

Profiling Applications to Choose the Right Computing Infrastructure plus Batch Management with HTCondor

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Follow Along at:

https://opensciencegrid.github.io/dosar/Materials/ Materials/





Some thoughts on the exercises

- It's okay to move ahead on exercises if you have time
- It's okay to take longer on them if you need to
- If you move along quickly, try the "On Your Own" sections and "Challenges"





Most important!

- Please ask me questions!
 - ...during the lectures
 - ...during the exercises
 - ...during the breaks
 - ...during the meals
 - ...over dinner
 - ...by email after we depart (kagross@iu.edu)
- If I don't know, I'll find the right person to answer your question.





Goals for this session

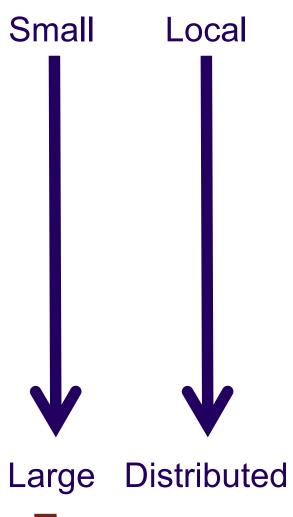
- Profiling your application
- Picking the appropriate resources
- Understand the basics of HTCondor







Let's take one step at a time



- Can you run one job on one computer?
- Can you run one job on another computer?
- Can you run 10 jobs on a set of computers?
- Can you run a multiple job workflow?
- How do we put this all together?

This is the path we'll take in the school



What does the user provide?

- A "headless job"
 - Not interactive/no GUI: how could you interact with 1000 simultaneous jobs?
- A set of input files
- A set of output files
- A set of parameters (command-line arguments)
- Requirements:
 - Ex: My job requires at least 2GB of RAM
 - Ex: My job requires Linux
- Control/Policy:
 - Ex: Send me email when the job is done
 - Ex: Job 2 is more important than Job 1
 - Ex: Kill my job if it runs for more than 6 hours



What does the system provide?

Methods to:

- Submit/Cancel job
- Check on state of job
- Check on state of available computers

Processes to:

- Reliably track set of submitted jobs
- Reliably track set of available computers
- Decide which job runs on which computer
- Manage a single computer
- Start up a single job



Gedankenexperiment

- Let's assume you have a 'large job'
 - What factors could make it large?
- Large Data Input or Output or both
- Needs to do heavy calculation
- Needs a lot of memory
- Needs to communicate with other jobs (whether required or not)
- Reads and writes a lot of data/files
- Heavy graphics processing
- Any combination of any of the above



There is no "One Size Fits All Solution"

- But some solutions are more "Open" than others.
 - Local Laptop/Desktop
 - Local Cluster
 - HPC System
 - Shared HTC Resources
 - Clouds





Why is HTC hard?

- The HTC system has to keep track of:
 - Individual tasks (a.k.a. jobs) & their inputs
 - Computers that are available
- The system has to recover from failures
 - There will be failures! Distributed computers means more chances for failures.
- You have to share computers
 - Sharing can be within an organization, or between orgs
 - So you have to worry about security
 - And you have to worry about policies on how you share
- If you use a lot of computers, you have to handle variety:
 - Different kinds of computers (arch, OS, speed, etc..)
 - Different kinds of storage (access methodology, size, speed, etc...)
 - Different networks interacting (network problems are hard to debug!)





Surprise! HTCondor does this (and more)

Methods to:

- Submit/Cancel job. condor_submit/condor_rm
- Check on state of job. condor_q
- Check on state of avail. computers. condor_status

Processes to:

- Reliably track set of submitted jobs. schedd
- Reliably track set of avail. computers. collector
- Decide which job runs on where. negotiator
- Manage a single computer startd
- Start up a single job starter



But not only Condor

- You can use other systems:
 - PBS/Torque
 - Oracle Grid Engine (né Sun Grid Engine)
 - LSF
 - SLURM
 - **–** ...
- But I won't cover them.
 - My experience is with Condor
 - My bias is with Condor
 - Overlays exist
- What should you learn at the school?
 - How do you think about Computing Resources?
 - How can you do your science with HTC?
 - For now, learn it with Condor, but you can apply it to other systems.





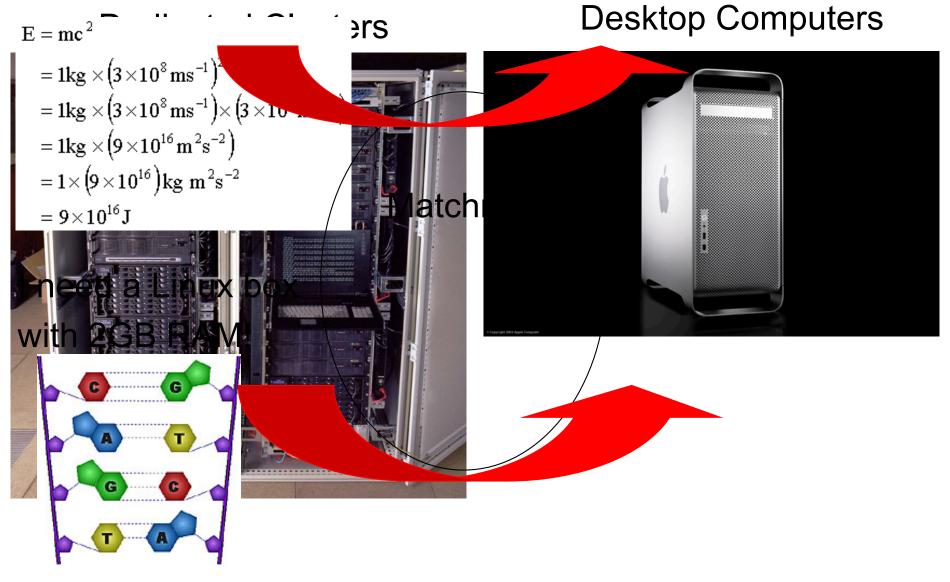
A brief introduction to Condor

- Please note, we will only scratch the surface of Condor:
 - We won't cover MPI, Master-Worker, advanced policies, site administration, security mechanisms, submission to other batch systems, virtual machines, cron, high-availability, computing on demand, containers.





I need a Mac!





Quick Terminology

- Cluster: A dedicated set of computers not for interactive use
- Pool: A collection of computers used by Condor
 - May be dedicated
 - May be interactive
- Remember:
 - Condor can manage a cluster in a machine room
 - Condor can use desktop computers
 - Condor can access remote computers
 - HTC uses all available resources



Matchmaking

- Matchmaking is fundamental to Condor
- Matchmaking is two-way
 - Job describes what it requires:
 - I need Linux && 8 GB of RAM
 - Machine describes what it requires:
 - I will only run jobs from the Physics department
- Matchmaking allows preferences
 - I need Linux, and I prefer machines with more memory but will run on any machine you provide me





Why Two-way Matching?

- Condor conceptually divides people into three groups:
 - Job submitters
 - Computer owners
 - Pool (cluster) administrator
- All three of these groups have preferences

May or may not be the same people





ClassAds

- ClassAds state facts
 - My job's executable is analysis.exe
 - My machine's load average is 5.6
- ClassAds state preferences
 - I require a computer with Linux
- ClassAds are extensible
 - They say whatever you want them to say







Example ClassAd

```
MyType = "Job" ←—String
TargetType = "Machine"
ClusterId = 1377 ← Number
Owner = "roy"
         = "analysis.exe"
Cmd
Requirements =  ← Expression
   (Arch == "INTEL")
&& (OpSys == "LINUX")
&& (Disk >= DiskUsage)
&& ((Memory * 1024)>=ImageSize)
```



Schema-free ClassAds

- Condor imposes some schema
 - Owner is a string, ClusterID is a number...
- But users can extend it however they like, for jobs or machines
 - AnalysisJobType = "simulation"
 - HasJava 1 6 = TRUE
 - ShoeLength = 10
- Matchmaking can use these attributes
 - Requirements = OpSys == "LINUX" && HasJava_1_6 == TRUE





Don't worry

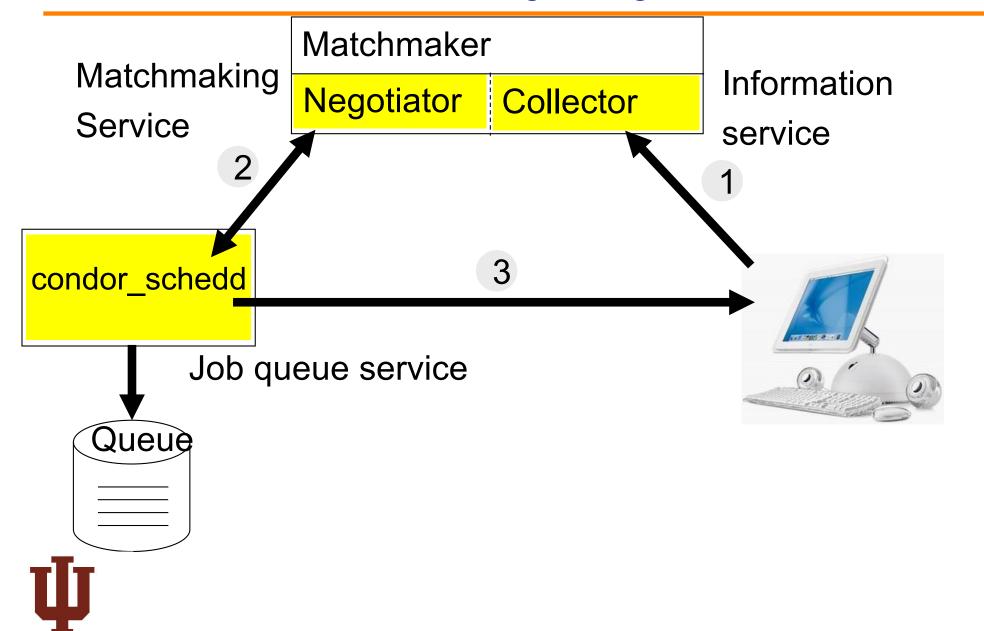
- You won't write ClassAds (usually)
 - You'll create a simple submit file
 - Condor will write the ClassAd
 - You can extend the ClassAd if you want to
- You won't write requirements (usually)
 - Condor writes them for you
 - You can extend them
 - In some environments you provide attributes instead of requirements expressions





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Matchmaking diagram





Why do jobs fail?

- The computer running the job fails
 - Or the network, or the disk, or the OS, or...
- Your job might be preempted:
 - Condor decides your job is less important than another, so your job is stopped and another started.
 - Not a "failure" per se, but it may feel like it to you.





Reliability

- When a job fails or is preempted:
 - It stays in the queue (on the schedd)
 - A note is written to the job log file
 - It reverts to "idle" state
 - It is eligible to be matched again
- Relax! Condor will run your job again





Access to data in Condor

- Option #1: Shared filesystem
 - Simple to use, but make sure your filesystem can handle the load
- Option #2: Condor's file transfer
 - Can automatically send back changed files
 - Atomic transfer of multiple files
 - Can be encrypted over the wire
 - Most common for small applications and data
- Option #3: Remote I/O



Condor File Transfer

- ShouldTransferFiles = YES
 - Always transfer files to execution site
- ShouldTransferFiles = NO
 - Rely on a shared filesystem
- ShouldTransferFiles = IF NEEDED
 - Will automatically transfer the files if needed

```
Universe = vanilla
Executable = my_job
Log = my_job.log
ShouldTransferFiles = YES
Transfer_input_files = dataset$(Process), common.data
Queue 600
```



Condor File Transfer with URLs

 Transfer_input_files can be a URL For example:

```
transfer input files = http://www.example.com/input.data
```





Clusters & Processes

- One submit file can describe lots of jobs
 - All the jobs in a submit file are a cluster of jobs
 - Yeah, same term as a cluster of computers
- Each cluster has a unique "cluster number"
- Each job in a cluster is called a "process"
- A Condor "job ID" is the cluster number, a period, and the process number ("20.1")
- A cluster is allowed to have one or more processes.
 - There is always a cluster for every job





The \$(Process) macro

- The initial directory for each job can be specified as run_\$(Process), and instead of submitting a single job, we use "Queue 600" to submit 600 jobs at once
- The \$(Process) macro will be expanded to the process number for each job in the cluster (0 599), so we'll have "run_0", "run_1", ...
 "run_599" directories
- All the input/output files will be in different directories!





Example of \$(Process)

```
# Example condor_submit input file that defines
# a cluster of 600 jobs with different directories
Universe = vanilla
Executable = my_job
Log = my_job.log
Arguments = -arg1 -arg2
Input = my_job.stdin
Output = my_job.stdout
Error = my_job.stderr
InitialDir = run_$(Process)
Queue 600
Creates job 3.0 ... 3.599
```





More \$(Process)

You can use \$(Process) anywhere:

```
Universe = vanilla
Executable = my_job
Log = my_job.$(Process).log
Arguments = -randomseed $(Process)
Input = my_job.stdin
Output = my_job.stdout
Error = my_job.stderr
InitialDir = run_$(Process)
Queue 600
```





Sharing a directory

- You don't have to use separate directories.
- \$(Cluster) will help distinguish runs

```
Universe = vanilla
Executable = my_job
Arguments = -randomseed $(Process)
Input = my_job.input.$(Process)
Output = my_job.stdout.$(Cluster).$(Process)
Error = my_job.stderr.$(Cluster).$(Process)
Log = my_job.$(Cluster).$(Process).log
Oueue 600
```





Not Only Programming Language

- You ran a C program earlier
- You can also run scripting languages such as bash, python, and perl
- You can also executing programs via the command like R





Day One Wrap Up Notes

- There are several different computing environments
- There is a very diverse set of computing jobs
- Matching jobs to resources is key to not wasting resources
- Not all of the available environments are open environments
- Research Computing is Complex



Quick UNIX Refresher Before We Start

- \$
- nano, vi, emacs, cat >, etc.
- source, module, chmod, ls





That was a whirlwind tour!

- Enough with the presentation: let's do some computing!
- Goal: Extend the diversity of our jobs and add some data to the mix.







Questions?

- Questions? Comments?
 - Feel free to ask me questions now or later:

Kyle Gross – <u>kagross@iu.edu</u>

Exercises start here:

https://opensciencegrid.github.io/dosar/Materials/



Presentations are also available from this URL