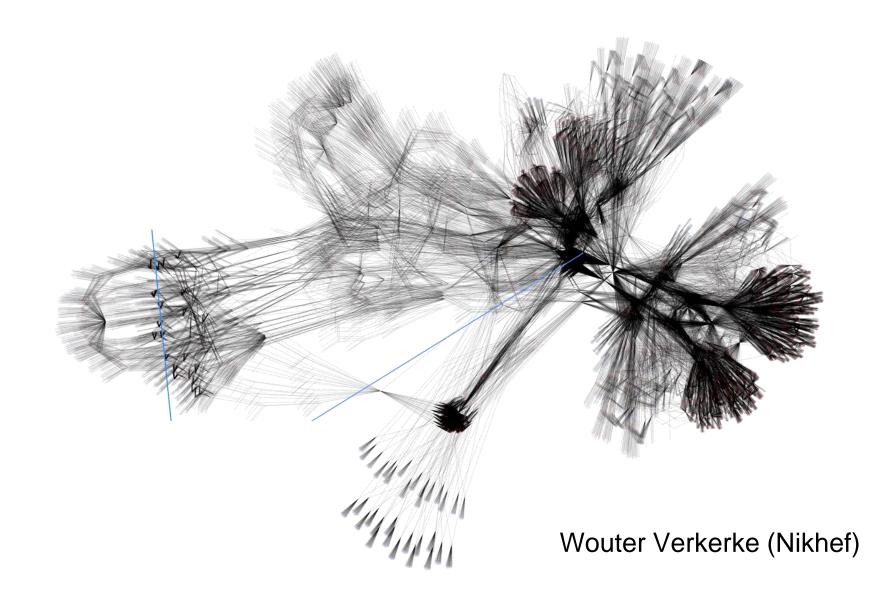
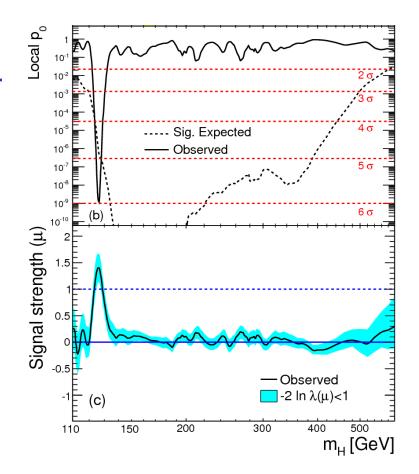
Statistical analysis tools for the Higgs discovery and beyond



What do you want to know?

- Physics questions we have...
 - Does the (SM) Higgs boson exist?
 - What is its production cross-section?
 - What is its boson mass?
- Statistical tests construct probabilistic statements: p(theo|data), or p(data|theo)
 - Hypothesis testing (discovery)
 - (Confidence) intervals
 Measurements & uncertainties
- Result: Decision based on tests
 - "As a layman I would now say: I think we have it"

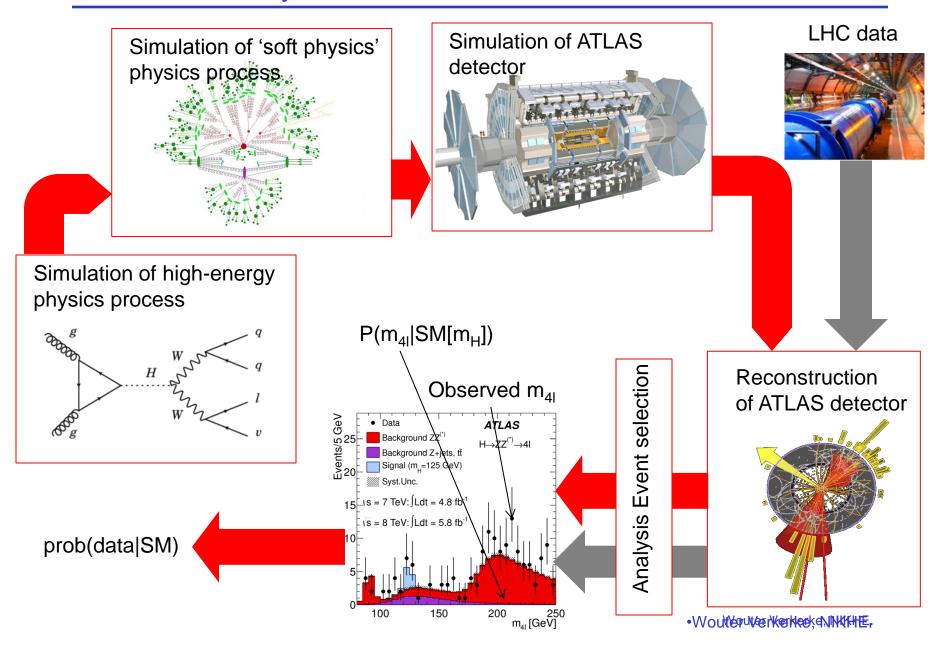




All experimental results start with the formulation of a model

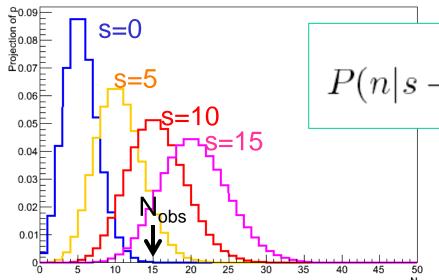
- Examples of HEP physics models being tested
 - SM with m(top)=172,173,174 GeV → Measurement top quark mass
 - SM with/without Higgs boson → Discovery of Higgs boson
 - SM with composite fermions/Higgs → Measurement of Higgs coupling properties
- Via chain of physics simulation, showering MC, detector simulation and analysis software, a physics model is reduced to a statistical model

The HEP analysis workflow illustrated



All experimental results start with the formulation of a

- Examples of HEP physics models being tested
 - SM with m(top)=172,173,174 GeV → Measurement top quark mass
 - SM with/without Higgs boson → Discovery of Higgs boson
 - SM with composite fermions/Higgs → Measurement of Higgs coupling properties
- Via chain of physics simulation, showering MC, detector simulation and analysis software, a physics model is reduced to a statistical model
- A statistical model defines p(data|theory) for all observable outcomes
 - Example of a statistical model for a counting measurement with a known background



$$P(n|s+b) = \frac{(s+b)^n}{n!}e^{-(s+b)}$$

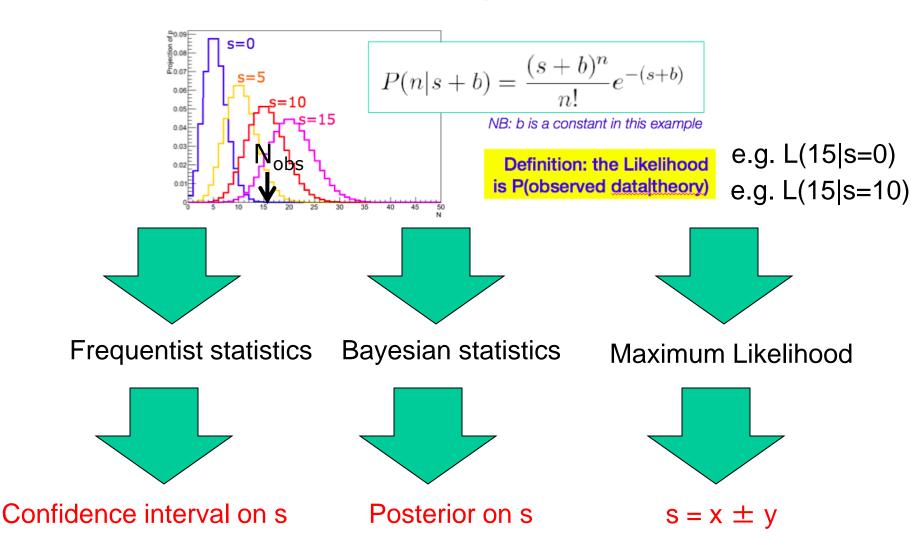
NB: b is a constant in this example

Definition: the Likelihood is P(observed data|theory)

Wouter Verkerke, NIKHEF

Everything starts with the likelihood

 All fundamental statistical procedures are based on the likelihood function as 'description of the measurement'



Everything starts with the likelihood

Frequentist statistics

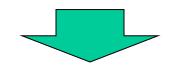
Bayesian statistics

Maximum Likelihood

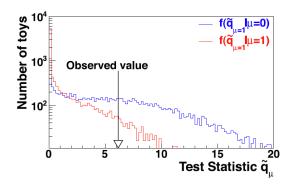


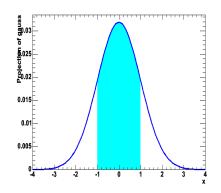
$$I_{m}(\vec{N}_{obs}) = \frac{L(\vec{N} \mid m)}{L(\vec{N} \mid \hat{m})}$$

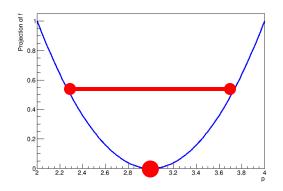
$$P(m) \sqcup L(x \mid m) \times p(m)$$



$$\frac{d \ln L(\vec{p})}{d\vec{p}}\bigg|_{p_i = \hat{p}_i} = 0$$





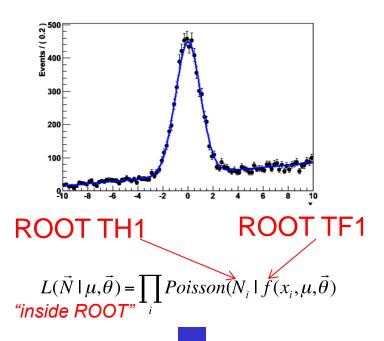


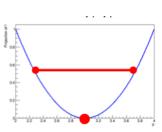




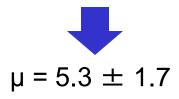


Gaussian + polynomial

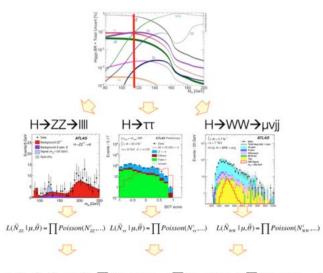




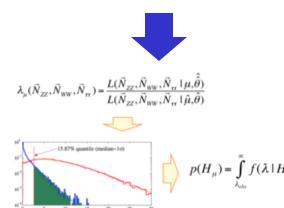
ML estimation of parameters μ,θ using MINUIT (MIGRAD, HESSE, MINOS)



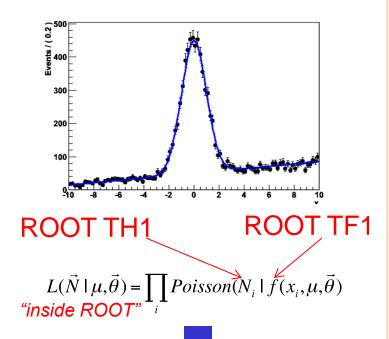
Higgs combination model

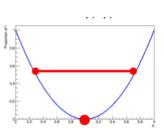


$$L(\vec{N}_{ZZ}, \vec{N}_{\pi}, \vec{N}_{WW} \mid \mu, \vec{\theta}) = \prod Poisson(N_{ZZ}^{i}, \ldots) \cdot \prod Poisson(N_{\pi}^{i}, \ldots) \cdot \prod Poisson(N_{WW}^{i}, \ldots) \cdot \ldots$$



Gaussian + polynomial





ML estimation of parameters μ,θ using MINUIT (MIGRAD, HESSE, MINOS)

$$\mu = 5.3 \pm 1.7$$

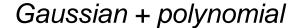
Likelihood Model orders of magnitude more complicated. Describes

- O(100) signal distributions
- O(100) control sample distr.
- O(1000) parameters representing syst. uncertainties

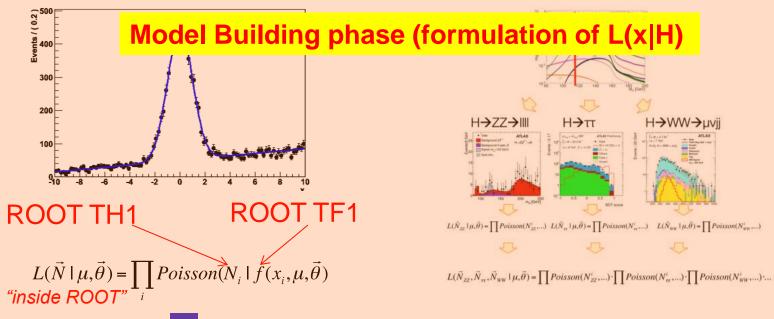


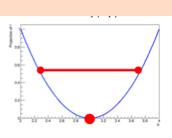


Frequentist confidence interval construction and/or p-value calculation not available as 'ready-to-run' algorithm in ROOT

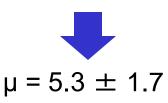


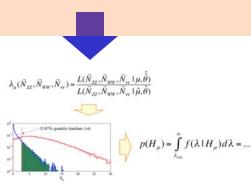
Higgs combination model



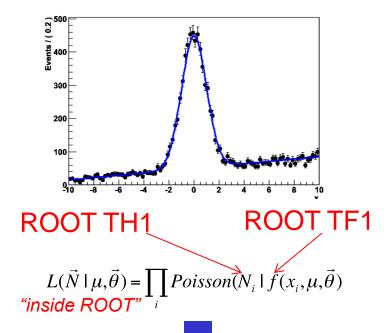


ML estimation of parameters μ,θ using MINUIT (MIGRAD, HESSE, MINOS)

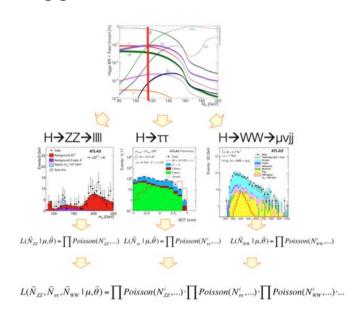


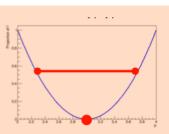


Gaussian + polynomial



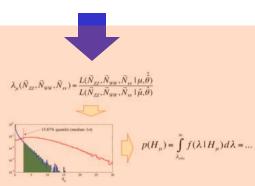
Higgs combination model





ML estimation of parameters μ,θ using MINUIT (MIGRAD, HESSE, MINOS)





Gaussian + polynomial

Higgs combination model

Design goal:

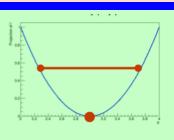
Separate **building of Likelihood model** as much as possible from statistical analysis **using the Likelihood model**

→ More modular software design

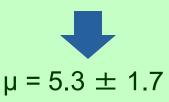
→ 'Plug-and-play with statistical techniques

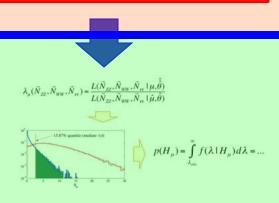
→ Factorizes work in collaborative effort

"il



ML estimation of parameters μ,θ using MINUIT (MIGRAD, HESSE, MINOS)





The idea behind the design of RooFit/RooStats/HistFactory

- Modularity, Generality and flexibility
- Step 1 Construct the likelihood function L(x|p)

RooFit, or RooFit+HistFactory

Step 2 – Statistical tests on parameter of interest p

Procedure can be Bayesian, Frequentist, or Hybrid), but always based on L(x|p)

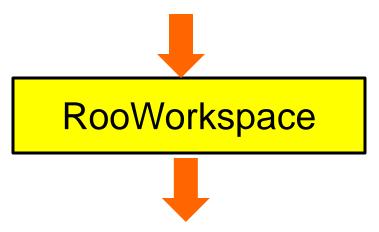
RooStats

 Steps 1 and 2 are conceptually separated, and in Roo* suit also implemented separately.

The idea behind the design of RooFit/RooStats/HistFactory

- Steps 1 and 2 can be 'physically' separated (in time, or user)
- Step 1 Construct the likelihood function L(x|p)

RooFit, or RooFit+HistFactory



Complete description of likelihood model, persistable in ROOT file (RooFit pdf function)

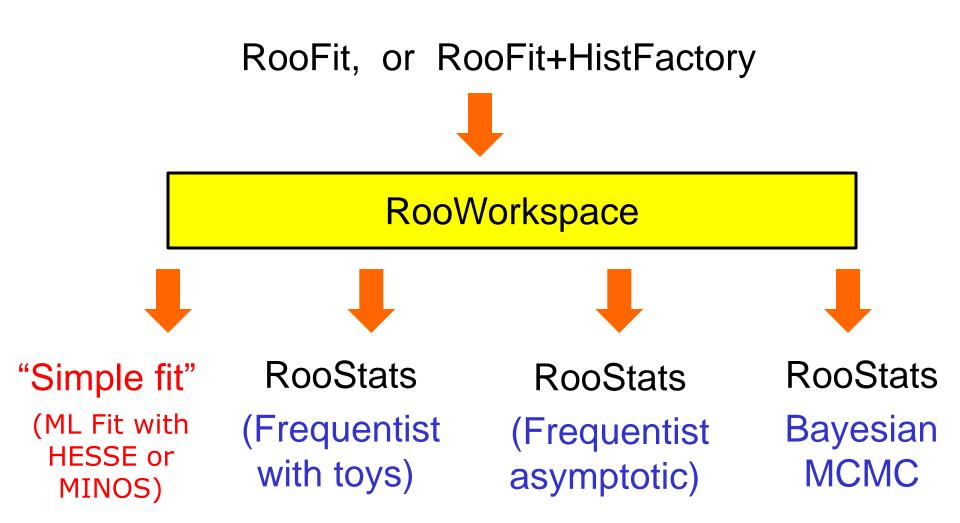
Allows full introspection and a-posteriori editing

Step 2 – Statistical tests on parameter of interest p

RooStats

The benefits of modularity

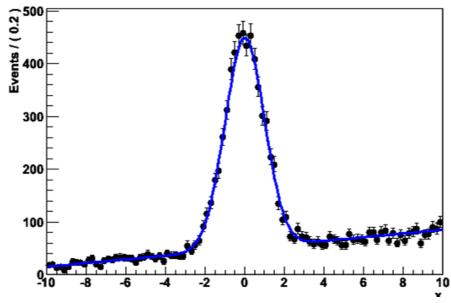
Perform different statistical test on exactly the same model



RooFit

RooFit – Focus: coding a probability density function

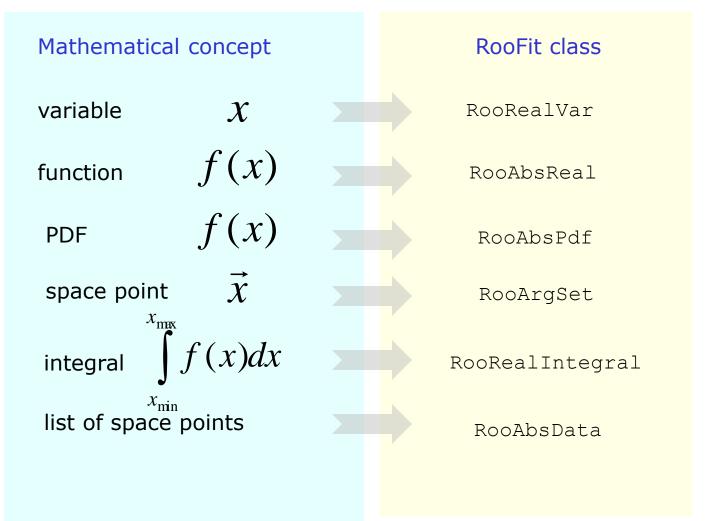
- Focus on one practical aspect of many data analysis in HEP: How do you formulate your p.d.f. in ROOT
 - For 'simple' problems (gauss, polynomial) this is easy
 - But if you want to do unbinned ML fits, use non-trivial functions, or work with multidimensional functions you quickly find that you need some tools to help you



 The RooFit project started in 1999 for data modeling needs for BaBar collaboration initially, publicly available in ROOT since 2003

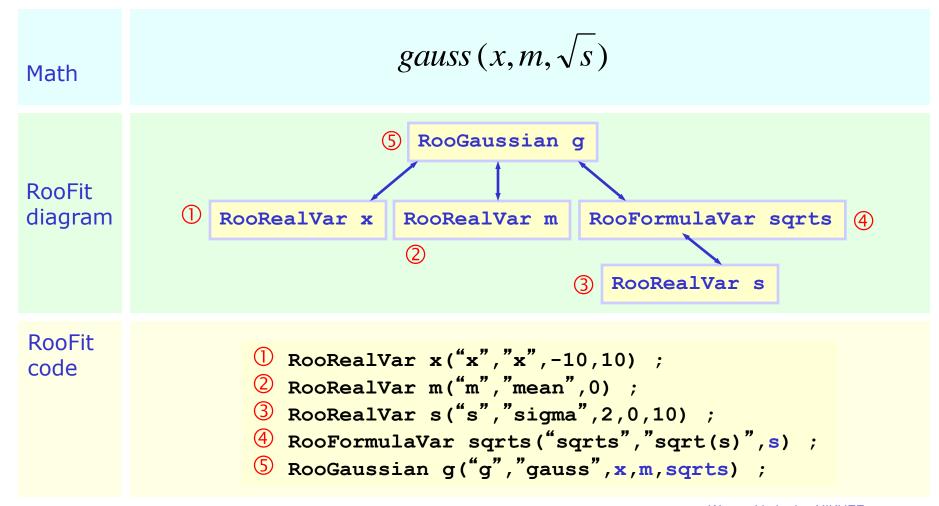
RooFit core design philosophy

Mathematical objects are represented as C++ objects



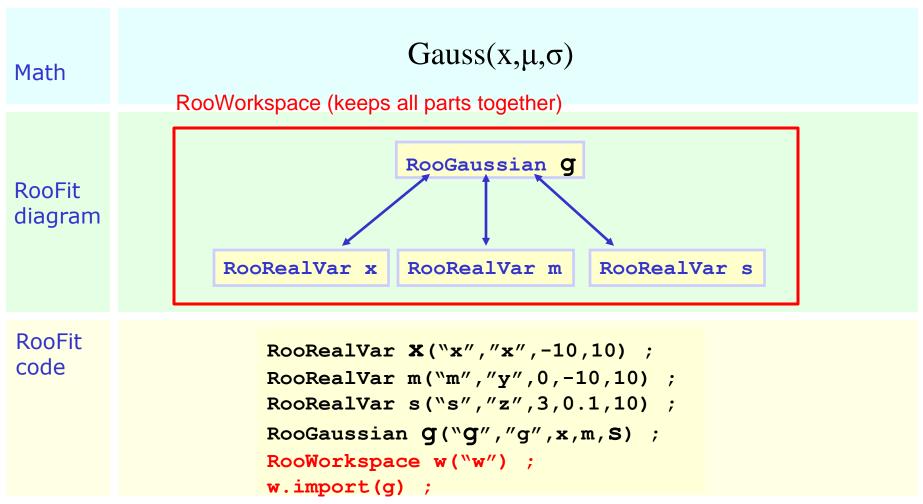
Data modeling – Constructing composite objects

Straightforward correlation between mathematical representation of formula and RooFit code



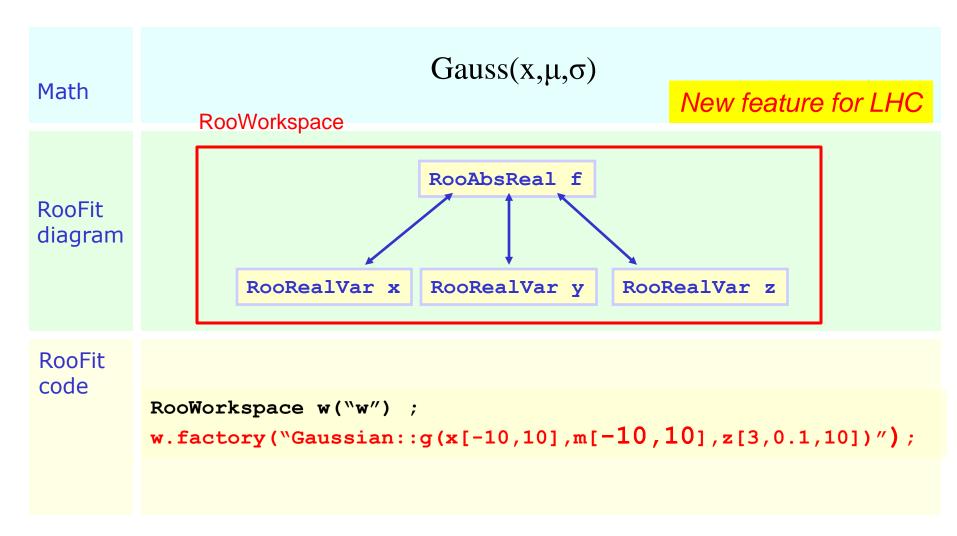
RooFit core design philosophy

 A special container class owns all objects that together build a likelihood function



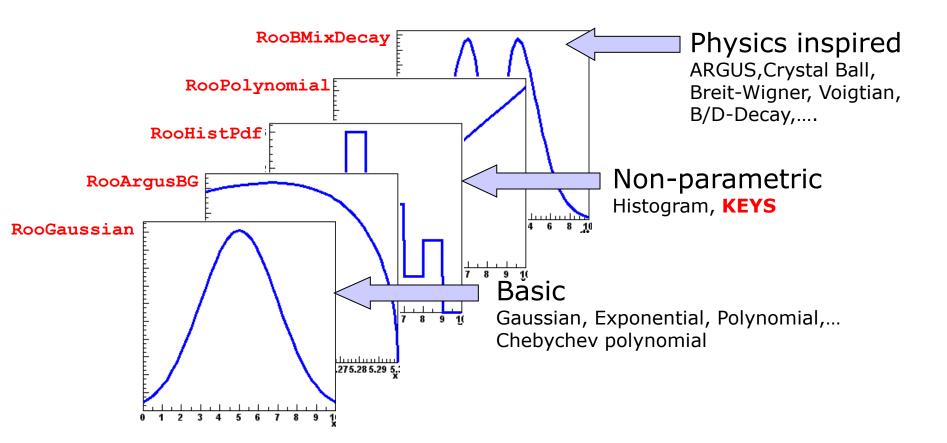
Populating a workspace the easy way – "the factory"

 The factory allows to fill a workspace with pdfs and variables using a simplified scripting language



Model building – (Re)using standard components

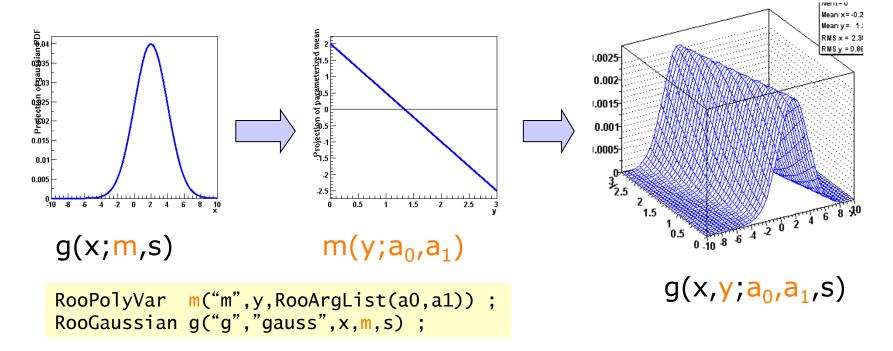
RooFit provides a collection of compiled standard PDF classes



Easy to extend the library: each p.d.f. is a separate C++ class

Model building – (Re)using standard components

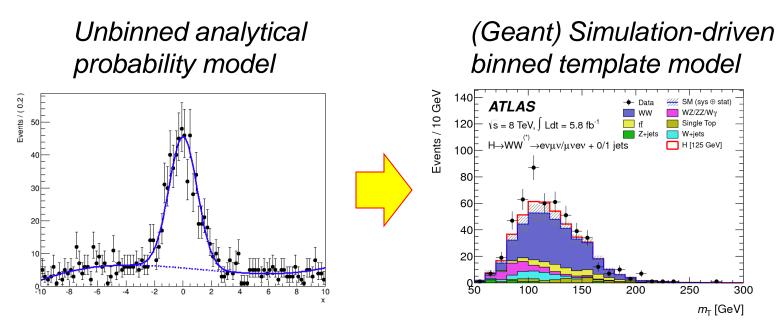
- Library p.d.f.s can be adjusted on the fly.
 - Just plug in any function expression you like as input variable
 - Works universally, even for classes you write yourself



Maximum flexibility of library shapes keeps library small

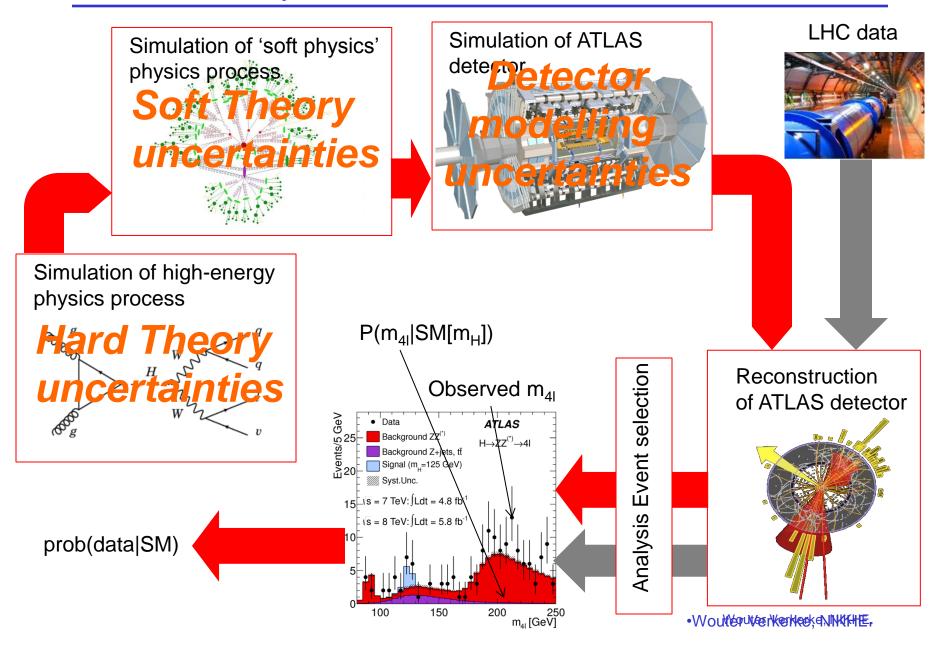
From empirical probability models to simulation-based models

 Large difference between B-physics and LHC hadron physics is that for the latter distributions usually don't follow simple analytical shapes



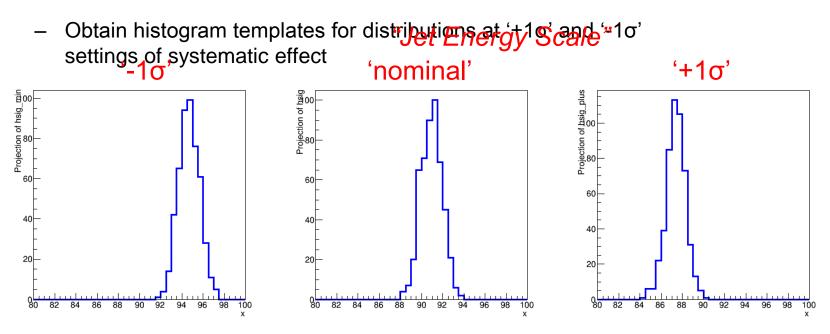
 But concept of simulation-driven template models can also extent to systematic uncertainties. Instead of empirically chosen 'nuisance parameters' (e.g. polynomial coefs) construct degrees of freedom that correspond to known systematic uncertainties

The HEP analysis workflow illustrated



Modeling of shape systematics in the likelihood

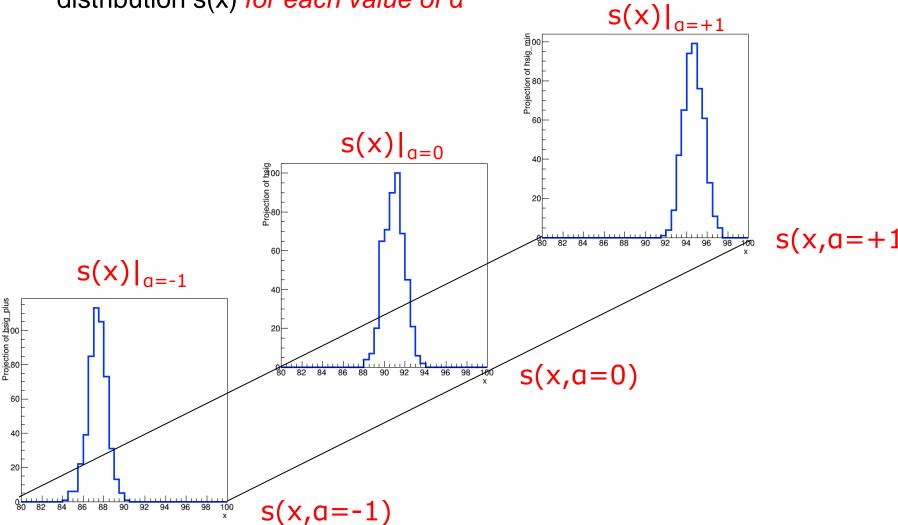
 Effect of any systematic uncertainty that affects the shape of a distribution can in principle be obtained from MC simulation chain



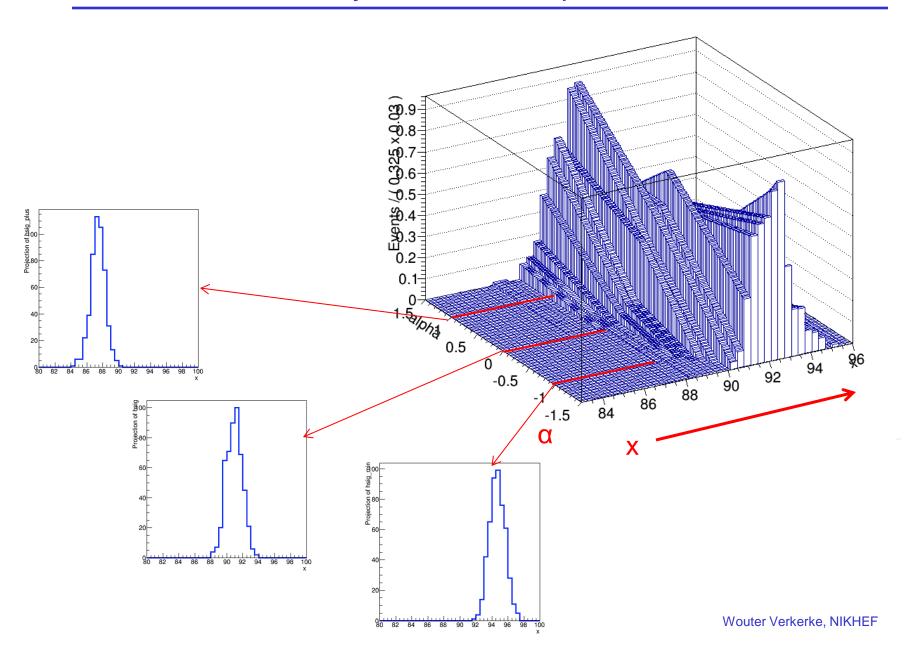
 Challenge: construct an empirical response function based on the interpolation of the shapes of these three templates.

Need to interpolate between template models

 Need to define 'morphing' algorithm to define distribution s(x) for each value of α



Visualization of bin-by-bin linear interpolation of distribution

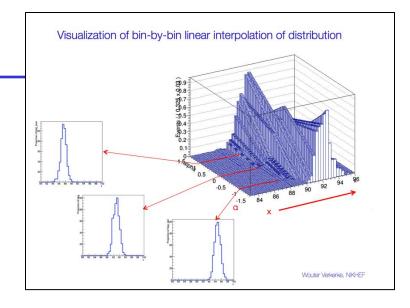


Example 2 : binned L with syst

 Example of template morphing systematic in a binned likelihood

$$s_{i}(\partial,...) = \int_{1}^{n} s_{i}^{0} + \partial \times (s_{i}^{+} - s_{i}^{0}) \quad " \quad \partial > 0$$

$$\downarrow \qquad \qquad \downarrow \qquad$$



```
// Import template histograms in workspace
w.import(hs_0,hs_p,hs_m);

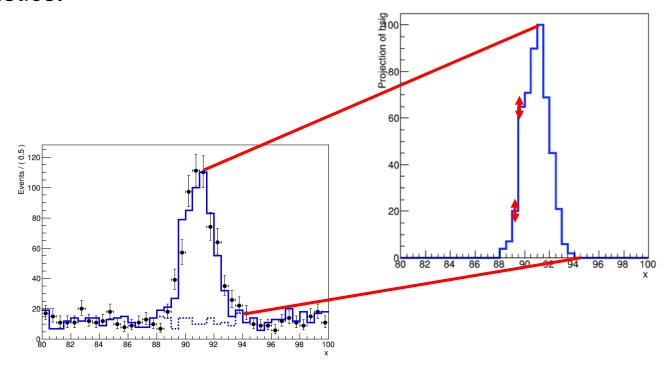
// Construct template models from histograms
w.factory("HistFunc::s_0(x[80,100],hs_0)");
w.factory("HistFunc::s_p(x,hs_p)");
w.factory("HistFunc::s_m(x,hs_m)");

// Construct morphing model
w.factory("PiecewiseInterpolation::sig(s_0,s_,m,s_p,alpha[-5,5])");

// Construct full model
w.factory("PROD::model(ASUM(sig,bkg,f[0,1]),Gaussian(0,alpha,1))");
```

Other uncertainties in MC shapes – finite MC statistics

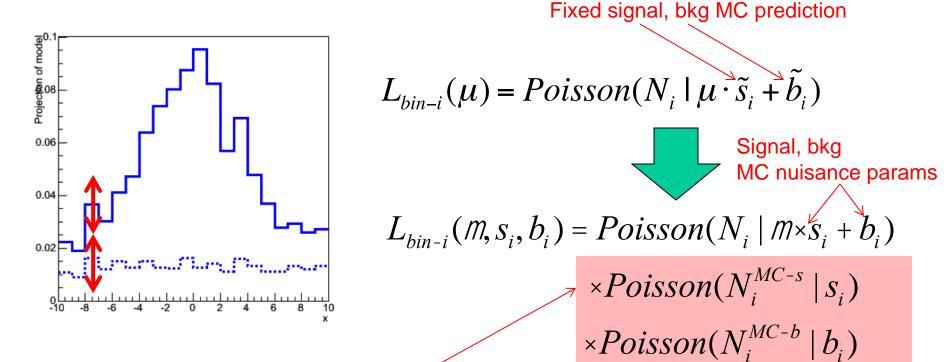
 In practice, MC distributions used for template fits have finite statistics.



Limited MC statistics represent an uncertainty on your model
 how to model this effect in the Likelihood?

Other uncertainties in MC shapes – finite MC statistics

- Modeling MC uncertainties: each MC bin has a Poisson uncertainty
- Thus, apply usual 'systematics modeling' prescription.
- For a single bin exactly like original counting measurement



Subsidiary measurement for signal MC ('measures' MC prediction s_i with Poisson uncertainty)

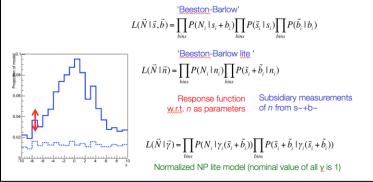
Code example – Beeston-Barlow

 Beeston-Barlow-(lite) modeling of MC statistical uncertainties

$$L(\vec{N} \mid \vec{g}) = \widetilde{O}_{bins} P(N_i \mid g_i(\tilde{s}_i + \tilde{b}_i)) \widetilde{O}_{bins} P(\tilde{s}_i + \tilde{b}_i \mid g_i(\tilde{s}_i + \tilde{b}_i))$$

Reducing the number NPs - Beeston-Barlow 'lite'

- Another approach that is being used is called 'BB' lite
- Premise: effect of statistical fluctuations on sum of templates is dominant → Use one NP per bin instead of one NP per component per bin



```
// Import template histogram in workspace
w.import(hs);

// Construct parametric template models from histograms
// implicitly creates vector of gamma parameters
w.factory("ParamHistFunc::s(hs)");

// Product of subsidiary measurement
w.factory("HistConstraint::subs(s)");

// Construct full model
w.factory("PROD::model(s,subs)");
```

Code example: BB + morphing

 Template morphing model with Beeston-Barlow-lite MC statistical uncertainties

$$s_{i}(\partial,...) = \int_{i}^{i} s_{i}^{0} + \partial \times (s_{i}^{+} - s_{i}^{0})$$
 " $\partial > 0$
 $s_{i}(\partial,...) = \int_{i}^{0} s_{i}^{0} + \partial \times (s_{i}^{0} - s_{i}^{-})$ " $\partial < 0$

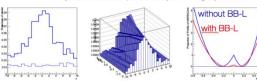
$$L(\vec{N} \mid \vec{s}, \vec{b}) = \underbrace{\tilde{O}}_{bins} P(N_i \mid g_i \times [s_i(a, s_i^-, s_i^0, s_i^+) + b_i]) \underbrace{\tilde{O}}_{bins} P(\tilde{s}_i + \tilde{b}_i \mid g_i \times [\tilde{s}_i + \tilde{b}_i]) G(0 \mid a, 1)$$

The interplay between shape systematics and MC systematics

• Commonly chosen practical solution $s_i(\alpha,...) = \begin{cases} s_i^0 + \alpha \cdot (s_i^+ - s_i^0) & \forall \alpha > 0 \\ s_i^0 + \alpha \cdot (s_i^0 - s_i^-) & \forall \alpha < 0 \end{cases}$

Morphing & MC response function Subsidiary measurements

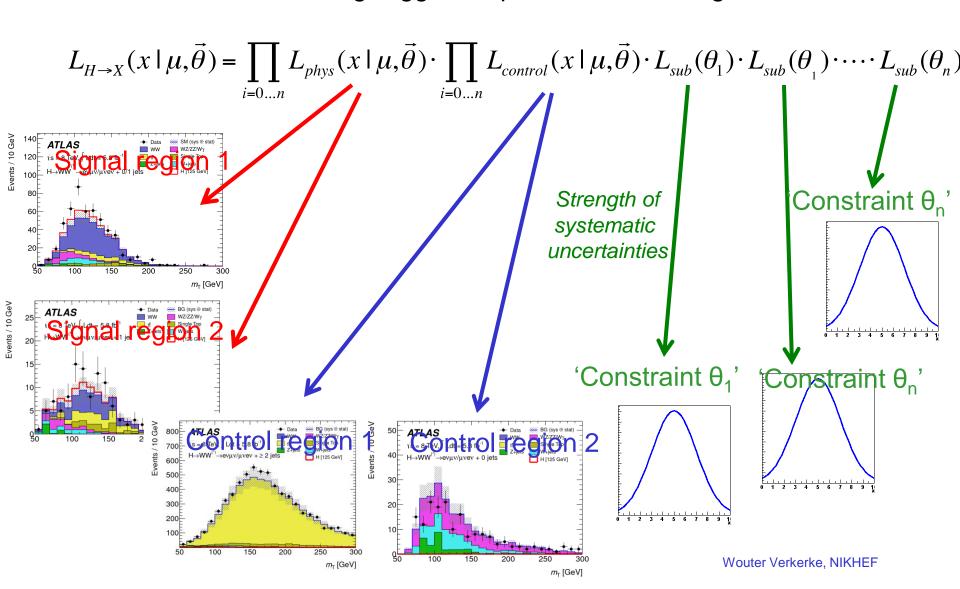
Models relative MC rate uncertainty for each bin w.r.t the nominal MC yield, even if morphed total yield is slightly different



- Approximate MC template statistics already significantly improves influence of MC fluctuations on template morphing
 - Because ML fit can now 'reweight' contributions of each bin

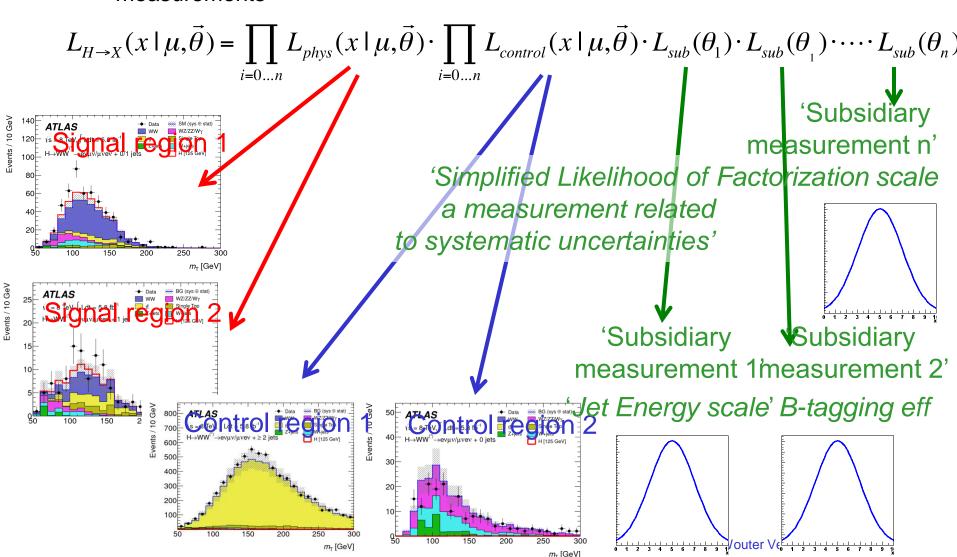
The structure of an (Higgs) profile likelihood function

Likelihood describing Higgs samples have following structure



The structure of an (Higgs) profile likelihood function

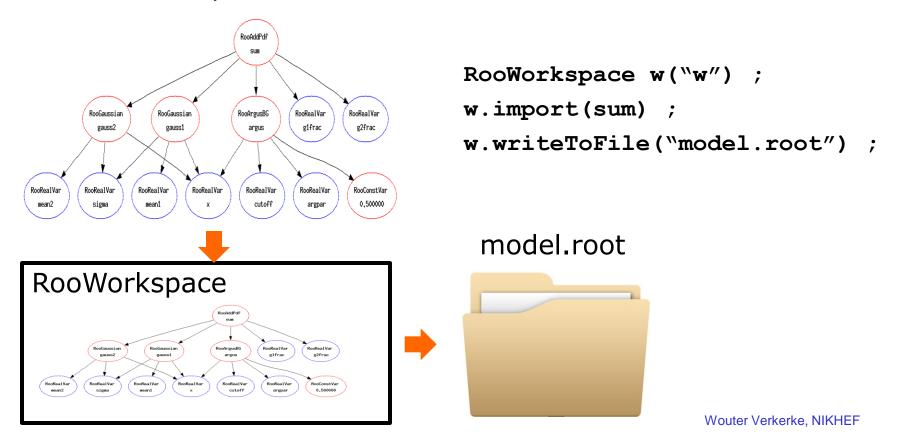
 A simultaneous fit of physics samples and (simplified) performance measurements



The Workspace

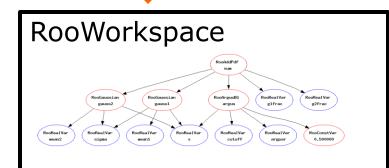
The workspace

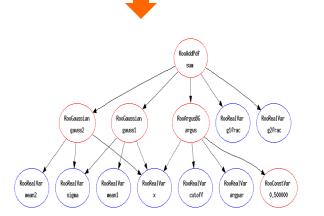
- The workspace concept has revolutionized the way people share and combine analysis
 - Completely factorizes process of building and using likelihood functions
 - You can give somebody an analytical likelihood of a (potentially very complex)
 physics analysis in a way to the easy-to-use, provides introspection, and is easy
 to modify.



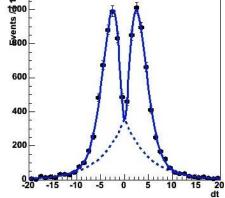
Using a workspace







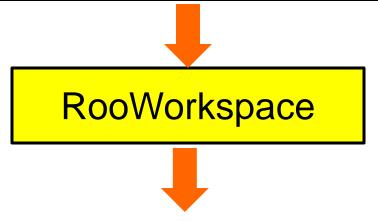
```
// Resurrect model and data
TFile f("model.root") ;
RooWorkspace* w = f.Get("w") ;
RooAbsPdf* model = w->pdf("sum") ;
RooAbsData* data = w->data("xxx") ;
// Use model and data
model->fitTo(*data) ;
RooPlot* frame =
         w->var("dt")->frame();
data->plotOn(frame) ;
model->plotOn(frame) ;
   <del>[0</del>00
```



The idea behind the design of RooFit/RooStats/HistFactory

• Step 1 – Construct the likelihood function L(x|p)

```
RooWorkspace w("w");
w.factory("Gaussian::sig(x[-10,10],m[0],s[1])");
w.factory("Chebychev::bkg(x,a1[-1,1])");
w.factory("SUM::model(fsig[0,1]*sig,bkg)");
w.writeToFile("L.root");
```



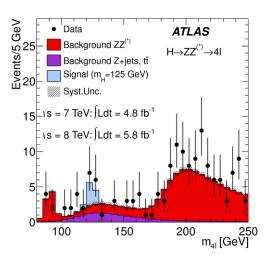
Complete description of likelihood model, persistable in ROOT file (RooFit pdf function) Allows full introspection and a-posteriori editing

Step 2 – Statistical tests on parameter of interest p

```
RooWorkspace* w=TFile::Open("L.root")->Get("w") ;
RooAbsPdf* model = w->pdf("model") ;
pdf->fitTo(data) ;
```

Example RooFit component model for realistic Higgs analysis

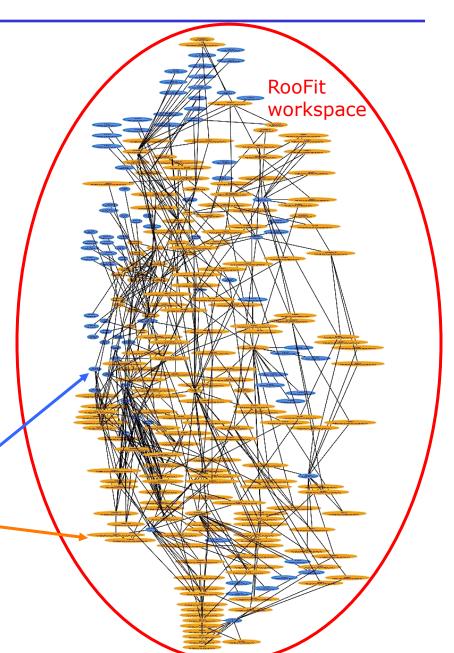
Likelihood model describing the ZZ invariant mass distribution including all possible systematic uncertainties



variables

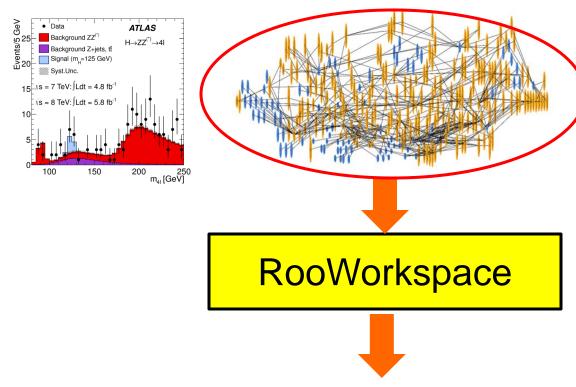
function objects

Graphical illustration of function components that call each other



Analysis chain identical for highly complex (Higgs)

• Step 1 – Construct the likelihood function L(x|p)



Complete description of likelihood model, persistable in ROOT file (RooFit pdf function) Allows full introspection and a-posteriori editing

Step 2 – Statistical tests on parameter of interest p

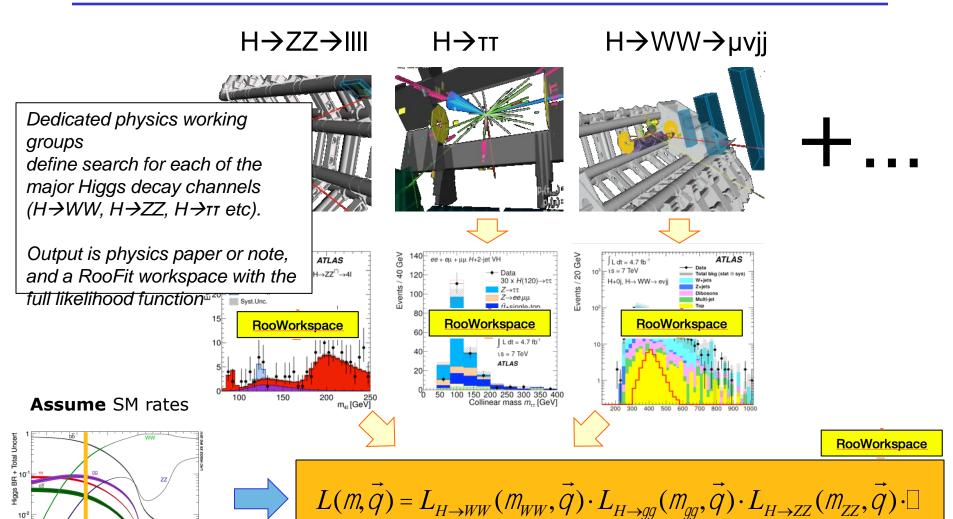
Workspaces power collaborative statistical modelling

- Ability to persist complete^(*) Likelihood models has profound implications for HEP analysis workflow
 - (*) Describing signal regions, control regions, and including nuisance parameters for all systematic uncertainties)
- Anyone with ROOT (and one ROOT file with a workspace) can re-run any entire statistical analysis out-of-the-box
 - About 5 lines of code are needed
 - Including estimate of systematic uncertainties
- Unprecedented new possibilities for cross-checking results, in-depth checks of structure of analysis
 - Trivial to run variants of analysis (what if 'Jet Energy Scale uncertainty' is 7% instead of 4%). Just change number and rerun.
 - But can also make structural changes a posteri. For example, rerun with assumption that JES uncertainty in forward and barrel region of detector are 100% correlated instead of being uncorrelated.

Collaborative statistical modelling

- As an experiment, you can effectively build a library of measurements, of which the full likelihood model is preserved for later use
 - Already done now, experiments have such libraries of workspace files,
 - Archived in AFS directories, or even in SVN....
 - Version control of SVN, or numbering scheme in directories allows for easy validation and debugging as new features are added
- Building of <u>combined</u> likelihood models greatly simplified.
 - Start from persisted components. No need to (re)build input components.
 - No need to know how individual components were built, or are internally structured. Just need to know meaning of parameters.
 - Combinations can be produced (much) later than original analyses.
 - Even analyses that were never originally intended to be combined with anything else can be included in joint likelihoods at a later time

Higgs discovery strategy – add everything together

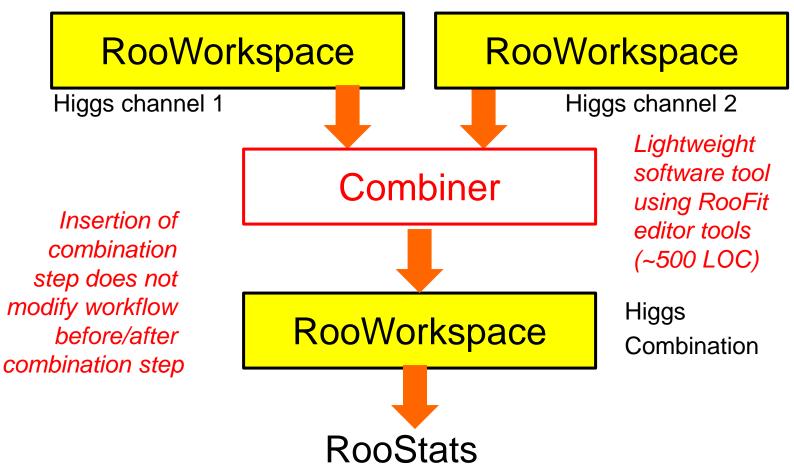


A small dedicated team of specialists builds a combined likelihood from the inputs. Major discussion point: naming of parameters, choice of parameters for systematic uncertainties (a physics issue, largely)

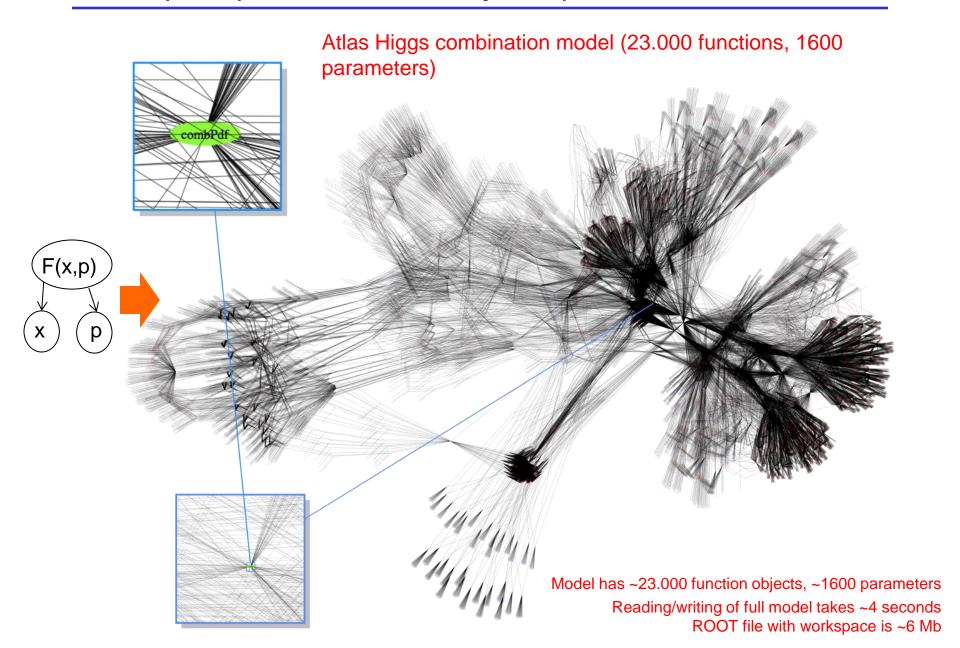
The benefits of modularity

Technically very straightforward to combine measurements

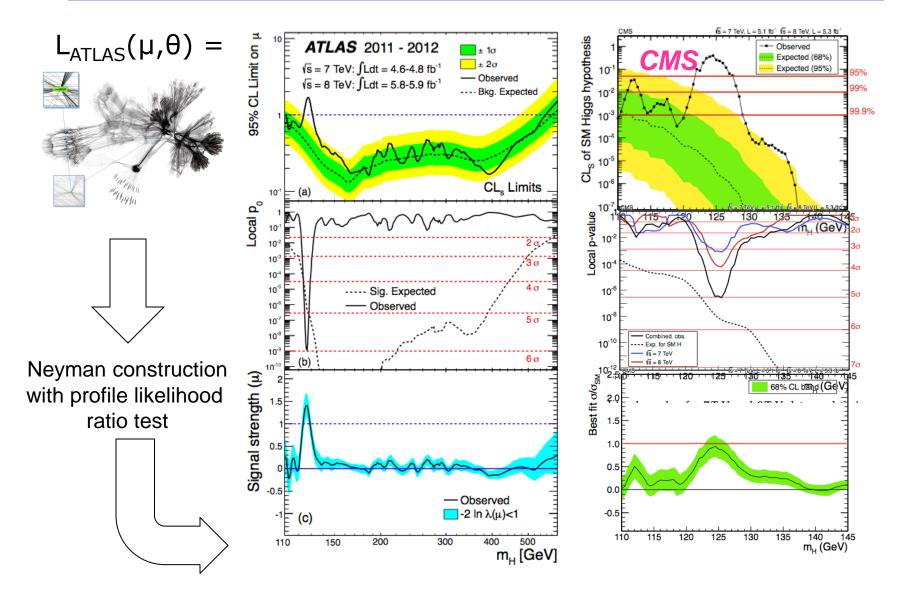
RooFit, or RooFit+HistFactory



Workspace persistence of *really* complex models works too!



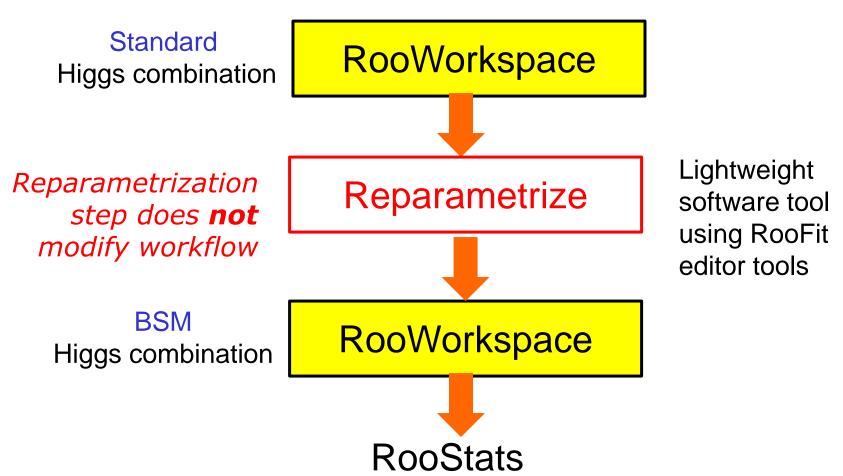
With these combined models the Higgs discovery plots were produced...



More benefits of modularity

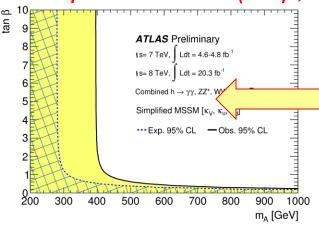
Technically very straightforward to reparametrize measurements

RooFit, or RooFit+HistFactory

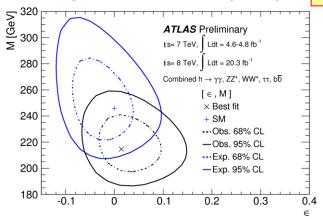


BSM Higgs constraints from reparametrization of SM Higgs Likelihood model

Simplified MSSM (tanβ,m_A)

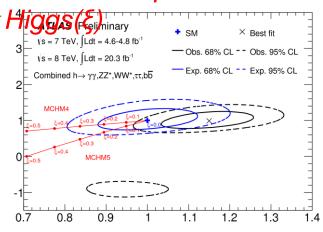


Imposter model(M,ε)



(ATLAS-CONF-2014-010)





Two Higgs Double Model (tanβ,cos(α-β))

Portal model (mx

DAMA/LIBRA (99.7% CL) CRESST (95% CL)

CDMS (95% CL) CoGeNT (90% CL)

XENON10 (90% CL)

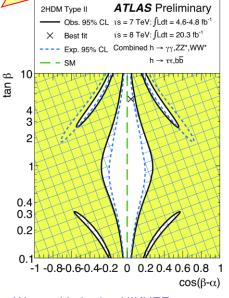
ATLAS Preliminary

 $\sqrt{s} = 7 \text{ TeV}$, $\int L dt = 4.6-4.8 \text{ fb}$

Majorana WIMP

m_χ [GeV]

 $\sqrt{s} = 8 \text{ TeV}, \int L dt = 20.3 \text{ fb}^{-1}$ $h \rightarrow yy, h \rightarrow ZZ^* \rightarrow 4l, h \rightarrow WW^* \rightarrow lyly, l$



Wouter Verkerke, NIKHEF

An excursion – Collaborative analyses with workspaces

- How can you reparametrize existing Higgs likelihoods in practice?
- Write functions expressions corresponding to new parameterization

$$\sigma(gg \to H) * BR(H \to \gamma \gamma) \sim \frac{\kappa_F^2 \cdot \kappa_\gamma^2(\kappa_F, \kappa_V)}{0.75 \cdot \kappa_F^2 + 0.25 \cdot \kappa_V^2}$$
w.factory("expr::mu_gg_func('(KF2*Kg2)/(0.75*KF2+0.25*KV2)', KF2,Kg2,KV2);

Import transformation in workspace, edit existing model

```
w.import(mu_gg_func) ;
w.factory("EDIT::newmodel(model,mu_gg=mu_gg_gunc)") ;
```

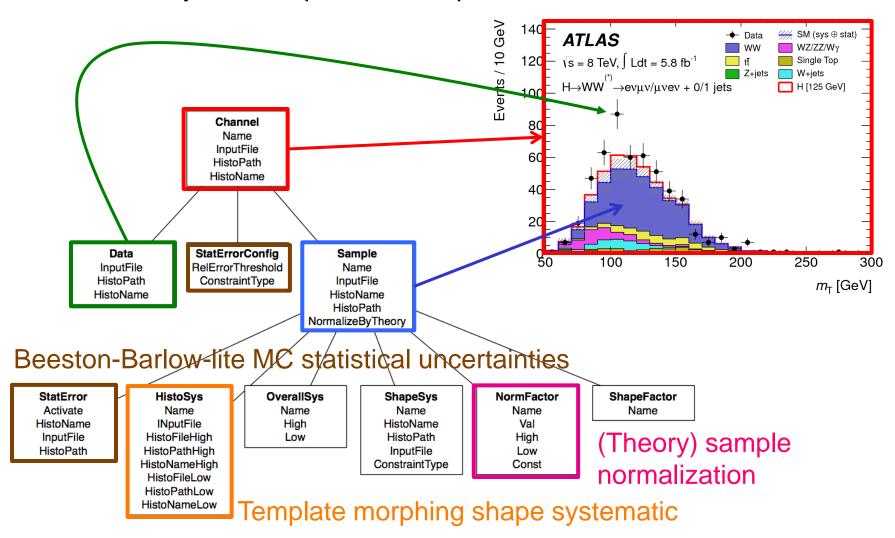
HistFactory

HistFactory – structured building of binned template models

- RooFit modeling building blocks allow to easily construct likelihood models that model shape and rate systematics with one or more nuisance parameter
 - Only few lines of code per construction
- Typical LHC analysis required modeling of 10-50 systematic uncertainties in O(10) samples in anywhere between 2 and 100 channels → Need structured formalism to piece together model from specifications. This is the purpose of HistFactory
- HistFactory conceptually similar to workspace factory, but has much higher level semantics
 - Elements represent physics concepts (channels, samples, uncertainties and their relation) rather than mathematical concepts
 - Descriptive elements are represented by C++ objects (like roofit),
 and can be configured in C++, or alternively from an XML file
- HistFactory builds a RooFit (mathematical) model from a physics model.

HistFactory elements of a channel

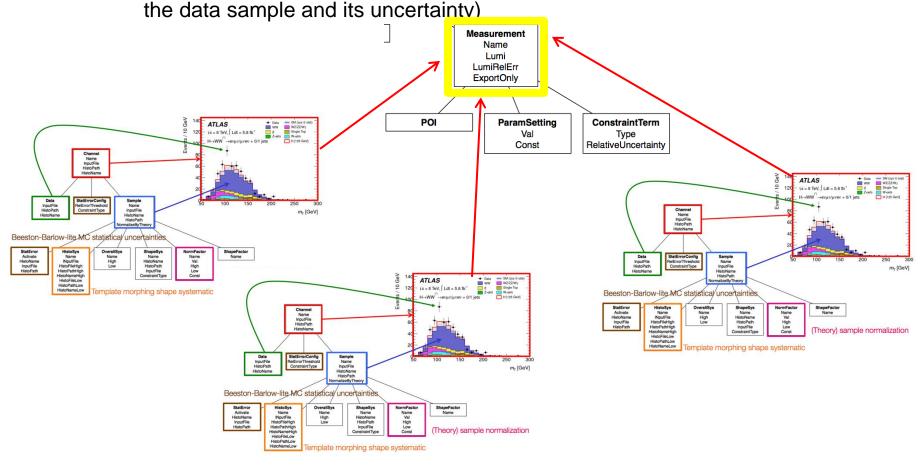
Hierarchy of concepts for description of one measurement channel



HistFactory elements of measurement

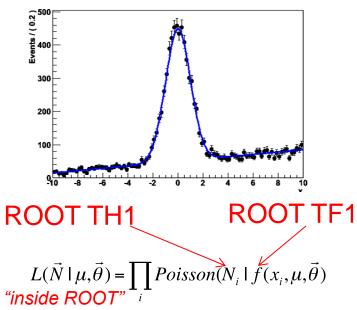
One or more channels are combined to form a measurement

 Along with some extra information (declaration of the POI, the luminosity of the data cample and its upportainty)



How is Higgs discovery different from a simple fit?

Gaussian + polynomial





Maximum Likelihood estimation of parameters μ,θ using MINUIT (MIGRAD, HESSE, MINOS)

$$\mu = 5.3 \pm 1.7$$

Likelihood Model orders of magnitude more complicated. Describes

- O(100) signal distantions
- O(100) tro' ple distr.
- O(1000 eters esenting syst. uncertainties



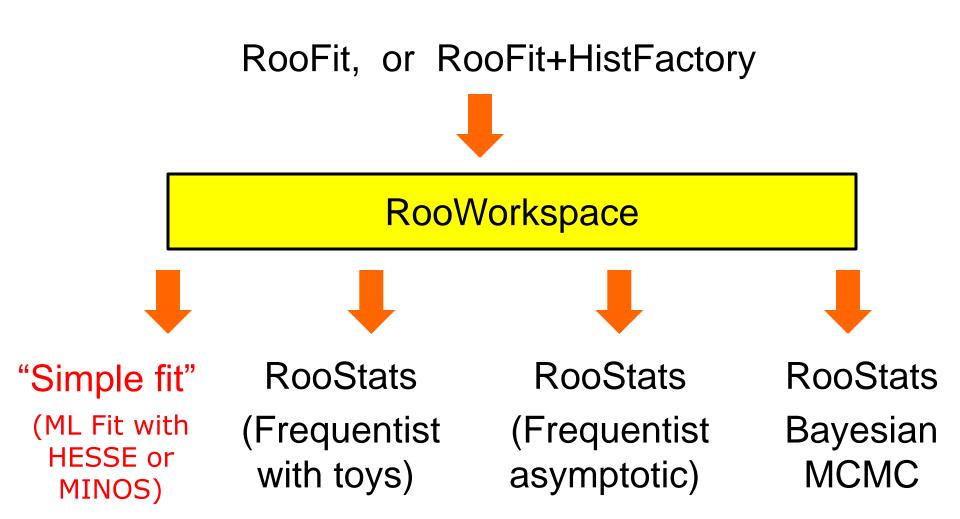


Frequentist confidence interval construction and/or p-value calculation not available as 'ready-to-run' algorithm in ROOT

RooStats

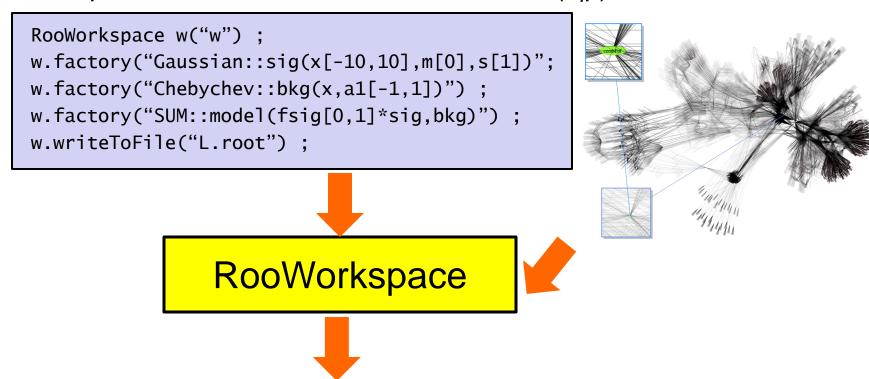
The benefits of modularity

Perform different statistical test on exactly the same model



Maximum Likelihood estimation as simple statistical analysis

Step 1 – Construct the likelihood function L(x|p)



Step 2 – Statistical tests on parameter of interest p

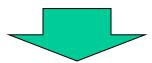
```
RooWorkspace* w=TFile::Open("L.root")->Get("w") ;
RooAbsPdf* model = w->pdf("model") ;
pdf->fitTo(data) ;
```

The need for fundamental statistical techniques

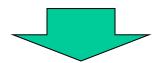
Frequentist statistics

Bayesian statistics

Maximum Likelihood



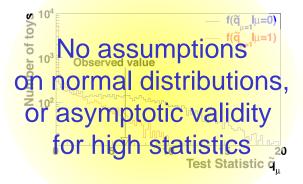
$$I_{m}(\vec{N}_{obs}) = \frac{L(\vec{N} \mid m)}{L(\vec{N} \mid \hat{m})}$$

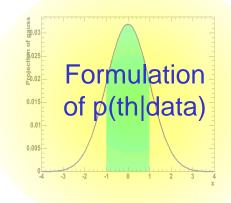


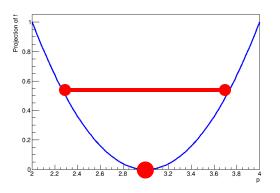
$$P(m) \sqcup L(x \mid m) \times p(m)$$



$$\frac{d \ln L(\vec{p})}{d\vec{p}}\bigg|_{p_i = \hat{p}_i} = 0$$





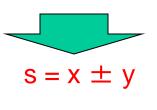




Confidence interval or p-value



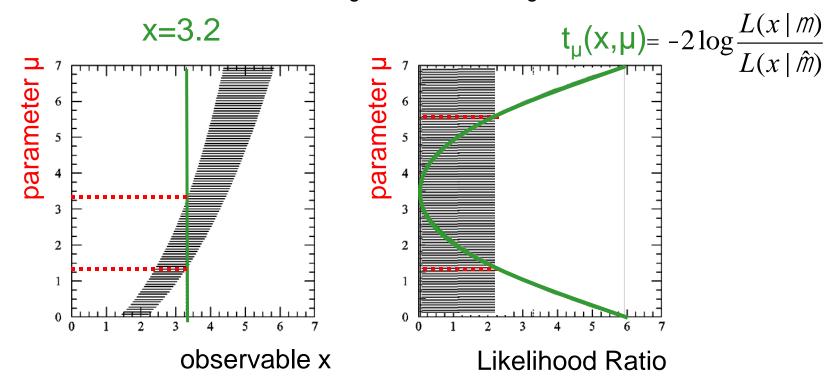
Posterior on s or Bayes factor



Wouter Verkerke, NIKHEF

But fundamental techniques can be complicated to

- Example of confidence interval calculation with Neyman construction
 - Need to construct 'confidence belt' using toy MC. Intersection observed data with belt defined interval in POI with guaranteed coverage



- Expensive, complicated procedure, but completely procedural once Likelihood and parameter of interest are fixed
 - → Can be wrapped in a tool that runs effectively 'out-of-the-box'

Running RooStats interval calculations 'out-of-the-box'

- Confidence intervals calculated with model
 - 'Simple Fit'

```
RooAbsReal* nll = myModel->createNLL(data) ;
RooMinuit m(*nll) ;
m.migrad() ;
m.hesse() ;
```

Feldman
 Cousins
 (Frequentist
 Confidence
 Interval)

```
FeldmanCousins fc;
fc.SetPdf(myModel);
fc.SetData(data); fc.SetParameters(myPOU);
fc.UseAdaptiveSampling(true);
fc.FluctuateNumDataEntries(false);
fc.SetNBins(100); // number of points to test per parameter
fc.SetTestSize(.1);
ConfInterval* fcint = fc.GetInterval();
```

Bayesian (MCMC)

```
UniformProposal up;
MCMCCalculator mc;
mc.SetPdf(w::PC);
mc.SetData(data); mc.SetParameters(s);
mc.SetProposalFunction(up);
mc.SetNumIters(100000); // steps in the chain
mc.SetTestSize(.1); // 90% CL
mc.SetNumBins(50); // used in posterior histogram
mc.SetNumBurnInSteps(40);
ConfInterval* mcmcint = mc.GetInterval();
```

But you can also look 'in the box' and build your own

Tool to calculate p-values for a given hypothesis $\int f(q_{\mu} \mid \mu') dq_{\mu}$

```
// create first HypoTest calculator (N.B null is s+b model)
FrequentistCalculator fc(*data, *bModel, *sbModel);
// configure ToyMCSampler and set the test statistics
ToyMCSampler *toymcs = (ToyMCSampler*)fc.GetTestStatSampler();
ProfileLikelihoodTestStat profll(*sbModel->GetPdf()) 
// for CLs (bounded intervals) use one-sided profile likelihood
profll.SetOneSided(true);
toymcs->SetTestStatistic(&profll);
HypoTestInverter calc(*fc) 
calc.UseCLs(true);
// configure and run the scan
calc.SetFixedScan(npoints,poimin,poimax);
HypoTestInverterResult * r = calc.GetInterval()
// get result and plot it
double upperLimit = r->UpperLimit();
double expectedLimit = r->GetExpectedUpperLimit(0);
HypoTestInverterPlot *plot = new HypoTestInverterPlot("hi","",r);
plot->Draw();
```

Offset advanced control over details of statistical procedure (use of CLS, choice of test statistic, boundaries...)

$$f(q_{\mu} | \mu')$$

 $q_{u,obs}$

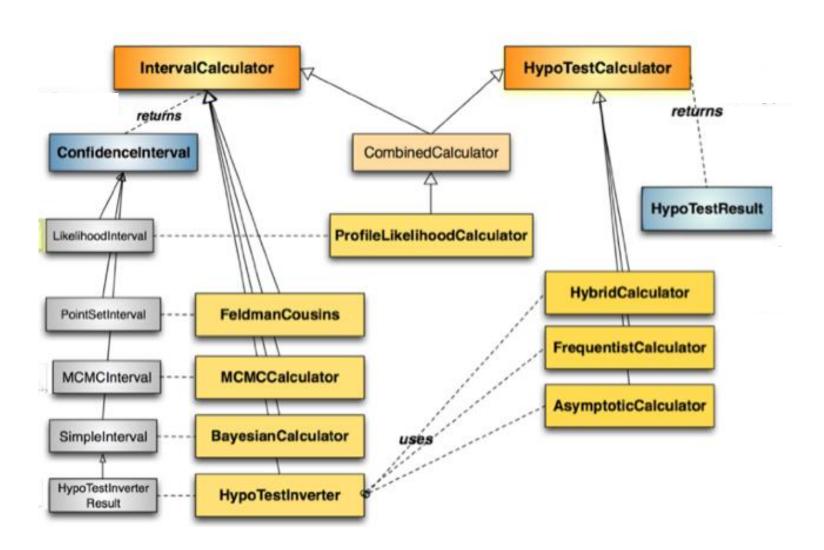
Tool to construct test statistic distribution

$$q_{\mu}(\mu')$$

The test statistic to be used for the calculation of p-values

Tool to construct interval from hypo test results

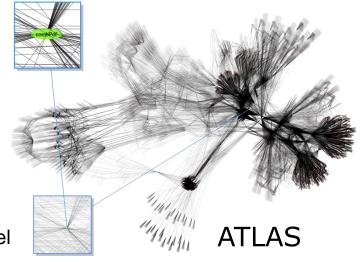
RooStats class structure

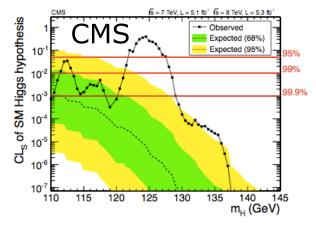


Summary

RooFit and RooStats allow you to perform advanced statistical data analysis

- LHC Higgs results a prominent example
- RooFit provides (almost) limitless model building facilities
 - Concept of persistable model workspace allows to separate model building and model interpretation
 - HistFactory package introduces structured model building for binned likelihood template models that are common in LHC analyses
- Concept of RooFit Workspace has completely restructured HEP analysis workflow with 'collaborative modeling'
- RooStats provide a wide set of statistical tests that can be performed on RooFit models
 - Bayesian, Frequentist and Likelihood-based test concepts





Wouter Verkerke, NIKHEF

The future - physics

- Many more high-profile RooFit/RooStats full likelihood combinations in the works
 - Combination of ATLAS and CMS Higgs results
 - CMS/LHC combination of rare B-decays
- But many more combinations are easily imaginable & feasible
 - Combination across physics domains (e.g. SUSY and Higgs, or Exotics and Higgs) → reparametrization allows to constrain parameters of BSM physics models that have features in both domains (e.g. 2 Higgs Doublet Models)
 - Incorporation of more sophisticated models for detector performance measurements (now often simple Gaussians).

Many ideas ongoing (e.g eigenvector diagonalization of calibration uncertainties across p_T bins \rightarrow less parameters with correlated subsidiary measurement), modeling of correlated effects between systematic uncertainties (e.g. Jet energy scales and flavor tagging)

The future - computing

- Technical scaling and performance generally unproblematic
 - MINUIT has been shown to still work with 10.000 parameters, but do you really need so much detail?
 - Persistence works miraculously well, given complexity of serialization problem
 - Algorithmic optimization of likelihood calculations works well
 - Likelihood calculations trivially parallelizable. But more work can be done here (e.g. portability of calculations to GPUs, taking advantage of modern processor architectures for vectorization)
 - Bayesian algorithms still need more development and tuning
- But physicists are very good and pushing performance and scalability to the limits
 - Generally, one keep adding features and details until model becomes 'too slow'
 - But if every Higgs channel reaches this point on its own, a channel combination is already 'way too slow' from the onset
 - Need to learn how to limit complexity → Prune irrelevant details from physics models, possibly a posteriori. Work in progress, some good ideas around
- Looking forward to LHC Run-2