

Intermediate Condor: DAGMan Monday, July 19th, 1:15pm

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Before we begin...

 Any questions on the lectures or exercises up to this point?





<u>DAGMan</u>

Directed Manager

Acyclic

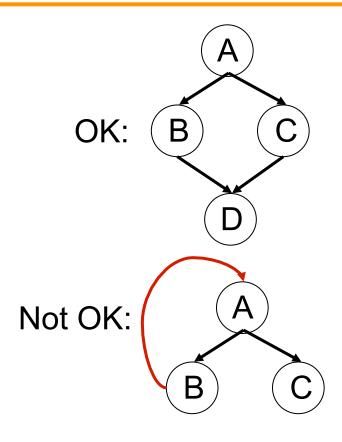
Graph

- DAGMan allows you to specify the dependencies between your Condor jobs, so it can manage them automatically for you.
- Example: "Don't run job B until job A has completed successfully."



What is a DAG?

- A DAG is the data structure used by DAGMan to represent these dependencies.
- Each job is a node in the DAG.
- Each node can have any number of "parent" or "children" nodes – as long as there are no loops!



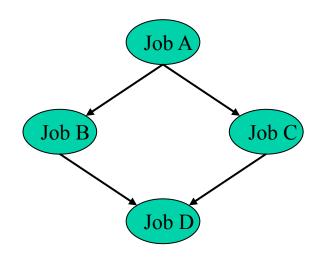


Defining a DAG

 A DAG is defined by a .dag file, listing each of its nodes and their dependencies:

```
Job A a.sub
Job B b.sub
Job C c.sub
Job D d.sub
```

Parent A Child B C Parent B C Child D



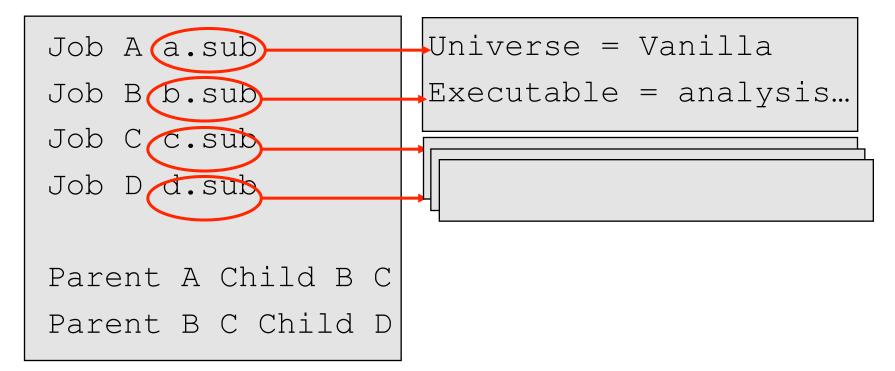


DAG Files....

The complete DAG has five files

One DAG File:

Four Submit Files:





Submitting a DAG

 To start your DAG, just run condor_submit_dag with your .dag file, and Condor will start a personal DAGMan process which to begin running your jobs:

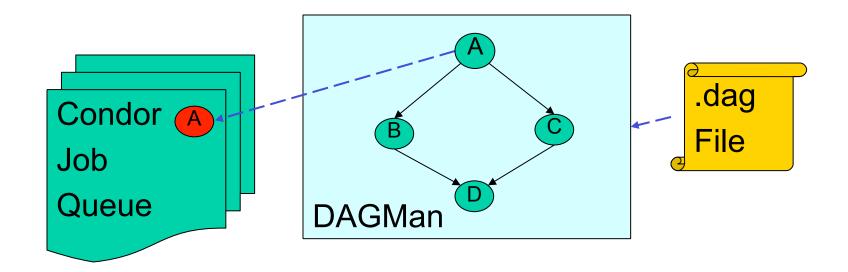
```
% condor_submit_dag diamond.dag
```

- condor_submit_dag submits a Scheduler Universe job with DAGMan as the executable.
- Thus the DAGMan daemon itself runs as a Condor job, so you don't have to baby-sit it.



Running a DAG

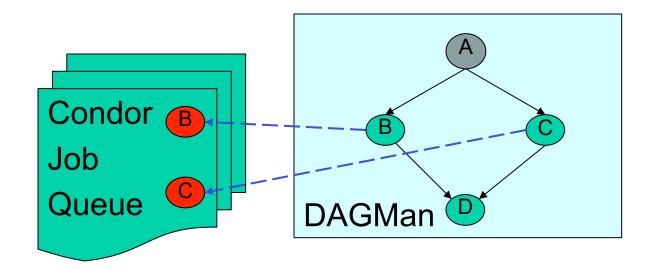
 DAGMan acts as a scheduler, managing the submission of your jobs to Condor based on the DAG dependencies.





Running a DAG (cont'd)

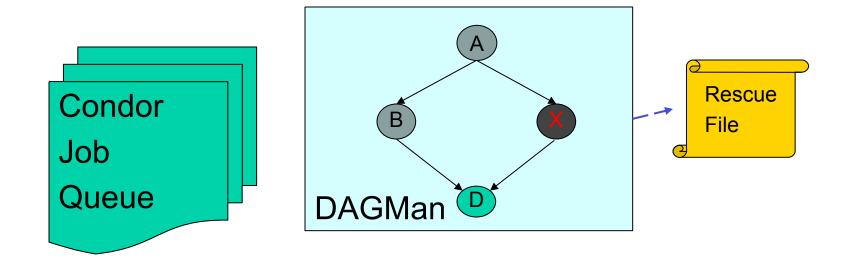
 DAGMan holds & submits jobs to the Condor queue at the appropriate times.





Running a DAG (cont'd)

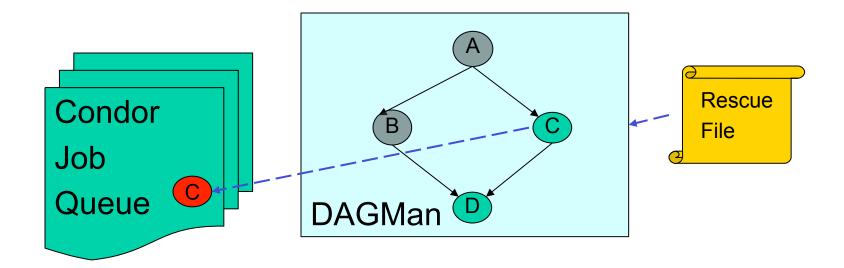
 In case of a job failure, DAGMan continues until it can no longer make progress, and then creates a "rescue" file with the current state of the DAG.





Recovering a DAG

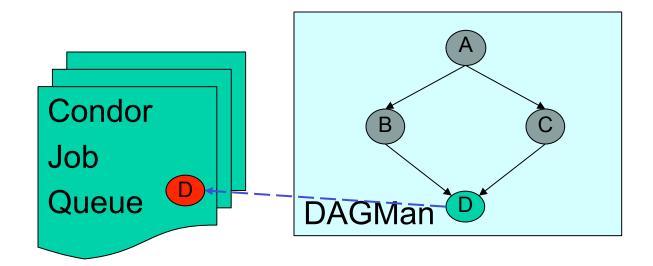
 Once the failed job is ready to be re-run, the rescue file can be used to restore the prior state of the DAG.





Recovering a DAG (cont'd)

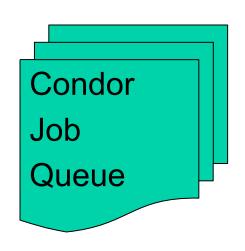
 Once that job completes, DAGMan will continue the DAG as if the failure never happened.

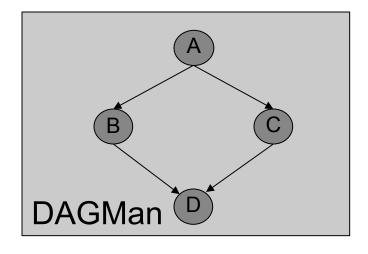




Finishing a DAG

 Once the DAG is complete, the DAGMan job itself is finished, and exits.

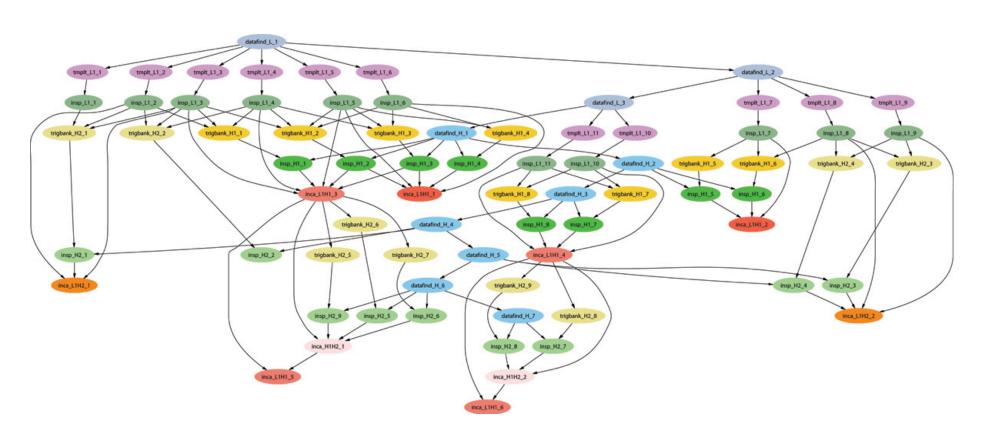








Example of a LIGO Inspiral DAG





Use of Condor by the LIGO Scientific Collaboration



- Condor handles 10's of millions of jobs per year running on the LDG, and up to 500k jobs per DAG.
- Condor standard universe check pointing widely used, saving us from having to manage this.
- At Caltech, 30 million jobs processed using 22.8 million CPU hrs. on 1324 CPUs in last 30 months.
- For example, to search 1 yr. of data for GWs from the inspiral of binary neutron star and black hole systems takes ~2 million jobs, and months to run on several thousand ~2.6 GHz nodes.



DAGMan & Fancy Features

- DAGMan doesn't have a lot of "fancy features"
 - No loops
 - Not much assistance in writing very large DAGs (script it yourself)
- Focus is on solid core
 - Add the features people need in order to run large DAGs well.
 - People build systems on top of DAGMan



DAGMan & Log Files

- For each job, Condor generates a log file
- DAGMan reads this log to see what has happened
- If DAGMan dies (crash, power failure, etc...)
 - Condor will restart DAGMan
 - DAGMan re-reads log file
 - DAGMan knows everything it needs to know



Advanced DAGMan Tricks

- Throttles and degenerative DAGs
- Sub-DAGs
- Pre and Post scripts: editing your DAG



Throttles

- Failed nodes can be automatically retried a configurable number of times
 - Can retry N times
 - Can retry N times, unless a node returns specific exit code
- Throttles to control job submissions
 - Max jobs submitted
 - Max scripts running



Degenerative DAG

- Submit DAG with:
 - -200,000 nodes







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- No dependencies
- Use DAGMan to throttle the jobs
 - Condor is scalable, but it will have problems if you submit 200,000 jobs simultaneously
 - DAGMan can help you get scalability even if you don't have dependencies

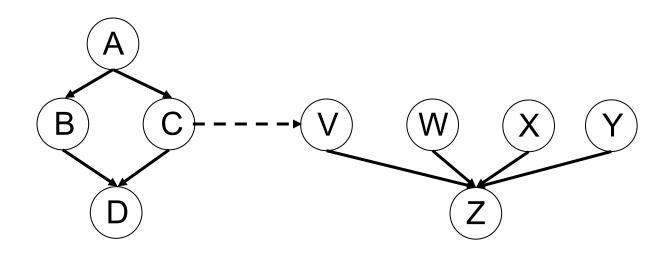


Sub-DAG

- Idea: any given DAG node can be another DAG
 - SUBDAG External Name DAG-file
- DAG node will not complete until recursive DAG finishes,
- Interesting idea: A previous node could generate this DAG node
- Why?
 - Simpler DAG structure
 - Implement a fixed-length loop
 - Modify behavior on the fly



Sub-DAG





DAGMan scripts

- DAGMan allows pre & post scripts
 - Don't have to be scripts: any executable
 - Run before (pre) or after (post) job
 - Run on the same computer you submitted from

Syntax:

```
JOB A a.sub

SCRIPT PRE A before-script $JOB

SCRIPT POST A after-script $JOB $RETURN
```



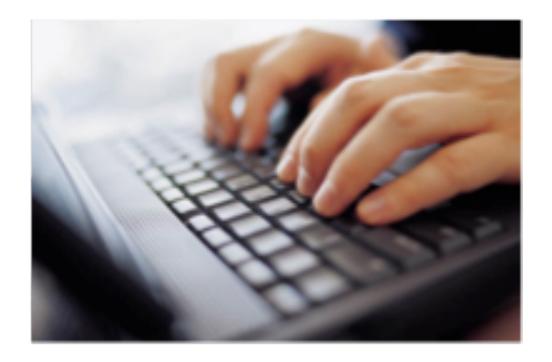
So What?

- Pre script can make decisions
 - Where should my job run? (Particularly useful to make job run in same place as last job.)
 - Generate Sub-DAG
 - Should I pass different arguments to the job?
 - Lazy decision making
 - Noop jobs
- Post script can change return value
 - DAGMan decides job failed in non-zero return value
 - Post-script can look at {error code, output files, etc} and return zero or non-zero based on deeper knowledge.



Let's try it out!

Exercises with DAGMan.



OSG Summer School 2010 25

Open Science Grid

Questions?

- Questions? Comments?
- Feel free to ask me questions later:
 Alain Roy <roy@cs.wisc.edu>
- Upcoming sessions
 - Now 3:15pm
 - Hands-on exercises
 - -3:15pm 3:30pm
 - Break
 - 3:30pm 5:00pm
 - Final Condor session

26