# RooFit: model serialization to JSON and other news

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### Introduction

- **RooFit**: C++ library for statistical data analysis in ROOT
  - Model specification and fitting to data (baseline RooFit)
  - Implements common statistical tests (RooStats)
  - Includes tools to specify complex binned models (HistFactory)
- Recent development focused on:
  - **Performance** boost (preparing for larger datasets of **HL-LHC**)
  - More **user friendly** interfaces and high-level tools
- **Topics** of today:
  - Quick overview on recent RooFit highlights
  - RooFit model serialization to JSON/YAML
  - RooFit and **automatic differentiation (AD)**
  - Statistical model **preservation**



### RooFit development areas

#### In which areas does RooFit evolve (besides bugfixes)?



- Not all areas are covered with the same level of activity
- Some areas started to be covered only recently (*automatic differentiation, interoperability*)

### New RooFit computation backend

- New computation backend for likelihood fits ("**BatchMode**") that makes use of vectorization and other optimizations
- Easy to enable in your fits:
  - o pdf.fitTo(data, BatchMode(true))
- **Significant speedup** of likelihood minimization in most RooFit tutorials (up to 7x, see plot on the right)
  - By next release (6.28) hopefully for all of them
- Please **try it out** and open GitHub issues if your fit results are wrong or the fit became slower

RooFit/HistFactory stress tests: speedup of NLL minimization by using BatchMode





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### Status of RooFit's BatchMode

- Architecture-specific accelerator libraries for key functions
  - Optimal one loaded at runtime, given current architecture
  - Now also includes **GPU version**! Try it out with pdf.fitTo(model, BatchMode("cuda"))
- Multithreading via ROOT::EnableImplicitMT()
- Huge speedup for unbinned fits with many events
- For large computation graphs with few events, BatchMode still has larger overhead than recursive evaluation
- Maybe this is exactly the kind model class Belle II uses?



RooFit: speedup in benchmark fits relative to scalar mode (1 million events)

### RooFit pythonizations

- PyROOT bindings **more pythonic** in 6.26
- Now you can for example:
  - use **Python keyword arguments** instead of RooFit command arguments
  - pass around Python sets or lists instead of RooArgSet or RooArgList
  - pass Python dictionaries to functions that take std::map<>
  - implicitly convert floats to RooConstVar in RooArgList/Set constructors
- All pythonizations are <u>documented</u>
- Some Pythonizations to help with C++/Python lifetime issue
  - Still there are memory leaks when returning owning pointers
- See also this <u>ROOT meeting presentation</u>

### *Example code from the <u>rf316 llratioplot.py</u> tutorial showcasing the pythonizations:*

```
# Create background pdf poly(x)*poly(y)*poly(z)
px = ROOT.RooPolynomial("px", "px", x, [-0.1, 0.004])
py = ROOT.RooPolynomial("py", "py", y, [0.1, -0.004])
pz = ROOT.RooPolynomial("pz", "pz", z)
bkg = ROOT.RooProdPdf("bkg", "bkg", [px, py, pz])
```

```
data = model.generate((x, y, z), 20000)
```

# Make plain projection of data and pdf on x observable
frame = x.frame(Title="Projection on X", Bins=40)
data.plotOn(frame)

### RooFit with NumPy, Pandas, and RDF

- ROOT v6.26 new converters between NumPy arrays/Pandas dataframes and RooDataSet/RooDataHist
  - No translation from RooDataHist to dataframe because histograms are in general multi-dimensional
  - Tutorial in <u>Python</u>
- New RooRealVar.bins() function to get RooFit
   bin boundaries as NumPy array
- Creating RooFit datasets from RDataFrame
  - Works for both RooDataSet and RooDataHist
  - Weighted filling still needs to be implemented
  - Tutorial in <u>C++</u> and <u>Python</u>

Example of exporting RooDataSet to Pandas:

from ROOT import RooRealVar, RooCategory, RooGaussian

| <pre>x = RooRealVar("x", "x", 0, 10) cat = RooCategory("cat", "cat",</pre> |    | x        | cat |
|----------------------------------------------------------------------------|----|----------|-----|
| {"minus": -1, "plus": +1})                                                 | 0  | 6.997865 | -1  |
|                                                                            | 1  | 7.211196 | -1  |
| <pre>mean = RooRealVar("mean", "mean",</pre>                               | 2  | 3.198248 | 1   |
|                                                                            | 3  | 5.015824 | 1   |
|                                                                            | 4  | 7.782388 | 1   |
|                                                                            |    |          |     |
| gauss = RooGaussian("gauss", "gauss",<br>x, mean, sigma)                   | 95 | 6.878027 | -1  |
|                                                                            | 96 | 0.475900 | 1   |
|                                                                            | 97 | 4.451101 | -1  |
| <pre>data = gauss.generate((x, cat), 100)</pre>                            | 98 | 3.481015 | -1  |
|                                                                            | 99 | 4.010105 | -1  |
| df = data.to pandas()                                                      |    | 1.25     | 80  |

100 rows × 2 columns

### Parallelized gradient calculation

- For many parameters, most fitting time is spent for the **numeric gradient computation** (re-evaluation after varying each parameter one at a time)
- Distributing the **gradient calculation over multiple processes** is a very general way to speed up fitting (see <u>ACAT 2019</u> presentation)
- Gradient parallelization is part of ROOT 6.26
- It comes together with **new likelihood classes** with improved performance for parallelization over entries





Figure from the ACAT 2019 presentation showcasing the scaling of the gradient parallelization for an ATLAS Higgs combination fit

### 

- The push towards publishing likelihoods is getting stronger
- pyhf has been extremely successful in attracting users
- important reason (among others): ability to define models in a declarative language
  - pyhf JSON is readable, editable, and feature-complete!
- however, limited to HistFactory use-case
  - no "complicated" models, only stacks of homogeneously binned histograms in non-overlapping regions

## A **round-trip-capable**, **human-readable declarative** format for statistical models was missing.

### RooWorkspace ≈ JSON/YAML

- Model-building tools require descriptive languages to define the model
- **JSON** or **YAML** is a well-readable standard industry format
- The new RooFit (6.26) includes a new
   <u>RooJSONFactoryWSTool</u> to import/export
   RooWorkspaces to JSON or YAML
- This can ease interoperability also with other statistics frameworks such as **<u>pyhf</u>** an **<u>zfit</u>**
- Serialization standard referred to as **HS3** (*HEP statistics serialization standard*)

### **Example on the right:** JSON for Gaussian signal with **RooArgusBG** background

```
"pdfs": {
         "signal": {
            "type": "Gaussian",
            "x": "mes", "mean": "sigmean", "sigma": "sigwidth"
        "background": {
            "type": "ARGUS",
            "mass": "mes", "resonance": 5.291,
            "slope": "argpar", "power": 0.5
        },
         "model": {
            "type": "pdfsum",
            "summands": [
                "signal",
                                     More info in this talk
                "background"
                                     on the ROOT users
            "coefficients": [
                "nsig",
                                     workshop 2022
                "nbkg"
             "tags": [
                "toplevel"
"variables": {
        "mes": { "value": 5.25, "min": 5.2, "max": 5.3 },
        "sigmean": { "value": -5.28, "min": 5.2, "max": 5.3 },
        "nsig": { "value": 200, "min": 0, "max": 10000 },
        "argpar": { "value": -20, "min": -100, "max": -1 },
        "nbkg": { "value": 800, "min": 0, "max": 10000 }
```

### The Implementation: RooJSONFactoryWSTool

- **Extensible system** to manage import and export of functions, pdfs and variables
- Two-layer approach:
  - Possibility to **plug in C++ code** for import/export of specific RooFit objects
  - For simpler classes, define mapping of JSON keys to RooFit constructor arguments (*import expressions*) or the RooAbsProxies (*export expressions*)
- More on how to do this in the <u>doxygen page</u>

#### **RooWorkspace to JSON:**

tool = ROOT.RooJSONFactoryWSTool(myworkspace)
tool.exportJSON("myworkspace.json")

```
Import Expressions
                               Export Expressions
                             'RooGaussian": {
"Gaussian": {
    "class": "RooGaussian",
                                 "type": "Gaussian",
    "arguments": [
                                 "proxies": {
                                     "mean": "mean",
        "mean".
        "sigma"
                                     "sigma": "sigma"
},
"Poisson": {
                              RooPoisson": {
    "class": "RooPoisson",
                                 "type": "Poisson",
    "arguments": [
                                  proxies":
        "х",
                                      'mean": "mean"
        mean
}.
```

#### JSON to RooWorkspace:

ws = ROOT.RooWorkspace("somename")
tool = ROOT.RooJSONFactoryWSTool(ws)
tool.importJSON("myworkspace.json")

### Standardizing the top-level JSON content

Going beyond the RooWorkspace, a JSON workspace standard should fulfill these criteria:

1. An arbitrary number of **likelihoods and models** should be storable

0 ?

- 2. **Combinations** of different likelihoods (binned and unbinned) should be very easy!
- 3. The JSON should be **easily manipulable** by different tools and also by hand

What can be written to the JSON should probably include:

o PDFs

• Parameters/parameter lists

- Functions
- Likelihoods/loss functions
- o Data

Parameter "snapshots"Metainfo

Ongoing **discussions** with **ATLAS users, pyhf** and **zfit developers** to explore possibility for a common standard, dubbed the <u>HEP statistics serialization standard</u>

This means the **JSON format** from the RooJSONFactoryWSTool is **not stable yet** 

### Automatic differentiation (AD) in RooFit

#### • **Gradient** of RooFit model essential for minimization

- RooFit uses numeric derivatives, varying one parameter at the time
- Using analytic gradients is much more efficient for many parameters
- We can use **automatic differentiation** techniques to get these gradients
- No code merged yet, but we investigate the following code towards AD:
  - C++ code generation form RooFit model to one C++ function and automatically differentiate with <u>clad</u> (or other source-code transformation fools like <u>Enzyme AD</u>)

#### Automatic Differentiation in ROOT - *Garima Singh* | 2<sup>nd</sup> MODE AD Workshop 14 Sept. 2022

### AD for binned likelihoods from HistFactory

Many binned likelihoods follow a similar pattern:

$$L(\vec{n}, \vec{a} \mid \vec{\eta}, \vec{\chi}) = \prod_{c \in \text{ channels } b \in \text{ bins}} \operatorname{Pois}(n_{cb} \mid \nu_{cb}(\vec{\eta}, \vec{\chi})) \prod_{\chi \in \vec{\chi}} c_{\chi}(a_{\chi} \mid \chi)$$
  
 $\vec{n} : \text{data}, \vec{a} : \text{auxiliary data} \qquad \text{product of Poisson terms} \qquad \text{constraints}$   
 $\vec{\eta} : \text{unconstrained parameters} \qquad \vec{\chi} : \text{ constrained parameters}$ 

HistFactory is a higher-level tool to build such likelihoods in RooFit.

Good model class for showing AD in RooFit:

- many parameters
- rich computation graph
- few normalization integrals







### Preliminary Results

Explicit Computation Graphs: An Example HistFactory Model



An example histogram fitting model with 2 bins and 2 channels, with 3 samples per channel. Based on the <u>hf\_001 example</u>.

### Preliminary Results

| <pre>double nll(double *in) {     double nomGammaB1 = 400;     double nomGammaB2 = 100;     double nominalLumi = 1;</pre> | straints defined as calls to respective 'evaluate's.                                                                                      | <pre>// cont double mu = 0; double temp;</pre>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           |
|---------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <pre>double constraint[3] {RooPoisson::poisson(r</pre>                                                                    | <pre>inomGammaB1, (nomGammaB1 * in[0])), inomGammaB2, (nomGammaB2 * in[1])), i[2], nominalLumi, 0.100000)};  Translated RooProdu NL</pre> | <pre>double nllSum = 0;<br/>unsigned int b1, b2, b3;<br/>for (int iB = 0; iB &lt; 2; iB++) {<br/>b1 = RooHistFunc::getBin(binBoundaries1, x[iB]);<br/>b2 = RooHistFunc::getBin(binBoundaries2, x[iB]);<br/>b3 = RooHistFunc::getBin(binBoundaries3, x[iB]);<br/>mu = 0;<br/>mu = 0;<br/>mu += sig[b1] * (in[3] * in[2]);<br/>mu += (bgk1[b2] * histVals[iB]) * (in[2] * 1.000000);<br/>mu += (bgk2[b3] * histVals[iB]) * (in[2] * 1.000000);<br/>mu += (bgk2[b3] * histVals[iB]) * (in[2] * 1.000000);<br/>temp = std::log((mu));<br/>nllSum -= -(mu) + weights[iB] * temp;<br/>} </pre> |
| <pre>for (int i = 0; i &lt; 3; i++) {     cnstSum -= std::log(constraint[i]); } // cont</pre>                             | Constraint sum.                                                                                                                           | <pre>return cnstSum + nllSum; }</pre>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |

### Automatic Differentiation in RooFit



### RooFit and analysis preservation

- RooFit has a few things going for it in the analysis preservation department:
  - The **RooWorkspace** is widely used in the community
  - Few software dependencies (only ROOT, which has only few dependencies itself)
  - We make **backwards** compatibility a **priority** (...at least since a few years)
  - No hurdles for fixing preservation issues upstream (as it's a HEP community project)
- But there are also problems:
  - Model specification tightly connected with implementation
  - The correct schema evolution of all RooFit objects is a big burden

The **JSON/YAML serializatio**n and **C++ code generation** approach can be useful to *overcome these shortcomings*!

Maybe we will have something

### Summary

- RooFit is **evolving** steadily
  - Support and development from **ROOT team** at CERN and **external contributors**
- Highlights of the recent version 6.26 are the **BatchMode** and the **Pythonizations** 
  - as well as the JSON serialization explained in more detail
- Status of the **JSON** to Workspace tool:
  - Support for most PDFs and functions, easily extensible by users
  - Standardization of JSON structure still in progress (HS3 project)
- Future developments will focus on **automatic differentiation** 
  - Approach of differentiable **C++ code generation** could benefit analysis preservation
  - Still at early R & D stage, don't expect much of this in upcoming ROOT 6.28 yet



#### Recent RooFit **presentations**:

- Talk at ACAT 2021
- RooFit talk at the <u>ROOT users workshop</u>
- RooFit talk at ICHEP 2022
- <u>Automatic differentiation in ROOT</u> (September 2022)

To get more info on the **JSON serialization**:

- The HEP statistics serialization standard <u>GitLab repository</u>
  - You can download the PDF as an artefact of the <u>CI pipeline</u>
- The landing page for the JSON tool in the RooFit documentation
- The <u>RooFit tutorial</u> that explains how to import a model from JSON
- For developers: the <u>RooFit HS3 README file</u> explaining how to extended model support