AN INTRODUCTION TO USING HTCOndor

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University of Wisconsin-Madison Center for High Throughput Computing (CHTC)







Agenda

- Today:
 - Introduction for users
 - Also useful for administrators ©
- > Plan for Tomorrow:
 - Session One (9:30am-11:00am)
 - Architecture/Administration Overview (90min)
 - Session Two (11:30am-1:00pm)
 - Open Question/Answer Session (45 min)
 - Monitoring (20 min)
 - What's New and What's Coming Up? (20 min)





Introduction

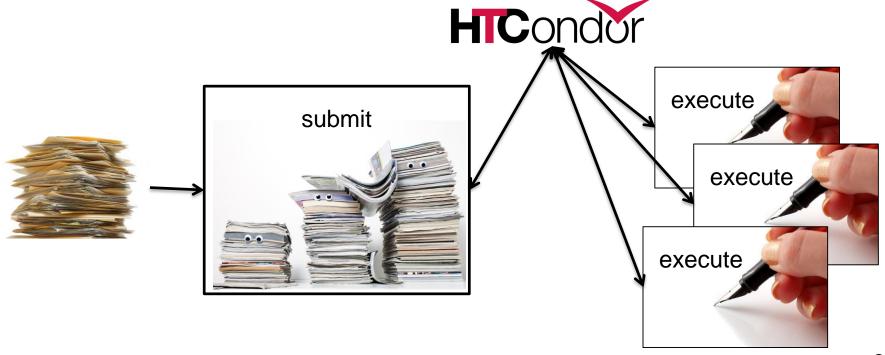
What is HTCondor?

 Software that schedules and runs computing tasks on computers

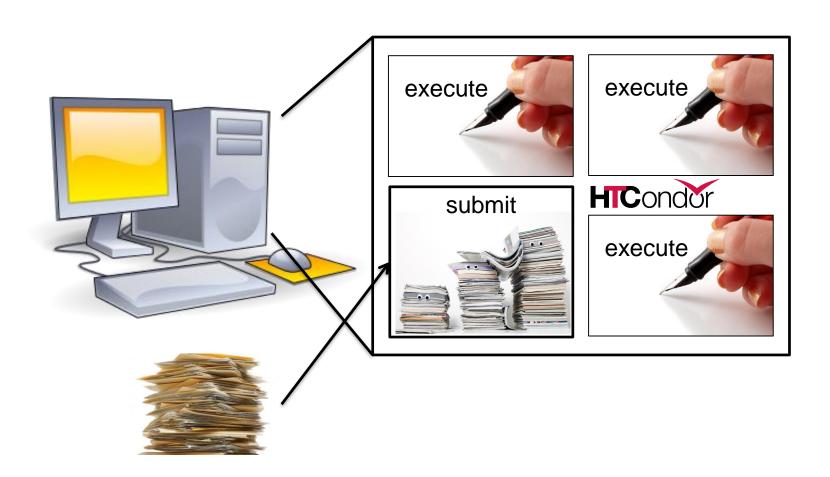


How It Works

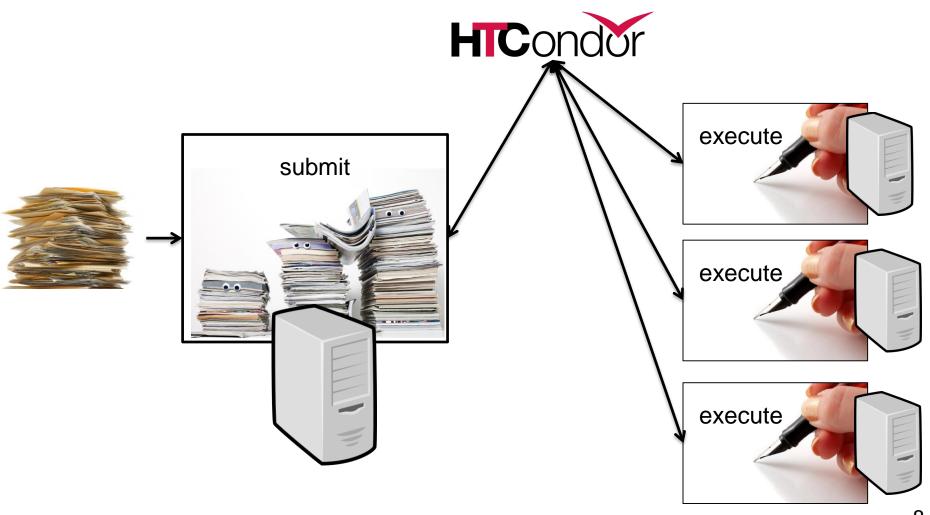
- Submit tasks to a queue (on a submit point)
- HTCondor schedules them to run on computers (execute points)



Single Computer



Multiple Computers



Why HTCondor?

- HTCondor manages and runs work on your behalf
- Schedule tasks on a single computer to not overwhelm the computer
- Schedule tasks on a group* of computers (which may/may not be directly accessible to the user)
- Schedule tasks submitted by multiple users on one or more computers

^{*}in HTCondor-speak, a "pool"

Why HTCondor, cont

- Open source software to enable distributed High Throughput Computing (HTC)
- Full featured, mature production system (1M+ LOC)
- Widely deployed
 - Used in production at hundreds of universities, government labs, commercial companies to manage compute clusters in science, engineering, finance, ...
 - Components used to federate compute clusters into campus grids and wide-area computing grids, e.g.
 Open Science Grid, WLCG, ...

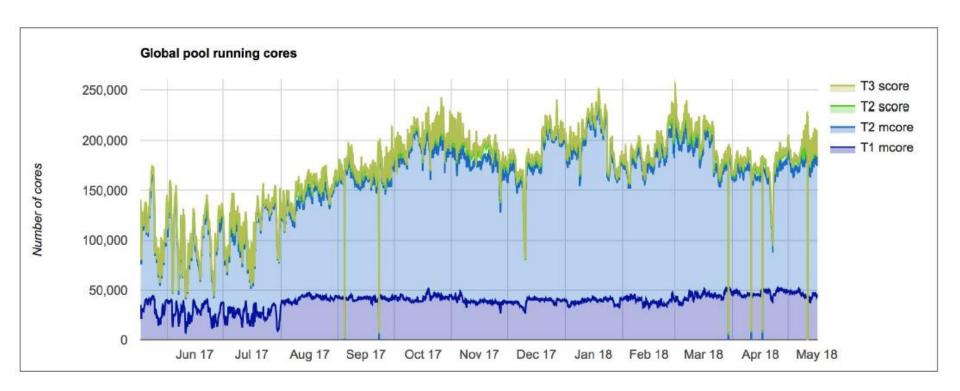
Open Science Grid



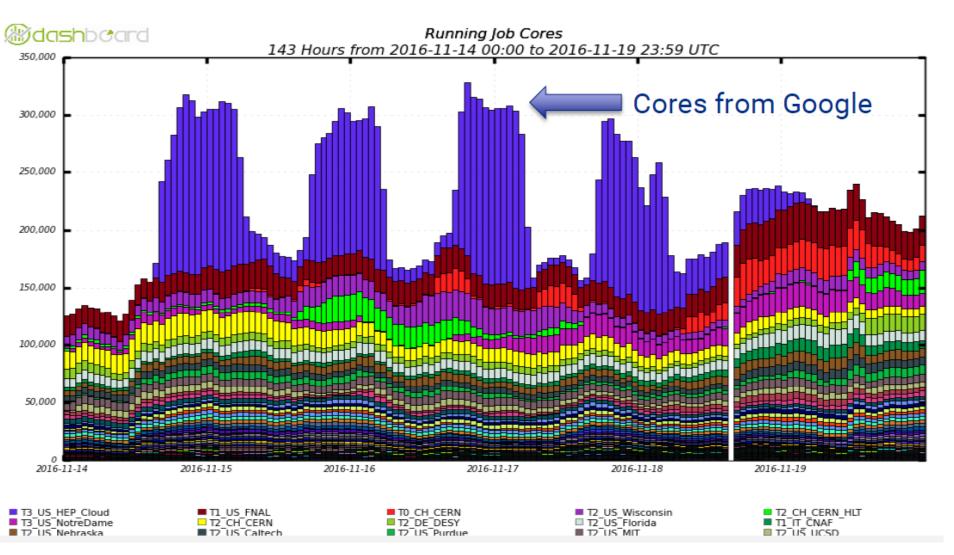
http://display.opensciencegrid.org

CMS Global Pool

 Dynamic cluster, ~200k - 300k cores pulled in from sites worldwide



Bursting into Google Cloud @ SC16



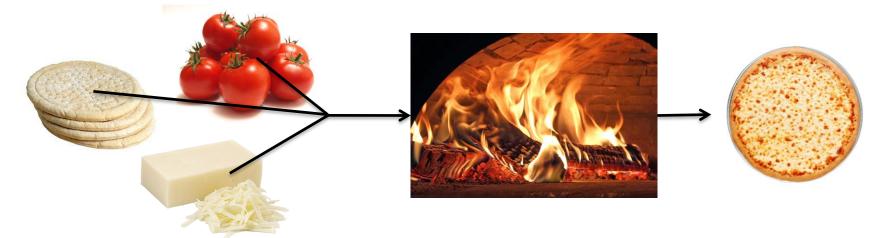
User-Focused Tutorial

- For the purposes of this tutorial, we are assuming that someone else has set up HTCondor on a computer/computers to create a HTCondor "pool".
- The focus of this talk is an introduction on how to get started running computational work on this system.

Running a Job with HTCondor

Jobs

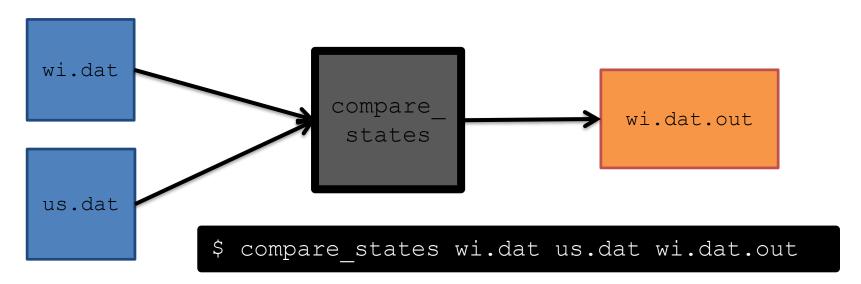
- A single computing task is called a "job"
- Three main pieces of a job are the input, executable (program) and output



 Executable must be runnable from the command line without any interactive input

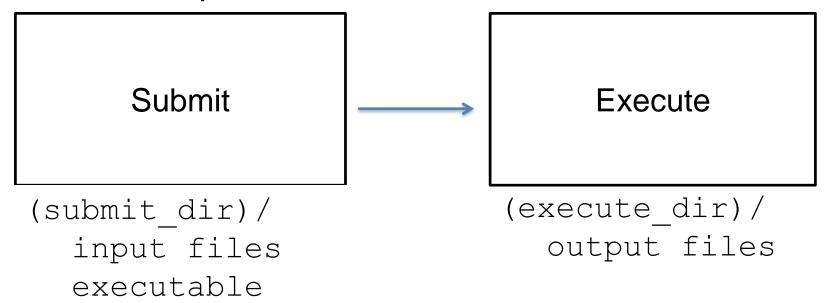
Job Example

 For our example, we will be using an imaginary program called "compare_states", which compares two data files and produces a single output file.



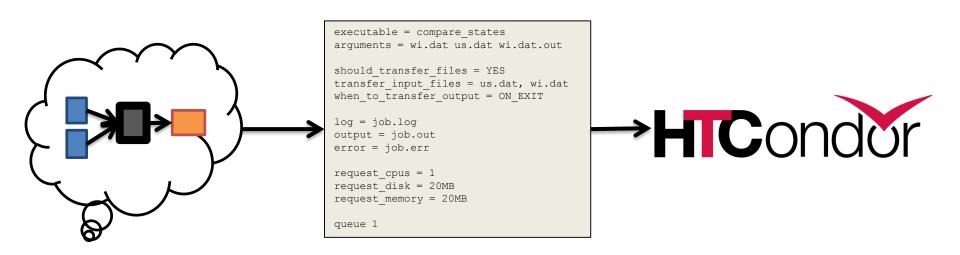
File Transfer

- What about files? Can use a shared file system, chirp, or file transfer mechanism.
- Our example will use HTCondor's file transfer :



Job Translation

 Submit file: communicates everything about your job(s) to HTCondor



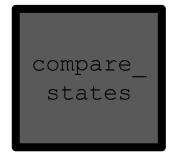
job.submit

```
executable = compare states
arguments = wi.dat us.dat wi.dat.out
should transfer files = YES
transfer input files = us.dat, wi.dat
when to transfer output = ON EXIT
log = job.log
output = job.out
error = job.err
request cpus = 1
request disk = 20MB
request memory = 20MB
queue 1
```

job.submit

```
executable = compare states
arguments = wi.dat us.dat wi.dat.out
should transfer files = YES
transfer input files = us.dat, wi.dat
when to transfer output = ON EXIT
log = job.log
output = job.out
error = job.err
request cpus = 1
request disk = 20MB
request memory = 20MB
queue 1
```

 List your executable and any arguments it takes.



 Arguments are any options passed to the executable from the command line.

\$ compare states wi.dat us.dat wi.dat.out

job.submit

```
executable = compare states
arguments = wi.dat us.dat wi.dat.out
should transfer files = YES
transfer input files = us.dat, wi.dat
when to transfer output = ON EXIT
log = job.log
output = job.out
error = job.err
request cpus = 1
request disk = 20MB
request memory = 20MB
queue 1
```

Indicate
 your input
 files.

wi.dat

us.dat

job.submit

```
executable = compare states
arguments = wi.dat us.dat wi.dat.out
should transfer files = YES
transfer input files = us.dat, wi.dat
when to transfer output = ON EXIT
log = job.log
output = job.out
error = job.err
request cpus = 1
request disk = 20MB
request memory = 20MB
queue 1
```

 HTCondor will transfer back all new and changed files (usually output) from the job.

wi.dat.out

job.submit

```
executable = compare states
arguments = wi.dat us.dat wi.dat.out
should transfer files = YES
transfer input files = us.dat, wi.dat
when to transfer output = ON EXIT
log = job.log
output = job.out
error = job.err
request cpus = 1
request disk = 20MB
request memory = 20MB
queue 1
```

- log: file created by HTCondor to track job progress
- output/err or: captures stdout and stderr

job.submit

```
executable = compare states
arguments = wi.dat us.dat wi.dat.out
should transfer files = YES
transfer input files = us.dat, wi.dat
when to transfer output = ON EXIT
log = job.log
output = job.out
error = job.err
request cpus = 1
request disk = 20MB
request memory = 20MB
queue 1
```

- Request the appropriate resources for your job to run.
- queue:
 keyword
 indicating
 "create a
 job."

Submitting and Monitoring

To submit a job/jobs:

```
condor_submit_submit_file_name
```

To monitor submitted jobs, use:

```
condor_q
```

```
$ condor_submit job.submit
Submitting job(s).
1 job(s) submitted to cluster 128.
```

More about condor_q

- By default condor_q shows:
 - user's job only (as of 8.6)
 - See everyone with "condor_q –allusers"
 - jobs summarized in "batches" (as of 8.6)
- Constrain with username, ClusterId or full JobId, which will be denoted [U/C/J] in the following slides

```
$ condor_q
-- Schedd: submit-5.chtc.wisc.edu: <128.104.101.92:9618?... @ 05/01/17 10:35:54

OWNER BATCH_NAME SUBMITTED DONE RUN IDLE TOTAL JOB_IDS

alice CMD: compare_states 5/9 11:05 ____ 1 128.0

1 jobs; 0 completed, 0 removed, 1 idle, 0 running, 0 held, 0 suspended
```

More about condor_q

To see individual job information, use:
 condor_q -nobatch

 We will use the -nobatch option in the following slides to see extra detail about what is happening with a job

Job Idle

Submit Node

```
(submit_dir)/
  job.submit
  compare_states
  wi.dat
  us.dat
  job.log
  job.out
  job.err
```

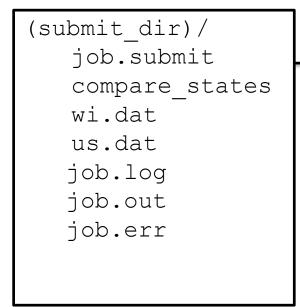
Job Starts by doing File Transfer

```
$ condor q -nobatch
-- Schedd: submit-5.chtc.wisc.edu : <128.104.101.92:9618?...
ΙD
            OWNER
                                    RUN TIME PRI SIZE CMD
                                   0+00:00:00 < 0
128.0
      alice
                      5/9 11:09
                                                     0.0 compare states wi.dat us.dat w
1 jobs; 0 completed, 0 removed, 0 idle, 1 running, 0 held, 0 suspended
```

wi.dat

us.dat

Submit Node



Execute Node

(execute dir)/ compare states

Job Running

Submit Node

```
(submit_dir)/
   job.submit
   compare_states
   wi.dat
   us.dat
   job.log
   job.out
   job.err
```

Execute Node

```
(execute_dir)/
   compare_states
   wi.dat
   us.dat
   stderr
   stdout
   wi.dat.out
```

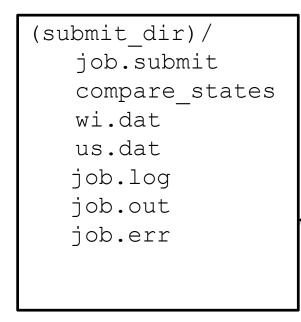
Job Completes

stderr

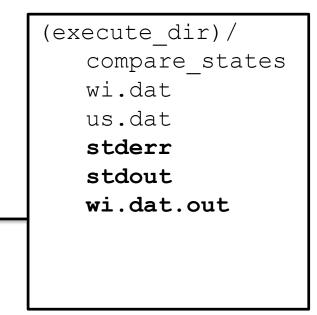
st.dout.

wi.dat.out

Submit Node



Execute Node



Job Completes (cont.)

```
$ condor_q -nobatch

-- Schedd: submit-5.chtc.wisc.edu : <128.104.101.92:9618?...
ID OWNER SUBMITTED RUN_TIME ST PRI SIZE CMD

0 jobs; 0 completed, 0 removed, 0 idle, 0 running, 0 held, 0 suspended</pre>
```

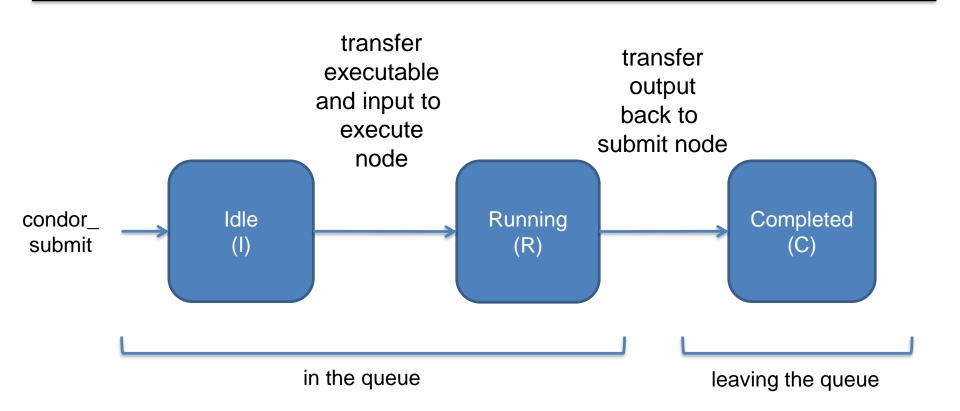
Submit Node

```
(submit_dir)/
   job.submit
   compare_states
   wi.dat
   us.dat
   job.log
   job.out
   job.err
   wi.dat.out
```

Log File

```
000 (128.000.000) 05/09 11:09:08 Job submitted from host:
<128.104.101.92&sock=6423 b881 3>
001 (128.000.000) 05/09 11:10:46 Job executing on host:
<128.104.101.128:9618&sock=5053 3126 3>
. . .
006 (128.000.000) 05/09 11:10:54 Image size of job updated: 220
    1 - MemoryUsage of job (MB)
    220 - ResidentSetSize of job (KB)
005 (128.000.000) 05/09 11:12:48 Job terminated.
    (1) Normal termination (return value 0)
        Usr 0 00:00:00, Sys 0 00:00:00 - Run Remote Usage
        Usr 0 00:00:00, Sys 0 00:00:00 - Run Local Usage
        Usr 0 00:00:00, Sys 0 00:00:00 - Total Remote Usage
        Usr 0 00:00:00, Sys 0 00:00:00 - Total Local Usage
    0 - Run Bytes Sent By Job
    33 - Run Bytes Received By Job
    0 - Total Bytes Sent By Job
    33 - Total Bytes Received By Job
    Partitionable Resources: Usage Request Allocated
       Cpus
                          : 14 20480 17203728
       Disk (KB)
      Memory (MB)
                                            2.0
                                                      2.0
```

Job States



Assumptions

- Aspects of your submit file may be dictated by infrastructure and configuration
- For example: file transfer
 - previous example assumed files would need to be transferred between submit/execute

```
should_transfer_files = YES
```

- not the case with a shared file system

```
should_transfer_files = NO
```

Shared file system

 If a system has a shared file system, where file transfer is not enabled, the submit directory and execute directory are the same.

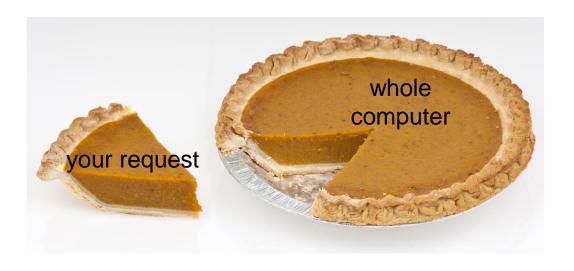
Submit

Execute

```
shared_dir/
input
executable
output
```

Resource Request

- Jobs are nearly always using a part of a computer, not the whole thing
- Very important to request appropriate resources (memory, cpus, disk) for a job



Resource Assumptions

- Even with reasonable default CPU, memory and disk requests, these may be too small!
- Important to run test jobs and use the log file to request the right amount of resources:
 - requesting too little: causes problems for your and other jobs; jobs might by held by HTCondor
 - requesting too much: jobs will match to fewer "slots"

Job Matching and Class Ad Attributes

The Central Manager

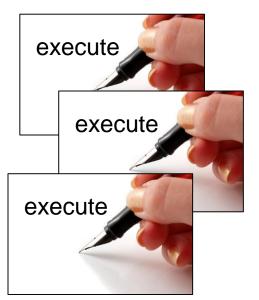
 HTCondor matches jobs with computers via a "central manager".







central manager



Class Ads

- HTCondor stores a list of information about each job and each computer.
- This information is stored as a "Class Ad"



Class Ads have the format:

AttributeName = value 4

can be a boolean, number, string, or expression

Job Class Ad

```
executable = compare_states
arguments = wi.dat us.dat wi.dat.out

should_transfer_files = YES
transfer_input_files = us.dat, wi.dat
when_to_transfer_output = ON_EXIT

log = job.log
output = job.out
error = job.err

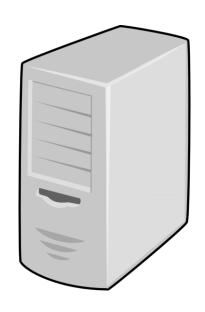
request_cpus = 1
request_disk = 20MB
request_memory = 20MB

queue 1
```

HTCondor configuration

```
RequestCpus = 1
Err = "job.err"
WhenToTransferOutput = "ON EXIT"
TargetType = "Machine"
Cmd =
"/home/alice/tests/htcondor week/compar
e states"
JobUniverse = 5
Iwd = "/home/alice/tests/htcondor week"
RequestDisk = 20480
NumJobStarts = 0
WantRemoteIO = true
OnExitRemove = true
TransferInput = "us.dat, wi.dat"
MyType = "Job"
Out = "job.out"
UserLog =
"/home/alice/tests/htcondor week/job.lo
a"
RequestMemory = 20
```

Computer "Machine" Class Ad



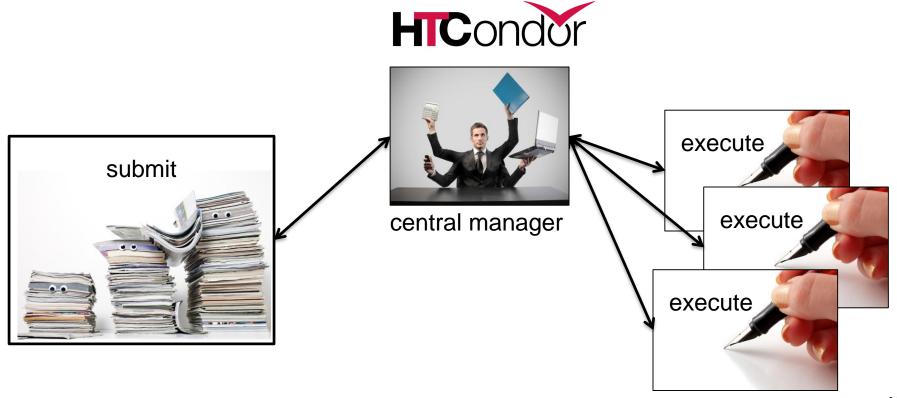
HTCondor configuration

```
HasFileTransfer = true
DynamicSlot = true
TotalSlotDisk = 4300218.0
TargetType = "Job"
TotalSlotMemory = 2048
Mips = 17902
Memory = 2048
UtsnameSysname = "Linux"
MAX PREEMPT = (3600 * 72)
Requirements = ( START ) && (
IsValidCheckpointPlatform ) && (
WithinResourceLimits )
OpSysMajorVer = 6
TotalMemory = 9889
HasGluster = true
OpSysName = "SL"
HasDocker = true
```

. . .

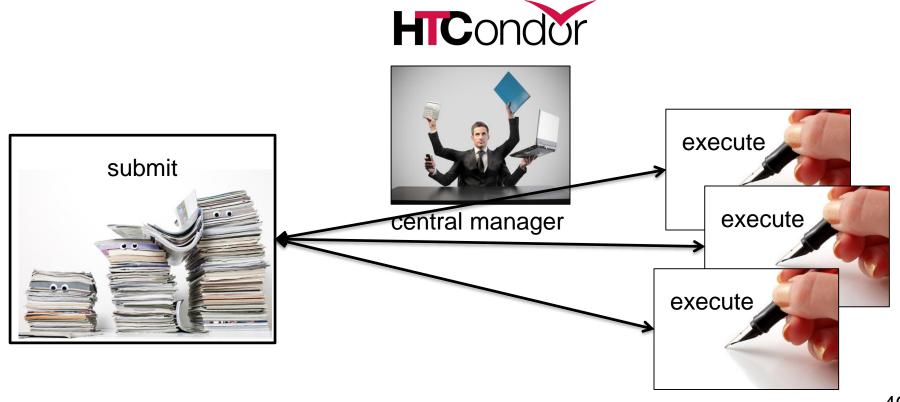
Job Matching

 On a regular basis, the central manager reviews Job resource requests and Machine Class Ads and matches jobs to computers.



Job Execution

 (Then the submit and execute points communicate directly.)



Class Ads for People

 Class Ads also provide lots of useful information about jobs and computers to HTCondor users and administrators



Finding Job Attributes

• Use the "long" option for condor_q condor_q -1 JobId

```
$ condor q -1 128.0
WhenToTransferOutput = "ON EXIT"
TargetType = "Machine"
Cmd = "/home/alice/tests/htcondor week/compare states"
JobUniverse = 5
Iwd = "/home/alice/tests/htcondor week"
RequestDisk = 20480
NumJobStarts = 0
WantRemoteIO = true
OnExitRemove = true
TransferInput = "us.dat, wi.dat"
MyType = "Job"
UserLog = "/home/alice/tests/htcondor week/job.log"
RequestMemory = 20
```

Some Useful Job Attributes

- UserLog: location of job log
- Iwd: Initial Working Directory (i.e. submission directory) on submit node
- MemoryUsage: maximum memory the job has used
- RemoteHost: where the job is running
- BatchName: attribute to label job batches
- ...and more

Selectively display specific attributes

Use the "auto-format" option:

```
condor q [U/C/J] -af Attribute1 Attribute2 ...
```

```
$ condor_q -af ClusterId ProcId RemoteHost MemoryUsage

17315225 116 slot1_1@e092.chtc.wisc.edu 1709
17315225 118 slot1_2@e093.chtc.wisc.edu 1709
17315225 137 slot1_8@e125.chtc.wisc.edu 1709
17315225 139 slot1_7@e121.chtc.wisc.edu 1709
18050961 0 slot1_5@c025.chtc.wisc.edu 196
18050963 0 slot1_3@atlas10.chtc.wisc.edu 269
18050964 0 slot1_25@e348.chtc.wisc.edu 245
18050965 0 slot1_23@e305.chtc.wisc.edu 196
18050971 0 slot1_6@e176.chtc.wisc.edu 220
```

Other Displays

See the whole queue (all users, all jobs)
 condor_q -all

```
$ condor q -all
-- Schedd: submit-5.chtc.wisc.edu : <128.104.101.92:9618?...
OWNER
        BATCH NAME
                     SUBMITTED
                                 DONE
                                               IDLE
                                                      HOLD
                                                            TOTAL JOB IDS
                                        RUN
                                                              1000 18888976.0 ...
        DAG: 128
                     5/9
alice
                         02:52
                                   982
                                                  89
bob
        DAG: 139 5/9 09:21
                                                               180 18910071.0 ...
        DAG: 219 5/9 10:31
                                                              1000 18911030.0 ...
alice
                                          997
bob
        DAG: 226
                 5/9 10:51
                                    10
                                                                44 18913051.0
        CMD: ce.sh 5/9 10:55
                                                                   18913029.0 ...
bob
                                                 998
alice
        CMD: sb
                     5/9
                          10:57
                                                                   18913030.0-999
```

condor_q Reminder

- Default output is batched jobs
 - Batches can be grouped manually using the JobBatchName attribute in a submit file:

```
JobBatchName = "CoolJobs"
```

- Otherwise HTCondor groups jobs automatically
- To see individual jobs, use:

Class Ads for Computers

as condor_q is to jobs, condor_status is to computers (or "machines")

<pre>\$ condor_status</pre>					
Name	OpSys	Arch State	Activity	LoadAv	Mem Actvty
slot1@c001.chtc.wisc.edu	LINU	X X86_64 Unclai	med Idle	0.000	673 25+01
slot1_1@c001.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+01
slot1_2@c001.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+01
slot1_3@c001.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+00
slot1_4@c001.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+14
slot1_5@c001.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	1024 0+01
slot1@c002.chtc.wisc.edu	LINU	X X86_64 Unclai	med Idle	1.000	2693 19+19
slot1_1@c002.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+04
slot1_2@c002.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+01
slot1_3@c002.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	0.990	2048 0+02
slot1@c004.chtc.wisc.edu	LINU	X X86_64 Unclai	med Idle	0.010	645 25+05
slot1_1@c004.chtc.wisc.edu	LINU	X X86_64 Claime	ed Busy	1.000	2048 0+01
Total	Owner Claimed	Unclaimed Matched Pr	eempting Bad	ckfill Dr	ain
X86 64/LINUX 10962	0 10340	613 0	0	0	9
X86 64/WINDOWS 2	2 0	0 0	0	0	0
<u> </u>					
Total 10964	2 10340	613 0	0	0	9

Machine Attributes

Use same options as condor_q:

```
condor_status -1 Slot/Machine
condor_status [Machine] -af Attribute1 Attribute2 ...
```

```
$ condor status -1 slot1 1@c001.chtc.wisc.edu
HasFileTransfer = true
COLLECTOR HOST STRING = "cm.chtc.wisc.edu"
TargetType = "Job"
TotalTimeClaimedBusy = 43334c001.chtc.wisc.edu
UtsnameNodename = ""
Mips = 17902
MAX PREEMPT = (3600 * (72 - 68 * (WantGlidein = ?= true)))
Requirements = (START) && (IsValidCheckpointPlatform) && (
WithinResourceLimits )
State = "Claimed"
OpSysMajorVer = 6
OpSysName = "SL"
```

Machine Attributes

 To summarize, use the "-compact" option condor_status -compact

\$ condor q -compact										
Machine	Plat	form	Slots	Cpus	Gpus	TotalGb	FreCpu	FreeGb	CpuLoad	ST
e007.chtc.wisc.edu	x64/	'SL6	8	8		23.46	0	0.00	1.24	Cb
e008.chtc.wisc.edu	x64/	'SL6	8	8		23.46	0	0.46	0.97	Cb
e009.chtc.wisc.edu	x64/	'SL6	11	16		23.46	5	0.00	0.81	* *
e010.chtc.wisc.edu	x64/	'SL6	8	8		23.46	0	4.46	0.76	Cb
matlab-build-1.chtc.wisc.ed	du x64/	'SL6	1	12		23.45	11	13.45	0.00	* *
matlab-build-5.chtc.wisc.ed	du x64/	'SL6	0	24		23.45	24	23.45	0.04	Ui
mem1.chtc.wisc.edu	x64/	'SL6	24	80		1009.67	8	0.17	0.60	* *
Total	Owner	Claimed	Unclair	med Ma	atched	Preempti	ing Back	fill Dra	ain	
x64/SL6 10416	0	9984	2	427	0		0	0	5	
x64/WinVista 2	2	0		0	0		0	0	0	
Total 10418	2	9984	2	427	0		0	0	5	

Submitting Multiple Jobs with HTCondor

Many Jobs, One Submit File

 HTCondor has built-in ways to submit multiple independent jobs with one submit file



Advantages

- Run many independent jobs...
 - analyze multiple data files
 - test parameter or input combinations
 - and more!
- ...without having to:
 - start each job individually
 - create separate submit files for each job

Multiple, Numbered, Input Files

```
job.submit
```

```
executable = analyze.exe
arguments = file.in file.out
transfer_input_files = file.in

log = job.log
output = job.out
error = job.err

queue
```

```
(submit_dir)/
analyze.exe
file0.in
file1.in
file2.in
```

 Goal: create 3 jobs that each analyze a different input file.

Multiple Jobs, No Variation

```
job.submit
```

```
executable = analyze.exe
arguments = file0.in file0.out
transfer_input_files = file.in

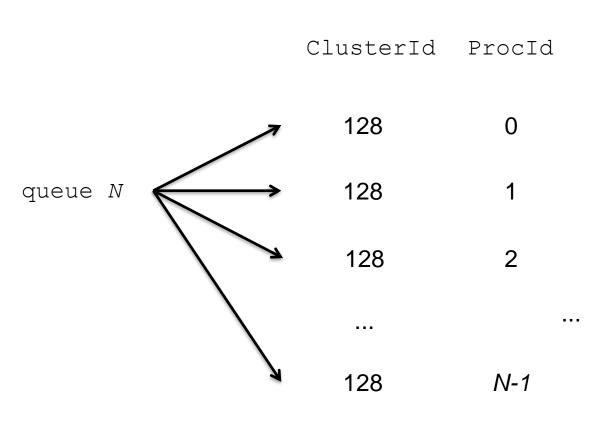
log = job.log
output = job.out
error = job.err

queue 3
```

```
(submit_dir)/
analyze.exe
file0.in
file1.in
file2.in
```

 This file generates 3 jobs, but doesn't use multiple inputs and will overwrite outputs

Automatic Variables



- Each job's
 ClusterId and
 ProcId numbers
 are saved as job
 attributes
- They can be accessed inside the submit file using:
 - \$(ClusterId)
 - \$(ProcId)

Job Variation

job.submit

```
executable = analyze.exe
arguments = file0.in file0.out
transfer_input_files = file0.in

log = job.log
output = job.out
error = job.err

queue
```

```
(submit_dir)/
analyze.exe
file0.in
file1.in
file2.in
```

 How to uniquely identify each job (filenames, log/out/err names)?

Using \$(Procld)

job.submit

```
executable = analyze.exe
arguments = file$(ProcId).in file$(ProcId).out
should_transfer_files = YES
transfer_input_files = file$(ProcId).in
when_to_transfer_output = ON_EXIT

log = job_$(ClusterId).log
output = job_$(ClusterId)_$(ProcId).out
error = job_$(ClusterId)_$(ProcId).err

queue 3
```

Use the \$ (ClusterId), \$ (ProcId)
 variables to provide unique values to jobs.*

Organizing Jobs

```
12181445_0.err 16058473_0.err 17381628_0.err 18159900_0.err 5175744_0.err 7266263_0.err 12181445_0.log 16058473_0.log 17381628_0.log 18159900_0.log 5175744_0.log 7266263_0.log 12181445_0.out 16058473_0.out 17381628_0.out 18159900_0.out 5175744_0.log 7266263_0.out 13609567_0.err 16060330_0.err 17381640_0.err 3446080_0.err 5176204_0.err 7266267_0.err 13609567_0.log 16060330_0.log 17381640_0.log 3446080_0.log 5176204_0.log 7266267_0.log 13609567_0.out 16060330_0.out 17381640_0.out 3446080_0.out 5176204_0.out 7266267_0.out 13612268_0.err 16254074_0.err 17381665_0.err 3446306_0.err 5295132_0.err 7937420_0.err 13612268_0.log 16254074_0.out 17381665_0.log 3446306_0.log 5295132_0.log 7937420_0.out 13630381_0.err 17134215_0.err 17381676_0.err 4347054_0.err 5318339_0.err 8779997_0.err 13630381_0.log 17134215_0.out 17381676_0.log 4347054_0.out 5318339_0.out 8779997_0.out 13630381_0.out 17134215_0.out 17381676_0.log 4347054_0.out 5318339_0.out 8779997_0.out
```



Shared Files

- HTCondor can transfer an entire directory or all the contents of a directory
 - transfer whole directory

```
transfer_input_files = shared
```

- transfer contents only

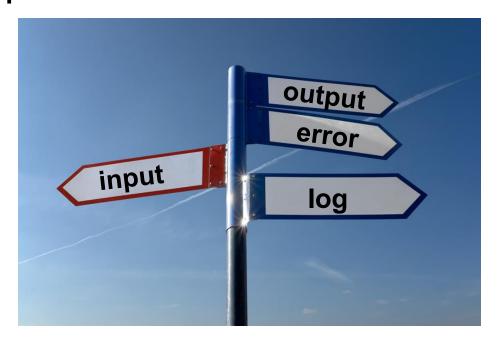
```
transfer_input_files = shared/
```

```
job.submit
shared/
    reference.db
    parse.py
    analyze.py
    cleanup.py
    links.config
```

 Useful for jobs with many shared files; transfer a directory of files instead of listing files individually

Organize Files in Sub-Directories

 Create sub-directories* and use paths in the submit file to separate input, error, log, and output files.



^{*} must be created before the job is submitted

Use Paths for File Type

```
(submit dir)/
```

```
input/
             file0.out
                                    log/
                                               err/
job.submit
                           file0.in
                                      job0.log
                                                 job0.err
            file1.out
analyze.exe
                           file1.in
                                      job1.log job1.err
             file2.out
                           file2.in
                                      job2.log
                                                 job2.err
```

job.submit

```
executable = analyze.exe
arguments = file$(Process).in file$(ProcId).out
transfer_input_files = input/file$(ProcId).in

log = log/job$(ProcId).log
error = err/job$(ProcId).err

queue 3
```

InitialDir

- Change the submission directory for each job using initialdir
- Allows the user to organize job files into separate directories.
- Use the same name for all input/output files
- Useful for jobs with lots of output files



Separate Jobs with InitialDir

```
(submit dir)/
```

```
job.submit
               job0/
                            job1/
                                         job2/
                 file.in
                              file.in
analyze.exe
                                          file.in
                 job.log
                              job.log
                                          job.log
                 job.err
                            job.err
                                          job.err
                 file.out
                              file.out
                                          file.out.
```

job.submit

```
executable = analyze.exe
initialdir = job$(ProcId)
arguments = file.in file.out
transfer_input_files = file.in

log = job.log
error = job.err

queue 3
```

Executable should be in the directory with the submit file, *not* in the individual job directories

Other Submission Methods

- What if your input files/directories aren't numbered from 0 - (N-1)?
- There are other ways to submit many jobs!



Submitting Multiple Jobs

```
executable = compare_states
arguments = wi.dat us.dat wi.dat.out

transfer_input_files = us.dat, wi.dat
queue 1
```

Replacing single job inputs

```
executable = compare_states
arguments = $(infile) us.dat $(infile).out

transfer_input_files = us.dat, $(infile)

queue ...
```

with a variable of choice

Possible Queue Statements

matching pattern	queue infile matching *.dat
in list	queue infile in (wi.dat ca.dat ia.dat)
from file	queue infile from state_list.txt wi.dat ca.dat ia.dat state_list.txt

Queue Statement Comparison

matching pattern	Natural nested looping, minimal programming, use optional "files" and "dirs" keywords to only match files or directories Requires good naming conventions,
in list	Supports multiple variables, all information contained in a single file, reproducible Harder to automate submit file creation
from file	Supports multiple variables, highly modular (easy to use one submit file for many job batches), reproducible Additional file needed

Using Multiple Variables

 Both the "from" and "in" syntax support using multiple variables from a list.

```
job.submit

executable = compare_states
arguments = -year $(option) -input $(file)

should_transfer_files = YES
when_to_transfer_output = ON_EXIT
transfer_input_files = $(file)

queue file,option from job_list.txt
```

```
job_list.txt
wi.dat, 2010
wi.dat, 2015
ca.dat, 2010
ca.dat, 2015
ia.dat, 2010
ia.dat, 2015
```

Other Features

Match only files or directories:

```
queue input matching files *.dat
queue directory matching dirs job*
```

Submit multiple jobs with same input data

```
queue 10 input matching files *.dat
```

– Use other automatic variables: \$ (Step)

```
arguments = -i $(input) -rep $(Step)
queue 10 input matching files *.dat
```

Testing and Troubleshooting

What Can Go Wrong?

- Jobs can go wrong "internally":
 - "job failed to run..."
 - something happens after the executable begins to run
- Jobs can go wrong from HTCondor's perspective:
 - A job can't be started at all ("failed to launch"),
 - Uses too much memory,
 - Has a badly formatted executable,
 - And more...

Reviewing Failed Jobs

 A job's log, output and error files can provide valuable information for troubleshooting

Log	Output	Error
 When jobs were submitted, started, and stopped Resources used Exit status Where job ran Interruption reasons 	Any "print" or "display" information from your program	Captured by the operating system

Reviewing Jobs

 To review a large group of jobs at once, use condor_history

As condor_q is to the present, condor_history is to the past

```
$ condor history alice
        OWNER
                 SUBMITTED
                                                        CMD
 ΙD
                             RUN TIME
                                        ST COMPLETED
                              0+00:07:37 C 5/11 16:00 /home/alice
189.1012 alice
                 5/11 09:52
189.1002 alice
              5/11 09:52
                             0+00:08:03 C 5/11 16:00 /home/alice
189.1081 alice
              5/11 09:52
                             0+00:03:16 C 5/11 16:00 /home/alice
189.944 alice
                5/11 09:52
                              0+00:11:15 C 5/11 16:00 /home/alice
189.659 alice
                              0+00:26:56 C 5/11 16:00 /home/alice
                 5/11 09:52
                 5/11 09:52
                             0+00:27:07 C 5/11 16:00 /home/alice
189.653 alice
189.1040 alice
                 5/11 09:52
                              0+00:05:15 C 5/11 15:59 /home/alice
189.1003 alice
              5/11 09:52
                              0+00:07:38 C 5/11 15:59 /home/alice
189.962 alice
                 5/11 09:52
                             0+00:09:36 C 5/11 15:59 /home/alice
189.961 alice
                 5/11 09:52
                              0+00:09:43 C
                                            5/11 15:59 /home/alice
189.898 alice
                 5/11 09:52
                              0+00:13:47 C
                                            5/11 15:59 /home/alice
```

"Live" Troubleshooting

To log in to a job where it is running, use:

```
condor_ssh_to_job JobId
```

```
$ condor_ssh_to_job 128.0
Welcome to slot1_31@e395.chtc.wisc.edu!
Your condor job is running with pid(s) 3954839.
```

Held Jobs

- HTCondor will put your job on hold if there's something YOU need to fix.
- A job that goes on hold is interrupted and kept from running again, but remains

submitted in the queue in the "H" state.



Diagnosing Holds

 If HTCondor puts a job on hold, it provides a hold reason, which can be viewed with:

condor_q -hold [-wide]

```
$ condor_q -hold -af HoldReason
Error from slot1_1@wid-003.chtc.wisc.edu: Job has gone over
  memory limit of 2048 megabytes.
Error from slot1_20@e098.chtc.wisc.edu: SHADOW at
  128.104.101.92 failed to send file(s) to <128.104.101.98:35110>: error
  reading from /home/alice/script.py: (errno 2) No such file or directory;
  STARTER failed to receive file(s) from <128.104.101.92:9618>
Error from slot1_11@e138.chtc.wisc.edu: STARTER
  at 128.104.101.138 failed to send file(s) to <128.104.101.92:9618>; SHADOW at
  128.104.101.92 failed to write to file /home/alice/Test_18925319_16.err:
  (errno 122) Disk quota exceeded
Error from slot1_38@e270.chtc.wisc.edu: Failed
  to execute '/var/lib/condor/execute/slot1/dir_2471876/condor_exec.exe' with
  arguments 2: (errno=2: 'No such file or directory')
```

Common Hold Reasons

- Job has used more memory than requested
- Incorrect path to files that need to be transferred
- Badly formatted bash scripts (have Windows instead of Unix line endings)
- Submit directory is over quota
- The admin has put your job on hold

Fixing Holds

 Job attributes can be edited while jobs are in the queue using:

```
condor_qedit [U/C/J] Attribute Value
```

```
$ condor_qedit 128.0 RequestMemory 3072
Set attribute "RequestMemory".
```

 If a job has been fixed and can run again, release it with:

```
condor_release [U/C/J]
```

```
$ condor_release 128.0
Job 18933774.0 released
```

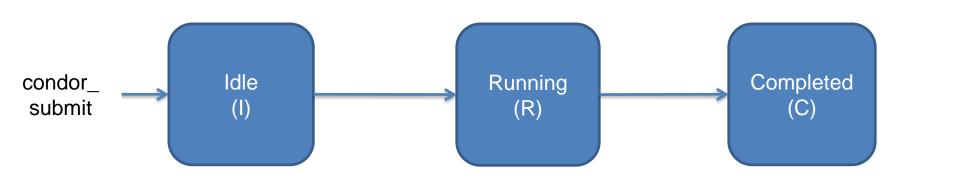
Holding or Removing Jobs

- If you know your job has a problem and it hasn't yet completed, you can:
 - Place it on hold yourself, with condor_hold [U/C/J]

```
$ condor_hold bob
All jobs of user "bob" have been held
$ condor_hold 128
All jobs in cluster 128 have been held
$ condor_hold 128.0
Job 128.0 held
```

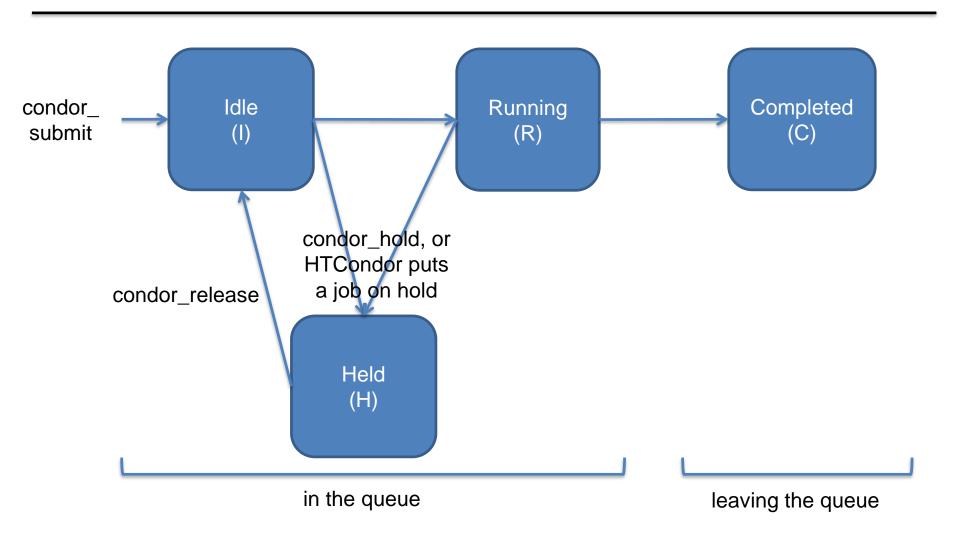
Remove it from the queue, using condor_rm [U/C/J]

Job States, Revisited

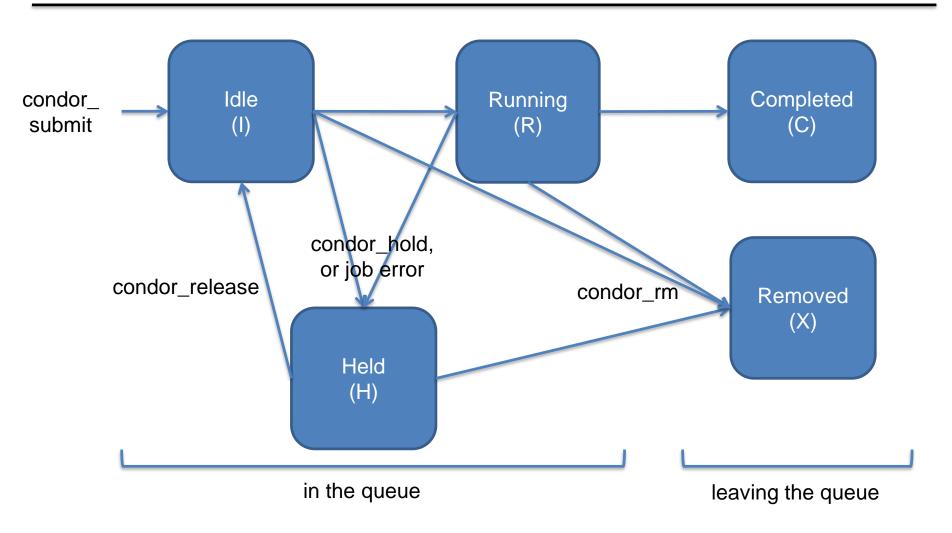


in the queue leaving the queue

Job States, Revisited



Job States, Revisited*



Some Use Cases and Mechanisms

Interactive Jobs

 An interactive job proceeds like a normal batch job, but opens a bash session into the job's execution directory instead of running an executable.

```
condor_submit -i submit_file
```

```
$ condor_submit -i interactive.submit
Submitting job(s).
1 job(s) submitted to cluster 18980881.
Waiting for job to start...
Welcome to slot1_9@e184.chtc.wisc.edu!
```

Useful for testing and troubleshooting

Output Handling

 Only transfer back specific files from the job's execution using transfer_ouput_files

```
(submit_dir)/
(execute_dir)/

condor_exec.exe
results-tmp-01.dat
results-tmp-03.dat
results-tmp-04.dat
results-tmp-05.dat
results-final.dat
```

condor_chirp

- What if you want to only read part of a file?
- What if you want to write records into an output file?

Use condor_chirp!

https://htcondor.readthedocs.io/en/stable/man-pages/condor_chirp.html

Can also edit job classad or add entries to the job event log file!

Self-Checkpointing

- By default, a job that is interrupted will start from the beginning if it is restarted.
- It is possible to implement selfcheckpointing, which will allow a job to restart from a saved state if interrupted.
- Self-checkpointing is useful for very long jobs, and being able to run on opportunistic resources.

Self-Checkpointing How-To

- Edit executable:
 - Atomically save intermediate states to a checkpoint file
 - Always check for a checkpoint file when starting
- Add HTCondor option that a) saves all intermediate/output files from the interrupted job and b) transfers them to the job when HTCondor runs it again

```
when_to_transfer_output = ON_EXIT_OR_EVICT
# Optional: also checkpoint if my job exits with
# a specified exit status
CheckpointExitCode = 77
```

Job Universes

 HTCondor has different "universes" for running specialized job types

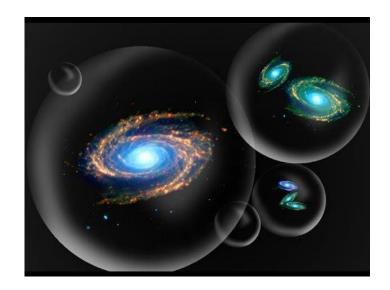
HTCondor Manual: Choosing an HTCondor Universe

- Vanilla (default)
 - good for most software

HTCondor Manual: Vanilla Universe

 Set in the submit file using:

universe = vanilla



Other Universes

- Local
 - Run jobs on the submit node
- Java
 - Built-in Java support
- e Java
- Executable is a jar file
- Grid
 - Delegate jobs to another scheduler
 - The basis for HTCondor-CE



Other Universes (cont.)

- Docker
 - Run jobs inside a Docker container
- VM
 - Run jobs inside a virtual machine
- Parallel
 - Used for coordinating jobs across multiple servers (e.g. MPI code)
 - Not necessary for single server multi-core jobs

Multi-CPU and GPU Computing

 Jobs that use multiple cores on a single computer can be run in the vanilla universe (parallel universe not needed):

```
request_cpus = 16
```

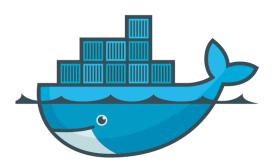
 If there are computers with GPUs, request them with:

```
request_gpus = 1
```

```
universe = docker
executable = /bin/my_executable
```

Executable comes either from submit machine or image

NOT FROM execute machine



```
universe = docker
executable = /bin/my_executable
docker_image = deb7_and_HEP_stack
```

Image is the name of the docker image stored on execute machine

```
universe = docker
executable = /bin/my executable
docker image = deb7 and HEP stack
transfer input files = some input
     HTCondor can transfer input files from
     submit machine into container
```

(same with output in reverse)

101

```
universe = docker
executable = /bin/my executable
arguments = arg1
docker image = deb7 and HEP stack
transfer input files = some input
output = out
error = err
log = log
queue
```

Automation

Automation

- After job submission, HTCondor manages jobs based on its configuration
- You can use options that will customize job management even further
- These options can
 automate when
 jobs are started,
 stopped, and removed.



Retries

- Problem: a small number of jobs fail with a known error code; if they run again, they complete successfully.
- Solution: If the job exits with the error code, leave it in the queue to run again

```
max retries = 3
```

Retries, cont.

 Can also combine with success_exit_code = < Integer > retry_until = < Integer | Expression >

```
executable = foo.exe
max_retries = 5
retry_untl = ExitCode >= 0
queue
```

Automatically Hold Jobs

- Problem: Your job should run in 2 hours or less, but a few jobs "hang" randomly and run for days
- Solution: Put jobs on hold if they run for over 2 hours, using a periodic_hold statement

```
periodic_hold = (JobStatus == 2) &&
  ((CurrentTime - EnteredCurrentStatus) > (60 * 60 * 2))

How long the job has been running, in seconds
2 hours
```

Automatically Release Jobs

- Problem (related to previous): A few jobs are being held for running long; they will complete if they run again.
- Solution: automatically release those held jobs with a periodic_release option, up to 5 times

```
periodic_release = (JobStatus == 5) &&
  (HoldReason == 3) && (NumJobStarts < 5)

job was put on hold
  by periodic_hold

job has started running
  less than 5 times</pre>
```

Automatically Remove Jobs

- Problem: Jobs are repetitively failing
- Solution: Remove jobs from the queue using a periodic remove statement



Automatic Memory Increase

- Putting all these pieces together, the following lines will:
 - request a default amount of memory (2GB)
 - put the job on hold if it is exceeded
 - release the the job with an increased memory request

```
request_memory = ifthenelse(isUndefined(MemoryUsage),
2048,(MemoryUsage * 3/2), 2048)
periodic_hold = (MemoryUsage >= ((RequestMemory) * 5/4 )) &&
(JobStatus == 2)
periodic_release = (CurrentTime - EnteredCurrentStatus) > 180) &&
(NumJobStarts < 5) && (HoldReasonCode == 3)</pre>
```

Relevant Job Attributes

- CurrentTime: current time
- EnteredCurrentStatus: time of last status change
- ExitCode: the exit code from the job
- HoldReasonCode: number corresponding to a hold reason
- NumJobStarts: how many times the job has gone from idle to running
- JobStatus: number indicating idle, running, held, etc.
- MemoryUsage: how much memory the job has used

General User Commands

- condor_submit
- condor_status
- condor_q
- condor_q -analyze
- condor_ssh_to_job
- condor_submit -i
- condor_hold / release
- condor_run
- condor_rm
- condor_prio
- condor_history
- condor_submit_dag
- condor_chirp

Submit new Jobs

View Pool Status

View Job Queue

Why job/machines fail to match?

Create ssh session to active job

Submit interactive job

Hold a job, or release a held job

Submit and block

Remove Jobs

Intra-User Job Prios

Completed Job Info

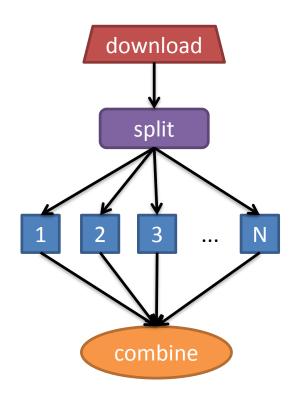
Submit new DAG workflow

Access files/ad from active job

Describing Workflows with DAGMan

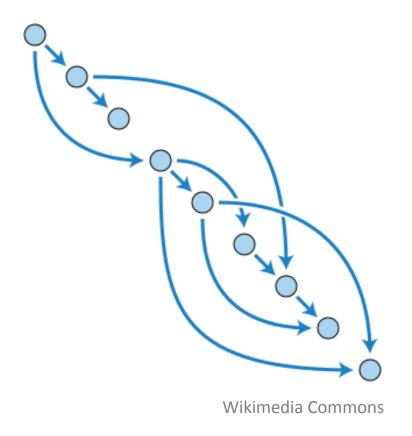
Workflows

- Problem: Want to submit jobs in a particular order, with dependencies between groups of jobs
- Solution: Write a DAG

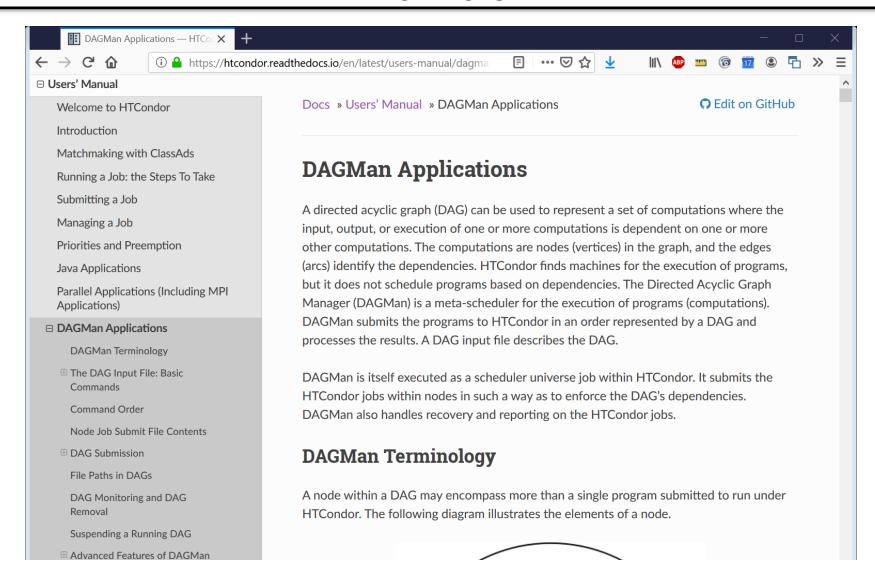


DAG = "directed acyclic graph"

- topological ordering of vertices ("nodes") is established by directional connections ("edges")
- "acyclic" aspect requires a start and end, with no looped repetition
 - can contain cyclic subcomponents, covered in later slides for workflows

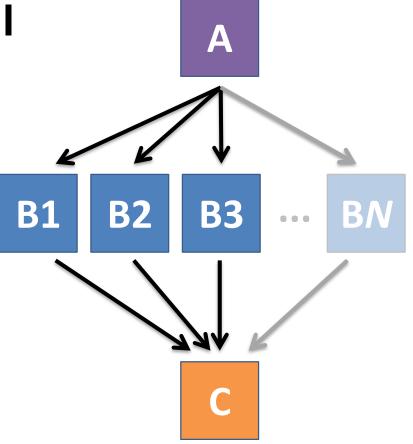


DAGMan in the HTCondor Manual



Simple Example for this Tutorial

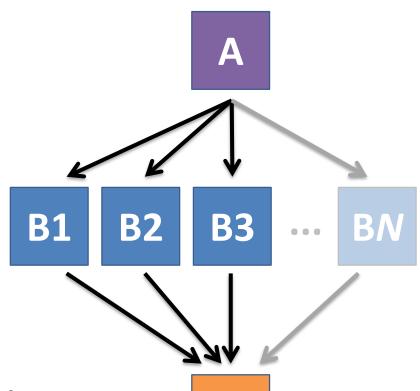
 The DAG input file will communicate the "nodes" and directional "edges" of the DAG



Basic DAG input file: JOB nodes, PARENT-CHILD edges

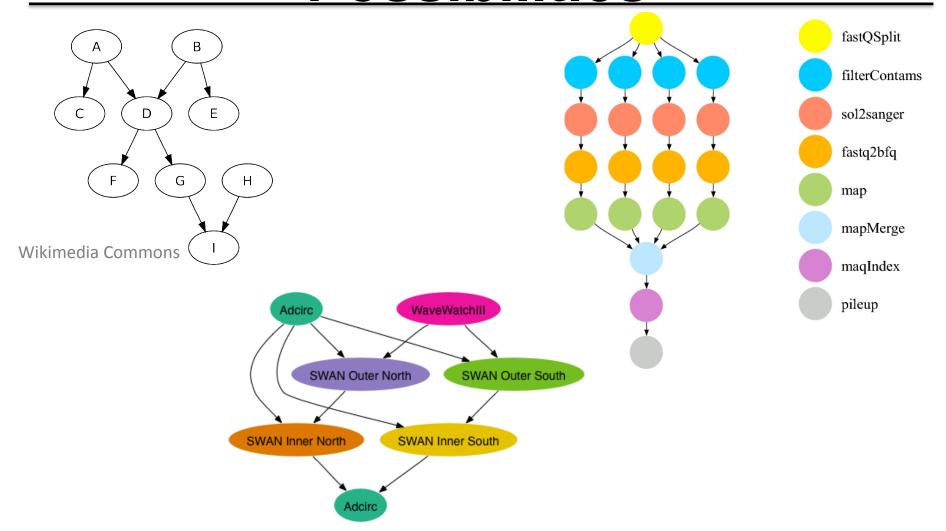
my.dag

```
JOB A A.sub
JOB B1 B1.sub
JOB B2 B2.sub
JOB B3 B3.sub
JOB C C.sub
PARENT A CHILD B1 B2 B3
PARENT B1 B2 B3 CHILD C
```

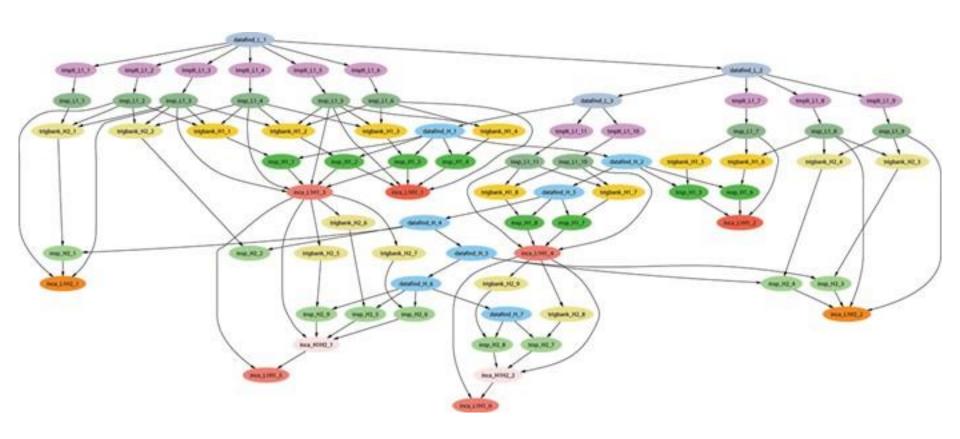


 Node names are used by various DAG features to modify their execution by DAG Manager.

Endless Workflow Possibilities



Endless Workflow Possibilities

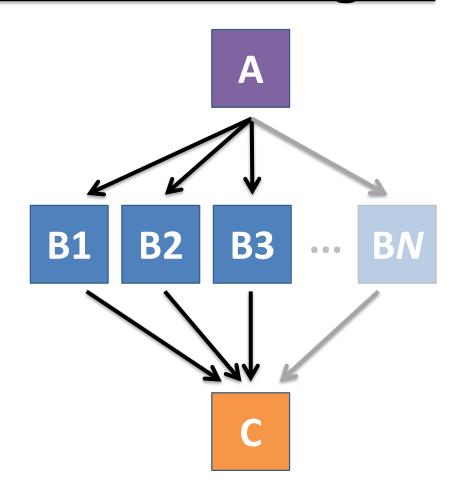


Submitting and Monitoring a DAGMan Workflow

Basic DAG input file: JOB nodes, PARENT-CHILD edges

my.dag

```
JOB A A.sub
JOB B1 B1.sub
JOB B2 B2.sub
JOB B3 B3.sub
JOB C C.sub
PARENT A CHILD B1 B2 B3
PARENT B1 B2 B3 CHILD C
```



Submitting a DAG to the queue

Submission command:

condor_submit_dag dag_file

```
$ condor_submit_dag my.dag

File for submitting this DAG to HTCondor : mydag.dag.condor.sub
Log of DAGMan debugging messages : mydag.dag.dagman.out
Log of HTCondor library output : mydag.dag.lib.out
Log of HTCondor library error messages : mydag.dag.lib.err
Log of the life of condor_dagman itself : mydag.dag.dagman.log

Submitting job(s).

1 job(s) submitted to cluster 87274940.
```

Jobs are automatically submitted by the DAGMan job

Seconds later, node A is submitted:

```
$ condor_q
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...

OWNER BATCH_NAME SUBMITTED DONE RUN IDLE TOTAL JOB_IDS
alice my.dag+128 4/30 18:08 _ _ _ 1 5 129.0

2 jobs; 0 completed, 0 removed, 1 idle, 1 running, 0 held, 0 suspended

$ condor_q -nobatch
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...

ID OWNER SUBMITTED RUN_TIME ST PRI SIZE CMD

128.0 alice 4/30 18:08 0+00:00:36 R 0 0.3 condor_dagman

129.0 alice 4/30 18:08 0+00:00:00 I 0 0.3 A_split.sh

2 jobs; 0 completed, 0 removed, 1 idle, 1 running, 0 held, 0 suspended
```

Jobs are automatically submitted by the DAGMan job

After A completes, B1-3 are submitted

```
$ condor q
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...
OWNER BATCH NAME SUBMITTED DONE RUN IDLE TOTAL JOB IDS
alice my.dag+128 4/30 8:08 1 3 5 129.0...132.0
4 jobs; 0 completed, 0 removed, 3 idle, 1 running, 0 held, 0 suspended
$ condor q -nobatch
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...
ID
      OWNER SUBMITTED RUN TIME ST PRI SIZE CMD
128.0 alice 4/30 18:08 0+00:20:36 R 0 0.3 condor dagman
130.0 alice 4/30 18:18 0+00:00:00 I 0 0.3 B run.sh
131.0 alice 4/30 18:18 0+00:00:00 I 0 0.3 B run.sh
132.0 alice 4/30 18:18 0+00:00:00 I 0 0.3 B run.sh
4 jobs; 0 completed, 0 removed, 3 idle, 1 running, 0 held, 0 suspended
```

Jobs are automatically submitted by the DAGMan job

After B1-3 complete, node C is submitted

```
$ condor_q
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...

OWNER BATCH_NAME SUBMITTED DONE RUN IDLE TOTAL JOB_IDS
alice my.dag+128 4/30 8:08 4 1 5 129.0...133.0
2 jobs; 0 completed, 0 removed, 1 idle, 1 running, 0 held, 0 suspended

$ condor_q -nobatch
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...

ID OWNER SUBMITTED RUN_TIME ST PRI SIZE CMD

128.0 alice 4/30 18:08 0+00:46:36 R 0 0.3 condor_dagman

133.0 alice 4/30 18:54 0+00:00:00 I 0 0.3 C_combine.sh
2 jobs; 0 completed, 0 removed, 1 idle, 1 running, 0 held, 0 suspended
```

Status files are Created at the time of DAG submission

(dag dir)/

```
A.sub B1.sub B2.sub
B3.sub C.sub (other job files)
my.dag my.dag.condor.sub my.dag.dagman.log
my.dag.dagman.out my.dag.lib.err my.dag.lib.out
my.dag.nodes.log
```

- *.condor.sub and *.dagman.log describe the queued DAGMan job process, as for all queued jobs
- *.dagman.out has detailed logging (look to first for errors)
- *.lib.err/out contain std err/out for the DAGMan job process
- *.nodes.log is a combined log of all jobs within the DAG

Removing a DAG from the queue

 Remove the DAGMan job in order to stop and remove the entire DAG:

```
condor_rm dagman_jobID
```

 Creates a rescue file so that only incomplete or unsuccessful NODES are repeated upon resubmission

```
$ condor_q
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...

OWNER BATCH_NAME SUBMITTED DONE RUN IDLE TOTAL JOB_IDS
alice my.dag+128 4/30 8:08 4 _ 1 6 129.0...133.0
2 jobs; 0 completed, 0 removed, 1 idle, 1 running, 0 held, 0 suspended
$ condor_rm 128
All jobs in cluster 128 have been marked for removal
```

Removal of a DAG results in a rescue file

```
(dag_dir)/
```

```
A.sub B1.sub B2.sub B3.sub C.sub (other job files)
my.dag my.dag.condor.sub my.dag.dagman.log
my.dag.dagman.out my.dag.lib.err my.dag.lib.out
my.dag.metrics my.dag.nodes.log my.dag.rescue001
```

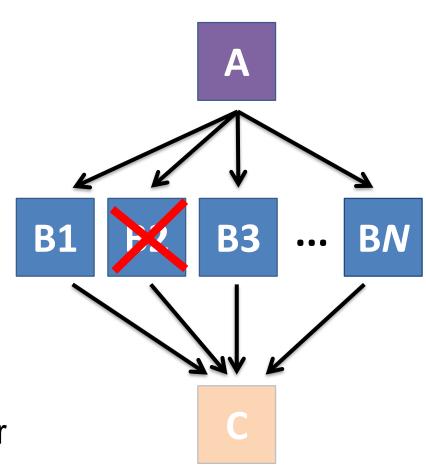
- Named dag file.rescue001
 - increments if more rescue DAG files are created
- Records which NODES have completed successfully
 - does not contain the actual DAG structure

Rescue Files For Resuming a Failed DAG

- A rescue file is created when:
 - a node fails, and after DAGMan advances through any other possible nodes
 - the DAG is removed from the queue (or aborted; covered later)
 - the DAG is halted and not unhalted (covered later)
- Resubmission uses the rescue file (if it exists)
 when the original DAG file is resubmitted
 - Override: condor_submit_dag dag_file -f

Node Failures Result in DAG Failure

- If a node JOB fails (non-zero exit code)
 - DAGMan continues to run other JOB nodes until it can no longer make progress
- Example at right:
 - B2 fails
 - Other B* jobs continue
 - DAG fails and exits after
 B* and before node C



Resolving held node jobs

```
$ condor_q -nobatch
-- Sched: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...
ID OWNER SUBMITTED RUN_TIME ST PRI SIZE CMD

128.0 alice 4/30 18:08 0+00:20:36 R 0 0.3 condor_dagman

130.0 alice 4/30 18:18 0+00:00:00 H 0 0.3 B_run.sh

131.0 alice 4/30 18:18 0+00:00:00 H 0 0.3 B_run.sh

132.0 alice 4/30 18:18 0+00:00:00 H 0 0.3 B_run.sh

4 jobs; 0 completed, 0 removed, 0 idle, 1 running, 3 held, 0 suspended
```

- Look at the hold reason (in the job log, or with 'condor q -hold')
- Fix the issue and release the jobs (condor_release)
 -OR- remove the entire DAG, resolve, then resubmit the DAG

DAG Completion

```
(dag dir)/
```

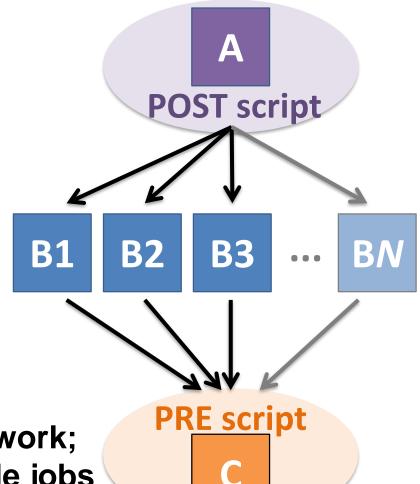
```
A.sub B1.sub B2.sub
B3.sub C.sub (other job files)
my.dag my.dag.condor.sub my.dag.dagman.log
my.dag.nodes.log my.dag.dagman.metrics
```

- *.dagman.metrics is a summary of events and outcomes
- *.dagman.log will note the completion of the DAGMan job
- *.dagman.out has detailed logging (look to first for errors)

Beyond the Basic DAG: Some Node-level Modifiers

PRE and POST scripts run on the submit server, as part of the node

```
my.dag
JOB A A.sub
SCRIPT POST A sort.sh
JOB B1 B1.sub
JOB B2 B2.sub
JOB B3 B3.sub
JOB C C.sub
SCRIPT PRE C tar it.sh
PARENT A CHILD B1 B2 B3
PARENT B1 B2 B3 CHILD C
```



 Use sparingly for lightweight work; otherwise include work in node jobs

RETRY failed nodes to overcome transient errors

 Retry a node up to N times if the exit code is non-zero:

RETRY node name N

Example:

```
JOB A A.sub

RETRY A 5

JOB B B.sub

PARENT A CHILD B
```

- See also: retry except for a particular exit code (UNLESS-EXIT), or retry scripts (DEFER)
- Note: Unnecessary for nodes (jobs) that can use max retries in the submit file

RETRY applies to whole node, including PRE/POST scripts

- PRE and POST scripts are included in retries
- RETRY of a node with a POST script uses the exit code from the POST script (not from the job)
 - POST script can do more to determine node success, perhaps by examining JOB output

Example:

```
SCRIPT PRE A download.sh
JOB A A.sub
SCRIPT POST A checkA.sh
RETRY A 5
```

SCRIPT Arguments and Argument Variables

```
JOB A A.sub
SCRIPT POST A checkA.sh my.out $RETURN
RETRY A 5
```

\$JOB: node name

\$JOBID: cluster.proc

\$RETURN: exit code of the job

\$PRE SCRIPT RETURN: exit code of PRE script

\$RETRY: current retry count

(more variables described in the manual)

<u>DAGMan Applications > DAG Input File > SCRIPT</u> <u>DAGMan Applications > Advanced Features > Retrying</u>

Modular Organization and Control of DAG Components

- Splices and SubDags
- Node Throttling
- Node Priorities
- Lots more in the Manual...

Additional Resources

HTCondor-Users Email List!

http://htcondor.org/mail-lists/

- Manual (and man pages) http://htcondor.readthedocs.org
- Nice HTCondor FAQs, examples, and documentation from our friends in Canary Islands:

https://is.gd/TjRvY8

 HTCondor HOWTO Recipes has FAQ on job submission http://wiki.htcondor.org/index.cgi/wiki?p=HowToAdminRecipes

THANK YOU!

ADDITIONAL DAGMAN SLIDES

Submit File Templates via VARS

 VARS line defines node-specific values that are passed into submit file variables

```
VARS node_name var1="value" [var2="value"]
```

 Allows a single submit file shared by all B jobs, rather than one submit file for each JOB.

my.dag

```
JOB B1 B.sub

VARS B1 data="B1" opt="10"

JOB B2 B.sub

VARS B2 data="B2" opt="12"

JOB B3 B.sub

VARS B3 data="B3" opt="14"
```

B.sub

```
...
InitialDir = $(data)
arguments = $(data).csv $(opt)
...
queue
```

SPLICE groups of nodes to simplify lengthy DAG files

my.dag

```
JOB A A.sub

SPLICE B B.spl

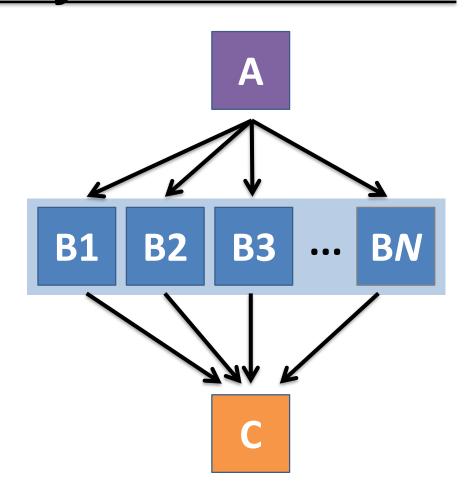
JOB C C.sub

PARENT A CHILD B

PARENT B CHILD C
```

B.spl

```
JOB B1 B1.sub
JOB B2 B2.sub
...
JOB BN BN.sub
```



Use nested SPLICEs with DIR for repeating workflow components

```
my.dag
```

```
JOB A A.sub DIR A

SPLICE B B.spl DIR B

JOB C C.sub DIR C

PARENT A CHILD B

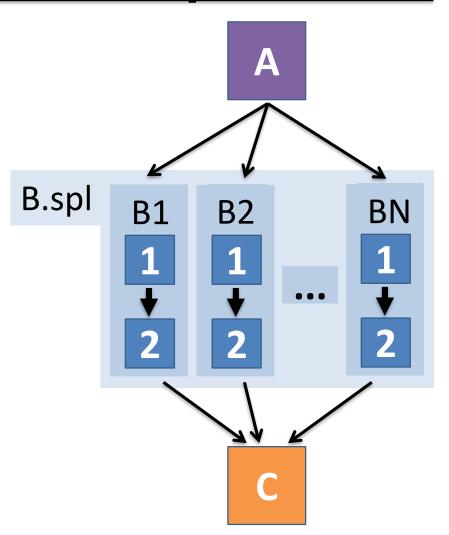
PARENT B CHILD C
```

B.spl

```
SPLICE B1 ../inner.spl DIR B1
SPLICE B2 ../inner.spl DIR B2
...
SPLICE BN ../inner.spl DIR BN
```

inner.spl

```
JOB 1 ../1.sub
JOB 2 ../2.sub
PARENT 1 CHILD 2
```



Use nested SPLICEs with DIR for repeating workflow components

```
my.dag
```

```
JOB A A.sub DIR A

SPLICE B B.spl DIR B

JOB C C.sub DIR C

PARENT A CHILD B

PARENT B CHILD C
```

B.spl

```
SPLICE B1 ../inner.spl DIR B1
SPLICE B2 ../inner.spl DIR B2
...
SPLICE BN ../inner.spl DIR BN
```

inner.spl

```
JOB 1 ../1.sub

JOB 2 ../2.sub

PARENT 1 CHILD 2
```

(dag dir)/

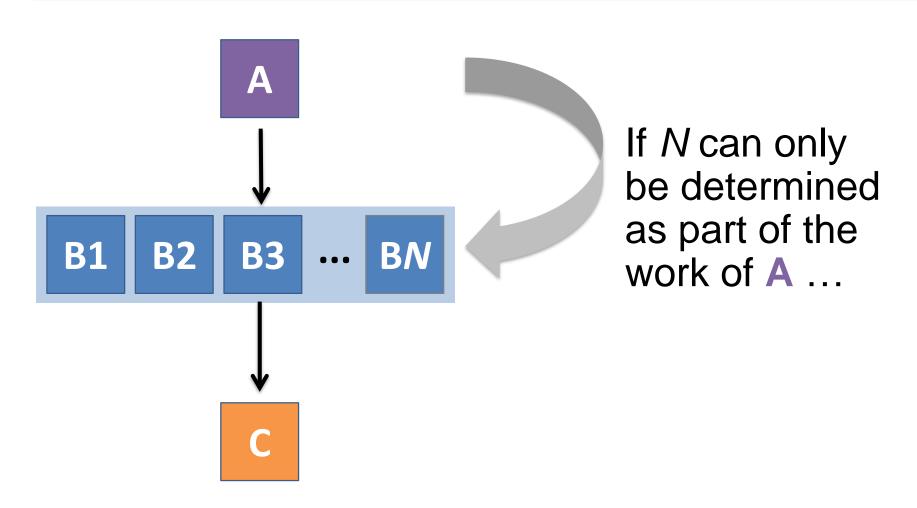
```
my.dag
A/ A.sub (A job files)
B/ B.spl inner.spl

1.sub 2.sub
B1/ (1-2 job files)
B2/ (1-2 job files)
...
BN/ (1-2 job files)
C/ C.sub (C job files)
```

More on SPLICE Behavior

- Upon submission of the outer DAG, nodes in the SPLICE(s) are added by DAGMan into the overall DAG structure.
 - A single DAGMan job is queued with single set of status files.
- Great for gradually testing and building up a large DAG (since a SPLICE file can be submitted by itself, as a complete DAG).
- SPLICE lines are not treated like nodes.
 - no PRE/POST scripts or RETRIES (though this may change)

What if some DAG components can't be known at submit time?



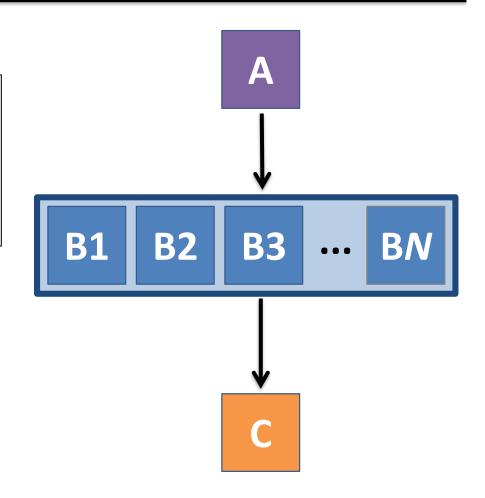
A SUBDAG within a DAG

my.dag

JOB A A.sub SUBDAG EXTERNAL B B.dag JOB C C.sub PARENT A CHILD B PARENT B CHILD C

B.dag (written by A)

JOB B1 B1.sub JOB B2 B2.sub JOB BN BN. sub



More on SUBDAG Behavior

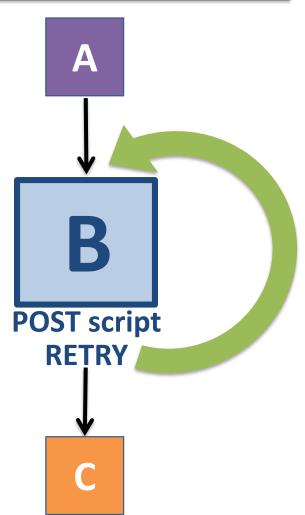
- WARNING: SUBDAGs should only be used (over SPLICES) when absolutely necessary!
 - Each SUBDAG EXTERNAL has it's own DAGMan job running in the queue.
- SUBDAGs are nodes (can have PRE/POST scripts, retries, etc.)
- A SUBDAG is not submitted until prior nodes in the outer DAG have completed.

Use a SUBDAG to achieve Cyclic Components within a DAG

- POST script determines whether another iteration is necessary; if so, exits non-zero
- RETRY applies to entire SUBDAG, which may include multiple, sequential nodes

my.dag

```
JOB A A.sub
SUBDAG EXTERNAL B B.dag
SCRIPT POST B iterateB.sh
RETRY B 1000
JOB C C.sub
PARENT A CHILD B
PARENT B CHILD C
```



DAG-level Control

Pause a running DAG with hold/release

Hold the DAGMan job process:

```
condor_hold dagman_jobID
```

- Pauses the DAG
 - No new node jobs submitted
 - Queued node jobs continue to run (including SUBDAGs), but no PRE/POST scripts
 - DAGMan jobs remains in the queue until released (condor_release) or removed

Pause a DAG with a halt file

- Create a file named <u>DAG_file.halt</u> in the same directory as the submitted DAG file
- Pauses the DAG
 - No new node jobs submitted
 - Queued node jobs, SUBDAGs, and POST scripts continue to run, but not PRE scripts
- DAGMan resumes after the file is deleted
 - If not deleted, the DAG creates rescue DAG file and exits after all queued jobs have completed

Throttle job nodes of large DAGs via DAG-level configuration

- If a DAG has many (thousands or more) jobs, performance of the submit server and queue can be assured by limiting:
 - Number of jobs in the queue
 - Number of jobs idle (waiting to run)
 - Number of PRE or POST scripts running
- Limits can be specified in a DAG-specific CONFIG file (recommended) or as arguments to condor submit dag

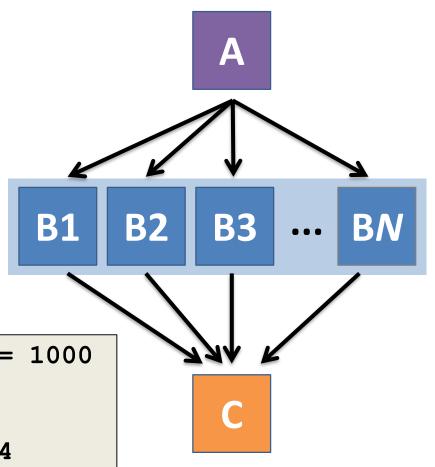
DAG-specific throttling via a **CONFIG** file

my.dag

JOB A A.sub SPLICE B B.dag JOB C C.sub PARENT A CHILD B PARENT B CHILD C CONFIG my.dag.config

my.dag.confiq

DAGMAN MAX JOBS SUBMITTED = 1000 DAGMAN MAX JOBS IDLE = 100 DAGMAN MAX PRE SCRIPTS = 4 DAGMAN MAX POST SCRIPTS = 4



Other DAGMan Features

Other DAGMan Features: Node-Level Controls

Set the PRIORITY of JOB nodes with:

```
PRIORITY node_name priority_value
```

 Use a PRE_SKIP to skip a node and mark it as successful, if the PRE script exits with a specific exit code:

```
PRE_SKIP node_name exit_code
```

Other DAGMan Features: Modular Control

- Append NOOP to a JOB definition so that its JOB process isn't run by DAGMan
 - Test DAG structure without running jobs (node-level)
 - Simplify combinatorial PARENT-CHILD statements (modular)
- Communicate DAG features separately with INCLUDE
 - e.g. separate file for JOB nodes and for VARS definitions, as part of the same DAG
- Define a CATEGORY to throttle only a specific subset of jobs

<u>DAGMan Applications > The DAG Input File > JOB</u>
<u>DAGMan Applications > Advanced Features > INCLUDE</u>
<u>DAGMan Applications > Advanced > Throttling by Category</u>

Other DAGMan Features: DAG-Level Controls

- Replace the <u>node_name</u> with <u>ALL_NODES</u> to apply a DAG feature to all nodes of the DAG
- Abort the entire DAG if a specific node exits with a specific exit code:

```
ABORT-DAG-ON node_name exit_code
```

 Define a FINAL node that will always run, even in the event of DAG failure (to clean up, perhaps).

```
FINAL node_name submit_file
```

<u>DAGMan Applications > Advanced > ALL NODES</u> <u>DAGMan Applications > Advanced > Stopping the Entire DAG</u> <u>DAGMan Applications > Advanced > FINAL Node</u>