

WORKFLOW MANAGEMENT

Thomas Uram
Argonne Leadership Computing Facility

WHY USE WORKFLOWS?

- Automate job submission
- Simplify computational campaigns
- Increase concurrency by disentangling data dependencies
- Robustness: Improve error handling and recovery (retries)
- Coscheduling of multiple resources
- Systematize data management
- Provenance/Metadata tracking
 - Validation
 - Reuse

WHAT IS A WORKFLOW?

- It depends who you ask
- Basically a collection of jobs to be run
 - Could be a sequence of individual jobs
 - Could be a sequence of varying numbers of jobs
- Available means of describing and running workflows
 - Script jobs
 - Job dependencies
 - Ensemble jobs
 - In situ
 - Custom workflows
 - Workflow software

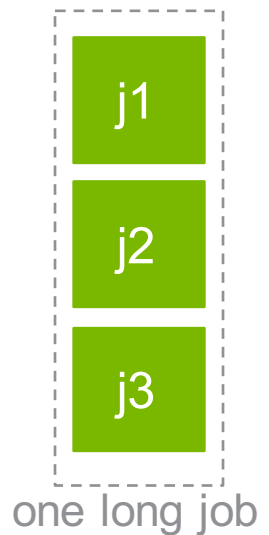
WORKFLOW VIA SCRIPT JOB

- You can submit a job that executes your application
 - `qsub -q prod -n 512 -t 10 -A yourproject application.exe`
- Alternatively, you can submit a script job that executes multiple applications sequentially
 - `qsub -q prod -n 512 -t 10 -A yourproject script.sh`

script.sh

```
runjob application.exe  
runjob application.exe
```

- With this approach, you wait in the queue one time to run your application multiple times
- However, small long jobs tend to stay in the queue longer than small short jobs
 - It may be better to submit individual jobs with dependencies



WORKFLOW VIA COBALT JOB DEPENDENCIES

- A simple way to achieve a linear workflow is simply to set dependencies between your jobs
 - `qsub -q prod -n 512 -t 10 -A yourproject a.out`
Job 12345 submitted
 - `qsub -q prod -n 512 -t 10 -A yourproject --dependencies 12345 a.out`
 - `qsub -q prod -n 512 -t 10 -A yourproject --dependencies 12345:12346 a.out`

- Is there an advantage to setting job dependencies?
 - Dependent jobs accumulate score more quickly
- How many jobs can be submitted at once?

multiple
independent
short
jobs

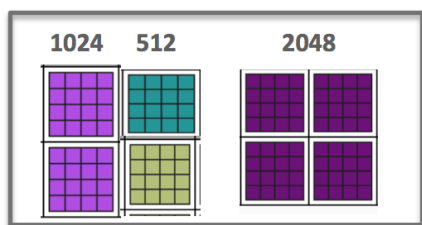
```
qstat -Q
```

Name	Users	Groups	MinTime	MaxTime	MaxRunning	MaxQueued	MaxUserNodes	MaxNodesPerNode
default	None	None	00:05:00	01:00:00	5	20	2048	1024

- On Mira, max queued is set to 20

WORKFLOW VIA ENSEMBLE JOBS: RUN BIGGER JOBS

Example of ensemble jobs



4K

	R00	R01	R02	R03	R04	R05	R06	R07	R08	R09	R0A	R0B	R0C	R0D	R0E	R0F
M1																
M0																
	R10	R11	R12	R13	R14	R15	R16	R17	R18	R19	R1A	R1B	R1C	R1D	R1E	R1F
M1																
M0																
	R20	R21	R22	R23	R24	R25	R26	R27	R28	R29	R2A	R2B	R2C	R2D	R2E	R2F
M1																
M0																

Minimum
partition size
on Mira

WORKFLOW VIA ENSEMBLE JOBS: RUN BIGGER JOBS

<http://trac.mcs.anl.gov/projects/cobalt/wiki/BGQUserComputeBlockControl>

```
#!/bin/bash
BLOCKS=`get-bootable-blocks --size 512 $COBALT_PARTNAME`

for BLOCK in $BLOCKS
do
    boot-block --block $BLOCK &
done
wait

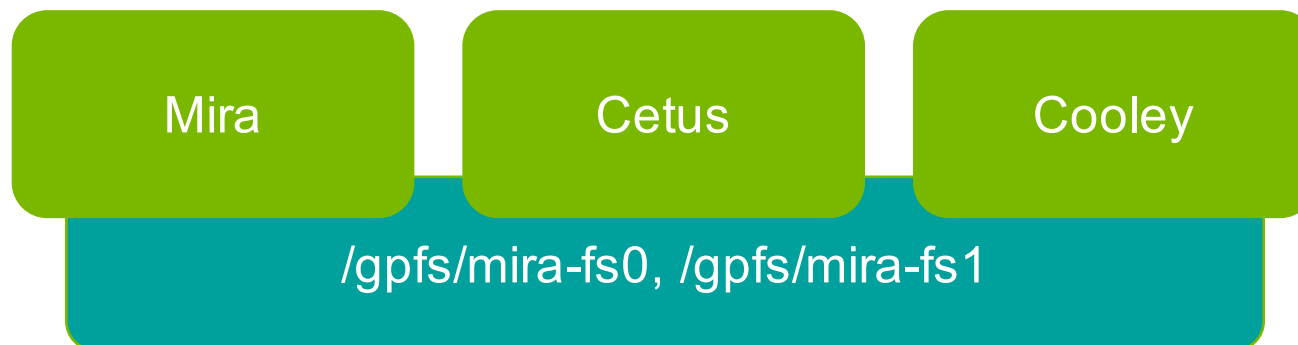
for BLOCK in $BLOCKS
do
    runjob --block $BLOCK : ./my_binary &
done
wait

for BLOCK in $BLOCKS
do
    boot-block --block $BLOCK --free &
done
wait
```

See Paul Rich's talk from
Monday re: Ensemble Jobs

WORKFLOW ACROSS ALCF SYSTEMS

- How can I set job dependencies across multiple resources (e.g. Mira and Cooley)?
 - ALCF does not provide a means of doing this currently
 - However, Cetus and Cooley mount the Mira filesystems. Analysis jobs run on Cooley without needing to transfer the data.

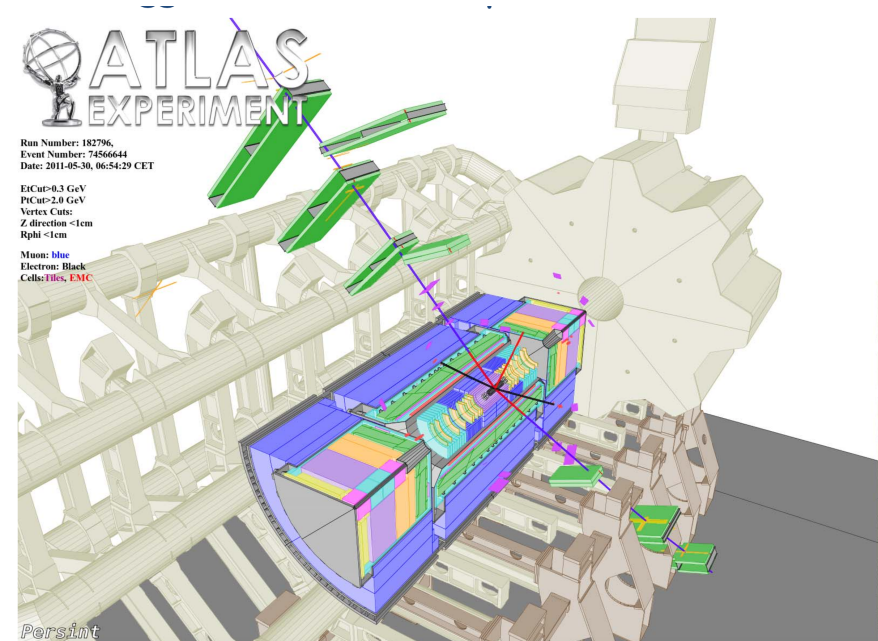


EXAMPLE: HEP EVENT GENERATION

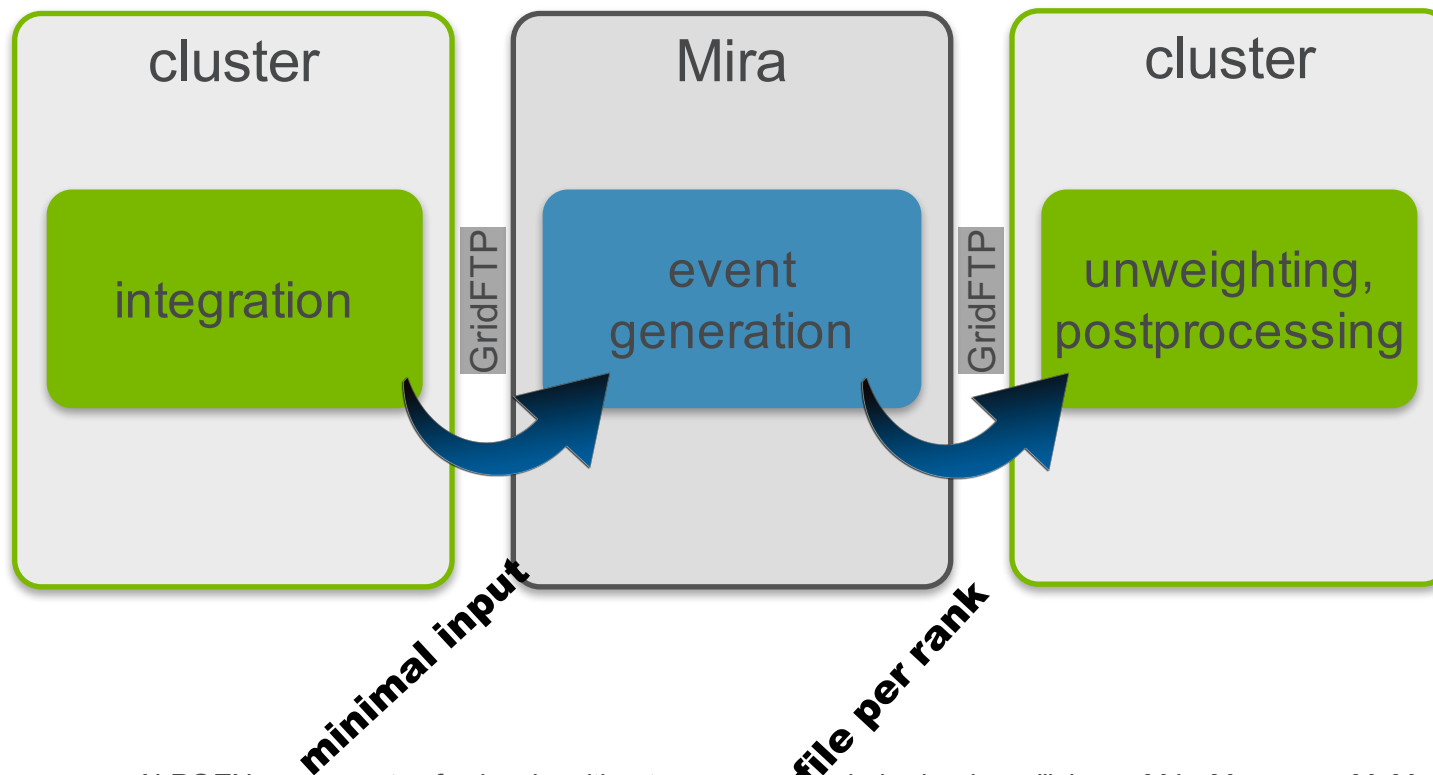
Monte Carlo-based generation of particle collision events such as occur in the ATLAS detector at the Large Hadron Collider, using Alpgen.

Consists of three stages:

1. Generation of phase-space integration grid
2. Generation of weighted events
3. Unweighting of events



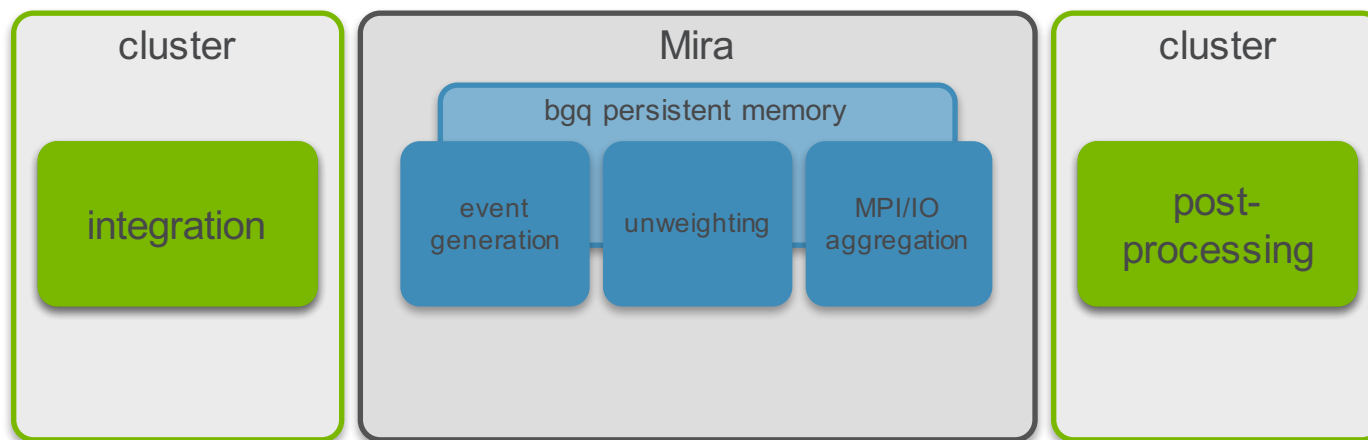
EXAMPLE: HEP EVENT GENERATION



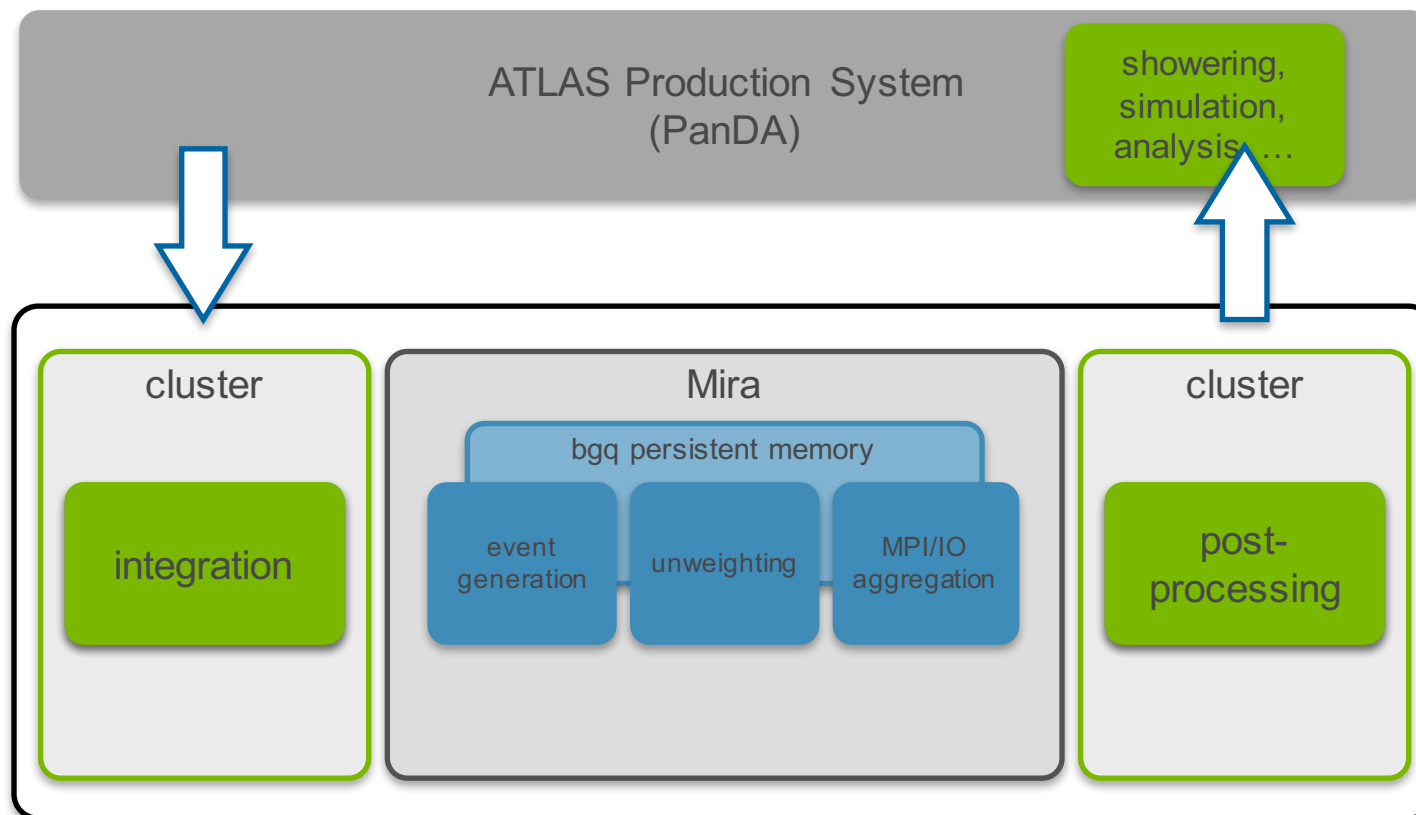
ALPGEN, a generator for hard multiparton processes in hadronic collisions, M.L. Mangano, M. Moretti, F. Piccinini, R. Pittau, A. Polosa, JHEP 0307:001,2003

EXAMPLE: HEP EVENT GENERATION

- Combine multiple application invocations in single script job
- Use persistent memory for exchanging data between invocations
- Aggregate data from persistent memory to filesystem



EXAMPLE: HEP EVENT GENERATION



COMPUTE-NODE PERSISTENT MEMORY

- Environment variable BG_PERSISTMEMSIZE sets MB of memory to use as ramdisk, mounted at /dev/persist
- Persistent memory persists for lifetime of (qsub) job
 - Specify BG_PERSISTMEMSIZE to runjob command
 - If persistent memory size changes between runjob executions, memory will be available but it will be cleared
- Caveat: Persistent memory clearly reduces the amount of memory available for the simulation; if you are operating near the limit of node memory, this solution is not for you

For more info on persistent memory, see
<http://www.redbooks.ibm.com/redbooks/pdfs/sg247948.pdf>

IN SITU ANALYSIS WORKFLOW

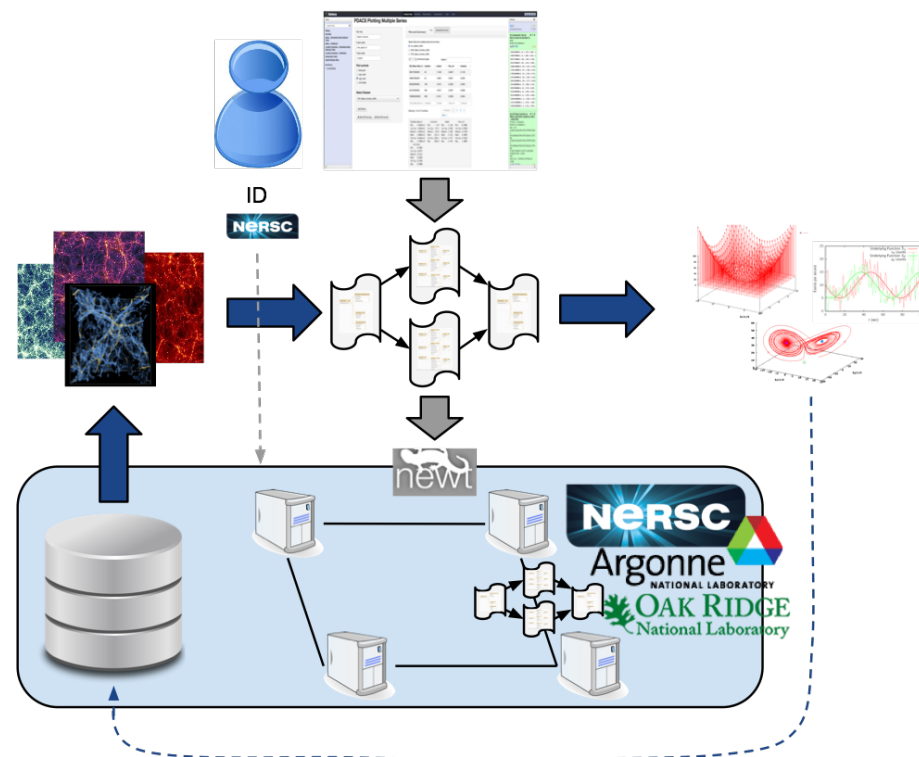
- In situ == running analysis inline with the simulation
- In situ analysis has several advantages
 - Operates on data already in memory; during postprocessing, reading large data from the filesystem can be expensive
 - Is able to perform analysis at the same scale as simulation
 - Operates on full simulation data sets before reduction as opposed to offline analysis on reduced data
 - How to reproduce analysis results based on simulation-time datasets that were not written to disk? Rerun the simulation.

HACC – COSMOTOOLS

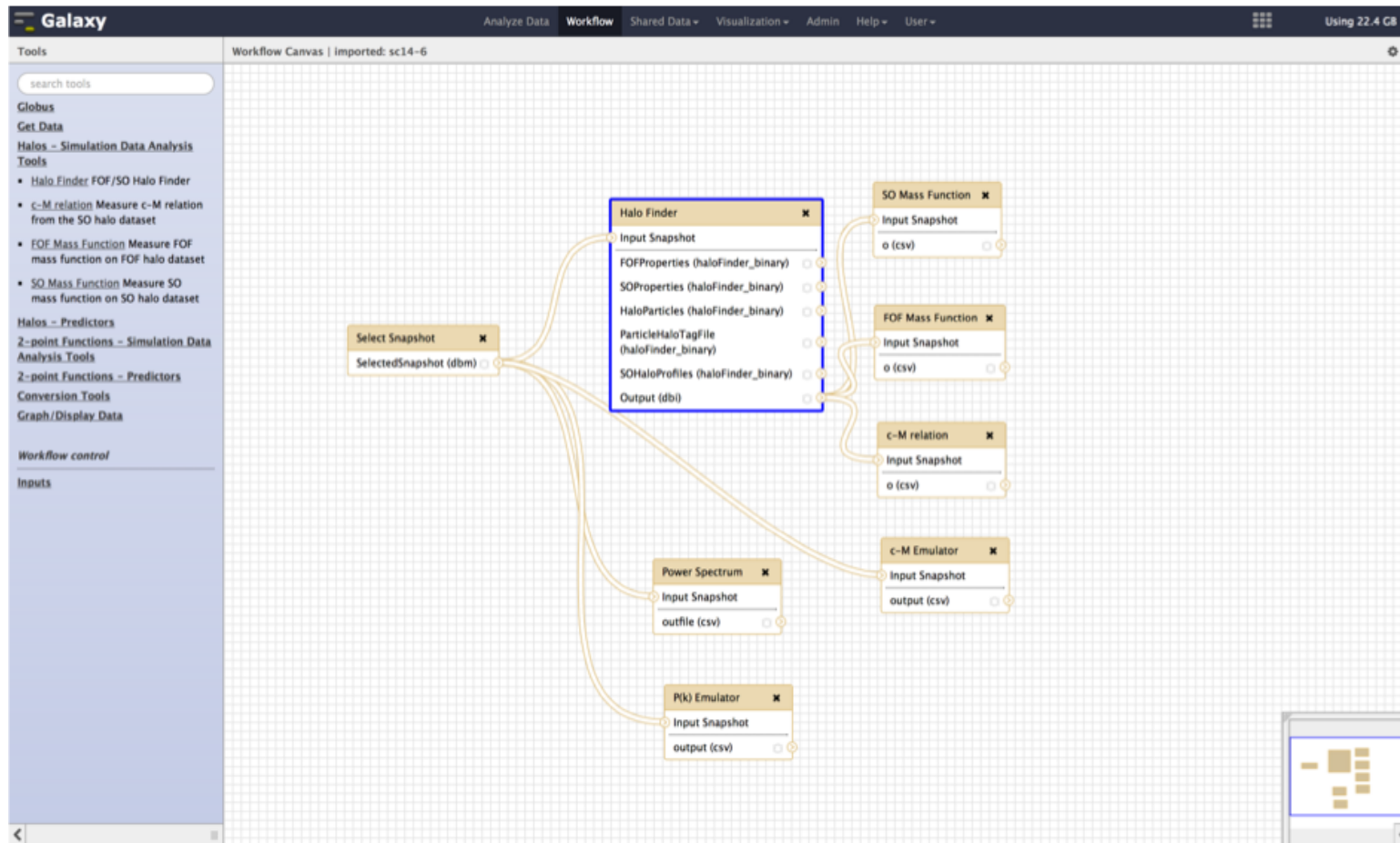
- Cosmotools is a configurable in-situ analysis framework
 - Simulation configuration specifies analyses that should be performed between time steps during the simulation, and their parameters
 - On execution, particle data is passed from the simulation to analysis routines for processing
 - When finished, the simulation proceeds to the next timestep
- Analysis runs prior to data reduction; potentially the only opportunity to analyze this data
- Efficient: avoids the need to subsequently load data from disk
- Delivers intermediate analysis results during (long) simulation runs

HACC – PDACS WORKFLOWS

- Analysis tools wrapped as strongly-typed modules
- Users construct workflows from modules using a graphical tool
- Workflows are executed on resources according to underlying configuration
- PDACS instances running against resources at Argonne, Oak Ridge, and NERSC



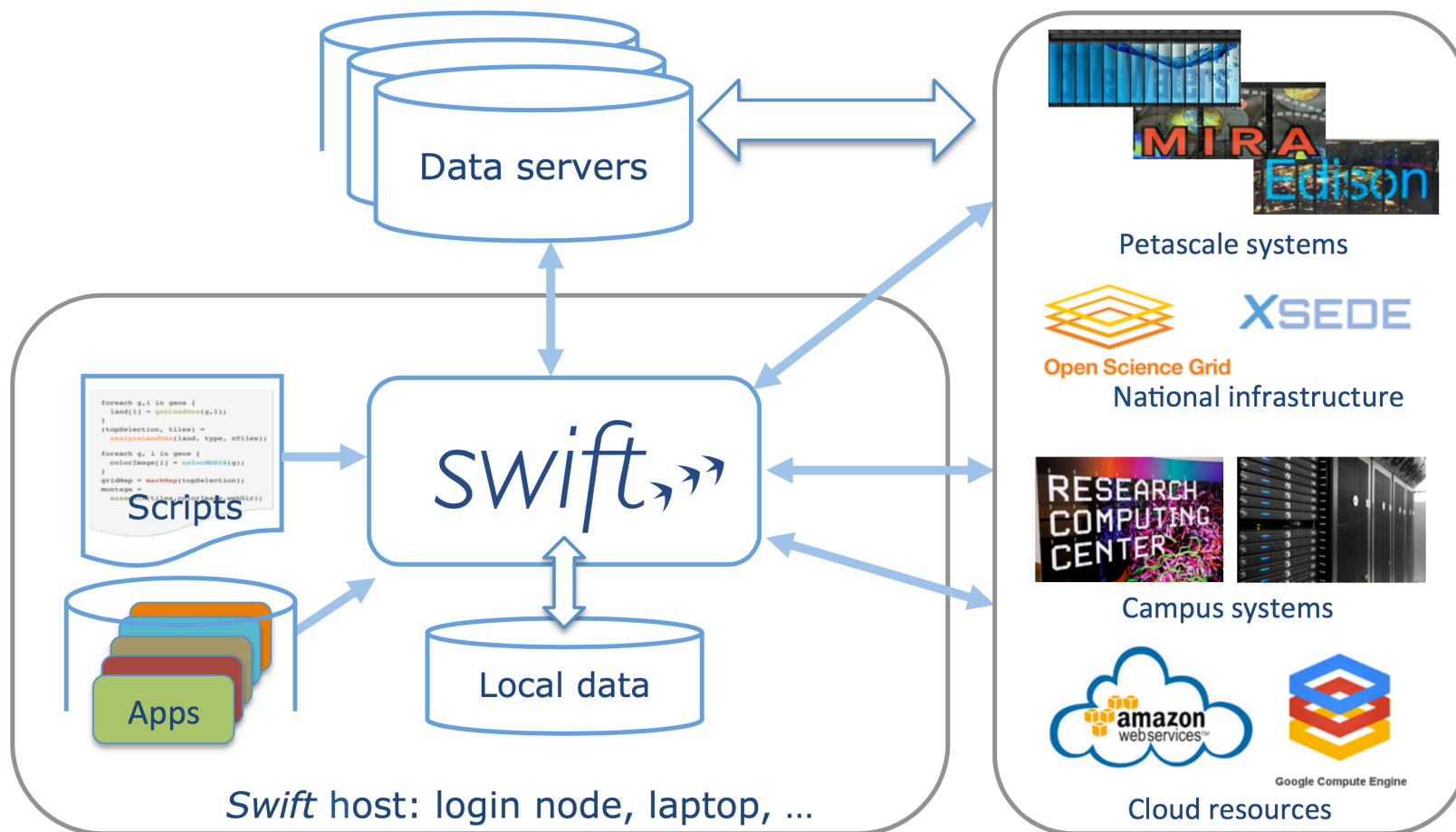
HACC – PDACS WORKFLOWS



WORKFLOW SOFTWARE TO CONSIDER

- Swift (<http://swift-lang.org>)
 - **C-like language** for expressing workflows by wrapping command-line applications or function calls
 - **Execution engine** for translating workflows into job submissions on compute resources, maximizing concurrency by exploiting data (in)dependencies
 - **(Not the Apple product)**
- Fireworks (<https://github.com/materialsproject/fireworks>)
 - Infrastructure for defining and running workflows

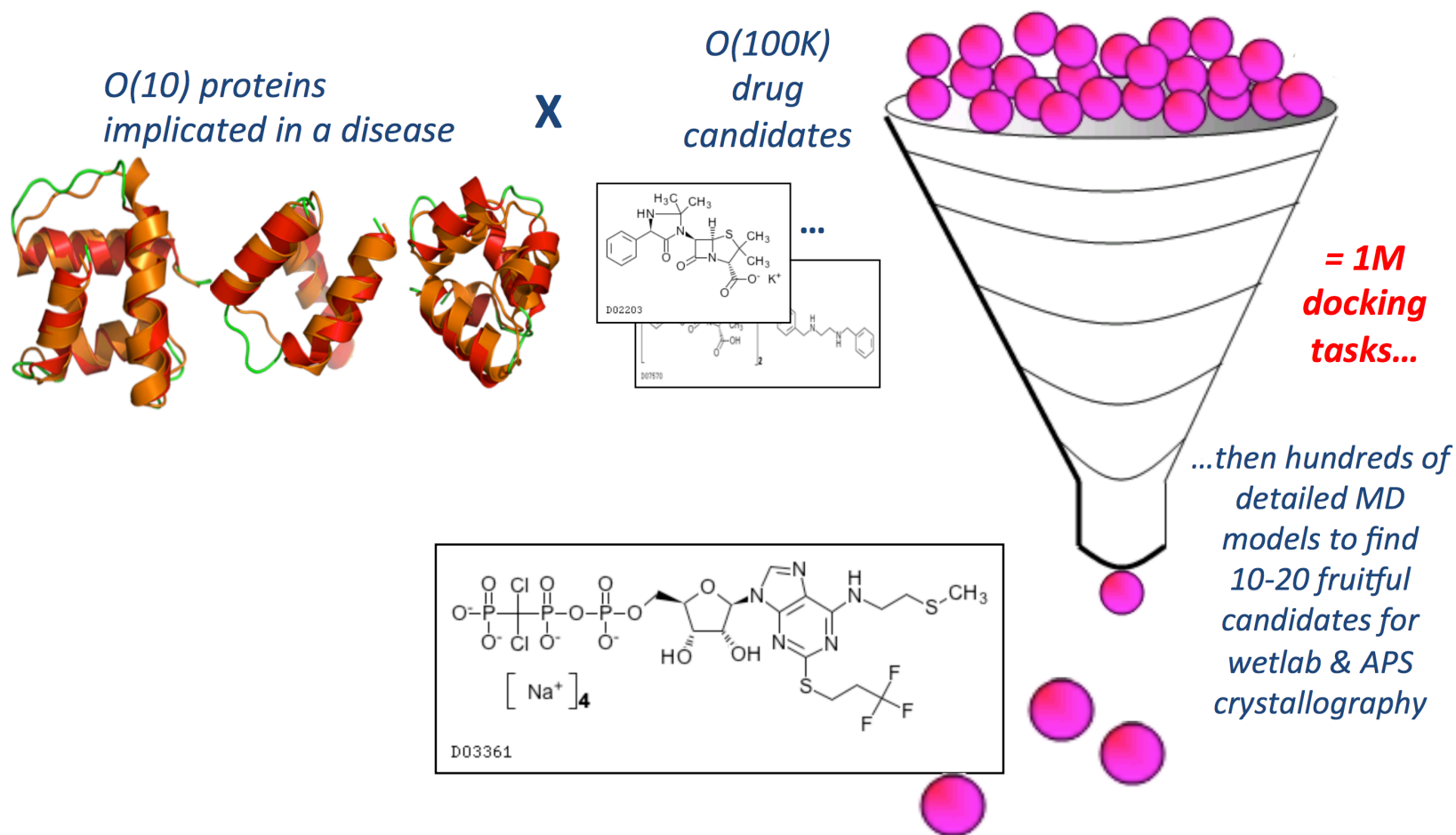
Swift enables execution of simulation campaigns across multiple HPC and cloud resources



See <https://www.alcf.anl.gov/swift>

When do you need HPC workflow?

Example application: protein-ligand docking for drug screening



Expressing this many task workflow in Swift

For protein docking workflow:

```
foreach p, i in proteins {  
    foreach c, j in ligands {  
        (structure[i,j], log[i,j]) =  
            dock(p, c, minRad, maxRad);  
    }  
    scatter_plot = analyze(structure)
```

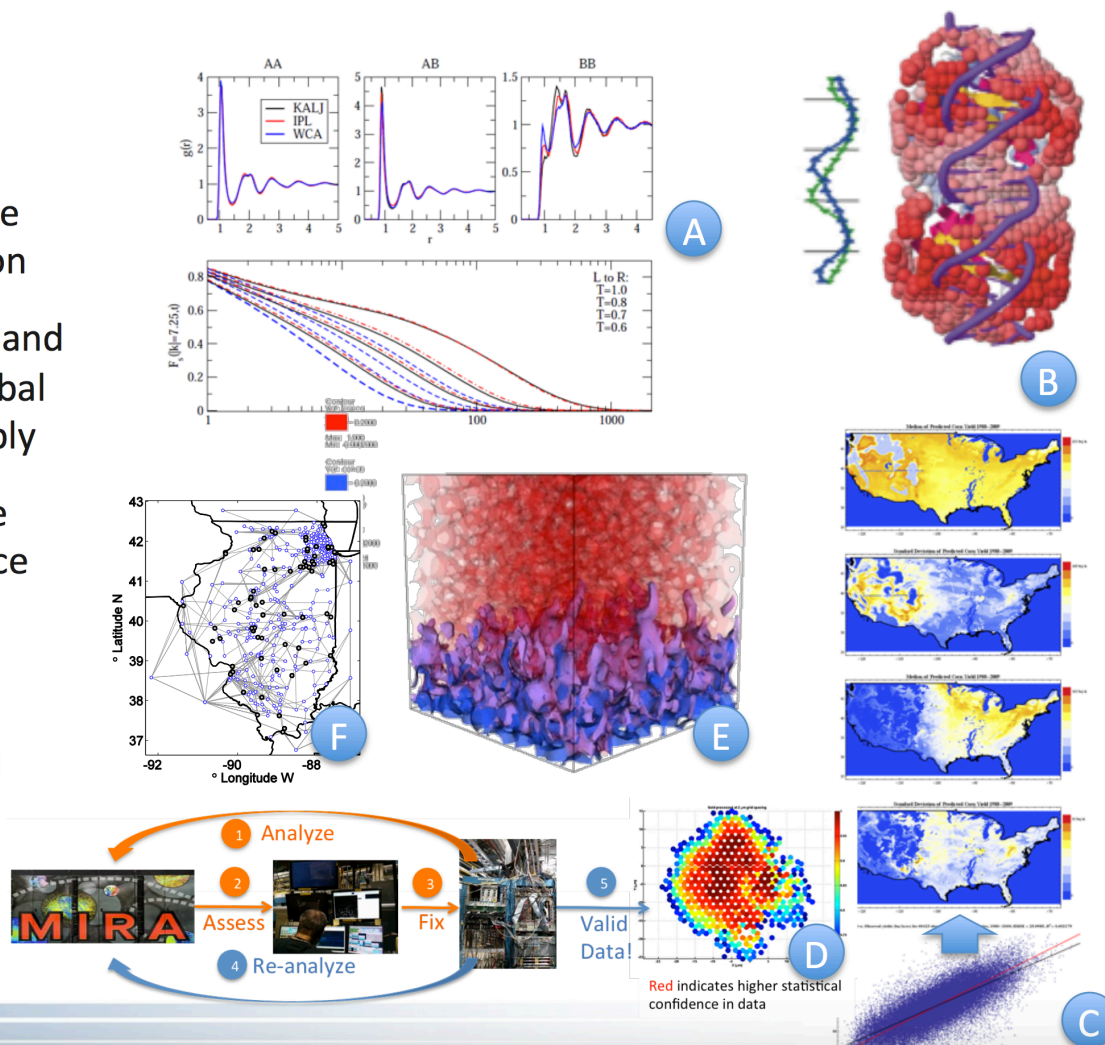
To run:

```
swift -site tukey,blues dock.swift
```

Large-scale applications using Swift

- A** Simulation of super-cooled glass materials
- B** Protein and biomolecule structure and interaction
- C** Climate model analysis and decision making for global food production & supply
- D** Materials science at the Advanced Photon Source
- E** Multiscale subsurface flow modeling
- F** Modeling of power grid for OE applications

All have published science results obtained using Swift



FireWorks v1.3.1

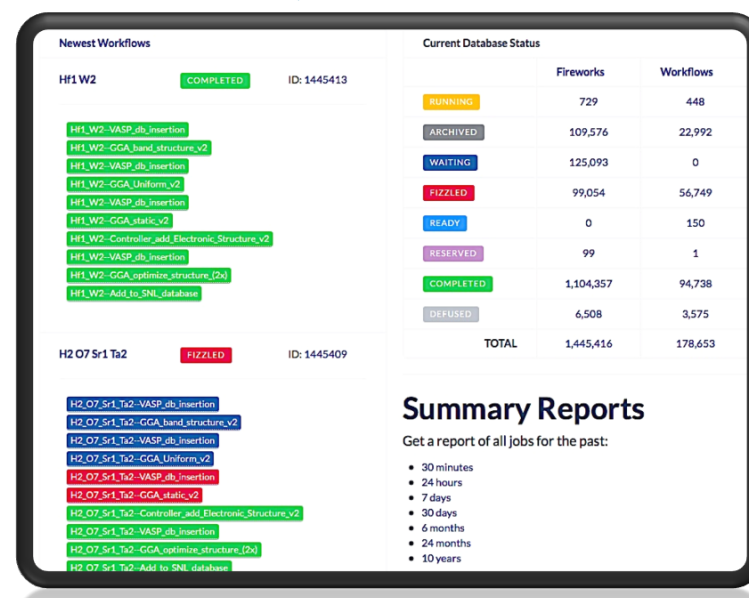
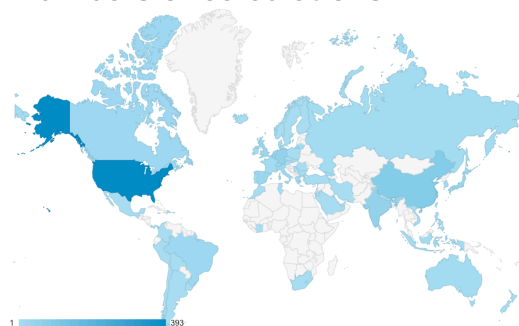
FireWorks is an application-agnostic workflow software for defining and executing large numbers of calculations

Usage and outreach:

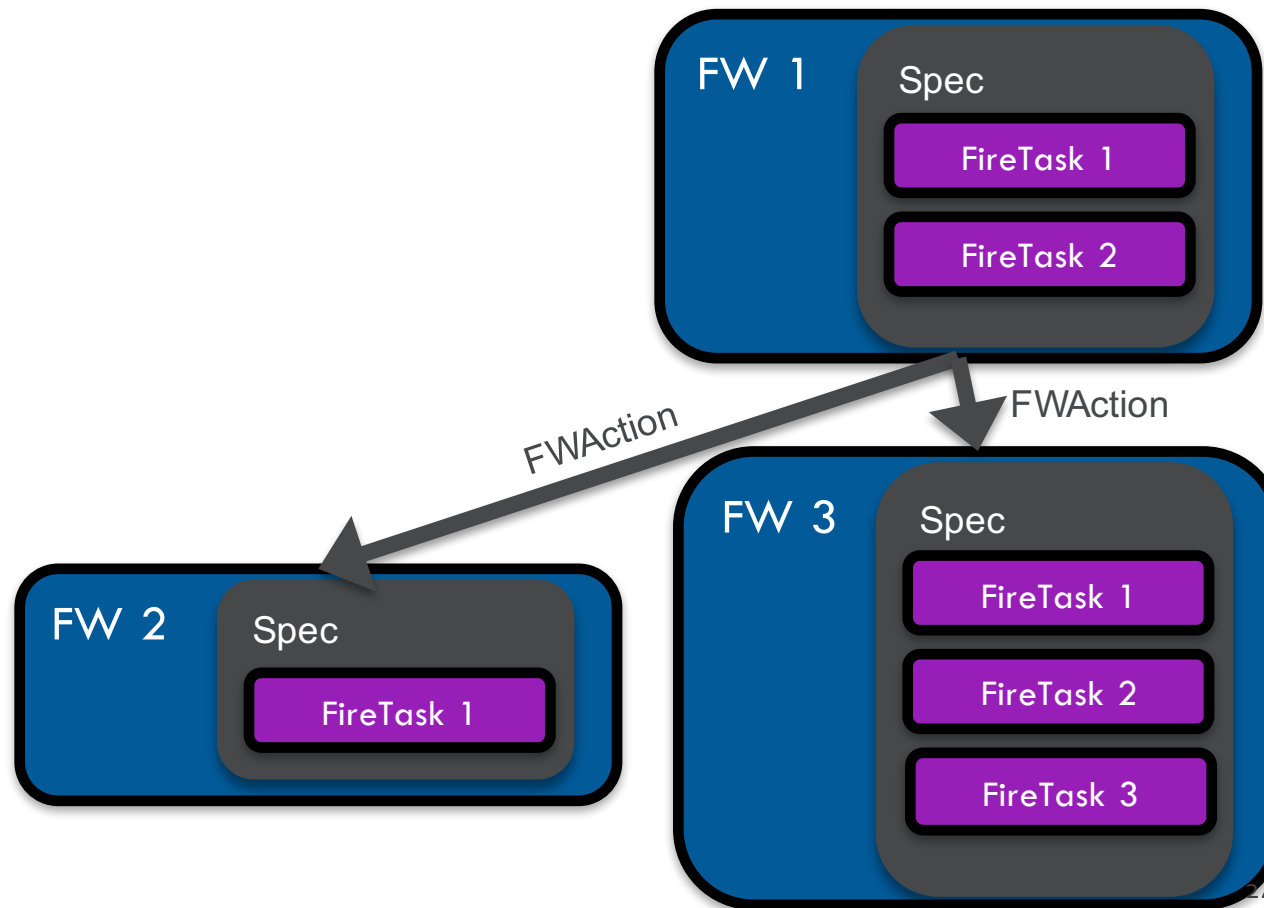
- Built w/Python+MongoDB. Open-source, pip-installable: <http://pythonhosted.org/FireWorks/>
- >7500 downloads per month
 - #1 Google hit for “Python workflow software”
 - #4 software hit for “scientific workflow software”
- 1 of 2 workflow softwares officially supported by NERSC
- Worldwide usage
- 97th percentile, scientific Python software impact (Depsy) paper published in C.S. journal

Major new features:

- Completely redesigned frontend,
- Auto-reporting compiled using Mongo aggregation framework
- Compatibility with Argonne Leadership Computing Facility scheduler (Cobalt)
- Major features such as task-level restart contributed by community



HOW WORKFLOWS ARE STRUCTURED



CODE EXAMPLE

```
from fireworks import Firework, Workflow, LaunchPad, ScriptTask
from fireworks.core.rocket_launcher import rapidfire

# set up the LaunchPad and reset it (first time only)
launchpad = LaunchPad()
launchpad.reset('', require_password=False)

# define the individual FireWorks and Workflow
fw1 = Firework(ScriptTask.from_str('echo "To be, or not to be,"'))
fw2 = Firework(ScriptTask.from_str('echo "that is the question:"'))
wf = Workflow([fw1, fw2], {fw1:fw2}) # set of FWs and dependencies

# store workflow in LaunchPad
launchpad.add_wf(wf)
# pull all jobs and run them locally
rapidfire(launchpad)
```

WORKFLOWS ARE SIMPLE JSON/YAML DOCUMENTS THAT HAVE VERY LITTLE “FLUFF”

(this is YAML, a bit prettier for humans but less pretty for computers)

```
fw:  
- fw_id: 1  
  spec:  
    _tasks:  
      - _fw_name: ScriptTask:  
        script: echo 'To be, or not to be,'  
- fw_id: 2  
  spec:  
    _tasks:  
      - _fw_name: ScriptTask  
        script: echo 'that is the  
question:'  
links:  
  1:  
    - 2  
metadata: {}
```

The same JSON document will produce the same result on any computer (with the same Python functions).

JSON + MONGODB MEANS YOU CAN STORE WORKFLOWS DIRECTLY AND MAKE RICH QUERIES

(this is YAML, a bit prettier for humans but less pretty for computers)

```
fw:  
- fw_id: 1  
  spec:  
    _tasks:  
    - _fw_name: ScriptTask:  
      script: echo 'To be, or not to be,'  
- fw_id: 2  
  spec:  
    _tasks:  
    - _fw_name: ScriptTask  
      script: echo 'that is the  
question:'  
links:  
  1:  
    - 2  
metadata: {}
```

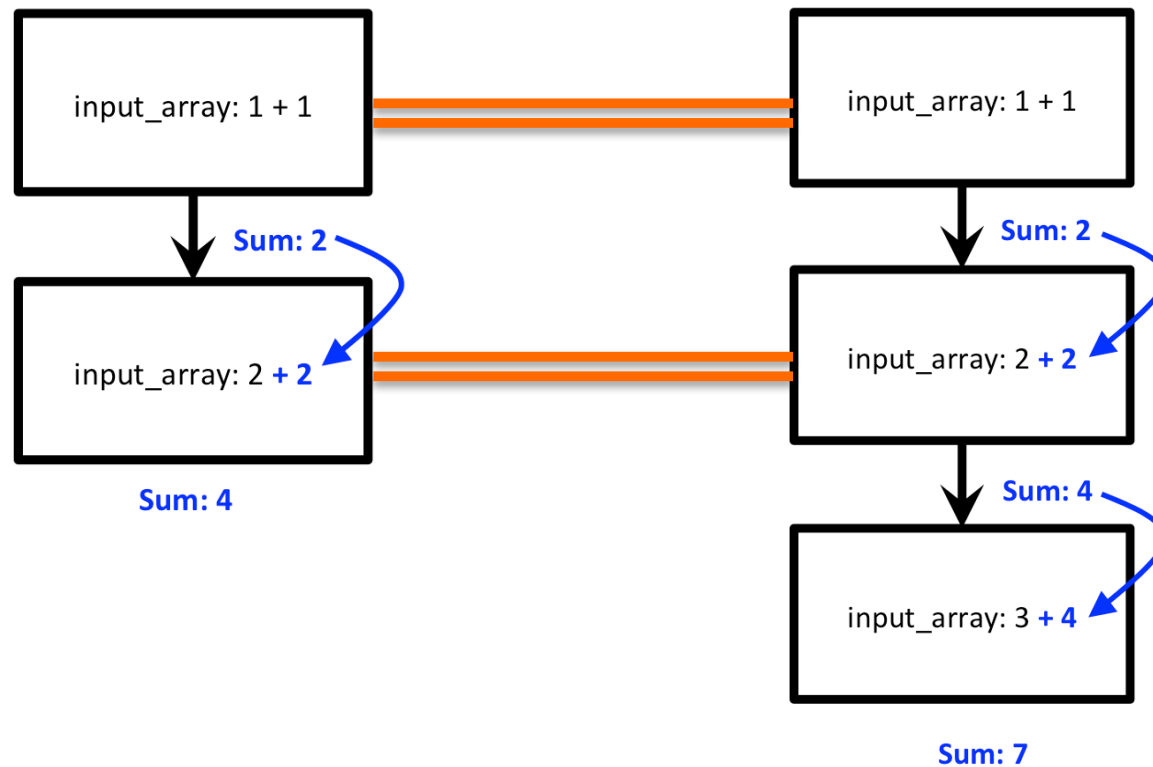


Just some of your search options:

- simple matches
- match in array
- greater than/less than
- regular expressions
- match subdocument
- Javascript function
- MapReduce...

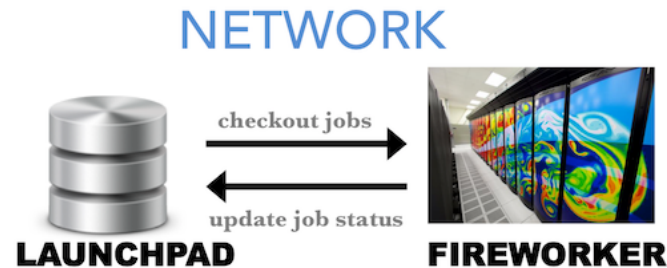
All for free, and all on the native workflow format!

JSON DUPLICATE CHECKING SIMPLE AND AUTOMATIC



“PULL” STRUCTURE (NOT PUSH) MAKES LIFE EASY: EXECUTION MODES ARE ALL VARIATIONS ON A THEME

- Theme: Worker machine pulls a job & runs it



- Variation 1:
 - different workers can be configured to pull different types of jobs via config + MongoDB
- Variation 2:
 - worker machines sort the jobs by a priority key and pull matching jobs the highest priority

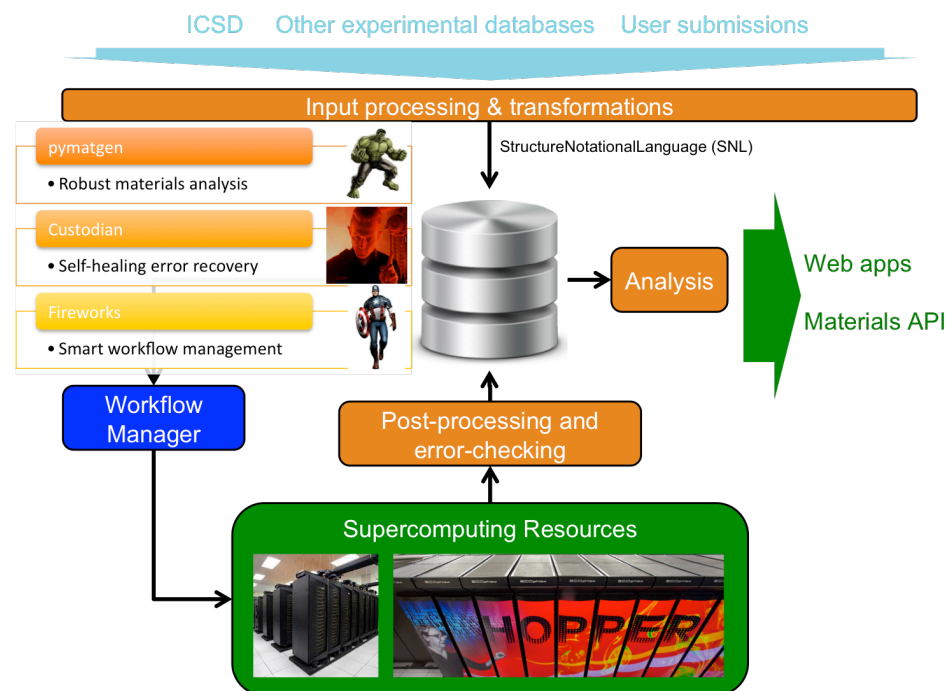
JCESR'S USAGE

JCESR leverages ALCF, MIT, and NERSC resources in a distributed fashion with the high-throughput (HT) infrastructure of the Materials Project to:

- Execute an electronic structure code (VASP, Qchem, Gaussian, AbInit, etc) with reasonable efficiency
- Monitor the calculations using the Fireworks workflow software to enable automatic restarts, saving of intermittent data, etc

Issues:

- ALCF does not host Mongo DB databases requiring users to setup and secure remote instances
- No method for telling a task to run outside the job scheduler even if there's no compute process



Automatic workflow of first-principles calculations