

## **Exercise/Hands-on/Lesson #9**

### **Scientific Data Analysis Lab course**

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(hands-on part)

- **Extended vs. non-extended ML fit : comparison**
- **Access to Maximum Likelihood fit information in RooFit**

(theory part)

**Profile Likelihood & MINOS**

# Bibliography

Inspired by part of the theory visualization & exercises by **Wouter Verkerke** :

<https://indico.cern.ch/event/72320/contributions/2082589/attachments/1037201/1478048/roofit-intro-roostats-v11a.pdf>

[https://indico.cern.ch/event/305391/contributions/701304/attachments/580262/798889/Verkerke\\_Statistics\\_L2.pdf](https://indico.cern.ch/event/305391/contributions/701304/attachments/580262/798889/Verkerke_Statistics_L2.pdf)

See also :

- his slides for the Ferrara School 2009: <https://www.nikhef.nl/~verkerke/ferrara>)

- his slides for IN2P3 School 2014: [https://indico.in2p3.fr/event/9742/contributions/50419/attachments/40828/50594/sos2014\\_systprof\\_v38.pdf](https://indico.in2p3.fr/event/9742/contributions/50419/attachments/40828/50594/sos2014_systprof_v38.pdf)

Of course a good reference book is : **Luca Lista, Statistical methods for Data Analysis in Particle Physics**, Springer, 2<sup>nd</sup> Ed.

In this Lab exercise we use CMS data used for the paper

<https://doi.org/10.1016/j.physletb.2014.05.055>

Physics Letters B 734 (2014) 261–281

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Physics Letters B

[www.elsevier.com/locate/physletb](http://www.elsevier.com/locate/physletb)

Observation of a peaking structure in the  $J/\psi\phi$  mass spectrum from  $B^\pm \rightarrow J/\psi\phi K^\pm$  decays

CMS Collaboration \*

CERN, Switzerland

## Retrieve binned data and plot

With reference to the code in the macro `yield.C` ...

- Get the histogram of the  $J/\psi(\mu\mu)\phi(KK)K$  invariant mass (the signal represents the 3-body decay  $B^\pm \rightarrow J/\psi \phi K^\pm$ ):

```
TFile f1("DatasetAandB_KaonTrackRefit_Bwin_new_21aug13.root", "READ");
TH1D *hist = (TH1D*)f1.Get("myJpsiKKKmass_all");
```

- Declare & initialize the variable to represent the invariant mass and prepare the corresponding RooPlot pointer:

```
RooRealVar y("y", "m(J/#psi #phi K)[GeV]", 5.15, 5.45);
RooPlot* yframe = y.frame("");
```

- Import the binned data by creating the RooDataHist object from the histogram and plot it:

```
RooDataHist BmassExt(hist->GetName(), hist->GetTitle(), RooArgSet(y), RooFit::Import(*hist, kFALSE));
BmassExt.plotOn(yframe);
//myC->cd(); // uncomment to plot
//yframe->Draw(); // uncomment to plot
```

## Build the negative log-likelihood (nll)

Build the model: - a gaussian for the signal (2 parameters: mass and width) ;  
- a Chebyshev of 2<sup>nd</sup> order (2 parameters) for the background.

Based on these two PDFs, build the full PDF to make an **extended fit**:

```
RooRealVar nsig("nsig","n. of signal cands",2500.,2000.,3800.);
RooRealVar nbkg("nbkg","n. of bkg cands",2000.,0.,200000.);
//
RooAddPdf model_extended("model_extended","gauss+cheby EXT",RooArgList(gausse,chebye),RooArgList(nsig,nbkg));
```

Create a function object that represents the negative-log-likelihood (nll) ...

... by using the method **RooAbsPdf::createNLL(RooAbsData&)**; the returned object is of type **RooAbsReal\***

```
RooAbsReal* nll = model_extended.createNLL(BmassExt);
```

In this way we explicitly constructed the likelihood (function of PDF/data combination) that **can be used as any RooFit function object**.

**Note:** likelihood can be created by a calculation that can be parallelized (suppose for instance on 4 cores):

```
RooAbsReal* nll = model_extended.createNLL(BmassExt, NumCPU(4));
```

# MINUIT session

Let us invoke **MINUIT** to perform the binned extended fit.

First we can create a **MINUIT** minimizer object:

```
Roofit m(*nll);
```

Calling `MIGRAD` we get the **central values** (*best estimates*) for the parameters when convergence is reached:

```
m.migrad();
```



```
MIGRAD FAILS TO FIND IMPROVEMENT
MIGRAD MINIMIZATION HAS CONVERGED.
MIGRAD WILL VERIFY CONVERGENCE AND ERROR MATRIX.
COVARIANCE MATRIX CALCULATED SUCCESSFULLY
FCN=-1.15354e+06 FROM MIGRAD      STATUS=CONVERGED      532 CALLS      533 TOTAL
                                EDM=0.00035446      STRATEGY= 1      ERROR MATRIX ACCURATE
EXT PARAMETER
NO.  NAME      VALUE      ERROR      STEP      FIRST
     NAME      VALUE      ERROR      SIZE      DERIVATIVE
  1  c0e      -2.25841e-01  5.55732e-03  4.09669e-06  -6.98549e+02
  2  c1e      -1.08452e-02  6.02446e-03  4.04505e-06  1.24095e+03
  3  mge      5.27943e+00  5.31398e-04  7.92138e-02  6.65397e-02
  4  nbkg      9.55524e+04  3.48579e+02  2.32333e-03  6.58743e-01
  5  nsig      2.92321e+03  1.70487e+02  9.92199e-02  -9.14745e-03
  6  wge      9.48373e-03  5.96383e-04  6.96549e-02  1.25538e-01
                                ERR DEF= 0.5
EXTERNAL ERROR MATRIX.      NDIM= 25      NPAR= 6      ERR DEF=0.5
  3.088e-05  3.173e-08  1.521e-07  -8.477e-02  8.560e-02  2.114e-07
  3.173e-08  3.629e-05  -6.934e-08  -4.304e-01  4.346e-01  1.017e-06
  1.521e-07  -6.934e-08  2.835e-07  -2.503e-03  2.542e-03  1.914e-08
 -8.477e-02  -4.304e-01  -2.503e-03  1.215e+05  -2.620e+04  -6.305e-02
  8.560e-02  4.346e-01  2.542e-03  -2.620e+04  2.942e+04  6.383e-02
  2.114e-07  1.017e-06  1.914e-08  -6.305e-02  6.383e-02  3.574e-07
PARAMETER CORRELATION COEFFICIENTS
NO.  GLOBAL      1      2      3      4      5      6
  1  0.10984  1.000  0.001  0.051 -0.044  0.090  0.064
  2  0.42489  0.001  1.000 -0.022 -0.205  0.421  0.282
  3  0.08630  0.051 -0.022  1.000 -0.013  0.028  0.060
  4  0.44039 -0.044 -0.205 -0.013  1.000 -0.438 -0.303
  5  0.71287  0.090  0.421  0.028 -0.438  1.000  0.622
  6  0.62527  0.064  0.282  0.060 -0.303  0.622  1.000
```

# Parabolic uncertainties

To recalculate the errors and the covariance matrix in an accurate way (still in parabolic assumption) we use HESSE, while central values (by Migrad) are conserved.

m.hesse ( ) ;



```
*****
** 18 **HESSE      3000
*****
COVARIANCE MATRIX CALCULATED SUCCESSFULLY
FCN=-1.15354e+06 FROM HESSE      STATUS=OK      40 CALLS      573 TOTAL
EDM=0.000362648      STRATEGY= 1      ERROR MATRIX ACCURATE

EXT PARAMETER
NO.  NAME      VALUE      ERROR      INTERNAL      INTERNAL
      NAME      VALUE      ERROR      STEP SIZE      VALUE
1  c0e      -2.25841e-01  5.55902e-03  1.63867e-07  -2.25841e-04
2  c1e      -1.08452e-02  6.05944e-03  1.61802e-07  -1.08452e-05
3  mge      5.27943e+00  5.30310e-04  3.16855e-03  9.53869e+00
4  nbkg      9.55524e+04  3.51136e+02  9.29332e-05  -4.44908e-02
5  nsig      2.92321e+03  1.74070e+02  3.96880e-03  -6.25739e+00
6  wge      9.48373e-03  6.08532e-04  2.78620e-03  2.50293e+01

ERR DEF= 0.5
EXTERNAL ERROR MATRIX,      NDIM= 25      NPAR= 6      ERR DEF=0.5
3.090e-05  6.228e-08  1.532e-07  -8.983e-02  8.986e-02  2.249e-07
6.228e-08  3.672e-05  -6.359e-08  -4.579e-01  4.580e-01  1.106e-06
1.532e-07  -6.359e-08  2.823e-07  -2.776e-03  2.778e-03  1.440e-08
-8.983e-02  -4.579e-01  -2.776e-03  1.233e+05  -2.775e+04  -6.864e-02
8.986e-02  4.580e-01  2.778e-03  -2.775e+04  3.069e+04  6.867e-02
2.249e-07  1.106e-06  1.440e-08  -6.864e-02  6.867e-02  3.722e-07
PARAMETER CORRELATION COEFFICIENTS
NO.  GLOBAL      1      2      3      4      5      6
1  0.11253  1.000  0.002  0.052 -0.046  0.092  0.066
2  0.43584  0.002  1.000 -0.020 -0.215  0.431  0.299
3  0.07496  0.052 -0.020  1.000 -0.015  0.030  0.044
4  0.45348 -0.046 -0.215 -0.015  1.000 -0.451 -0.320
5  0.72800  0.092  0.431  0.030 -0.451  1.000  0.643
6  0.64446  0.066  0.299  0.044 -0.320  0.643  1.000
```

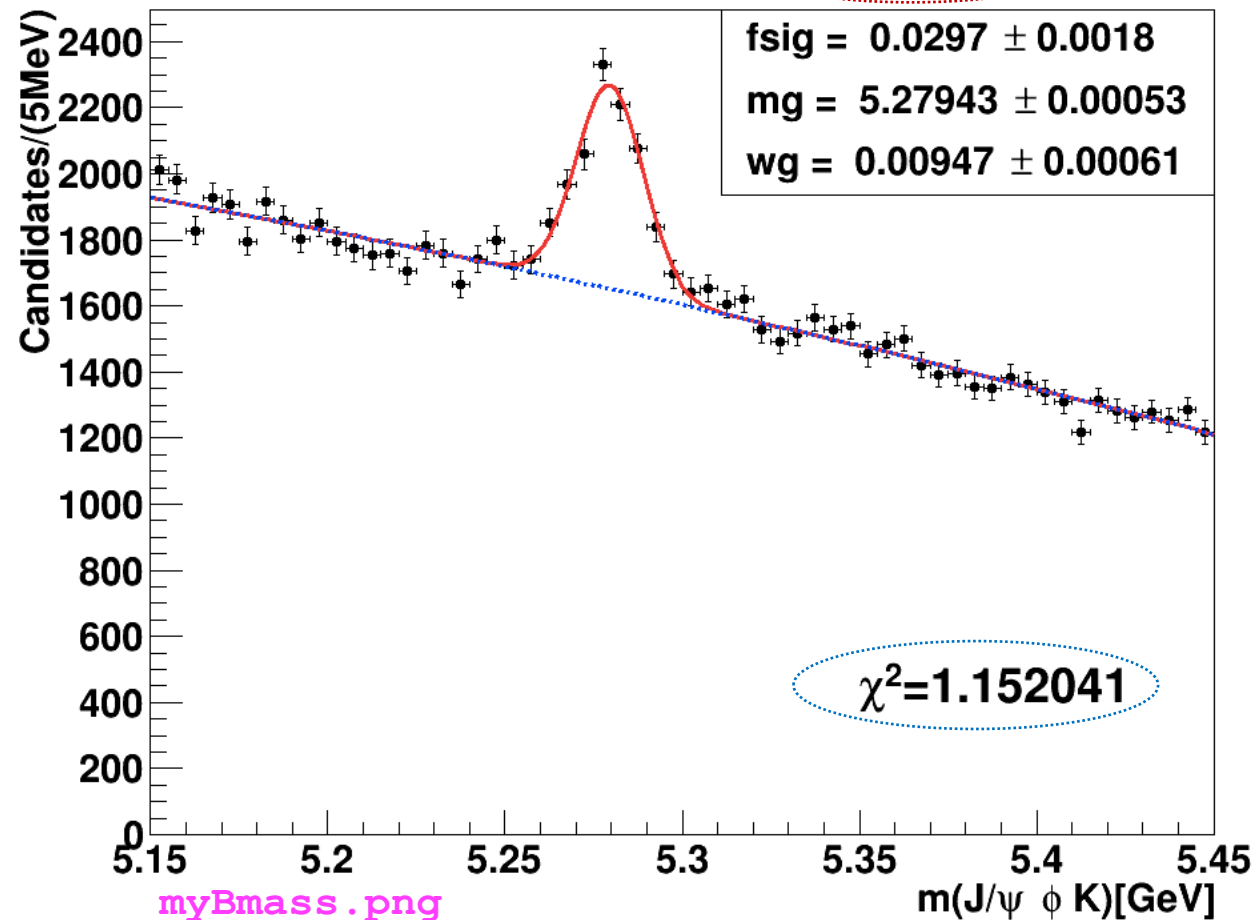
# Extended vs not-extended fits: a comparison - I

Not extended fit : just fsig

$$n_{sig} = 2921.3 \pm 174.9$$

$$\begin{aligned} f_{sig} &= 0.0297 \pm 0.0018 \\ m_g &= 5.27943 \pm 0.00053 \\ w_g &= 0.00947 \pm 0.00061 \end{aligned}$$

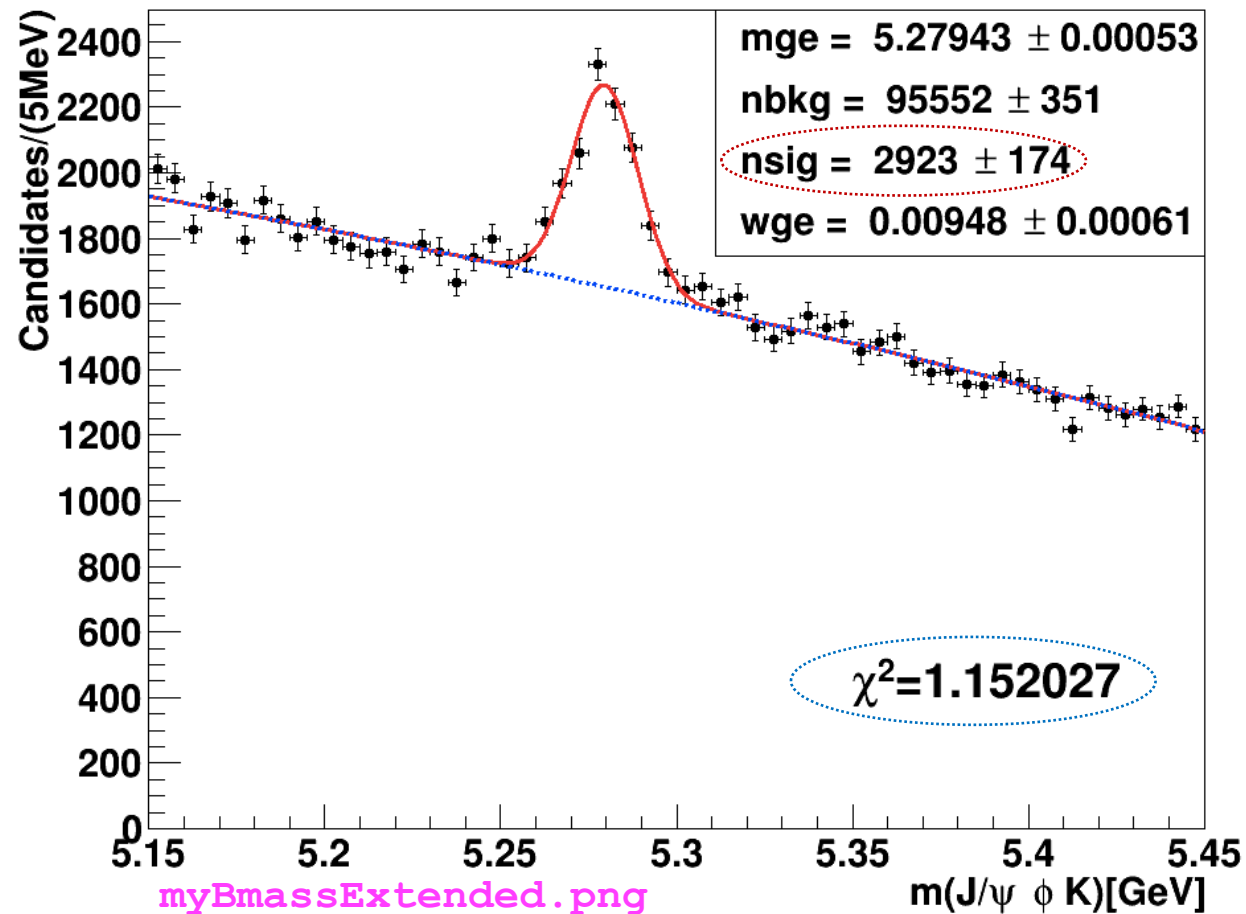
$$\chi^2 = 1.152041$$



Extended fit : nsig and nbkg

$$\begin{aligned} m_{ge} &= 5.27943 \pm 0.00053 \\ n_{bkg} &= 95552 \pm 351 \\ n_{sig} &= 2923 \pm 174 \\ w_{ge} &= 0.00948 \pm 0.00061 \end{aligned}$$

$$\chi^2 = 1.152027$$



Difference can be hardly appreciated: mass and width are about identical (see also next slide)! Also they have very similar  $\tilde{\chi}_{fit}^2$   
Extended fit has the advantage to provide directly as output also the number of B+ candidates ( $n_{sig}$ )

## Extended vs not-extended fits: a comparison - II

### NOT-extended

```
*****
** 9 **HESSE      2500
*****
COVARIANCE MATRIX CALCULATED SUCCESSFULLY
FCN=-119793 FROM HESSE      STATUS=OK      31 CALLS      359 TOTAL
EDM=1.33432e-05  STRATEGY= 1  ERROR MATRIX ACCURATE

EXT PARAMETER
NO.  NAME      VALUE      ERROR      INTERNAL  INTERNAL
STEP SIZE  VALUE
1  c0      -2.25828e-01  5.55891e-03  2.64024e-07  -2.25828e-04
2  c1      -1.09184e-02  6.05941e-03  5.21376e-08  -1.09184e-05
3  fsig     2.96640e-02  1.77569e-03  7.21237e-05  -1.91699e+00
4  mg       5.27943e+00  5.30028e-04  1.01863e-03  -2.85771e+02
5  wg       9.47402e-03  6.07750e-04  8.94182e-04  9.41424e+01
-----
ERR DEF= 0,5
EXTERNAL ERROR MATRIX.  NDIM= 25  NPAR= 5  ERR DEF=0,5
3.090e-05  6.145e-08  -9.119e-07  1.530e-07  2.245e-07
6.145e-08  3.672e-05  -4.649e-06  -6.319e-08  1.105e-06
-9.119e-07  -4.649e-06  3.153e-06  -2.829e-08  -6.963e-07
1.530e-07  -6.319e-08  -2.829e-08  2.820e-07  1.421e-08
2.245e-07  1.105e-06  -6.963e-07  1.421e-08  3.712e-07
PARAMETER CORRELATION COEFFICIENTS
NO.  GLOBAL  1  2  3  4  5
1  0.11250  1.000  0.002 -0.092  0.052  0.066
2  0.43581  0.002  1.000 -0.432 -0.020  0.299
3  0.69297 -0.092 -0.432  1.000 -0.030 -0.644
4  0.07459  0.052 -0.020 -0.030  1.000  0.044
5  0.64456  0.066  0.299 -0.644  0.044  1.000
-----
```

$$m = (5279,43 \pm 0.53)MeV$$

$$\sigma = (9,4740 \pm 0,6078)MeV$$

### Extended

```
** 18 **HESSE      3000
*****
COVARIANCE MATRIX CALCULATED SUCCESSFULLY
FCN=-1.15354e+06 FROM HESSE  STATUS=OK      40 CALLS      573 TOTAL
EDM=0.000362648  STRATEGY= 1  ERROR MATRIX ACCURATE

EXT PARAMETER
NO.  NAME      VALUE      ERROR      INTERNAL  INTERNAL
STEP SIZE  VALUE
1  c0e      -2.25841e-01  5.55902e-03  1.63867e-07  -2.25841e-04
2  c1e      -1.08452e-02  6.05944e-03  1.61802e-07  -1.08452e-05
3  mge      5.27943e+00  5.30310e-04  3.16855e-03  9.53869e+00
4  nbkg     9.55524e+04  3.51136e+02  9.29332e-05  -4.44908e-02
5  nsig     2.92321e+03  1.74070e+02  3.96880e-03  -6.25739e+00
6  wge      9.48373e-03  6.08532e-04  2.78620e-03  2.50293e+01
-----
ERR DEF= 0,5
EXTERNAL ERROR MATRIX.  NDIM= 25  NPAR= 6  ERR DEF=0,5
3.090e-05  6.228e-08  1.532e-07  -8.983e-02  8.986e-02  2.249e-07
6.228e-08  3.672e-05  -6.359e-08  -4.579e-01  4.580e-01  1.106e-06
1.532e-07  -6.359e-08  2.823e-07  -2.776e-03  2.778e-03  1.440e-08
-8.983e-02  -4.579e-01  -2.776e-03  1.233e+05  -2.775e+04  -6.864e-02
8.986e-02  4.580e-01  2.778e-03  -2.775e+04  3.069e+04  6.867e-02
2.249e-07  1.106e-06  1.440e-08  -6.864e-02  6.867e-02  3.722e-07
PARAMETER CORRELATION COEFFICIENTS
NO.  GLOBAL  1  2  3  4  5  6
1  0.11253  1.000  0.002  0.052 -0.046  0.092  0.066
2  0.43584  0.002  1.000 -0.020 -0.215  0.431  0.299
3  0.07496  0.052 -0.020  1.000 -0.015  0.030  0.044
4  0.45348 -0.046 -0.215 -0.015  1.000 -0.451 -0.320
5  0.72800  0.092  0.431  0.030 -0.451  1.000  0.643
6  0.64446  0.066  0.299  0.044 -0.320  0.643  1.000
```

$$m = (5279,43 \pm 0.53)MeV$$

$$\sigma = (9,4837 \pm 0,6085)MeV$$



# Asymmetric uncertainties

To get asymmetric error (central values and parabolic are the same) for a specific parameter, like `nsig`:

`m.minos (nsig) ;` 

To additionally print the result just do: `nsig.Print() ;` 

```
*****
** 23 **MINOS      3000      5
*****
FCN=-1.15354e+06 FROM MINOS  STATUS=SUCCESSFUL  132 CALLS      705 TOTAL
                    EDM=0.000362648  STRATEGY= 1  ERROR MATRIX ACCURATE
EXT PARAMETER
NO.  NAME      VALUE      PARABOLIC  MINOS ERRORS
      NAME      VALUE      ERROR      NEGATIVE   POSITIVE
 1  c0e      -2.25841e-01  5.55902e-03
 2  c1e      -1.08452e-02  6.05944e-03
 3  mge       5.27943e+00  5.30310e-04
 4  nbkg      9.55524e+04  3.51136e+02
 5  nsig      2.92321e+03  1.74070e+02 -1.74275e+02  1.76453e+02
 6  wge       9.48373e-03  6.08532e-04
ERR DEF= 0.5
RooRealVar::nsig = 2923.21 +/- (-174.275,176.453) L(2000 - 3800)
```

To get asymmetric error for all the parameters :

`m.minos () ;` 

Note: asymmetric errors can slightly change if you execute MINOS for 1 or all parameters

(in this case only ... upper uncertainty changes)

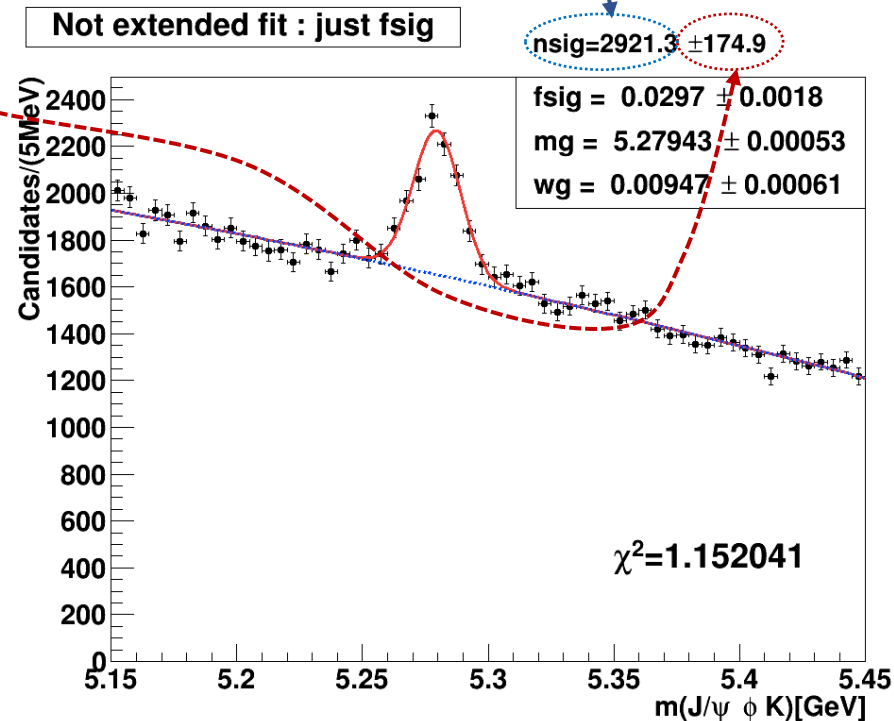
```
*****
** 23 **MINOS      3000
*****
FCN=-1.15354e+06 FROM MINOS  STATUS=SUCCESSFUL  702 CALLS      1275 TOTAL
                    EDM=0.000362648  STRATEGY= 1  ERROR MATRIX ACCURATE
EXT PARAMETER
NO.  NAME      VALUE      PARABOLIC  MINOS ERRORS
      NAME      VALUE      ERROR      NEGATIVE   POSITIVE
 1  c0e      -2.25841e-01  5.55902e-03 -5.54067e-03  5.57848e-03
 2  c1e      -1.08452e-02  6.05944e-03 -6.12080e-03  6.00134e-03
 3  mge       5.27943e+00  5.30310e-04 -5.28542e-04  5.34442e-04
 4  nbkg      9.55524e+04  3.51136e+02 -3.50289e+02  3.52220e+02
 5  nsig      2.92321e+03  1.74070e+02 -1.74275e+02  1.76448e+02
 6  wge       9.48373e-03  6.08532e-04 -5.99390e-04  6.22600e-04
ERR DEF= 0.5
RooRealVar::nsig = 2923.21 +/- (-174.275,176.448) L(2000 - 3800)
```

# How-to-calculate the # of signal candidates in a not extended fit

After executing the *not extended* fit ...

```
// double cand_s = fsig.getVal()*myEntries; // in case you want to use it later as a variable
//
cout << "\n # of entries = " << myEntries << " of which # signal candidates is = " << fsig.getVal()*myEntries << " +/- " << fsig.getError()*myEntries << endl;

// -- let me write the # of candidates estimated by the fit (via fsig):
TLatex* myLatCands = new TLatex(5.318,2600.,Form("nsig=%.1f",fsig.getVal()*myEntries));
TLatex* myLatCands1 = new TLatex(5.38,2600.,Form("#pm%.1f",fsig.getError()*myEntries)); // it rounds as expected
myLatCands->SetTextSize(0.038);
myLatCands1->SetTextSize(0.038);
xframeChi2->addObject(myLatCands);
xframeChi2->addObject(myLatCands1);
...
```



## How-to-calculate the normalized chi-squared $\tilde{\chi}_{fit}^2$ - I

In the code of the macro, I propose **two different ways to extract the  $\tilde{\chi}_{fit}^2$**  of the binned ML fit.

We apply both for the *not extended* fit. **We will prefer the 2<sup>nd</sup> approach and choose it for the *extended* fit.**

```
// Note: will try two approaches
//
// -- 1st approach
//
// --https://root.cern.ch/doc/master/classRooPlot.html
// Syntax: chiSquare (const char *pdfname, const char *histname, int nFitParam=0) const
// Description : Calculate and return reduced chi-squared between a curve and a histogram.
// Syntax: chiSquare (int nFitParam=0) const
// Description: Shortcut for RooPlot::chiSquare(const char* pdfname, const char* histname, int nFitParam=nullptr)
RooPlot* xframeChi2 = x.frame("");
Bmass.plotOn(xframeChi2); // histogram (type RooDataHist)
model.plotOn(xframeChi2,RooFit::LineColor(kRed)); // curve
//
RooArgSet* floatParams = model.getParameters(Bmass);
int numFreeParams = floatParams->getSize();
cout << "\n # of free fit params is = " << numFreeParams << endl; (A)
//
// normalized chiSquared is given by chi2 divided by ndof = (# bins of the fit range - #free params)
// though, the following is given already normalized!
Double_t chi2Norm = xframeChi2->chiSquare(numFreeParams);
cout << "\n the Chi2 for the not-extended fit is = " << chi2Norm << endl; (B)
//
```

# How-to-calculate the normalized chi-squared $\tilde{\chi}_{fit}^2$ - II

```
// -- 2nd approach
//
RooAbsReal* chi2 = model.createChi2(Bmass,Range(begin,end));
// if extended ... add Extended(true): createChi2(Bmass,Range(begin,end),Extended(true))
// because in this way the prediction of the total number of events is taken from the extended pdf and not from the histogram
cout << "\n chi2 (not normalized) is = " << chi2->getVal() << endl; (C)
// maybe this is the best way but be careful with the parameters you pass to the createChi2 function
// since it does not divide by ndf, i.e. the number of non-zero bins minus the number of parameters;
// you have to do this manually afterwards!
//
int nDOF = nBins - numFreeParams;
cout << "\n nDOF is = " << nDOF << endl; (D)
double chi2NormCalc = (chi2->getVal()) / nDOF;
cout << "\n chi2 (normalized by manual calculation) is = " << chi2NormCalc << endl; (E)
//
// Still, this method doesn't get it right if your pdf is continuous,
// because it just takes the pdf value in the bin center.
// If this is a problem for you, you can consider wrapping the pdf in a RooBinSamplingPdf - to be explored -
// which turns the continuous pdf into a binned pdf where the values for each bin are obtained by integration.
// However, this can be important only in case of steep functions
// where value at the center of the bin and integral over the bin may differ; this is not the case here!
```

this is done for the extended fit !

```
# of free extended-fit params is = 6
nDOF is = 54
chi2 (normalized by manual calculation) is = 1.15203
```

We get on screen@ execution time:

```
# of free fit params is = 5
the Chi2 for the not-extended fit is = 1.158
-----
chi2 (not normalized) is = 63.3622
nDOF is = 55
chi2 (normalized by manual calculation) is = 1.15204
```

(A)

(B)

(C)

(D)

(E)

1<sup>st</sup> approach

2<sup>nd</sup> approach

To print on the plot:

```
TLatex* myLatChi2 = new TLatex(5.35,400.,Form("#chi^{2}=%f",chi2NormCalc));
xframeChi2->addObject(myLatChi2);
```

# Correlation Matrix

It is possible to save the status of the fit, including the information about the covariance matrix:

```
RooFitResult* fitres = m.save();
```

It is possible to visualize the correlation matrix:

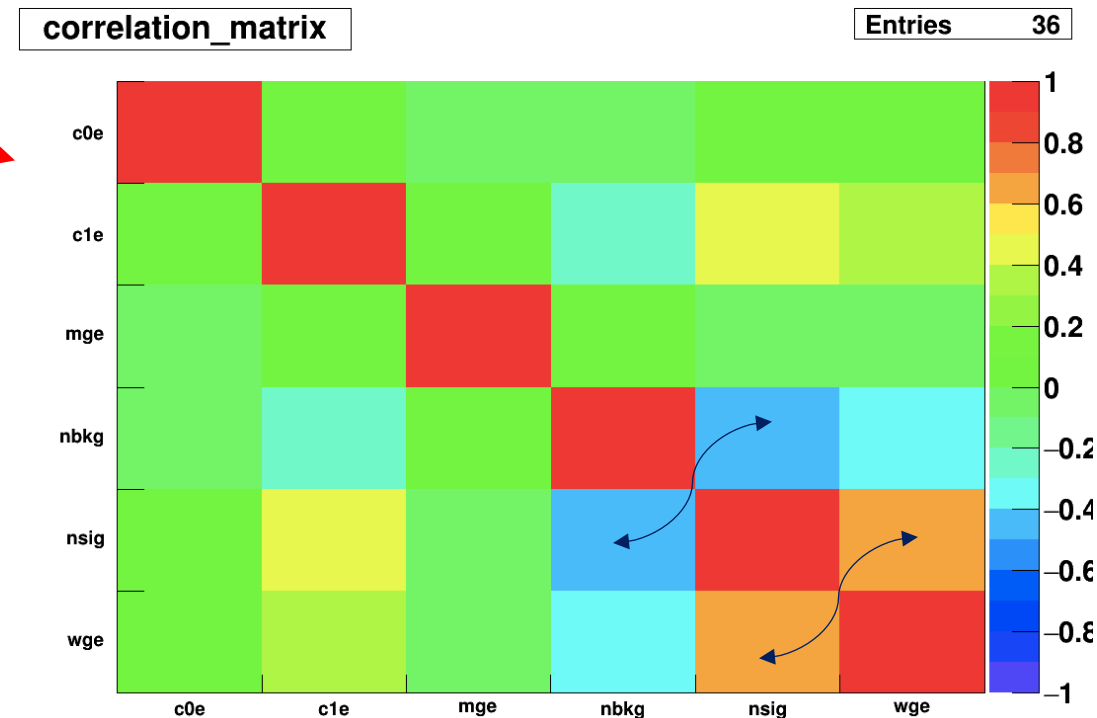
```
gStyle->SetPalette(1); //- for better color choice  
fitres->correlationHist()->Draw("colz");
```

## Note:

- anticorrelation between `nsig` and `nbkg` as expected
- correlation between `nsig` and width of Gaussian `wge`

Note: in general, if correlations are very strong (i.e.  $> 0.9$ ) the model may become unstable and it may be worthwhile to fix one of the parameters in the fit.

If the strong correlation is between two nuisance parameters, this is not a problem. Instead, when a parameter-of-interest is correlated with a nuisance one, it must be avoided to fix the nuisance parameter because the risk is to strongly under-estimate the uncertainty on the physical parameter!



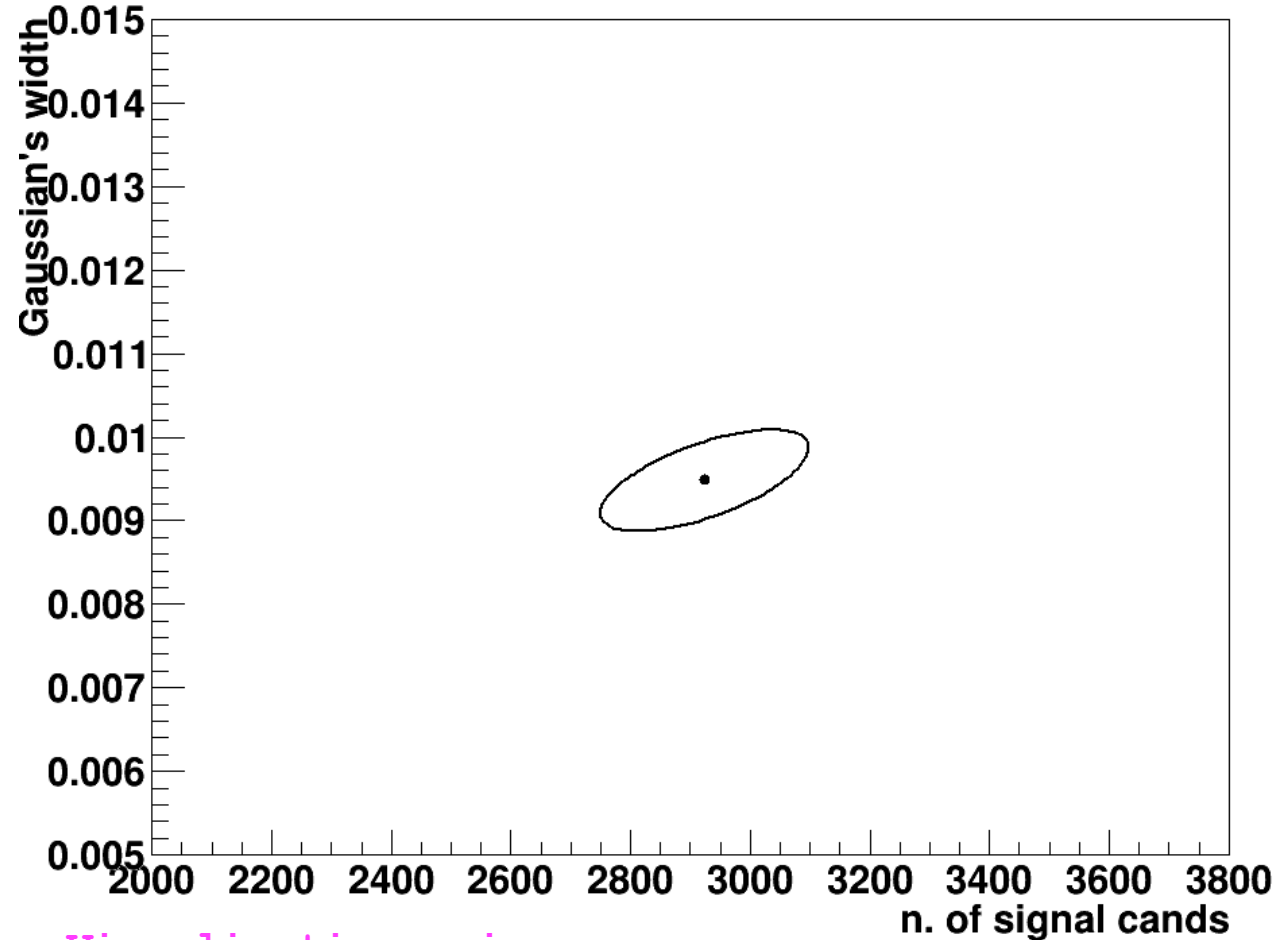
myCorrelationMatrix.png

# Visualization of correlated errors - I

It is also possible to visualize errors & correlation matrix elements:

```
RooPlot* paramFrame = new RooPlot(nsig,wge);  
fitres->plotOn(paramFrame,nsig,wge);  
paramFrame->SetTitleOffset(1.38,"Y");  
paramFrame->Draw();  
myC->SaveAs("./pdfParamVisualization_nsig_wge.png");
```

where this example shows a certain level of correlation.



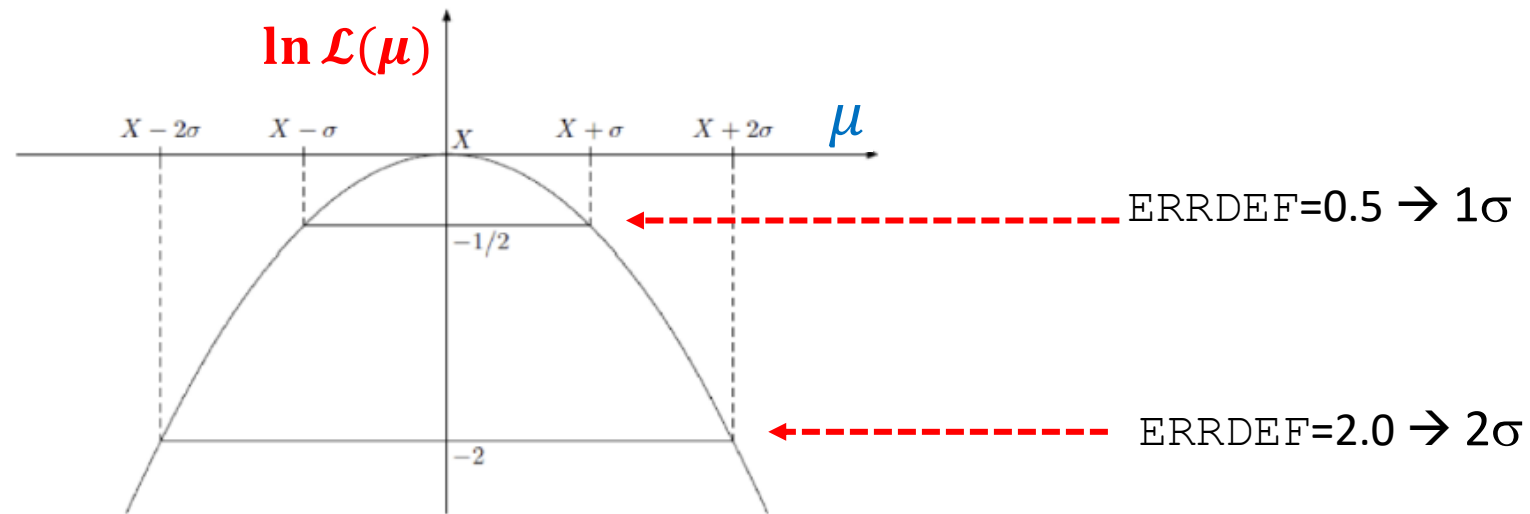
pdfParamVisualization\_nsig\_wge.png

## Visualization of correlated errors - II

But why  $ERRDEF=0.5$  and  $2.0$  are considered? This is a reminder.

Well, do not forget that a PDF can be converted into a Likelihood function  $\mathcal{L}$  by “exchanging” the vector of observations  $\vec{x}$  with the vector of parameters  $\vec{\theta}$  !

For only one parameter, say  $\mu$ , the likelihood is a function of it, namely  $\mathcal{L}(\mu)$ , and  **$\ln \mathcal{L}(\mu)$  is a parabola!**



Note : if you put the “-” in front of it, thus getting the neg-log-likelihood,  $-\ln \mathcal{L}(\mu)$ , the parabola changes sign and “points” upwards instead of downwards.

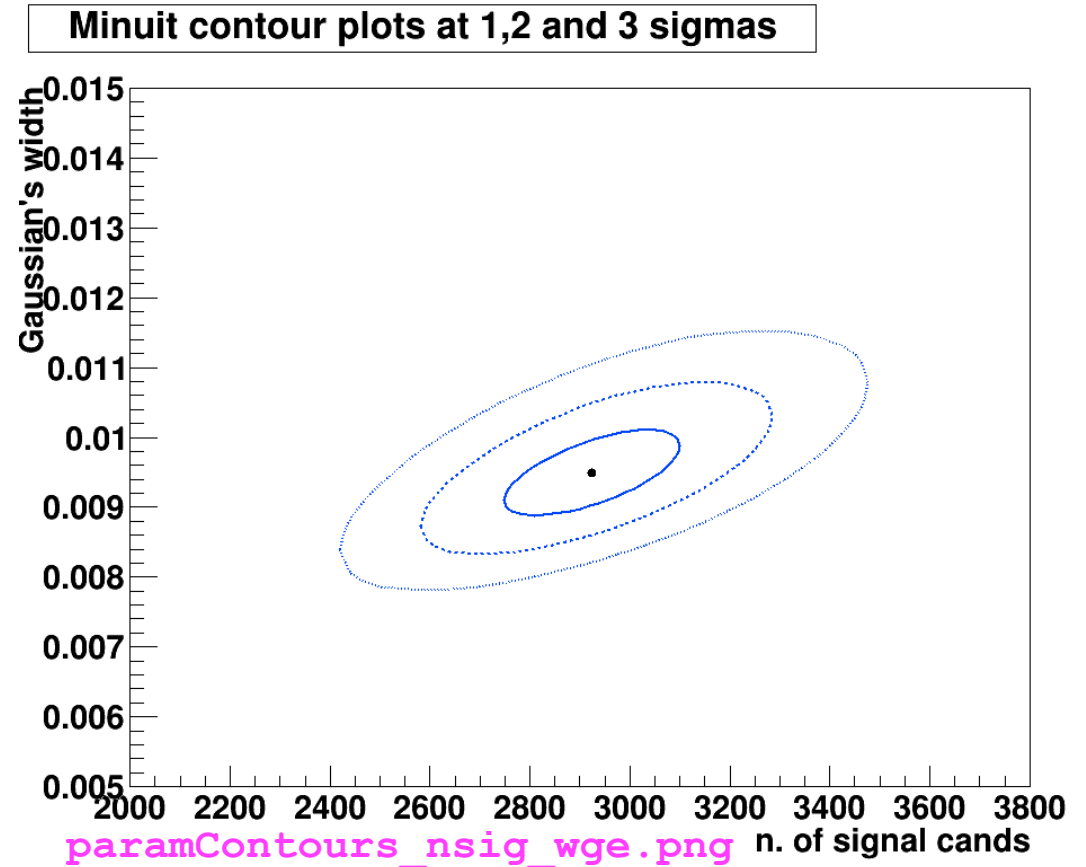
**Extension:** Now suppose we’ve 2 parameters of interest; in this case you can imagine a paraboloid instead of a parabola with different aperture when projecting in 1-dim. The “multivariate” uncertainty is then represented by an elliptic contour.



## Visualization of correlated errors - III

```
RooPlot* contourFrame = m.contour(nsig,wge,1,2,3,0,0,0); // gives the 3 contours obtained for 1, 2 and 3 sigmas
contourFrame->SetTitle("Minuit contour plots at 1,2 and 3 sigmas");
contourFrame->SetTitleOffset(1.38,"Y");
contourFrame->Draw();
myC->SaveAs("./paramContours_nsig_wge.png");
```

It starts the MNCONTOUR calculation of 50 point on three contours (for  $ERRDEF = 0.5, 2.0$  and  $4.5$ ). Each point is identified by a pair of values of parameters 5 ( $nsig$ ) and 6 ( $wge$ ) on the scatter plot.



As you can check ... the three sets of 50 pairs of values are printed on the screen @ execution time.  
Note: the contour with  $ERRDEF=0.5$  is the same one obtained earlier with a different command.



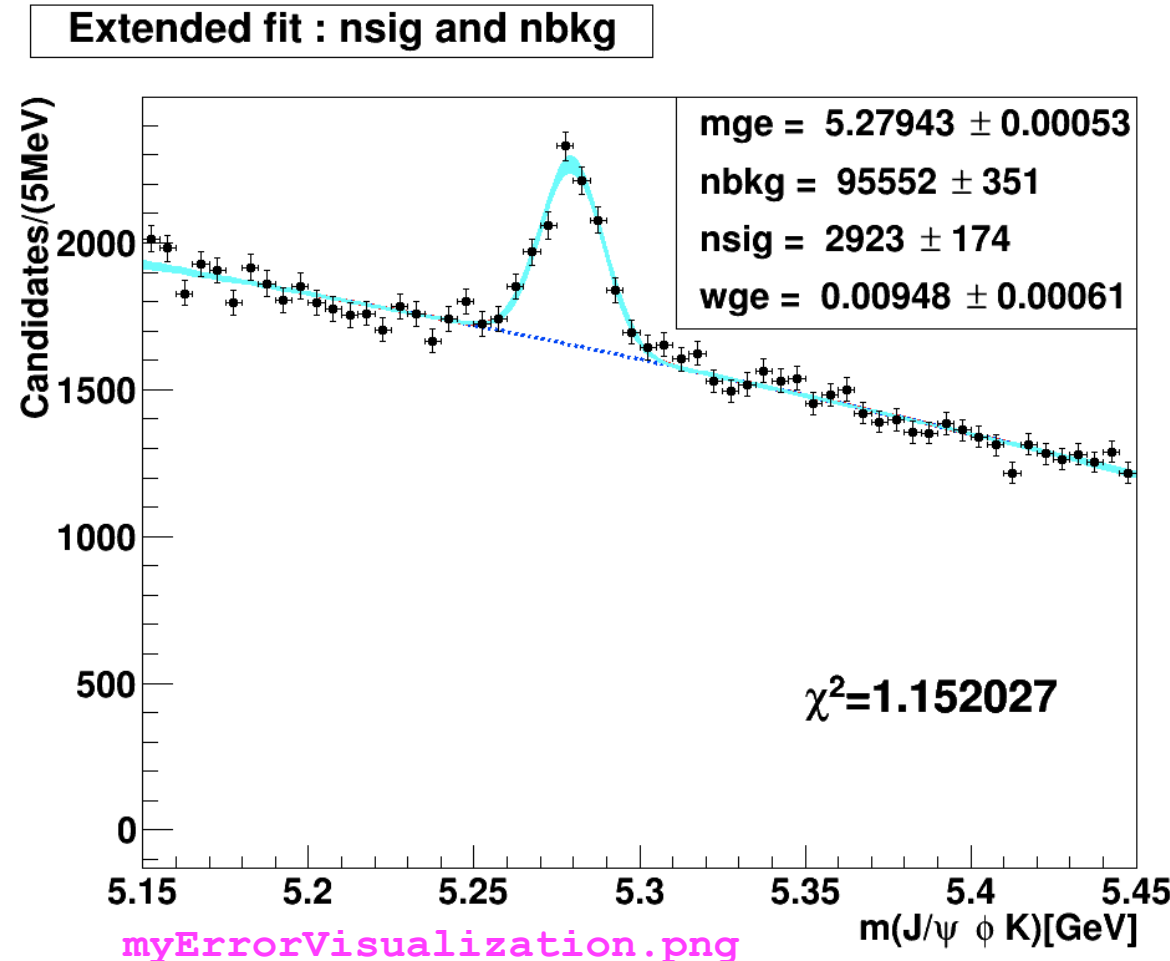
# Visualization of the fit uncertainty

It is possible to propagate the errors (stored in the covariance matrix of a fit result) to a PDF projection:

```
model_extended.plotOn(yframe, VisualizeError(*fitres));  
yframe->Draw();
```

To get the points' errors over the cyan shaded region describing the uncertainty we need to add the following two lines (to get the "trick" done):

```
BmassExt.plotOn(yframe);  
yframe->Draw("Esame");
```



# Visualization of the fit log-likelihood function and of the Profile Likelihood ratio

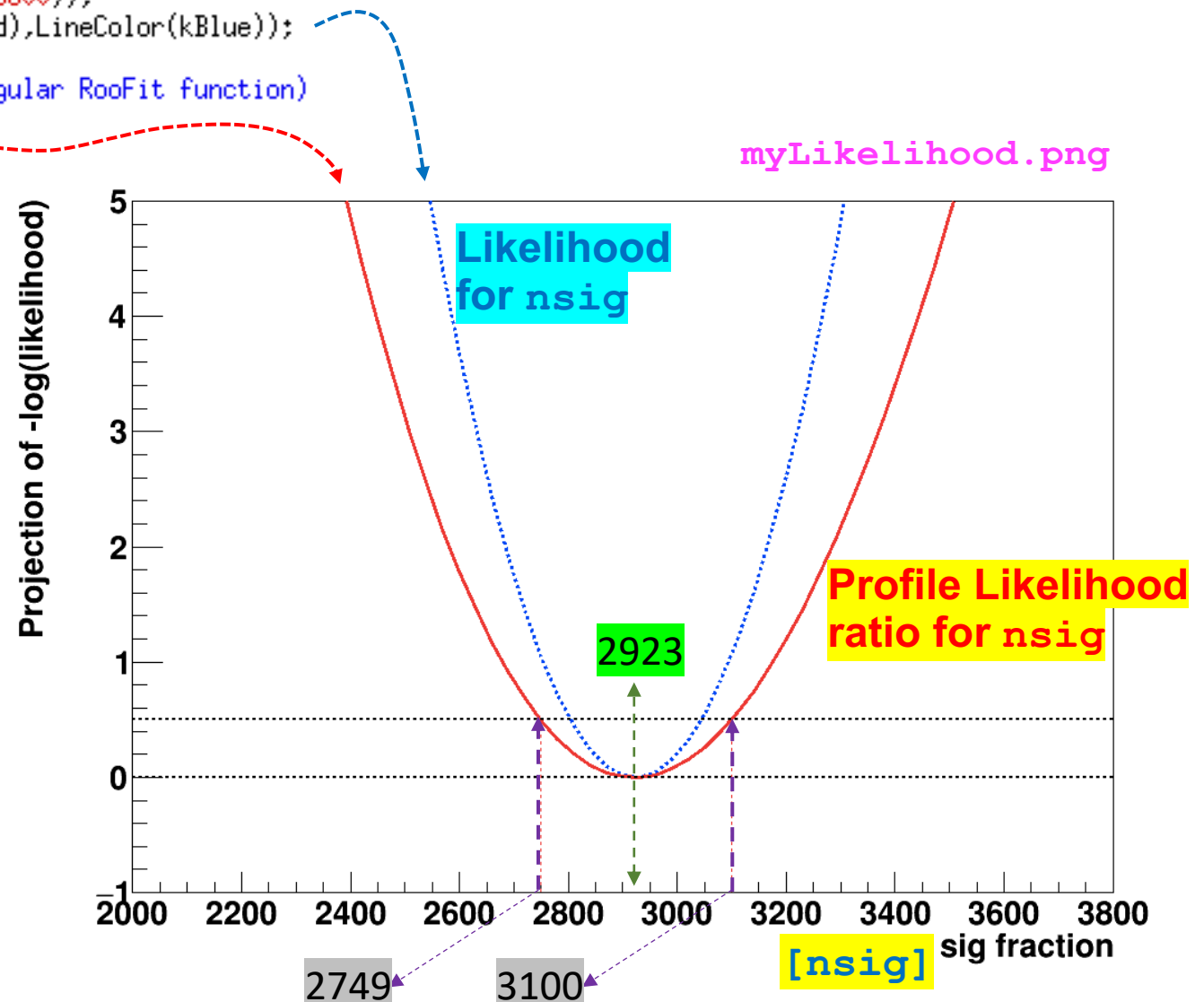
We can obtain the best estimate for **nsig** & the MINOS uncertainty **corresponding** to the interval provided by the PL ratio :

```
// plot the likelihood as a function of the parameter of interest (here nsig):  
RooPlot* nsig_frame = nsig.frame(RooFit::Bins(60),RooFit::Range(2000,3800));  
nll->plotOn(nsig_frame,RooFit::ShiftToZero(),RooFit::LineStyle(kDashed),LineColor(kBlue));  
//  
// make the Profile Likelihood ratio (that can be represented as a regular RooFit function)  
RooAbsReal* pll_nsig = nll->createProfile(nsig);  
pll_nsig->plotOn(nsig_frame,RooFit::ShiftToZero(),LineColor(kRed));  
nsig_frame->SetMinimum(-1);  
nsig_frame->SetMaximum(5);  
nsig_frame->Draw();
```

From MIGRAD: 2923.2

From MINOS: (2923.2)  $- 174.3 + 176.4$   
(slightly asymmetric)

Overall interval:  $\cong [2749, 3100]$



# Code of the macro `yield.C` - I

```
////////////////////////////////////
// To run it:
// root> .L yield.C
// root> main()
//
////////////////////////////////////

#include <TRoot.h>
#include <TFile.h>
#include <TH1.h>
#include <TF1.h>
#include <TF2.h>
#include <TFormula.h>
#include <TStyle.h>
#include <TCanvas.h>
#include <TProfile.h>
#include <TString.h>
#include <TLine.h>
#include <TPad.h>
#include <TMath.h>
#include <TLatex.h>
#include <TLegend.h>
#include <iostream>
#include <TColor.h>
#include "TAxis.h"
#include <TPaveLabel.h>

using namespace RooFit;

////////////////////////////////////---inizio main

int main() {

  gROOT->SetStyle("Plain");
  gStyle->SetOptStat(10);
  gStyle->SetTitleOffset(1.2, "");
  //
  TCanvas* myC = new TCanvas("myC", "Plots", 1000, 800);
  //
  //////////////////////////////////////
  //
  TFile f1("DatasetAandB_KaonTrackRefit_Bwin_new_21aug13.root", "READ");
  TH1D *hist = (TH1D*)f1.Get("myJpsiKKKmass_all");
  // in alternativa c'e' anche l'istogramma myJpsiKKKmass_tight
  //
  //--- # of bins:
  int nBins = hist->GetNbinsX();
  int entries = hist->GetEntries(); // this contains also overflow so I recalculate by hand as follows:
  int myEntries;
  for (int l=0; l<60; l++)
  {
    myEntries += hist->GetBinContent(l+1);
  }
  //
  //////////////////////////////////////
}
```

## Code of the macro `yield.C` - II

```
////////////////////////////////////
//
RooRealVar x("x","m(J/#psi #phi K)[GeV]",5.15,5.45);
RooDataHist Bmass(hist->GetName(),hist->GetTitle(),RooArgSet(x),RooFit::Import(*hist, kFALSE));
//
Float_t begin = 5.15;
Float_t end = 5.45;
//
RooPlot* xframe = x.frame("");
Bmass.plotOn(xframe);
////
myC->cd();
//
// -- SIGNAL
RooRealVar mg("mg","Gaussian's mean",5.28,5.275,5.285);
RooRealVar wg("wg","Gaussian's width",0.010,0.005,0.015);
RooGaussian gauss1("gauss1","Gauss(x,mg,wg)",x,mg,wg);
// -- BKG
RooRealVar c0("c0","1st coeff",0.5,-1000.,1000.);
RooRealVar c1("c1","2nd coeff",-0.5,-1000.,1000.);
//--RooRealVar c2("c2","3rd coeff",0.1,-1000.,1000.); // 2nd order degree is enough here
RooChebychev cheby("cheby","Chebyshev",x,RooArgList(c0,c1)); // 2 coeff. means 2nd order polynomial
//
// -- TOTAL pdf : f*gauss1 + (1-f)*cheby
RooRealVar fsig("fsig","narrow fraction",0.05,0.0,1.0);
RooAddPdf model("model","gauss1+cheby",RooArgList(gauss1,cheby),fsig); // configured in this way this is not extended
//
// -- Execute FIT
// model.fitTo(Bmass,RooFit::Minos(kTRUE)); // let us give explicitly also the FitRange:
model.fitTo(Bmass,RooFit::Minos(kTRUE),Range(begin,end));
model.plotOn(xframe,RooFit::LineColor(kRed));
model.plotOn(xframe,RooFit::Components(cheby),RooFit::LineStyle(kDashed));
model.paramOn(xframe,Parameters(RooArgSet(mg,wg,fsig)),Layout(0.53,0.9,0.9)); // 3rd is up
//
// double candS = fsig.getVal()*myEntries; // in case you want to use it later as a variable
//
cout << "\n # of entries = " << myEntries << " of which # signal candidates is = " << fsig.getVal()*myEntries << " +/- " << fsig.getError()*myEntries << endl;
//
xframe->SetTitle("Not extended fit : just fsig and (1-fsig)");
xframe->Draw();
//
```

## Code of the macro `yield.C` - III

```
//  
// -- calculate a chiSquared (from a curve and a histogram in a RooPlot)  
//=====
```

//  
// Note: will try two approaches  
//  
// -- 1st approach  
//  
// --<https://root.cern.ch/doc/master/classRooPlot.html>  
// Syntax: chiSquared (const char \*pdfname, const char \*histname, int nFitParam=0) const  
// Description : Calculate and return reduced chi-squared between a curve and a histogram.  
// Syntax: chiSquared (int nFitParam=0) const  
// Description: Shortcut for RooPlot::chiSquared(const char\* pdfname, const char\* histname, int nFitParam=nullptr)

```
RooPlot* xframeChi2 = x.frame("");  
Bmass.plotOn(xframeChi2); // histogram (type RooDataHist)  
model.plotOn(xframeChi2,RooFit::LineColor(kRed)); // curve  
//  
RooArgSet* floatPars = model.getParameters(Bmass);  
int numFreeParams = floatPars->getSize();  
cout << "\n # of free fit params is = " << numFreeParams << endl;  
//  
// normalized chiSquared is given by chi2 divided by ndof = (# bins of the fit range - #free params)  
// though, the following is given already normalized!  
Double_t chi2Norm = xframeChi2->chiSquared(numFreeParams);  
cout << "\n the Chi2 for the not-extended fit is = " << chi2Norm << endl;  
//
```

# Code of the macro `yield.C` - IV

```
//  
// -- 2nd approach  
//  
RooAbsReal* chi2 = model.createChi2(Bmass,Range(begin,end));  
// if extended ... add Extended(true): createChi2(Bmass,Range(begin,end),Extended(true))  
// because in this way the prediction of the total number of events is taken from the extended pdf and not from the histogram  
cout << "\n chi2 (not normalized) is = " << chi2->getVal() << endl;  
// maybe this is the best way but be careful with the parameters you pass to the createChi2 function  
// since it does not divide by ndf, i.e. the number of non-zero bins minus the number of parameters;  
// you have to do this manually afterwards!  
//  
int nDOF = nBins - numFreeParams;  
cout << "\n nDOF is = " << nDOF << endl;  
double chi2NormCalc = (chi2->getVal()) / nDOF;  
cout << "\n chi2 (normalized by manual calculation) is = " << chi2NormCalc << endl;  
//  
// Still, this method doesn't get it right if your pdf is continuous,  
// because it just takes the pdf value in the bin center.  
// If this is a problem for you, you can consider wrapping the pdf in a RooBinSamplingPdf - to be explored -  
// which turns the continuous pdf into a binned pdf where the values for each bin are obtained by integration.  
// However, this can be important only in case of steep functions  
// where value at the center of the bin and integral over the bin may differ; this is not the case here!  
//  
model.plotOn(xframeChi2,RooFit::Components(cheby),RooFit::LineStyle(kDashed));  
model.paramOn(xframeChi2, Parameters(RooArgSet(mg,wg,fsig)), Layout(0.53,0.9,0.9));  
//  
TLatex* myLatChi2 = new TLatex(5.35,400.,Form("#chi^{2}=%f",chi2NormCalc));  
xframeChi2->addObject(myLatChi2);  
//  
// -- let me write the # of candidates estimated by the fit (via fsig):  
TLatex* myLatCands = new TLatex(5.318,2600.,Form("nsig=%1f",fsig.getVal()*myEntries));  
TLatex* myLatCands1 = new TLatex(5.38,2600.,Form("#pm%1f",fsig.getError()*myEntries)); // it rounds as expected  
myLatCands->SetTextSize(0.038);  
myLatCands1->SetTextSize(0.038);  
xframeChi2->addObject(myLatCands);  
xframeChi2->addObject(myLatCands1);  
//  
xframeChi2->SetTitle("Not extended fit : just fsig");  
xframeChi2->SetYTitle("Candidates/(5MeV)");  
xframeChi2->SetTitleOffset(1.32,"Y");  
xframeChi2->Draw();  
//  
myC->SaveAs("./myBmass.png");  
//gSystem->Sleep(20000);  
//  
myC->Update();  
myC->cd();  
//
```





## Code of the macro `yield.C` - VI

```
//
// -- fit is done but now we want to derive more info from the RooFitResult object
//
model_extended.plotOn(yframe, VisualizeError(*fitres));
yframe->Draw();
BmassExt.plotOn(yframe);
yframe->Draw("Esame");
myC->SaveAs("./myErrorVisualization.png");
myC->Update();
myC->cd();
//
//--
////RooAbsPdf* paramPDF = fitres->createHessePdf(RooArgSet(nsig,wge)); //not working
//
RooPlot* paramFrame = new RooPlot(nsig,wge);
fitres->plotOn(paramFrame,nsig,wge);
paramFrame->SetTitleOffset(1.38,"Y");
paramFrame->Draw();
myC->SaveAs("./pdfParamVisualization_nsig_wge.png"); // it gives just a visualization with the 1sigma ellipse
myC->Update();
myC->cd();
//
// -- the following is more useful than the previous
//
RooPlot* contourFrame = m.contour(nsig,wge,1,2,3,0,0,0); // gives the 3 contours obtained for 1, 2 and 3 sigmas
contourFrame->SetTitle("Minuit contour plots at 1,2 and 3 sigmas");
contourFrame->SetTitleOffset(1.38,"Y");
contourFrame->Draw();
myC->SaveAs("./paramContours_nsig_wge.png");
myC->Update();
myC->cd();
//
```



## Code of the macro `yield.C` - VII

```
//  
////////// now plot Likelihood and Profile Likelihood Ratio functions :  
//  
myC->Divide(1,1);  
//  
// plot the likelihood as a function of the parameter of interest (here nsig):  
RooPlot* nsig_frame = nsig.frame(RooFit::Bins(60),RooFit::Range(2000,3800));  
nll->plotOn(nsig_frame,RooFit::ShiftToZero(),RooFit::LineStyle(kDashed),LineColor(kBlue));  
//  
// make the Profile Likelihood ratio (that can be represented as a regular RooFit function)  
RooAbsReal* pll_nsig = nll->createProfile(nsig);  
pll_nsig->plotOn(nsig_frame,RooFit::ShiftToZero(),LineColor(kRed));  
nsig_frame->SetMinimum(-1);  
nsig_frame->SetMaximum(5);  
nsig_frame->Draw();  
//  
TLine *line0 = new TLine(2000,0,3800,0);  
line0->SetLineColor(1);  
line0->SetLineWidth(2);  
line0->SetLineStyle(2);  
line0->Draw("same");  
//  
TLine *line05 = new TLine(2000,0.5,3800,0.5);  
line05->SetLineColor(1);  
line05->SetLineWidth(2);  
line05->SetLineStyle(2);  
line05->Draw("same");  
//  
TLine *lineN1 = new TLine(3100,-1.,3100,0.5);  
lineN1->SetLineColor(2);  
lineN1->SetLineWidth(1);  
lineN1->SetLineStyle(2);  
lineN1->Draw("same");  
//  
TLine *lineN2 = new TLine(2749,-1.,2749,0.5);  
lineN2->SetLineColor(2);  
lineN2->SetLineWidth(1);  
lineN2->SetLineStyle(2);  
lineN2->Draw("same");  
//  
myC->SaveAs("./myLikelihood.png");  
myC->Update();  
myC->cd();  
//  
delete myC;  
//  
gROOT->Reset();  
gROOT->Clear();  
//  
return 0;  
}
```

## Connection between **MINOS uncertainties** & **Profile Likelihood**

In the next slides this connection will be investigated & explained.

# Profile Likelihood

In the next slides this connection will be argued/explained.

Firstly, remember the difference between these two concepts:

- **POI(s)** = **parameter(s) of interest**: parameter(s) of theoretical model (we assume predicts distribution of observed variables)

- **NPs** = **nuisance parameters**: additional unknown parameters, appearing together with the POI(s), that represent the effect of the detector response (resolutions, miscalibrations, ...), the presence of background, ...

Typically they can represent systematic uncertainties & can be usually determined from simulation or data control samples.

Let's assume for simplicity to have a POI  $\mu$  and a set of NPs  $\vec{\theta}$  (i.e. all parameter are treated as NPs with exception of  $\mu$ ).

The **likelihood function** is written as:  $\mathcal{L}(\vec{x}; \mu, \vec{\theta})$ . To easy the notation we drop the  $\vec{x}$  and write simply  $\mathcal{L}(\mu, \vec{\theta})$ .

The so-called **profile likelihood** is constructed following this prescription:

- for a **given value of the POI**  $\bar{\mu}$  derive the **ML estimates**  $\hat{\hat{\theta}}(\bar{\mu})$  (it's a *conditional ML estimate*; fit with  $\mu$  fixed to a constant value  $\bar{\mu}$ )

- thus the maximum likelihood for a given value of  $\bar{\mu}$  is  $\mathcal{L}_{max}(\bar{\mu}, \hat{\hat{\theta}}(\bar{\mu}))$ ;

- recalculating (CPU expensive) for each value of  $\mu$  (scan of  $\mu$  values) we get a truly function of  $\mu$ :  $\mathcal{L}_{max}(\mu, \hat{\hat{\theta}}(\mu))$   
which is the **likelihood function maximized w.r.t. all the NPs and** is called **profile likelihood** !

## Profile Likelihood ratio

On the other hand it is always possible to maximize the likelihood getting the best estimates (fit values) of  $\mu$  and  $\vec{\theta}$  corresponding to the observed data  $\vec{x}$ :  $\hat{\mu}$  and  $\hat{\vec{\theta}}$ . Thus the maximized likelihood is:  $\mathcal{L}_{max}(\hat{\mu}, \hat{\vec{\theta}})$

At this point we can consider the **Profile Likelihood ratio**:  $\lambda(\mu) = \frac{\mathcal{L}_{max}(\mu, \hat{\vec{\theta}}(\mu))}{\mathcal{L}_{max}(\hat{\mu}, \hat{\vec{\theta}})}$  (that does not depend on the NPs  $\vec{\theta}$ )

This ratio is used in the convenient test statistic  $t_\mu = -2 \ln \lambda(\mu)$ . Dropping the obvious “max” index:  $\lambda(\mu) = \frac{\mathcal{L}(\mu, \hat{\vec{\theta}}(\mu))}{\mathcal{L}(\hat{\mu}, \hat{\vec{\theta}})}$

In other words the profile likelihood ratio **substitutes** the ordinary likelihood ratio, in the test statistics  $t_\mu = -2 \ln \lambda(\mu)$ , **when** we have to deal with nuisance parameters:

$$\lambda(\mu) = \frac{\mathcal{L}(\mu)}{\mathcal{L}(\hat{\mu})} \quad \text{Maximum Likelihood} \quad \longrightarrow \quad \lambda(\mu) = \frac{\mathcal{L}(\mu, \hat{\vec{\theta}}(\mu))}{\mathcal{L}(\hat{\mu}, \hat{\vec{\theta}})} \quad \text{Maximum Likelihood}$$

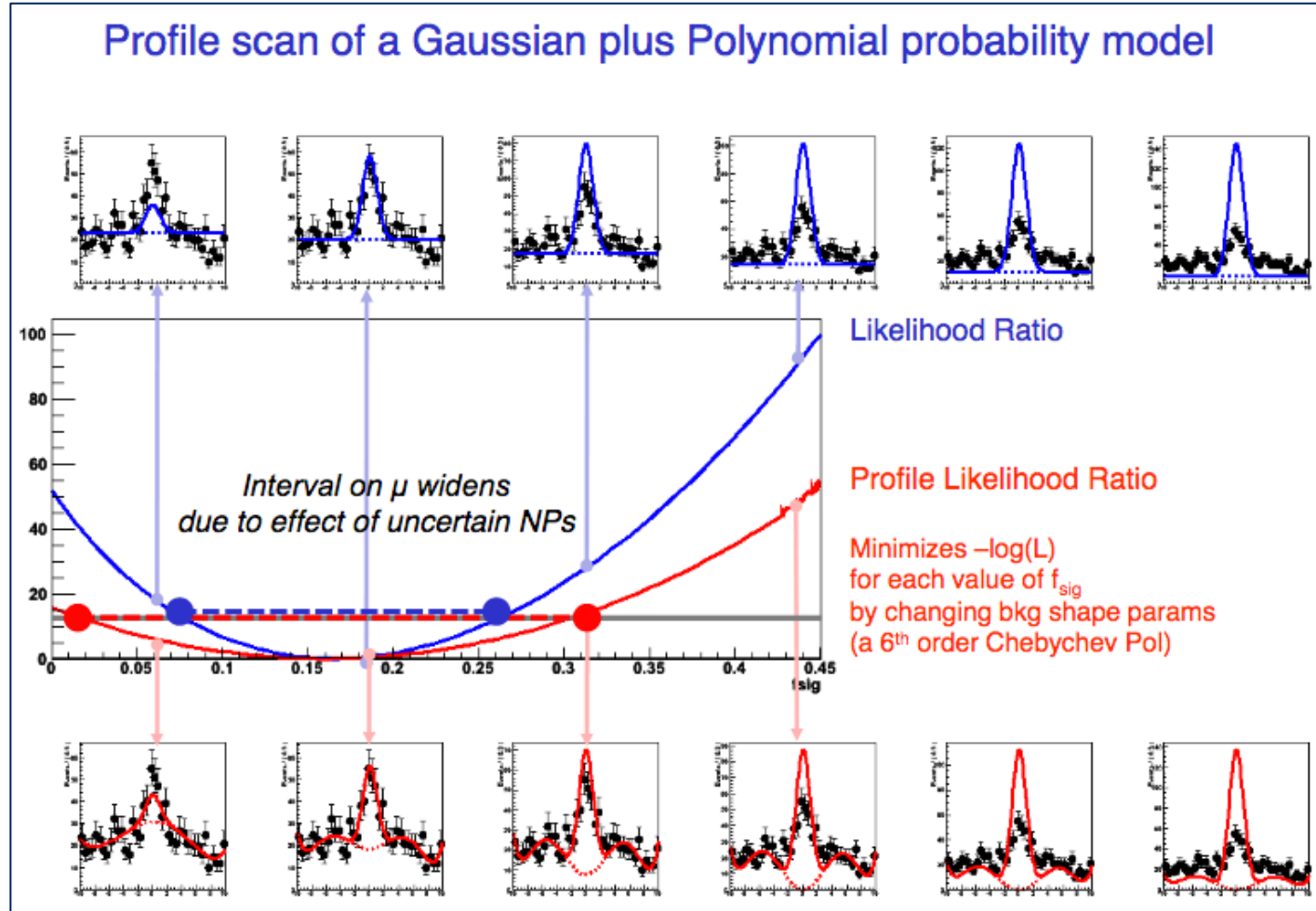
Maximum Likelihood for a given  $\mu$

Comments on the **Profile Likelihood approach**:

- it is **computationally challenging** because it requires to perform the minimization of the likelihood w.r.t. **all** the nuisance parameters for every point in the profile likelihood curve (see also next slide that illustrates this)
- the minimization can be difficult because of the possibly strong correlation among POIs and NPs or multiple/local minima

# How to obtain a Profile Likelihood

For visualization purposes have a look at this figure illustrating the scan of  $\mu$  values in order to obtain  $\mathcal{L}(\mu, \hat{\hat{\theta}}(\mu))$  :



# Profile Likelihood & Contours - I

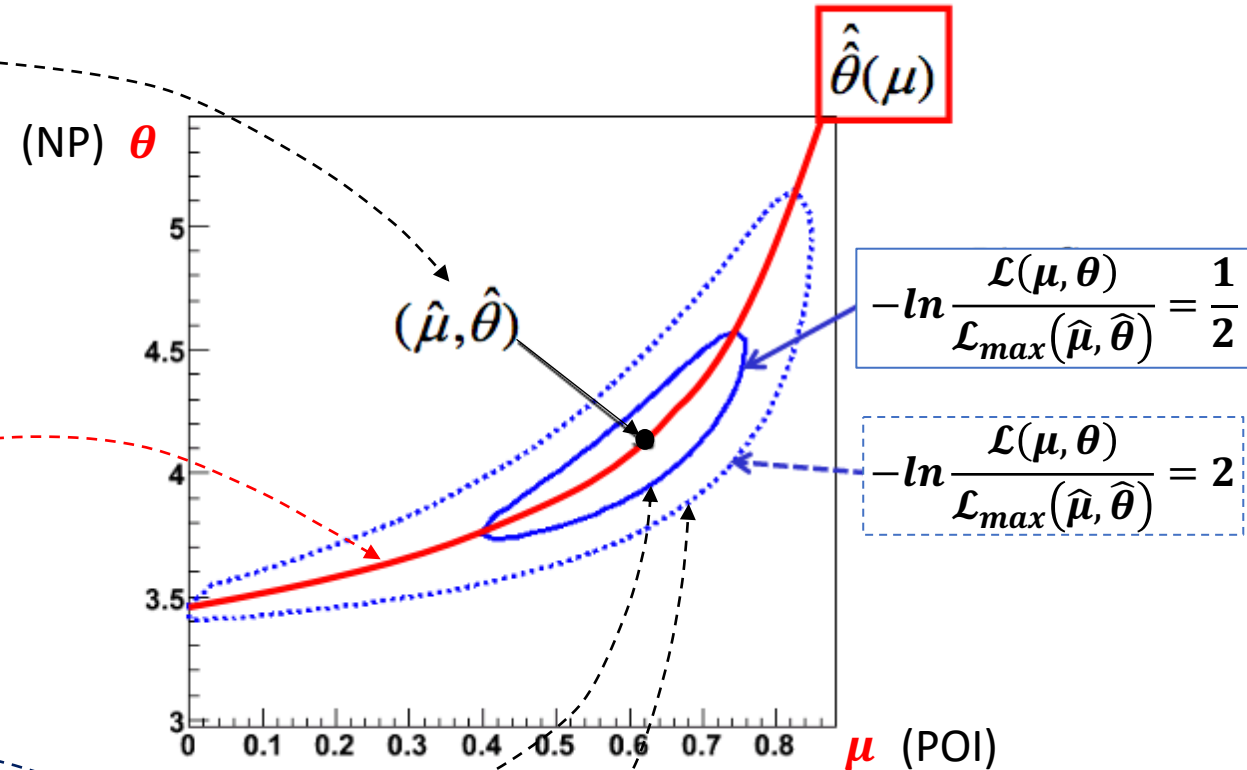
For illustration purposes let us consider one POI ( $\mu$ ) and one NP ( $\theta$ ) in order to visualize the profiling.

Firstly let us design/locate the **(black) point** representing their best estimates that maximize the likelihood:  $(\hat{\mu}, \hat{\theta})$

$$(\mu, \theta) \equiv (\hat{\mu}, \hat{\theta}) \Rightarrow \mathcal{L} \equiv \mathcal{L}_{max}(\hat{\mu}, \hat{\theta})$$

Secondly let's design the **red curve** that represents those points  $(\mu, \hat{\theta}(\mu))$  for which  $\mathcal{L} \equiv \mathcal{L}_{max}(\mu, \hat{\theta}(\mu))$

...corresponding to a subset of subsequently given/fixed values of  $\mu$  which includes also the special value  $\bar{\mu} \equiv \hat{\mu}$ .



Finally we can design the contour curves at  $1\sigma$  and  $2\sigma$  with respect to  $(\hat{\mu}, \hat{\theta})$  the maximum (minimum) of the (negative) likelihood. This is discussed in detail in the next slide.

## Profile Likelihood & Contours - II

In particular the first contour corresponds to a set of parameters such that:  $-2\ln\mathcal{L}(\mu, \theta) = -2\ln\mathcal{L}_{max}(\hat{\mu}, \hat{\theta}) + 1$

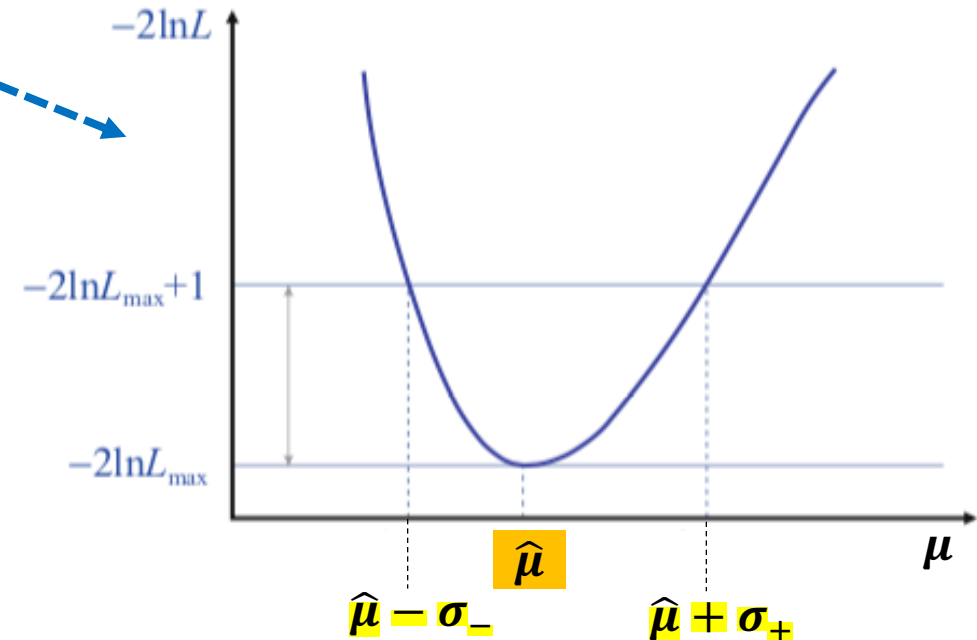
Indeed in the simplest case (only one POI & no NPs) one has:

$$-2\ln\mathcal{L}(\mu) \equiv -2\ln\mathcal{L}_{max}(\hat{\mu}) + 1$$

$$\iff 2\ln\mathcal{L}(\mu) - 2\ln\mathcal{L}_{max}(\hat{\mu}) = -1$$

$$\iff 2\ln\frac{\mathcal{L}(\mu)}{\mathcal{L}_{max}(\hat{\mu})} = -1 \iff -\ln\frac{\mathcal{L}(\mu)}{\mathcal{L}_{max}(\hat{\mu})} = +\frac{1}{2}$$

Note that in general **the uncertainty** (and thus the  $1\sigma$  interval) can be **asymmetric** (as depicted).



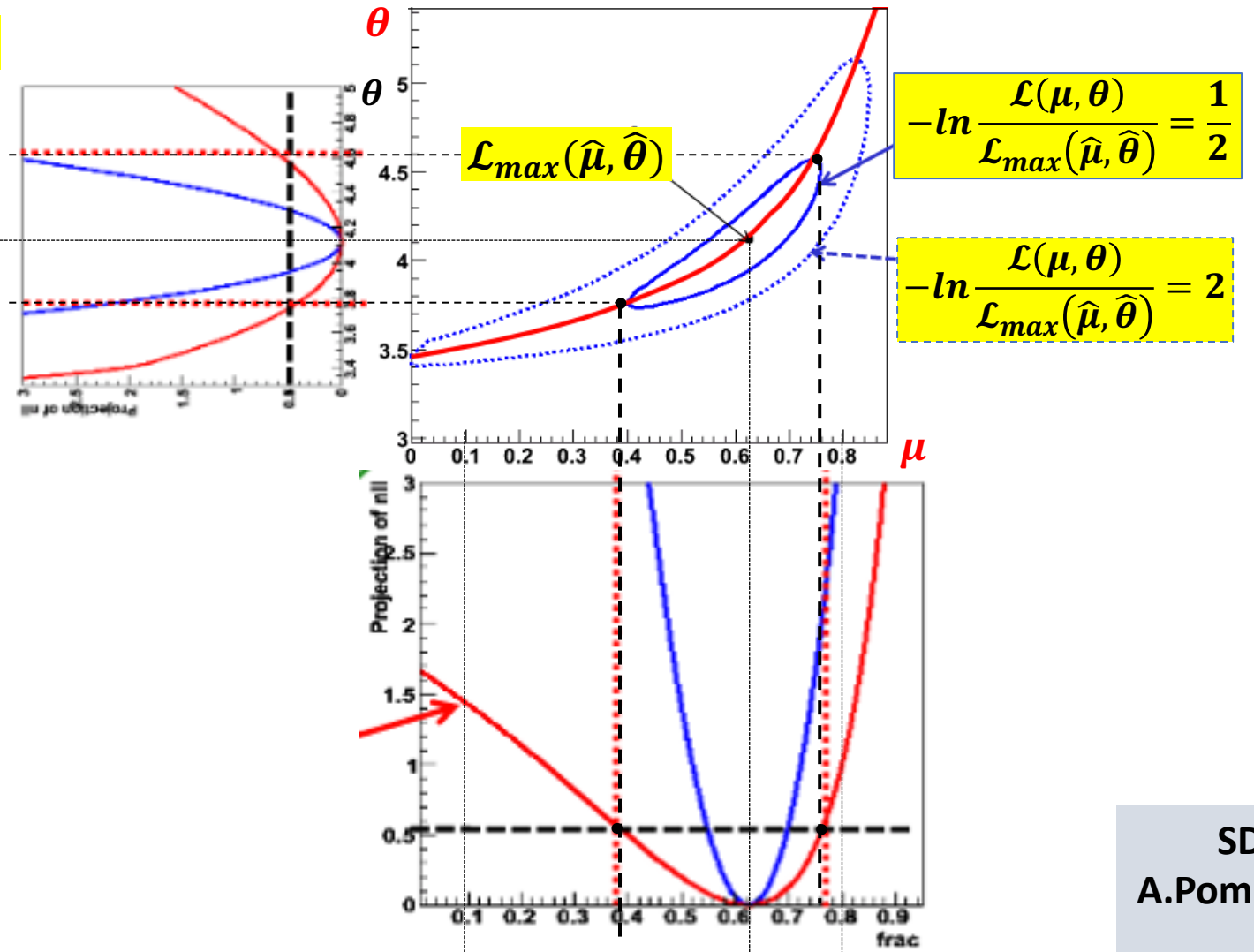
When there is also one NP one gets 2D contours (see next slide) and

$$-\ln\frac{\mathcal{L}(\mu)}{\mathcal{L}_{max}(\hat{\mu})} \quad \text{becomes} \quad -\ln\frac{\mathcal{L}(\mu, \theta)}{\mathcal{L}_{max}(\hat{\mu}, \hat{\theta})}$$

# Profile Likelihood & Contours - III

When there is also one POI and one NP one gets **2D contours**, here designed with the 2 projections:

Clearly **the correct  $1\sigma$  interval for the POI is given by the projection of the contour** (and not by the -marginalized - likelihood, that is the blue projection, which ignores the effect of the presence of the NP). It can be demonstrated that **this confidence interval provides the correct coverage in the frequentistic approach**.



**The overall uncertainty is in general asymmetric!**



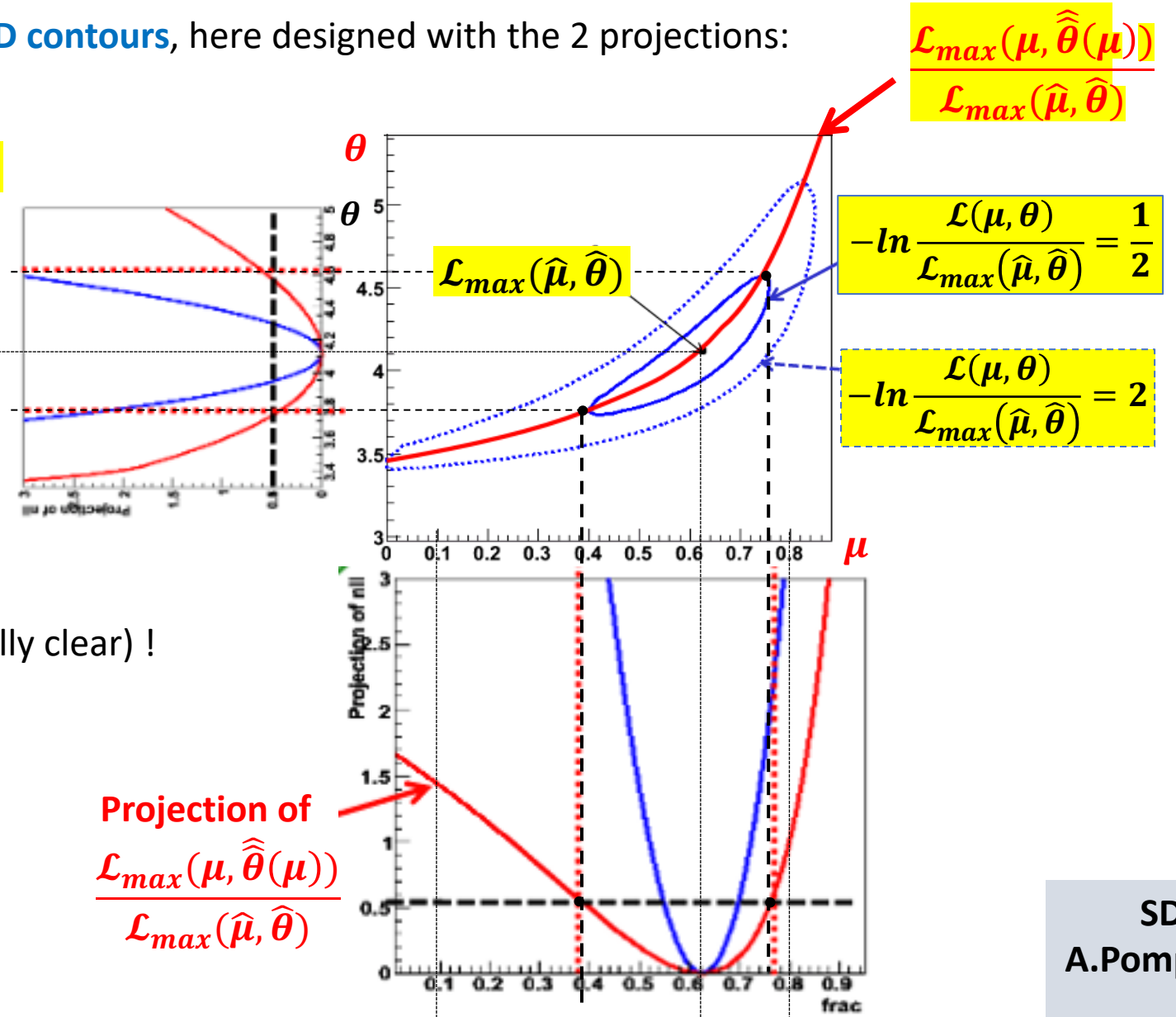
# Same confidence interval provided by Profile Likelihood & Contours

When there is also one POI and one NP one gets **2D contours**, here designed with the 2 projections:

Clearly **the correct  $1\sigma$  interval for the POI is given by the projection of the contour** (and not by the -marginalized - likelihood, that is the blue projection, which ignores the effect of the presence of the NP). It can be demonstrated that **this confidence interval provides the correct coverage in the frequentistic approach.**

It is also crucial to know that **this interval is the same provided by (the projection of) the Profile Likelihood ratio** based on  $\mathcal{L}_{max}(\mu, \hat{\theta}(\mu))$  (as visually clear) !

Indeed the addition of NP(s) broadens the shape of the Profile Likelihood as a function of the POI compared with the case where NP(s) are not added. As a consequence, **the uncertainty on the POI increases when NPs - that usually model sources of systematic uncertainties - are included.** **The overall uncertainty is in general asymmetric!**

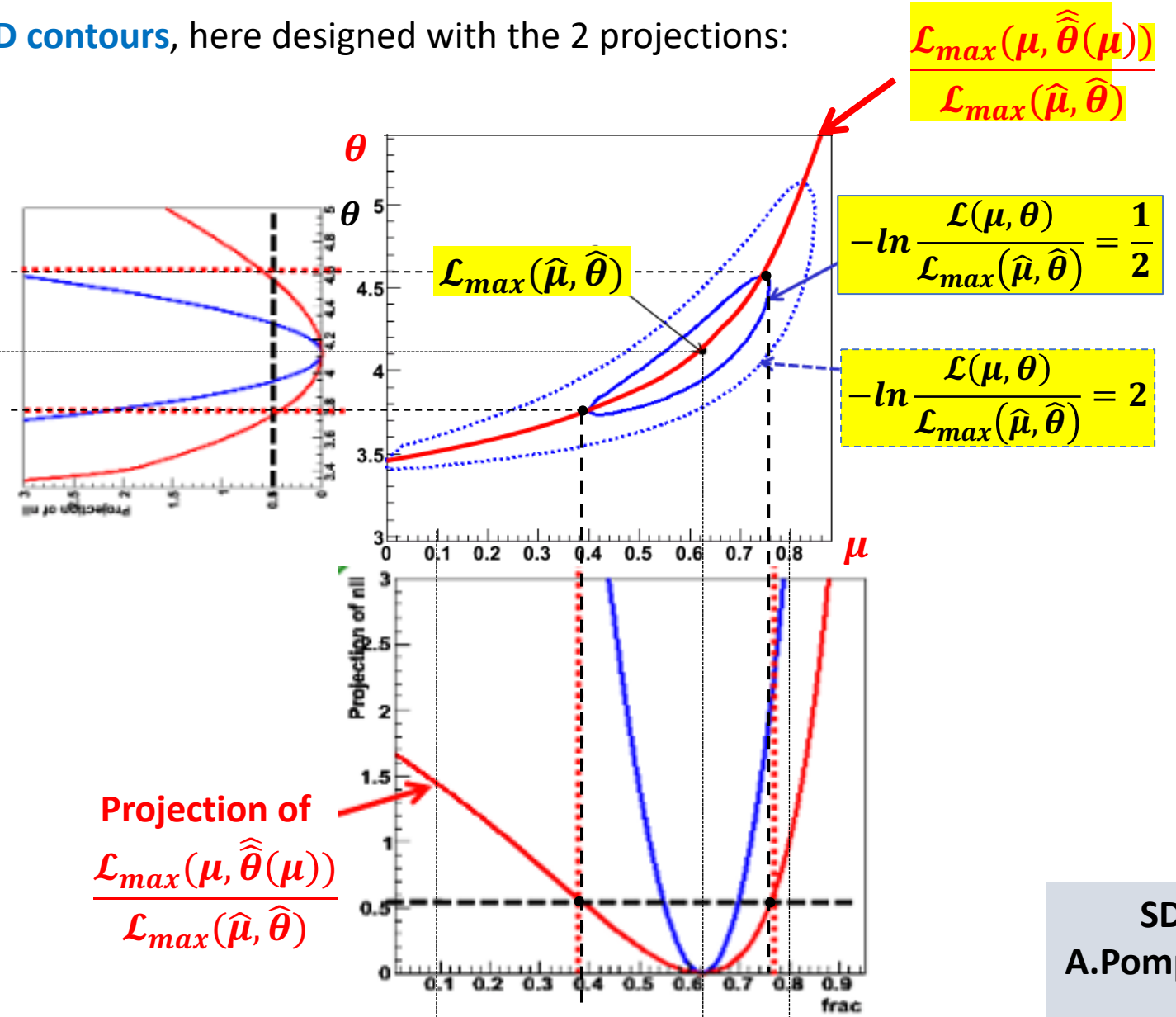


# MINOS uncertainties by likelihood scan

When there is also one POI and one NP one gets **2D contours**, here designed with the 2 projections:

Moreover :

The MINOS method (called by MINUIT) determines the overall uncertainties (in general asymmetric) based on the *likelihood scan* namely on the  $-2\ln\mathcal{L}(\mu)$  scan used to determine the  $1\sigma$  contour.



# MINOS uncertainties by likelihood scan

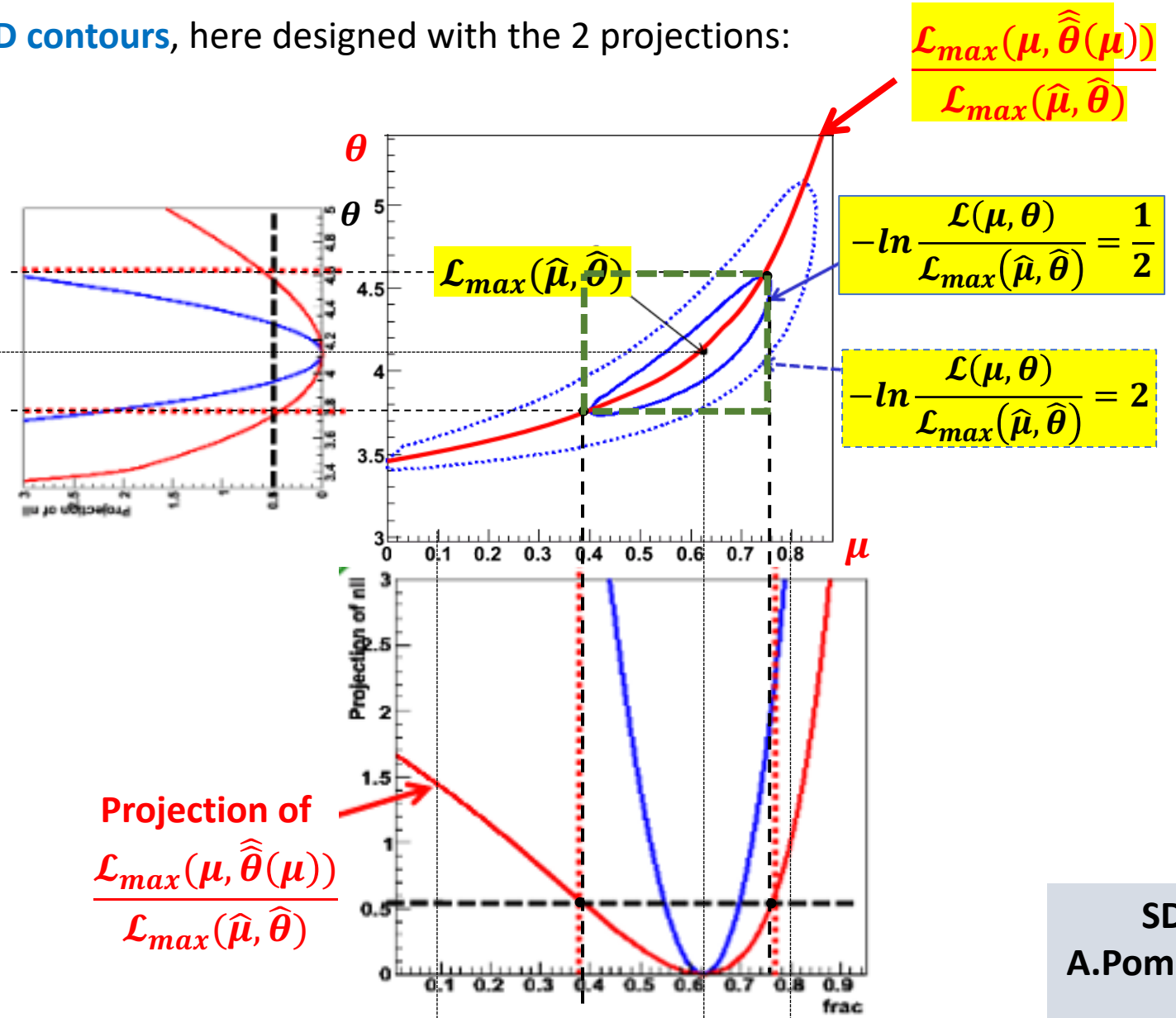
When there is also one POI and one NP one gets **2D contours**, here designed with the 2 projections:

Moreover :

The MINOS method (called by MINUIT) determines the overall uncertainties (in general asymmetric) based on the *likelihood scan* namely on the  $-2\ln\mathcal{L}(\mu)$  scan used to determine the  $1\sigma$  contour.

The MINOS errors can be visualized with the size of the green bounding box around the contour

given by 
$$-\ln \frac{\mathcal{L}(\mu, \theta)}{\mathcal{L}_{max}(\hat{\mu}, \hat{\theta})} = \frac{1}{2} !$$



# MINOS uncertainties by likelihood scan

When there is also one POI and one NP one gets **2D contours**, here designed with the 2 projections:

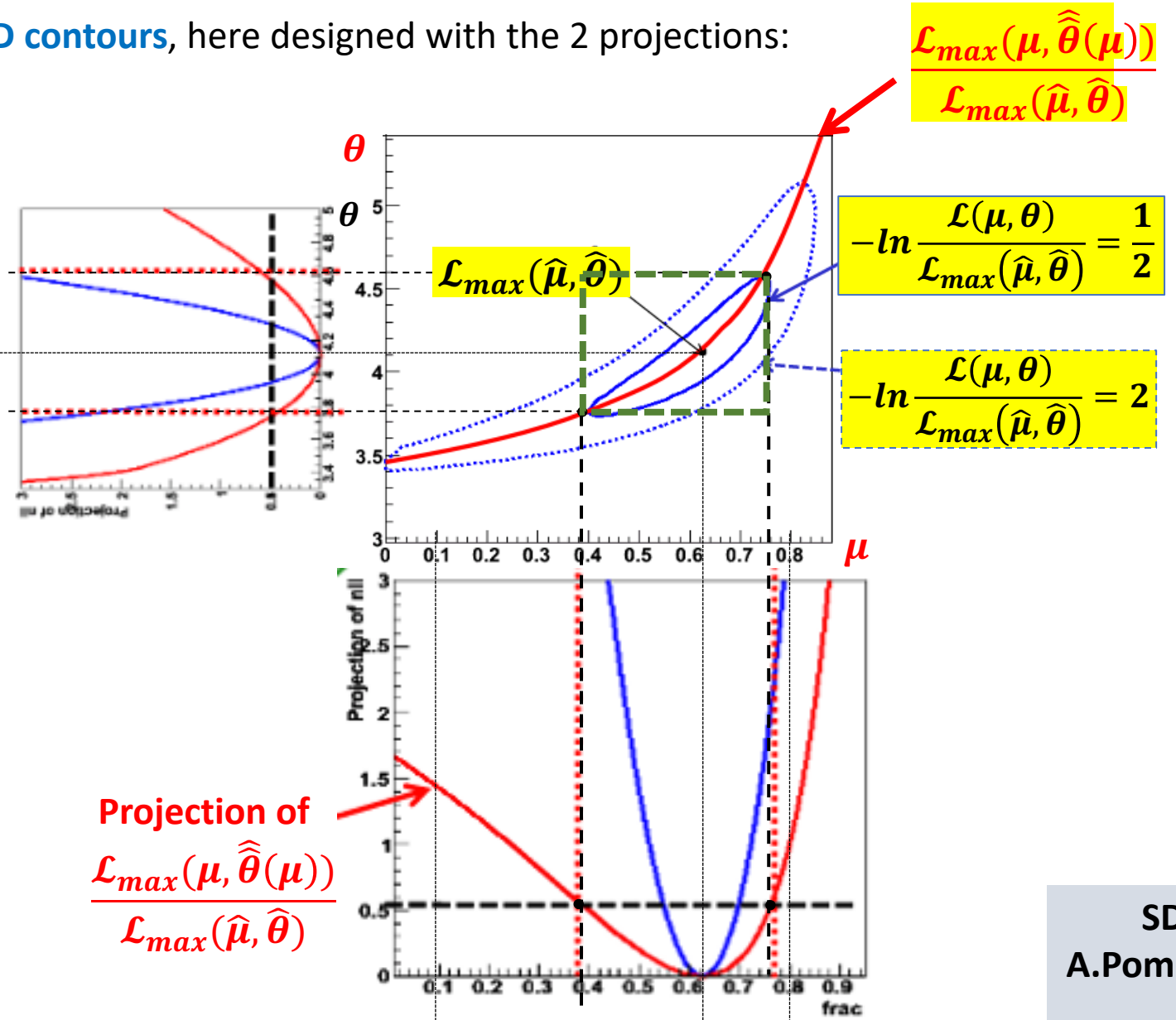
Moreover :

The MINOS method (called by MINUIT) determines the overall uncertainties (in general asymmetric) based on the *likelihood scan* namely on the  $-2\ln\mathcal{L}(\mu)$  scan used to determine the  $1\sigma$  contour.

The MINOS errors can be visualized with the size of the green bounding box around the contour

$$-\ln \frac{\mathcal{L}(\mu, \theta)}{\mathcal{L}_{max}(\hat{\mu}, \hat{\theta})} = \frac{1}{2} !$$

In the gaussian case/regime  $-2\ln\mathcal{L}(\mu)$  can have a parabolic shape and the uncertainty of the POI is symmetric, or “close to symmetric” if parabolic approximation is good and contour is an ellipsis. However - in general - the coverage is usually improved performing the likelihood scan instead of the parabolic approximation (given by HESSE).



# Correspondence between MINOS uncertainties & Profile Likelihood intervals

Summarizing : the MINOS algorithm provides the same (asymmetric) uncertainties given by the Profile Likelihood ratio

For both ... the resulting confidence interval is satisfactorily “covered”.

Let us remind that in the frequentist approach:

For a large fraction of repeated experiments - usually 68.27% - the unknown true value of  $\mu$  is contained in the confidence interval  $[\hat{\mu} - \sigma, \hat{\mu} + \sigma]$ . The fraction is meant in the limit of infinitely large number of repetitions of the experiment, and  $\hat{\mu}$  &  $\sigma$  may vary from one experiment to the other, being the result of a measurement in each experiment.

- **Coverage**: property of the estimated interval to contain the true value in 68.27% of the experiments.

- **Confidence level** : the reference *probability level* usually taken as 68.27%.

Interval estimates that have a larger (or smaller) probability of containing the true value, compared to the desired confidence level, are said to **overcover** (or **undercover**).

It is important to know that the resulting confidence interval from the Profile Likelihood construction will have exact coverage for the points  $(\mu, \hat{\theta}(\mu))$ ; elsewhere it might be over- or under- covering.

We conclude stating that: in the asymptotic regime (very large number of experiments) the MINOS algorithm provides the (asymmetric) uncertainties used in the definition of the frequentist confidence intervals !

## Frequentist confidence intervals when NP are present

Exact confidence intervals are difficult when nuisance parameters are present:

- intervals should cover for any value of NPs (technically difficult)
- typically there can be a significant over-coverage

The approach to use the Profile Likelihood ratio guarantees the coverage at the measured values of NPs (only !)

- technically replace Likelihood ratio with Profile Likelihood ratio
- computationally more intensive but still very tractable

Asymptotically confidence intervals constructed with Profile Likelihood ratio correspond to MINOS likelihood ratios intervals

- as the distribution of the Profile Likelihood becomes asymptotically independent of  $\theta$  the coverage for all values of  $\theta$  is restored !