

Workflows with HTCondor's DAGMan

Monday PM, Lecture 1

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Questions so far?



Goals for this Session

- Why Create a Workflow?
- Describing workflows as directed acyclic graphs (DAGs)
- Workflow execution via DAGMan (DAG Manager)



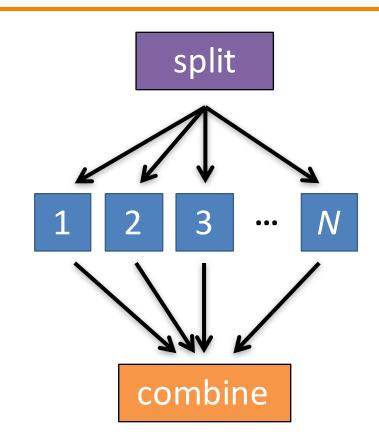
WHY WORKFLOWS? WHY DAGS?



Automation!

 Objective: Submit jobs in a particular order, automatically.

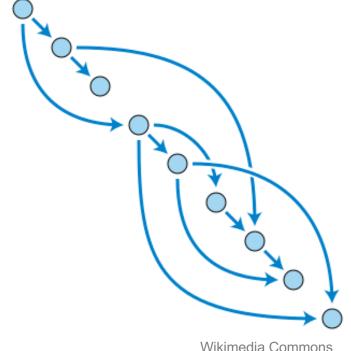
 Especially if: Need to replicate the same workflow multiple times in the future.





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 DAG workflows





DESCRIBING WORKFLOWS WITH DAGMAN



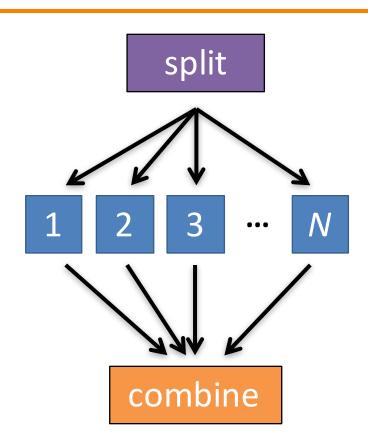
DAGMan in the HTCondor Manual





An Example HTC Workflow

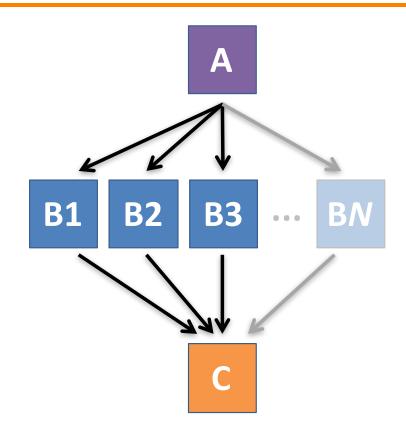
 User must communicate the "nodes" and directional "edges" of the DAG





Simple Example for this Tutorial

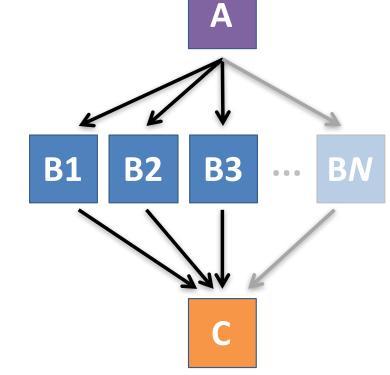
The DAG input file
 will communicate the
 "nodes" and directional
 "edges" of the DAG





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

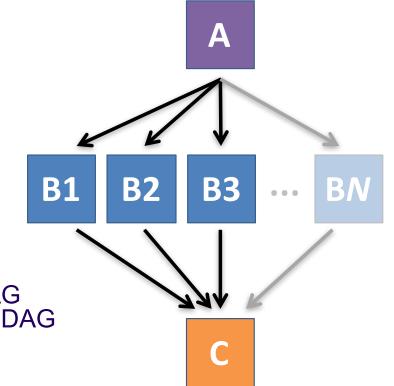
JOB **B3** B3.sub

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.





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
```

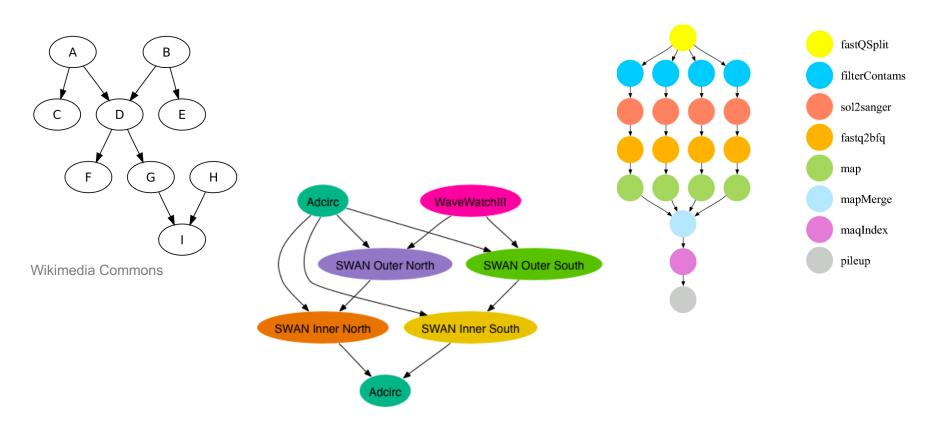
```
(dag_dir)/
```

```
A.sub B1.sub
B2.sub B3.sub
C.sub my.dag
(other job files)
```

- Node names and filenames can be anything.
- Node name and submit filename do not have to match.

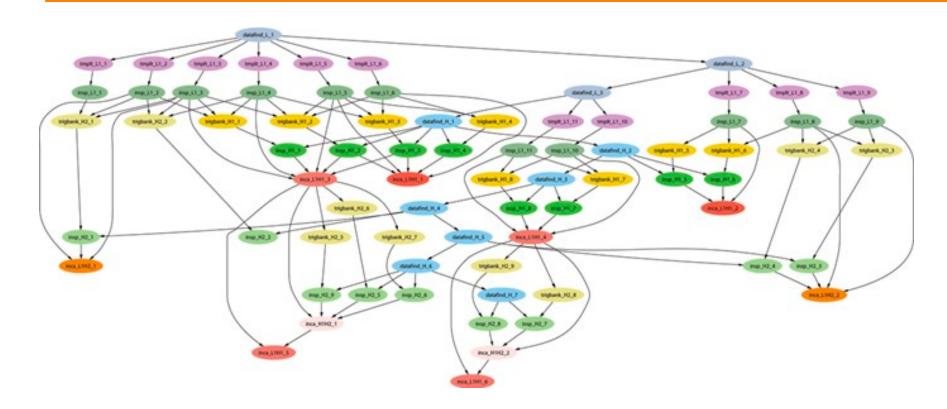


Endless Workflow Possibilities



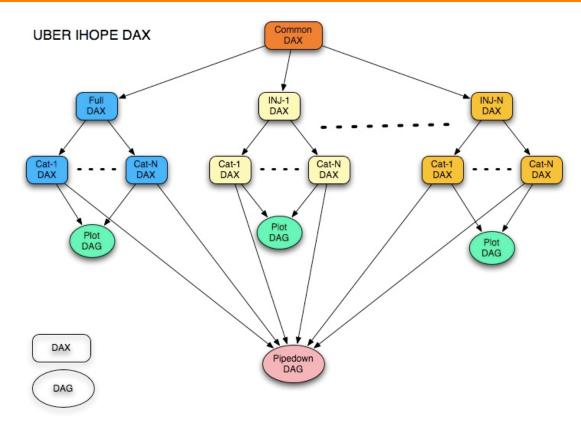


Endless Workflow Possibilities





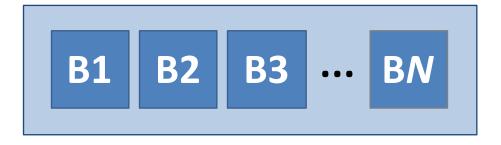
Repeating DAG Components!!



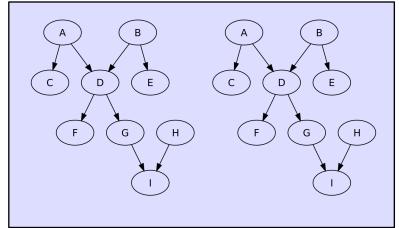


DAGs are also useful for nonsequential work

'bag' of HTC jobs



disjointed workflows





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

my.dag

JOB A A.sub

JOB **B1** B1.sub

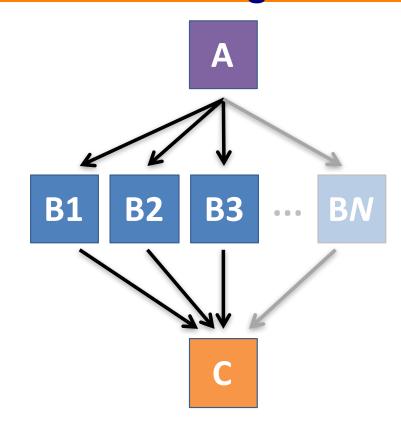
JOB **B2** B2.sub

JOB B3 B3.sub

JOB C C.sub

PARENT A CHILD B1 B2 B3

PARENT B1 B2 B3 CHILD C





SUBMITTING AND MONITORING A DAGMAN WORKFLOW



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.
```



A submitted DAG creates and DAGMan job in the queue

- DAGMan runs on the submit server, as a job in the queue
- At first:

```
$ condor q
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...
        BATCH NAME
OWNER
                      SUBMITTED DONE
                                        RUN
                                              IDLE
                                                   TOTAL
                                                          JOB IDS
alice my.dag+128  4/30 18:08
                                                          0.0
1 jobs; 0 completed, 0 removed, 0 idle, 1 running, 0 held, 0 suspended
$ condor q -nobatch
-- Schedd: submit-3.chtc.wisc.edu : <128.104.100.44:9618?...
       OWNER
                SUBMITTED
                             RUN TIME ST PRI SIZE CMD
 ID
128.0 alice 4/30 18:08 0+00:00:06 R 0
                                             0.3 condor dagman
1 jobs; 0 completed, 0 removed, 0 idle, 1 running, 0 held, 0 suspended
```



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?...
       BATCH NAME SUBMITTED DONE RUN IDLE
OWNER
                                            TOTAL JOB IDS
alice
      my.dag+128 4/30 18:08
                                                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?...
       OWNER
               SUBMITTED
ID
                            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 any other jobs
- *.dagman.out has DAGMan-specific 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

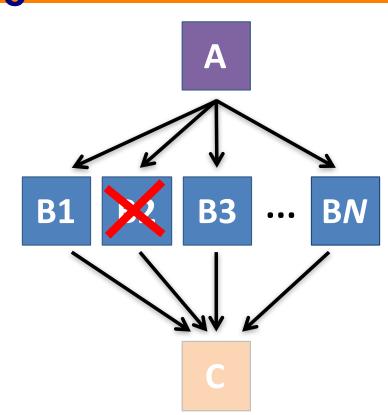
- A rescue file is created when:
 - a node fails, and after DAGMan advances through any other possible nodes
 - the DAG is removed from the queue (or aborted; covered later)
 - the DAG is halted and not unhalted (covered later)
- Resubmission uses the rescue file (if it exists) when the original DAG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



Node Failures Result in DAG

Failure

- If a node JOB fails (nonzero 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
-- 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 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</pre>
```

- 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 (remember the automatic rescue DAG file!)



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.dagman.out my.dag.lib.err my.dag.lib.out

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)



YOUR TURN!



Exercises!

- Ask questions!
- Lots of instructors around

- Coming up:
 - Now-3:15 Hands-on Exercises
 - -3:15-3:30 Break
 - 3:30 5:00 Workflows 2 & Hands-on