

Workflows with HTCondor's DAGMan

Monday, Lecture 4

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Questions so far?



Goals for this Session

- Describing workflows as directed acyclic graphs (DAGs)
- Workflow execution via DAGMan (DAG Manager)
- Node-level options in a DAG
- Modular organization of DAG components
- Additional DAGMan Features



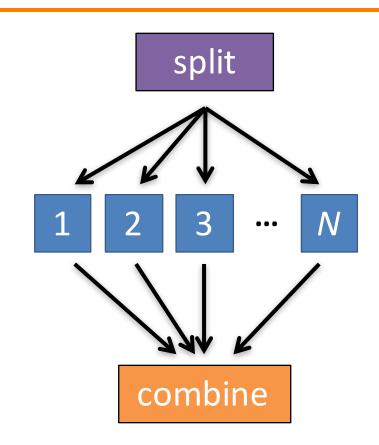
WHY WORKFLOWS? WHY DAGS?



Automation!

 Objective: Submit jobs in a particular order, automatically.

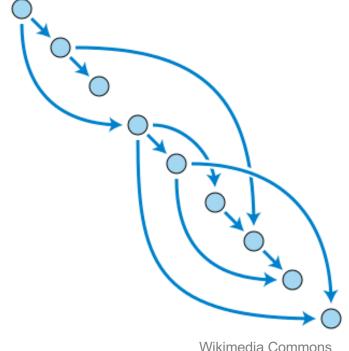
 Especially if: Need to replicate the same workflow multiple times in the future.





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



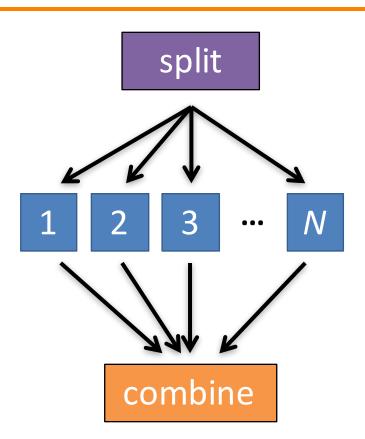
DAGMan in the HTCondor Manual





An Example HTC Workflow

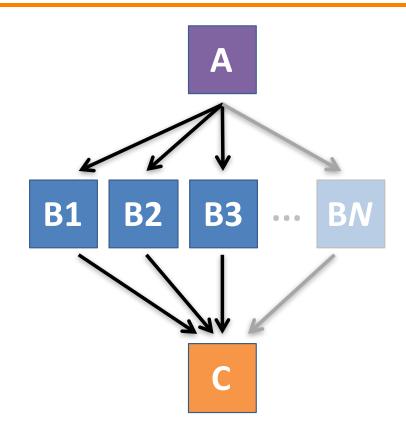
 User must communicate the "nodes" and directional "edges" of the DAG





Simple Example for this Tutorial

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

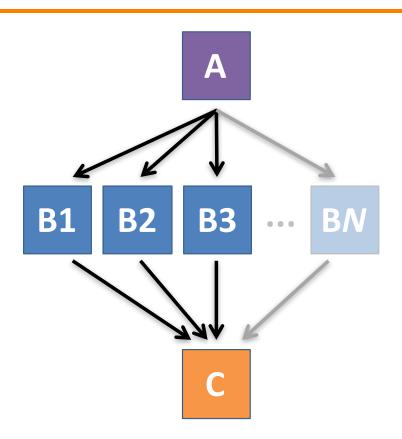




Simple Example for this Tutorial

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

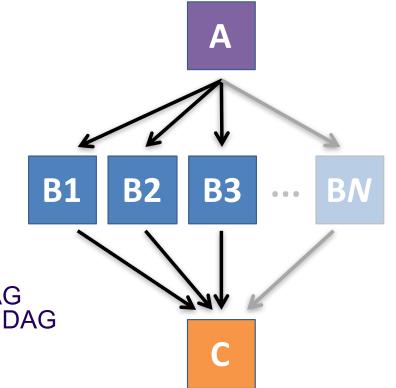
JOB **B3** B3.sub

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
```

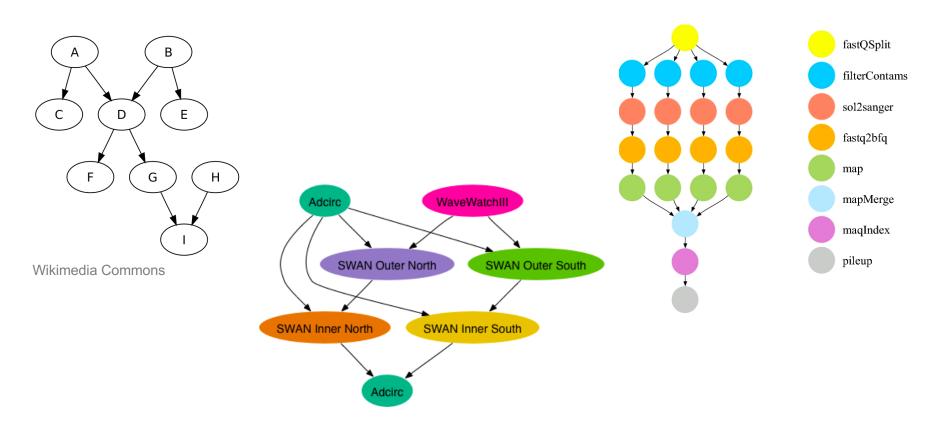
```
(dag_dir)/
```

```
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.

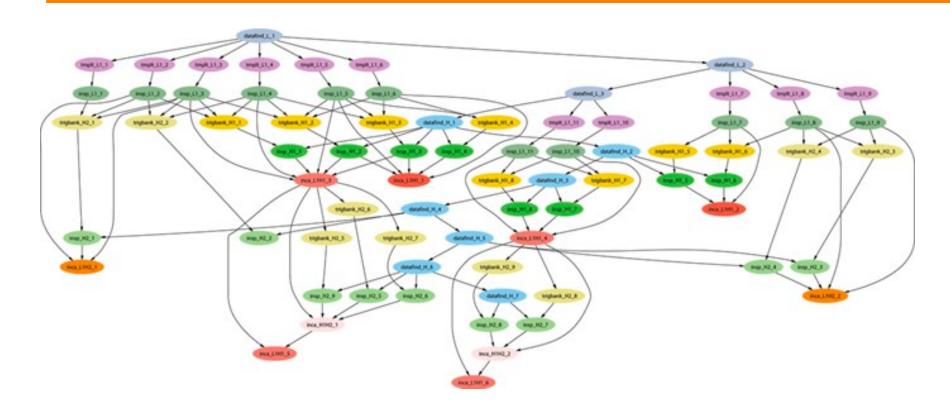


Endless Workflow Possibilities



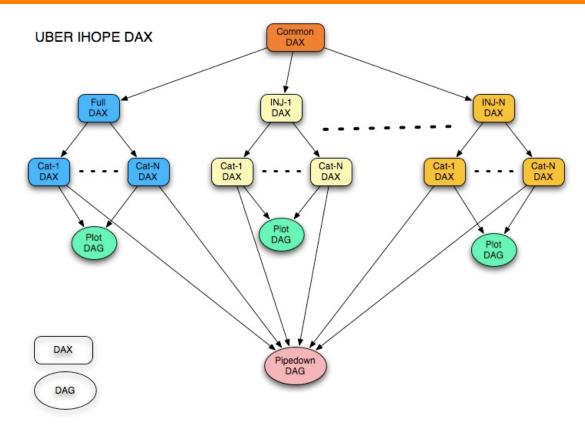


Endless Workflow Possibilities





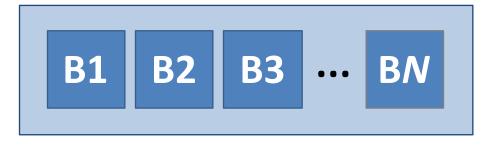
Repeating DAG Components!!



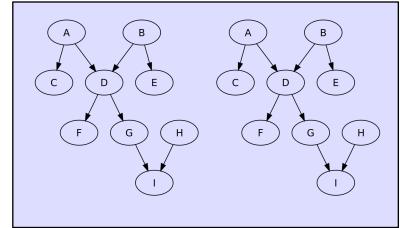


DAGs are also useful for nonsequential work

'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

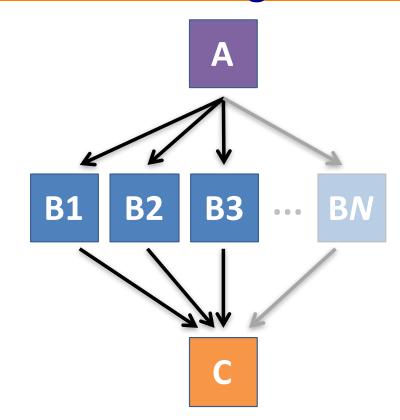
JOB **B2** B2.sub

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

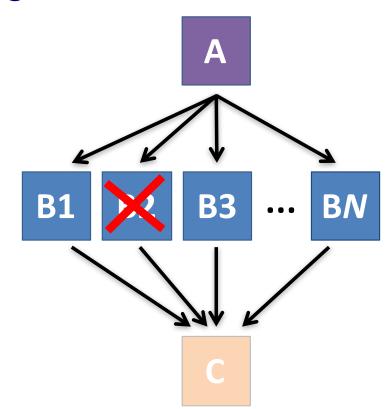
- A rescue file is created when:
 - a node fails, and after DAGMan advances through any other possible nodes
 - the DAG is removed from the queue (or aborted; covered later)
 - the DAG is halted and not unhalted (covered later)
- Resubmission uses the rescue file (if it exists) when the original DAG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

- 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 (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)



BEYOND THE BASIC DAG: NODE-LEVEL MODIFIERS



Default File Organization

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
```

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

 What if you want to organize files into other directories?



Node-specific File Organization with *DIR*

DIR sets the submission directory of the node

my.dag

```
JOB A A.sub DIR A

JOB B1 B1.sub DIR B

JOB B2 B2.sub DIR B

JOB B3 B3.sub DIR B

JOB C C.sub DIR C

PARENT A CHILD B1 B2 B3

PARENT B1 B2 B3 CHILD C
```

(dag_dir)/

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



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

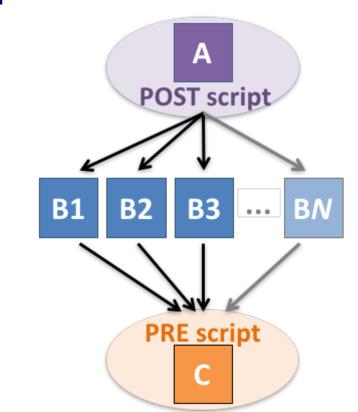
JOB C C.sub

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





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 node

\$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> DAGMan Applications > Advanced Features > Retrying



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

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

DAGMan Applications > Advanced Features > Retrying
DAGMan Applications > DAG Input File > SCRIPT



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:

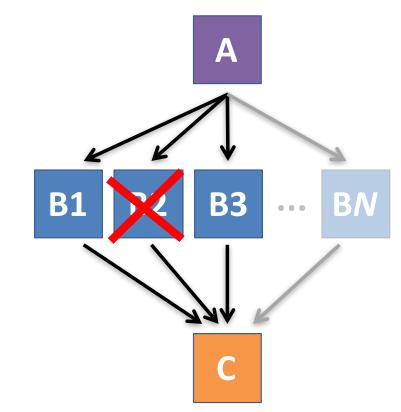
```
SCRIPT PRE A download.sh
JOB A A.sub
SCRIPT POST A checkA.sh
RETRY A 5
```

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



Best Control Achieved with One Process per JOB Node

- While submit files can 'queue' many processes, a single process per submit file is best for DAG JOBs
 - Failure of any process in a
 JOB node results in failure of
 the entire node and
 immediate removal of other
 processes in the node.
 - RETRY of a JOB node retries the entire submit file.





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
```



MODULAR ORGANIZATION OF DAG COMPONENTS

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SPLICE groups of nodes to simplify lengthy DAG files

my.dag

```
JOB A A.sub

SPLICE B B.spl

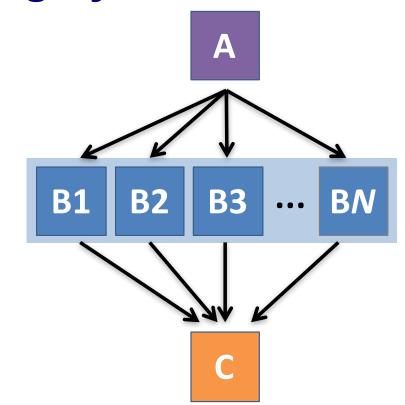
JOB C C.sub

PARENT A CHILD B

PARENT B CHILD C
```

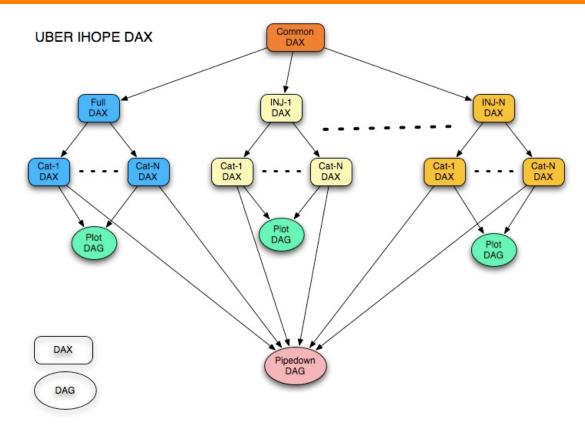
B.spl

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





Repeating DAG Components!!





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
```

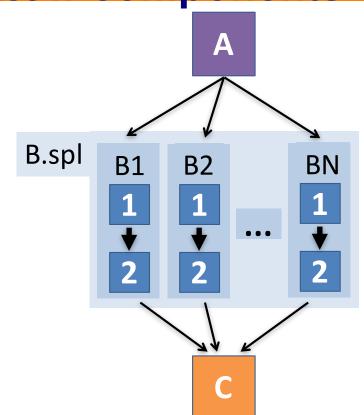
B.spl

PARENT B CHILD C

```
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

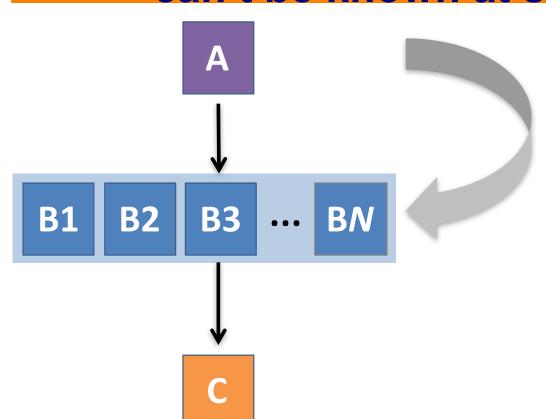
PARENT 1 CHILD 2
```

(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 \text{ job files})
     C.sub (C job files)
```



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



If *N* can only be determined as part of the work of **A** ...



A SUBDAG within a DAG

my.dag

```
JOB A A.sub

SUBDAG EXTERNAL B B.dag

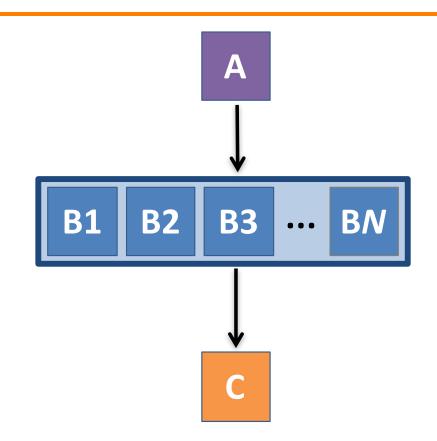
JOB C C.sub

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
```





Much More at the end of the presentation and in the HTCondor Manual!!!

https://research.cs.wisc.edu/htcondor/manual/current/2_Users_Manual.html



YOUR TURN!

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DAGMan Exercises!

- Ask questions!
- Lots of instructors around

- Coming up:
 - now–5:00pm Hands-On Exercises
 - 5:00pm on On Your Own

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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)



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, on the submit server.
- SUBDAGs are nodes in the outer DAG (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 a Cyclic Component 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

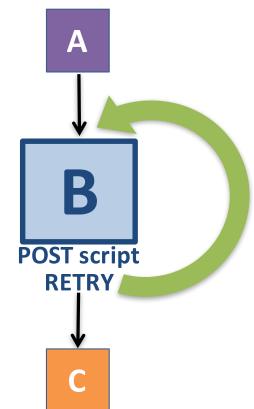
SCRIPT POST B iterateB.sh

RETRY B 1000

JOB C C.sub

PARENT A CHILD B

PARENT B CHILD C
```





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>

DAGMan Applications > Advanced > Throttling by Category



Other DAGMan Features: DAG-Level Controls

- Replace the node_name with ALL_NODES 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>