

A Python package for distributed ROOT RDataFrame analysis

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PyHEP 2021

ROOT

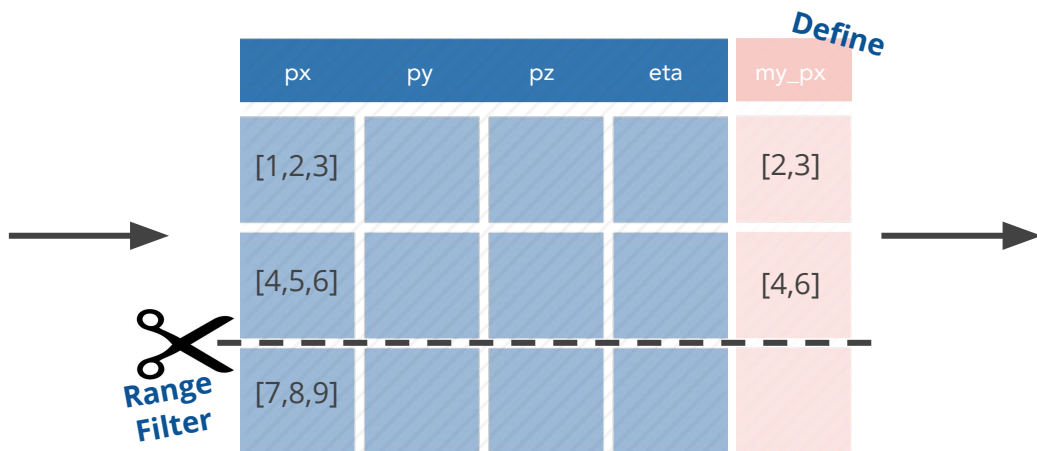
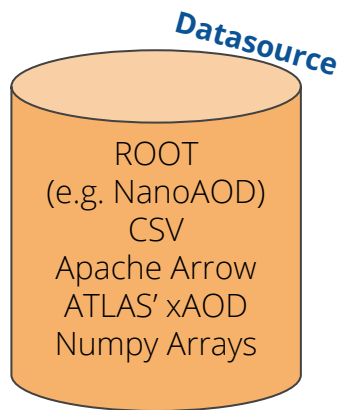
Data Analysis Framework

<https://root.cern>

1. RDataFrame: ROOT's declarative interface for analysis
2. Distributed Analysis with RDataFrame
 - a. Programming model
 - b. Backends
3. Demo on SWAN
 - a. Dimuon analysis with RDataFrame
 - b. Local, Spark and Dask: execution and monitoring
4. Performance tests
5. Summary



RDataFrame analysis interface



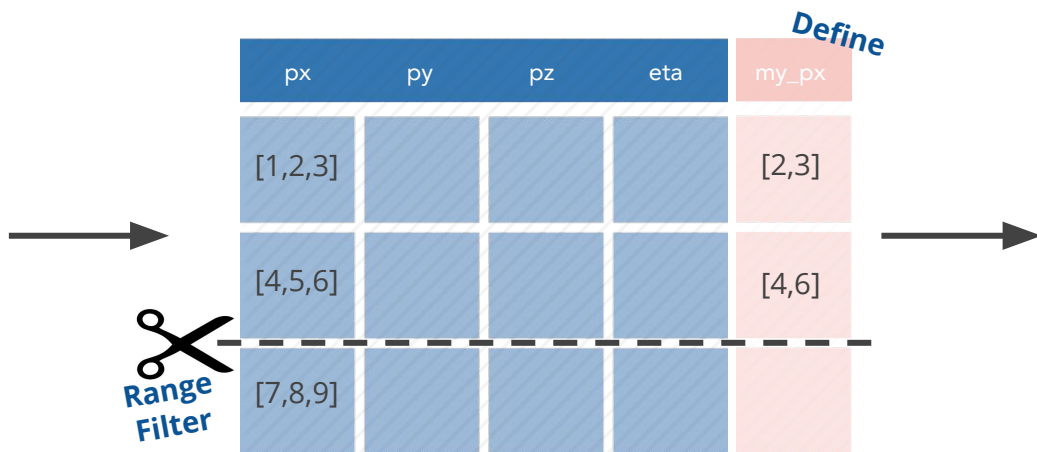
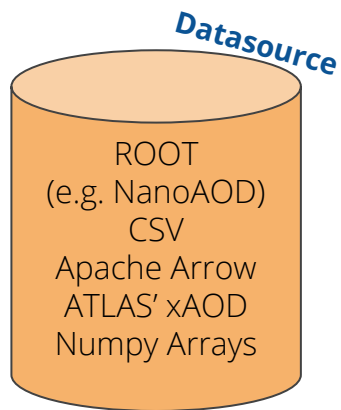
```
# enable multi-threading  
ROOT.EnableImplicitMT()  
df = ROOT.RDataFrame(dataset)
```

```
df = df.Range(2)  
    .Define("my_px", "px[eta > 0]")
```

```
# filled in a single pass  
h1 = df.Histo1D("my_px", "w")  
h2 = df.Histo1D("px", "w")
```



RDataFrame analysis interface



- histograms, profiles
- new ROOT files
- cut-flow reports
- data reductions (mean, sum,..)
- any user-defined operation

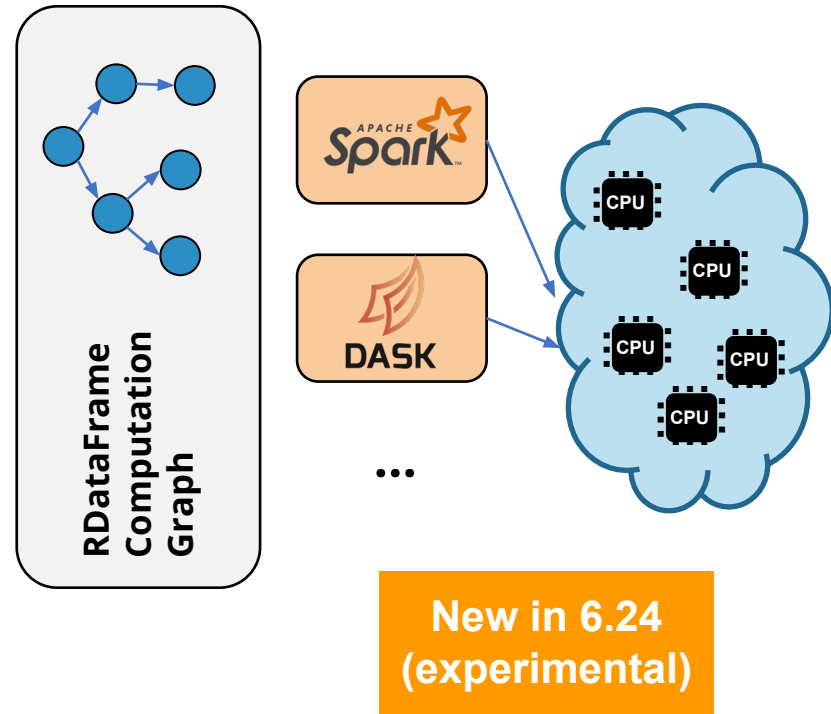
Goal

Best performance and ease of use across the board,
for most (all?) HEP analysis use cases

[RDataFrame reference guide](#)

[RDataFrame tutorials](#)

- Enable **interactive large-scale distributed** data analysis with RDataFrame
- Can run with different schedulers
 - **Spark**
 - **Dask**
 - ...
- Analysis from start to end in a **single interface**





Programming Model

Local

```
from ROOT import RDataFrame
```

Importing RDataFrame

```
df = RDataFrame('treename', 'filename.root')
```

Constructing RDataFrame

Rest of application

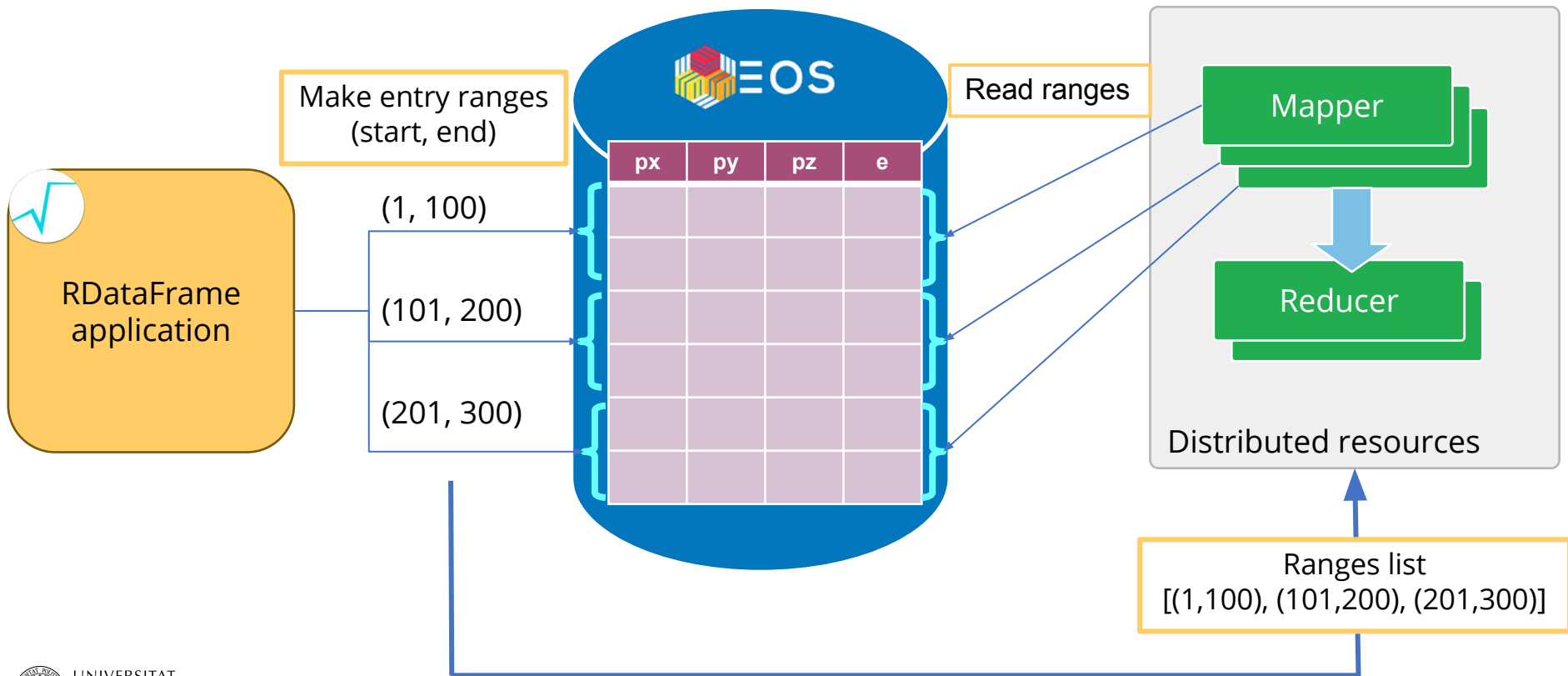
```
df2 = df.Filter(...).Define(...)  
h1 = df2.Histo1D(...)  
h1.Draw()
```

Distributed

```
import ROOT  
RDataFrame = \  
ROOT.RDF.Experimental.Distributed.Dask.RDataFrame
```

```
from dask.distributed import Client  
df = RDataFrame('treename', 'filename.root',  
                daskclient = Client('tcp://hostname:port'))
```

Behind distributed RDataFrame



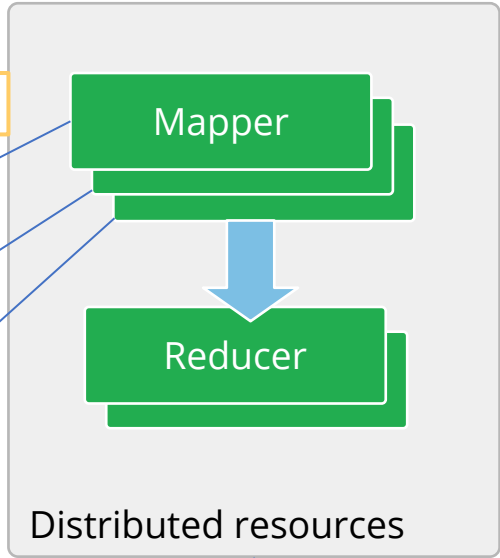
RDataFrame application

Make entry ranges (start, end)

- (1, 100)
- (101, 200)
- (201, 300)

px	py	pz	e

Read ranges

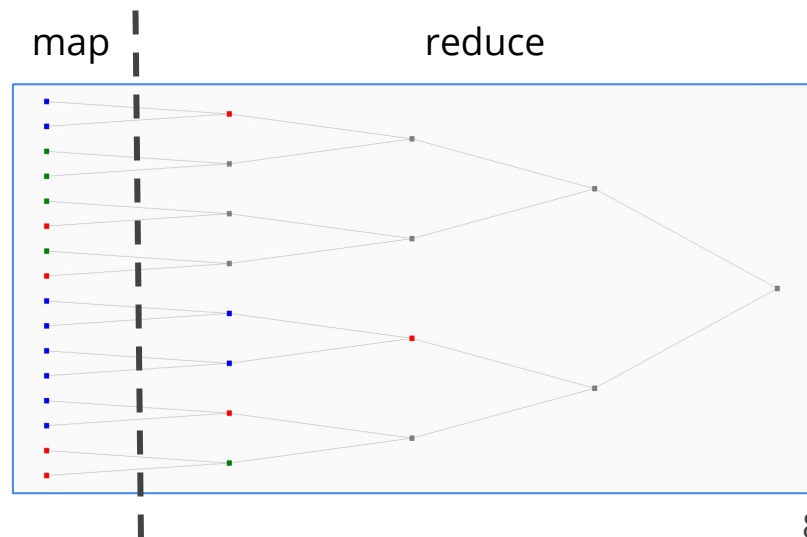


Ranges list [(1,100), (101,200), (201,300)]

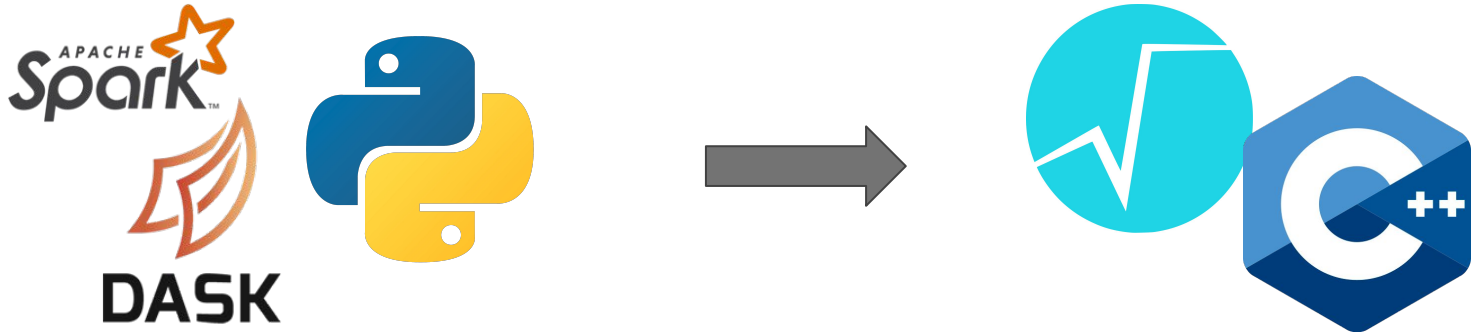


Dask vs Spark for distributed RDataFrame:

- ▶ `dask.delayed` vs `spark.map` & `spark.treeReduce`
- ▶ Same MapReduce tree pattern, different API
- ▶ Spark processes in stages
 - first map, then reduce
- ▶ Dask can reduce two completed mappers while others are still being processed



- ▶ RDataFrame interfaces (in Python) with schedulers such as Dask or Spark for task distribution
- ▶ RDataFrame tasks run mostly in **C++** via PyROOT
 - I/O and event loop execute in C++



Notebooks on GitHub



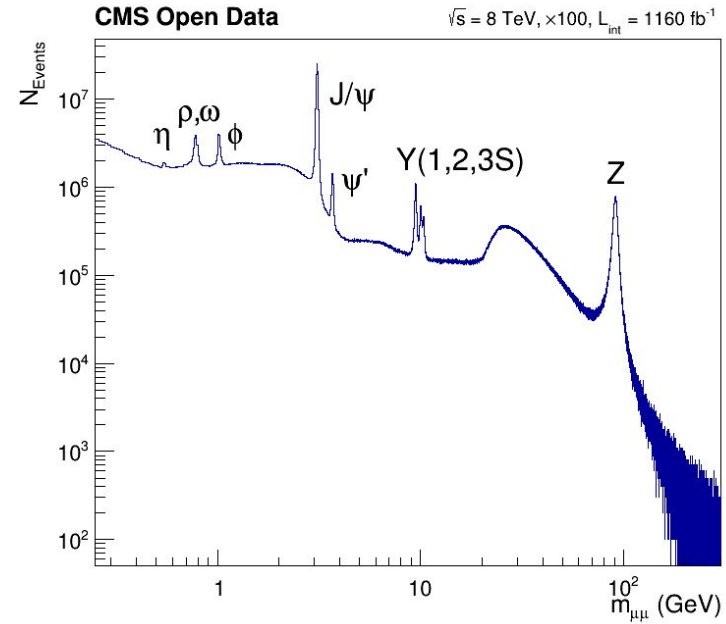


Cluster of CERN OpenStack VMs:

- ▶ 13 VMs
 - 4 cores, 7.3 GB RAM, 40 GB spinning disk
- ▶ 1 reserved for the Dask scheduler / Spark master
- ▶ 12 Dask/Spark workers → 48 cores, 88 GB RAM

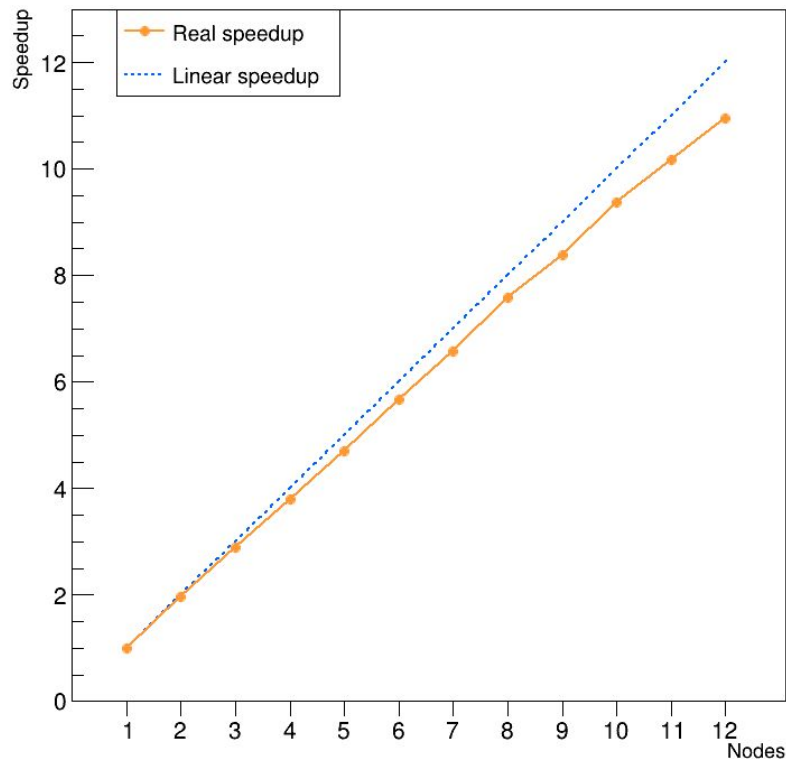
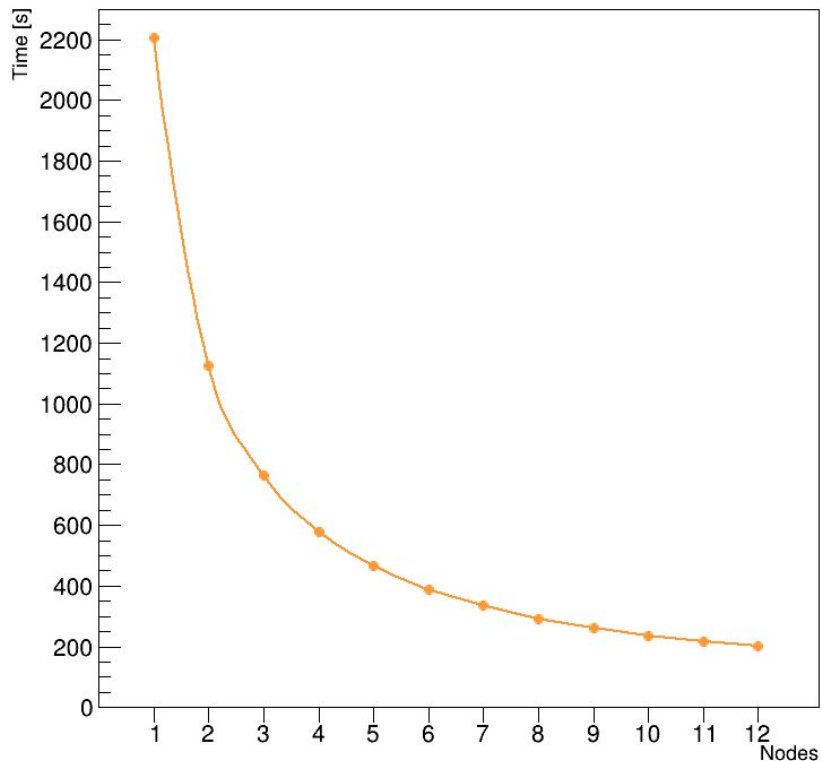
Test setup: dataset

- ▶ Dimuon analysis
- ▶ 100x original dataset
 - 210 GB
 - ZLIB compressed
- ▶ All data is read and processed in the analysis



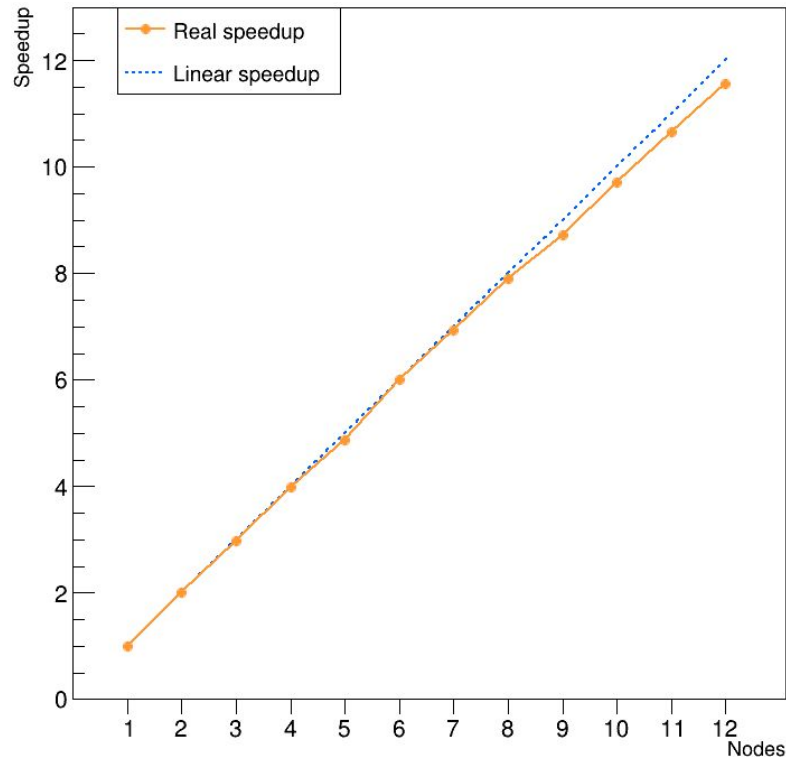
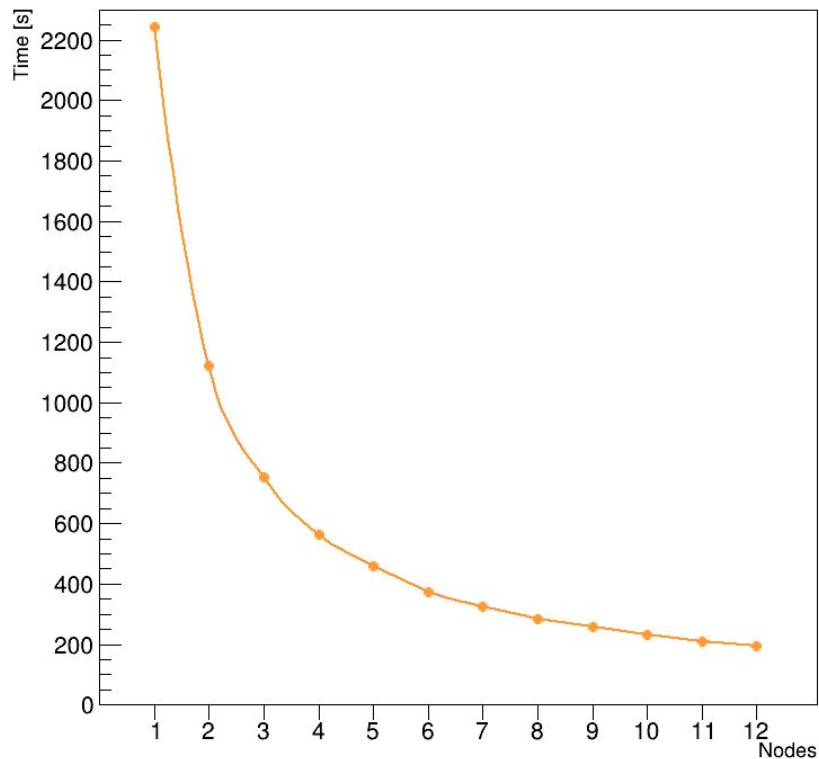


Dimuon analysis performance (Dask)



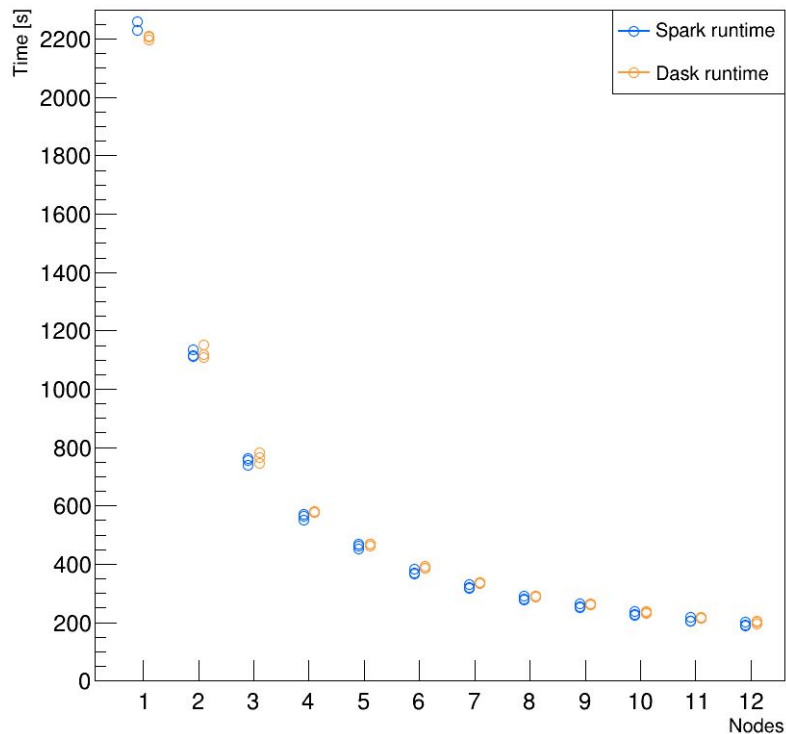


Dimuon analysis performance (Spark)





Spark vs Dask comparison



- ▶ RDataFrame: ROOT's declarative interface for data analysis
- ▶ Since ROOT 6.24: also **distributed!**
 - Requires minimal changes to original (local) application
 - Task scheduling with Spark or Dask (for now!)
- ▶ Useful links
 - [RDataFrame reference guide](#)
 - [Tutorial: distributed RDataFrame with Spark](#)