

# Analysis Benchmarks Where are the bottlenecks?

Luke Kreczko (CMS/LZ) 24th March 2020



# First ECHEP workshop

During first workshop we had a lot of emphasis on processing speed

- Some known inefficiencies in generators (e.g. negative weights, lexical\_cast)
- Benefits of SIMD, trading accuracy for speed in simulation (fast-sim)
- Alternative architectures (GPU, FPGA, etc) to accelerate software in HEP

Improvements usually identified through profiling & benchmarks

In some cases a lot of expert knowledge went into improvements

# One possible 6 month plan

- Step 1: Identify several benchmark analyses (1 month)
  - Open data, coordinate between experiments
- Step 2: Implement using existing tools (2.5 months)
  - Coffea or FAST-HEP
  - Directly with PARSL or DASK where data not amenable to columnar approach
  - Deliverable: where do these existing tools struggle for different analyses / experiments
- Step 3: Understand caching requirements (2.5 months)
  - Timing without caching
  - Deliverable: Profiling for second run with caching
  - Deliverable: Size of caching required
- Step 4 (extension): Explore caching between analyzers (?)

(the near future is Python)

# Python for accelerated development

- High level programming → fast to try something
- Compact language → fewer lines of code, fewer bugs
- Quick refactoring
- Access to Big data tools → ML + distributed processing
- Full-Stack prototyping (even FPGA)
- Can be easy to change architectures (e.g.  $CPU \rightarrow GPU$  with <u>numba/tensorflow</u>)

#### Numba: Python just-in-time compiler

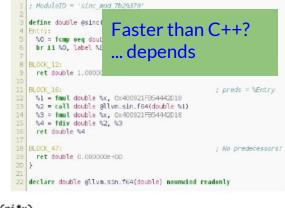
- Few 'array-oriented' compilers though common use case and hardware optimizations exist.
- Wasn't possible few years ago, Python faster than your C++ code.

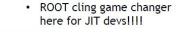
@vectorize
def sinc(x):

if x==0.0: return 1.0

else:

return sin(x\*pi)/(pi\*x)





https://zenodo.org/record/1418513#.XniKnoj7TAQ

# Step 1: Identify several benchmark analyses

Start with some small subtasks for guidance

# **HEP Data analysis**

Compared to dedicated efforts in generators and simulation, data analysis is more varied

- Analysis methods & algorithms can differ widely between groups
- Code is usually written by non-experts
- Mixture of using experiment software frameworks and <u>standalone code</u>

Difficult to utilize advanced techniques or new architectures

# Analysis benchmarks

First two benchmarks <u>implemented</u> - meant to highlight big differences between individual analysis steps

- Selecting events with at least 4 jets with pt > 30 GeV and |eta| < 2.4 (e.g. a loose skim)
  - $\circ \quad 0 \text{ to N jets per event} \rightarrow \text{loops of depth 2}$
  - Uses <u>CMS Top Quark open data</u> (6,423,106 events)
- Categorise ttbar decay channel
  - 10 different decay channels (from full hadronic to full leptonic)
  - Need to traverse genparticle decay chain  $\rightarrow$  loops of depth 3 & many comparisons
  - Uses private MC due to missing information in Open Data (1,441,999 events)

Processed on CERN GitLab CI, data are copied to local disk

## Selection results (6,423,106 events)

method	HDD (CERN CI)	SSD	comment
numpy	18.5 s	8.4 s	Advanced python
Loop depth 1	53.2 s	30.5 s	Advanced python
Loop depth 2	214.5 s	128.2 s	Beginner python
C++ loop depth 2	6.6 s	4.4 s	Advanced ROOT

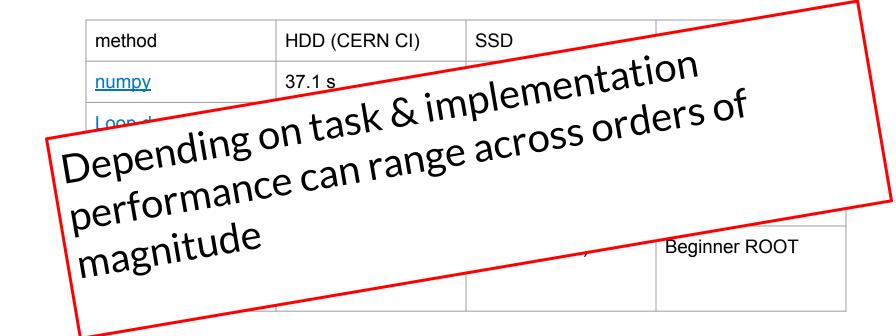
Advanced ROOT: using ROOT features that are not that common (yet) but recommended (e.g. TTreeReaderArray) - ROOT usage is stuck "in the past" due to inheritance of macros

## (fresh) Decay channel results (1,441,999 events)

method	HDD (CERN CI)	SSD	comment
numpy	37.1 s	12.5 s	Advanced python
Loop depth 3	1436.8 s	822.4 s	Beginner python
C++ loop depth 3	8.9 s	4.3 s (18.5 s)	Advanced ROOT
<u>C++ GetEntry</u>	308.9 s	148.3 s	Beginner ROOT
<u>C++ GetEntry +</u> disabling unused branches		15.7 s (51.2s)	Beginner ROOT

Note 1: Arrays stored in NanoAOD do not work well with SetBranch method (output is wrong for some events)) Note 2: Using namespaces in the ROOT macro, increases processing time (???)

### (fresh) Decay channel results (1,441,999 events)



Note 1: Arrays stored in NanoAOD do not work well with SetBranch method (output is wrong for some events)) Note 2: Using namespaces in the ROOT macro, increases processing time (???)

# Step 2: Implement using existing tools

# Moving from implementation to description

FAST-HEP tools offer another approach

- Describe what you need, tools will do the rest
- Optimizations can happen in the background
  - This includes architecture selection & parallel processing (local & distributed)
- What is the fraction of covered use-cases?
  - We know it works for some CMS, LZ and DUNE analyses
    - Including a range of distributed backends (e.g. from Coffea project)
  - Performance issues with nested types (e.g. vector<vector>>)
    - Awkward-array 1.0 will fix these
  - $\circ$  Multi-tree input (e.g. CMS L1 trigger prototyping, standard LZ analyses) not yet functional

# **CMS Efforts overview**

### CMS Analysis Facility/real-time data query system

#### Steps 3 & 4 have certainly overlap

- Analysis with Apache Spark
  - Caching capabilities
- <u>Real-time data query system</u>
  - Explores fast data query and caching
  - Shows big difference depending on how the data are accessed
  - code transformation performance should be similar to <u>TTreeReaderArray</u>
- Now all under <u>ServiceX</u>
  - Use Kafka for streaming
  - Cache results for instant replay

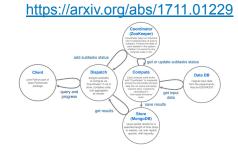
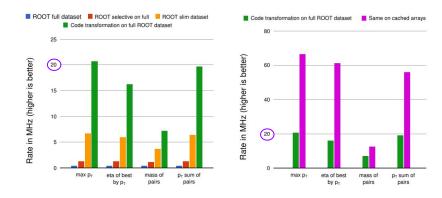


Figure 2. Schematic for distributed query processing to minimize cache misses (see text).



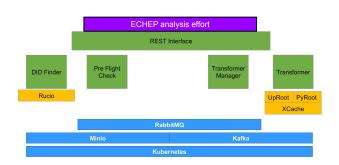
## **New CMS effort**

Software and Computing R&D and code modernization/performance

- Just started: <u>First meeting</u> on the 12th of March
- Many areas targeted
  - Not sure how much of it is public knowledge (or allowed to be)
- But, due to the nature of the effort:
  - Significant overlap with ECHEP

# Summary

#### **Questions and Outlook**



#### First benchmarks implemented

- What is the best analysis to highlight current bottleneck?
- Are these the same between ATLAS & CMS?
- Do we have something similar for LHCb (and other experiments)?

Next: implement with existing (high-level) tools

- FAST-HEP + Coffea (Parsl backend)
- Current example highlight the difficulty of replacing for-loops

#### Other efforts exist

- Mostly going in a similar direction
  - Minimize overlap, maximize syngery
- User  $\rightarrow$  ServiceX for steps 3+4 might be an option

# **Backup slides**

# **Other thoughts**

(as a by-product of step 2)

# **Expertise vs automation**

Other than being an expert in C++, Python & <distributed computing system of your experiment, department, funding agency's choice > what can be done?

Automation:

- E.g. CERN CI has access to EOS
  - can a mini-version of the analysis be run in CI (tests & profiling)?
  - Do we have to provide skeleton analysis or training for this?
- Is there a way to test this in the submission systems?
  - $\circ \quad \mathsf{CI} \mathop{\rightarrow} \mathsf{submission} \ \mathsf{system} \ \mathsf{for} \ \mathsf{scaling} ?$

# TODO

Do we have to worry about producing plots?

• It can be time-consuming, but usually is not

Statistical analysis (RooFit, PyHF, Minuit) can take a lot of time

• Which tool is the best (fast & reliable)?

Putting everything into ECHEP context:

Analysis is fraction X of total computing budget, following route Y would reduce usage by Z.  $\rightarrow$  does this translate into analysis improvements, more resources for other analyses or just saving money?

## **Other items**

HSF simulation on non-LHC simulation requirements: <u>https://indico.cern.ch/event/899153/</u>

# Draft roadmap

Attempt to measure analysis bottlenecks

- Focus on specific algorithms within different analyses
  - Compare SIMD/GPU to typical implementations
- Provide training material for advanced algorithms
  - Translating an analysis into SIMD or different architectures is not trivial
- Advertise tools (e.g. FAST-HEP, Coffea) that can improve transitions for non-experts

Focus on Python (compare to C++) as it is a friendlier language