

Intermediate Condor: Workflows

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Before we begin...

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





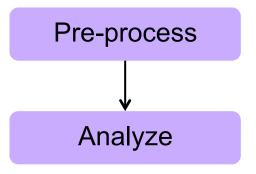
Workflows

- What if you have a complex set of programs to run for your science?
- For example:
 - You want to analyze a set of images
 - Each image needs to be pre-processed
 - Each image needs to be analyzed
 - You need to summarize the results of all the analyses
 - Each of these is a separate program

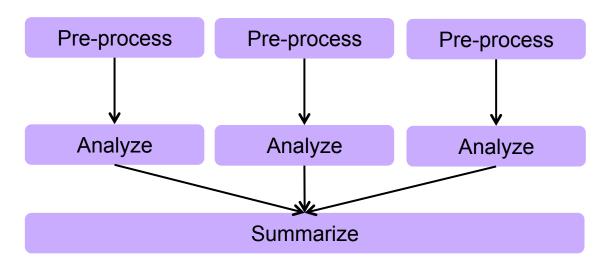


Workflows

One Image:



Three Images:





Workflows: definition

Definition 1:

A set of steps to complete a complex task

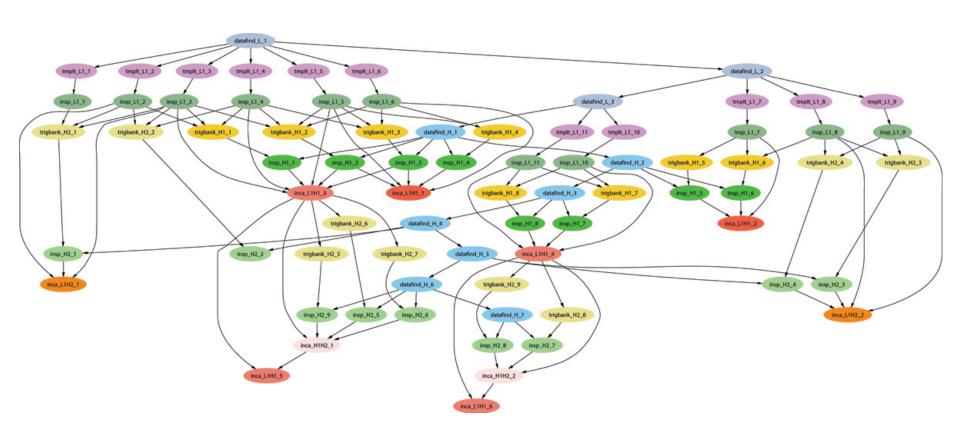
Definition 2:

A graph of jobs to run: some jobs need to run before others while other jobs can run in parallel





Example of a LIGO Inspiral DAG



LIGO

Use of Condor by the LIGO Scientific Collaboration

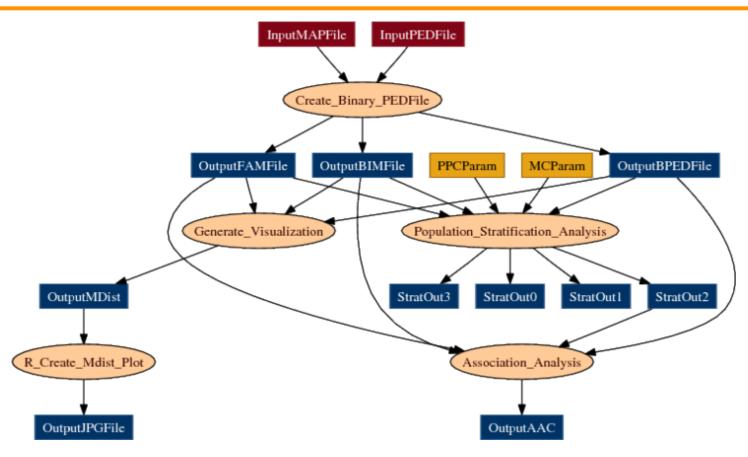


- Condor handles 10's of millions of jobs per year running on the LDG, and up to 500k jobs per DAG.
- Condor standard universe check pointing widely used, saving us from having to manage this.
- At Caltech, 30 million jobs processed using 22.8 million CPU hrs. on 1324 CPUs in last 30 months.
- For example, 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.

(Statement from 2010—"last 30 months" isn't from now. Also, I think they do up to 1 million jobs per DAG now.)



Example workflow: Bioinformatics



From Mason, Sanders, State (Yale)

http://pegasus.isi.edu/applications/association_test



LEGEND

Sub Workflow

Compute Job

Example workflow: Astronomy

Remote Tile Setup - Runs mDAGGalacticPlane on Compute Site to generate the DAX and Replica Catalog with input URL pointing to cache at ISI - Runs for 45 minutes Local Tile Setup - Prepares the environment for running the montage sub workflow . 1 ... 900 Montage 5 Montage 5 Degree Workfow Degree Workfow

Remote Extra Cleanup Job - Delete extra untracked files

mBgModel

mBackground

mlmgTbl

mAdd

mShrink

mJpeg

mProjectPP

mDiffFit

mConcatFit

Montage Galactic Plane Workflow

From Berriman & Good (JPAC)

http://pegasus.isi.edu/applications/galactic-plane



DAGMan

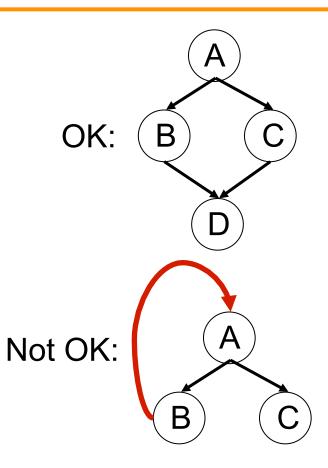
- DAGMan:
 - Directed Acyclic Graph (DAG) Manager (Man)
- Allows you to specify the dependencies between your Condor jobs
- Manages the jobs and their dependencies

That is, it manages a workflow of Condor jobs



What is a DAG?

- A DAG is the 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!



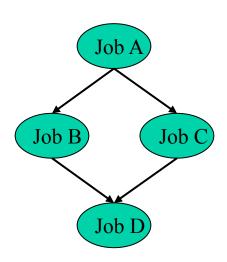


Defining a DAG

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

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

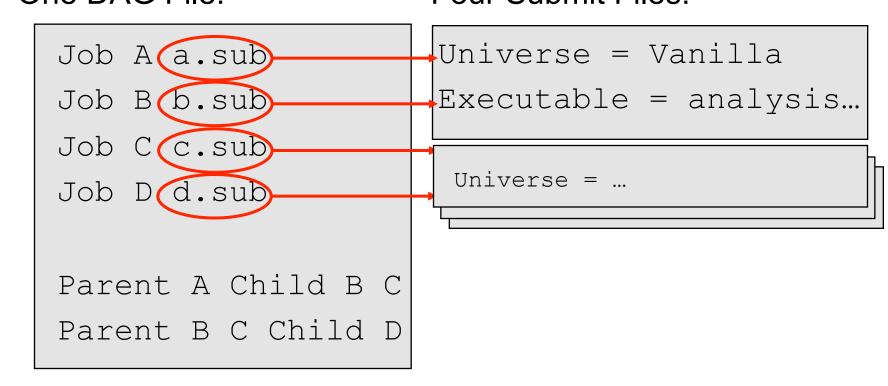
Parent A Child B C Parent B C Child D





DAG Files....

This 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 DAGMan process to manage your jobs:

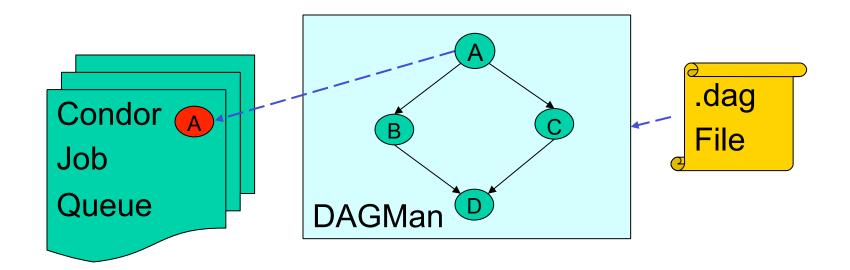
```
% condor submit dag diamond.dag
```

- 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

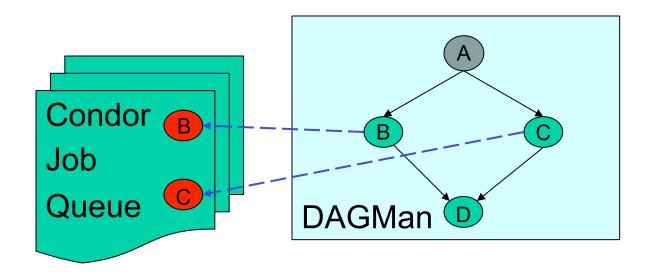
 DAGMan acts as a scheduler, managing the submission of your jobs to Condor based on the DAG dependencies





Running a DAG (cont'd)

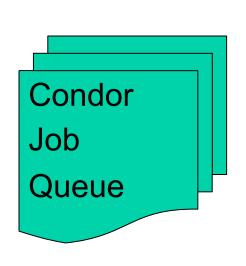
- DAGMan submits jobs to Condor at the appropriate times
- For example, after A finishes, it submits B & C

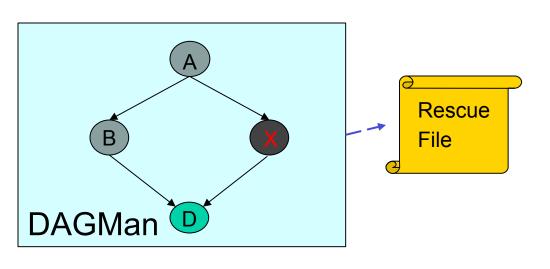




Running a DAG (cont'd)

- A job fails if it exits with a non-zero exit code
- In case of a job failure, DAGMan runs other jobs until it can no longer make progress, and then creates a "rescue" file with the current state of the DAG

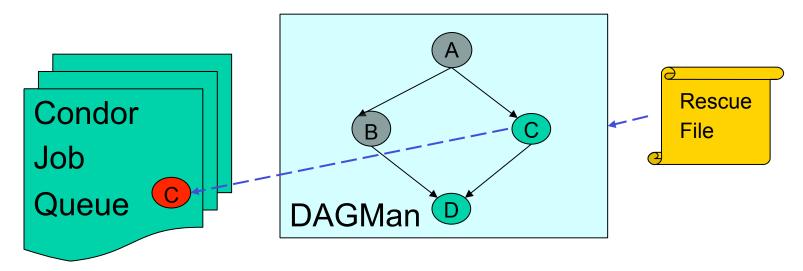






Recovering a DAG

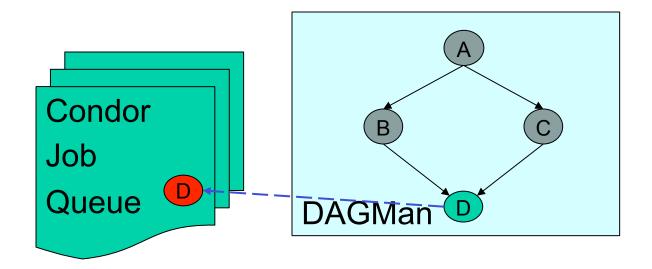
- Once the failed job is ready to be re-run, the rescue file can be used to restore the prior state of the DAG
 - Another example of reliability for HTC!





Recovering a DAG (cont'd)

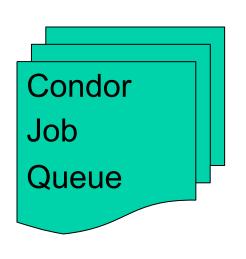
 Once that job completes, DAGMan will continue the DAG as if the failure never happened

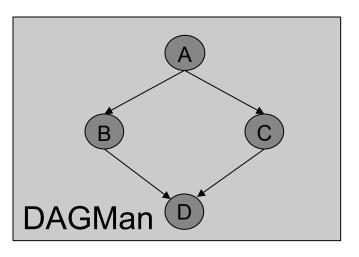




Finishing a DAG

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







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



Related Software

Pegasus: http://pegasus.isi.edu/

- Writes DAGs based on abstract description
- Runs DAG on appropriate resource (Condor, OSG, EC2...)
- Locates data, coordinates execution
- Uses DAGMan, works with large workflows

Makeflow: http://nd.edu/~ccl/software/makeflow/

- User writes make file, not DAG
- Works with Condor, SGE, Work Queue...
- Handles data transfers to remote systems
- Does not use DAGMan



DAGMan: Reliability

- 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
 - Principle: DAGMan can recover state from files and without relying on a service (Condor queue, database...)
- Recall: HTC requires reliability!



Advanced DAGMan Tricks

- Throttles
- DAGs without dependencies
- Sub-DAGs
- Pre and Post scripts: editing your DAG



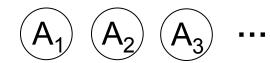
Throttles

- Failed nodes can be automatically retried a configurable number of times
 - Helps recover from jobs that crash some percentage of the time
- Throttles to control job submissions
 - Max jobs submitted
 - Max scripts running
 - These are important when working with large DAGs



DAGs without dependencies

- Submit DAG with:
 - -200,000 nodes



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

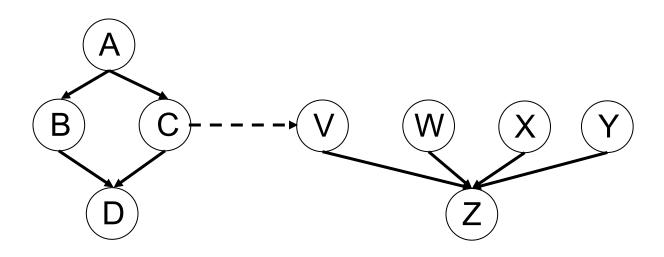


Sub-DAG

- Idea: any given DAG node can be another DAG
 - SUBDAG External Name DAG-file
- DAG node will not complete until sub-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
 - Run before (pre) or after (post) job
 - Run on the same computer you submitted from
 - Don't have to be scripts: any executable

Syntax:

```
JOB A a.sub

SCRIPT PRE A before-script $JOB

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



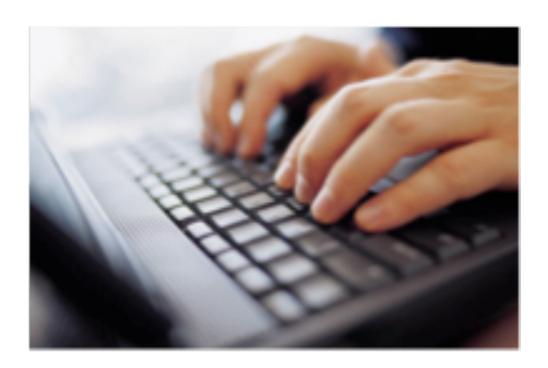
So What?

- Pre script can make decisions
 - Where should my job run? (Particularly useful to make job run in same place as last job.)
 - What should my job do?
 - Generate Sub-DAG
- 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.





Questions?

- Questions? Comments?
- Feel free to ask me questions later:
 - Rob Quick rquick@iu.edu