



(US) CMS Analysis Facilities

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Analysis Facilities are not new



CAF specs

The intention of the CAF specs from 1997 seem to have been to define central analysis facilities that would provide a level of "user satisfaction" similar to Run I.

**JAN
27
2004**

The CMS User Analysis Farm (UAF)

- [Introduction](#)
- [News](#)
- [How to get access to the cluster?](#)

Analysis Facilities are not old



The Evolution of HEP Analysis Facilities

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V0, 8/31/2020

Analysis Facilities for Late-Stage Analysis

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Snowmass Letter of Interest - Analysis Facilities - CompF5

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Why evolve Analysis Facilities?

- Traditionally in HEP we have done **row-wise** analysis
 - Process one event at a time serially
 - Scripts and batch jobs (and now also notebooks)
 - We need to continue to support this
- Trending towards **column-wise** (tidy/big data) analyses
 - **Low-latency** access to CPU resources
 - Fast access to a significant amount of **disk**
 - Large investment from industry
- Low latency implies $<100\%$ resource utilization
 - **Coscheduling** analysis with other computation is appealing

Core elements for analysis facilities

- As a foundation layer, use a **containerized infrastructure** (Kubernetes-based).
- A **Jupyterhub** deployment as the glass
- Support “typical” (**row-wise**) analysis needs
- Support columnar analysis low latency application (e.g. Apache Spark, **Dask**, ...) and frameworks (**Coffea**)
- Deploy **analysis services** (ServiceX, Skyhook, pyhf, ...)
- Packaging and redeployment
 - e.g. Demonstrate running FNAL AF at a US CMS Tier 2

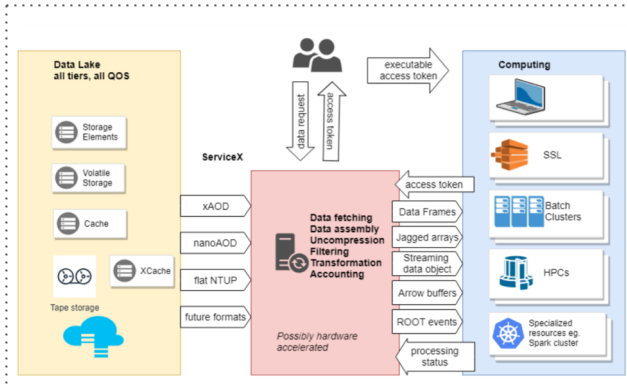
Coffea: a columnar analysis framework for HEP



- Developed by the US CMS research program
 - Driven by Lindsey Gray, Nick Smith @ **Fermilab**
- Provides a **user interface** for columnar analysis with missing pieces of the stack filled in get users from ROOT files to histograms; some of this is transitional
- Components include
 - Lookup tools
 - Experiment-specific corrections in array programming context
 - Histogram tools (API will be *hist* package + mplhep)
 - NanoEvents
 - Wrap “flat TTree” as high-level lazy-access awkward array; enables cross-references, self-references, vector semantics, dynamic attributes, ...
 - Processors
 - Wrap analysis code (utilizing the above) and take care of map-reduce

Analysis Services – a sampling

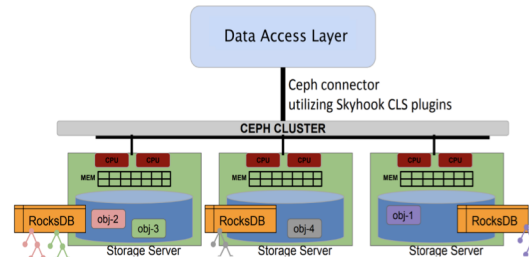
ServiceX



ServiceX provides user level ntuple production

- Converts experiment-specific datasets to columns (e.g. NanoAOD)
- Enable simple cuts or simple derived columns and fields
 - Heavy-weight analysis will still happen via some separate processing toolchain (like CRAB)

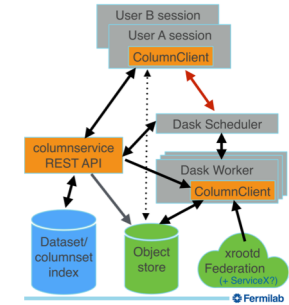
Skyhook DM



SkyhookDM converts ROOT files to columnar format

- Ceph-side C++ plugins transition from on-disk format to desired memory format
- Uses Dask workers to distribute data to clients
- Data delivered as Arrow tables or via Arrow Dataset API

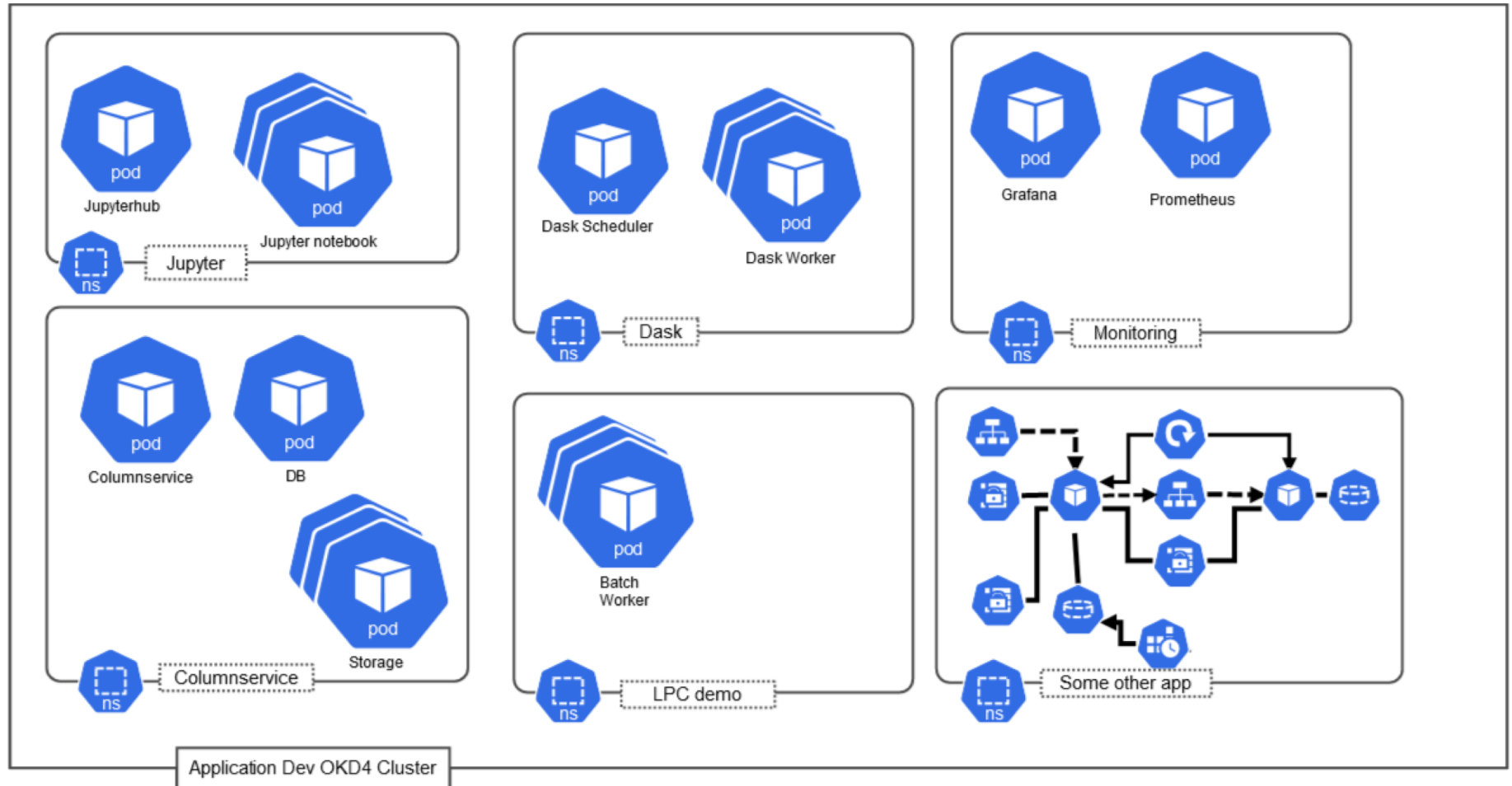
Columnservice



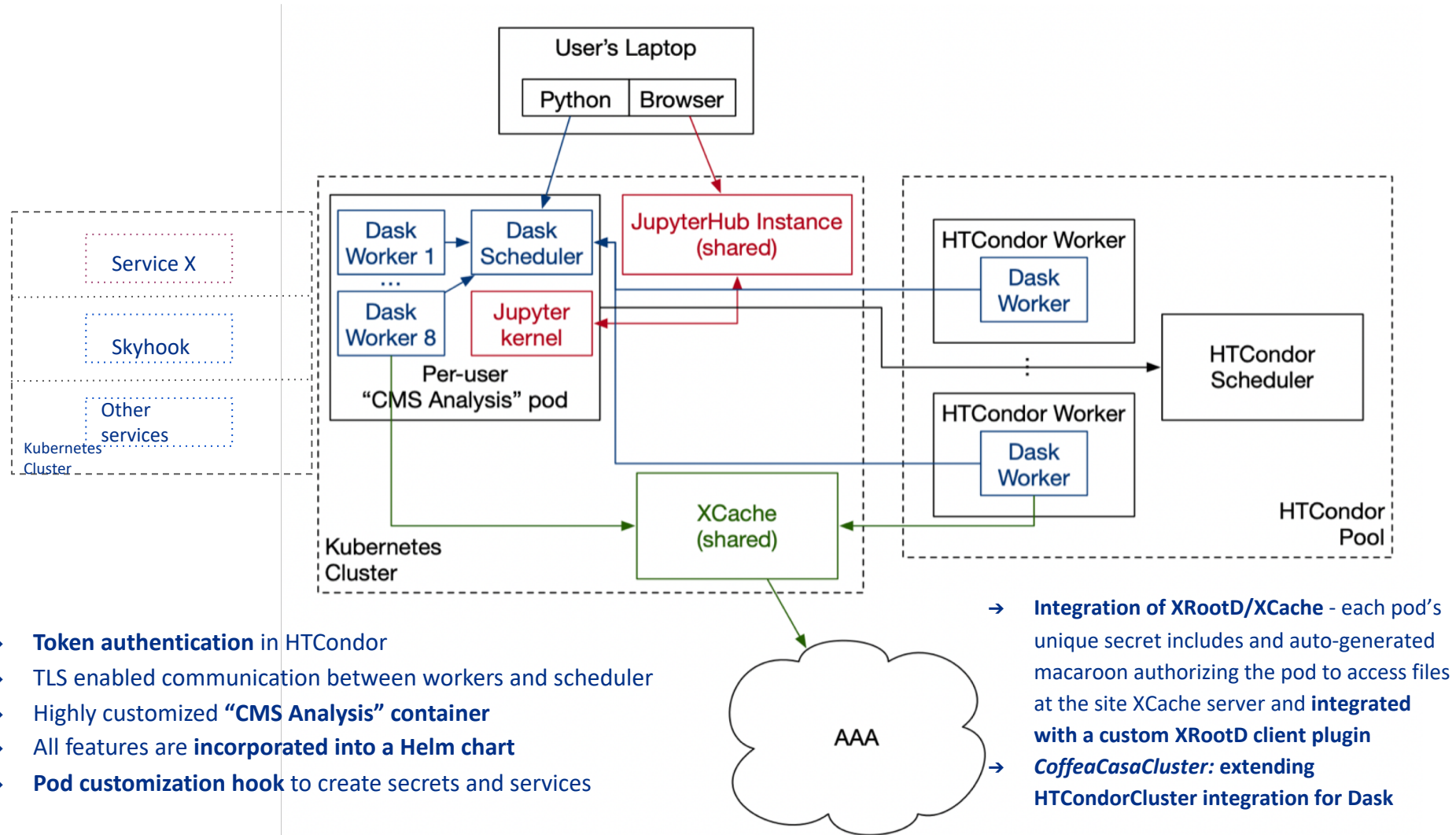
Columnservice caches columnar data for multiple users

- Shared input cache at column granularity
- Derived columns only constructed and cached on access
- Unified metadata and dataset schema database

Fermilab Analysis Facility: pre-prototype

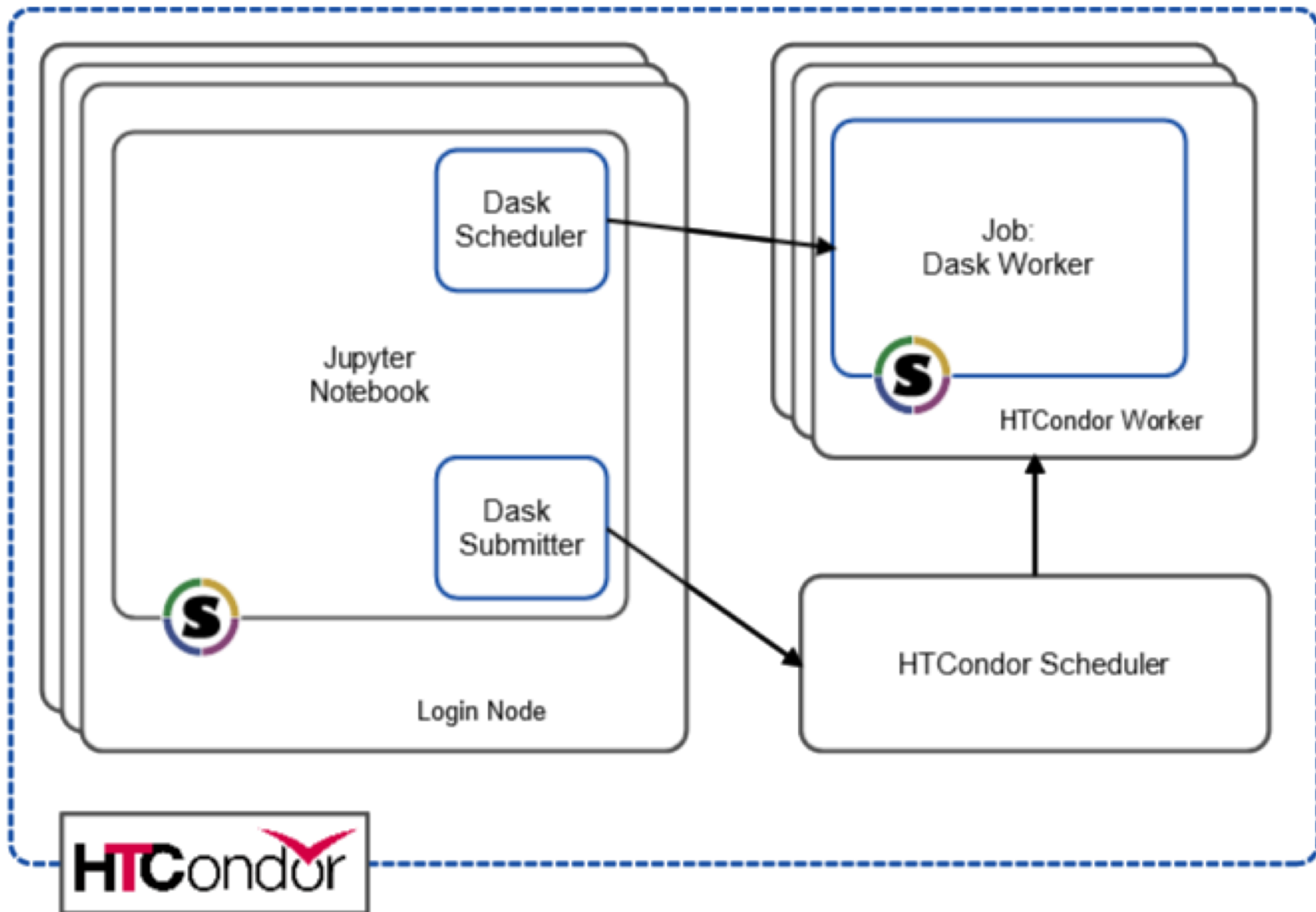


Analysis Facility “Coffea-casa” @ UNL



- **Integration of XRootD/XCache** - each pod's unique secret includes and auto-generated macaroon authorizing the pod to access files at the site XCache server and **integrated with a custom XRootD client plugin**
- **CoffeaCasaCluster: extending HTCondorCluster integration for Dask**

Fermilab interim solution – Dask on HTCondor



Fermilab Analysis Facility plans – adding elasticity

