

# Creating Custom Work Queue Applications



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# Makeflow vs. Work Queue

- Makeflow
  - Directed Acyclic Graph programming model.
  - Static structure known in advance.
  - All communication through files on disk.
- Work Queue
  - Submit-Wait programming model.
  - Dynamic structure decided at run-time.
  - Communicate through buffers or files.
  - More detailed knowledge of how tasks ran.

# Work Queue API

```
#include "work_queue.h"

queue = work_queue_create();

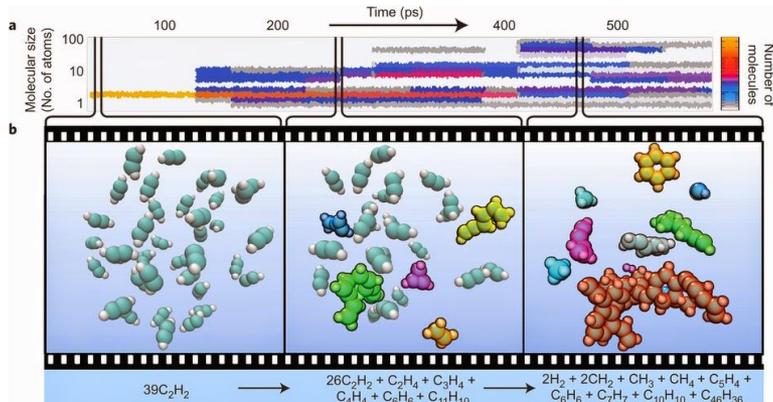
while( not done ) {
    while (more work ready) {
        task = work_queue_task_create();
        // add some details to the task
        work_queue_submit(queue, task);
    }

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

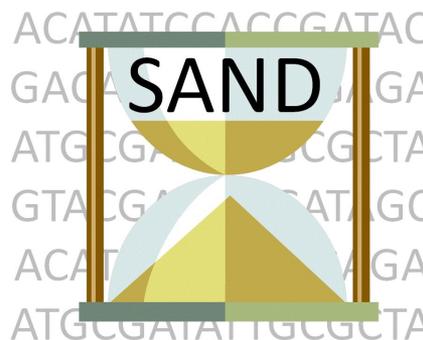
<http://ccl.cse.nd.edu/software/workqueue>

# Work Queue Applications

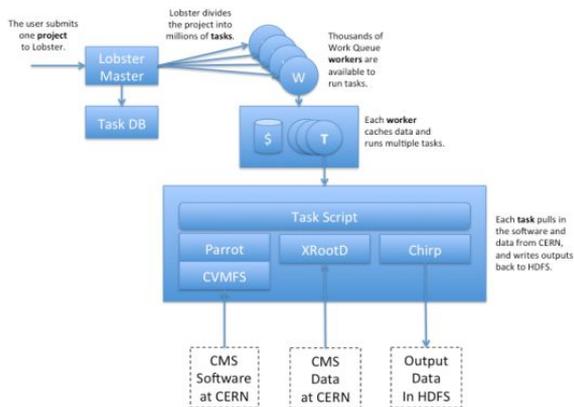
## Nanoreactor MD Simulations



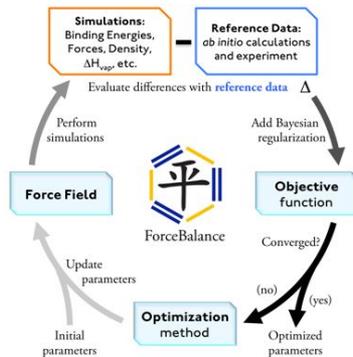
## Scalable Assembler at Notre Dame



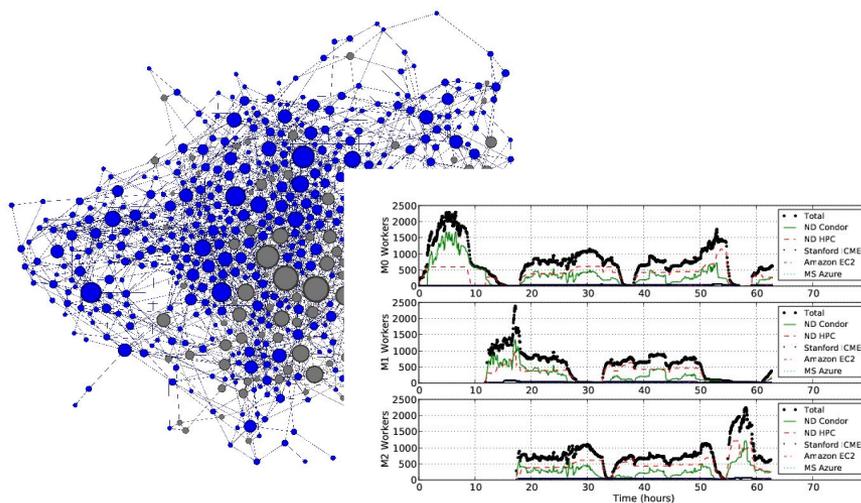
## Lobster HEP



## ForceBalance



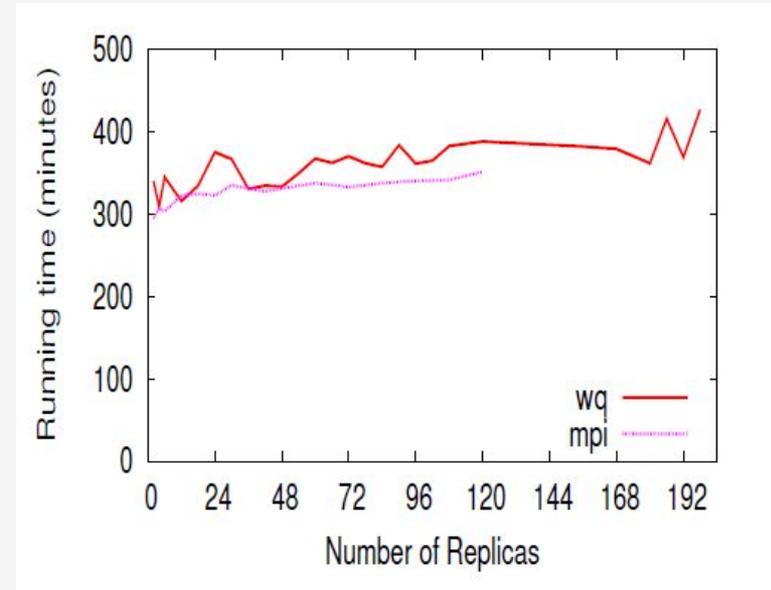
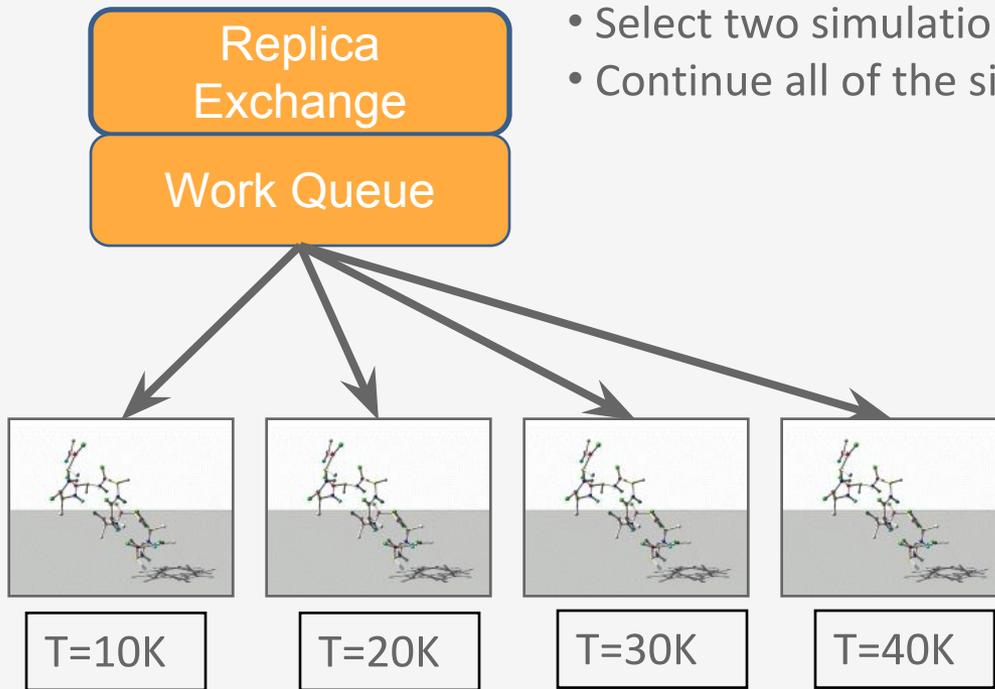
## Adaptive Weighted Ensemble



# Replica Exchange

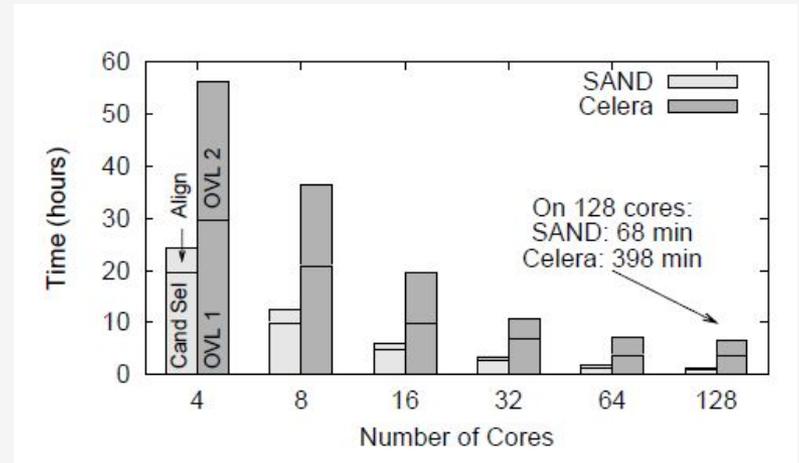
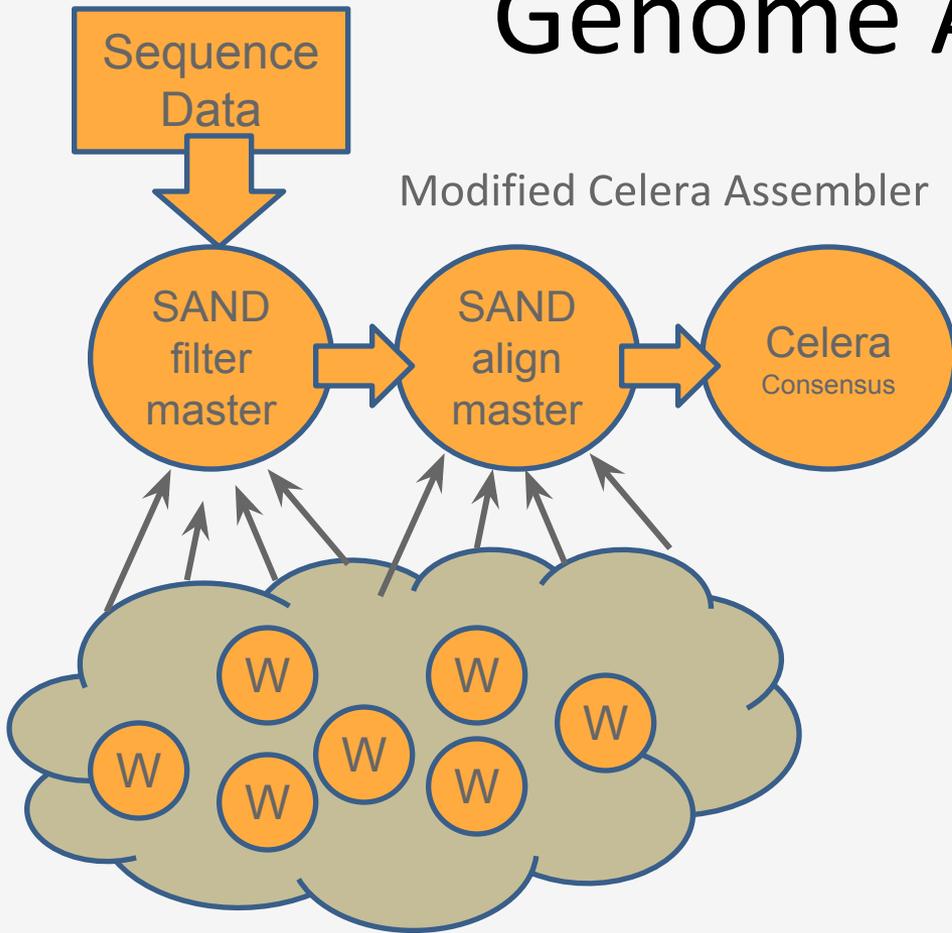
## Simplified Algorithm:

- Submit N short simulations at different temps.
- Wait for all to complete.
- Select two simulations to swap.
- Continue all of the simulations.



Dinesh Rajan, Anthony Canino, Jesus A Izaguirre, and Douglas Thain,  
Converting A High Performance Application to an Elastic Cloud Application, Cloud Com 2011.

# Genome Assembly

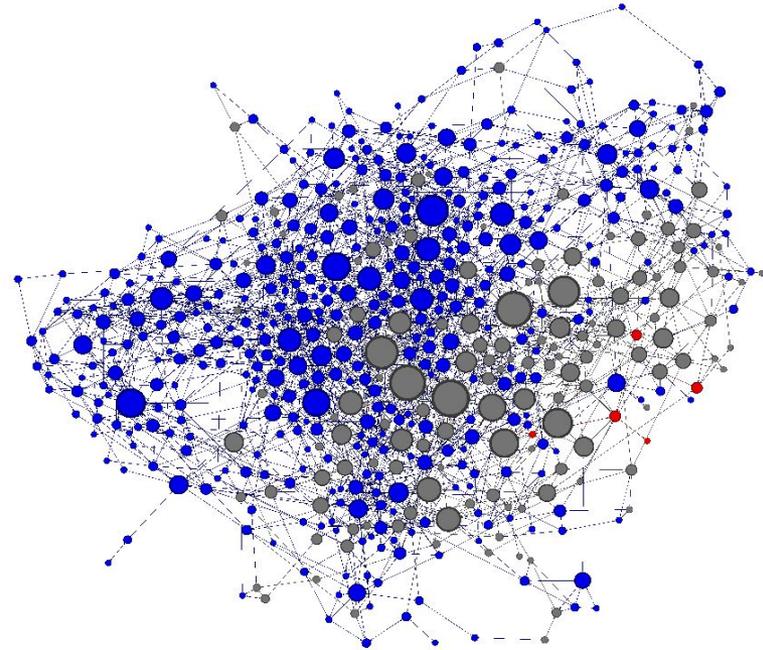
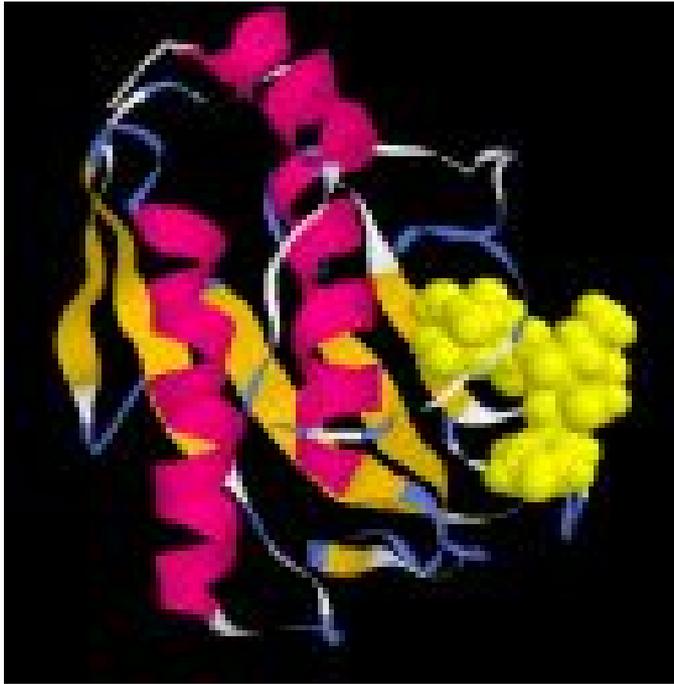


Using WQ, we could assemble a human genome in 2.5 hours on a collection of clusters, clouds, and grids with a speedup of 952X.

Christopher Moretti, Andrew Thrasher, Li Yu, Michael Olson, Scott Emrich, and Douglas Thain,  
**A Framework for Scalable Genome Assembly on Clusters, Clouds, and Grids,**  
*IEEE Transactions on Parallel and Distributed Systems*, 2012

# Adaptive Weighted Ensemble

Proteins fold into a number of distinctive states, each of which affects its function in the organism.

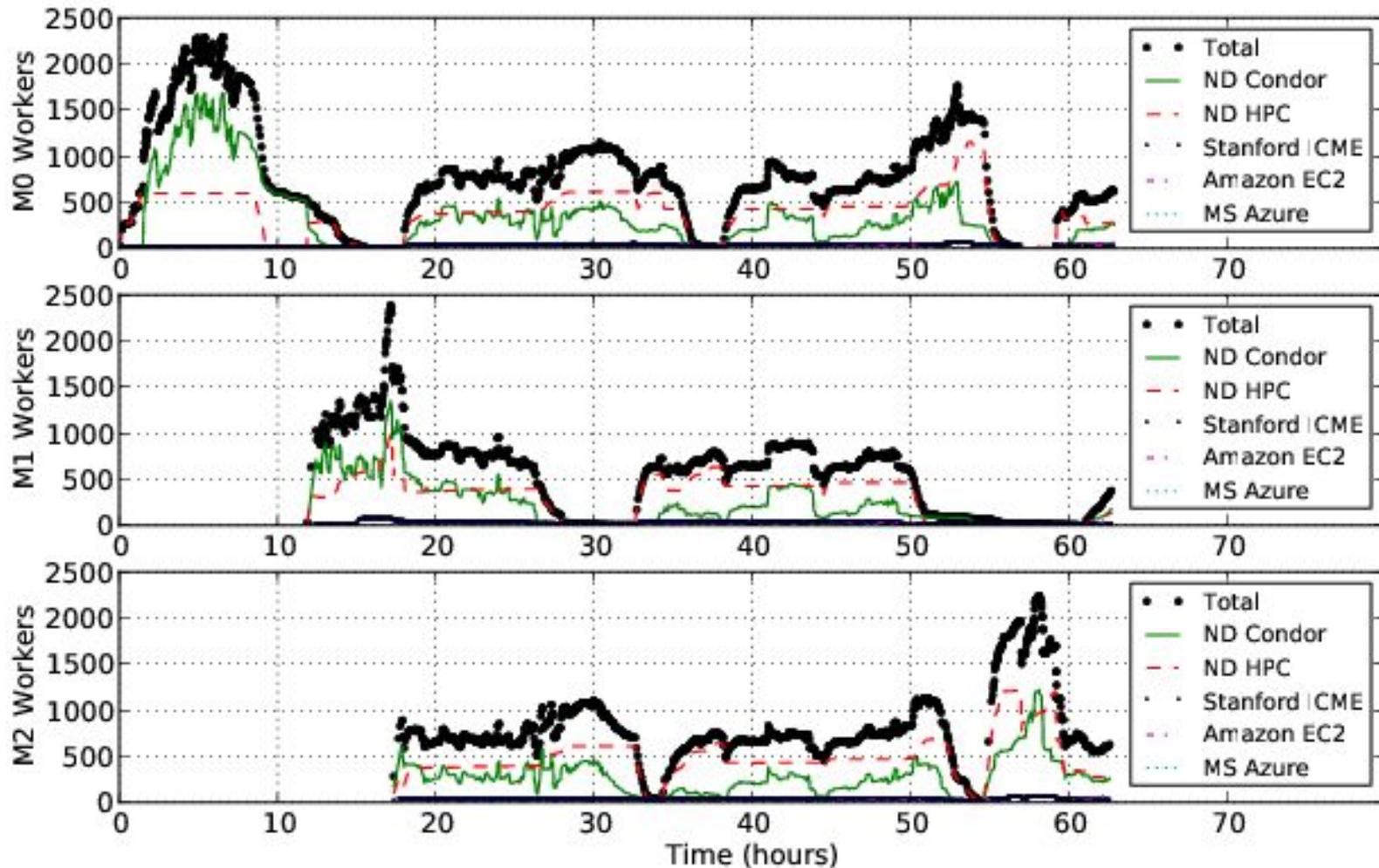


How common is each state?

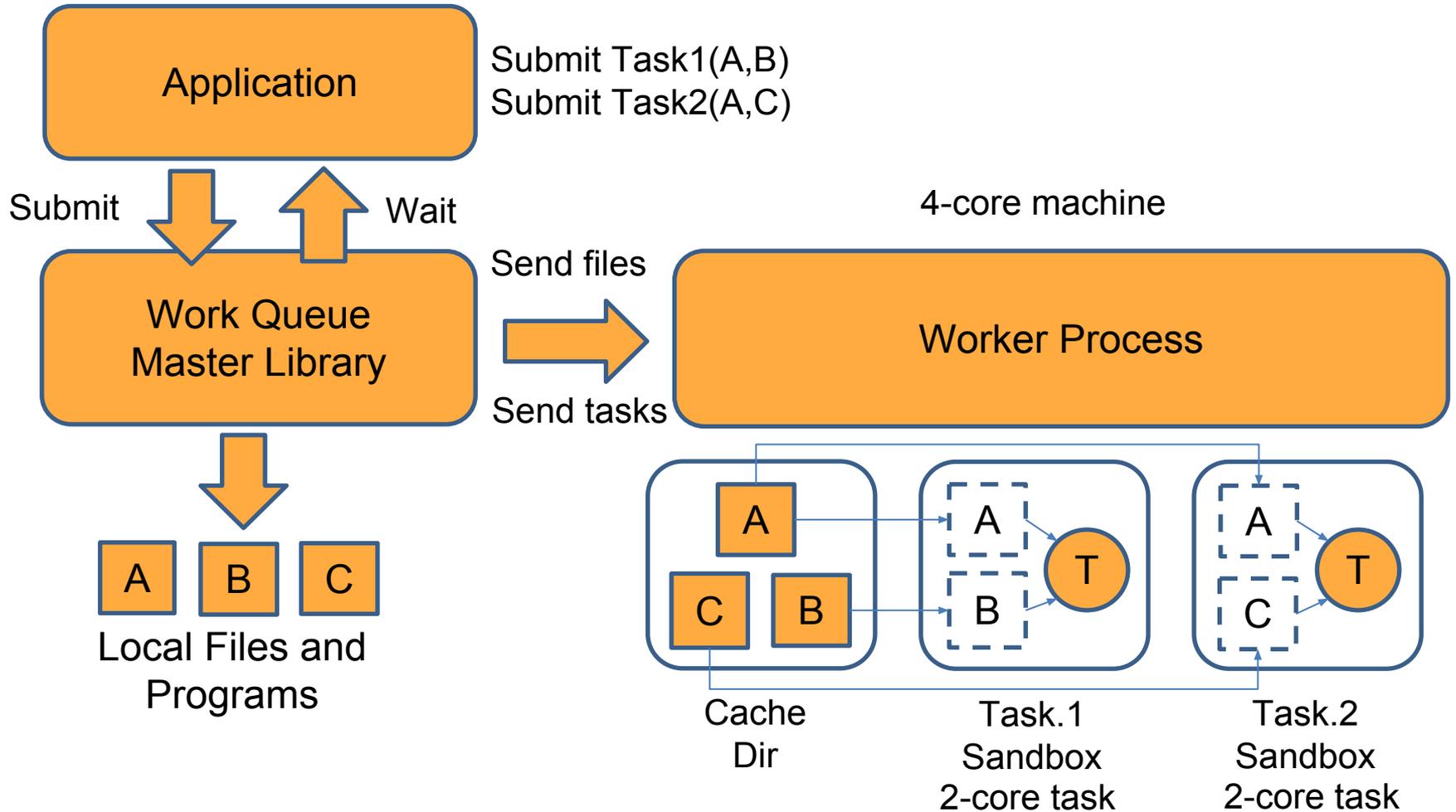
How does the protein transition between states?

How common are those transitions?

# AWE on Clusters, Clouds, and Grids



# Work Queue Architecture



# Basic Queue Operations

```
#include "work_queue.h"
struct work_queue *queue;
struct work_queue_task *task;

// Creates a new queue listening on a port, use zero to pick any port.
queue = work_queue_create( port );
// Submits a task into a queue. (non-blocking)
work_queue_submit( queue, task );
// Waits for a task to complete, returns the complete task.
task = work_queue_wait( queue, timeout );
// Returns true if there are no tasks left in the queue.
work_queue_empty( queue );
// Returns true if the queue is hungry for more tasks.
work_queue_hungry( queue );
```

# Basic Task Operations

```
#include "work_queue.h"
struct work_queue_task *task;

// Create a task that will run a given Unix command.
task = work_queue_task_create( command );

// Indicate an input or output file needed by the task.
work_queue_task_specify_file( task, name, remote_name, type, flags );

// Indicate an input buffer needed by the task.
work_queue_task_specify_buffer( task, data, length, remote_name, flags);

// Destroy the task object.
work_queue_task_delete( task );
```

# Run One Task in C

```
#include "work_queue.h"

struct work_queue *queue;
struct work_queue_task *task;

queue = work_queue_create( 0 );
work_queue_specify_name( "myproject" );

task = work_queue_task_create("sim.exe -p 50 in.dat >out.txt");
/// Missing: Specify files needed by the task.
work_queue_submit( queue, task );

while(!work_queue_empty(queue)) {
    task = work_queue_wait( queue, 60 );
    if(task) work_queue_task_delete( task );
}
```

# Run One Task in Perl

```
use work_queue;  
  
$queue = work_queue_create( 0 );  
  
work_queue_specify_name( "myproject" );  
  
$task = work_queue_task_create("sim.exe -p 50 in.dat >out.txt");  
### Missing: Specify files needed by the task.  
  
work_queue_submit( $queue, $task );  
  
while(!work_queue_empty($queue)) {  
    $task = work_queue_wait( $queue, 60 );  
    if($task) work_queue_task_delete( $task );  
}
```

# Run One Task in Python

```
from work_queue import *

queue = WorkQueue( port = 0 )

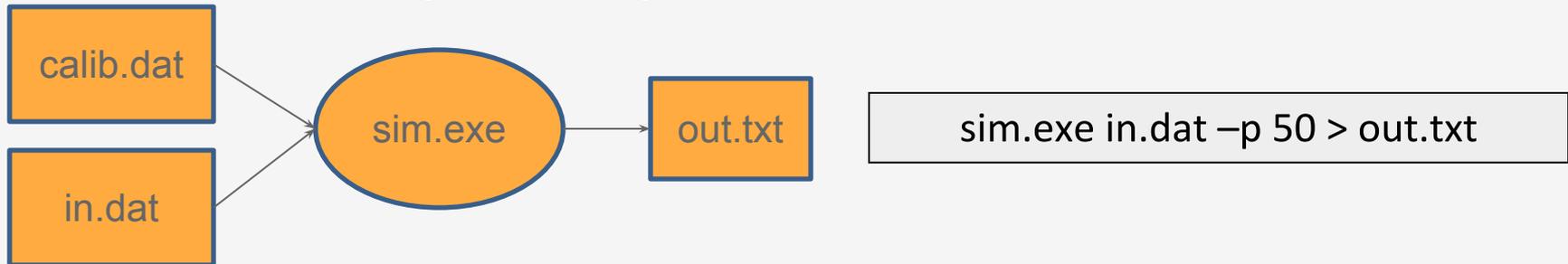
queue.specify_name( "myproject" );

task = Task("sim.exe -p 50 in.dat >out.txt")

### Missing: Specify files needed by the task.
queue.submit( task )

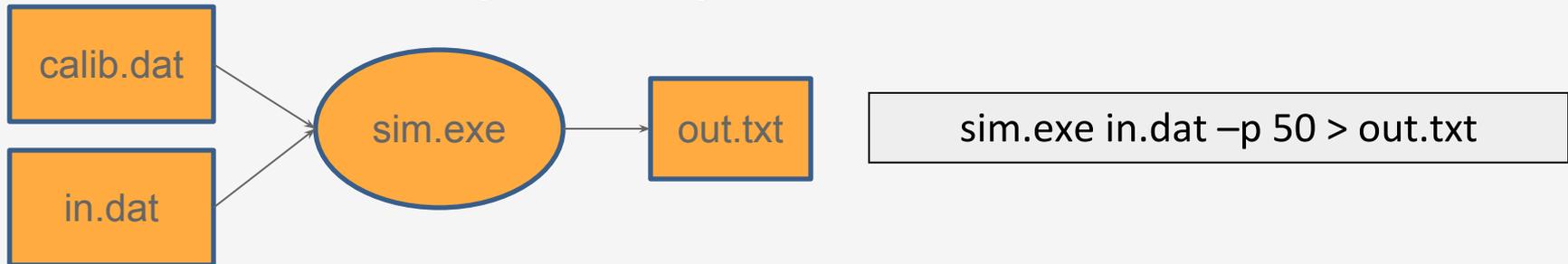
While not queue.empty():
    task = queue.wait(60)
```

# C: Specify Files for a Task



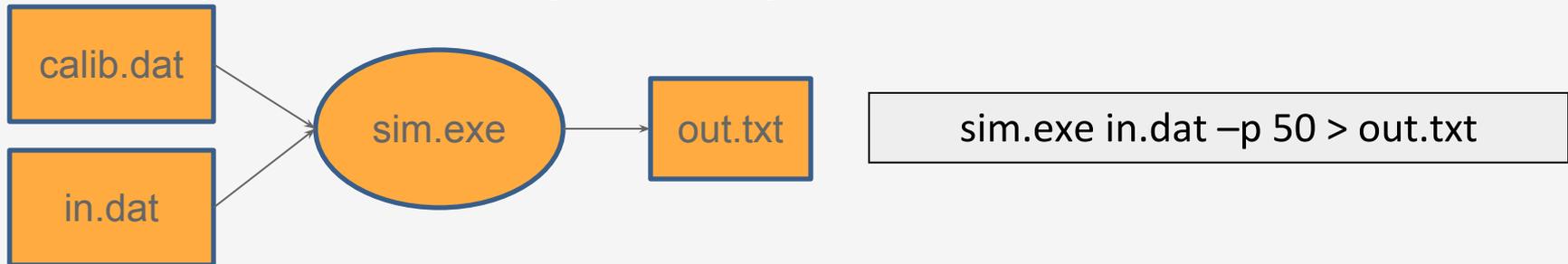
```
work_queue_task_specify_file( $task, "in.dat", "in.dat",  
    $WORK_QUEUE_INPUT, $WORK_QUEUE_NOCACHE );  
  
work_queue_task_specify_file($task, "calib.dat", "calib.dat",  
    $WORK_QUEUE_INPUT, $WORK_QUEUE_NOCACHE );  
  
work_queue_task_specify_file( $task, "out.txt", "out.txt",  
    $WORK_QUEUE_OUTPUT, $WORK_QUEUE_NOCACHE );  
  
work_queue_task_specify_file( $task, "sim.exe", "sim.exe",  
    $WORK_QUEUE_INPUT, $WORK_QUEUE_CACHE );
```

# Perl: Specify Files for a Task



```
work_queue_task_specify_file( $task, "in.dat", "in.dat",  
    $WORK_QUEUE_INPUT, $WORK_QUEUE_NOCACHE );  
  
work_queue_task_specify_file($task, "calib.dat", "calib.dat",  
    $WORK_QUEUE_INPUT, $WORK_QUEUE_NOCACHE );  
  
work_queue_task_specify_file( $task, "out.txt", "out.txt",  
    $WORK_QUEUE_OUTPUT, $WORK_QUEUE_NOCACHE );  
  
work_queue_task_specify_file( $task, "sim.exe", "sim.exe",  
    $WORK_QUEUE_INPUT, $WORK_QUEUE_CACHE );
```

# Python: Specify Files for a Task



```
task.specify_file( "in.dat", "in.dat",  
                  WORK_QUEUE_INPUT, cache = False )  
  
task.specify_file( "calib.dat", "calib.dat",  
                  WORK_QUEUE_INPUT, cache = False )  
  
task.specify_file( "out.txt", "out.txt",  
                  WORK_QUEUE_OUTPUT, cache = False )  
  
task.specify_file( "sim.exe", "sim.exe",  
                  WORK_QUEUE_INPUT, cache = True )
```



You must state  
all the files  
needed by the command.

# Running a Work Queue Program

```
gcc work_queue_example.c -o work_queue_example  
-I $HOME/cctools/include/cctools  
-L $HOME/cctools/lib  
-lwork_queue -ldttools -lm
```

```
./work_queue_example  
Listening on port 8374 ...
```

In another window:

```
./work_queue_worker master.host.name.org 8374
```



# ... for Perl

```
setenv PERL5LIB ${PERL5LIB}: (no line break)  
    ${HOME}/cctools/lib/perl5/site_perl
```

```
./work_queue_example.pl  
Listening on port 8374 ...
```

In another window:

```
./work_queue_worker master.host.name.org 8374
```



# ... for Python

```
setenv PYTHONPATH ${PYTHONPATH}: (no line break)  
    ${HOME}/cctools/lib/python2.6/site-package
```

```
./work_queue_example.py
```

```
Listening on port 8374 ...
```

In another window:

```
./work_queue_worker master.host.name.org 8374
```



# Start Workers Everywhere

Submit workers to Condor:

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

Submit workers to SGE:

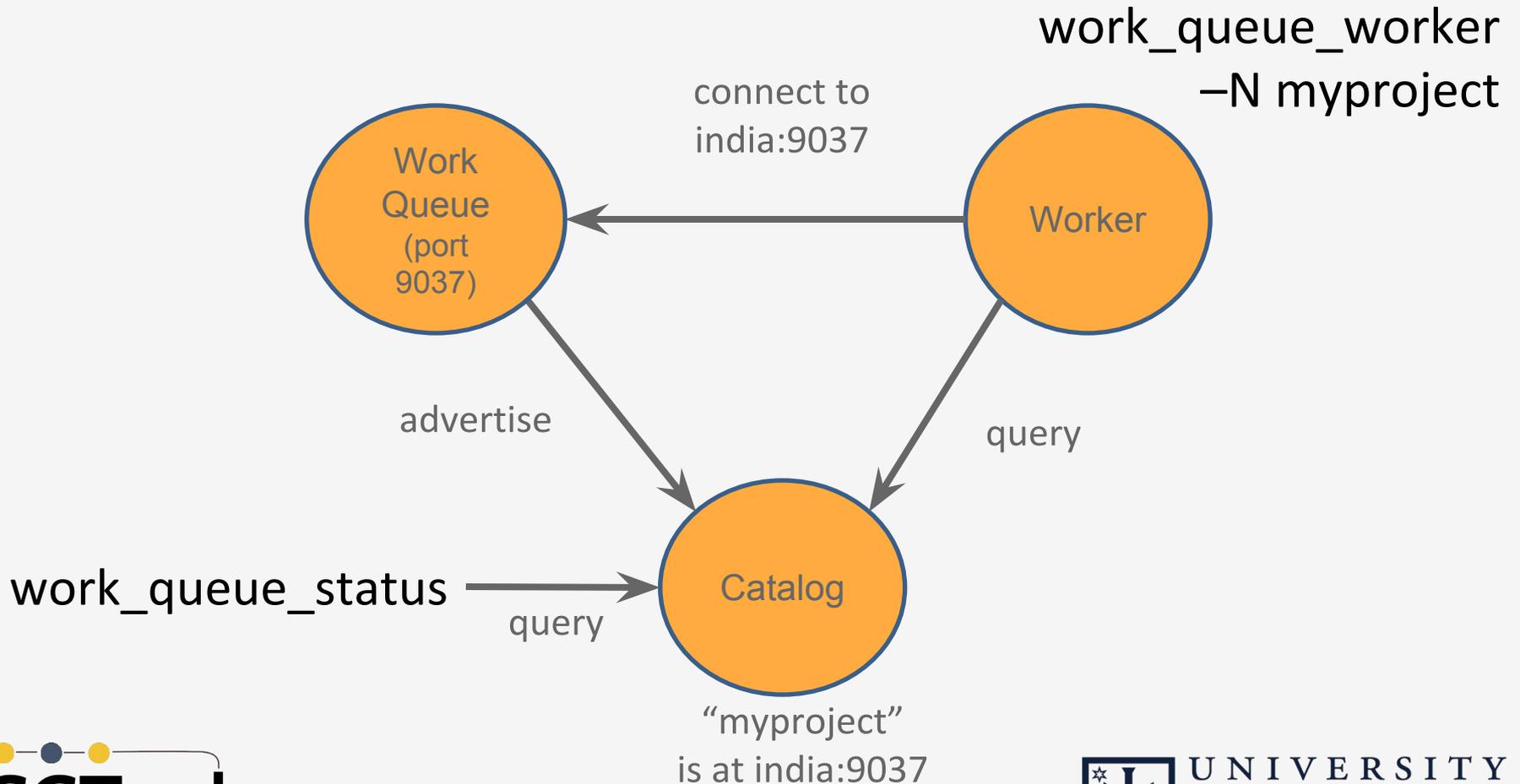
```
sgesubmit_workers master.hostname.org 8374 25
```

Submit workers to Torque:

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



# Use Project Names



# Specify Project Names in Work Queue

Specify Project Name for Work Queue master:

C:

```
work_queue_specify_name (q, "myproject");
```

Perl:

```
work_queue_specify_name ($q, "myproject");
```

Python:

```
q.specify_name ("myproject")
```

# Start Workers with Project Names

Start one worker:

```
$ work_queue_worker -N myproject
```

Start many workers:

```
$ sge_submit_workers -N myproject 5
```

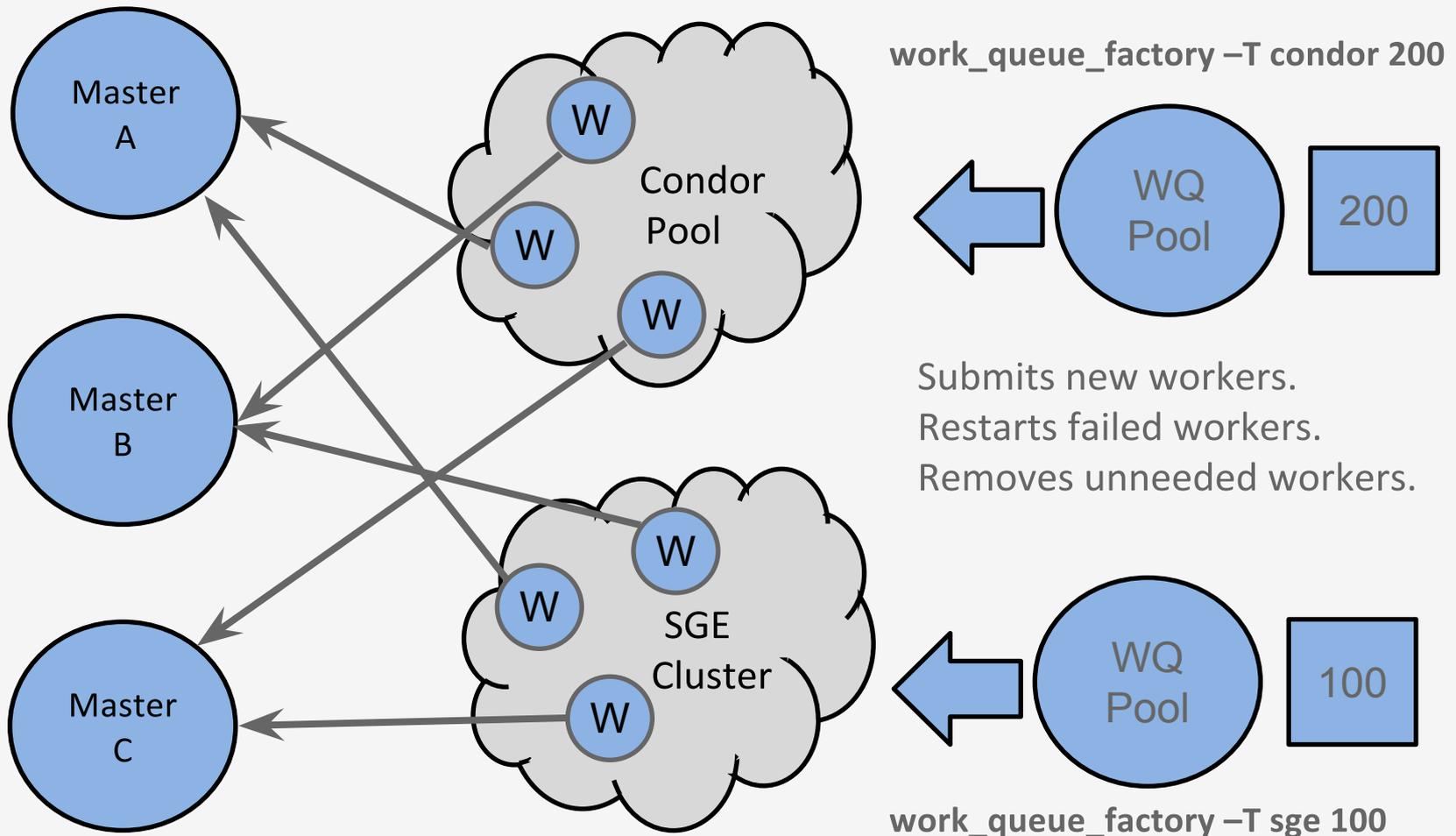
```
$ condor_submit_workers -N myproject 5
```

```
$ torque_submit_workers -N myproject 5
```

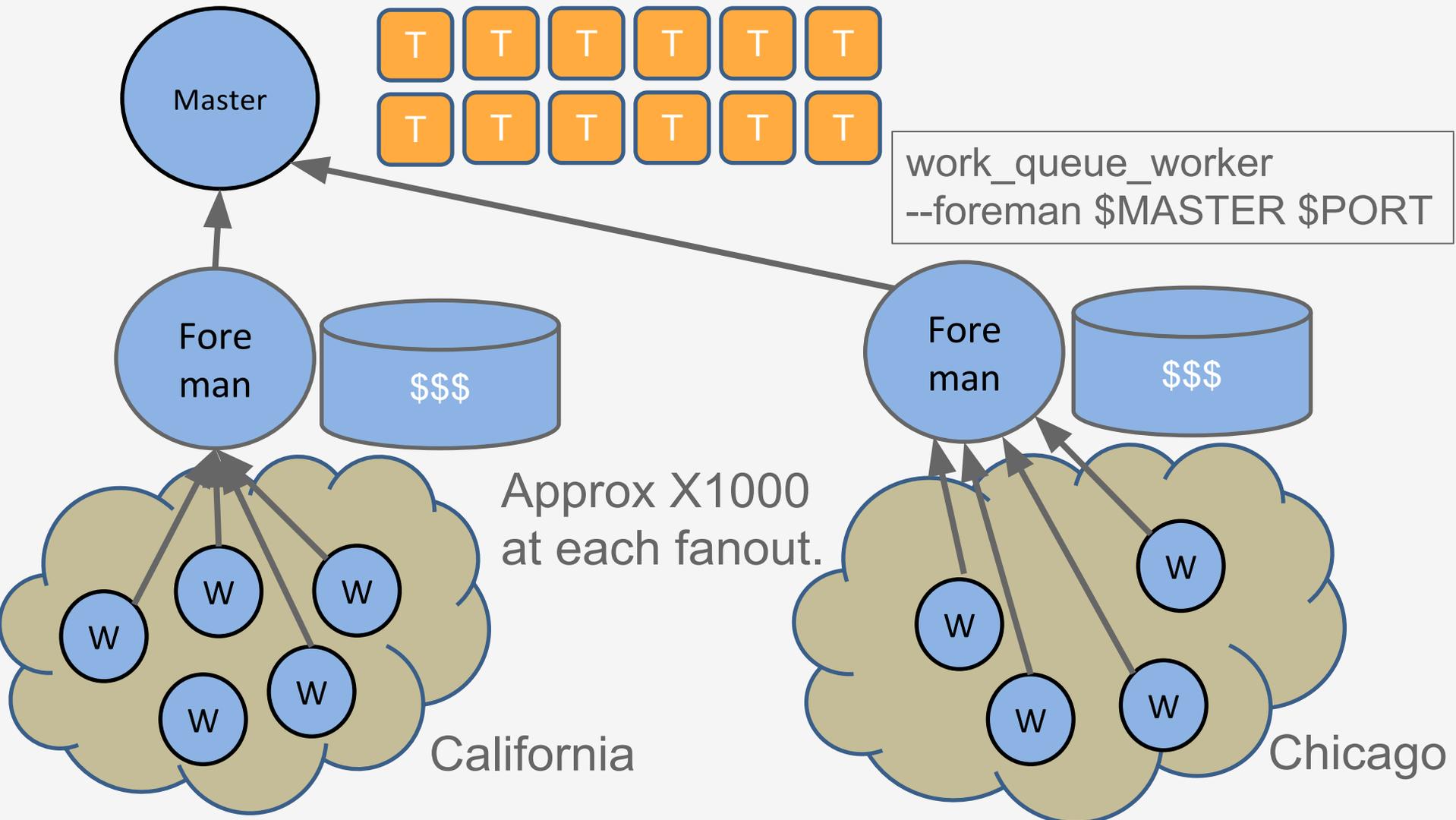
# Advanced Features (in the docs)

- Submit / remove tasks by tag / name.
- Auto reschedule tasks that take too long.
- Send in-memory data as a file.
- Log and graph system performance
- Much more!

# Managing Your Workforce



# Using Foremen



# Multi-Slot Workers

