



The Case for Columnar Analysis (a two-part series)

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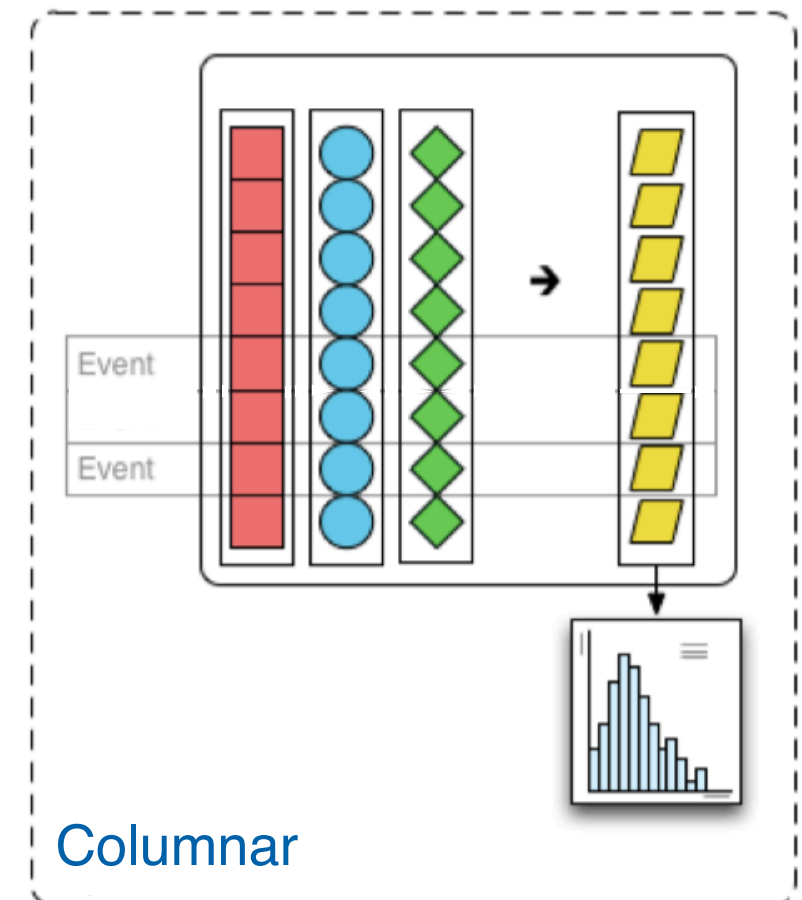
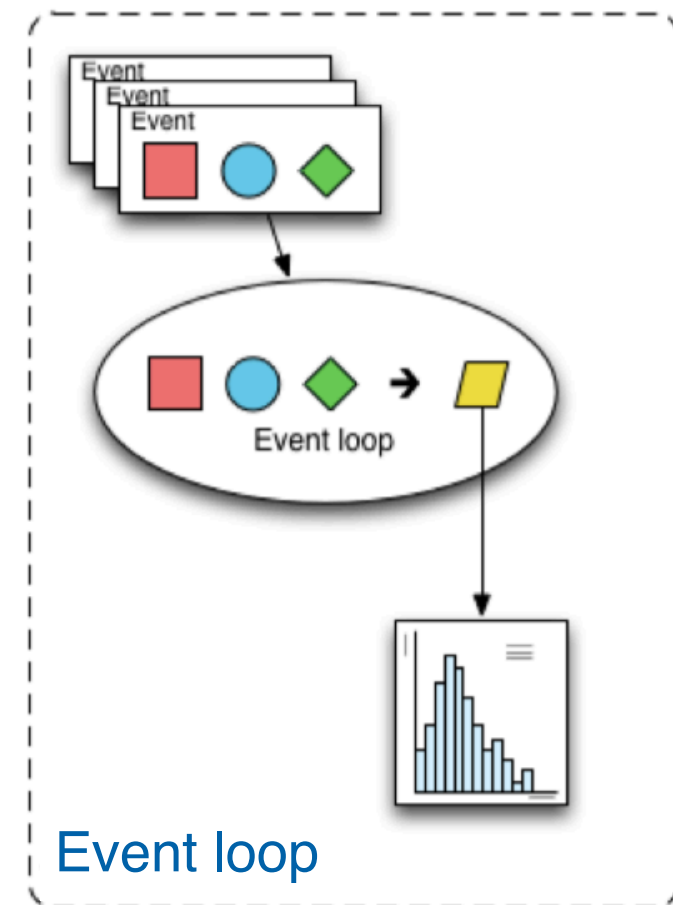
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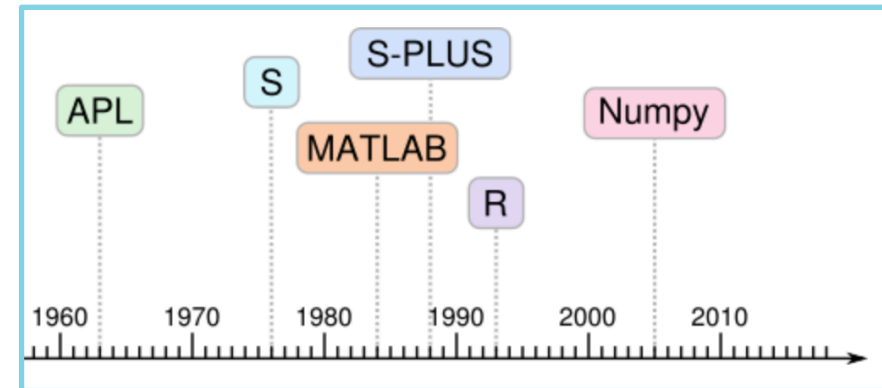
Prologue: terminology

- Event loop analysis:
 - Load relevant values for a specific event into local variables
 - Evaluate several expressions
 - Store derived values
 - Repeat (explicit outer loop)
- Columnar analysis:
 - Load relevant values for many events into contiguous arrays
 - Nested structure (array of arrays) → flat content + offsets
 - This is how TTree works!
 - Evaluate several **array programming** expressions
 - Implicit *inner* loops
 - Store derived values



Prologue: technology

- Array programming:
 - Simple, composable operations
 - Extensions to manipulate offsets
 - *Not declarative* but towards goal
- Awkward array programming:
 - Extension of numpy syntax
 - Variable-length dimensions: “jagged arrays”
 - View SoA as AoS, familiar object syntax, e.g. `p4.pt()`
 - References, masks, other useful extensions
 - See [awkward](#), talk by J. Pivarski at [ACAT2019](#)
- Coffea framework:
 - Prototype analysis framework utilizing columnar approach
 - Provide lookup tools, histogramming, other ‘missing pieces’ usually 1
 - See [fnal-column-analysis-tools](#)
 - Functionality will be factorized as it matures



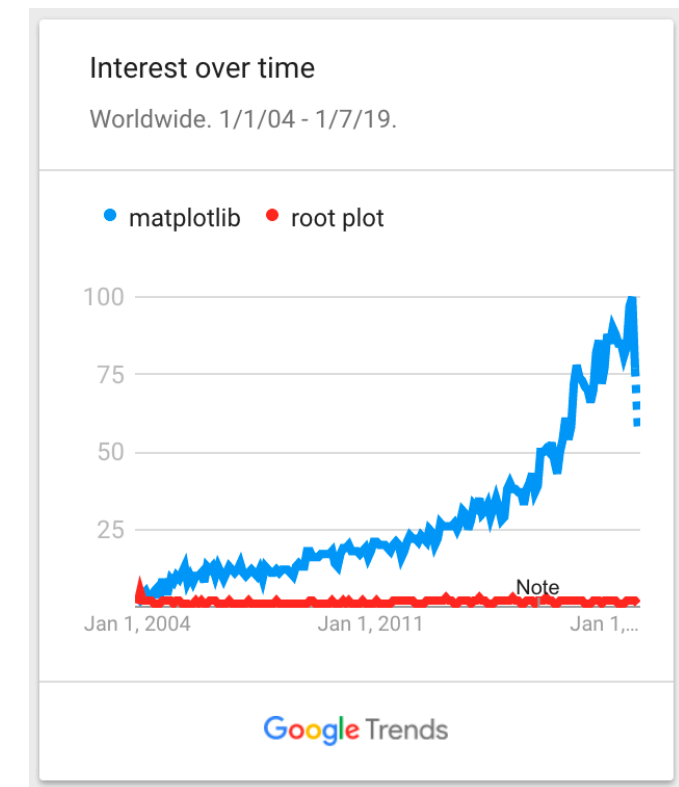
Awkward
Array



Part I: Analyzer Experience

User experience

- Unsurprisingly, #1 user priority
 - Any working analysis code can scale up...for now
 - c.f. usage of PyROOT event loops despite dismal performance
 - (this will *never* change)
- Fast learning curve for scientific python stack
 - Excellent ‘google-ability’
 - The quality and quantity of off-the-shelf components is impressive—many analysis tool implementations contain very little original code
 - Essentially all functions available in a vectorized form
- Challenge: re-frame problem in array programming primitives rather than imperative style (for+if)
 - User [interviews](#) conducted:
 - “its different, not necessarily harder”
 - “easier to read than write” ?!



Code samples I

- Idea of what Z candidate selection can look like
- Python allows very flexible interface, under-the-hood data structure is columnar

```
ele = electrons[(electrons.p4.pt > 20) &
                 (np.abs(electrons.p4.eta) < 2.5) &
                 (electrons.cutBased >= 4)]

mu = muons[(muons.p4.pt > 20) &
            (np.abs(muons.p4.eta) < 2.4) &
            (muons.tightId > 0)]
```

- Selects good candidates (per-entry selection)

```
ee = ele.distincts()
mm = mu.distincts()
em = ele.cross(mu)
```

- Creates pair combinatorics (creates new pairs array, also jagged)

```
channels['ee'] = good_trigger & (ee.counts == 1) & (mu.counts == 0)
channels['mm'] = good_trigger & (mm.counts == 1) & (ele.counts == 0)
channels['em'] = good_trigger & (em.counts == 1) & (ele.counts == 1) & (mu.counts == 1)
```

- Selects good events, partitioning by type (per-event selection)

```
dileptons['ee'] = ee[(ee.i0.pdgId*ee.i1.pdgId == -11*11) & (ee.i0.p4.pt > 25)]
dileptons['mm'] = mm[(mm.i0.pdgId*mm.i1.pdgId == -13*13)]
dileptons['em'] = em[(em.i0.pdgId*em.i1.pdgId == -11*13)]
```

- Selects good pairs, partitioning by type (per-entry selection on pairs array)

Code samples II

- Enable expressive abstractions without python interpreter overhead
 - e.g. storing boolean event selections from systematic-shifted variables in named bitmasks: each add() line operates on O(100k) events

```
shiftSystematics = ['JESUp', 'JESDown', 'JERUp', 'JERDown']
shiftedQuantities = {'AK8Puppijet0_pt', 'pfmet'}
shiftedSelections = {'jetKinematics', 'jetKinematicsMuonCR', 'pfmet'}
for syst in shiftSystematics:
    selection.add('jetKinematics'+syst, df['AK8Puppijet0_pt_'+syst] > 450)
    selection.add('jetKinematicsMuonCR'+syst, df['AK8Puppijet0_pt_'+syst] > 400.)
    selection.add('pfmet'+syst, df['pfmet_'+syst] < 140.)
```

- Columnar analysis is a [lifestyle brand](#)
 - Opens up scientific python ecosystem. e.g. interpolator from 2D ROOT histogram:

```
def centers(edges):
    return (edges[:-1] + edges[1:])/2

h = uproot.open("histo.root")["a2dhisto"]
xedges, yedges = h.edges
xcenters, ycenters = np.meshgrid(centers(xedges), centers(yedges))
points = np.hstack([xcenters.flatten(), ycenters.flatten()])
interp = scipy.interpolate.LinearNDInterpolator(points, h.values.flatten())
x, y = np.array([1,2,3]), np.array([3., 1., 15.])
interp(x, y)
```

- Don't want linear interpolation? Try one of several [other options](#)

Domain of applicability

- Domain of applicability depends on:
 - Complexity of algorithms
 - Size of per-event input state
- Examples:
 - JEC (binned parametric function): use binary search, masked evaluation: **columnar ok**
 - Object gen-matching, cross-cleaning: $\min(\text{metric}(\text{pairs of offsets}))$: **columnar ok**
 - Deterministic annealing PV reconstruction: large input state, iterative: **probably not**
- How far back can columnar go?
 - *Missing array programming primitives not a barrier, can always implement our own*

Event loop

Columnar

Event Reconstruction

1 MB/evt

Complex algorithms
operating on large per-
event input state

Inter-event SIMD

Analysis Objects

40-400 kB/evt

Fewer complex
algorithms, smaller per-
event input state

Filtering & Projection

(skimming & slimming)
1 kB/evt

Few complex
algorithms, $O(1 \text{ column})$
input state

Empirical PDFs

(histograms)
No event scaling

Trivial operations

Scalability

- Present a unified data structure to analysis function or class
 - Dataframe of awkward arrays
 - Decouple data delivery system from analysis system
- We can run real-world analyses at a range of scales
 - With home-grown and commercial scheduler software
- Lessons learned so far:
 - Fast time-to-backtrace as important as time-to-insight, keep in mind for analysis facilities!
 - Physics-driven bookkeeping (dataset names, cross sections, storage of derived data, etc.) is nontrivial in all cases, *needs to be decoupled*
 - Inherently higher memory footprint, solved by adjusting partitioning (chunking) scheme
 - Tradeoff with data delivery overhead

<i>Data delivery system</i>	<i>Z peak wall-time throughput</i>	<i>Subjective ‘ease of use’</i>
<i>uproot on laptop</i>	<i>~ 100 kHz</i>	<i>5/5</i>
<i>uproot + xrootd + multiprocessing</i>	<i>~ 250 kHz @ 10 cores *</i>	<i>5/5</i>
<i>uproot + condor jobs</i>	<i>Arbitrary</i>	<i>3/5</i>
<i>striped system</i>	<i>~ 10 MHz @ 100 cores</i>	<i>2/5</i>
<i>Apache spark</i>	<i>~ 1 MHz @ 100 cores **</i>	<i>4/5</i>

Part II: Technical Underpinnings

Theoretical Motivation

- Aligned with strengths of modern CPUs
 - Simple instruction kernels aid pipelining, branch prediction, and pre-fetching
 - Event loop = input data controlling instruction pointer = less likely to exploit all three!
 - *Unnecessary work is cheaper than unusable work*
- Inherently SIMD-friendly
 - Event loop cannot leverage SIMD unless inter-event data sufficiently large
- In-memory data structure *exactly* matches on-disk serialized format
 - Event loop must transform data structure - significant overhead
 - Memory consumption managed by chunking (event groups, or baskets)
- Array programming kernels form computation graph
 - Could allow query planning, automated caching, non-trivial parallelization schemes

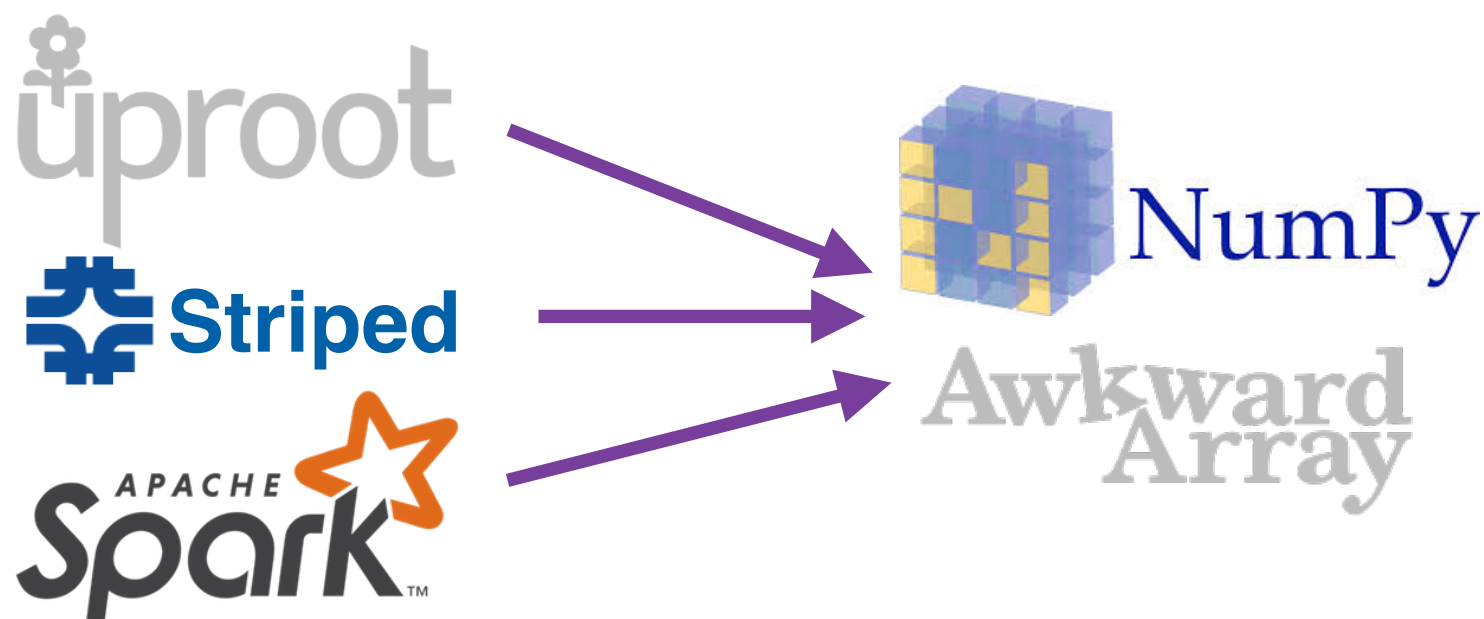
The Coffea framework

- Column Object Framework For Effective Analysis:
 - Prototype analysis framework utilizing columnar approach
 - Provides object-class-style view of underlying arrays
 - Implements typical recipes needed to operate on NANOAOB-like nTuples
 - One monolith for now: [fnal-column-analysis-tools](https://github.com/Scirra/fnal-column-analysis-tools)
 - Functionality will be factorized into targeted packages as it matures
- Realized using scientific python ecosystem
 - numpy: general-purpose array manipulation library
 - numba: uses llvm to JIT-compile python code, understands numpy
 - Work ongoing to extend to awkward arrays as well
 - scipy: large library of specialized functions
 - cloudpickle: serialize arbitrary python objects, even function signatures
 - matplotlib: python visualization library

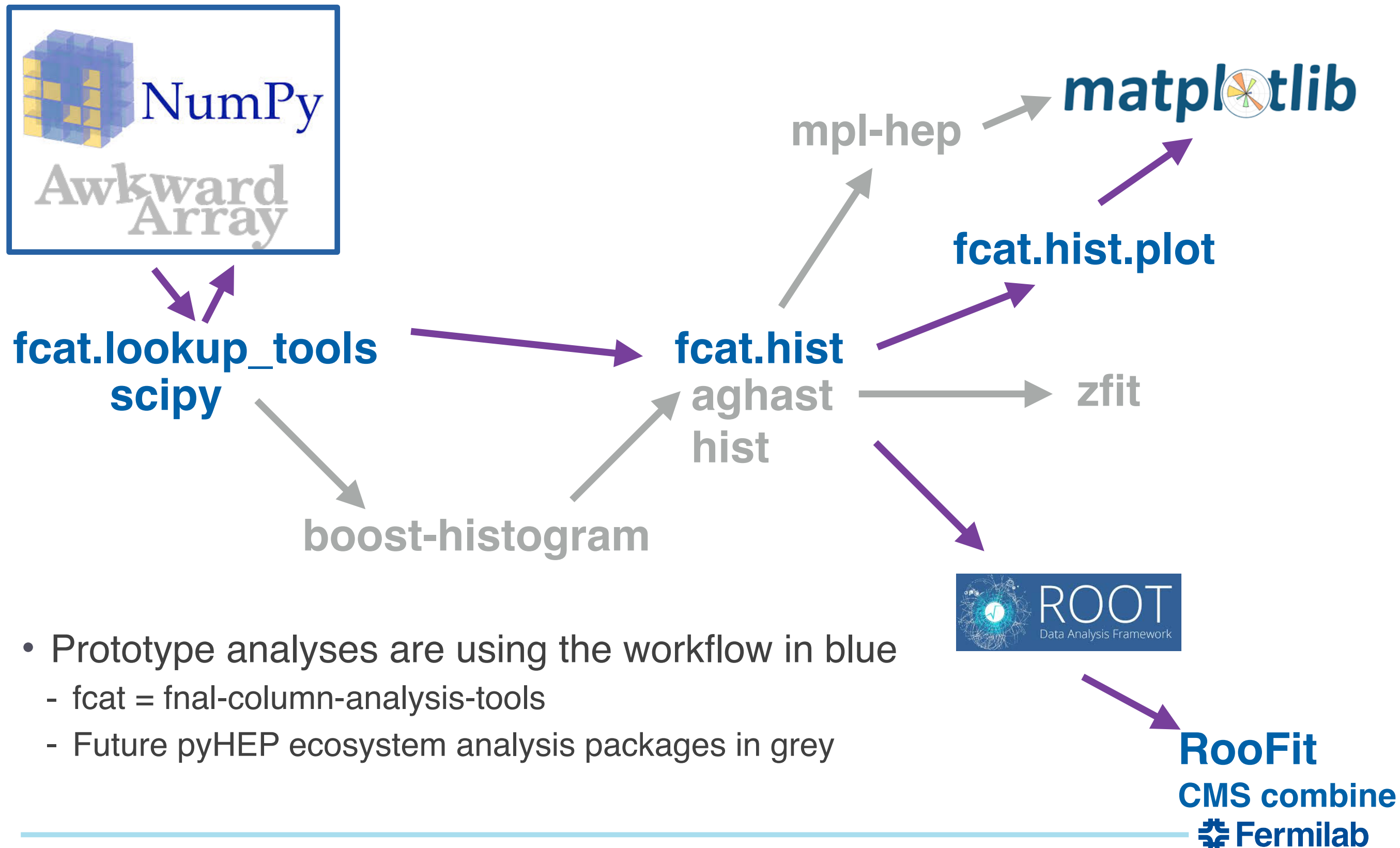


Factorized Data Delivery

- [Uproot](#)
 - Direct conversion from TTree to numpy arrays and/or awkward JaggedArrays
- [Striped](#)
 - NoSQL database delivers 'stripes': numpy arrays
 - Re-assemble awkward structure via object counts + content
 - memcached layer, python job scheduler, ~150 core cluster
 - Derived columns persistable
- [Spark](#)
 - Interface using vectorized UDF (user-defined function)
 - Currently restricted to intermediate pandas format (pyarrow UDF to be implemented)
 - Derived columns persistable



Package ecosystem



Performance

- Z peak benchmark

- Includes many typical corrections: lumimask, PU correction, ID scale factors, flavor-categorized
- 350 lines jupyter notebook, 25 columns accessed
- 6 μ s/evt/thread (125 kHz) wall time
 - ROOT C++ TBranch::GetEntry(): ~1.5x faster

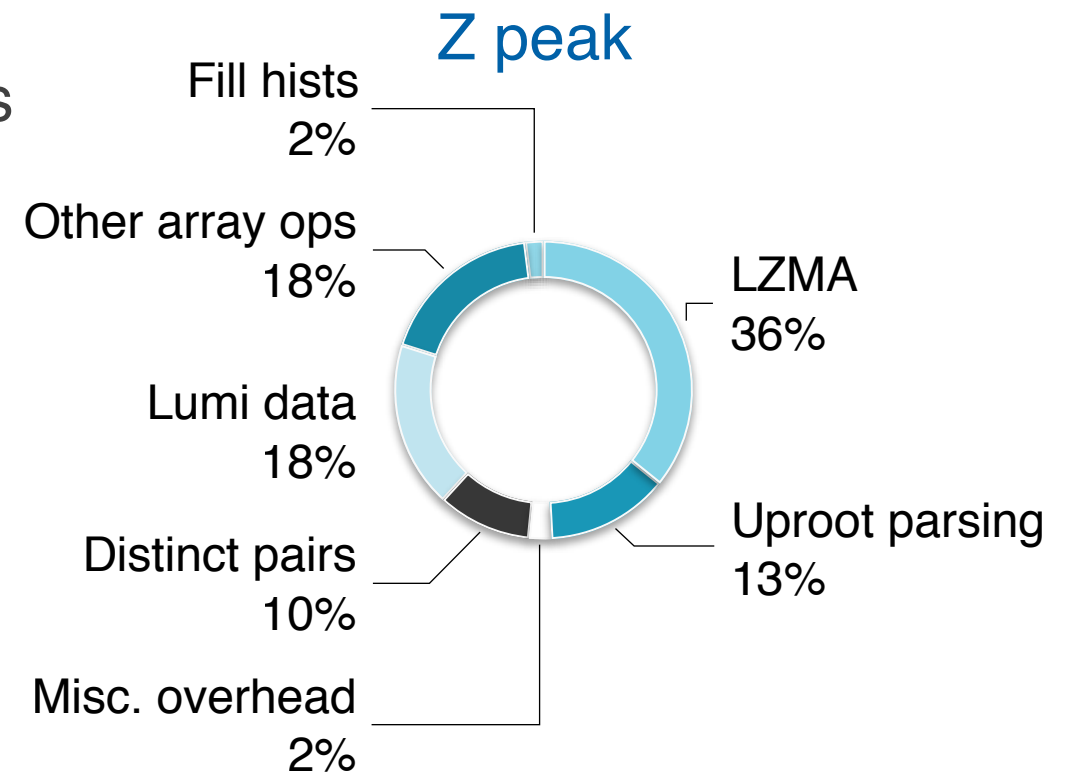
- Two prototype analyses

- “end-to-end” = NanoAOD-like nTuple to templates
- Varies from 30-150 μ s/evt/thread
- Already being used to steer analysis, present results in analysis group meetings

- Many inefficiencies known

- Can be removed with further development in awkward and helper libraries

```
def run(self, events, job):
    times = OrderedDict()
    times['start'] = time.time()
    if self.weights_eval is None:
        self.weights_eval = cloudpickle.loads(atlib.decompress(job['weights_eval']))
    if self.lumimask is None:
        self.lumimask = cloudpickle.loads(atlib.decompress(job['lumimask']))
    dataset = job['dataset']
    hints = job['hints']
    isLocalData = False
    good_trigger = np.zeros(events.nevents, dtype=bool)
    if dataset in self.triggers:
        isLocalData = True
        # local data, prevent overlaps
        for trigger in self.triggers[dataset]:
            good_trigger |= getattr(events, trigger)
        # apply lumimask post certified lumimask
        good_trigger &= lumimask(events.run, events.luminosityBlock)
        # record lumi processed
        # is uproot job, full lumi completion guaranteed by full-file processing
        # is striped job, partial lumi completion is not traceable at present
        if dataset == "Doublet":
            hints['lumi_sum'] = LumList(events.run, events.luminosityBlock)
        elif dataset == "Doublet":
            hints['lumi_sum'] = LumList(events.run, events.luminosityBlock)
        else:
            # MC, OR all
            for trigger in self.triggers.values():
                for trigger in hints:
                    good_trigger |= getattr(events, trigger)
            times['lumimask_trigger'] = time.time()
            genw = np.sign(events.LHEWeight_original*0.000707)
            electrons = PhysicalColumnGroup(events, "Electron", [(k, k for k in self.electron_cols)])
            electrons = jaggedFromColumnGroup(electrons)
            electrons['weight'] = weights_eval['electronFactor_weight_pt0']*(electrons.pt.eta, electrons.pt.pt)
            muons = PhysicalColumnGroup(events, "Muon", [(k, k for k in self.muon_cols)])
            muons = jaggedFromColumnGroup(muons)
            muons['weight'] = weights_eval['muonFactor_weight_pt0']*(muons.pt.eta, muons.pt.pt)
            e = electrons[electrons.pt > 20] & (np.abs(electrons.pt.eta) < 2.5) & (electrons.cutBased == 4)
            mu = muons[muons.pt > 20] & (np.abs(muons.pt.eta) < 2.4) & (muons.cutBased == 0)
            times['good leptons'] = time.time()
            em = e.discriminate()
            em = mu.discriminate()
            em = e.cross(mu)
            dileptons = {}
            dileptons['em'] = em(em.i0.pdgId==em.i1.pdgId == -11*11) & (em.i0.pt > 25)
            dileptons['mm'] = em(em.i0.pdgId==em.i1.pdgId == -13*13)
            dileptons['em'] = em(em.i0.pdgId==em.i1.pdgId == -11*13)
            times['good pairs'] = time.time()
            channels = {}
            channels['ee'] = good_trigger & (em.counts == 1) & (mu.counts == 0)
            channels['mm'] = good_trigger & (em.counts == 1) & (mu.counts == 0)
            channels['em'] = good_trigger & (em.counts == 1) & (mu.counts == 1)
            times['channels'] = time.time()
            dups = np.zeros(events.nevents, dtype=bool)
            tot = 0
            for channel, cut in channels.items():
                reands = dileptons[channel][cut]
                dups |= cut
                tot += cut.sum()
            if not isLocalData:
                weight = np.array(1)
                weight = reands.i0['weight'] * reands.i1['weight'] * genw[cut]
                hints['genw'].fill(dataset=dataset, genw=genw)
                hints['reands_pt'].fill(dataset=dataset, channel=channel, reands_pt=reands.i0.pt.flatten(), reands_pt=reands.i1.pt.flatten(), weight=weight.flatten())
            hints['xMass'].fill(dataset=dataset, channel=channel, mass=reands.pt.mass.flatten(), weight=weight.flatten())
            if dups.sum() != tot:
                raise Exception("Double-counting events!")
            times['plots'] = time.time()
            #profiling info
            t = list(times.values())
            for i, name in enumerate(times):
                if i==0: continue
                dt = t[i] - t[i-1]
                hints['profile'].fill(op=name, dt=dt*(dt/(len(events.run.count)))
            hints['profile'].fill(op='total', dt=dt*(dt/(len(events.run.count)))
            job.send(hints)
```



Future Directions

- As Coffea (& underlying libraries) matures, invite beta testers
 - I encourage everyone to try uproot+numpy now
- Target first release this summer
 - Two full analysis implemented
 - Data delivery mechanisms fully separated
 - User interface improvements and documentation
- Far future: analysis facility
 - This feeds towards the dream of a “short time-to-insight” “analysis as a service” facility
 - Tendering bids for additional buzzwords
 - Array programming allows easier construction of computation graphs
 - Query planning can detect common patterns and execute them once
 - By removing manual cache management, we can optimize throughput and storage
- First, lets see if we are happy and productive with the columnar approach
 - So far, the answer appears to be yes