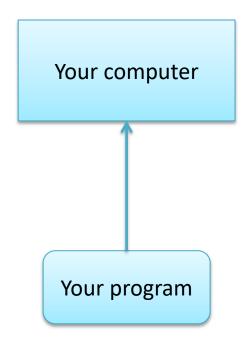
# An overview of batch processing

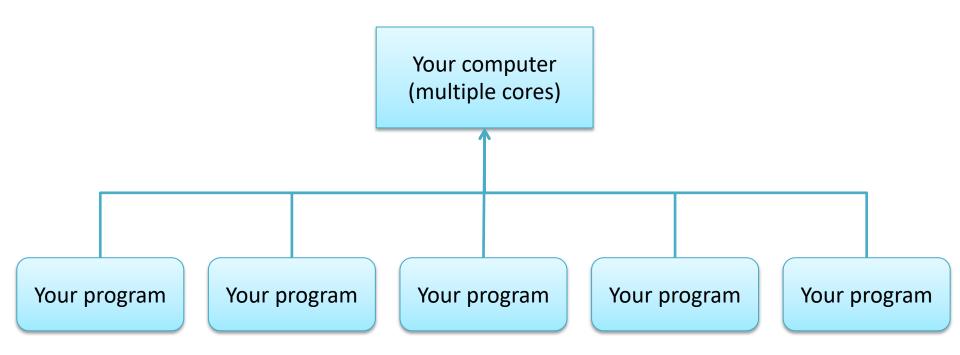
30-May-2024

#### One-on-one

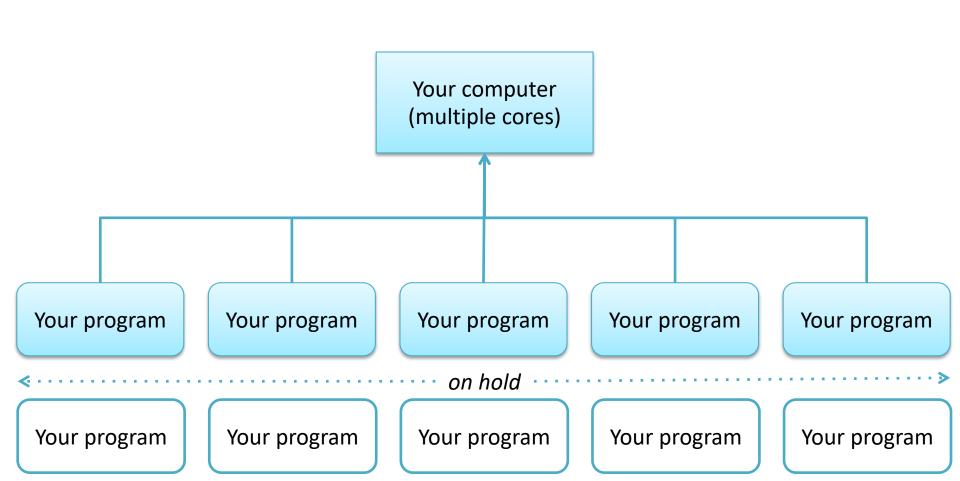


One way to speed things up within ROOT: multi-threaded code (RDataFrame, PROOF) Your computer (multiple cores) Your program (multiple threads) One thread One thread One thread One thread One thread Outside of ROOT, you can have multiple programs on a single computer (UNIX command "at")

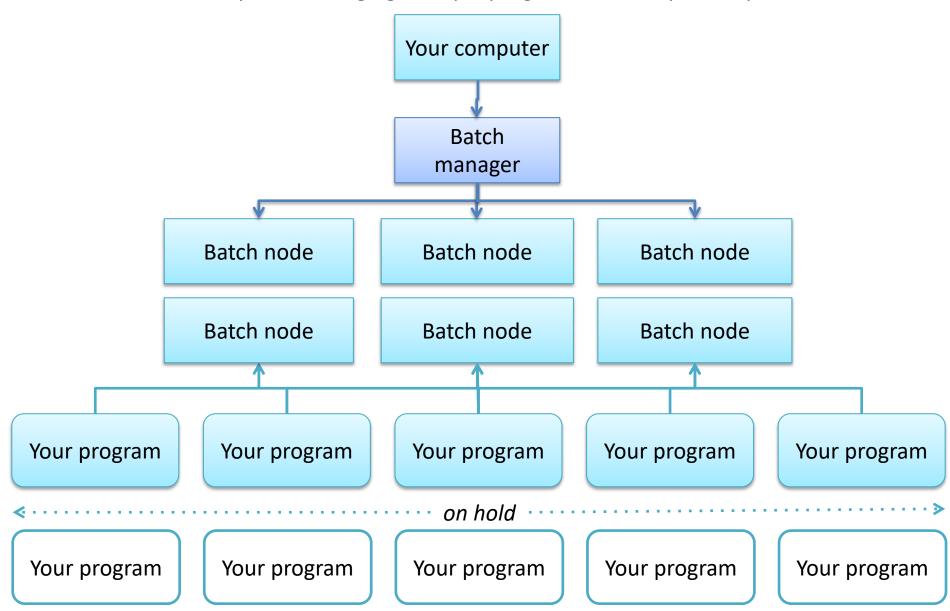
You have to remember not to submit too many jobs or you'll overload the computer



A batch system managing multiple programs on a single computer (UNIX command "batch")



A batch system managing multiple programs on multiple computers



The standard software for managing batch systems in scientific computing is HTCondor (or just Condor)

#### Main web page

http://research.cs.wisc.edu/htcondor/

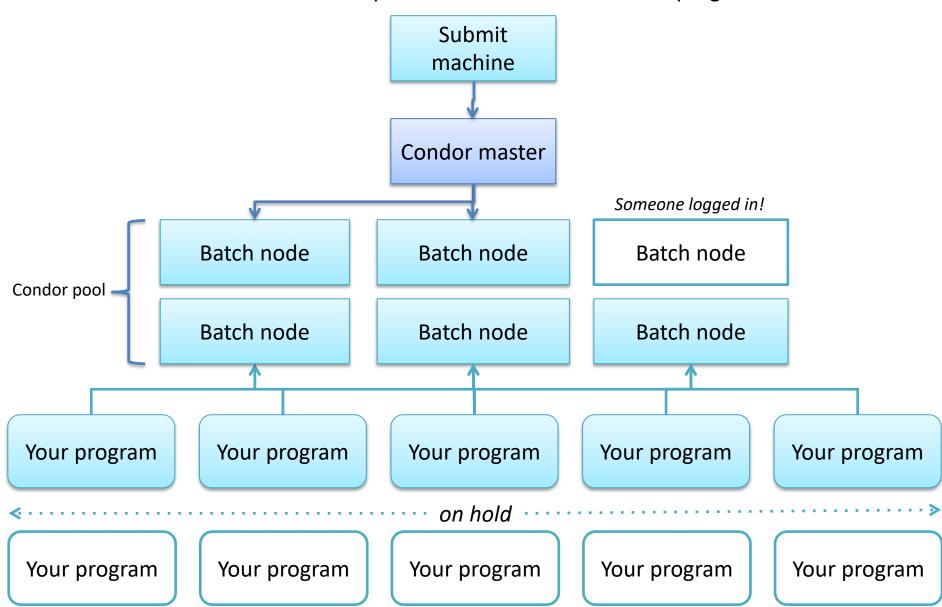
#### Quick start

https://htcondor.readthedocs.io/en/latest/users-manual/quick-start-guide.html

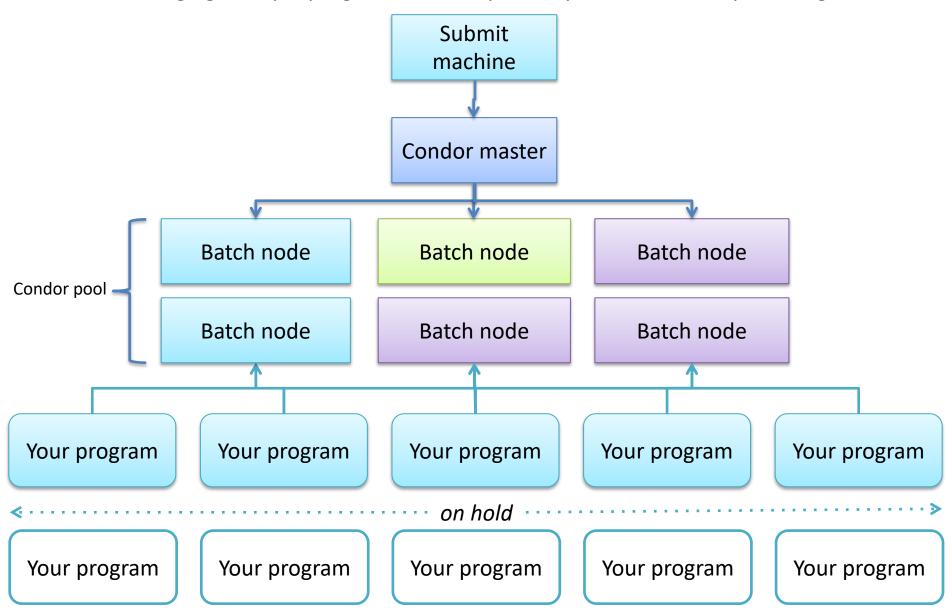
#### Full manual

https://htcondor.readthedocs.io/en/latest/

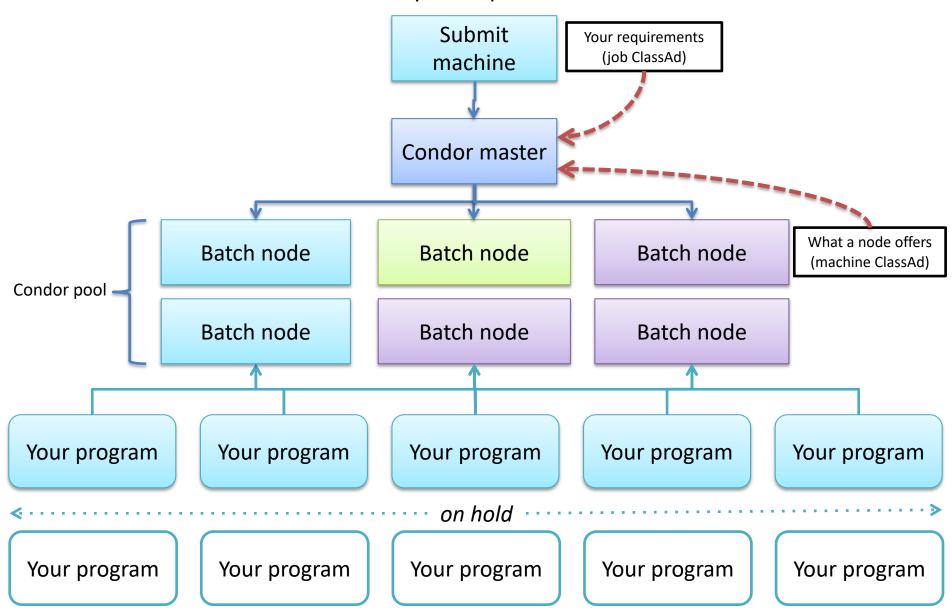
 Stick to the "vanilla" universe; the "standard" universe won't work for ROOT or any other particle-physics software (so you don't need condor\_compile). Condor will halt a queue in favor of an interactive program



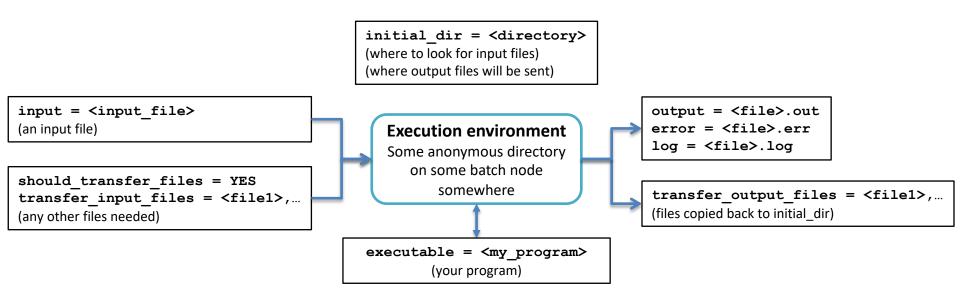
Condor managing multiple programs on multiple computers with multiple configurations



Condor uses "ClassAds" to match your requirements with what each node offers



- Condor can't do everything for you.
- Think about input files (including programs) and output files and how they'll be accessed.
- Think about disk space. "df -h" and "du -shx \*" can help.
- Fun fact: The particle-physics Condor pools can't see your home directory!
- Moral: Let condor transfer your files... when possible.

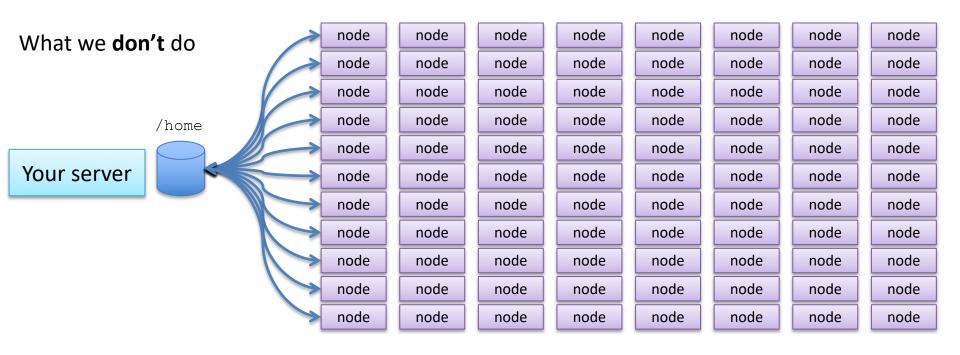


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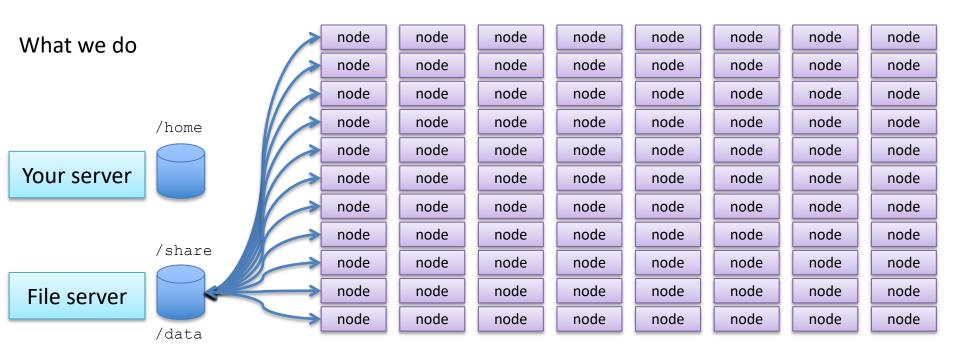
When you can't let condor transfer your files, here are disk-sharing methods outside of condor:

- NFS used at Nevis
- CVMFS Fermilab and CERN
- Grid, BlueArc only used at Fermilab

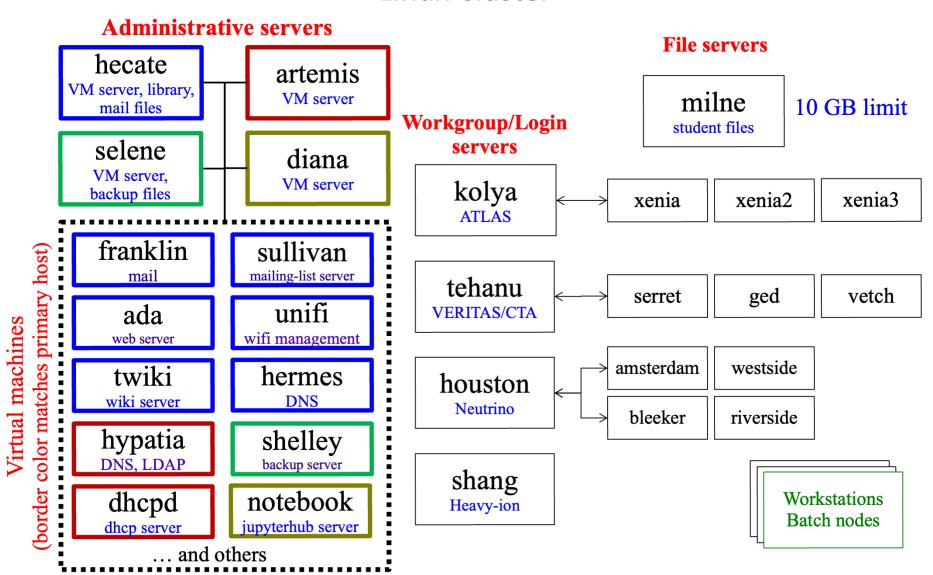
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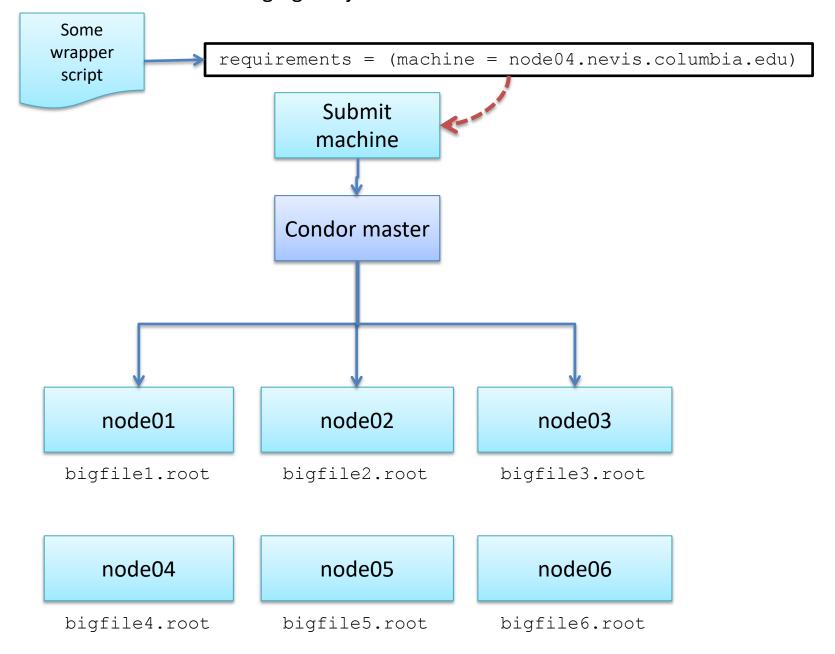


# Particle-Physics Computer Systems Linux Cluster



https://twiki.nevis.columbia.edu/twiki/bin/view/Main/LinuxClusterhttps://twiki.nevis.columbia.edu/twiki/bin/view/Main/ListOfMachineshttps://twiki.nevis.columbia.edu/twiki/bin/view/Main/ListOfMachineshttps://

#### Bringing the job to the data



# Final tips

- Split up your task so each condor job takes 20-60 minutes
- If your job must be preempted, it will have to run from the beginning on the same machine that cancelled the job
- Test your job with one process before submitting it for 10,000 processes!

### Resources

Main web page

http://research.cs.wisc.edu/htcondor/

#### Quick start

https://htcondor.readthedocs.io/en/latest/users-manual/quick-start-guide.html

**Full manual** 

https://htcondor.readthedocs.io/en/latest/

Nevis particle-physics condor guide

https://twiki.nevis.columbia.edu/twiki/bin/view/Nevis/Condor

Basic Condor@Nevis tutorial

https://www.nevis.columbia.edu/~seligman/rootclass/html/appendix/batch/index.html