

Introduction to Makeflow and Work Queue



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Go to <http://ccl.nd.edu> and Click on ACIC Tutorial

The Cooperative Computing Lab

[Software](#) | [Download](#) | [Manuals](#) | [Forum](#) | [Papers](#)

Go to the **ACIC 2017 Tutorial** on Makeflow and Work Queue, Nov 14th and 16th!

About the CCL

We design software that enables our collaborators to easily harness large scale distributed systems such as clusters, clouds, and grids. We perform fundamental computer science research that enables new discoveries through computing in fields such as physics, chemistry, bioinformatics, biometrics, and data mining.

CCL News and Blog

- [Automatic job sizing for maximum throughput](#) (26 Oct 2017)
- [Makeflow Feature: JX Representation](#) (18 Oct 2017)
- [Announcement: CCTools 6.2.0 released](#) (09 Oct 2017)
- [2017 DISC Summer REU Conclusion](#) (30 Aug 2017)
- [Announcement: CCTools 6.1.6 released](#) (29 Aug 2017)
- [Talk at ScienceCloud Workshop](#) (27 Jun 2017)
- [Congrats to Ph.D Graduates](#) (22 May 2017)
- [Announcement: CCTools 6.1.0. released](#) (17 May 2017)
- [Makeflow and Mesos Paper at CCGrid 2017](#) (05 May 2017)
- [\(more news\)](#)



Community Highlight

Lifemapper is a high-throughput, webservice-based, single- and multi-species modeling and analysis system designed at the Biodiversity Institute and Natural History Museum, University of Kansas. Lifemapper was created to compute and web publish, species distribution models using available online species occurrence data. Using the Lifemapper platform, known species localities georeferenced from museum specimens are combined with climate models to predict a species' "niche" or potential habitat availability, under current day and future climate change scenarios. By assembling large numbers of known or predicted species distributions, along with phylogenetic and biogeographic data, Lifemapper can analyze biodiversity, species communities, and evolutionary influences at the landscape level.



Lifemapper has had difficulty scaling recently as our projects and analyses are growing exponentially. For a large proof-of-concept project we deployed on the XSEDE resource Stampede at TACC, we integrated Makeflow and Work Queue into the job workflow. Makeflow simplified job dependency management and reduced job-scheduling overhead, while Work Queue scaled our computation capacity from hundreds of simultaneous CPU cores to thousands. This allowed us to perform a sweep of computations with various parameters and high-resolution inputs producing a plethora of outputs to be analyzed and compared. The experiment worked so well that we are now integrating Makeflow and Work Queue into our core infrastructure. Lifemapper benefits not only from the increased speed and efficiency of computations, but the reduced complexity of the data management code, allowing developers to focus on new analyses and leaving the logistics of job dependencies and resource allocation to these tools.



The Cooperative Computing Lab

- We **collaborate with people** who have large scale computing problems in science, engineering, and other fields.
- We **operate computer systems** on the O(10,000) cores: clusters, clouds, grids.
- We **conduct computer science research** in the context of real people and problems.
- We **develop open source software** for large scale distributed computing.

<http://ccl.cse.nd.edu>



Outline

Tuesday, Nov 14th

- Thinking Opportunistically
- Overview of the Cooperative Computing Tools
- Makeflow
- Makeflow + Work Queue
- Hands-On Tutorial

Thursday, Nov 16th

- Makeflow Features
 - Resource Management
 - Containers
- Work Queue API
- Hands-On Tutorial



Thinking Opportunistically



Opportunistic Computing

- Much of scientific computing is done in conventional computing centers with a fixed operating environment with professional sysadmins.
- But, there exists a large amount of computing power available to end users that is not prepared or tailored to your specific application:
 - National HPC facility
 - Campus-level cluster and batch system.
 - Volunteer computing systems: Condor, BOINC, etc.
 - Cloud services.
- Can we effectively use these systems for “long tail” scientific computing?

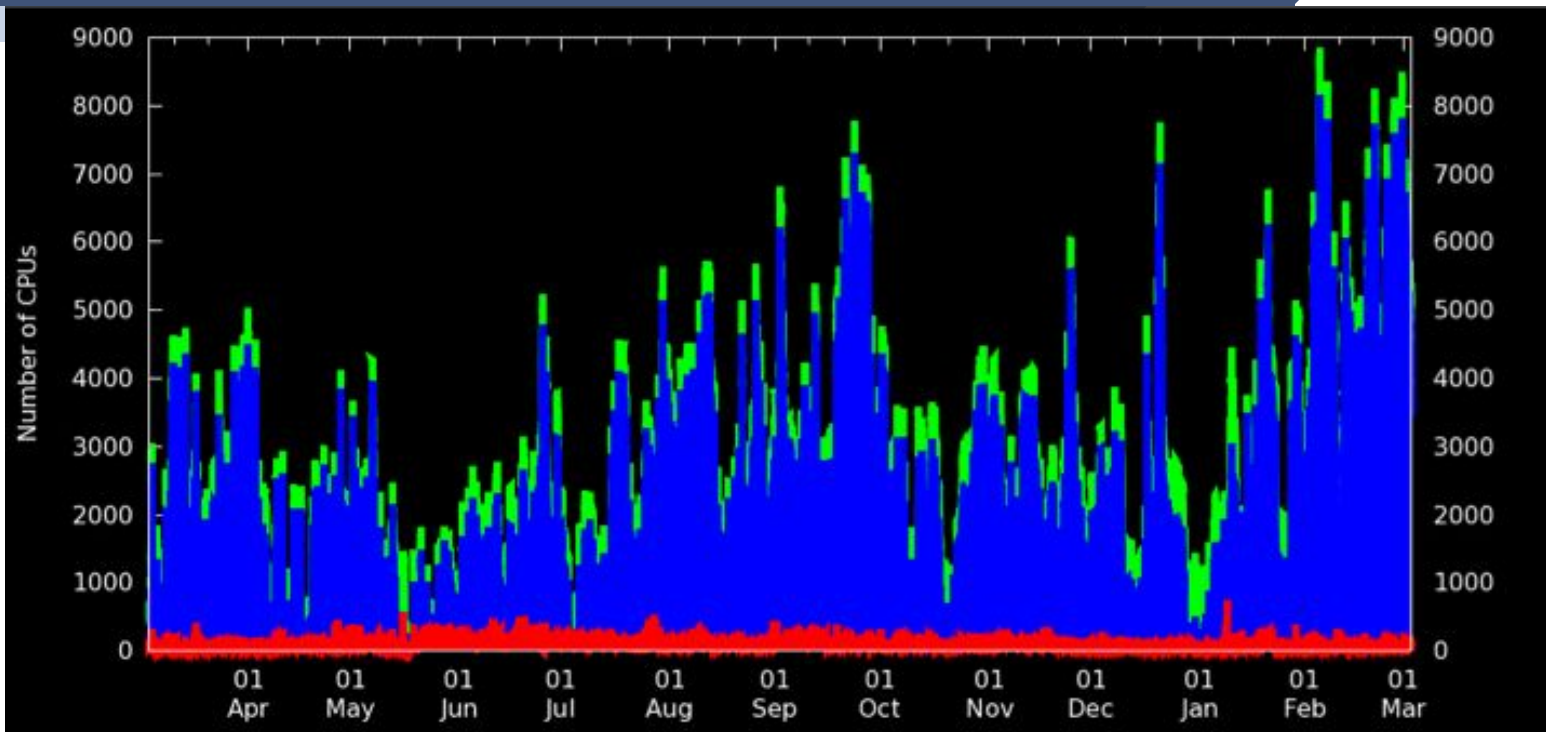


Opportunistic Challenges

- When borrowing someone else's machines, you cannot change the OS distribution, update RPMs, patch kernels, run as root...
- This often puts important technology just out of reach of the end user, e.g.:
 - FUSE might be installed, but without setuid binary.
 - Docker might be available, but you aren't a member of the required Unix group.
- The resource management policies of the hosting system may work against you:
 - Preemption due to submission by higher priority users.
 - Limitations on execution time and disk space.
 - Firewalls only allow certain kinds of network connections.

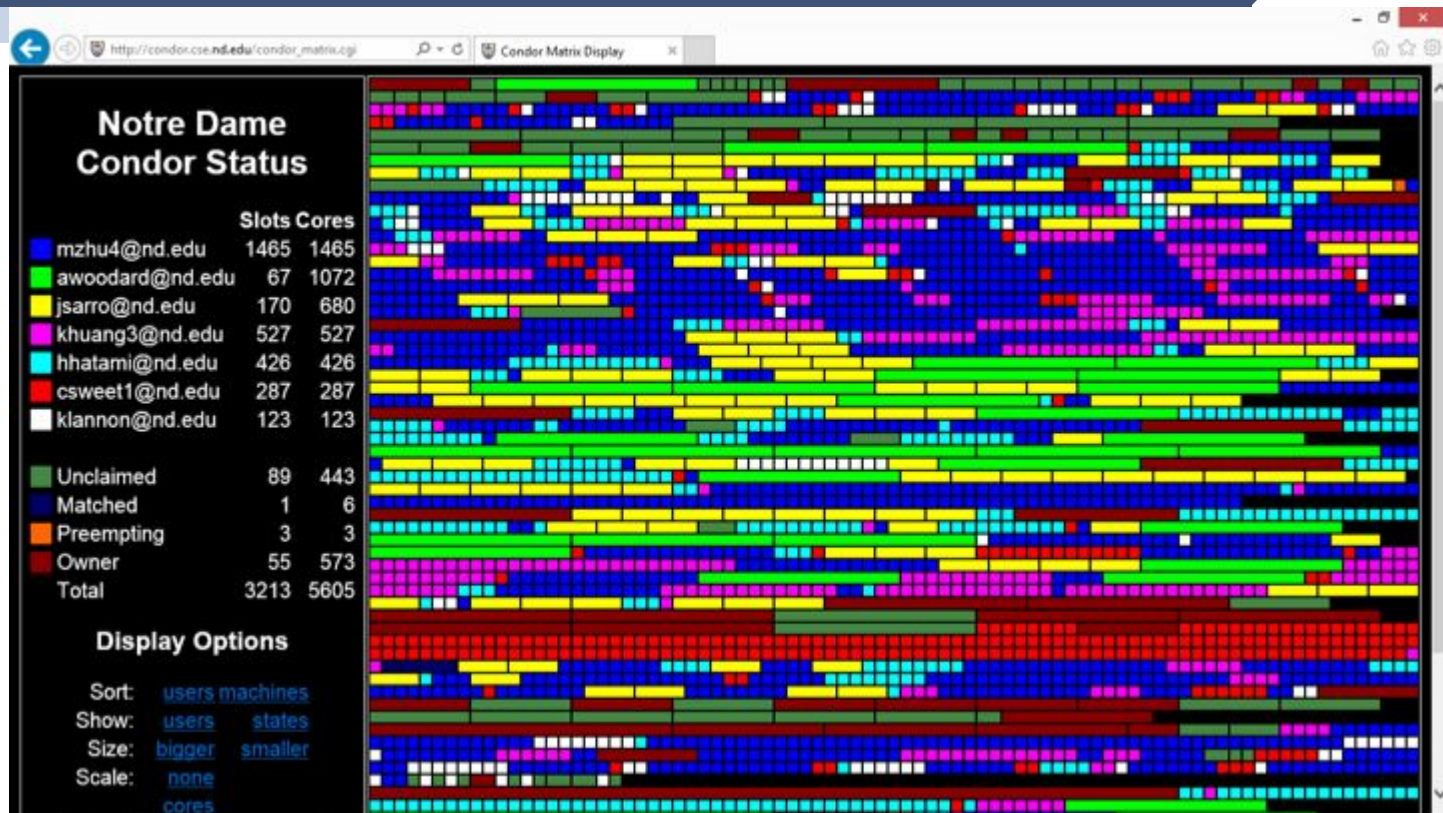


Backfilling HPC with Condor at Notre Dame





Users of Opportunistic Cycles





I can get as many machines
on the cloud/grid as I want!

How do I organize my application
to run on those machines?



Cooperative Computing Tools



Our Philosophy

- Harness all available resources: desktops, clusters, clouds, and grids.
- Make it easy to scale up from one desktop to national scale infrastructure.
- Provide familiar interfaces that make it easy to connect existing apps together.
- Allow portability across operating systems, storage systems, middleware...
- Make simple things easy, and complex things possible.
- **No special privileges required.**



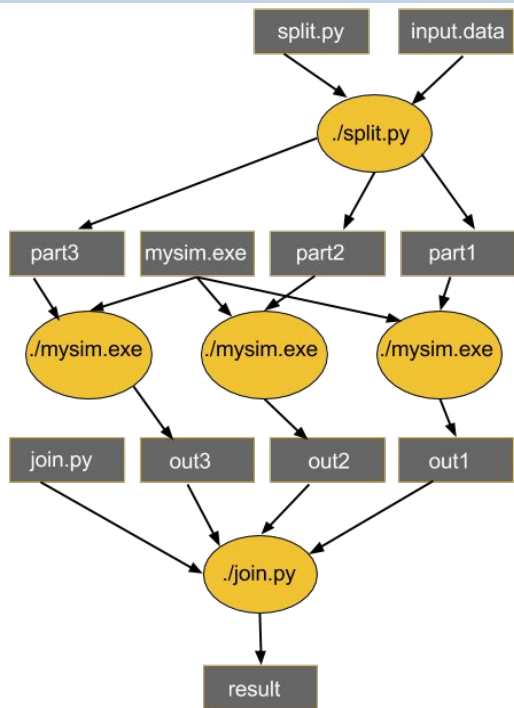
A Quick Tour of the CCTools

- Open source, GNU General Public License.
- Compiles in 1-2 minutes, installs in \$HOME.
- Runs on Linux, Solaris, MacOS, FreeBSD, ...
- Interoperates with many distributed computing systems.
 - Condor, SGE, Torque, Globus, iRODS, Hadoop...
- Components:
 - Makeflow – A portable workflow manager.
 - Work Queue – A lightweight distributed execution system.
 - Parrot – A personal user-level virtual file system.
 - Chirp – A user-level distributed filesystem.

<http://ccl.cse.nd.edu/software>



MAKEFLOW (MAKE + WORKFLOW)



- Provides portability across batch systems.
- Enable parallelism (but not too much!)
- Fault tolerance at multiple scales.
- Data and resource management.

Makeflow

Local

Condor

SGE

Work
Queue

<http://ccl.cse.nd.edu/software/makeflow>



Work Queue API

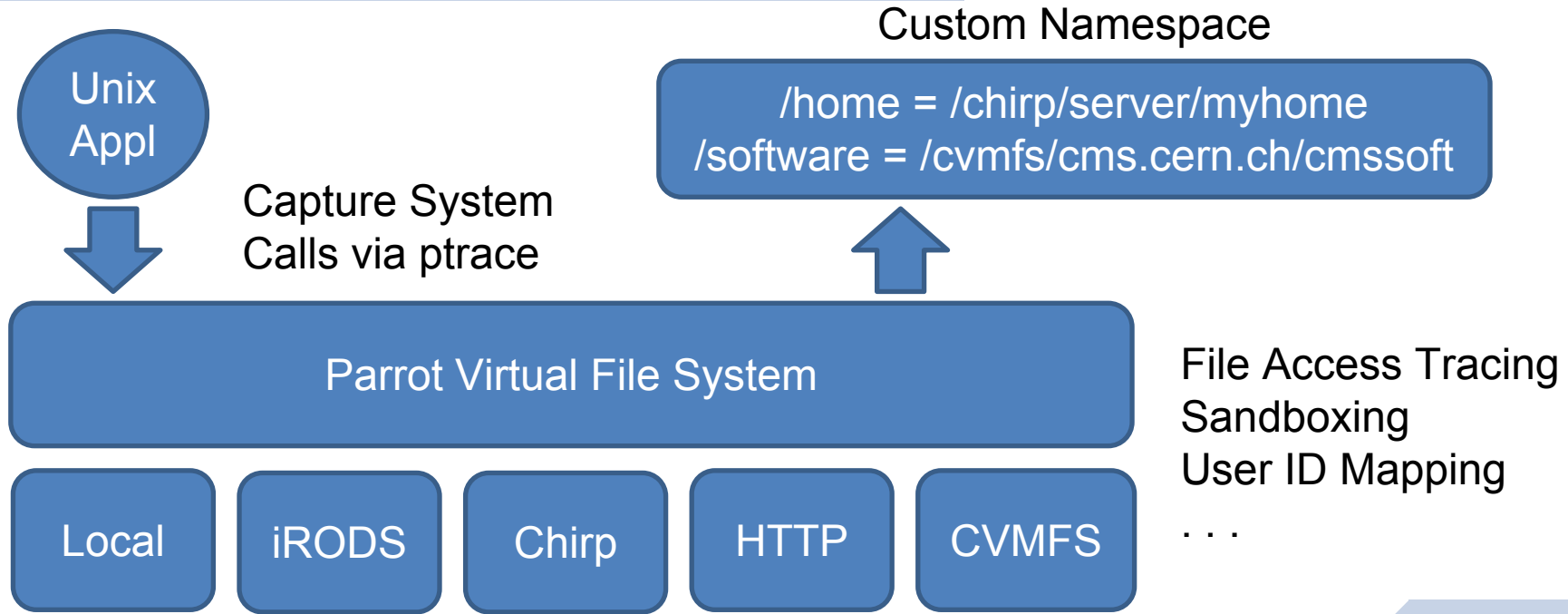
```
#include "work_queue.h"
while( not done ) {

    while (more work ready) {
        task = work_queue_task_create();
        // add some details to the task
        work_queue_submit(queue, task);
    }

    task = work_queue_wait(queue);
    // process the completed task
}
```



Parrot Virtual File System





Lots of Documentation

The screenshot shows a web browser with three tabs. The first tab is 'The Cooperative Comput...'. The second tab is 'Makeflow = Make + Work'. The third tab is 'makeflow(1)'. The address bar shows 'ccl.cse.nd.edu/software/manuals/man/makeflow.html'. The page content includes sections for NAME, SYNOPSIS, DESCRIPTION, and OPTIONS.

makeflow(1)

NAME

makeflow - workflow engine for executing distributed workflows

SYNOPSIS

makeflow [options] <dagfile>

DESCRIPTION

Makeflow is a workflow engine for distributed computing. It accepts a specification of a large amount of work to be performed, and runs it on remote machines in parallel where possible. In addition, **Makeflow** is fault-tolerant, so you can use it to coordinate very large tasks that may run for days or weeks in the face of failures. **Makeflow** is designed to be similar to Make, so if you can write a Makefile, then you can write a **Makeflow**.

You can run a **Makeflow** on your local machine to test it out. If you have a multi-core machine, then you can run multiple tasks simultaneously. If you have a Condor pool or a Sun Grid Engine batch system, then you can send your jobs there to run. If you don't already have a batch system, **Makeflow** comes with a system called Work Queue that will let you distribute the load across any collection of machines, large or small.

OPTIONS

When **makeflow** is ran without arguments, it will attempt to execute the workflow specified by the **Makeflow** dagfile using the local execution engine.

Commands

-c, --clean Clean up: remove logfile and all targets.
-f, --summary-log <file>
Write summary of workflow to file.

<http://ccl.cse.nd.edu>

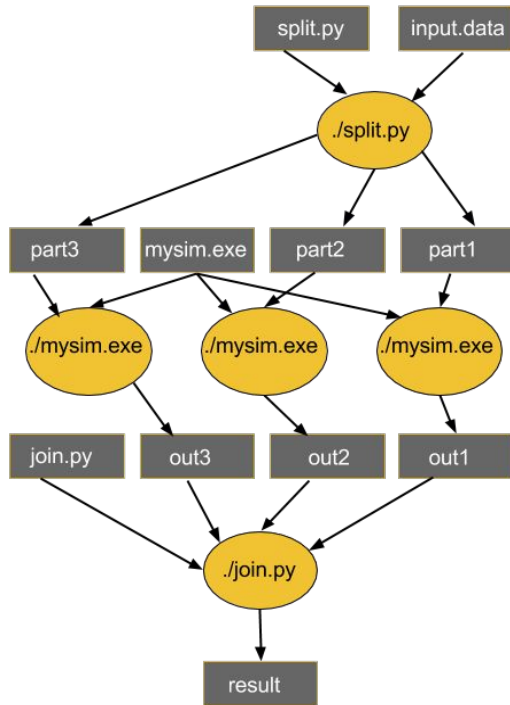


Makeflow

A Portable Workflow System



MAKEFLOW (MAKE + WORKFLOW)



- Provides portability across batch systems.
- Enable parallelism (but not too much!)
- Trickle out work to batch system
- Fault tolerance at multiple scales.
- Data and resource management.



Local

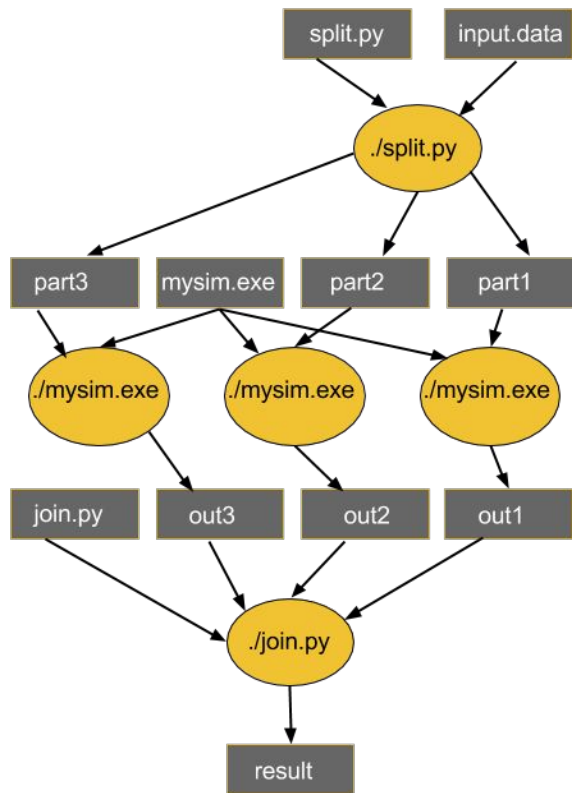
Condor

SGE

Work
Queue



MAKEFLOW (MAKE + WORKFLOW) BASED OFF AN OLD IDEA: MAKEFILES



part1 part2 part3: input.data split.py
./split.py input.data

out1: part1 mysim.exe
./mysim.exe part1 >out1

out2: part2 mysim.exe
./mysim.exe part2 >out2

out3: part3 mysim.exe
./mysim.exe part3 >out3

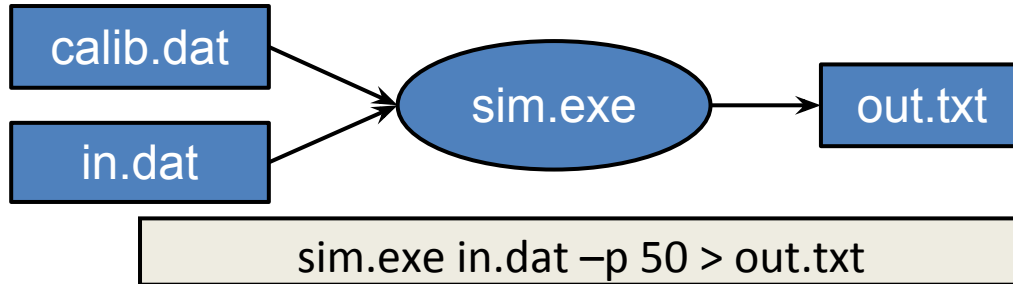
result: out1 out2 out3 join.py
./join.py out1 out2 out3 > result



Makeflow Syntax

[output files] : [input files]
[command to run]

One Rule



out.txt : in.dat **calib.dat sim.exe**
sim.exe in.data -p 50 > out.txt



Makeflow Syntax : sims.mf

```
out.10 : in.dat calib.dat sim.exe  
sim.exe -p 10 in.data > out.10
```

```
out.20 : in.dat calib.dat sim.exe  
sim.exe -p 20 in.data > out.20
```

```
out.30 : in.dat calib.dat sim.exe  
sim.exe -p 30 in.data > out.30
```



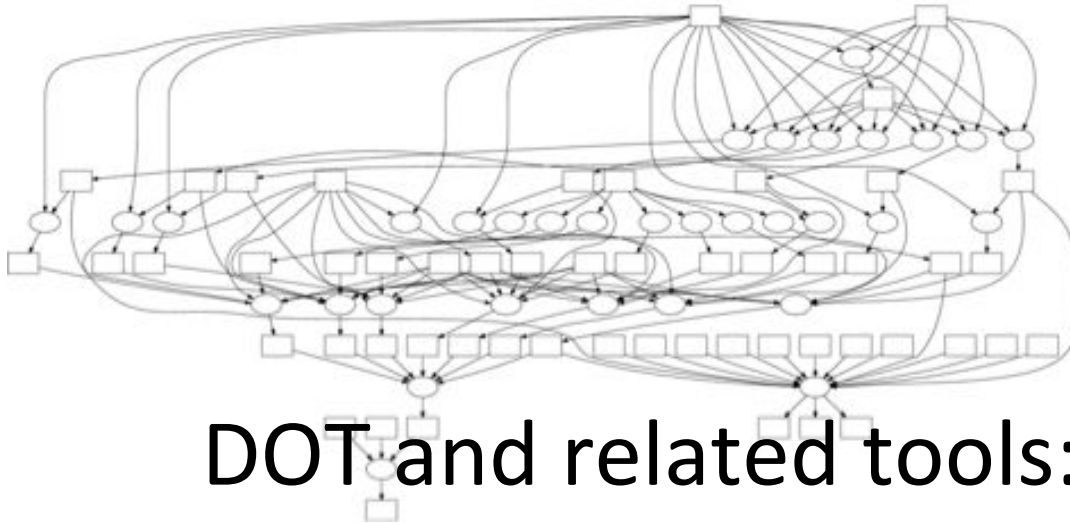
How to run a Makeflow

- Run a workflow locally (multicore?)
 - `makeflow -T local sims.mf`
- Clean up the workflow outputs:
 - `makeflow -c sims.mf`
- Run the workflow on Torque:
 - `makeflow -T torque sims.mf`
- Run the workflow on Condor:
 - `makeflow -T condor sims.mf`



Visualization with DOT

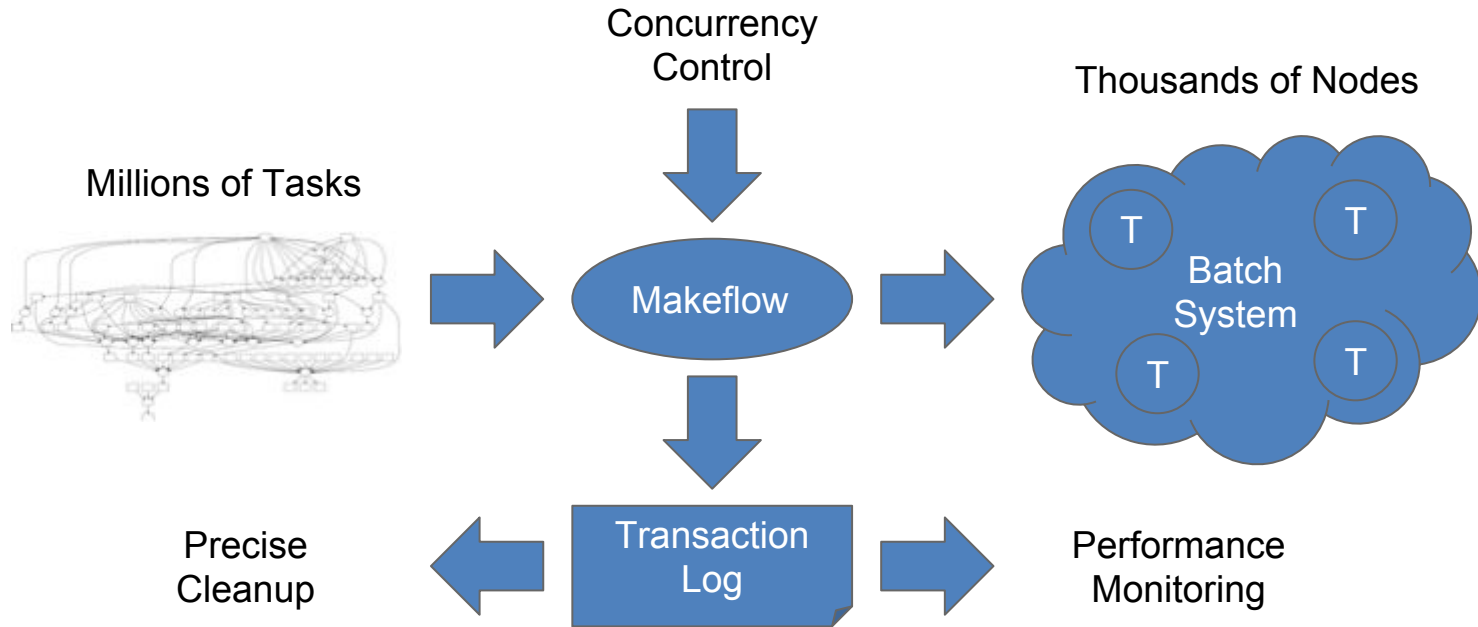
- `makeflow_viz -D example.mf > example.dot`
- `dot -T gif < example.dot > example.gif`



DOT and related tools:
<http://www.graphviz.org>



Makeflow Shapes a Workflow





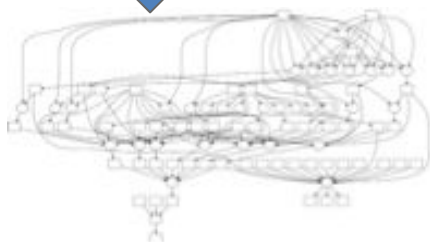
Example: Biocompute Portal



BLAST
SSAHA
SHRIMP
EST
MAKER
...

Progress
Bar

Generate
Makeflow

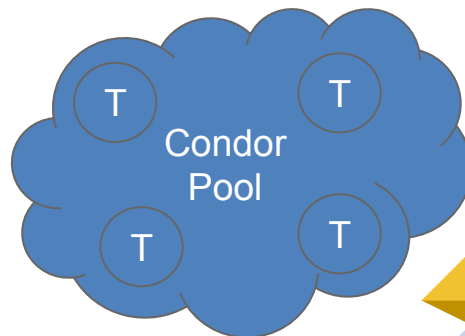


Transaction
Log

Run
Makeflow

Makeflow

Update
Status





Makeflow Applications

- Bioinformatics
- Biometrics
- High Energy Physics

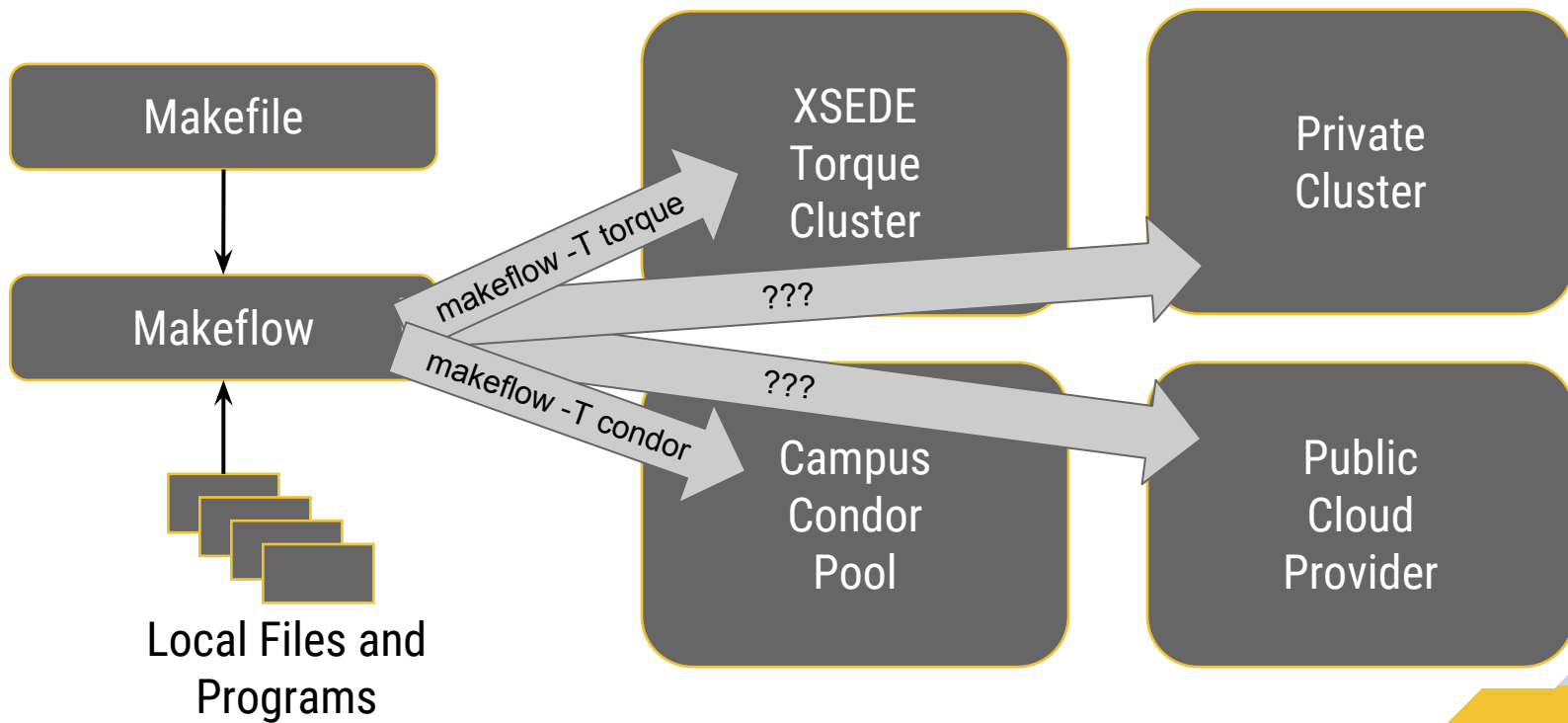


Makeflow + Work Queue

A Portable Workflow System

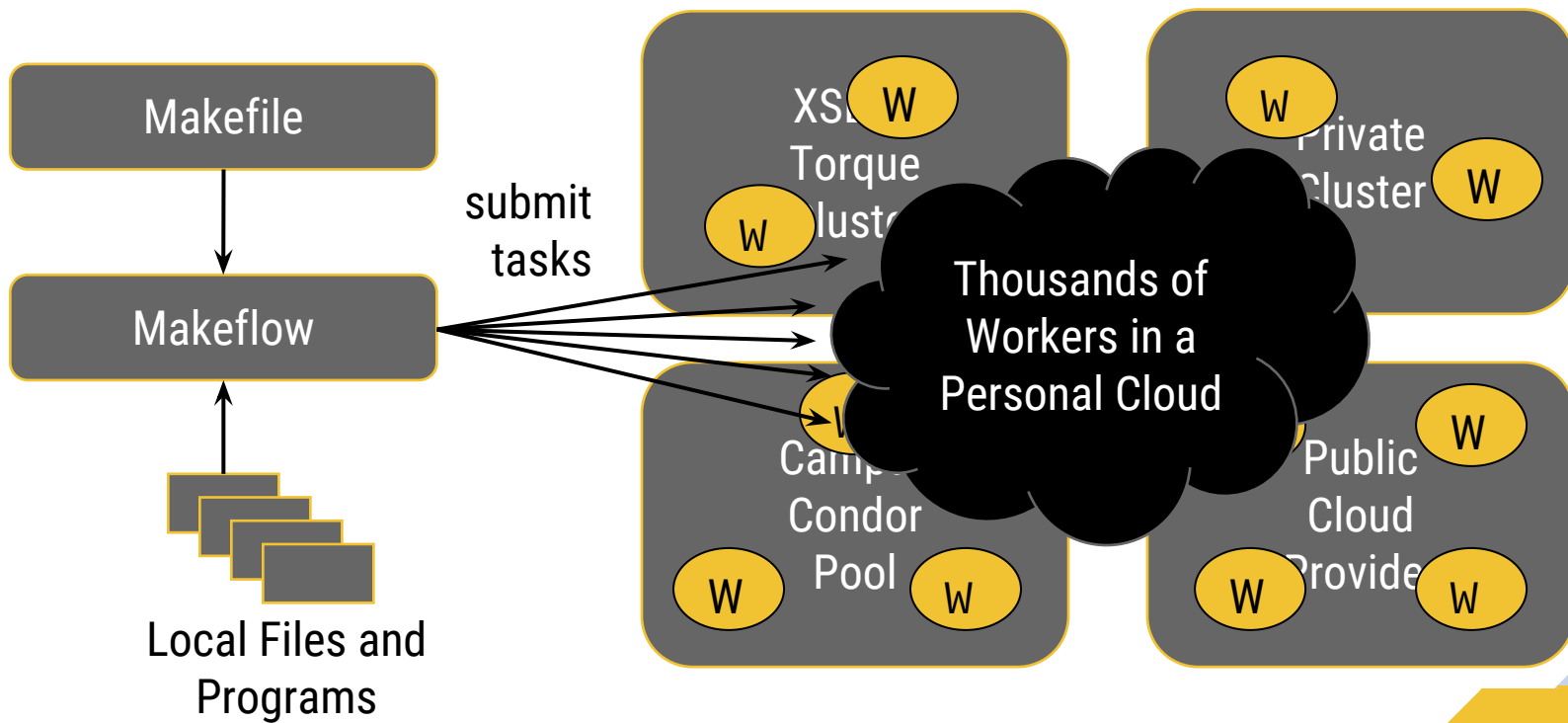


MAKEFLOW



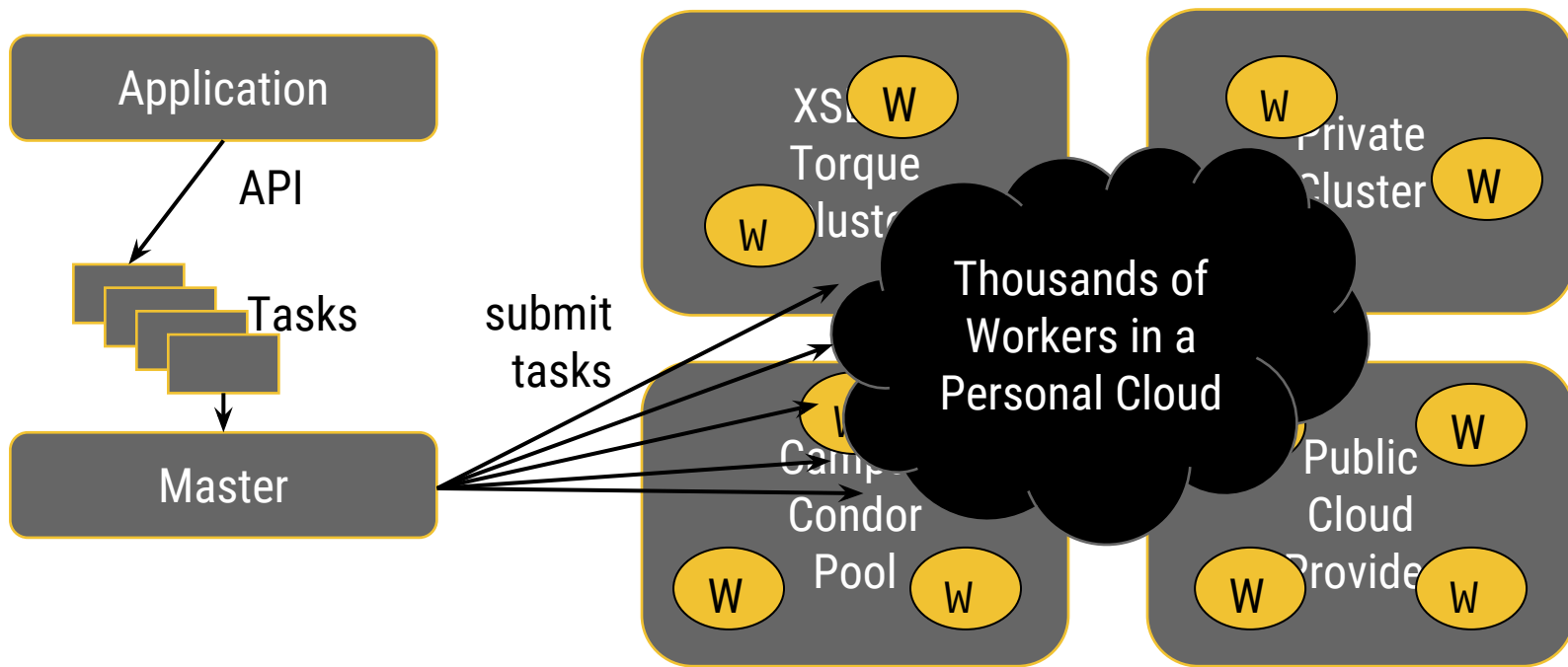


MAKEFLOW + WORK QUEUE





WORK QUEUE





Advantages of Work Queue

- Harness multiple resources simultaneously.
- Hold on to cluster nodes to execute multiple tasks rapidly.
 - (ms/task instead of min/task)
- Scale resources up and down as needed.
- Better management of data, with local caching for data intensive tasks.
- Matching of tasks to nodes with data.



Makeflow and Work Queue

To start the Makeflow

```
% makeflow -T wq sims.mf
```

Could not create work queue on port 9123.

```
% makeflow -T wq -p 0 sims.mf
```

Listening for workers on port 8374...

To start one worker:

```
% work_queue_worker master.hostname.org 8374
```



Start 25 Workers in Batch System

Submit workers to Condor:

```
condor_submit_workers master.hostname.org 8374 25
```

Submit workers to SGE:

```
sgc_submit_workers master.hostname.org 8374 25
```

Submit workers to Torque:

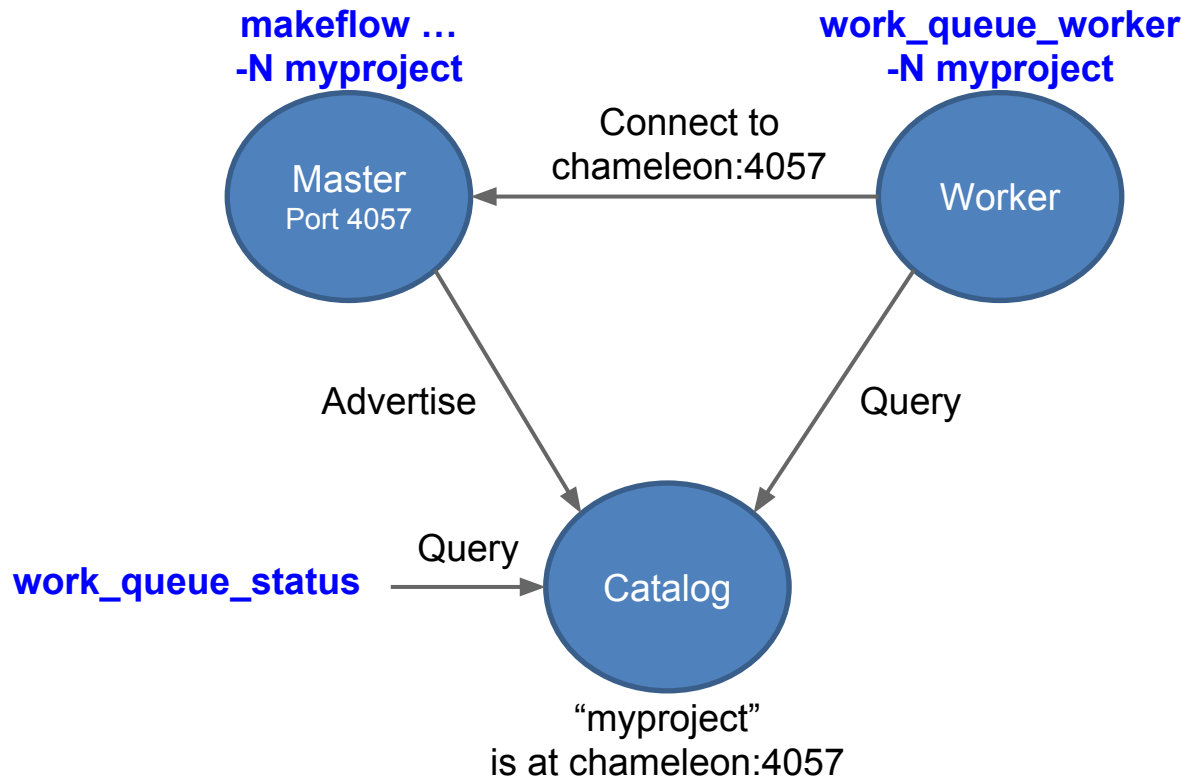
```
torque_submit_workers master.hostname.org 8374 25
```



**Keeping track of port
numbers gets old fast...**



Project Names





Project Names

Start Makeflow with a project name:

```
% makeflow -T wq -N myproject sims.mf
```

Listening for workers on port XYZ...

Start one worker:

```
% work_queue_worker -N myproject
```

Start many workers:

```
% torque_submit_workers -N myproject 5
```



work_queue_status

```
wizard.cse.nd.edu - PuTTY
% ./work_queue_status
PROJECT          NAME                PORT  WAITING  BUSY  COMPLETE  WORKERS
awe-fip35         fahnd04.crc.nd.edu    1024   719     1882  1206967   1882
hfeng-gromacs-10ps lclsstor01.crc.nd.edu 1024   4980     0    1280240   111
hfeng2-ala5       lclsstor01.crc.nd.edu 1025   2404    140   1234514   140
forcebalance      leeping.Stanford.EDU  5817   1082     26    822       26
forcebalance      leeping.Stanford.EDU  9230     0        3    147       3
fg-tutorial       login1.futuregrid.tacc 1024     3        0     0        0
% █
```



Advantages of Work Queue

- MF +WQ is fault tolerant in many different ways:
 - If Makeflow crashes (or is killed) at any point, it will recover by reading the transaction log and continue where it left off.
 - Makeflow keeps statistics on both network and task performance, so that excessively bad workers are avoided.
 - If a worker crashes, the master detects failure and restarts the task elsewhere.
 - Workers can be added and removed at any time during workflow execution.
 - Multiple masters with the same project name can be added and removed while the workers remain.
 - If the worker sits idle for too long (default 15m) it will exit, so as not to hold resources idle.



Alternative Makeflow Formats

Utilizing JSON/JX for easier scripting

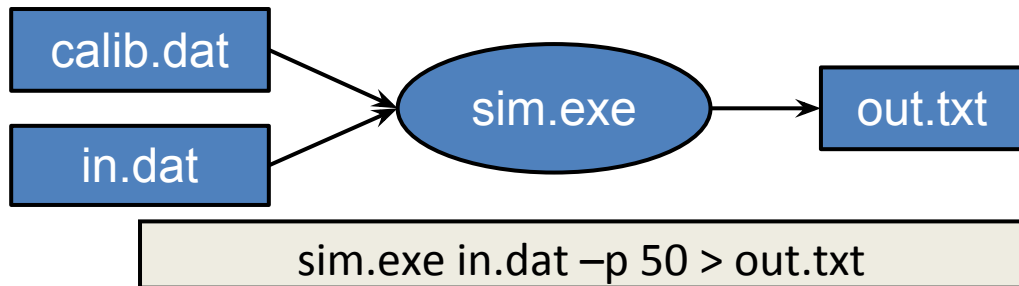


Makeflow JSON Syntax

- Verbose flexible structure
- Familiar structure
- Consists of four items:
 - ▶ "categories": Object<Category>
 - ▶ "default_category": String
 - ▶ "environment": Object<String>
 - ▶ "rules": Array<Rule>



Makeflow JSON Syntax



```
{  
  "outputs": [{"path": "out.txt"}],  
  "inputs": [ {"path": "in.dat"}, {"path": "calib.dat"}, {"path": "sim.exe"}]  
  "command": "sim.exe -p 50 in.data > out.txt",  
}
```



Makeflow JSON Syntax

```
{  
  "outputs": [{"path": "out_10.txt"}],  
  "inputs": [ {"path": "in.dat"}, {"path": "calib.dat"}, {"path": "sim.exe"}]  
  "command": "sim.exe -p 10 in.data > out_10.txt",  
},  
{  
  "outputs": [{"path": "out_20.txt"}],  
  "inputs": [ {"path": "in.dat"}, {"path": "calib.dat"}, {"path": "sim.exe"}]  
  "command": "sim.exe -p 20 in.data > out_20.txt",  
},...
```



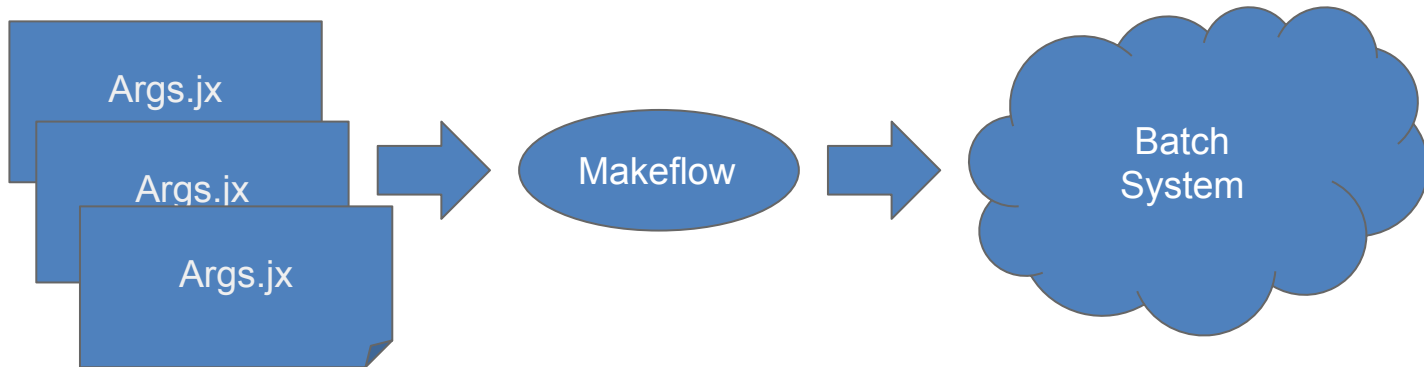
Makeflow JSON Rule

- "inputs": Array<File>
- "outputs": Array<File>
- "command": String
- "local_job": Boolean
- "category": String
- "resources": Resources
- "allocation": String
- "environment": Object<String>



Makeflow JX Syntax

- Allows for more compact makeflows.
 - Provides functions for expanding tasks: range, variables, etc...
- Can be used as templates in conjunction with an arguments file.
- Useful for consistently structure data and different data.





Makeflow JX Syntax

```
{  
  "outputs": [{"path": format("out_%d.txt", i)}],  
  "inputs": [ {"path": "in.dat"}, {"path": "calib.dat"}, {"path": "sim.exe"}]  
  "command": format("sim.exe -p %d in.data > out_%d.txt", i),  
} for i in range(10, 30, 10),
```



How to run a Makeflow

- Run a workflow from json
 - `makeflow --json sims.json`
- Clean up the workflow outputs:
 - `makeflow -c --json sims.json`
- Run the workflow from jx:
 - `makeflow --jx sims.jx`
- Run the workflow with jx and args:
 - `makeflow --jx sims.jx --jx-args args.jx`



Resource Management

Allowing tasks to share resources



Why Manage Resource?

- More accurate accounting and provisioning.
- Allows for multi-tenant situations.
- Provides consistent resources to tasks.
 - Prevents slower execution.
 - Mitigate failures from under provisioning.

How can this happen?



Makeflow Resource Specification

- Category
 - ▷ Cores
 - ▷ Memory
 - ▷ Disk

```
...  
CATEGORY=analysis  
DISK=1024  
MEMORY=1024  
CORES=1  
  
out1: part1 mysim.exe  
      ./mysim.exe part1 >out1  
  
out2: part2 mysim.exe  
      ./mysim.exe part2 >out2  
  
...
```



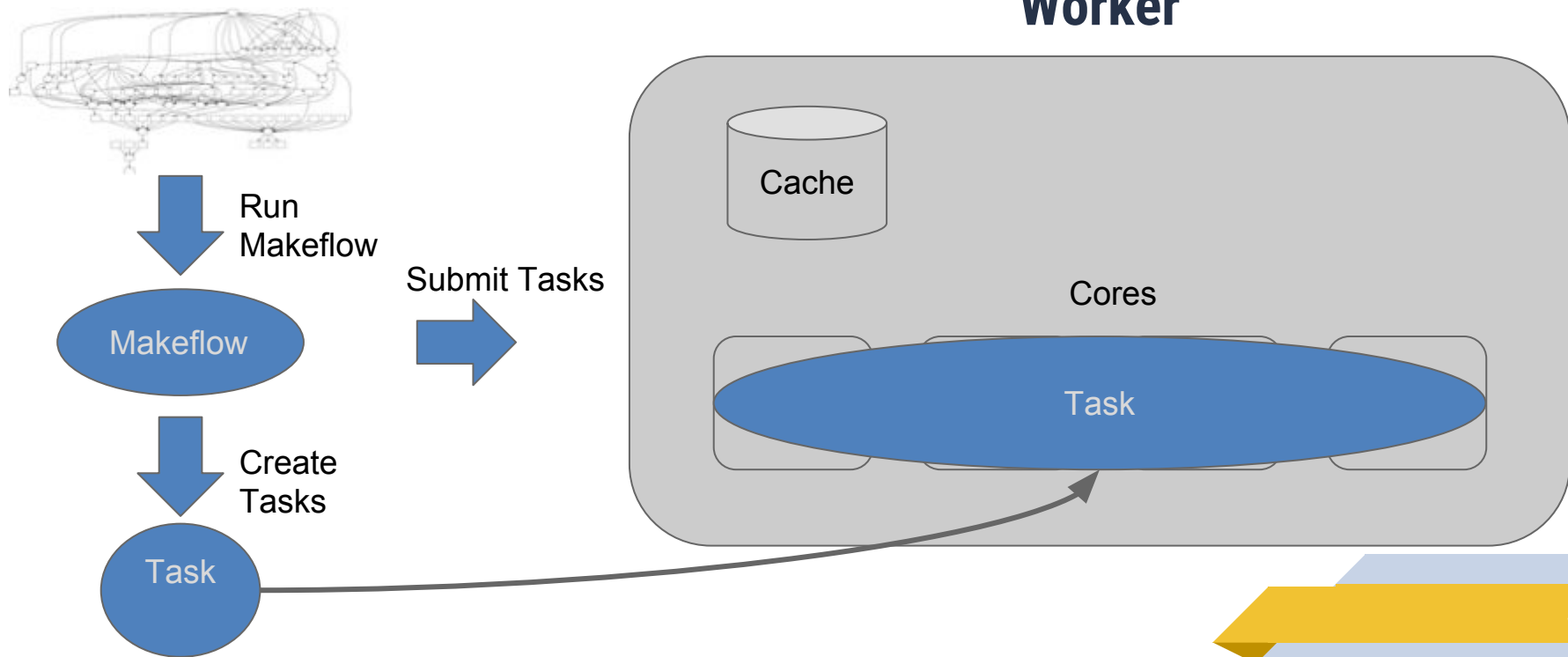
Makeflow Resource Specification

- Category
 - ▷ Cores
 - ▷ Memory
 - ▷ Disk

```
...  
CATEGORY=analysis  
DISK=1024  
MEMORY=1024  
CORES=1  
  
out1: part1 mysim.exe  
./mysim.exe part1 >out1  
  
...  
  
CATEGORY=join  
DISK=2048  
MEMORY=2048  
CORES=2  
  
result: out1 out2 out3 join.py  
./join.py out1 out2 out3 > result
```

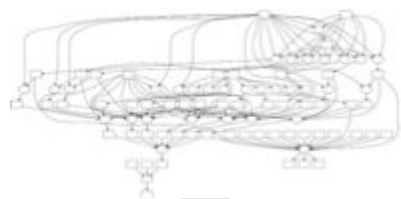


Work Queue Workers





Work Queue Multi-tenant Workers



Run
Makeflow

Makeflow

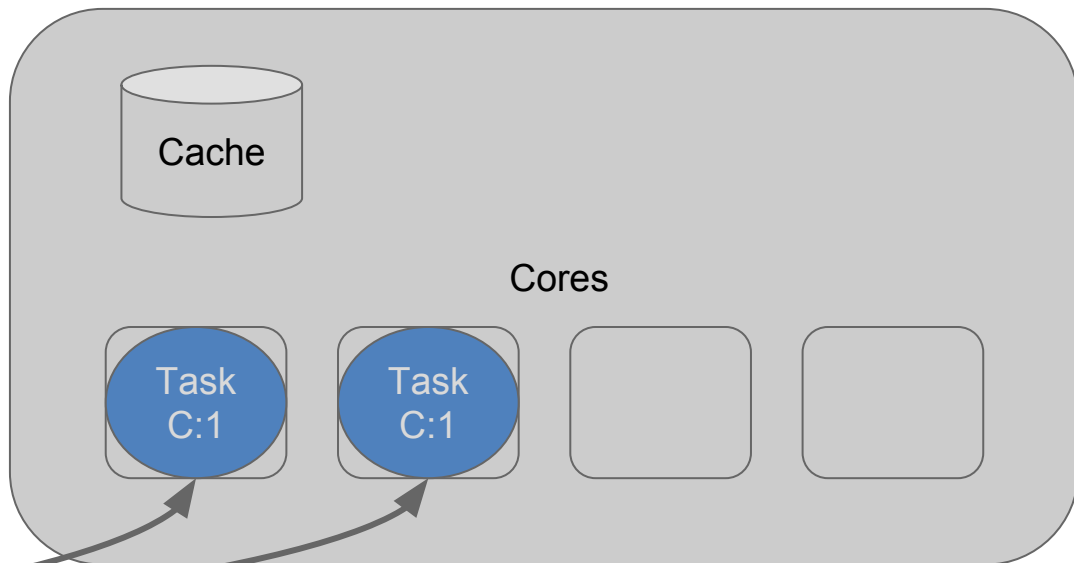
Create Labeled
Task

Task
C:1

Task
C:1

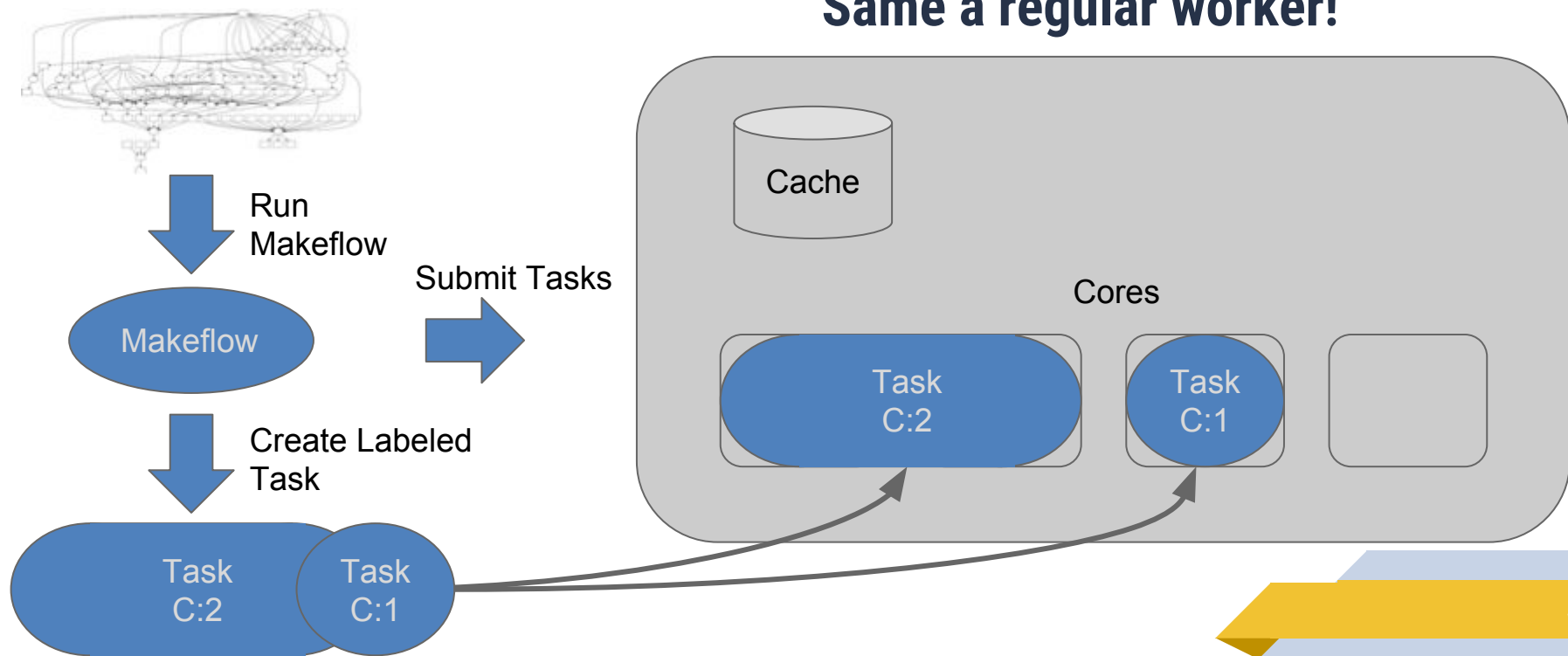
Submit Tasks

Same a regular worker!





Work Queue Multi-tenant Workers





Resource Monitor

- Watches process to ensure correct resource usage
- Evict jobs that act outside of resource allocation
- Report actual usage for future calibration
- Can be used in conjunction with Makeflow to automate an accurate image size.



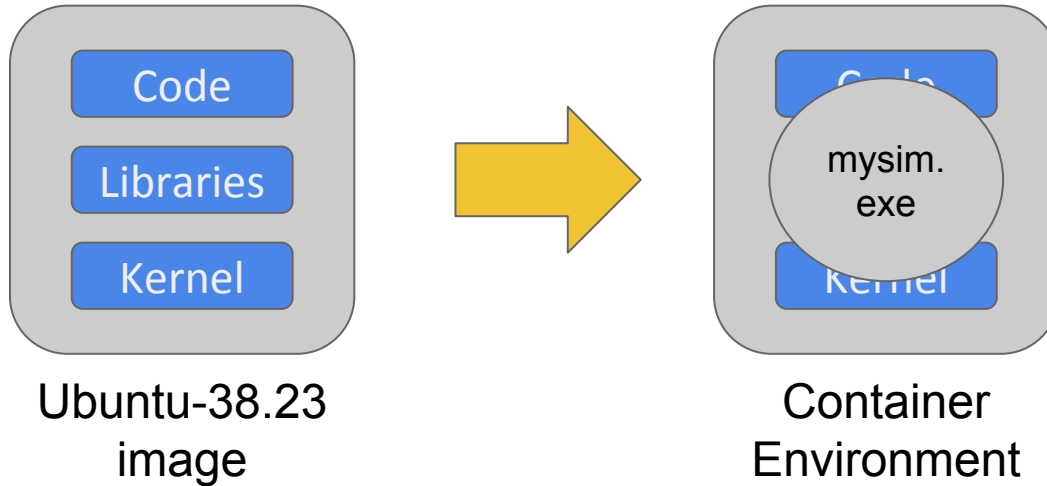
Container Integration

Providing consistent environments



Containers Create Precise Execution Environments

```
docker run ubuntu-38.23 mysim.exe
```



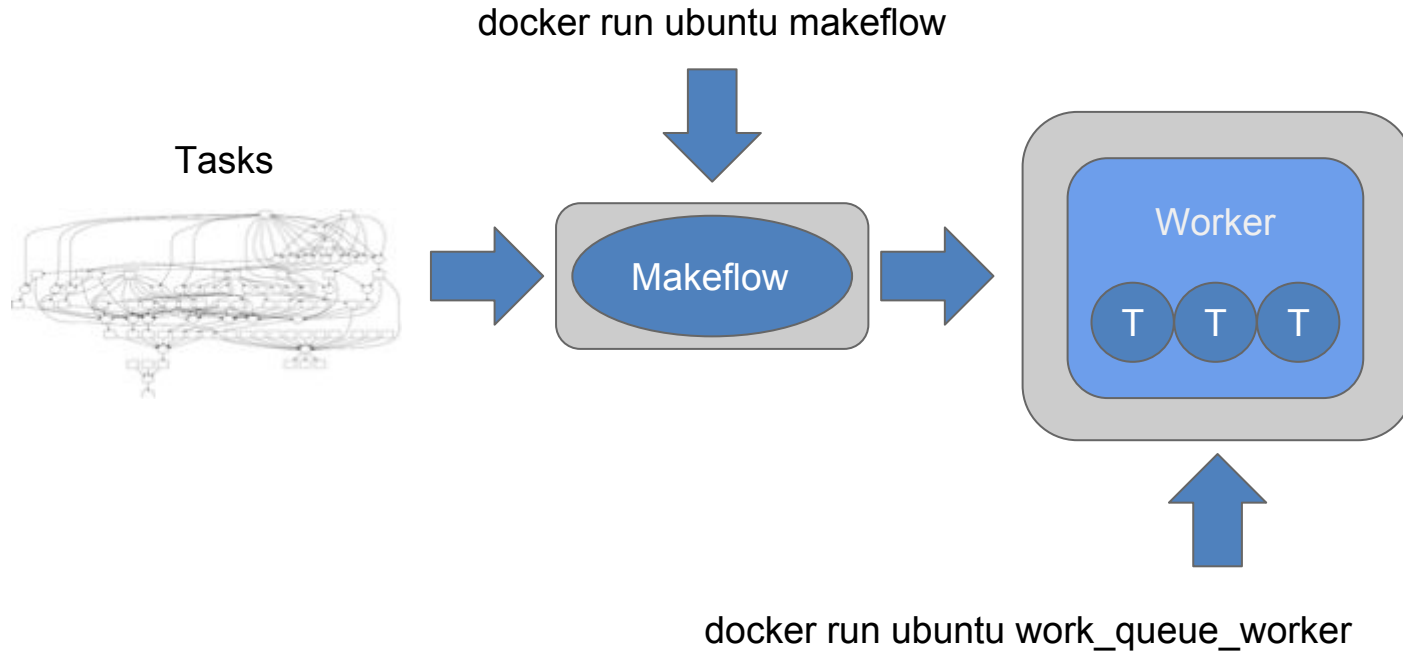


Approaches to Containers with Makeflow

- Approach 1:
 - Create containers for starting MF and WQ, then let them run as normal.
 - You are responsible for moving container images responsibly.
- Approach 2:
 - Let MF create containers as needed for each task.
 - Provides more control over moving container images.
 - Sending and starting up containers for each task.

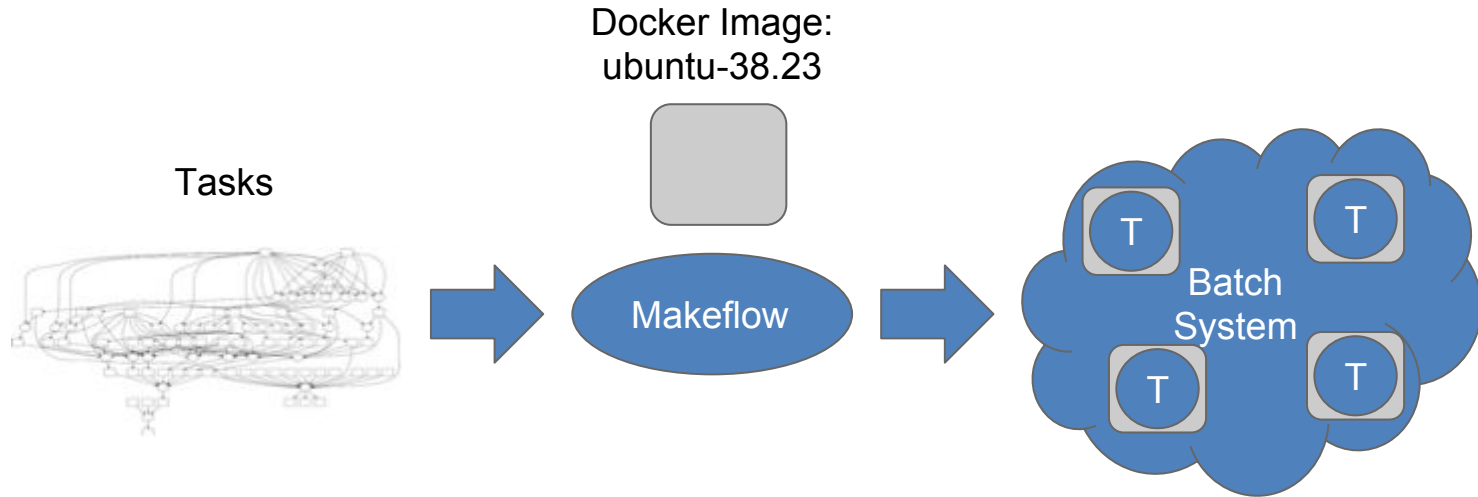


Approach 1: Container for MF/WQ





Approach 2: Container for Each Task



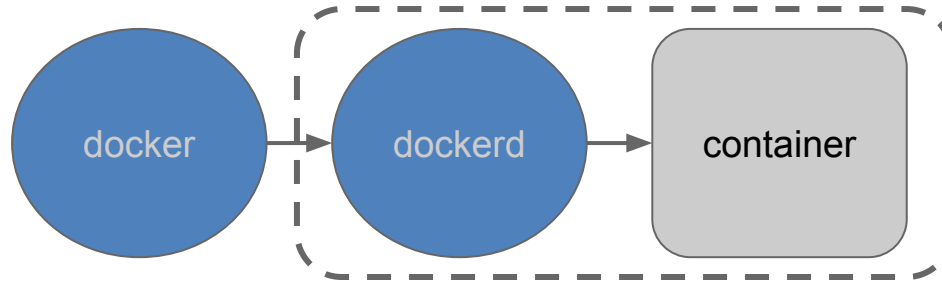
```
makeflow --docker ubuntu-38.23 -T sge . . .
```

Container Technology is Evolving



docker.io

docker run ubuntu command

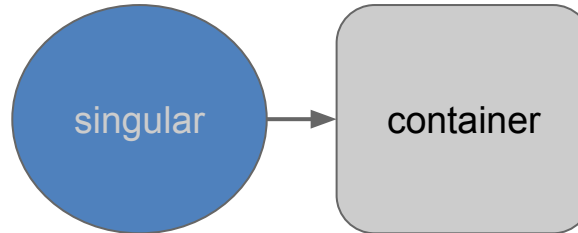


Installed service running as root



singularity.lbl.gov

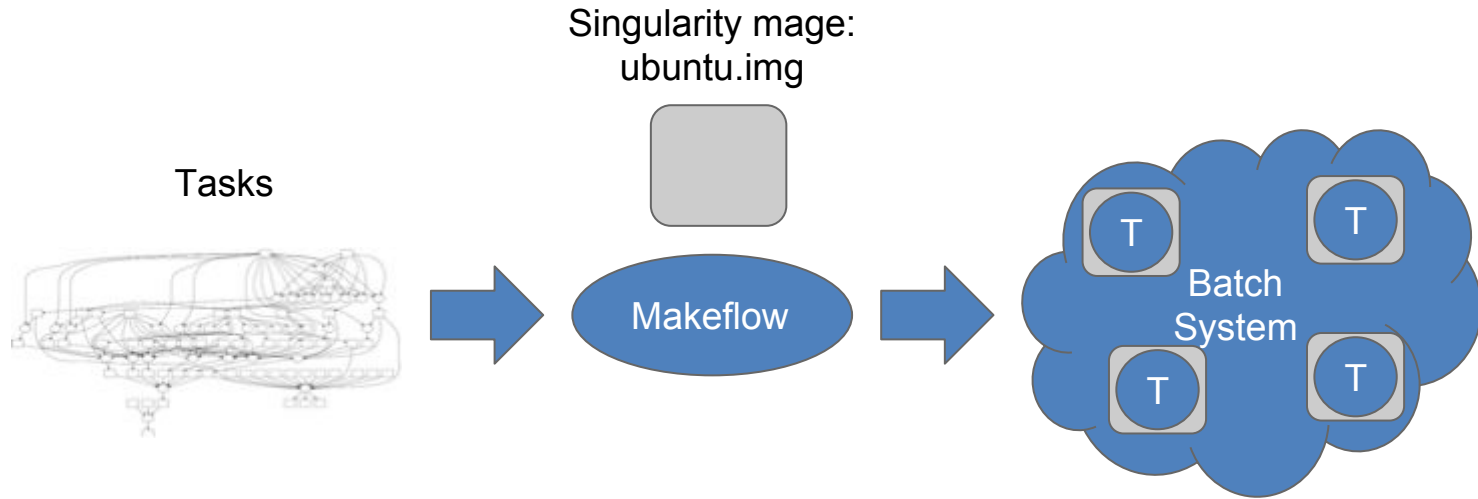
singularity exec ubuntu command



Container runs directly as a child process
(still needs setuid tool, though)



Approach 2 using Singularity



```
makeflow --singularity ubuntu.img -T sge ...
```



Cloud Operation

Methods to Deploying

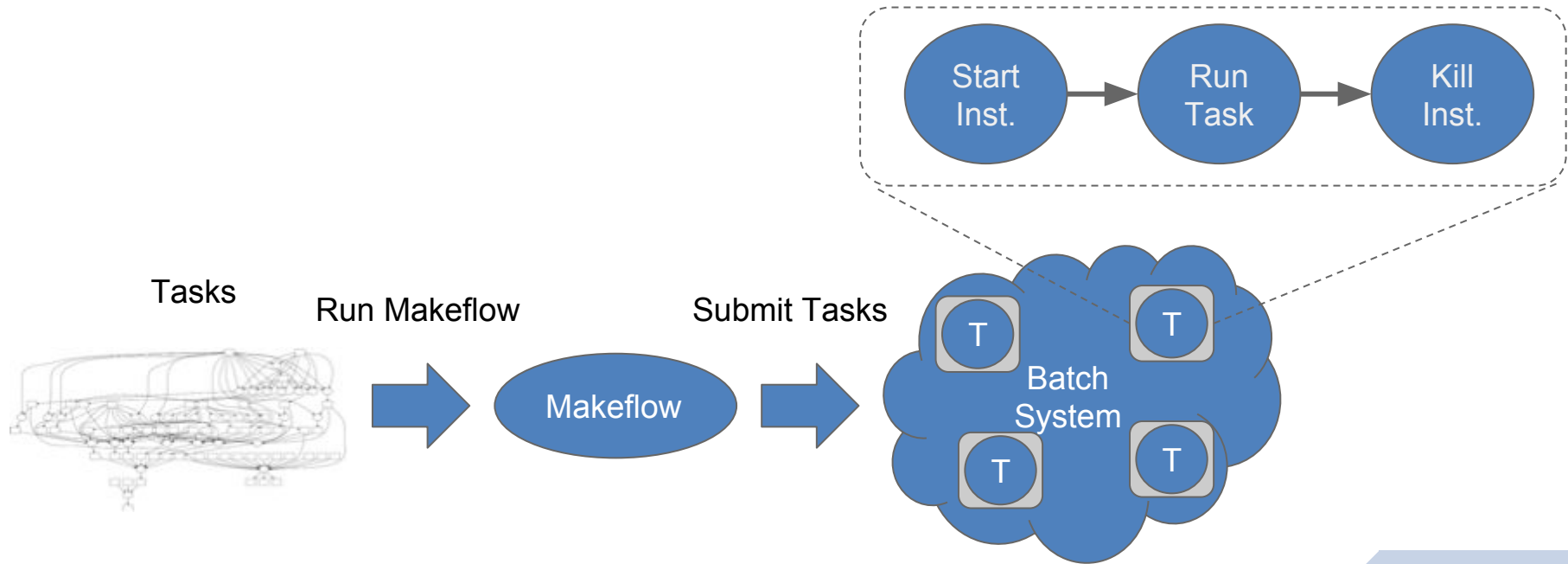


Approaches to Cloud Provisioning with Makeflow

- Approach 1:
 - MF creates unique instance for each task.
 - Provides complete isolation between tasks.
 - Requires startup and tear-down time of instances.
- Approach 2:
 - Create instances and run WQ Workers on them, submitting to WQ from MF.
 - Relies on WQ for task isolation, but caches shared files.
 - Instance management relies on the user.



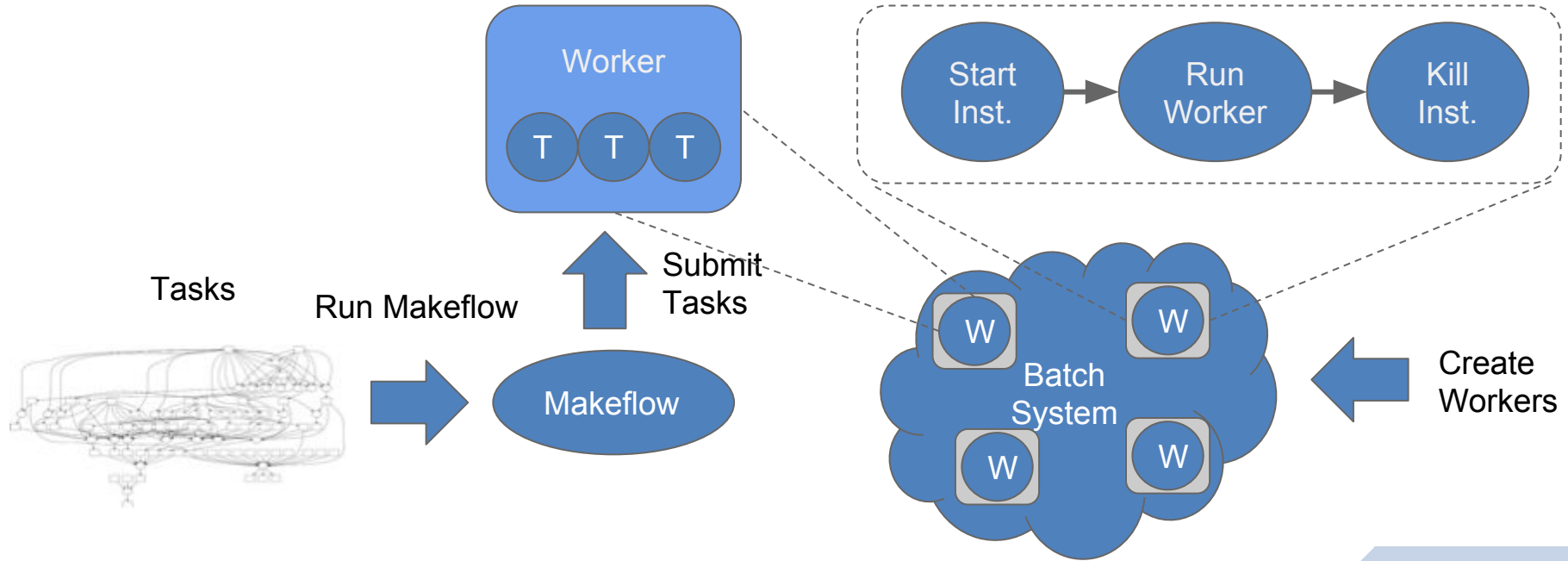
Approach 1: Individual instances per task



```
makeflow -T amazon --amazon-config my.config ...
```



Approach 2: Individual instances per worker



```
work_queue_factory -T amazon --amazon-config my.config
```



Questions?

Nick Hazekamp

Email : nhazekam@nd.edu

CCL Home : <http://ccl.cse.nd.edu>

Tutorial Link : <http://ccl.cse.nd.edu/software/tutorials/acic17>