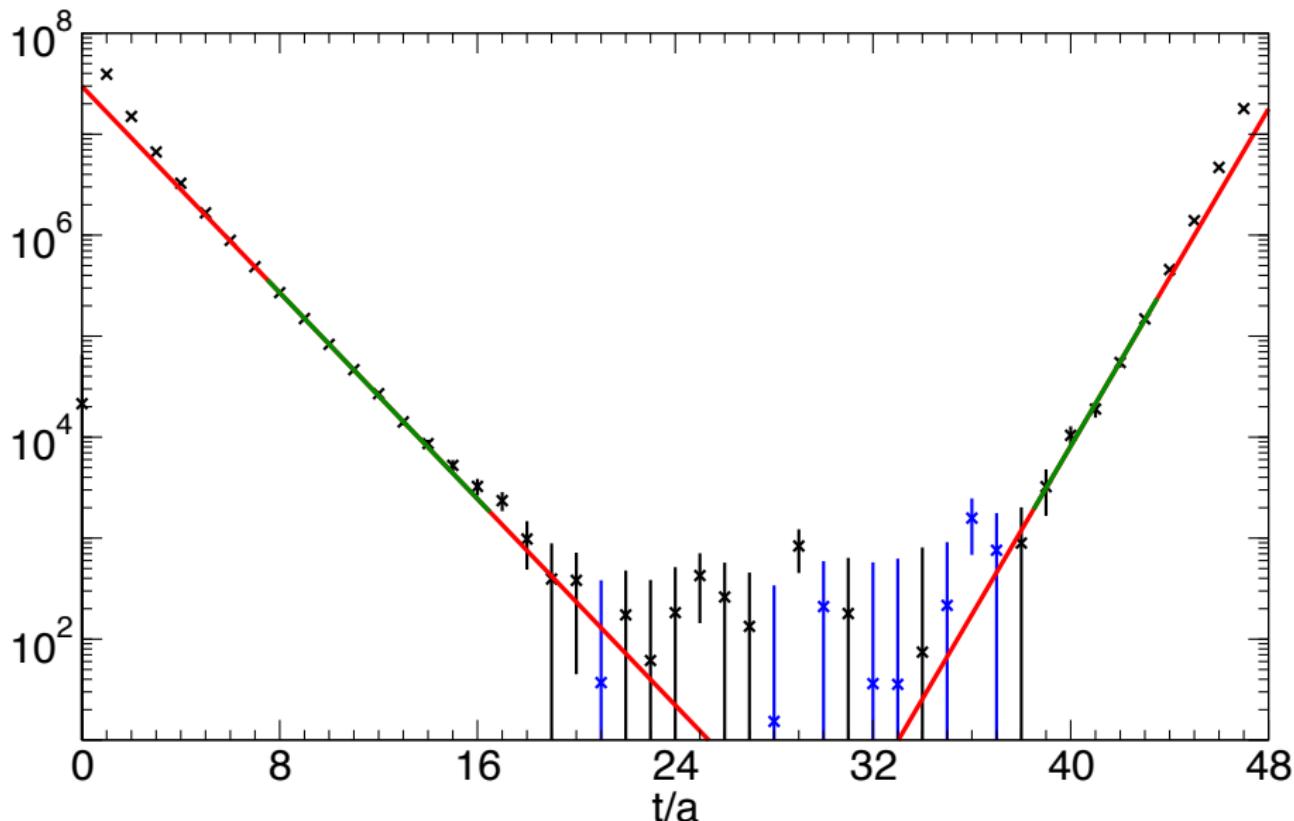
# Example pion fit



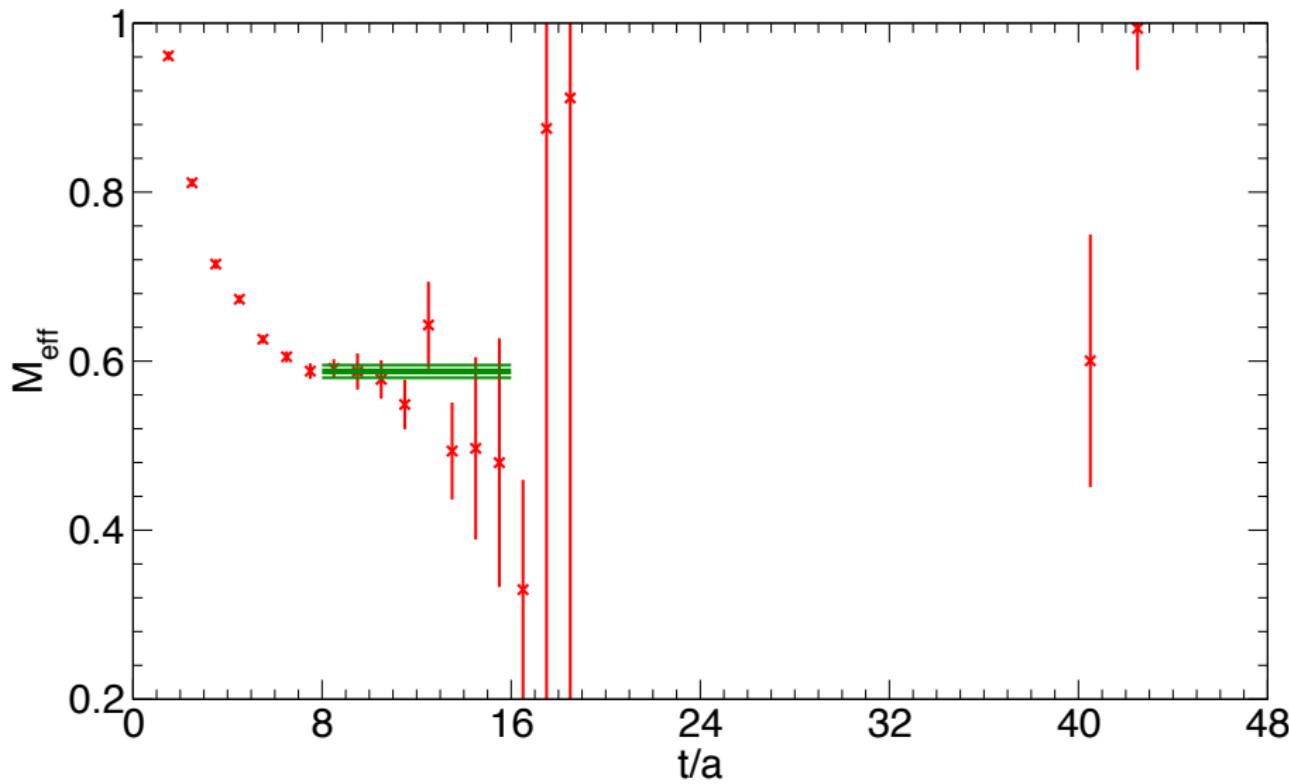
## Example pion fit



# Example nucleon fit



# Example nucleon fit

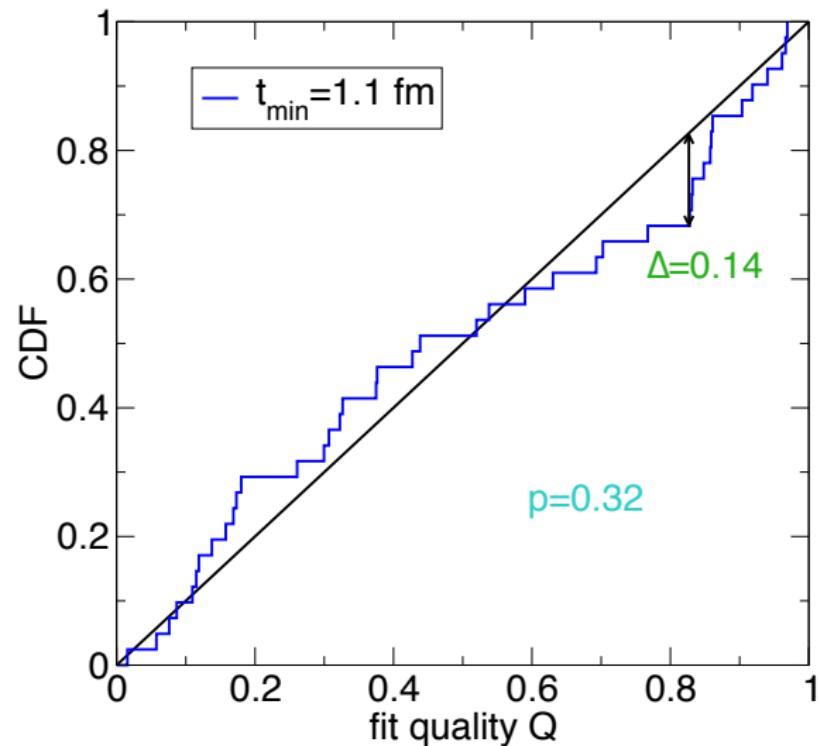


# Fit results

From a fit we in principle get 3 things:

- ✓ The most likely value of the fit parameters
  - Values of the parameters at  $\chi^2 \rightarrow \min$
- ✓ Standard errors of the parameters  
(more generally, confidence regions)
  - Contours of constant  $\Delta\chi^2 = \chi^2 - \chi^2_{\min}$
- ✓ The quality of the fit
  - From  $Q = \frac{\Gamma(\frac{n}{2}, \frac{\chi^2}{2})}{\Gamma(\frac{n}{2})} = \frac{\int_{\chi^2/2}^{\infty} t^{\frac{n}{2}-1} e^{-t} dt}{\int_0^{\infty} t^{\frac{n}{2}-1} e^{-t} dt}$
  - $Q$ : probability that - given the model - the data are at least as far off the prediction as the real data
  - ☞  $Q$  should be a flat random value  $\in [0, 1]$

# Example Q distribution



- Need many ensembles
- Plot CDF
- KS test flat distribution

$P(\Delta > \text{observed})$ :

$$p(\Delta(\sqrt{N} + 0.12 + \frac{0.11}{\sqrt{N}}))$$

with

$$p(x) = \sum_j \frac{(-)^{j-1} 2}{e^{-2j^2 x^3}}$$

# Correlations

For uncorrelated data,  $\Sigma$  is diagonal

$$C_{st} = \frac{\Sigma_{st}}{\sigma(c_s)\sigma(c_t)}$$

Typical (estimated) correlation matrix  $C$  for a correlator:

1.0000	0.9963	0.9840	0.9746	0.9509
0.9963	1.0000	0.9912	0.9801	0.9595
0.9840	0.9912	1.0000	0.9934	0.9846
0.9746	0.9801	0.9934	1.0000	0.9912
0.9509	0.9595	0.9846	0.9912	1.0000

Eigenvalues:

4.9224	0.0661	0.0059	0.0041	0.0014
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# Problems with correlations

The structure of the covariance matrix can be problematic

- Covariance matrix determined statistically
- In  $C^{-1}$ , small modes dominate
- Smallest modes have large errors

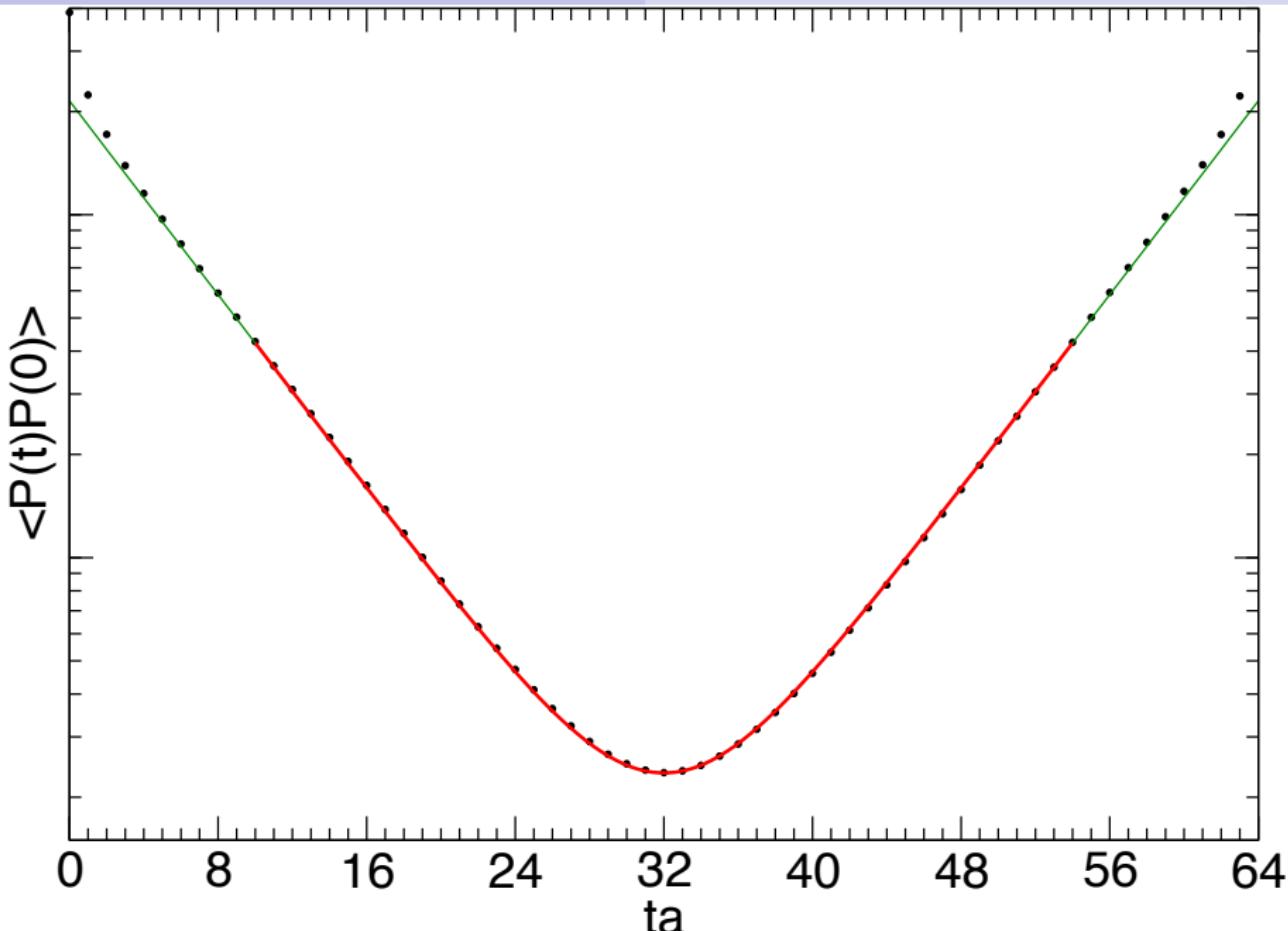
One can:

- Do an uncorrelated fit:  $\Sigma$  diagonal
- Truncate small eigenmodes
  - Truncate them (optionally correct diagonal)
  - Average them (Michael, Mc Kerrell, 1994)

Problem:  $Q$  and parameter errors useless

- Need to be determined in some other way

### Correlator fits



# Computing errors

When you make  $N$  measurements  $a_i$ , you compute

- the estimate of the expectation value

$$\langle a \rangle = \frac{1}{N} \sum_{i=1}^N a_i$$

- the estimated error of the expectation value

$$\sigma_{\langle a \rangle}^2 = \frac{1}{N} \frac{1}{N-1} \sum_{i=1}^N (a_i - \langle a \rangle)^2$$

From  $\mathcal{O}(100)$  configs, we get one mass measurement!  
Do we have to repeat this  $\mathcal{O}(100)$  times to estimate  $\sigma^2$ ?

# Resampling

No! We can resample our ensemble:

- Given  $N$  configs  $c_i$  and the full ensemble  $E = \{1, \dots, N\}$
- Given an observable  $O(A)$  on an arbitrary Ensemble  $A$
- We can produce one resampled ensemble  $B_1$  by drawing with repetition  $N$  configs from  $E$
- We actually draw  $N_B$  resampled ensembles  $B_i$
- We compute  $\overline{O} = O(E)$  and  $O_i = O(B_i)$

The distribution of  $O_i$  mimics independent measurements!

$$\sigma_O^2 \approx \sigma^2(O_i) \quad \langle O \rangle \approx \overline{O} + \overline{O} - \langle O_i \rangle$$

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$$\sigma_O^2 \approx \sigma^2(O_i) \quad \langle O \rangle \approx \bar{O} + \text{XXXX}$$

Usually better not to correct (stability)

# Jackknife

Jackknife is similar to bootstrap:

- Cut the ensemble  $E$  into  $N_J$  same size blocks
- Form  $N_J$  resampled ensembles  $J_i$  by leaving out one block from  $E$  at a time
- Compute  $\bar{O} = O(E)$  and  $O_i = O(J_i)$

$$\sigma_O^2 \approx (N_J - 1)\sigma^2(O_i) \quad \langle O \rangle \approx \bar{O} + (N_J - 1) (\bar{O} - \langle O_i \rangle)$$

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$$\sigma_O^2 \approx (N_J - 1)\sigma^2(O_i)$$

$$\langle O \rangle \approx \bar{O} + (\overline{X_{J_i}} - \bar{O})$$

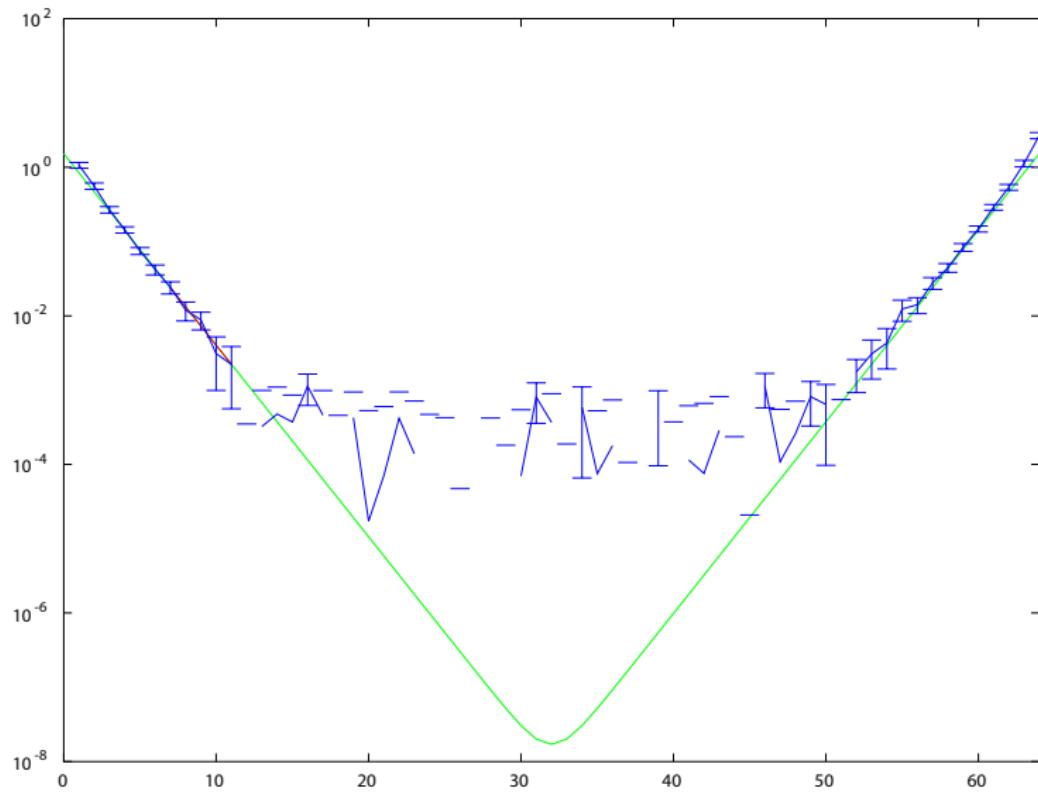
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# Using bootstrap

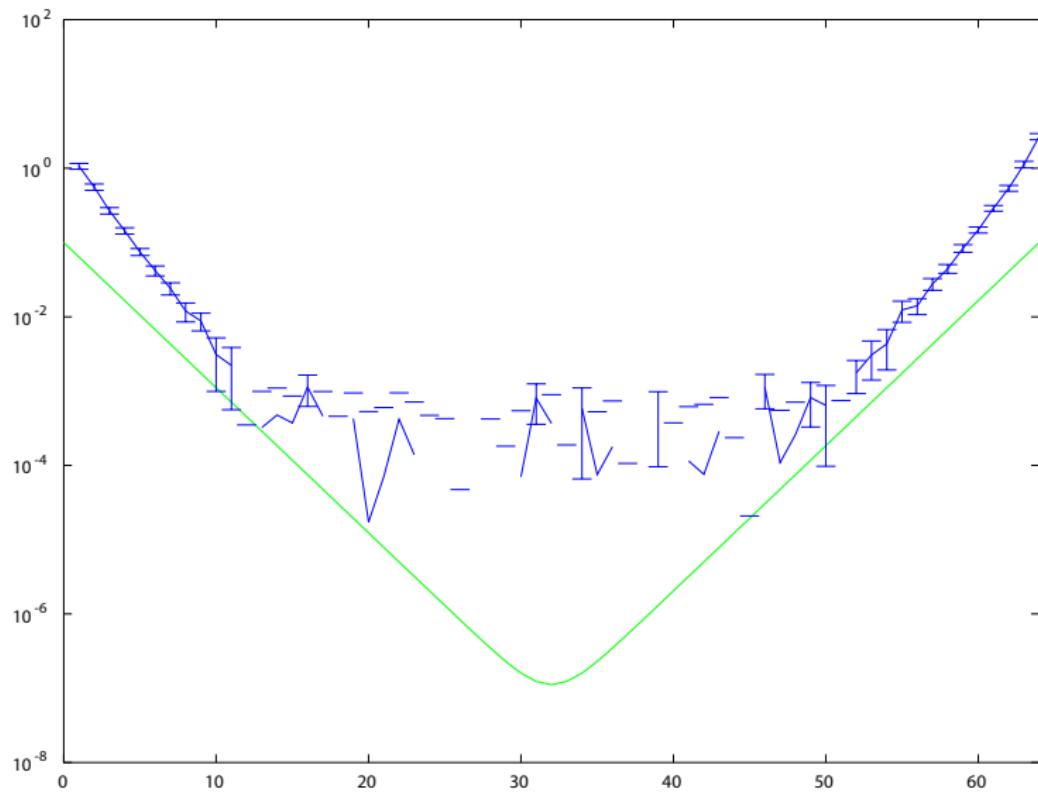
## Some practical notes:

- Use bootstrap if you can (more expensive though)
- Choose  $N_B$  as large as you can
- Do the **complete analysis** within the bootstrap
  - This does even include averaging over different analyses procedures for systematics etc.
  - Only exception are estimates of global ensemble properties like e.g. (co-)variances needed for fits within the bootstrap.
    - nesting bootstraps usually not necessary
- Not necessary if  $O$  is linear:  $\sigma_{JN} \equiv \sigma_{\text{naive}}$
- You may extract more information from distribution of  $O_i$ 
  - Confidence intervals, percentiles, etc.

# Rho propagator



# Rho propagator



# Fits with x-errors

A typical analysis situation:

- We have collected data at different bare quark masses
- We want to make a prediction at the physical point (for simplicity we ignore continuum and infinite volume)

How do we proceed?

- Define the physical point (e.g.  $M_\pi$ )
- Extrapolate target observable there

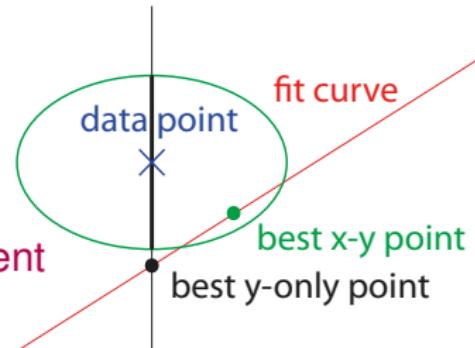
$M_\pi$  is not a parameter!

# X-errors

Fitting data with errors in the x-axis:

- add each x-value as a fit parameter
- constrain each x-value with measurement

Uncorrelated case:



$$\chi^2 \rightarrow \chi^2 + \sum_i (\textcolor{red}{x}_i - \textcolor{green}{p}_i)^2 / \sigma_{x_i}^2$$

Generalization with full covariance matrix

- ☞ Big covariance matrices lead to uncontrolled fits  
Mandatory to eliminate spurious correlations

# Correlated errors

Special case:  $x_i, y_i$  correlated, but uncorrelated with  $x_j, y_j$   $i \neq j$

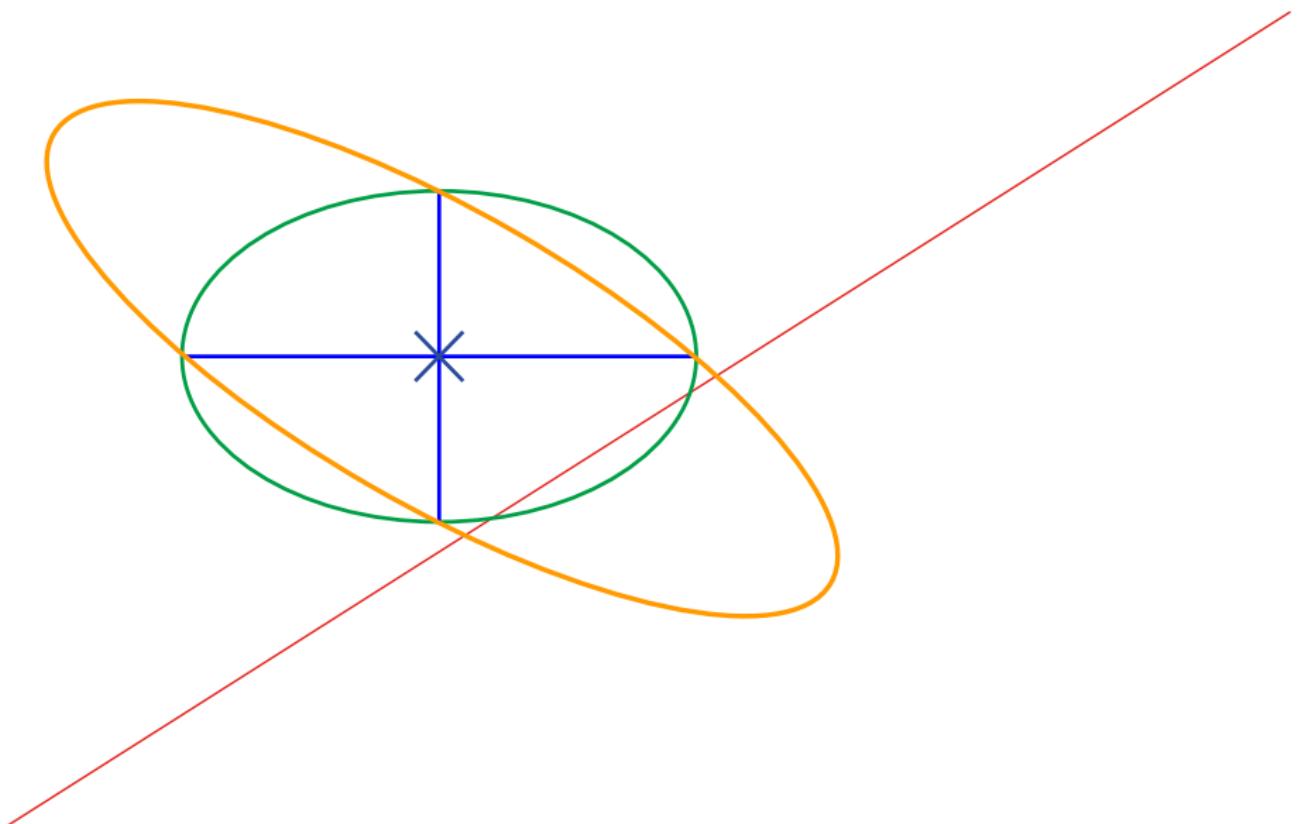
- Appears naturally in fit of independent ensembles
- Covariance matrix reduces to block diagonal form

Contribution to  $\chi^2$ :

$$\chi^2 \supset \chi_i^2 \begin{pmatrix} \Delta x & \Delta y \end{pmatrix} \begin{pmatrix} \Sigma_{xx}^{-1} & \Sigma_{xy}^{-1} \\ \Sigma_{xy}^{-1} & \Sigma_{yy}^{-1} \end{pmatrix} \begin{pmatrix} \Delta x \\ \Delta y \end{pmatrix}$$

- $\chi_i^2$  constant along an ellipse
- Covariance  $\Sigma_{xy}^{-1}$  tilts the axis
- ✓ Including x-errors can never increase  $\chi_i^2$
- ✓ Including x-errors does not change  $n$  (d.o.f.)

# Error ellipses



# General strategy

Sometimes subsets of data points are correlated

- 3 independent ensembles at each of 3 lattice spacings
- A measurement of each of the 3 lattice spacings  $a_i$

How do you extrapolate the observable  $M$  to the continuum?

- Form  $M = M_{\text{lat}}/a_i$  for each ensemble
- Error on  $M = M_{\text{lat}}/a_i$  is combination of error on  $M_{\text{lat}}$  and  $a_i$
- ✗ Introduces correlations between independent ensembles

# General strategy

Sometimes subsets of data points are correlated

- 3 independent ensembles at each of 3 lattice spacings
- A measurement of each of the 3 lattice spacings  $a_i$

How do you extrapolate the observable  $M$  to the continuum?

- Form  $M = M_{\text{lat}}/a_i$  for each ensemble
  - Error on  $M = M_{\text{lat}}/a_i$  error on  $M_{\text{lat}}$ , ignore  $a_i$
- ✗ Lattice spacing error not accounted for

# General strategy

Sometimes subsets of data points are correlated

- 3 independent ensembles at each of 3 lattice spacings
- A measurement of each of the 3 lattice spacings  $a_i$

How do you extrapolate the observable  $M$  to the continuum?

- Introduce a fit parameter  $\hat{a}_i$  for each lattice spacing
- Constrain  $\hat{a}_i$  with measurement
- Fit  $M_{\text{lat}} = M\hat{a}_i$  for each ensemble

# Combined fit quality

When doing your continuum/chiral/infinite volume fit

- Data points are often results of fits themselves
- How do you compute the quality of cascaded fits?

Theoretical ideal (not feasible):

- Do one big fit

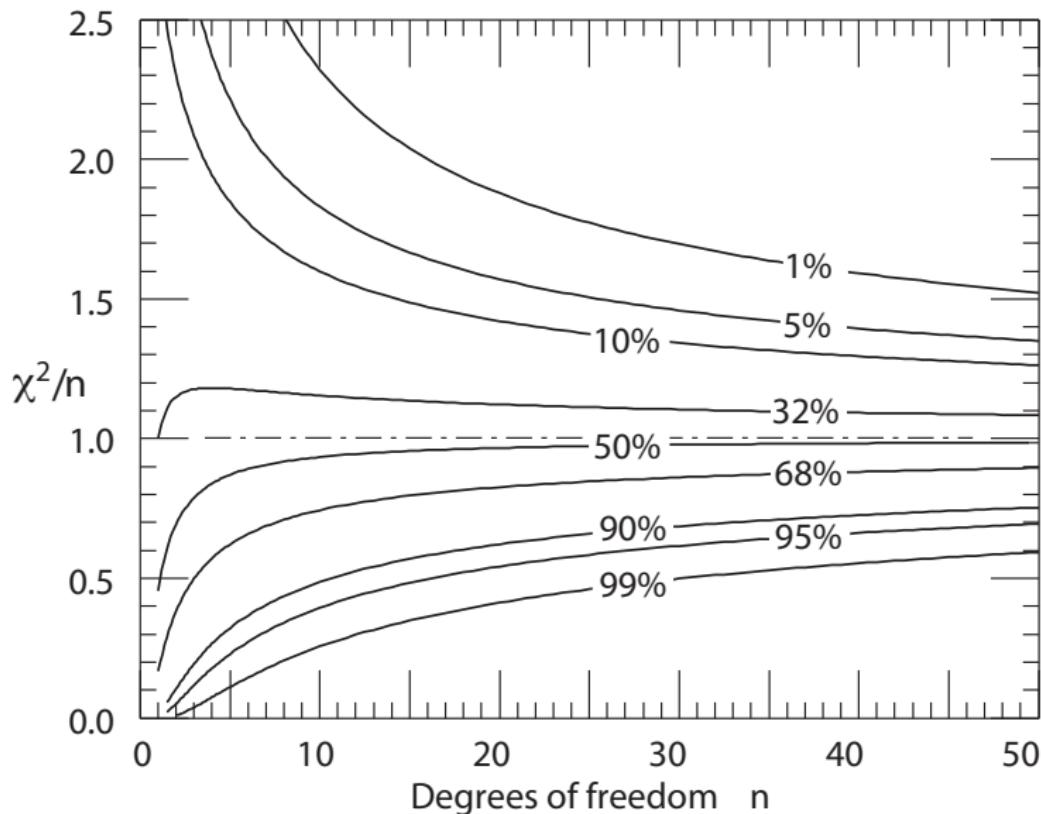
All original fits worked fully correlated:

- Sum  $\chi^2$  and d.o.f. of all fits  $\rightarrow Q$

Original fits not fully correlated:

- Treat data points as input, just compute  $Q$  of final fit

# Fit quality



(PDG, 2012)

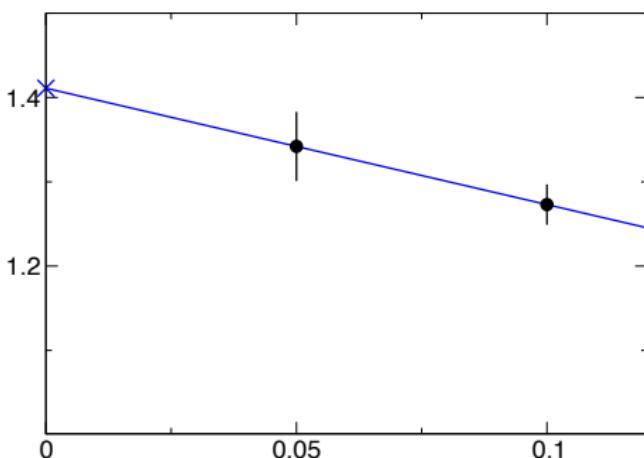
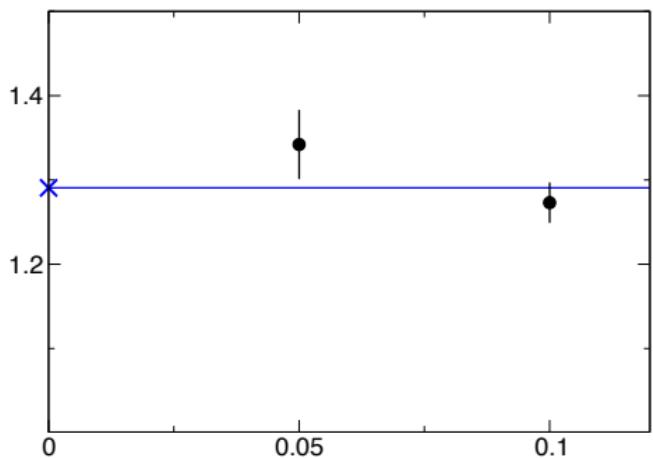
# Which fit is better?

The following slides compare 2 fits each

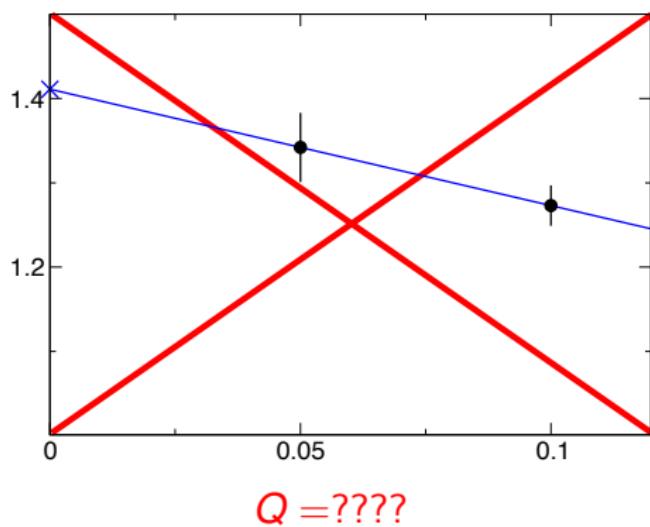
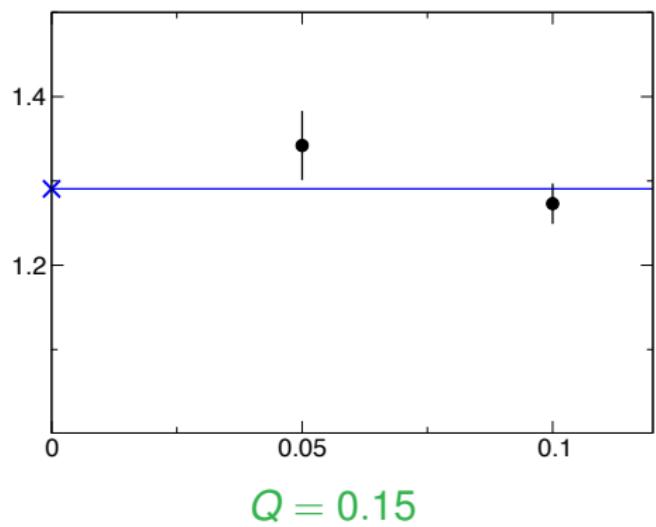
All data are uncorrelated

Which fit can be trusted more?

# Which fit is better?

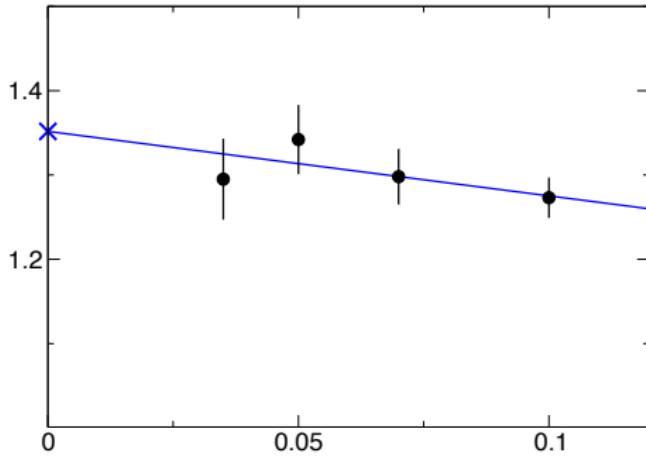
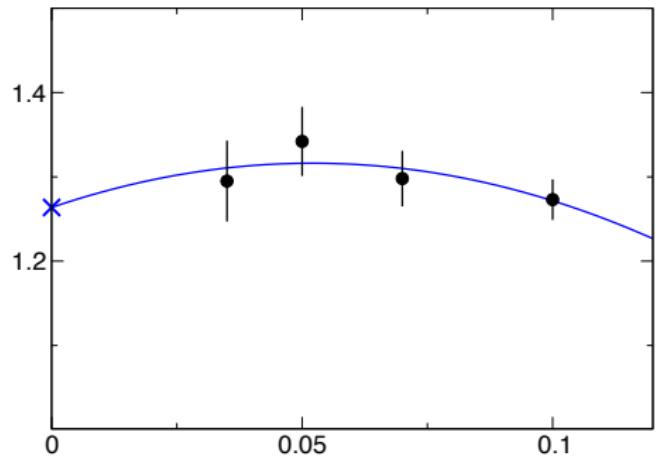


# Which fit is better?

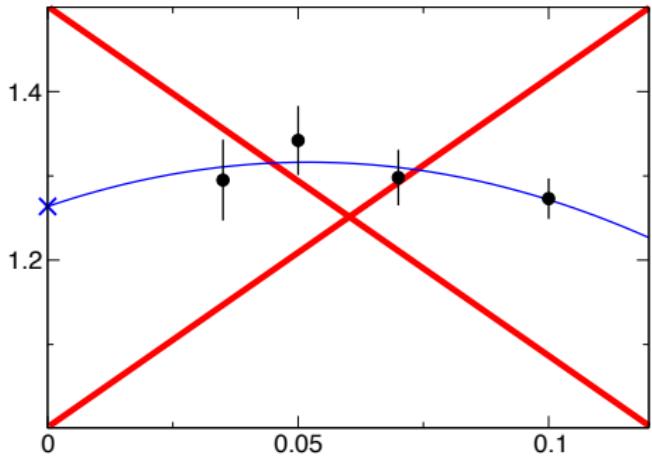


Never leave 0 d.o.f., you loose control over fit quality

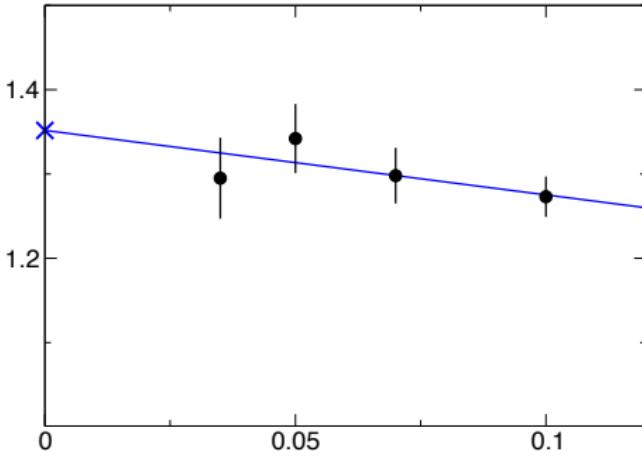
# Which fit is better?



# Which fit is better?



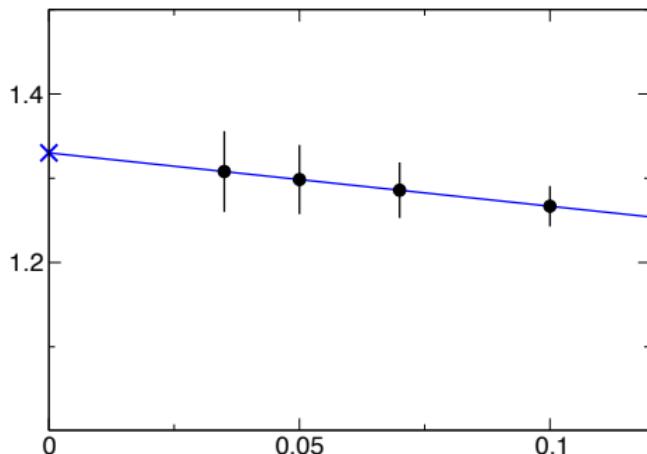
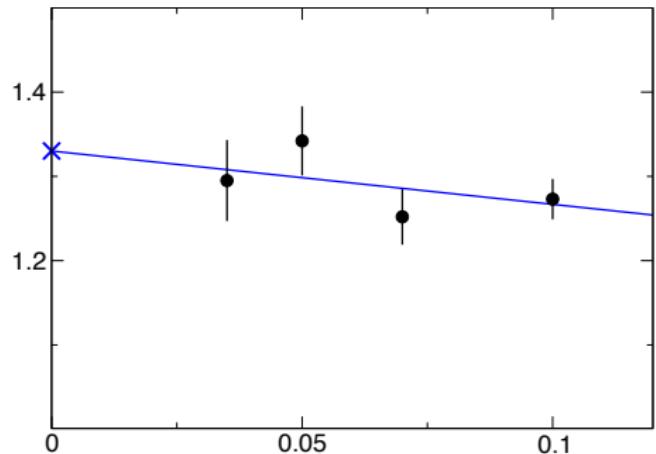
$$Q = 0.42$$



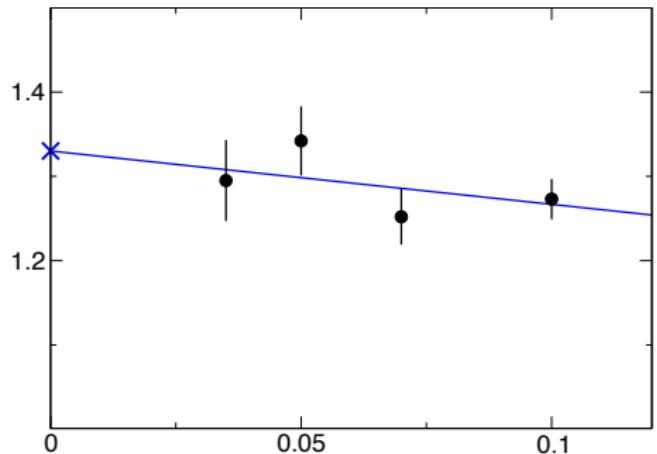
$$Q = 0.64$$

- Do not try to extract too much from the data
- The displayed data have no sensitivity towards a curvature term. It is compatible with 0.

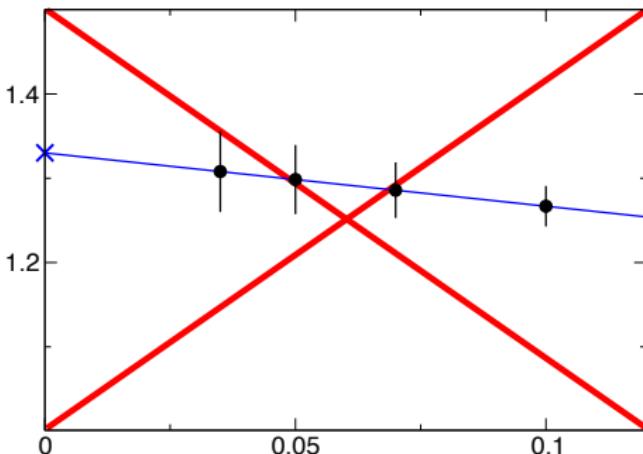
# Which fit is better?



# Which fit is better?



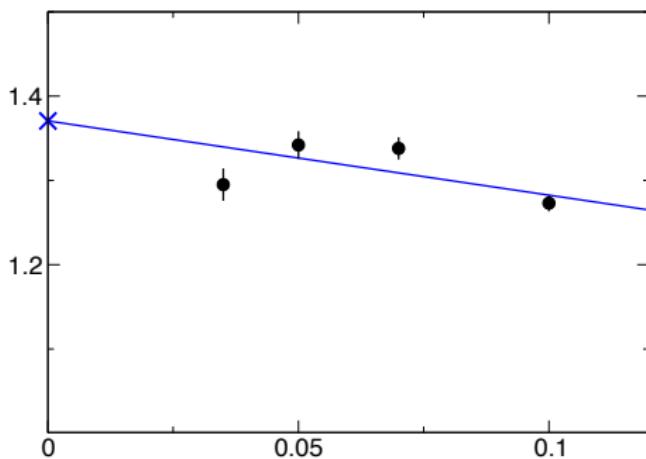
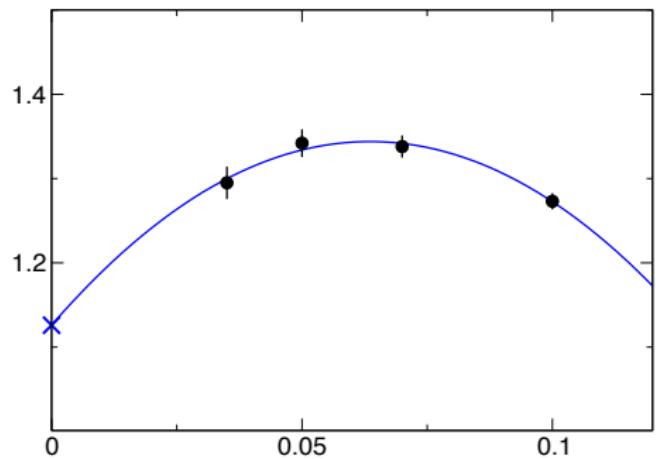
$$Q = 0.31$$



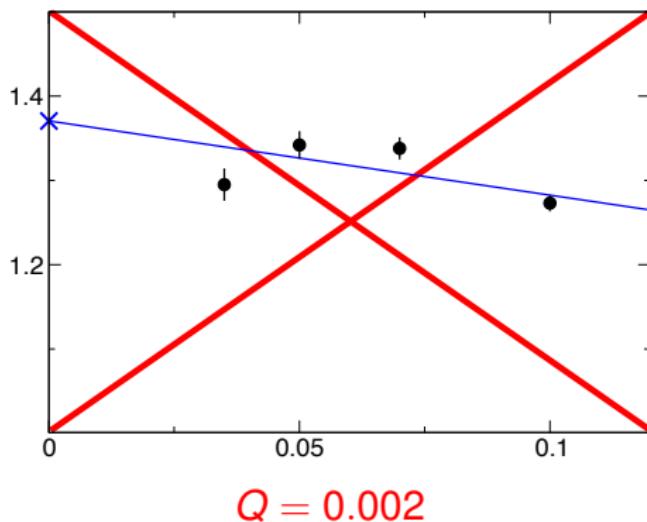
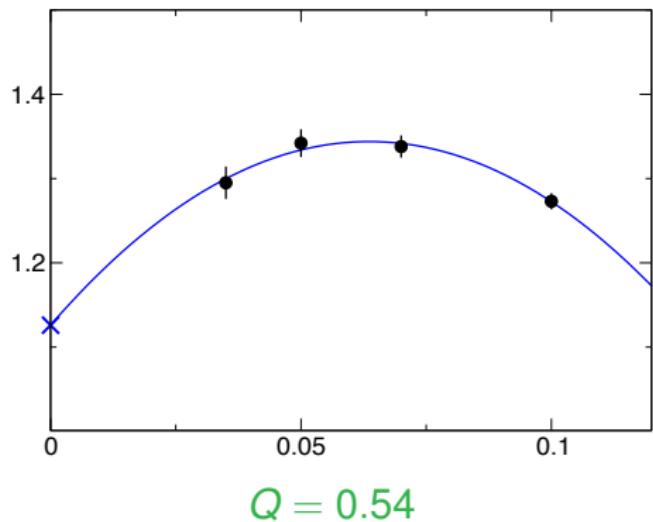
$$Q = 1.00$$

- $1 - Q = 8 \times 10^{-13} \rightarrow$  winning the lottery is more probable than having a result this good by chance
- Data are suspicious (unrecognized correlation)

# Which fit is better?



# Which fit is better?



Linear modell is not sufficient for these data

# Practical hints

Some hints for numerically minimizing a complex  $\chi^2$  function

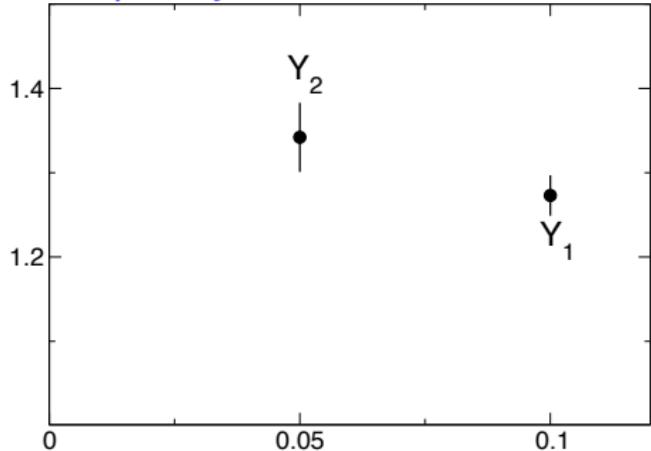
- Give reasonable starting values
  - Solver might find a wrong minimum or crash
- Build up your fit parameter by parameter
  - Start with all but the most relevant parameters constrained
  - Minimize the constrained fit first
  - When it has converged, free one more parameter
- Check pulls and bootstrap samples for outliers
  - A good fit can identify problematic input data
- Always look at the fit to check it does fit the data

# Systematics

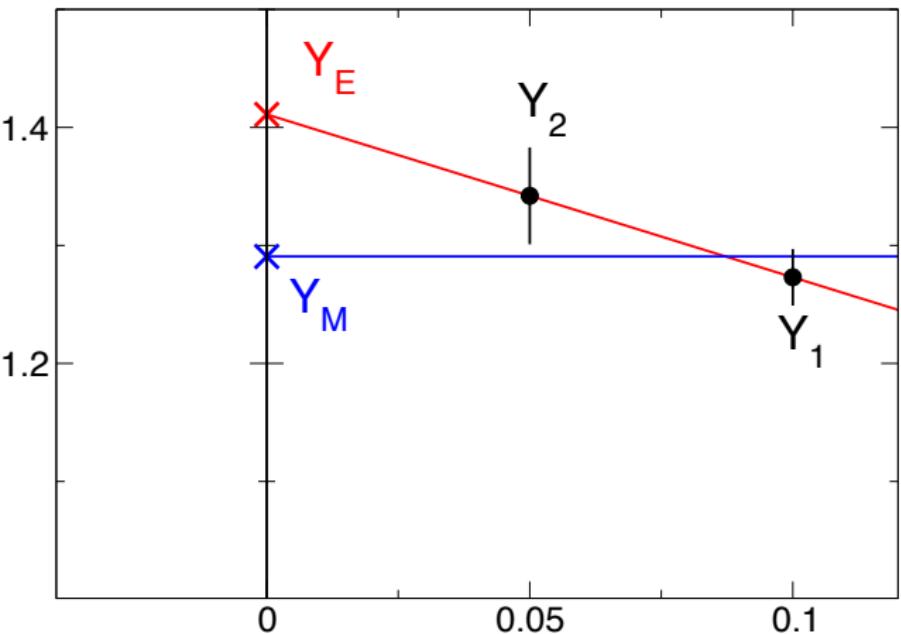
How do we compute the systematic error?

- We don't
- Systematics can only be estimated
- There is no single correct procedure

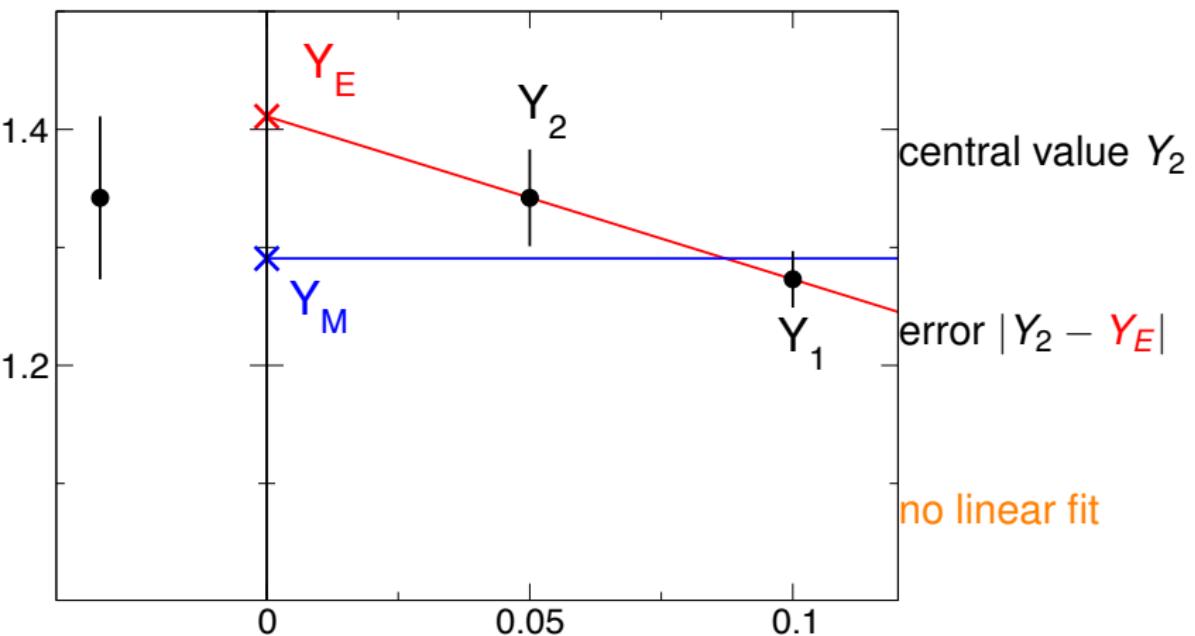
Example: systematic error of  $x \rightarrow 0$



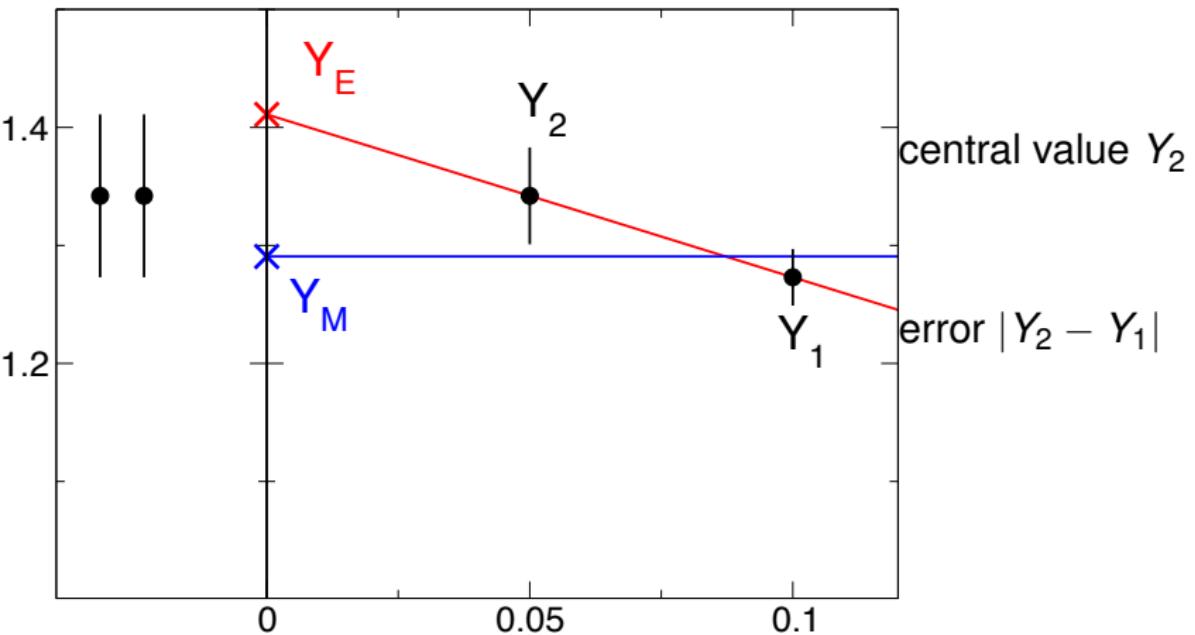
# Simple estimates



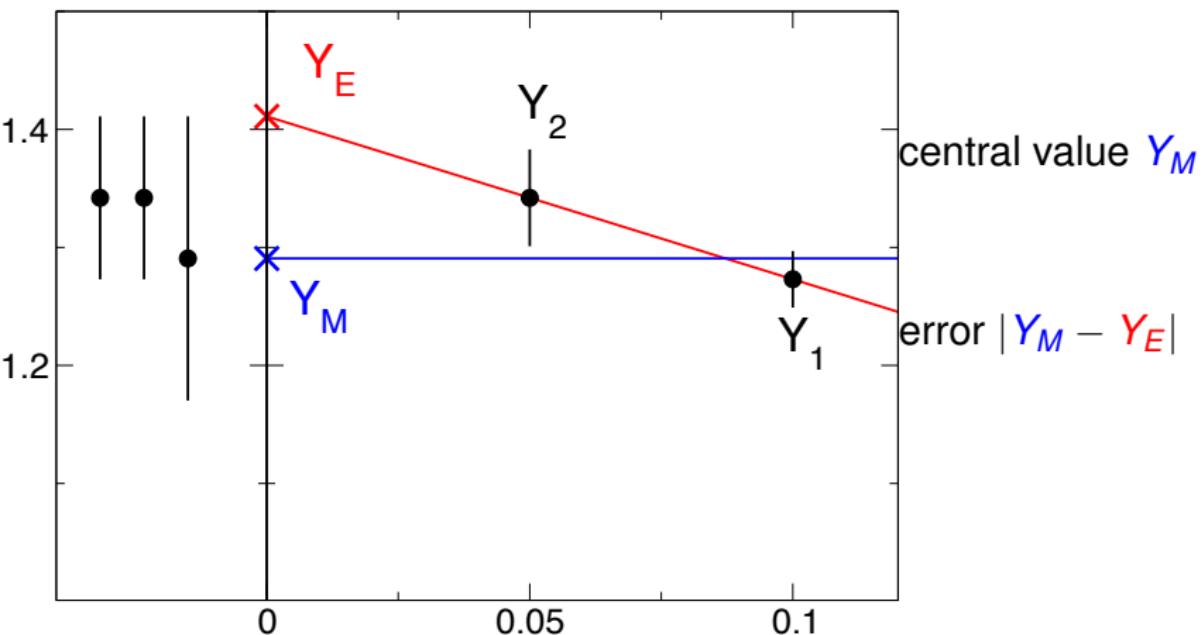
# Simple estimates



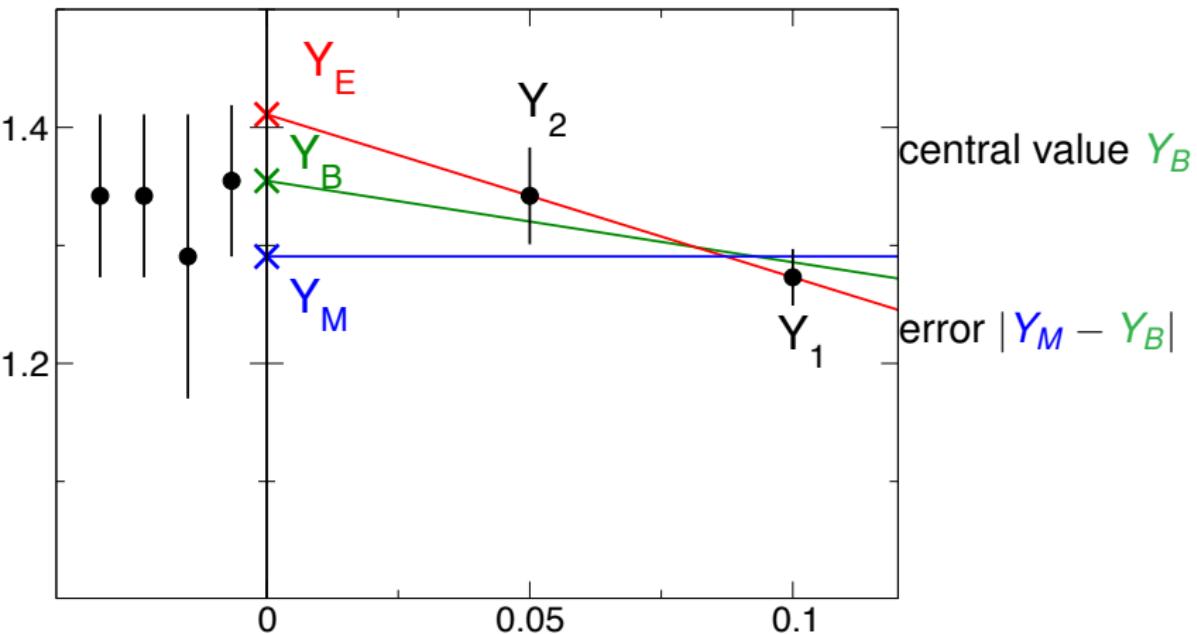
# Simple estimates



# Simple estimates



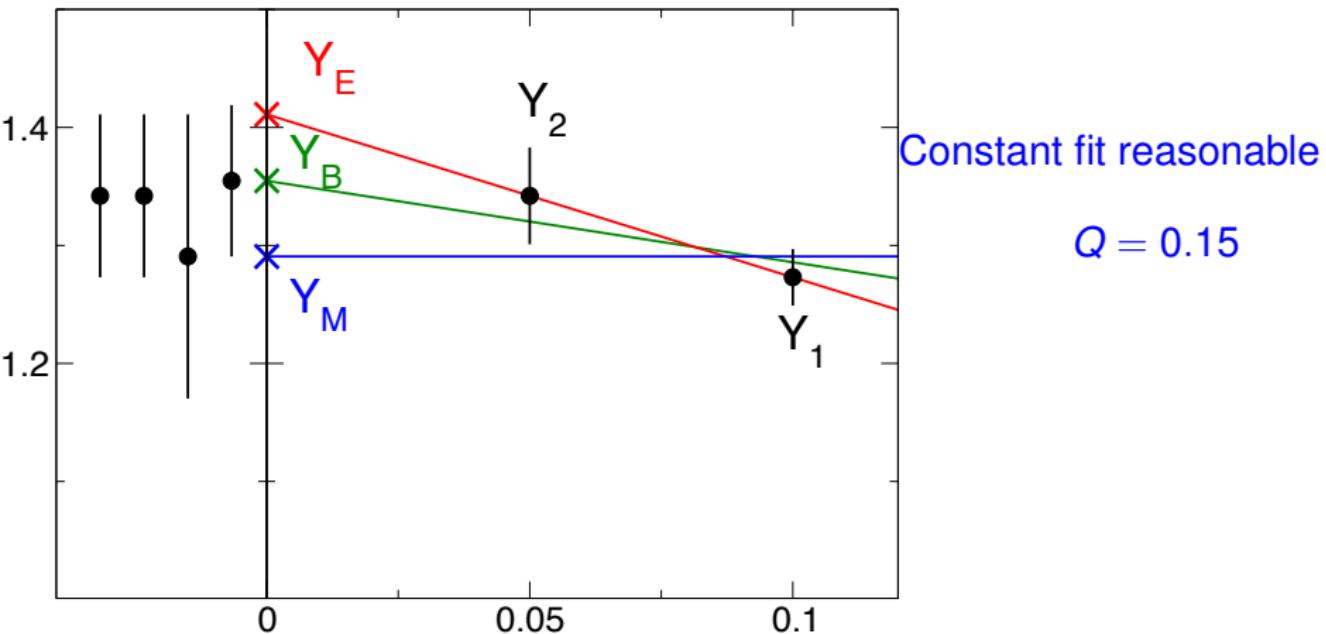
# Simple estimates



You can do a linear fit if you have prior knowledge on the slope

☞ Constraint on slope is an additional data point

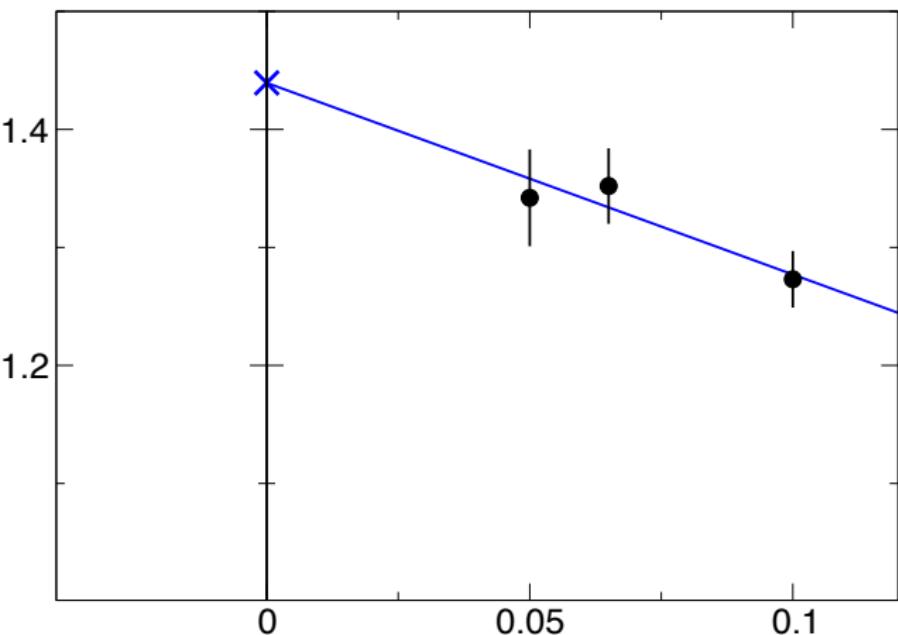
# Simple estimates



These are estimates for what systematics?

☞ Neglecting first order (linear) corrections to constant

# Simple estimates



One more data point: error on linear term is now statistical

☞ Now we need to estimate systematic due to higher orders

# Systematics

One conservative strategy for systematics:

- Identify all higher order effects you have to neglect
- For each of them:
  - Repeat the entire analysis treating this one effect differently
  - Add the spread of results to systematics
- Important:
  - Do not do suboptimal analyses
  - Do not double-count analyses

make sure there are no unknown unknowns

# Combining results

How to determine the spread of results?

- Stdev or  $1\sigma$  confidence interval of results
- Can weight it with fit quality  $Q$

Information theoretic optimum: Akaike Information Criterion

- Information content of a fit depends on how well data are described per fit parameter
- Information lost wrt. correct fit  $\propto$  cross-entropy  $J$
- Compute information cross-entropy  $J_m$  of each fit  $m$
- Probability that fit is correct  $\propto e^{J_m}$

# Akaike Information Criterion

- $N$  measurements  $\Gamma_i$  from unknown pdf  $g(\Gamma)$
- Fit model  $f(\Gamma|\Theta)$  with parameters  $\Theta$
- Cross-entropy ( $\sim$  Kullback-Leibler divergence)

$$J_m = J(g, f_m[\Theta]) = \int d\Gamma g(\Gamma) \ln(f(\Gamma|\Theta))$$

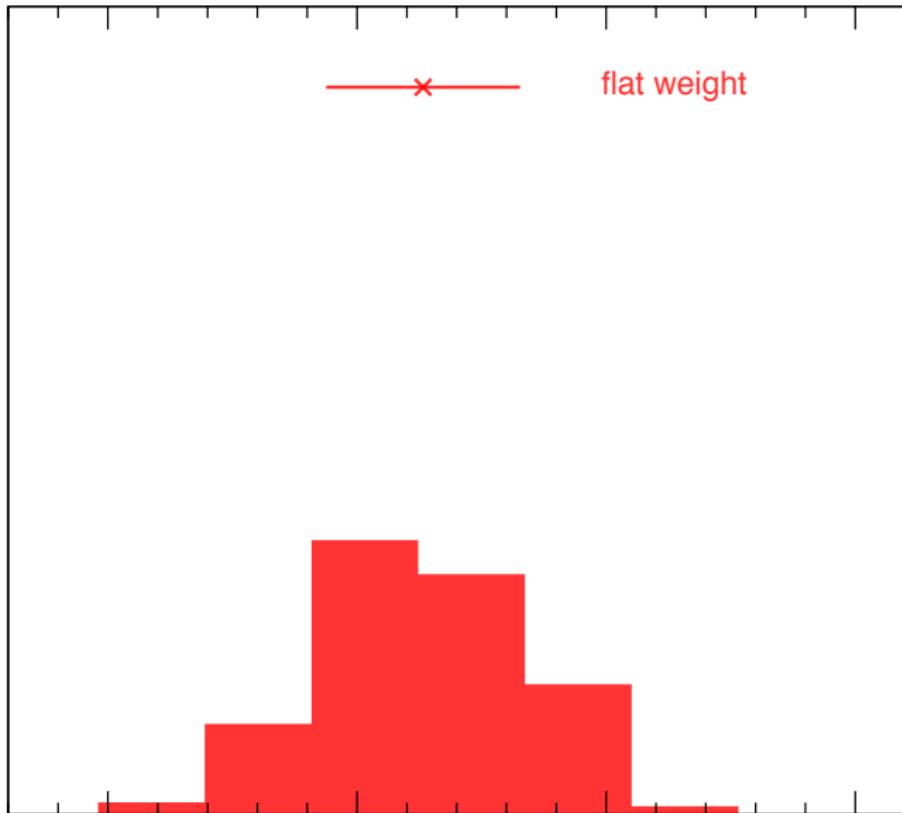
- For  $N \rightarrow \infty$  and  $f$  close to  $g$ :

$$J_m = -\frac{\chi_m^2}{2} - p_m$$

where  $p_m$  is the number of fit parameters

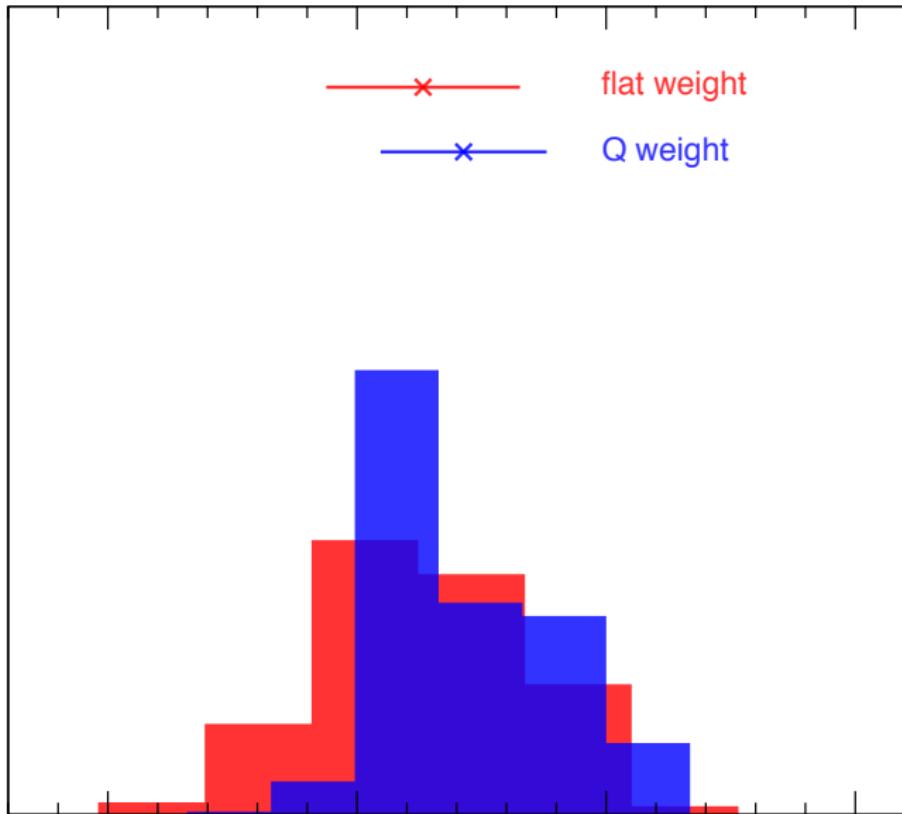
Is this the only correct method?

# Combining results



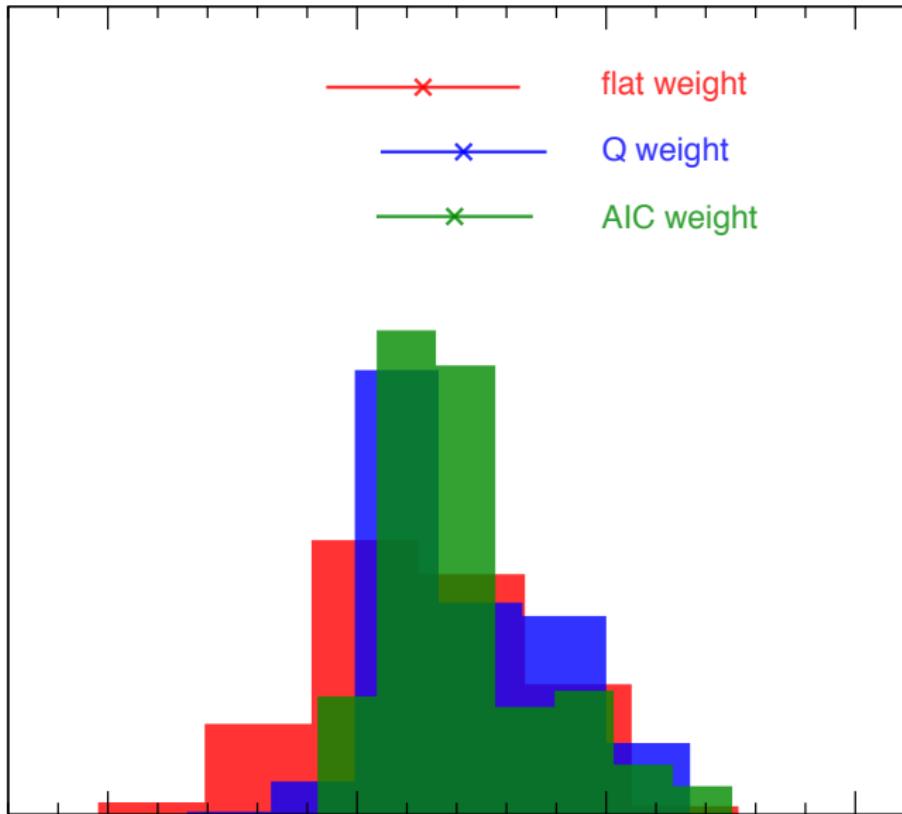
- AIC suppresses strongly
- Other weights more conservative
- Agreement is excellent crosscheck

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Let's practice!