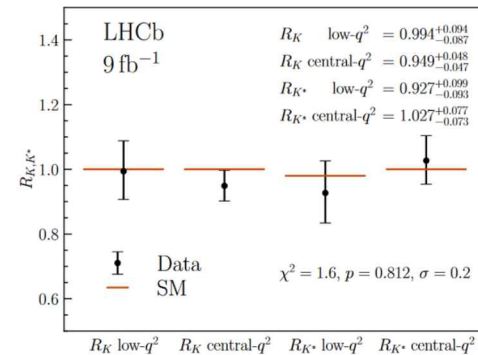
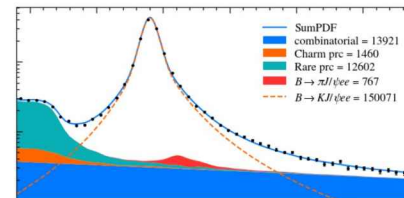
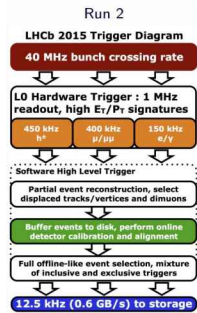
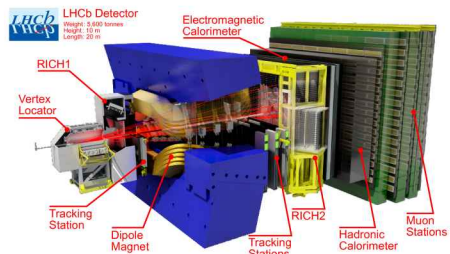
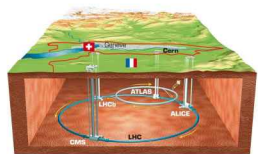
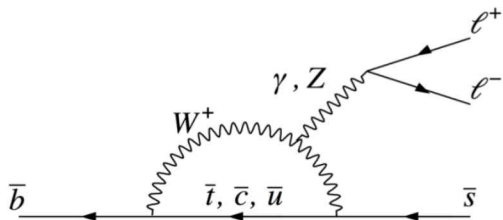


Model fitting in Python with zfit and Scikit-HEP

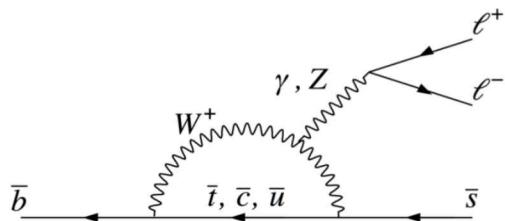
Jonas Eschle

jonas.eschle@cern.ch

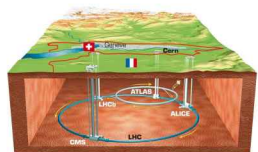
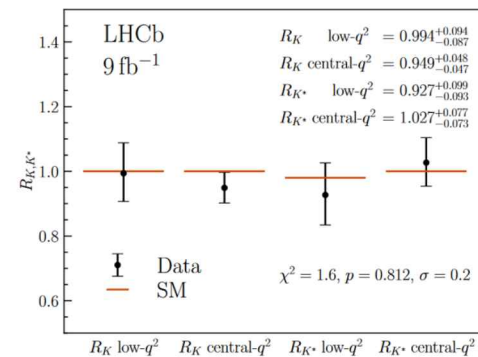
HEP Analysis



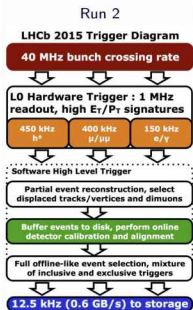
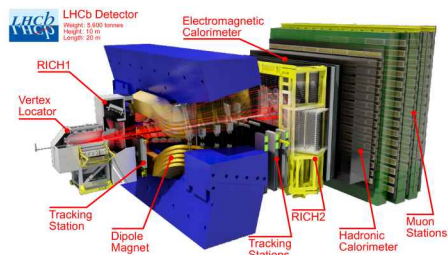
HEP Analysis



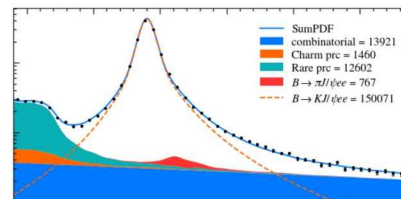
Lots of code



Lots of code

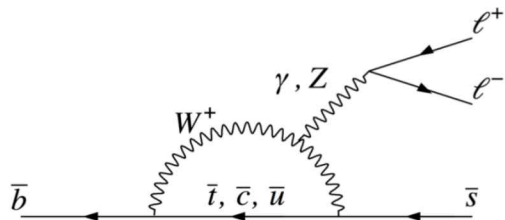


Lots of code

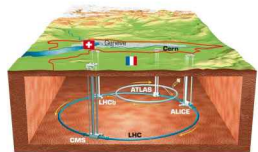
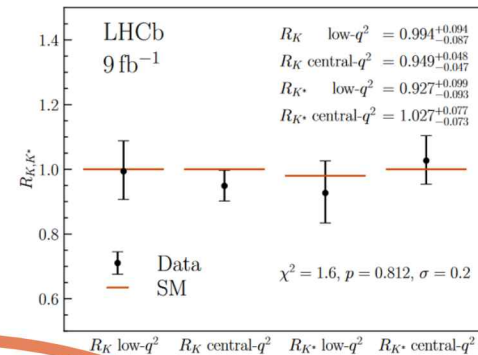


Lots of code

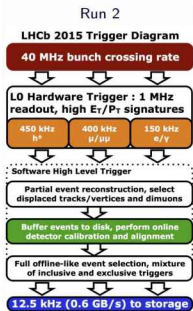
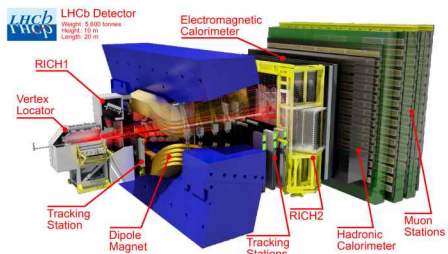
HEP Analysis



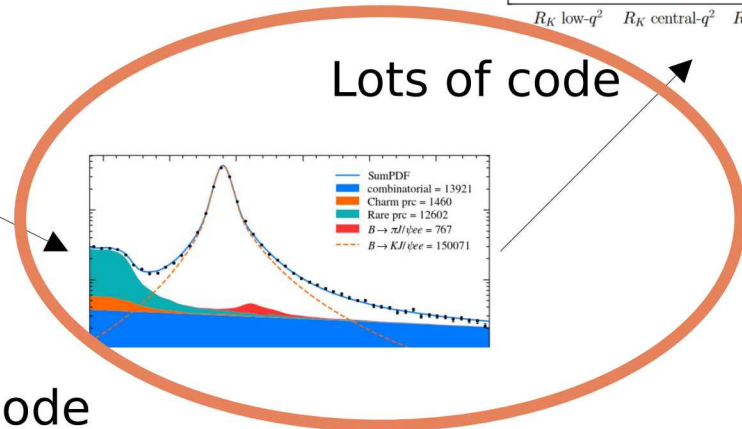
Lots of code



Lots of code



Lots of code



Lots of code

End-user analysis

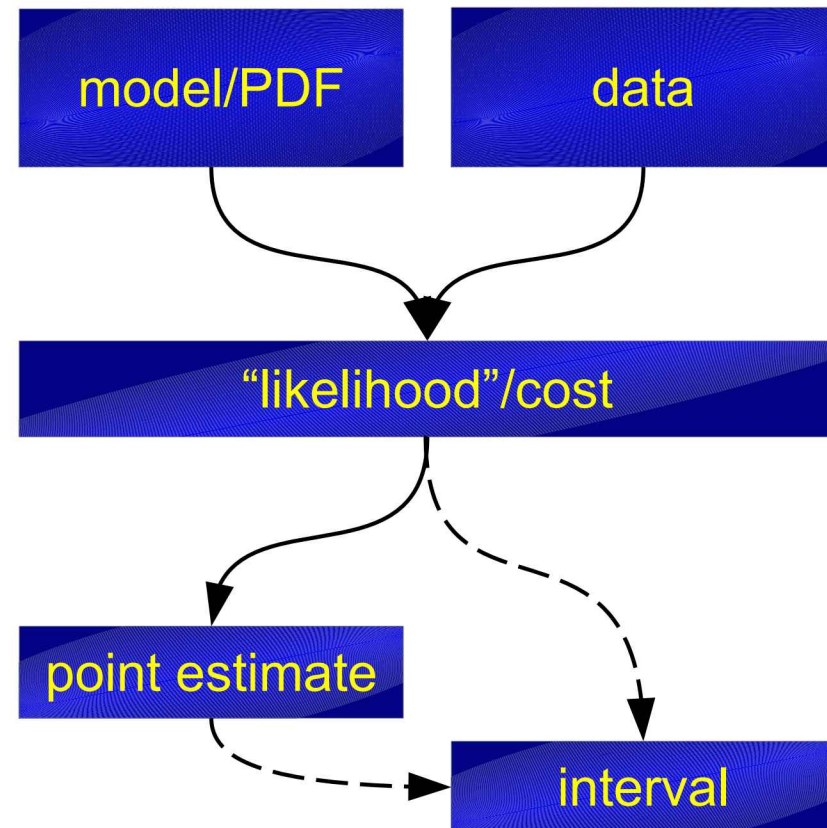
Focus on «fitting»

«Fitting»

«statistical inference»

Variety of possibilities

Focus on likelihood based
(as is very common in HEP)



Historic landscape



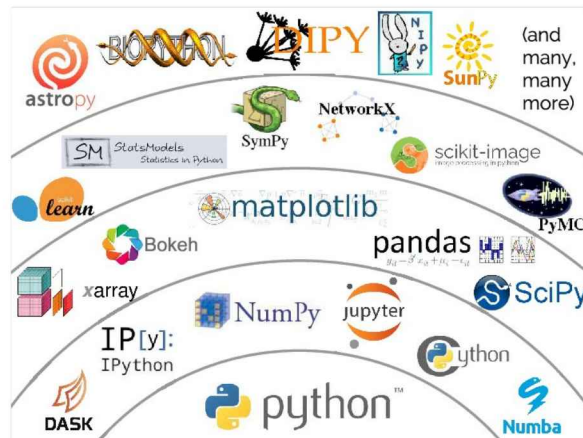
Analyses transition from C++ to Python

Many, non-monolithic packages

Talk by Eduardo



Philosophy: extend/build on existing ecosystem



2018



Historic landscape



Analyses transition from C++ to Python

Many, non-monolithic packages

Talk by Eduardo



HEP fitting libraries still in C++

Strong libraries, but mediocre bindings, not «pythonic»

2018



Historic landscape



Analyses transition from C++ to Python

Many, non-monolithic packages

Talk by Eduardo



HEP fitting libraries still in C++

Strong libraries, but mediocre bindings, not «pythonic»

Why even move to Python?

2018



Historic landscape

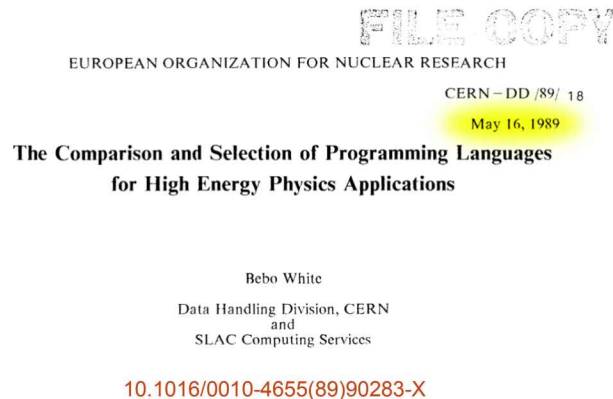


Analyses transition from C++ to Python Many, non-monolithic packages

Talk by Eduardo



HEP fitting libraries still in C++
Strong libraries, but mediocre bindings, not «pythonic»



Zanella [32] has said " If HEP wishes to keep to its level of achievement, credibility and excellence, then it needs an injection of bright young computer-wise scientists and engineers." This means that HEP cannot become "an island." HEP applications must be able to utilize "state of the art" facilities in all areas of applicability including data processing. HEP must be able to take advantage of the technological advancements in other arenas of science and engineering. Many of these advancements are occurring in fields which are presently *not software compatible* with HEP. Much of the work being done in embedded systems with Ada or telecommuni-

2018



Historic landscape



Analyses transition from C++ to Python

Many, non-monolithic packages

Talk by Eduardo

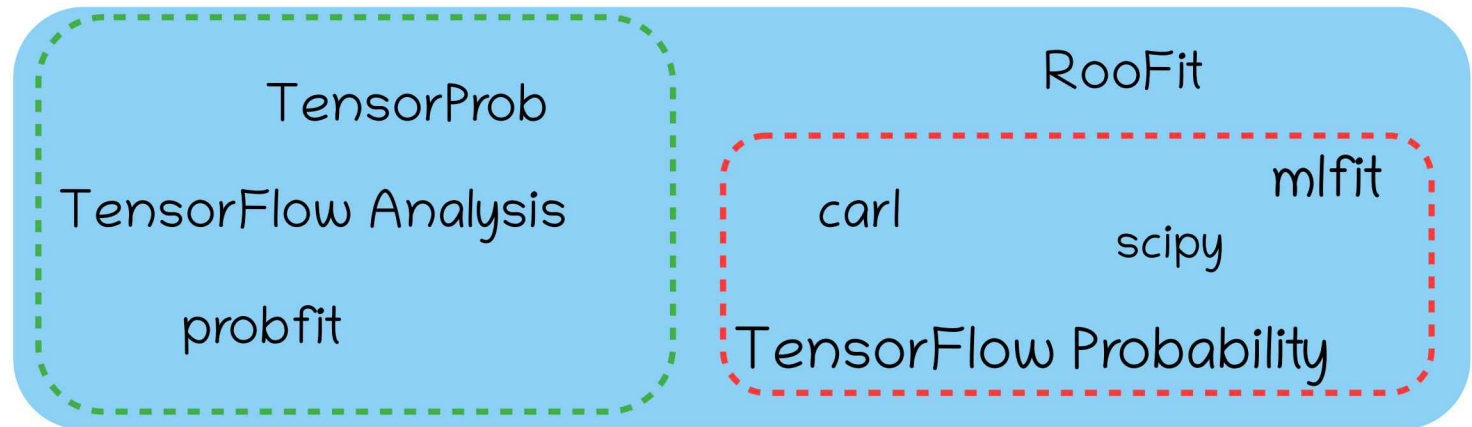


HEP fitting libraries still in C++

Strong libraries, but mediocre bindings, not «pythonic»

Python packages

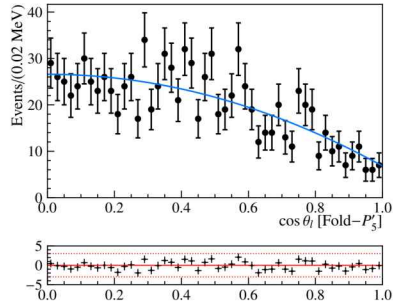
- no advanced features
- «Python too slow»



2018



HEP Model Fitting in Python



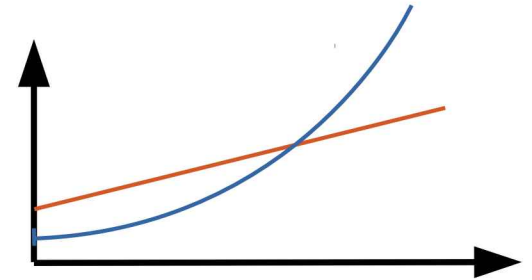
HEP

advanced features,
simply extendable



Scalable

large data, complex models



Pythonic  NumPy  python™
integrate into ecosystem, stable API

History



5 years ago

first presentation
HSF/WLCG
Workshop

A presentation slide for zfit. The background is dark with a complex, glowing pattern of blue and white lines, resembling a particle detector or a data visualization. The text is white and orange. The logo 'zfit' is large and white, with 'scalable pythonic fitting' below it. The name 'Jonas Eschle' is in large white font, with his email 'jonas.eschle@cern.ch' below it. The text 'In collaboration with A. Puig, R. S. Coutinho, N. Serra' is in a smaller white font. The University of Zurich logo and 'University of Zurich UZH' are on the left. The FNSNF logo and 'SWISS NATIONAL SCIENCE FOUNDATION' are at the bottom left.

zfit
scalable pythonic fitting

Jonas Eschle
jonas.eschle@cern.ch

In collaboration with
A. Puig, R. S. Coutinho, N. Serra

 **University of Zurich** ^{UZH}

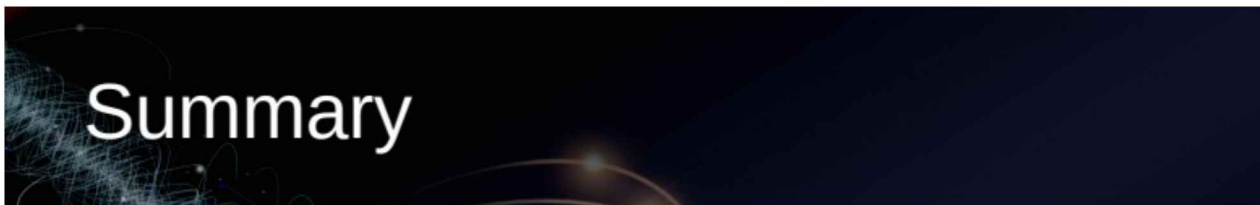
 **FNSNF**
SWISS NATIONAL SCIENCE FOUNDATION

History



5 years ago

first presentation
HSF/WLCG
Workshop



- Beta stage, **usable!** (already used in LHCb analyses)
 - Not feature complete, but API stabilizing
- Contributions in form of *feedback and criticism* very welcome
 - API, use-cases, bugs,...
 - Any crazy idea!

It's about a reliable library

not about

«we need to get it working fast»

Different kind of fits



- Binned (*vs histfactory*) vs unbinned
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»

Different kind of fits

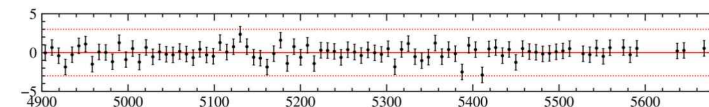
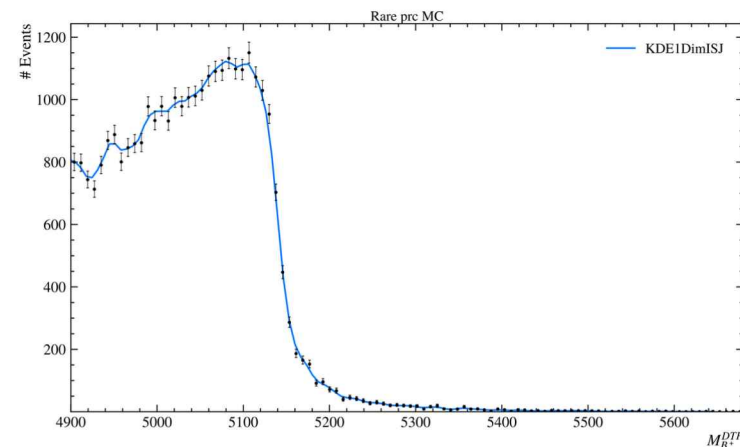
- Binned (*vs histfactory*) vs unbinned
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»
- Template vs analytic
 - Shape from (simulation) sample vs closed-form function

Different kind of fits

- Binned (*vs histfactory*) vs unbinned
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»
- Template vs analytic
 - Shape from (simulation) sample vs closed-form function
- *Analytical vs numerical normalization*
 - *Bin or closed-form integral vs numerical*

Different kind of fits

- Binned (*vs histfactory*) vs **unbinned**
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»
- **Template** vs analytic
 - Shape from (simulation) sample vs closed-form function
- **Analytical** vs **numerical normalization**
 - *Bin or closed-form integral vs numerical*



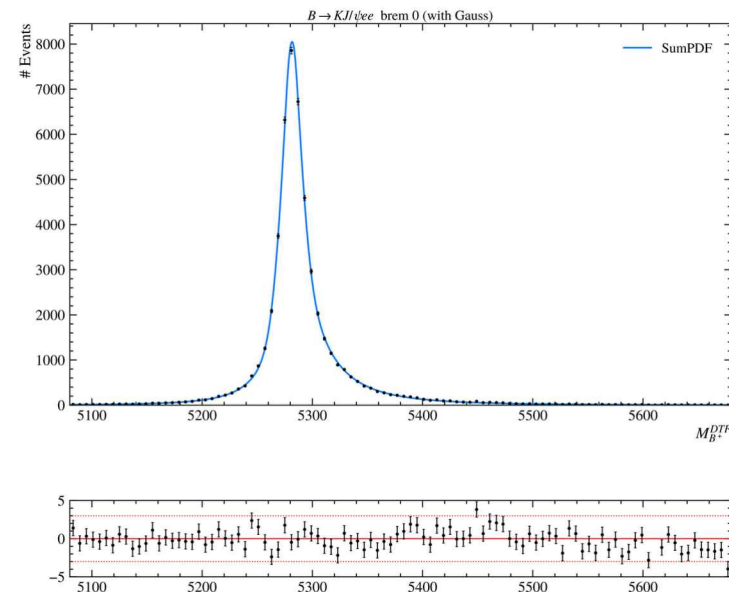
KDE

Gaussian kernel → analytic norm

ISJ → numeric norm

Different kind of fits

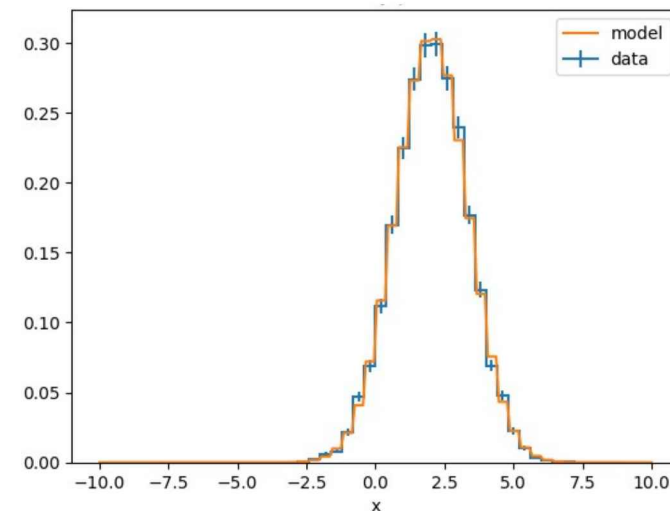
- Binned (*vs histfactory*) vs **unbinned**
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»
- Template vs **analytic**
 - Shape from (simulation) sample vs closed-form function
- **Analytical** vs *numerical normalization*
 - *Bin or closed-form integral vs numerical*



Double Crystalball

Different kind of fits

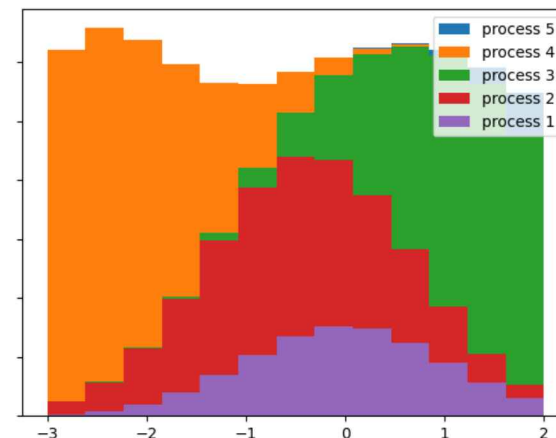
- **Binned** (*vs histfactory*) vs unbinned
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»
- Template vs **analytic**
 - Shape from (simulation) sample vs closed-form function
- **Analytical** vs *numerical normalization*
 - *Bin or closed-form integral vs numerical*



**(binned) Gaussian
fit to histogram**

Different kind of fits

- **Binned** (vs *histfactory*) vs unbinned
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of probabilities
 - Binned data: «counting experiments»
- **Template** vs analytic
 - Shape from (simulation) sample vs closed-form function
- **Analytical** vs *numerical normalization*
 - *Bin or closed-form integral vs numerical*

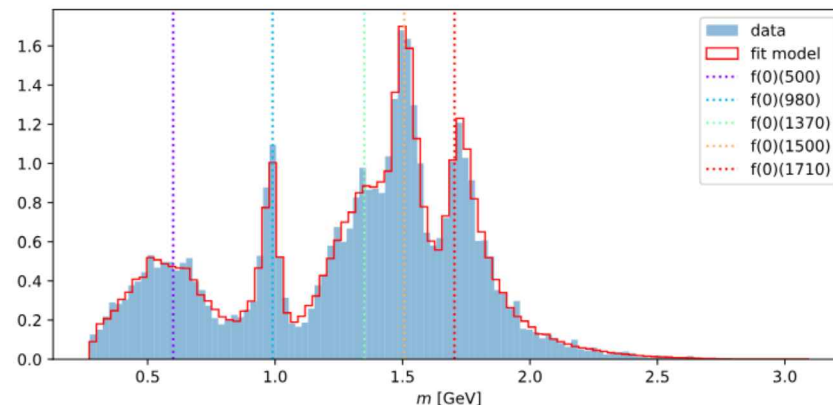


Stacked histograms PDFs

- One extreme: HistFactory model (pyhf)
 - Template, binned, analytic normalization
 - Assumption: Bins «free-standing», not next to each other
- «Closed-world» fitter
 - Limited scope, specialized on 80%+ use-case in CMS/ATLAS
 - extremely powerful/tested, serializable

Different kind of fits

- Binned (vs *histfactory*) vs unbinned
 - Refers to data, cost/loss/likelihood and PDF
 - Unbinned data: product of PDFs
 - Binned data: «counting experiments»
- Template vs analytic
 - Shape from (simulation) sample vs closed-form function
- Analytical vs numerical normalization
 - Bin or closed-form integral vs numerical



Amplitude (partial wave) analysis
Angular analysis

Partial wave analysis



- The other extreme: amplitude analysis (**ComPWA**, ...)
 - Unbinned, analytic, numerical normalisation
 - Description of observable based on amplitude, can be 1k + lines
- Fitting is also hard
 - Fitting time (~100 parameters): hours/days, up to weeks (one fit)
 - Bottleneck: evaluation of PDF

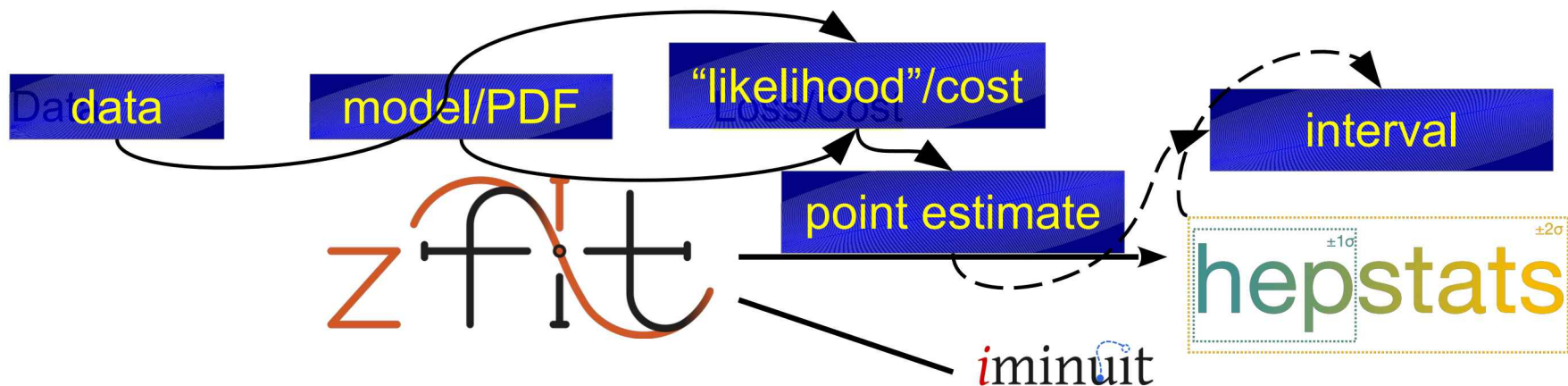
Statistical inference landscape



Closed-world
HistFactory-like



Open-world
Binned,
unbinned,
mixed



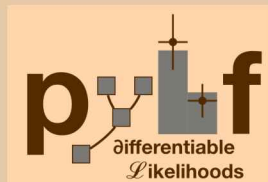
Statistical inference landscape



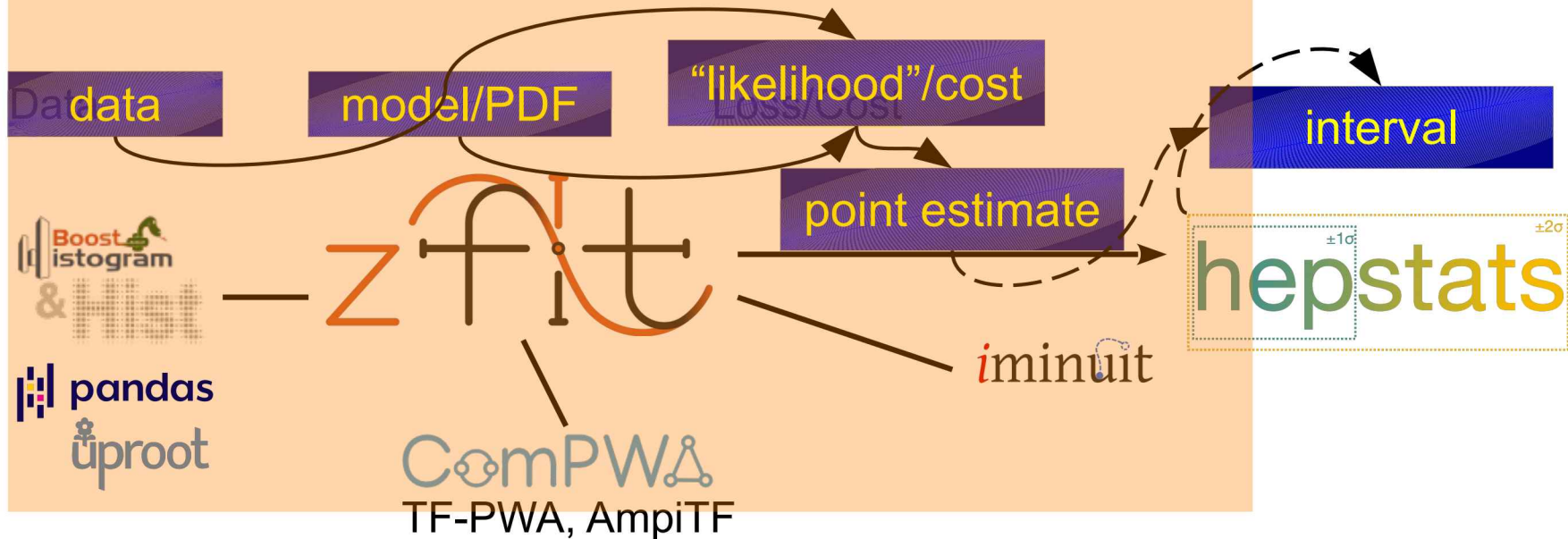
Closed-world
HistFactory-like



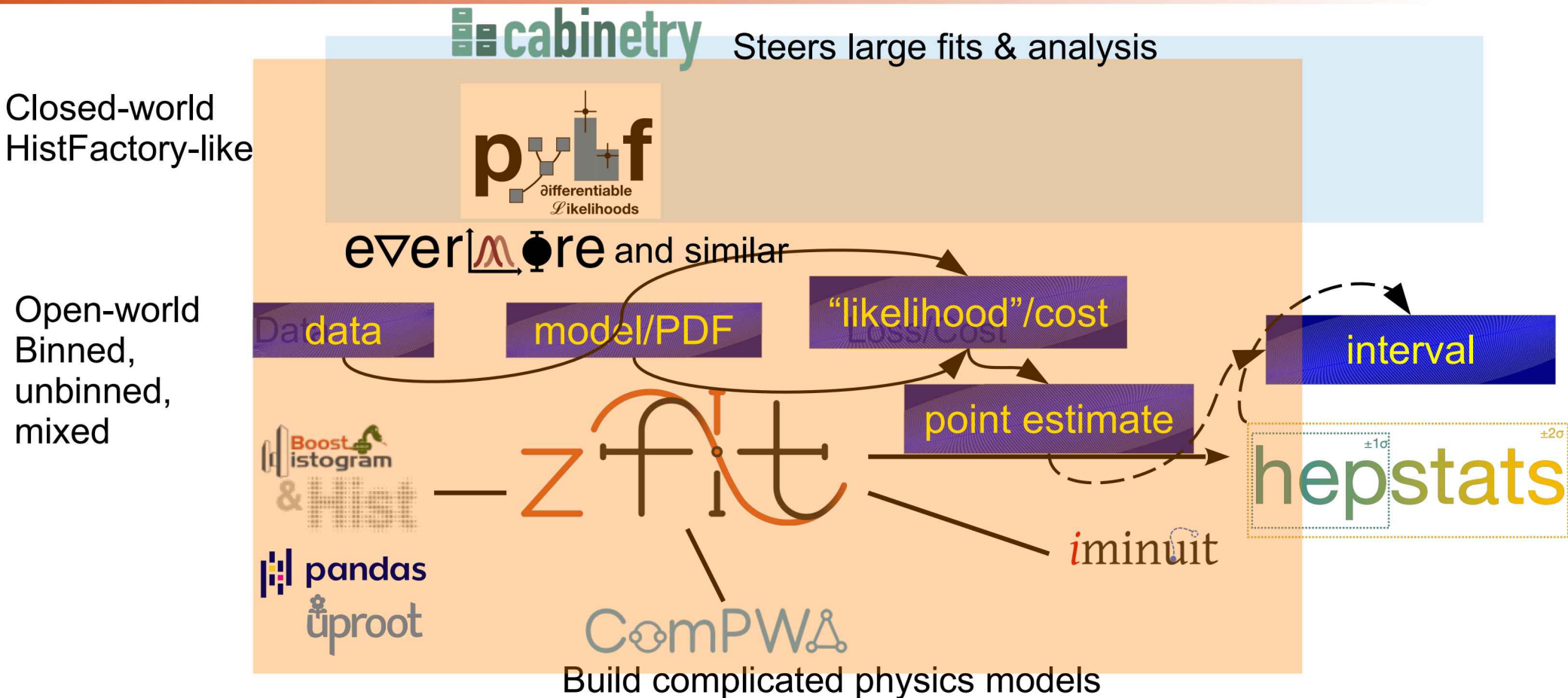
Steers large fits & analysis



Open-world
Binned,
unbinned,
mixed



Statistical inference landscape



Statistical inference landscape

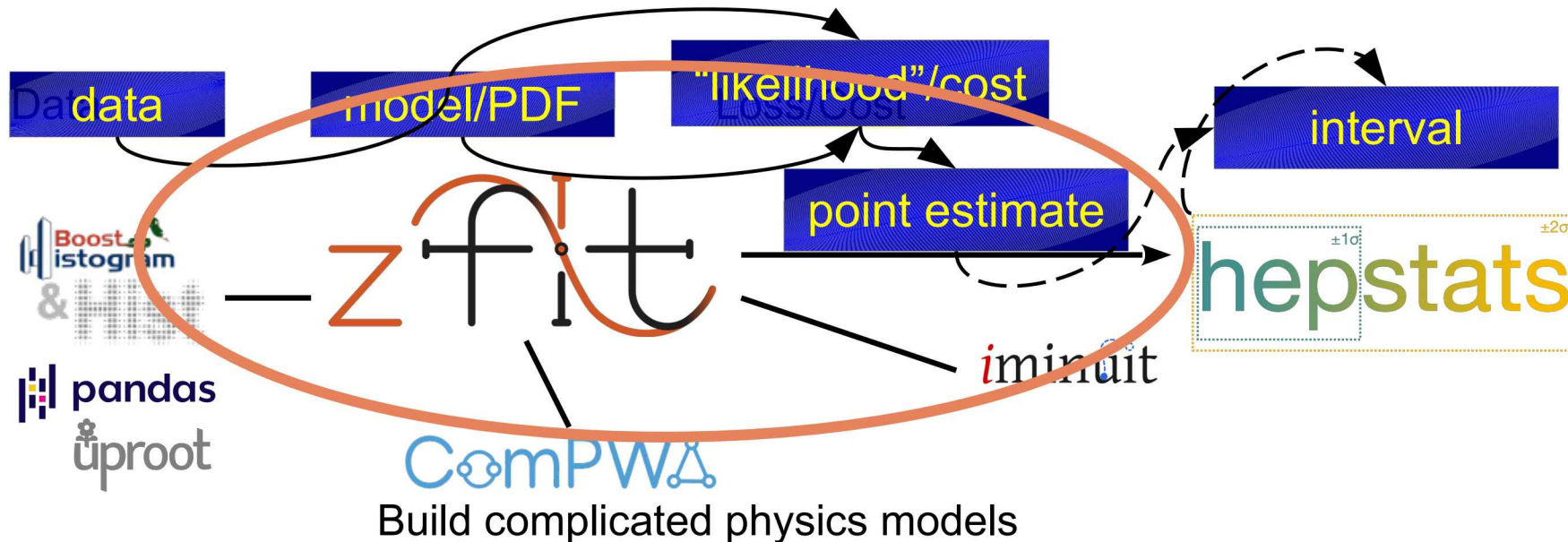


Steers large fits & analysis

Closed-world
HistFactory-like



Open-world
Binned,
unbinned,
mixed



Historic landscape



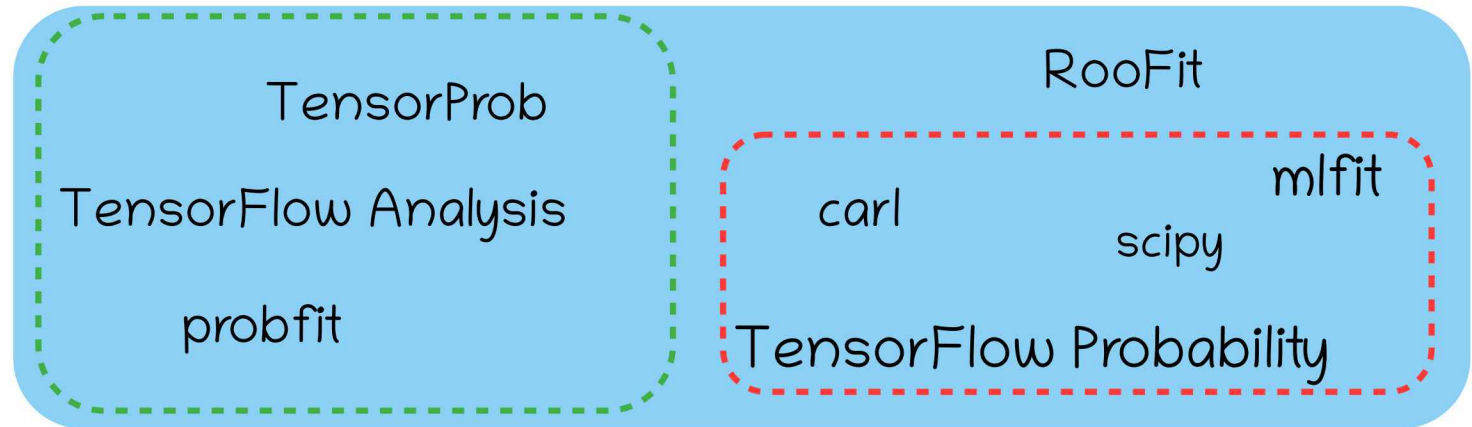
Analyses transition from C++ to Python
Many, non-monolithic packages

Talk by Eduardo



HEP fitting libraries still in C++
Strong libraries, **but mediocre bindings, not «pythonic»**

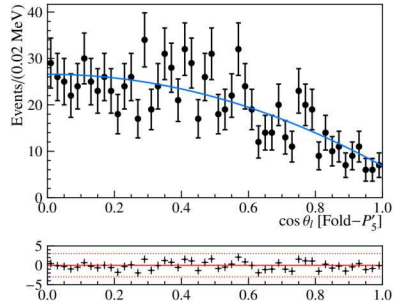
Python packages
- no advanced features
- «Python too slow»



2018



HEP Model Fitting in Python



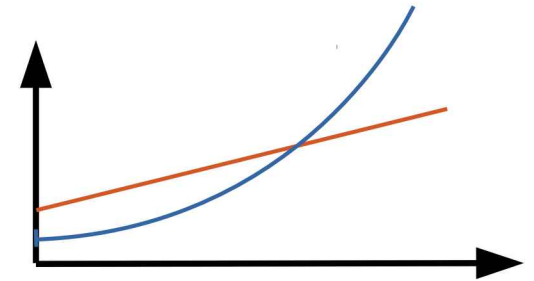
HEP

advanced features,
simply extendable



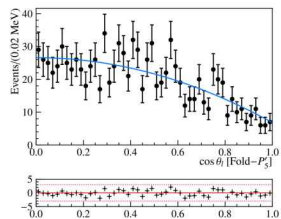
Scalable

large data, complex models



Pythonic  NumPy  python™
integrate into ecosystem, stable API

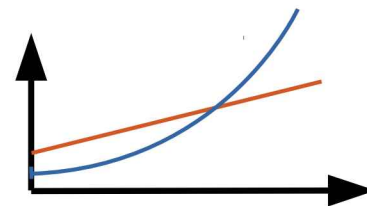
HEP Model Fitting in Python



HEP
advanced features,
simply extendable



Scalable
large data, complex models



Pythonic
integrate into ecosystem, stable API

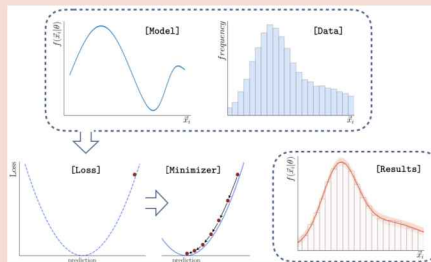


NumPy

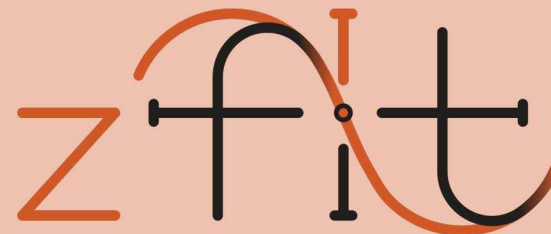


python™

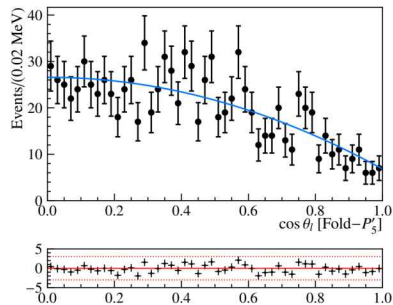
API & Workflow



Computing backend



HEP Model Fitting in Python



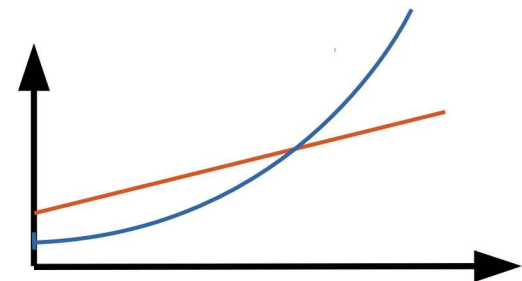
HEP

advanced features,
simply extendable

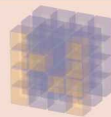


Scalable

large data, complex models



Pythonic



NumPy



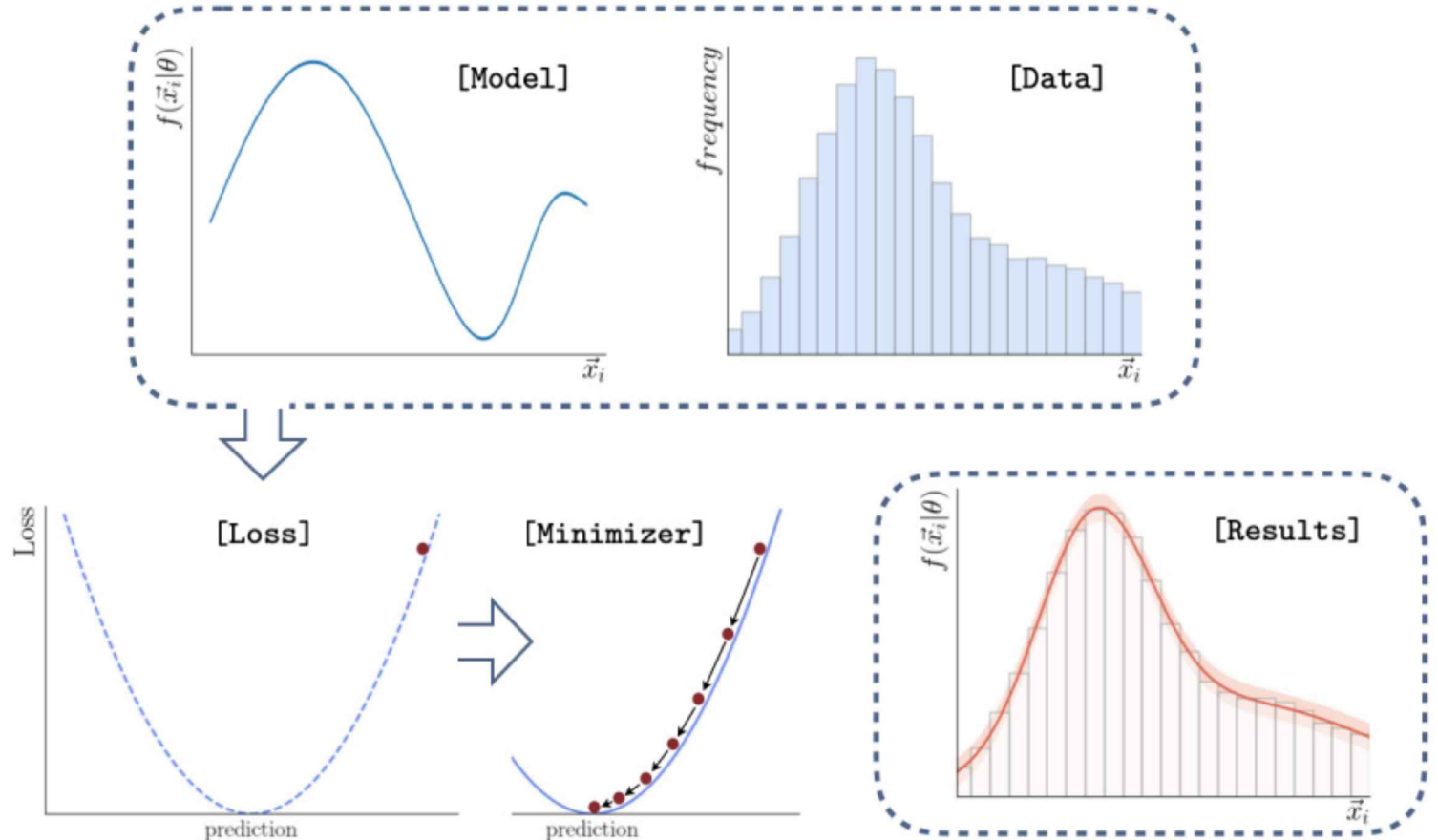
python™

integrate into ecosystem, stable API

API & Workflow

Five maximally independent parts

"Fits look always the same"



Complete fit

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

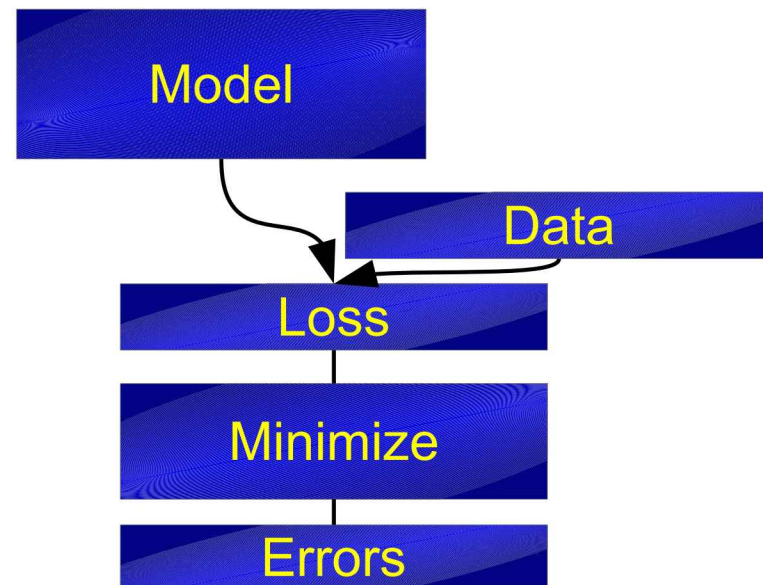
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Model

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

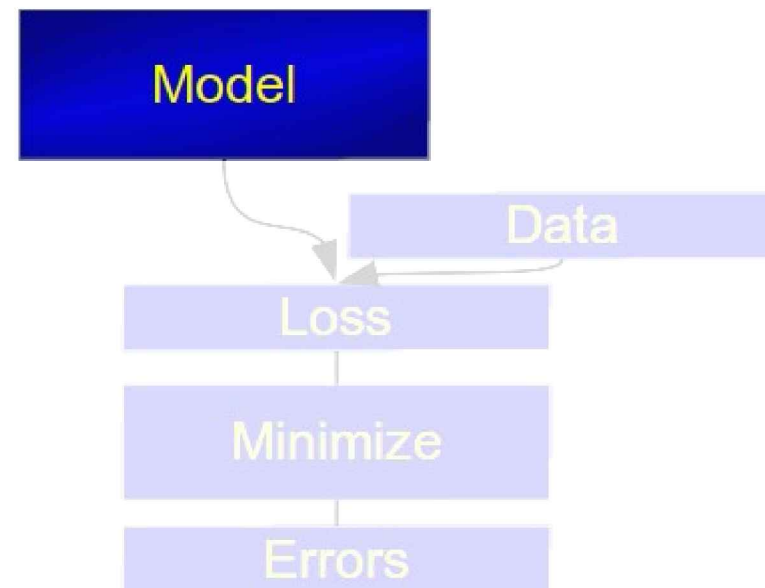
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Data

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

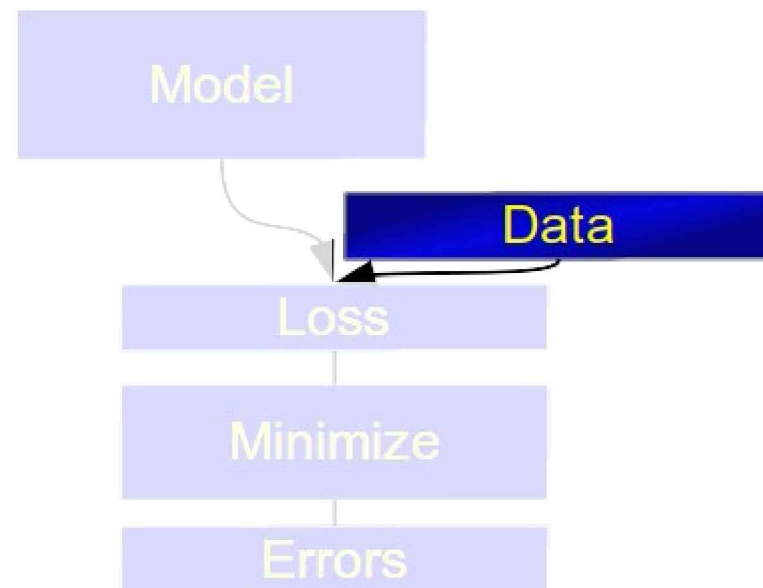
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Loss

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

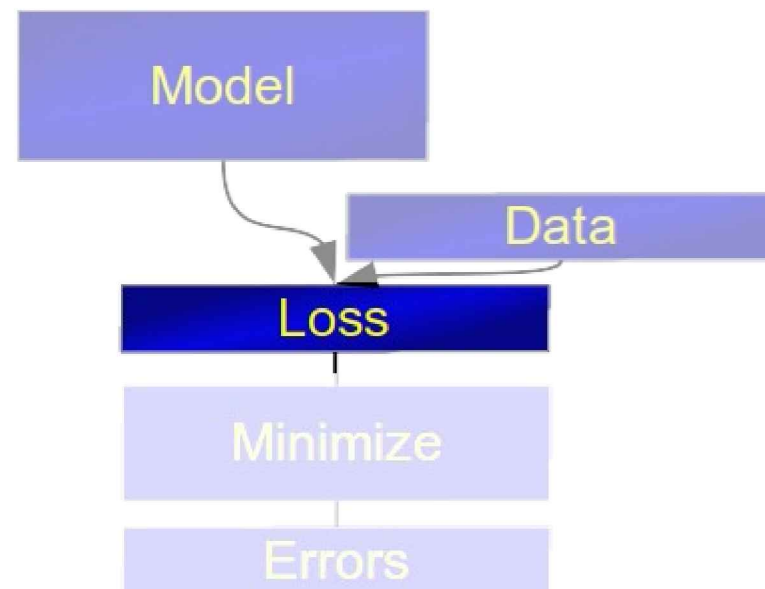
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Minimization

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

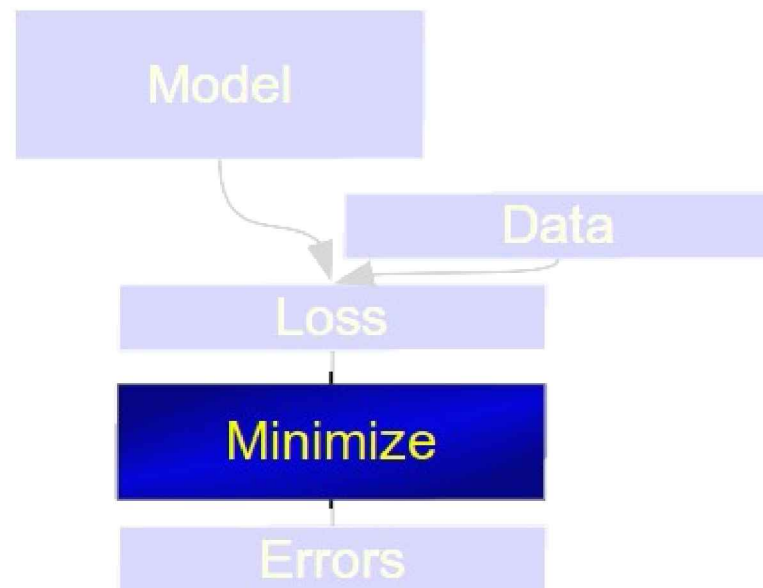
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Result

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

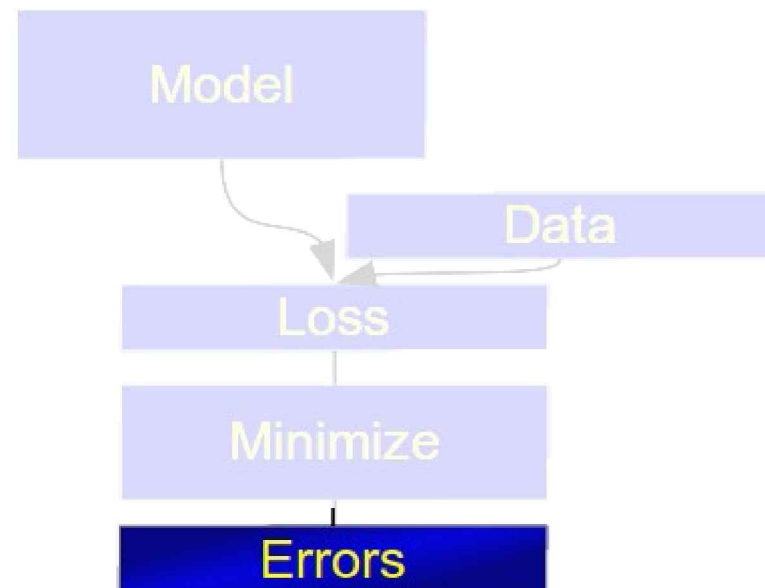
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Basic API example



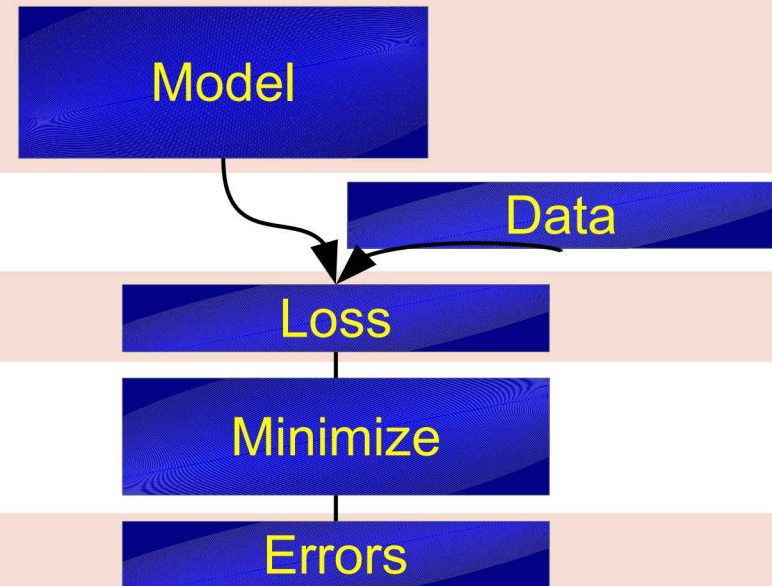
```
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)
```

```
data = zfit.Data.from_numpy(obs=obs, array=normal_np)
```

```
nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)
```

```
minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)
```

```
param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Basic API example

Going binned

```
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)
obs_binned = obs.with_binning(30)
gauss_binned = gauss.to_binned(obs_binned)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)
data_binned = data.to_binned(obs_binned)
nll = zfit.loss.BinnedNLL(model=gauss_binned, data=data_binned)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```

Model

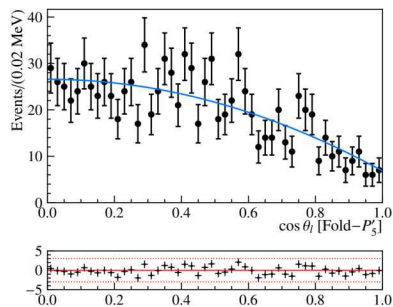
Data

Loss

Minimize

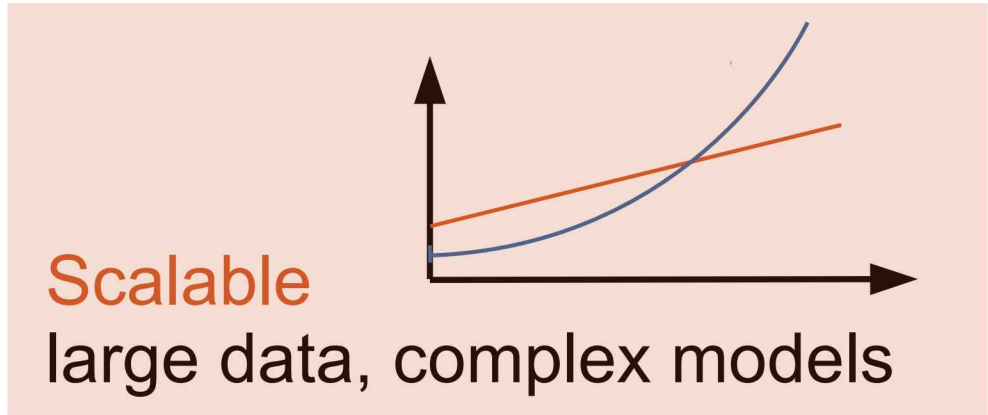
Errors

HEP Model Fitting in Python



HEP

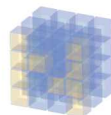
advanced features,
simply extendable



Scalable

large data, complex models

Pythonic



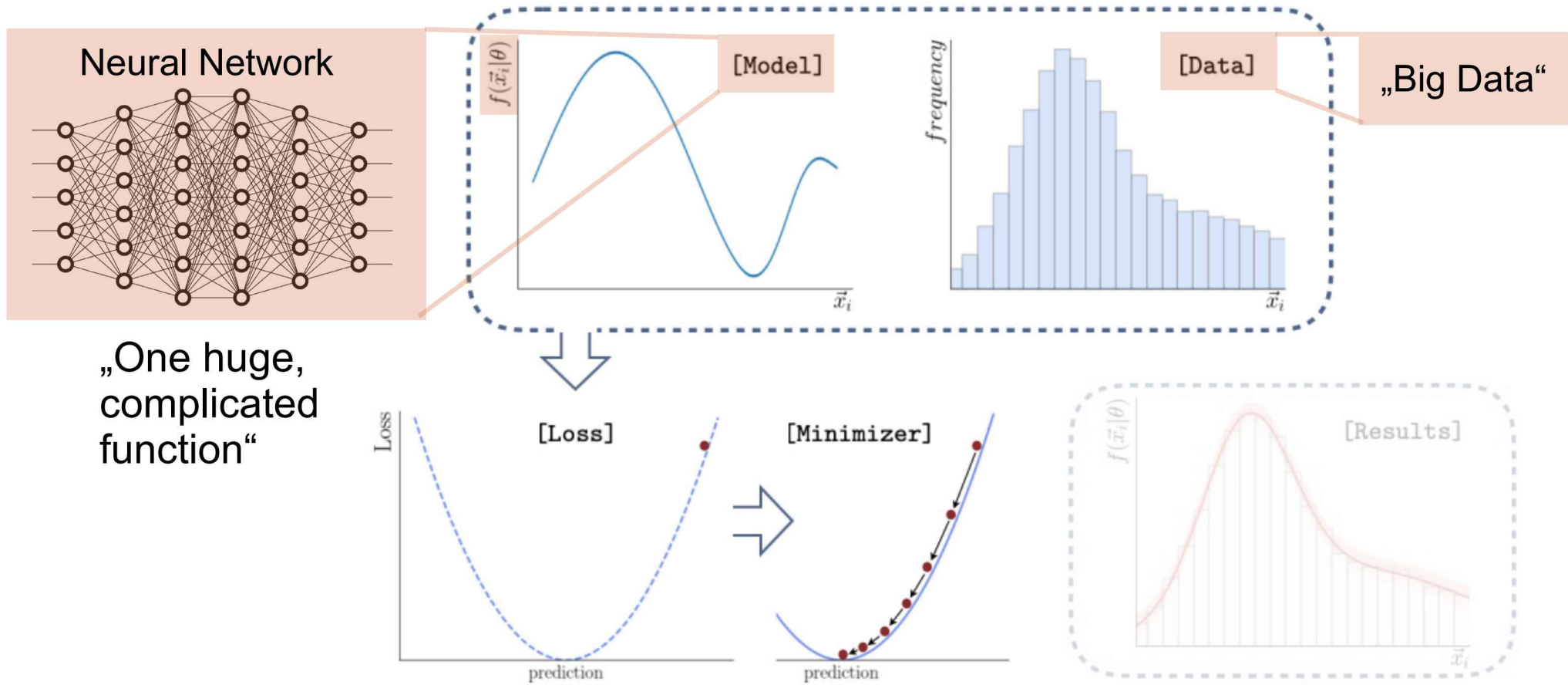
NumPy



python™

integrate into ecosystem, stable API

Deep Learning



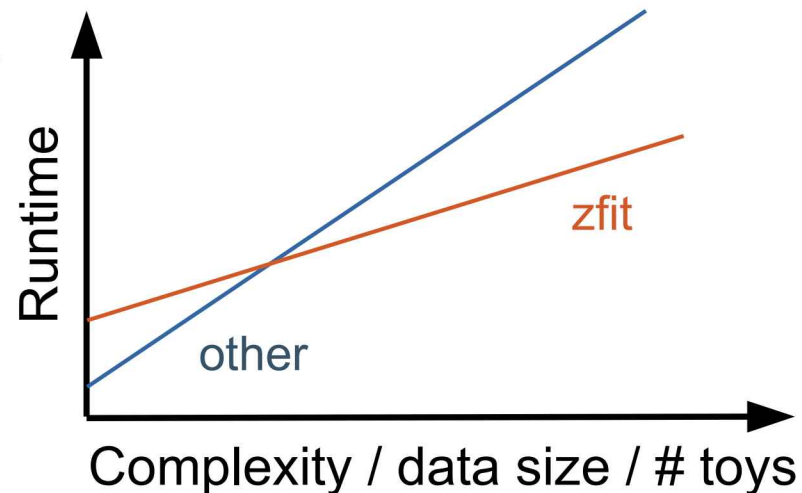
Scalable: Performance

Use same backend as ML uses

- Numpy-like backend TensorFlow (JAX)
 - JIT compiled, CPU or GPU
 - Automatic gradient

Single, simple fit "slow"

- 0.01 or 1 sec not relevant
- 1 or 10 hours relevant

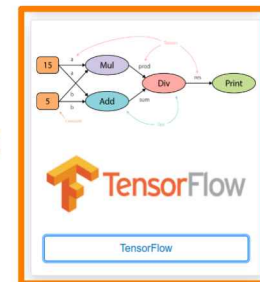


```
import zfit.z.numpy as znp
```

```
ar = znp.linspace(0, 1, 10)
```

```
sin = znp.sin(ar)
```

```
sum_sin = znp.sum(sin)
```

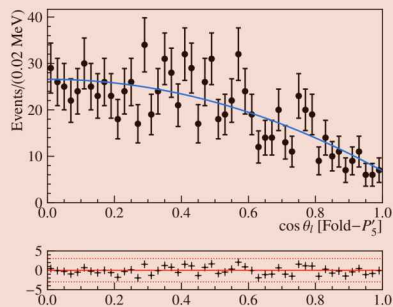


Delegating the workload



	C++ library	Numpy based	zfit
HEP specific content/API			
Models		SciPy	TF Probability
Gradients	CLAD		
Computational optimizations			
Parallelization/GPU		Numba NumPy	intel NVIDIA
Low level handling		python	

HEP Model Fitting in Python



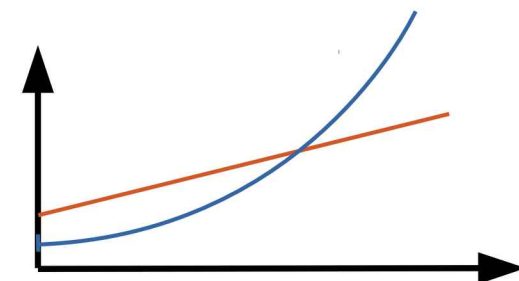
HEP

advanced features,
simply extendable

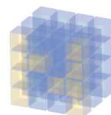


Scalable

large data, complex models



Pythonic



NumPy



python™

integrate into ecosystem, stable API

Complete fit

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

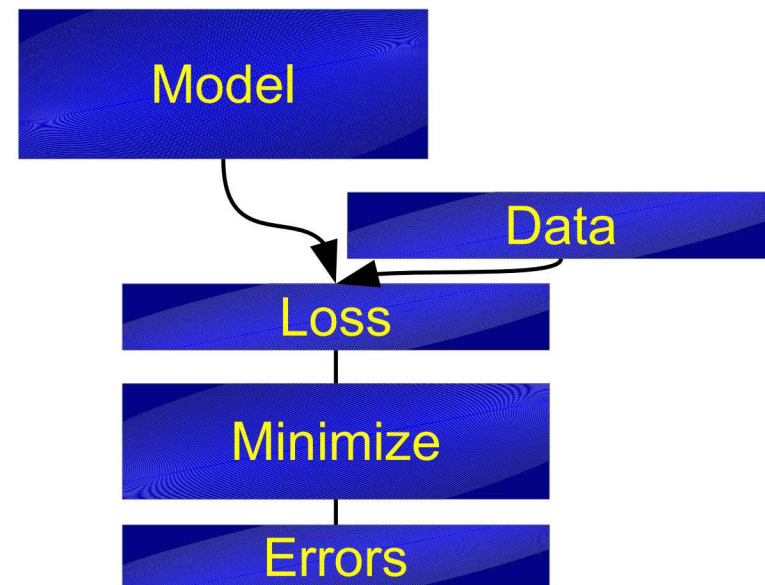
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Model

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

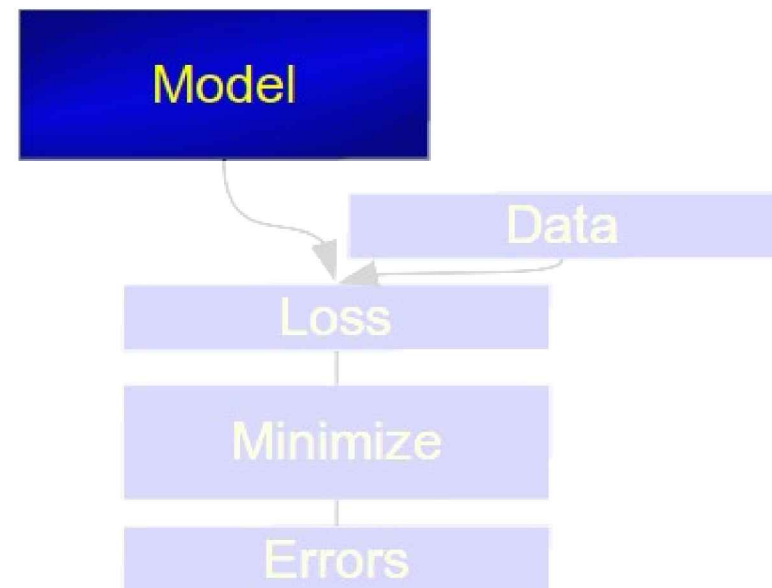
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```

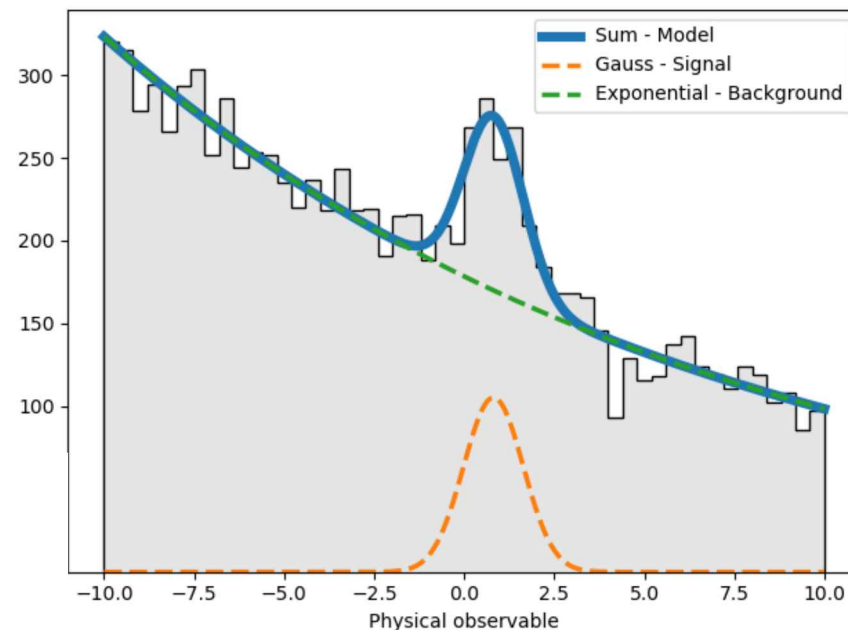


Example: Mass fit

- Sum, Product, (*Convolution*)
- Gauss, (double) Crystalball,...
- Exponential, Polynomials,...
- Histograms, SplineInterpolation,...

```
lambd = zfit.Parameter("lambda", -0.06, -1, -0.01)  
frac = zfit.Parameter("fraction", 0.3, 0, 1)
```

```
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)  
exponential = zfit.pdf.Exponential(lambd, obs=obs)  
model = zfit.pdf.SumPDF([gauss, exponential], fracs=frac)
```



Example: Mass fit

- Sum, Product, (*Convolution*)
- Gauss, (double) Crystalball,...
- Exponential, Polynomials,...
- Histograms, SplineInterpolation,...



```
lambda = zfit.Parameter("lambda", -0.06, -1, -0.01)
frac = zfit.Parameter("frac", 0.1, 0, 1)
```

```
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)
exponential = zfit.pdf.Exponential(lambda=lambda, obs=obs)
model = zfit.pdf.SumPDF([gauss, exponential], fracs=frac)
```

Good for out-of-the-box but...
does not cover even closely all HEP PDFs

Custom PDF



```
from zfit import z
from zfit.z import numpy as znp
```

```
class CustomPDF(zfit.pdf.ZPDF):
```

```
    _PARAMS = ['alpha']
```

```
    def _unnormalized_pdf(self, x):
```

```
        data = z.unstack_x(x)
```

```
        alpha = self.params['alpha']
```

```
        return znp.exp(alpha * data)
```



implement custom function

Custom PDF



```
from zfit import z
from zfit.z import numpy as znp
```

```
class CustomPDF(zfit.pdf.ZPDF):
```

```
    _PARAMS = ['alpha']
```

```
    def _unnormalized_pdf(self, x):
```

```
        data = z.unstack_x(x)
```

```
        alpha = self.params['alpha']
```

```
        return znp.exp(alpha * data)
```

```
custom_pdf = CustomPDF(obs=obs, alpha=0.2)
```

```
integral = custom_pdf.integrate(limits=(-1, 2))
```

```
sample = custom_pdf.sample(n=1000)
```

```
prob = custom_pdf.pdf(sample)
```

} use functionality of model

```
from zfit import z
from zfit.z import numpy as znp
```

```
class CustomPDF(zfit.pdf.ZPDF):
```

```
    _PARAMS = ['alpha']
```

```
    def _unnormalized_pdf(self, x):
```

```
        data = z.unstack_x(x)
```

```
        alpha = self.params['alpha']
```

```
        return znp.exp(alpha * data)
```

```
custom_pdf = CustomPDF(obs=obs, alpha=0.2)
```

```
integral = custom_pdf.integrate(limits=(-1, 2))
```

```
sample = custom_pdf.sample(n=1000)
```

```
prob = custom_pdf.pdf(sample)
```

} use functionality of model

Example of zfit Base Classes

Can also override:

- integrate → `_integrate`
- pdf → `_pdf`
- sample → `_sample`

Or register integral

Arbitrary analytic shapes



```
class P5pPDF(zfit.pdf.ZPDF):
    _PARAMS = ['FL', 'AT2', 'P5p']
    _N_OBS = 3

    def unnormalized_pdf(self, x):
        FL = self.params['FL']
        AT2 = self.params['AT2']
        P5p = self.params['P5p']
        costheta_l, costheta_k, phi = ztf.unstack_x(x)

        sintheta_k = tf.sqrt(1.0 - costheta_k * costheta_k)
        sintheta_l = tf.sqrt(1.0 - costheta_l * costheta_l)

        sintheta_2k = (1.0 - costheta_k * costheta_k)
        sintheta_2l = (1.0 - costheta_l * costheta_l)

        sin2theta_k = (2.0 * sintheta_k * costheta_k)
        cos2theta_l = (2.0 * costheta_l * costheta_l - 1.0)

        pdf = ((3.0 / 4.0) * (1.0 - FL) * sintheta_2k +
              FL * costheta_k * costheta_k +
              (1.0 / 4.0) * (1.0 - FL) * sintheta_2k * cos2theta_l +
              -1.0 * FL * costheta_k * costheta_k * cos2theta_l +
              (1.0 / 2.0) * (1.0 - FL) * AT2 * sintheta_2k *
              sintheta_2l * tf.cos(2.0 * phi) + tf.sqrt(FL * (1 - FL))
              * P5p * sin2theta_k * sintheta_l * tf.cos(phi))

    return pdf
```

For example, create amplitude with **ComPWA** and **zfit**

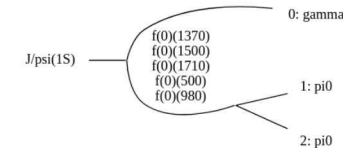
Amplitude analysis with zfit

► Show code cell content

Formulating the model

```
import grules
reaction = grules.generate_transitions(
    initial_state="J/psi(1S)", [-1, +1],
    final_state=["gamma", "pi0", "pi0"],
    allowed_intermediate_particles=["f(0)"],
    allowed_interaction_types=["strong", "EM"],
    formalism="helicity",
)
```

► Show code cell source



```
import ampform
from ampform.dynamics.builder import (
    create_non_dynamic_with_ff,
    create_relativistic_breit_wigner_with_ff,
)
model_builder = ampform.get_builder(reaction)
```

Binned models

- Success story of Universal Histogram Interface (UHI)
- Modelled after/ compatible with **boost-histogram/hist/UHI**
 - Axes, names,



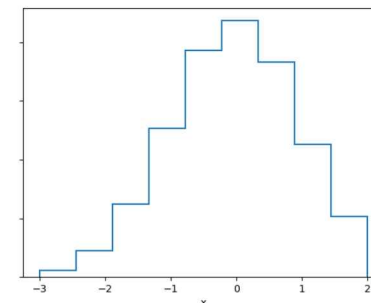
```
h = hist.Hist(hist.axis.Regular(3, -3, 3, name="x", flow=False),
              hist.axis.Regular(2, -5, 5, name="y", flow=False))
```

```
x = np.random.randn(1_000_000)
y = 0.5 * np.random.randn(1_000_000)
h.fill(x=x, y=y)
```

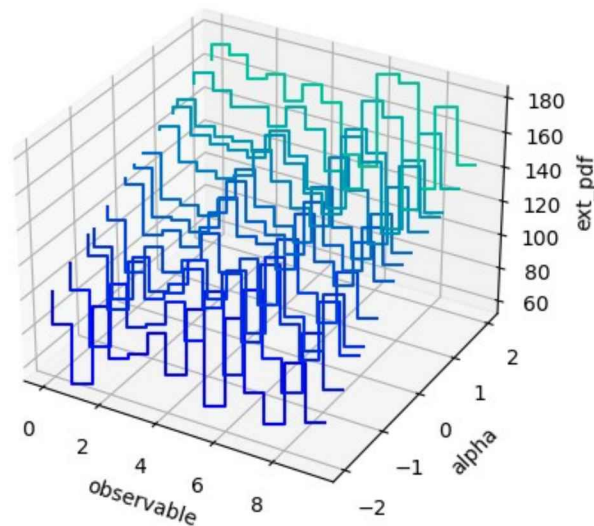
```
pdf = zfit.pdf.HistogramPDF(data=h)
```

```
mplhep.histplot(h_back)
```

```
...and back
h_back = pdf.to_hist()
```

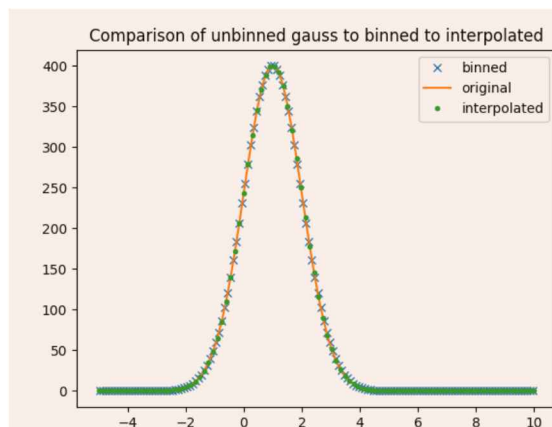


More histograms

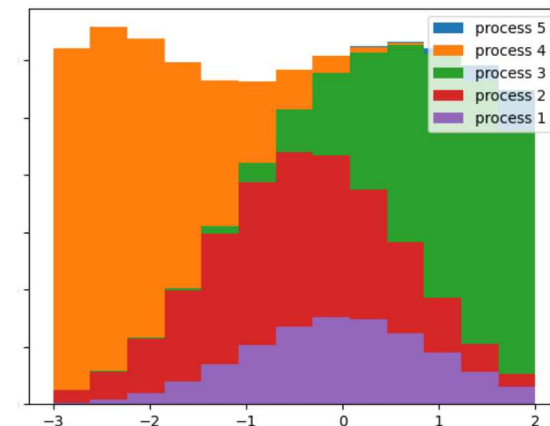


Shape modifier

```
pdf_syst = zfit.pdf.BinwiseScaleModifier(pdf, modifiers=True)
```



Unbinned → binned → interpolated



```
pdfs = [zfit.pdf.HistogramPDF(h) for h in histos]
alpha = zfit.Parameter('alpha', 0, -5, 5)
morph = SplineMorphingPDF(alpha=alpha, hists=pdfs)
```

```
pdfs = [zfit.pdf.HistogramPDF(h) for h in histos]
sumpdf = zfit.pdf.BinnedSumPDF(pdfs)
```

Complete fit: Data

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

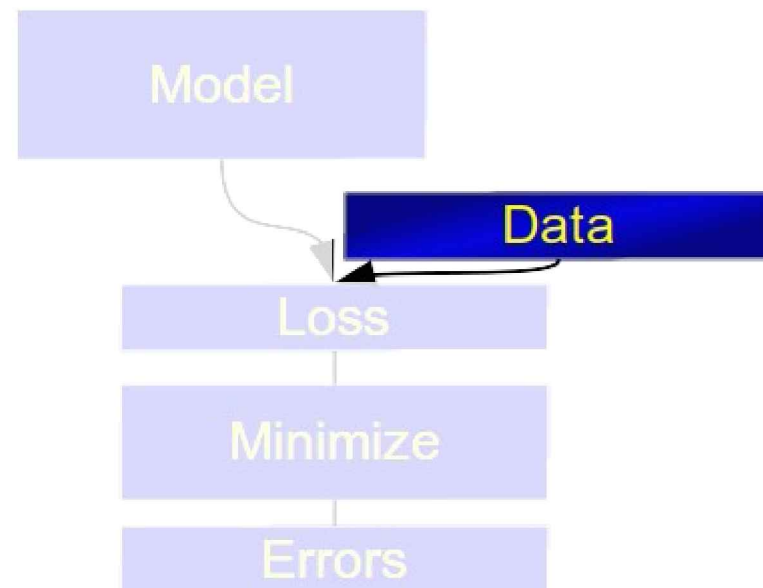
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Complete fit: Data



- From different sources
 - Hist, numpy, Pandas, ROOT, ...
 - Can directly be given

Use the HEP/Python ecosystem for preprocessing

- Sampled from a model (toy studies)

```
data = model.create_sampler(n_sample, limits=obs)
```

- UHI compatible!

```
binneddata = binnedpdf.sample()  
mplhep.histplot(binneddata)
```

Complete fit: Loss

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

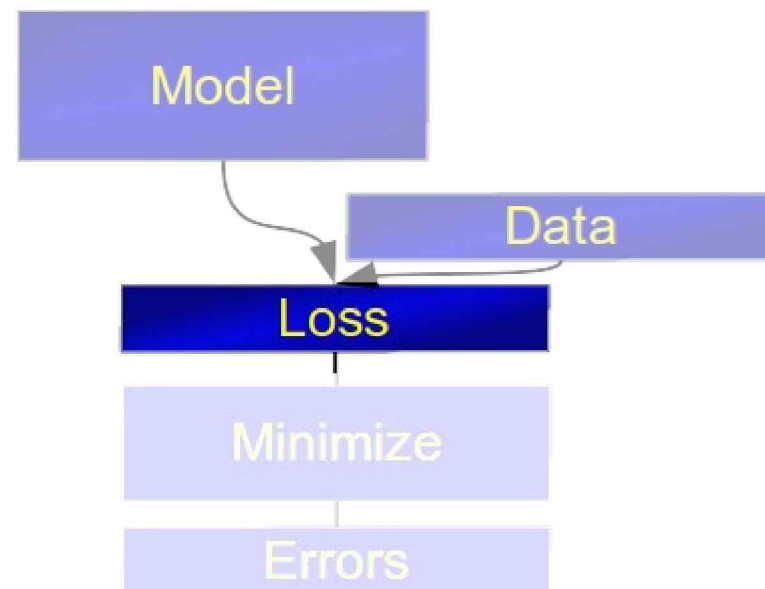
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

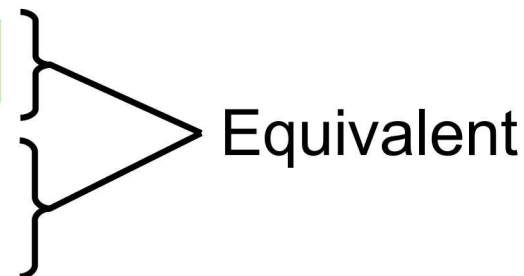
minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



```
nll_simultaneous = zfit.loss.UnbinnedNLL(model=[gauss1, gauss2],  
                                          data=[data1, data2])
```

```
nll1 = zfit.loss.UnbinnedNLL(model=gauss1, data=data1)  
nll2 = zfit.loss.UnbinnedNLL(model=gauss2, data=data2)  
nll_simultaneous2 = nll1 + nll2
```



(arbitrary) constraints supported, added to loss

```
constr = GaussianConstraint(params=params, observation=observed, uncertainty=sigma)  
nll = zfit.loss.BinnedNLL(model=model, data=data, constraint=constr)
```

Directly compatible with iminuit

Complete fit: Minimization

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

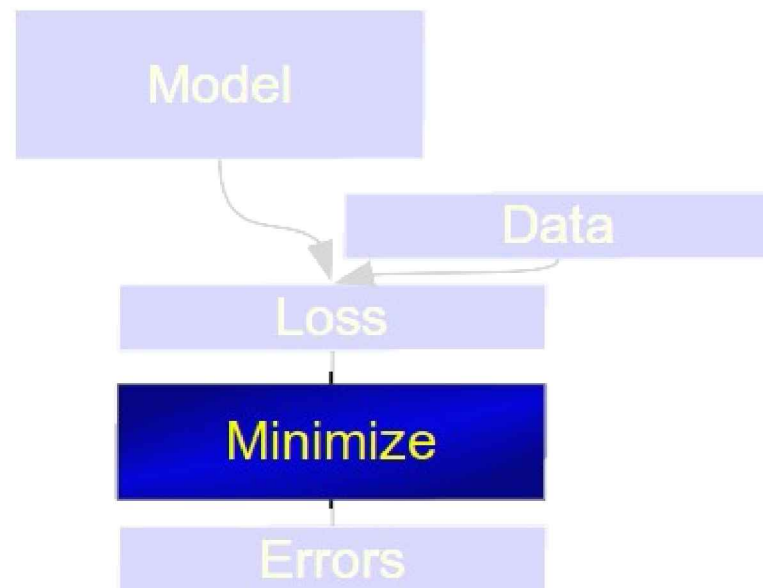
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



Minimize

- Problem: many, non-unified minimizer APIs
 - SciPy interface "a bit messy", different convergence criterion, etc...
- Unified API: zfit minimizers, simply switch

```
minimizer = zfit.minimize.IpyoptV1()  
minimizer = zfit.minimize.Minuit()  
minimizer = zfit.minimize.ScipyTrustConstrV1()  
minimizer = zfit.minimize.NLoptLBFGSV1()
```

- Can use zfit loss, but also ***pure Python function***

```
result = minimizer.minimize(func, params)
```

Complete fit: Result

```
normal_np = np.random.normal(2., 3., size=10_000)

obs = zfit.Space("x", limits=(-2, 3))

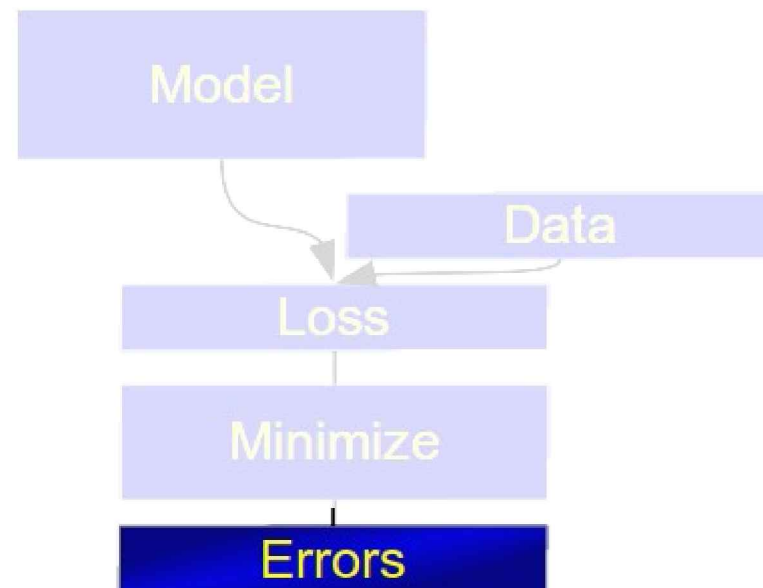
mu = zfit.Parameter("mu", 1.2, -4, 6)
sigma = zfit.Parameter("sigma", 1.3, 0.5, 10)
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)

data = zfit.Data.from_numpy(obs=obs, array=normal_np)

nll = zfit.loss.UnbinnedNLL(model=gauss, data=data)

minimizer = zfit.minimize.Minuit()
result = minimizer.minimize(nll)

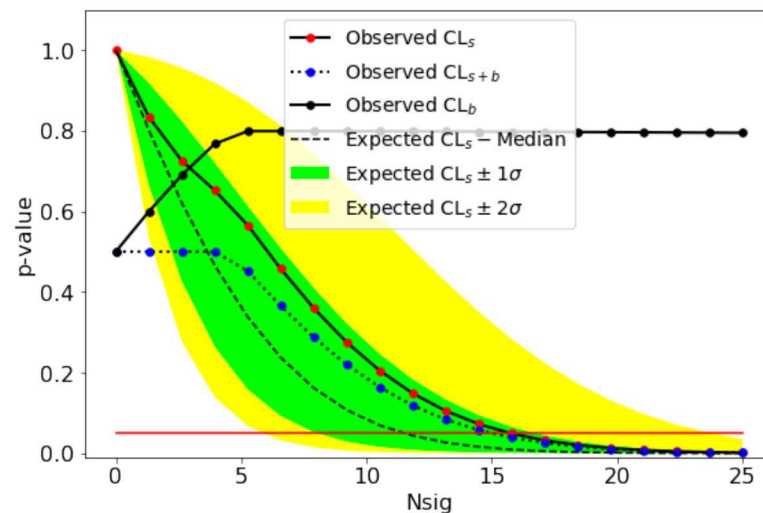
param_errors = result.hesse()
param_errors_asymmetric, new_result = result.errors()
```



- Inference library for hypothesis tests
- Takes model, data, loss from zfit
- sWeights, CI, limits, ...
- asymptotic or toys calculator



```
calculator = AsymptoticCalculator(loss, minimizer)
poinull = POIarray(Nsig, np.linspace(0.0, 25, 20))
poialt = POI(Nsig, 0)
ul = UpperLimit(calculator, poinull, poialt)
ul.upperlimit(alpha=0.05, CLs=True)
```



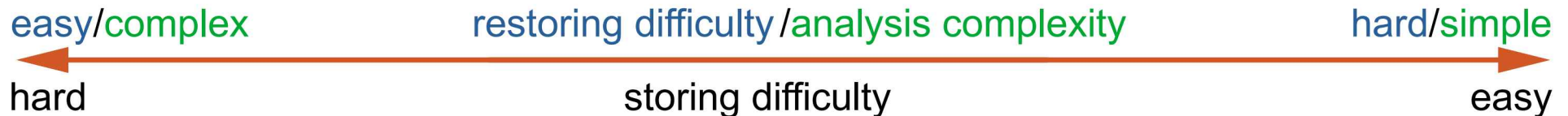
HS³

HEP Statistics Serialization Standard

Preservation and interoperability



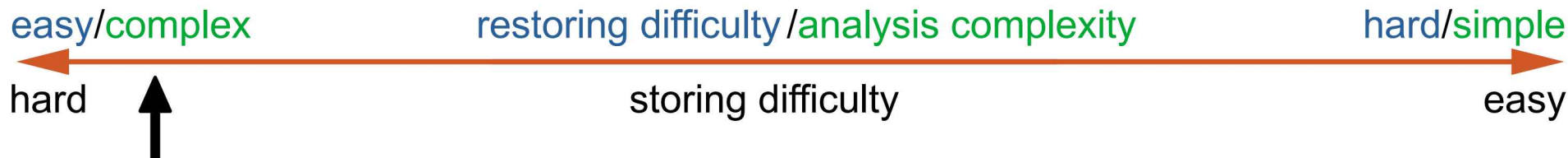
- Goal: restore and exchange likelihood information
 - Use different frameworks (connect across languages)
 - Easily modify likelihoods (theorists!)
- Question: how? Which format?
 - Human-readable vs binary, scripts vs description, virtual machines vs software dependencies, paper vs electronic,...



HEP Statistics Serialization Standard

Human-readable & preservable format for HEP statistics

- By RooFit, zfit, pyhf and bat.jl; developing stage
- Explore and define common ground
 - What is a Gaussian/Gauss/Normal? Sum? Variable?
 - Defining channels and nuisance parameter (HistFactory-like)
- Best effort base: «What works for all, works»

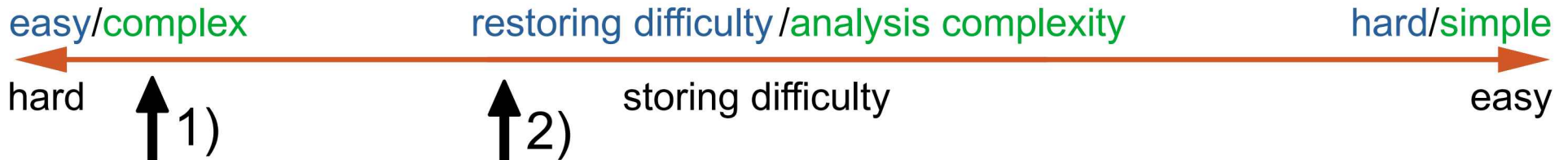


zfit serialization - HS3



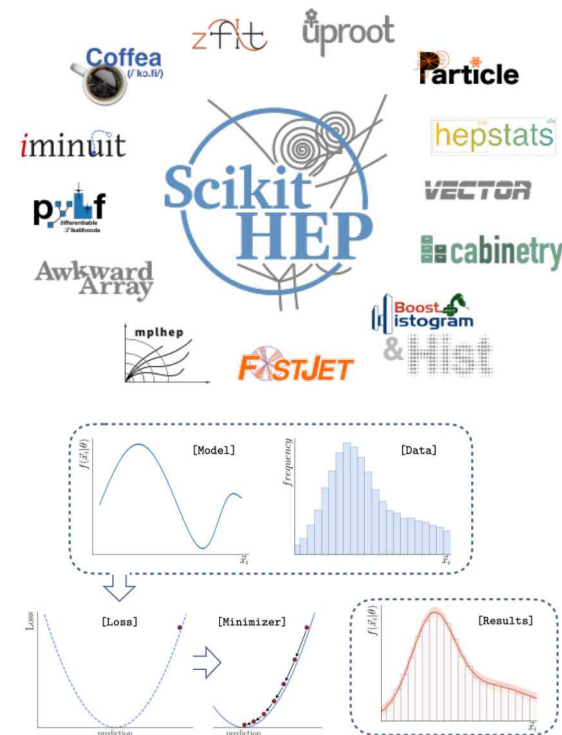
- 1) Can dump/load (some) PDFs HS3-like
- 2) Custom functions currently pickled, in future as SymPy/Mathematica
 - **Project** already achieved that
 - ComPWA for amplitudes in Sympy

```
'pdfs': {'SumPDF': {'pdfs': [{'extended': 'n_sig',  
    'mu': 'mu',  
    'sigma': 'sigma',  
    'type': 'Gauss',  
    'x': 'x'},  
    {'extended': 'n_bkg',  
    'lam': 'lambda',  
    'type': 'Exponential',  
    'x': 'x'}]},  
    'type': 'SumPDF'}},  
'variables': {'lambda': {'max': -0.009999999776482582,  
    'min': -1.0,  
    'name': 'lambda',  
    'step_size': 0.001,  
    'value': -0.06294756382703781},
```



Conclusion

- Python HEP fitting ecosystem built from multiple libraries
- zfit «open-world» fitter, well integrated with the ecosystem
- Interoperability & building on existing Python scientific ecosystem is key
- HS³ likelihood serialization standard



Conclusion

build stable model fitting ecosystem for HEP

- Recent addition of binned/mixed fits
- Human-readable serialization

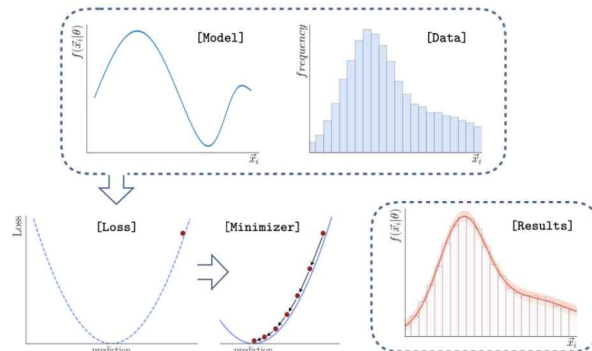
HS3 JSON serialization (WIP)

Pickling of results

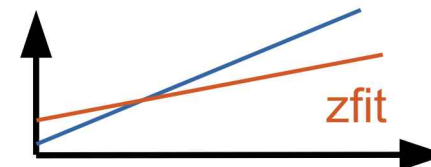
Custom dumping simple

Serialization of toys

- Planning for zfit V2



```
'pdfs': {'SumPDF': {'pdfs': [{'extended': 'n_sig',
                             'mu': 'mu',
                             'sigma': 'sigma',
                             'type': 'Gauss',
                             'x': 'x'},
                              {'extended': 'n_bkg',
                             'lam': 'lambda',
                             'type': 'Exponential',
                             'x': 'x'}],
         'type': 'SumPDF'}}
```



Backup Slides

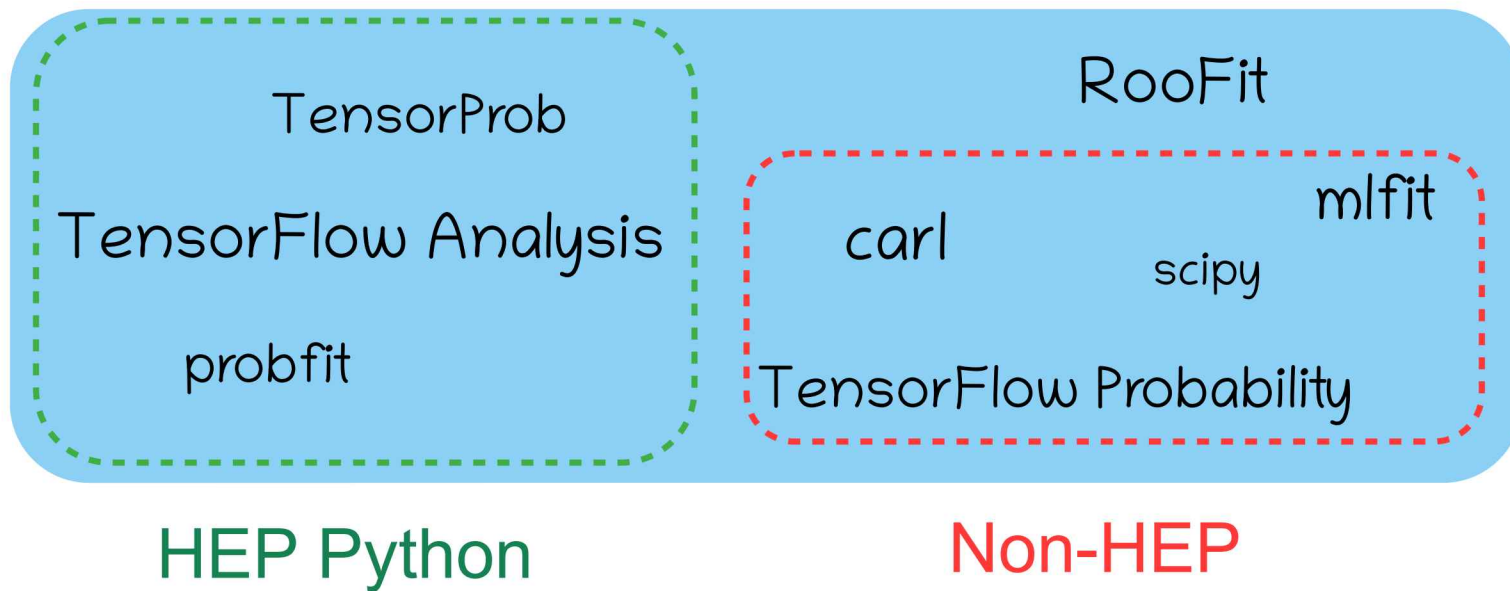
- Backend & TF
- Amplitude
- K*ll toys
- K*mumu Wilson coeffs
- Other fitting packages
- Zfit (associated) packages
- Zfit project
- Zfit elements examples

- Extended fits, Chi2, binned, unbinned, mixed
- PDFs convertible binned ↔ unbinned (including to hist), mixed
- Multidimensional
- Any backend supported (numpy-like), optimal with TF currently
- Sample from PDF
- Arbitrary constraints (custom made)
- Custom PDF: define shape → auto normalized, sampling etc.
- Automatic/numerical gradient
- Different minimizers, optimized API
- JIT/eager support

Fitting in Python



A lot of projects are around



Backend & TensorFlow

Backend: a comparison

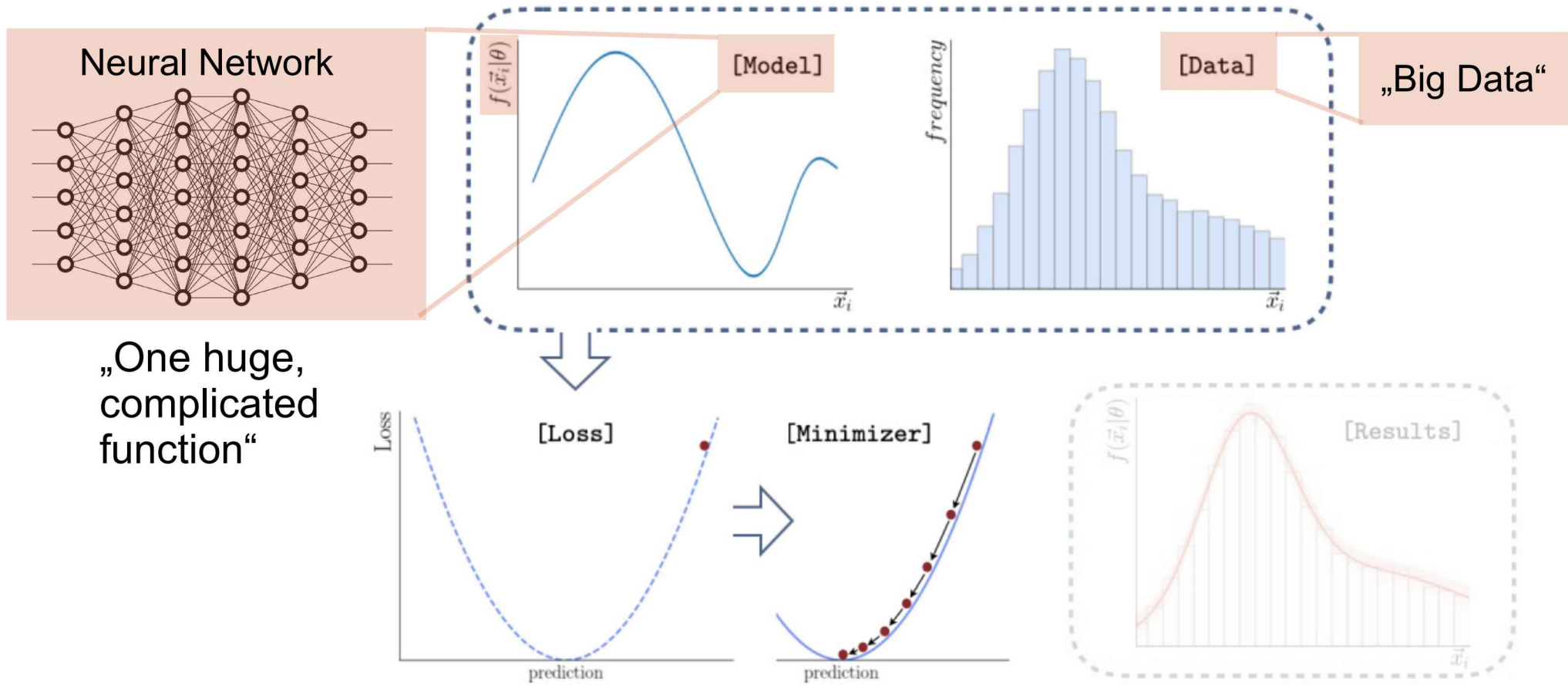


- TensorFlow: supports the most features to this day
- PyTorch: missing advanced math (complex support, ...)
- Numpy/SciPy: Too slow, no gradient, no GPU
- JAX: very promising, but *no globals (cache,...)*, only static known shapes (adaptive algorithms, accept-reject...), only JAX/Numpy arrays compatible
- SymPy: limited to mathematical expressions (no control-flow,...) but can convert to any other backend (used by TensorWaves)

Deep Learning

lessons for model fitting

Deep Learning



Can we express model fitting as
static graphs?

Yes!

- 1) Definition of computation, shape etc. (add static knowledge)
- 2) Compilation of the graph
- 3) Execution of computation (re-use optimized graph)

Inside TF, hidden to end-user

HPC: the more is know *before* the execution, the better

TensorFlow takes care of *how* to use this knowledge

... do not have to be constant!

Parameters

Can change their value

Random numbers

Generate newly on every graph execution: MC integration,...

Control flow (if, while)

Steer the execution: Accept-reject sampling (while), etc.

Static, not constant

Deep Learning vs. Model Fitting



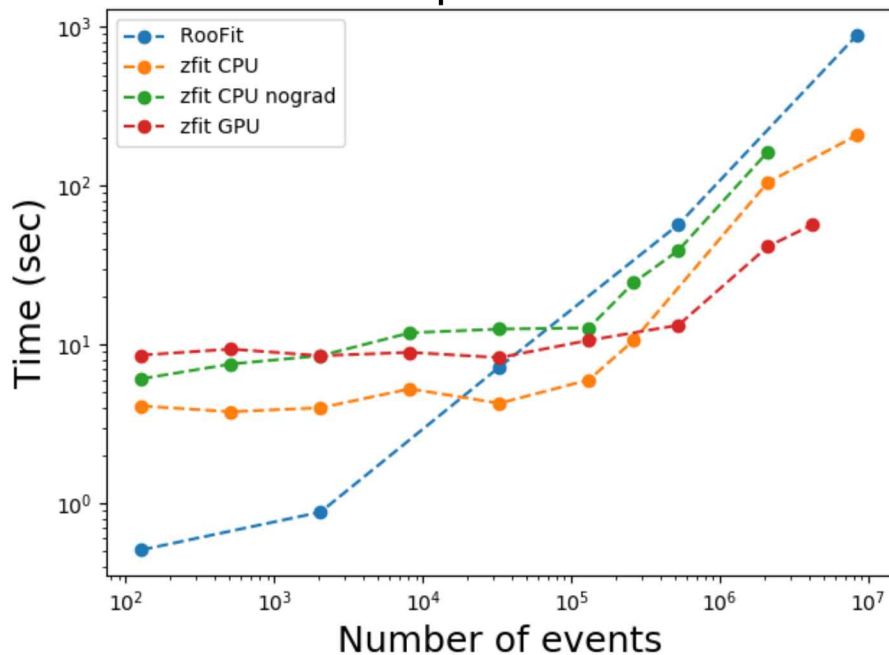
Similarity	Complicated Models	Large Data	Composed loss	Minimization	Results and uncertainties
HEP	Non-trivial functions	Whole Dataset	simultaneous, constraints	Global min, 2 nd derivative algorithm	Hesse, profiling
Deep Learning	Combine many, trivial functions	Many, small Batches	<i>Anything!</i> (GANs, RL,...)	Local (!) min, 1 th derivative, many steps	None
Conclusion					

Scalability: Performance

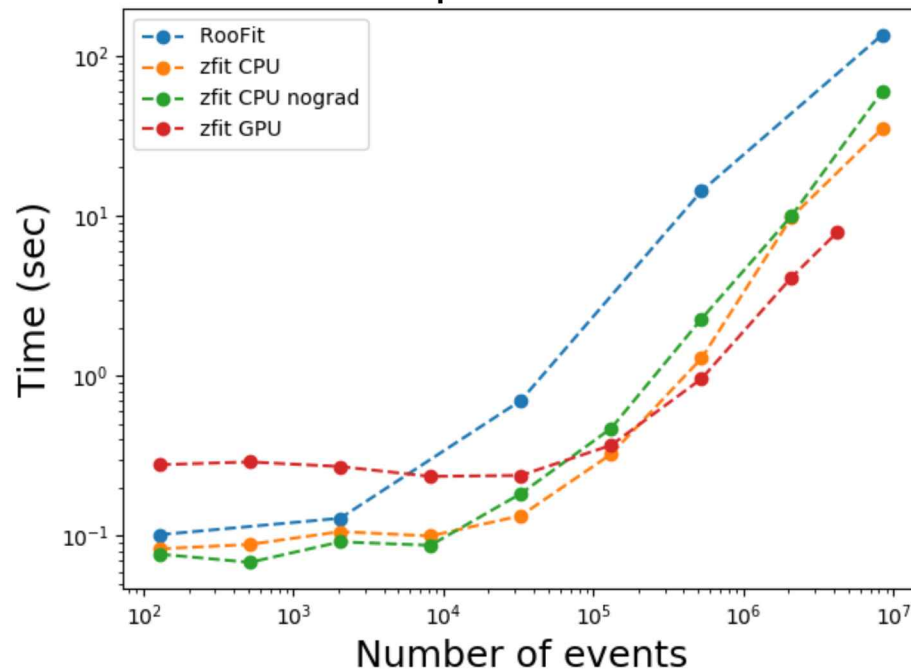


Fitting time (lower is better): **RooFit** vs. **zfit**

9 free parameters



2 free parameters



Amplitude

Angular toys

$B^0 \rightarrow K^{*0} l^+ l^-$ angular: toy study



Sensitivity study

- draw toys (sample) from PDF
- Fit to sample

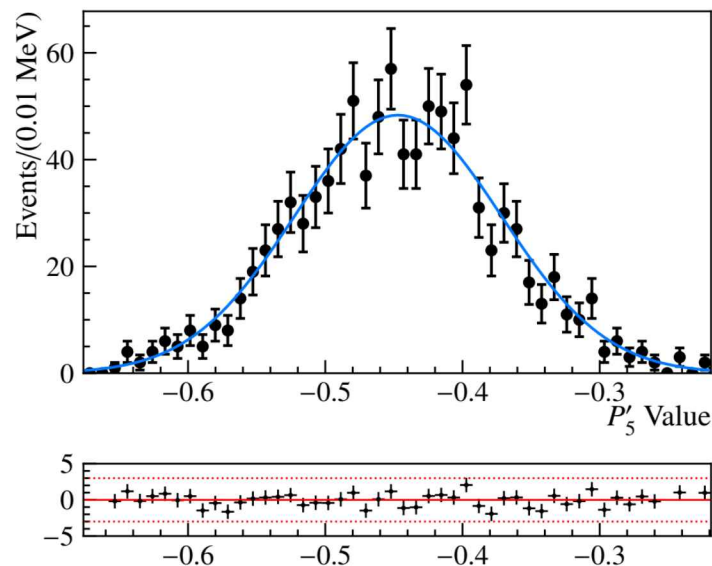
```
for i in range(ntoys):  
  
    # set initial sampling values  
    for param in params:  
        param.set_value(...)  
  
    sampler.resample()  
  
    # set random initial values  
    for param in params:  
        param.set_value(...)  
  
    result = minimizer.minimize(nll)  
  
    if result.converged:  
        ...
```

$B^0 \rightarrow K^{*0} l^+ l^-$ angular: toy study

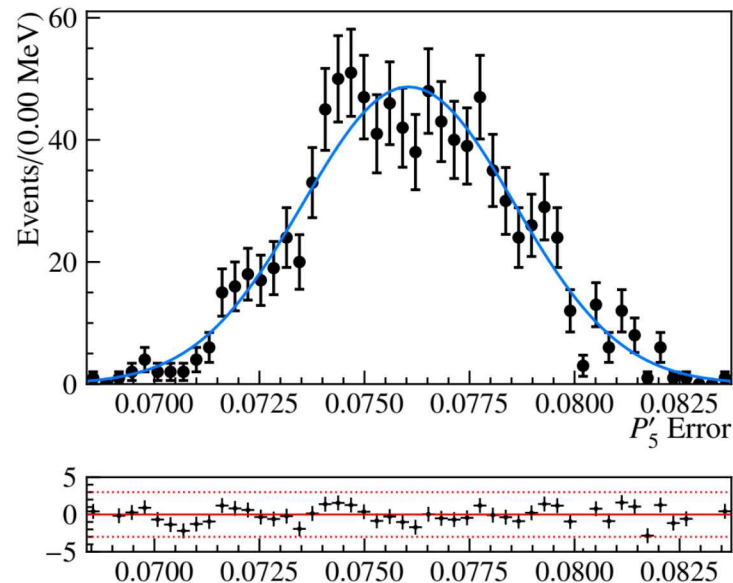


Result of toy study

P5' value



P5' error



Extending with a mass shape

```
# Create mass pdf
mu = zfit.Parameter("mu", 5279, 5200, 5400)
sigma = zfit.Parameter("sigma", 30, 0, 300)
a0 = zfit.Parameter("a0", 1.0, 0, 10)
a1 = zfit.Parameter("a1", 1.0, 0, 10)
n0 = zfit.Parameter("n0", 5, 0, 10)
n1 = zfit.Parameter("n1", 5, 0, 10)

mass = zfit.Space("mass", limits=(4900, 5600))

massPDF = zfit.pdf.DoubleCB(obs=mass, mu=mu, sigma=sigma,
|                             alphas=a0, nls=n0, alphas=a1, nrs=n1)

pdf = massPDF * angularPDF
```

Build model

Fitting libraries and comparison

Python model fitting in HEP



- **Scalable:** large data, complex models
- **Pythonic:** use Python ecosystem/language
- Specific HEP functionality:
 - Normalization: specific range, numerical integration,...
 - Composition of models
 - Multiple dimensions
 - Custom models
 - Non-trivial loss (constraints, simultaneous,...)

Probit, TensorProb,...

- Lack **generality** and extendibility
- “experimental”, but great proof of concept
 - API and Python in general
 - Computational backends (e.g. Cython, TensorFlow)
 - Building an ecosystem (iminuit,...)

} **General impression** in comparison with other HEP packages

Scipy, Imfit, TensorFlow Probability,...

- Lack of specific HEP features
 - *Normalization: specific range, numerical integration,...*
 - *Composition of models*
 - *Multiple dimensions*
 - *Custom models*
- Irrelevant functionality supported in API
 - Survival function, ...

zfit related packages

- Package for phasespace generation of particles
- Covers functionality of TGenPhaseSpace (and more)
- Pure Python (& TensorFlow), integrates seamless with zfit

```
pion = GenParticle('pi+', PION_MASS)
kaon = GenParticle('K+', KAON_MASS)
kstar = GenParticle('K*', KSTARZ_MASS).set_children(pion, kaon)
gamma = GenParticle('gamma', 0)
bz = GenParticle('B0', B0_MASS).set_children(kstar, gamma)

weights, particles = bz.generate(n_events=1000)
```

Zfit: project description

Establish a stable API

- High level libraries (statistics, plotting,...)
 - „code against an **interface**, not an implementation“
- **Replace each component**
 - Allow other libraries to implement custom parts

**Many discussions with community
to avoid splitting/duplication**

Pythonic



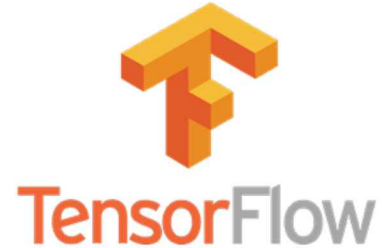
- Pure Python («pip install zfit»)
- Integrated into python ecosystem
 - Load ROOT files (`uproot`, no ROOT dependence!)
 - Use Minuit for minimization (`iminuit`)
 - Data preprocessing with Pandas DataFrame
 - Plotting with matplotlib
 - High level statistics (`lauztat`, more WIP)
- Extendable classes
 - e.g. custom PDF



Scalable: TensorFlow



- Deep Learning framework by Google
- Modern, declarative graph approach
- Built for highly parallelized, fast communicating CPU, GPU, TPU,... clusters
- Built to use «Big Data»



Zfit library examples

Minimize Python function



```
def func(x):  
    x = np.array(x) # make sure it's an array  
    return np.sum((x - 0.1) ** 2 + x[1] ** 4)
```

```
func.errordef = 0.5
```

```
params = [1, -3, 2, 1.4, 11]
```

```
result = minimizer.minimize(func, params)
```

Model, loss building

sum of two pdfs

```
sum_pdf = zfit.pdf.SumPDF([gauss, exponential], fracs=frac)
```

shared parameters

```
mu_shared = zfit.Parameter("mu_shared", 1., -4, 6)
```

```
gauss1 = zfit.pdf.Gauss(mu=mu_shared, sigma=sigma1, obs=obs)  
gauss2 = zfit.pdf.Gauss(mu=mu_shared, sigma=sigma2, obs=obs)
```

simultaneous loss

```
nll1 = zfit.loss.UnbinnedNLL(model=gauss1, data=data1)  
nll2 = zfit.loss.UnbinnedNLL(model=gauss2, data=data2)  
nll_simultaneous2 = nll1 + nll2
```

From
classical

to more
TensorFlow

Model, loss building

Simple combinations

```
func_n = zfit.func.ZFunc(...) # pseudo code  
func = func_1 + func_2 * func_3
```

Composite Parameter

```
pdf = zfit.pdf.Gauss(mu=tensor1, sigma=4)
```

Custom Loss

```
loss = zfit.loss.SimpleLoss(lambda: tensor_loss)
```

=> use all of zfit functionality like minimizers

up to pure
TensorFlow

Model building

```
obs = zfit.Space("x", limits=(-10, 10))
```

```
mu = zfit.Parameter("mu", 1, -4, 6)  
sigma = zfit.Parameter("sigma", 1, 0.1, 10)  
lambda = zfit.Parameter("lambda", -1, -5, 0)  
frac = zfit.Parameter("fraction", 0.5, 0, 1)
```

} parameters

```
gauss = zfit.pdf.Gauss(mu=mu, sigma=sigma, obs=obs)  
exponential = zfit.pdf.Exponential(lambda, obs=obs)
```

} models

Simultaneous fit

```
mu_shared = zfit.Parameter("mu_shared", 1., -4, 6)
sigma1 = zfit.Parameter("sigma_one", 1., 0.1, 10)
sigma2 = zfit.Parameter("sigma_two", 1., 0.1, 10)
```

```
gauss1 = zfit.pdf.Gauss(mu=mu_shared, sigma=sigma1, obs=obs)
gauss2 = zfit.pdf.Gauss(mu=mu_shared, sigma=sigma2, obs=obs)
```

} shared parameters

```
nll_simultaneous = zfit.loss.UnbinnedNLL(model=[gauss1, gauss2],
                                          data=[data1, data2])
```

```
nll1 = zfit.loss.UnbinnedNLL(model=gauss1, data=data1)
nll2 = zfit.loss.UnbinnedNLL(model=gauss2, data=data2)
nll_simultaneous2 = nll1 + nll2
```

} Completely equivalent