

# Spark-like and query-like analysis systems and tools

(what works, what doesn't work, what's needed, and what we're building to fill that need)

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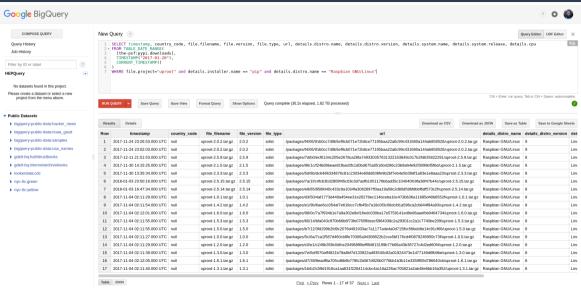
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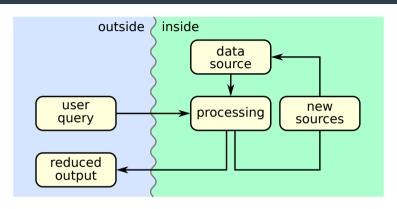
Consider this request: "Please submit a GRID job to plot the muon  $p_T$  spectrum. Then we'll think about what we want to look at next." Is that unreasonable?

# An example of what I have in mind: Google BigQuery

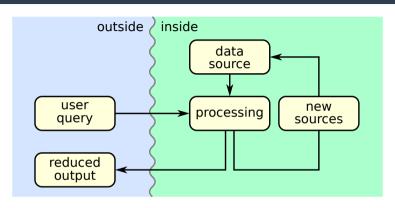






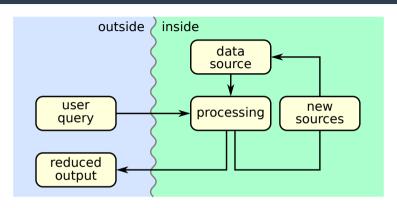






The "reduced output" must be small enough to download quickly (e.g. histograms or highly skimmed tables). If not, the system ceases to be "exploratory."

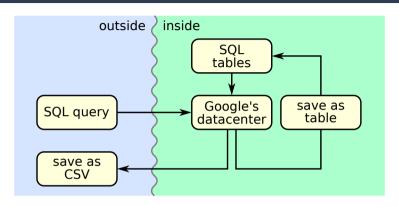




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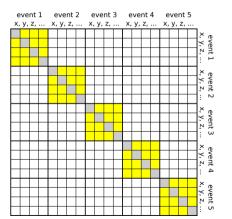
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  - However, *for every one of these*, you quickly encounter "not implemented yet" messages when you try to use them for HEP events.
- ▶ Reason #3: User interface. The web form is fun, but we need queries embedded in an interactive, programmable environment with plotting and statistics libraries: ROOT or Python or both (probably both).

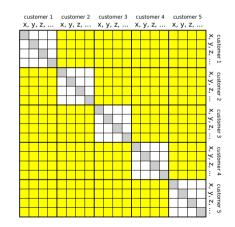
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# HEP analysis is simpler

HEP functions never have to cross events. Common elsewhere: e.g. market basket analysis.



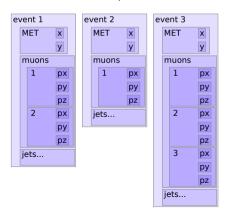


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# HEP analysis is more complex

HEP data are variable-length, nested data structures, and we typically need to loop over combinations of particles.



In many fields, data are not considered ready for analysis until they're in a tabular form. (Earlier steps are called "tidying.")

	column 1	column 2	column 3
row 1			
row 2			
row 3			
row 4			
row 5			
row 6			

# HEP is both simpler and more complex than relational analytics



	independent events	all-to-all shuffle
tabular data	basic spreadsheets	relational analytics
variable-length, nested data	HEP analysis	graph analytics



#### Identically typed, variable-length, nested data can be split into columns:

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- ▶ **Share cache:** *column* popularity distribution is steeper than the *file* popularity distribution; most popular columns could remain in memory for all users.
- ▶ **Avoid touching disk:** sorted partitions do not need to be fully read if a cut is being applied to the sorted variable (most likely  $p_T$ ).



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- $\sqrt{\ }$  Array fetching is through an arbitrary dict-like object: may defer to disk, network, or per-column cache.
- $\sqrt{}$  OAMap has enough indirection that different particle types can be *independently* sorted by their own  $p_T$ s. Thus, a filter like "muon  $p_T > X$  and jet  $p_T > Y$ " may touch disk for only half the muon data and half the jet data.

# OAMap example



```
>>> import uproot
>>> import oamap.source.root
>>> url = "http://scikit-hep.org/uproot/examples/HZZ.root"
>>> events = uproot.open(url)["events"].oamap()
>>> events.schema.content["muons"].show()
List (
  starts = 'NMuon',  # schema maps object attributes to array names
  stops = 'NMuon',
  content = Record( # at all levels of nesting
    fields = {
      'px': Primitive(dtype('float32'), data='Muon_Px'),
      'py': Primitive(dtype('float32'), data='Muon_Py'),
      'pz': Primitive(dtvpe('float32'), data='Muon Pz'),
      'energy': Primitive(dtype('float32'), data='Muon_E'),
      'charge': Primitive(dtype('int32'), data='Muon_Charge'),
      'isolation': Primitive(dtype('float32'), data='Muon Iso')
    }))
```

# OAMap example



#### The dataset looks like a nested Python list.

#### But it is generated on demand from arrays.

```
"NMuon": array([2, 1, 2, ..., 1, 1, 1], dtype=int32)
"Muon_Px": array([-52.899456, 37.73778, -0.81645936, ...,
-29.756786, 1.1418698, 23.913206], dtype=float32)
```

# OAMap example

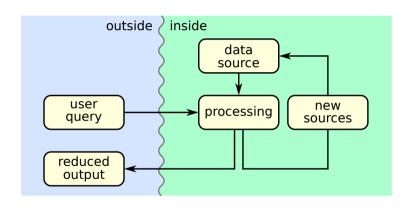


It can also be included in compiled code with no change in syntax.

```
>>> import numpy
>>> import numba
>>> import oamap.compiler
>>> @numba.njit
                                # declares the following function to be compiled
... def compute (events, out):
   i = 0
    for event in events: # "event" and "event.muons" are a compiler fiction
           if len(event.muons) == 2:
                mu1. mu2 = event.muons[0]. event.muons[1]
                px = mu1.px + mu2.px
               pv = mu1.pv + mu2.pv
. . .
               pz = mu1.pz + mu2.pz
               energy = mul.energy + mul.energy
. . .
               out[i] = sqrt(energy**2 - px**2 - py**2 - pz**2)
. . .
                i += 1
. . .
>>>  out = numpy.empty(1371)
>>> compute (events, out) # compilation and array-fetching happen on first call
>>> 011t
array([90.22780609, 74.74654388, 89.75765991, ..., 92.06494904,
       85.44384003, 75.960617071)
```

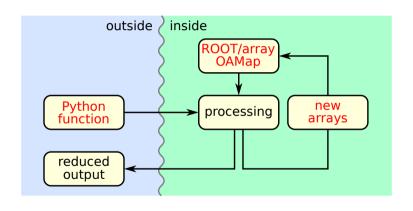
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- ▶ I had been missing an important fact: most of Pandas's functionality is in its handling of *indexes*.
  - ► TTrees/events are indexed only by entry or run/event number.
  - ▶ Pandas indexes may be non-contiguous, non-numeric, intervals/durations, multi-component, . . . , and every operation maintains consistent indexes.

# What should our reduced output be?



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- ▶ Pandas has more in common with *histograms* than it does with *event sources*.

### Just an example...



I'll illustrate these operations through a series of examples. Consider a small DataFrame with string arrays as row and column indexes:

Using the stack method on this data pivots the columns into the rows, producing a Series:





#### import pandhist

```
# define bins in many dimensions; we'll think about how to plot later
muonhist = (pandhist)
            .bin(100, 0, 500, "mass")
            .cut("q1*q2 < 0")
            .irrbin([0.2, 0.5], "mt1")
            .irrbin([0.2, 0.5], "mt2")
            .fillable()) # creates a fillable Pandas DataFrame
for muons, charge, mt2activity in uproot.iterate(
  "RA2Analysis/*.root", "TreeMaker2/PreSelection",
  ["Muons", "Muons charge", "Muons MT2Activity"], outputtype=tuple):
    for i in range(len(muons)):
        if len(charge[i]) == 2:
            mu1, mu2 = muons[i]
            q1, q2 = charge[i]
            mt1, mt2 = mt2activitv[i]
            # fill method has an argument for each variable
            muonhist.fill((mu1 + mu2).mass, q1, q2, mt1, mt2)
                                                                    17 / 25
```

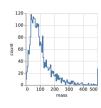
>>> muonhist



```
count
                                                     # in this example, count is
mass q1*q2 < 0 \text{ mt1} mt2
                                                     # the only column; the rest
[-\inf, 0.0) fail [-\inf, 0.2) [-\inf, 0.2) 0.0
                                                    # is a hierarchical index
                                               0.0
                                                     # (mass, q1*q2 < 0, mt1, mt2)
                                  [0.2, 0.5)
                                  [0.5, inf) 0.0
                       [0.2, 0.5) [-inf, 0.2) 0.0
                                  [0.2, 0.5) 0.0
                                                    # if we asked for weights,
                                 [0.5, inf) 0.0
                                                    # sumw and sumw2 would be
                       [0.5, inf) [-inf, 0.2) 0.0
                                                     # separate columns
                                  [0.2, 0.5) 0.0
                                  [0.5, inf)
                                               0.0
                      [-\inf, 0.2) [-\inf, 0.2) 0.0
                                                     # if we asked for a profile,
             pass
                                  [0.2, 0.5)
                                               0.0
                                                     # we'd get sum(v) and sum2(v)
                                  [0.5, inf) 0.0
                       [0.2, 0.5) [-inf, 0.2)
                                               0.0
                                  [0.2, 0.5) 0.0
                                                     # but most of the work is done
                                 [0.5, inf) 0.0
                                                     # by the hierarchical index
                       [0.5, inf) [-inf, 0.2) 0.0
                                  [0.2, 0.5)
                                               0.0
                                  [0.5. inf)
                                               0.0
[1836 rows x 1 columns]
```

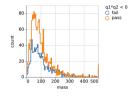


>>> pandhist.steps("mass").data(muonhist)



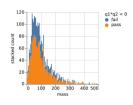


>>> pandhist.steps("mass").overlay("q1\*q2 < 0").data(muonhist)



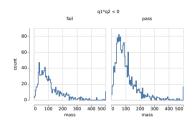


>>> pandhist.area("mass").stack("q1\*q2 < 0").data(muonhist)



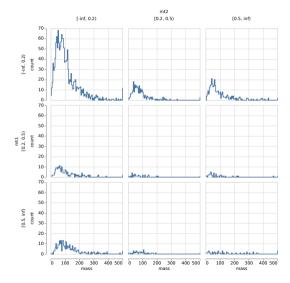


>>> pandhist.steps("mass").column("q1\*q2 < 0").data(muonhist)



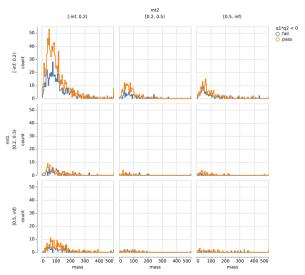


>>> pandhist.steps("mass").row("mt1").column("mt2").data(muonhist)

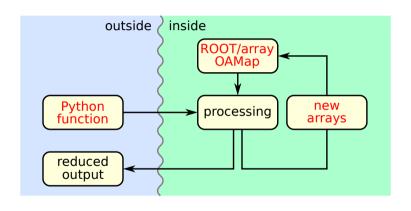




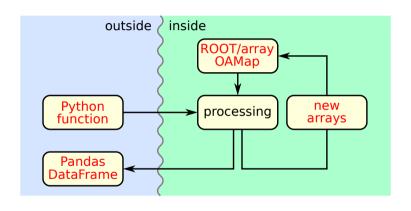
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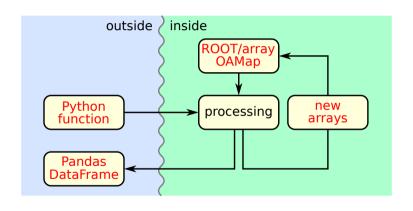


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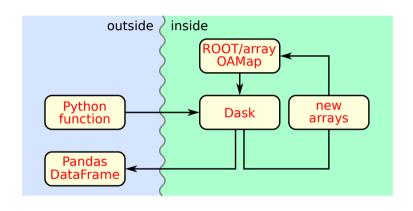
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- ► Thanat Jatuphattharachat (summer student) investigated raw Zookeeper coordination, but this is getting close to DIY.
- ▶ Dask looks like a good choice: it's in the C++/Python world, general enough to piece together what we need out of basic parts.

(More about this in the concurrency session.)













#### Again, no need for HEP to invent.

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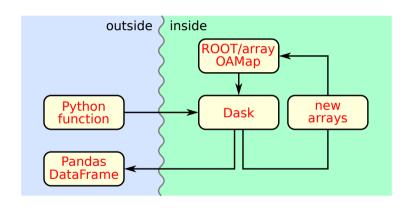


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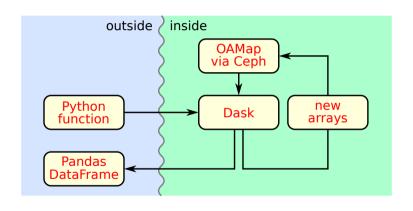


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  - Lets us experiment both ways.
  - We could use the static function to place executable tasks, improving data locality...









### Conclusions



Though an analysis service should use as many open source parts as possible, some functionality is missing and we need to build our own.

#### Software products developed along the way:

uproot	Quickly access ROOT branches as arrays.	mature, in use
oamap	Translate between object-oriented Python and low-level array operations.	ready for testing
pandhist	Reinterpretation of Pandas DataFrames as superhistograms. (Plotting in Vega-Lite.)	experimental
vegascope	Browser-based TCanvas for Vega/Vega-Lite, so that you don't have to use Jupyter if you don't want to.	done (simple)
?	OAMap as a collection in Dask.	
?	Column manager (for sharing data among datasets).	
?	All-in-one environment for query-based analysis.	