

# The Landscape of Academic Research Computing

Rob Quick <rquick@iu.edu>

Chief Operations Officer - Open Science Grid

Manager High Throughput Computing

Some Slides Contributed by the University of Wisconsin  
HTCondor Team and Scot Kronenfeld





# Let's jump right in...

$$\begin{aligned}
 E_n^{(1)} &= V_{nn} \\
 E_n^{(2)} &= \frac{|V_{nk_2}|^2}{E_{nk_2}} \\
 E_n^{(3)} &= \frac{V_{nk_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_3}} - V_{nn} \frac{|V_{nk_3}|^2}{E_{nk_3}^2} \\
 E_n^{(4)} &= \frac{V_{nk_4} V_{k_4 k_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_3} E_{nk_4}} - \frac{|V_{nk_4}|^2 |V_{nk_2}|^2}{E_{nk_4}^2 E_{nk_2}} - V_{nn} \frac{V_{nk_4} V_{k_4 k_3} V_{k_3 n}}{E_{nk_3}^2 E_{nk_4}} - V_{nn} \frac{V_{nk_4} V_{k_4 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_4}^2} + V_{nn}^2 \frac{|V_{nk_4}|^2}{E_{nk_4}^3} \\
 &= \frac{V_{nk_4} V_{k_4 k_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_3} E_{nk_4}} - E_n^{(2)} \frac{|V_{nk_4}|^2}{E_{nk_4}^2} - 2V_{nn} \frac{V_{nk_4} V_{k_4 k_3} V_{k_3 n}}{E_{nk_3}^2 E_{nk_4}} + V_{nn}^2 \frac{|V_{nk_4}|^2}{E_{nk_4}^3} \\
 E_n^{(5)} &= \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 k_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_3} E_{nk_4} E_{nk_5}} - \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 n} |V_{nk_2}|^2}{E_{nk_4}^2 E_{nk_5} E_{nk_2}} - \frac{V_{nk_5} V_{k_5 k_2} V_{k_2 n} |V_{nk_2}|^2}{E_{nk_2} E_{nk_5}^2 E_{nk_2}} - \frac{|V_{nk_5}|^2 V_{nk_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_5}^2 E_{nk_2} E_{nk_3}} \\
 &\quad - V_{nn} \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 k_3} V_{k_3 n}}{E_{nk_3}^2 E_{nk_4} E_{nk_5}} - V_{nn} \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_4}^2 E_{nk_5}} - V_{nn} \frac{V_{nk_5} V_{k_5 k_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_3} E_{nk_5}^2} + V_{nn} \frac{|V_{nk_5}|^2 |V_{nk_3}|^2}{E_{nk_5}^2 E_{nk_3}^2} + 2V_{nn} \frac{|V_{nk_5}|^2 |V_{nk_2}|^2}{E_{nk_5}^3 E_{nk_2}} \\
 &\quad + V_{nn}^2 \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 n}}{E_{nk_4}^3 E_{nk_5}} + V_{nn}^2 \frac{V_{nk_5} V_{k_5 k_3} V_{k_3 n}}{E_{nk_3}^2 E_{nk_5}^2} + V_{nn}^2 \frac{V_{nk_5} V_{k_5 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_5}^3} - V_{nn}^3 \frac{|V_{nk_5}|^2}{E_{nk_5}^4} \\
 &= \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 k_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_3} E_{nk_4} E_{nk_5}} - 2E_n^{(2)} \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 n}}{E_{nk_4}^2 E_{nk_5}} - \frac{|V_{nk_5}|^2 V_{nk_3} V_{k_3 k_2} V_{k_2 n}}{E_{nk_5}^2 E_{nk_2} E_{nk_3}} \\
 &\quad - 2V_{nn} \left( \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 k_3} V_{k_3 n}}{E_{nk_3}^2 E_{nk_4} E_{nk_5}} - \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 k_2} V_{k_2 n}}{E_{nk_2} E_{nk_4}^2 E_{nk_5}} + \frac{|V_{nk_5}|^2 |V_{nk_3}|^2}{E_{nk_5}^2 E_{nk_3}^2} + 2E_n^{(2)} \frac{|V_{nk_5}|^2}{E_{nk_5}^3} \right) \\
 &\quad + V_{nn}^2 \left( 2 \frac{V_{nk_5} V_{k_5 k_4} V_{k_4 n}}{E_{nk_4}^3 E_{nk_5}} + \frac{V_{nk_5} V_{k_5 k_3} V_{k_3 n}}{E_{nk_3}^2 E_{nk_5}^2} \right) - V_{nn}^3 \frac{|V_{nk_5}|^2}{E_{nk_5}^4}
 \end{aligned}$$

# Who Am I?



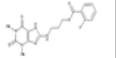


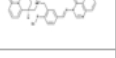



- Chief Operations Officer of the Open Science Grid and the Software Assurance Marketplace
- Manager High Throughput Computing (IU)
- IU Institutional OSG PI
- PI – Robust PID Project
- Co-PI Science Node – (Formally - International Science Grid This Week)
- Co-chair CODATA School of Research Data Science Task Group
- External Advisory Board for EGI
- Member of the Organizational Advisory Board for RDA

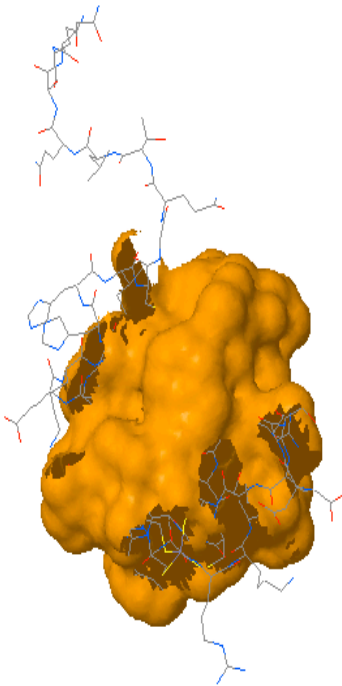
# Protein Docking Project at the IU School of Medicine

- SPLINTER - Structural Protein-Ligand Interactome
- Used autodock-vina – “...open-source program for drug discovery, molecular docking and virtual screening...”
- First run in 2013 - docked ~3900 Proteins with 5000 Ligands for a total of ~19M docked pairs.
- Submitted via command line to Condor using Pegasus on the OSG-XSEDE submission node
- Infrastructure is set and new runs can be easily started

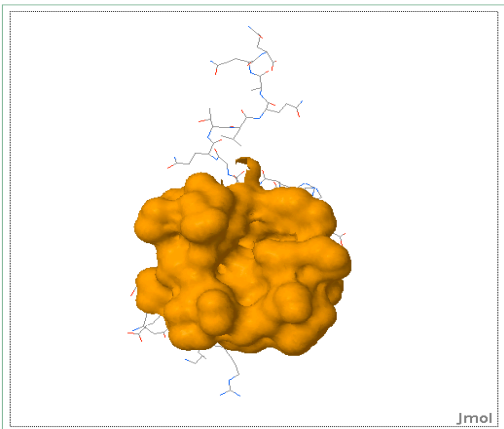


•Various rotations of Protein CBFA2T1 (Cyclin-D-related protein) (Eighty two protein) (Protein ETO) (Protein MTG8) (Zinc finger MYND domain-containing protein 2)

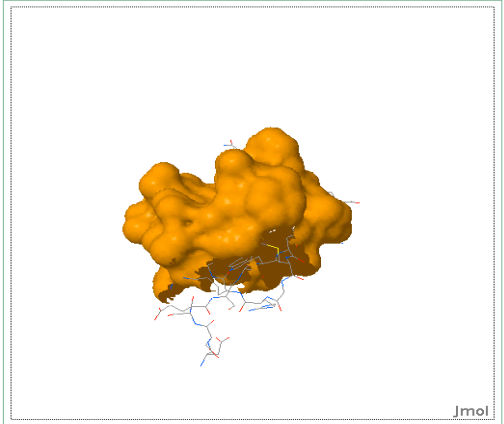
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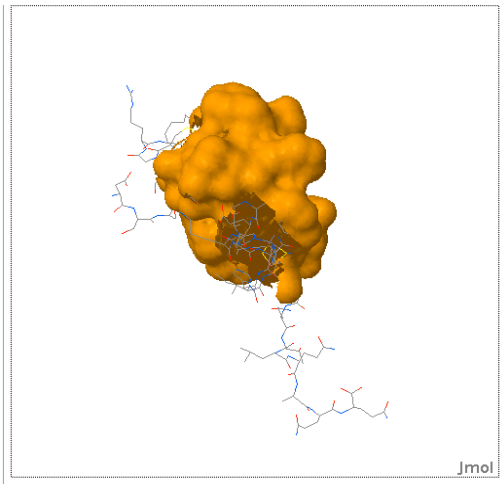
Jmol



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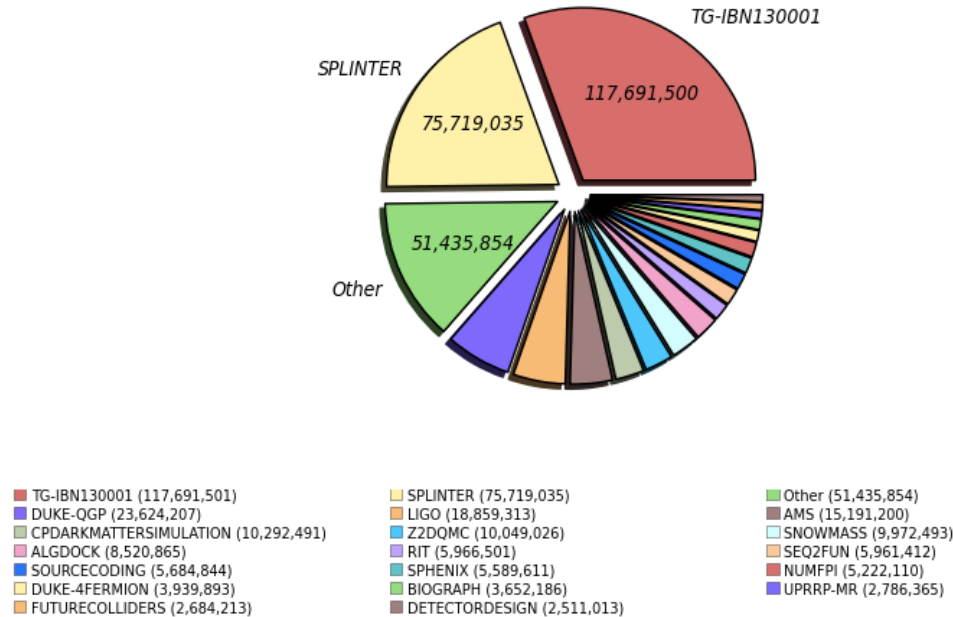


Jmol



# Some Numbers

Wall Hours by VO (Sum: 385,354,132 Hours)  
1309 Days from Week 00 of 2013 to Week 31 of 2016

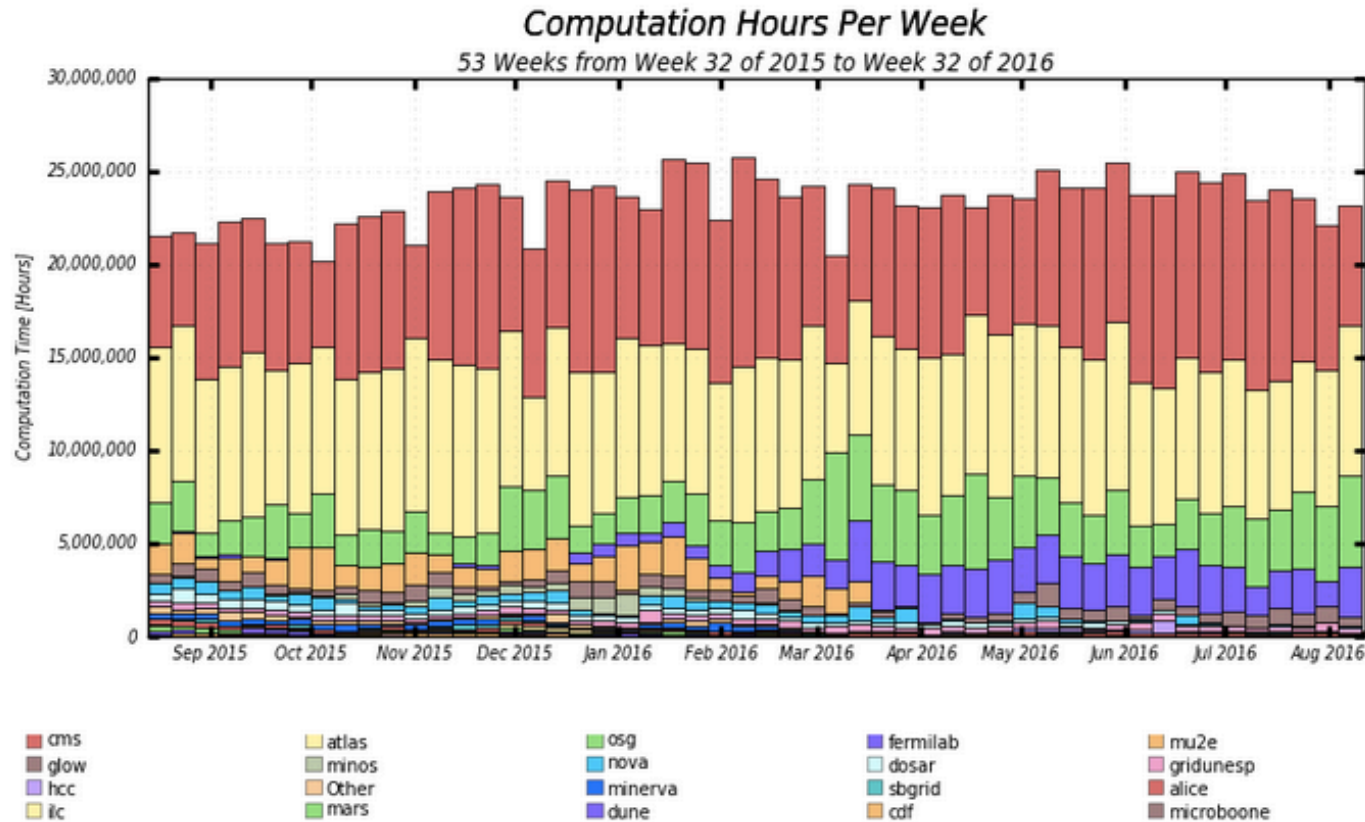


- Amazon EC2 Computing \$0.073/hour
- \$5.5M Compute Only
- Data Transfer and Storage Not Included



# At what scale?

## Job Activity



Maximum: 25,760,175 Hours, Minimum: 284,123 Hours, Average: 22,932,758 Hours, Current: 284,123 Hours



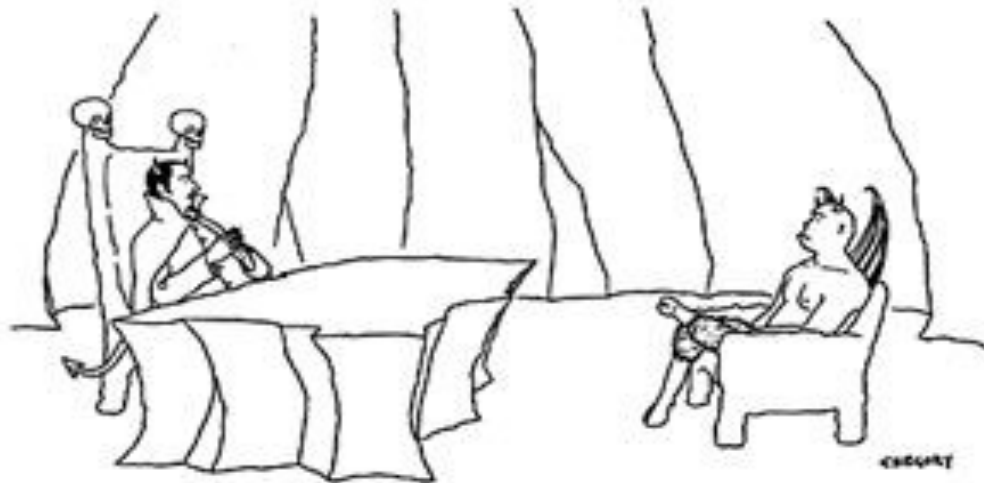
# Follow Along at:

<https://opensciencegrid.github.io/dosar/Materials/TriesteMaterials/>



# Overview of day

- Lectures alternating with exercises
  - Emphasis on lots of exercises
  - Hopefully overcome PowerPoint fatigue



*"I need someone well versed in the art of torture—do you know PowerPoint?"*

# Some thoughts on the exercises

- It's okay to move ahead on exercises if you have time
- It's okay to take longer on them if you need to
- If you move along quickly, try the “On Your Own” sections and “Challenges”

# Most important!

- Please ask questions!
  - ...during the lectures
  - ...during the exercises
  - ...during the breaks
  - ...during the meals
  - ...over dinner
  - ...via email after we depart ([rquick@iu.edu](mailto:rquick@iu.edu))
- If I don't know, I'll find the right person to answer your question.

# Goals for this session

- Define Local, Clustered, High Throughput Computing (HTC), High Performance Computing (HPC), Cloud Computing (XaaS), and Containers
- Shared, Allocated, and Purchased
- What is HTCondor? And why are we using it?

# The setup: You have a problem

- Your science computing is complex!
  - Monte carlo, image analysis, genetic algorithm, simulation...
- It will take a year (CPU time) to get the results on your laptop, but your paper is due in a week.
- What do you do?

# Option 1: Wait a year





# Option 2: Local Clustered Computing

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- Easy access to additional nodes
- Local support for porting to environment (maybe)
- Often a single type of resource
- Often running at capacity





# Option 3: Use a “supercomputer” aka High Performance Computing(HPC)

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- “Clearly, I need the best, fastest computer to help me out”
- Maybe you do...
  - Do you have a highly parallel program?
    - i.e. individual modules must communicate
  - Do you require the fastest network/disk/memory?
- Are you willing to:
  - Port your code to a special environment?
  - Request and wait for an allocation?





# Option 4: Use lots of commodity computers

- Instead of the fastest computer, lots of individual computers
- May not be fastest network/disk/memory, but you can access a lot of them
- Job can be broken down into separate, independent pieces
  - If I give you more computers, you run more jobs
  - You care more about total quantity of results than instantaneous speed of computation
- This is **high-throughput computing**





# Option 5: Buy (or Borrow) some computing from a Cloud Provider

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- Unlimited resources (if you can afford them)
- Full administrative access to OS of the resources you 'buy'
- Specialized VM images reducing effort in porting
- XaaS Business Model



# These are All Valid Options

- Remember the problem you have one month to publish results for your conference
  - Option 1: You will miss your deadline
  - Option 2: You might miss your deadline – But if your lucky you'll make it (or if you know the admin)
  - Option 3: If you have parallelized code and can get an allocation you have a good chance
  - Option 4: If you can serialize your workflow you have a good chance
  - Option 5: You can meet your deadline for a price. Though academic clouds are becoming more available.

# Computing Infrastructures

- Local Laptop/Desktop – Short jobs with small data
- Local Cluster – Larger jobs and larger data but subject to availability
- HPC – Prime performance with parallelized code
- HTC – Sustained computing over a long period for serialized
- Cloud – Need deeper permission on an OS and have deeper pockets



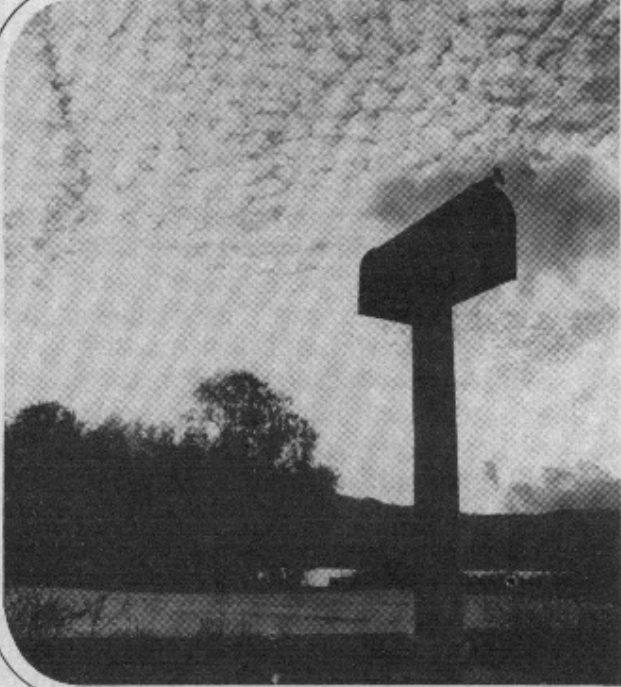
# Why focus on high-throughput computing? (HTC)

- An approach to distributed computing that focuses on long-term throughput, not instantaneous computing power
  - We don't care about operations per second
  - We care about operations per year
- Implications:
  - Focus on reliability
  - Use all available resources
    - Any Linux based machine can participate



*Only that shall happen  
Which has happened,  
Only that occur  
Which has occurred;  
There is nothing new  
Beneath the sun!*

Ecclesiastes Chapter 1 verse 9



What is a "Distributed"  
Data Processing System?

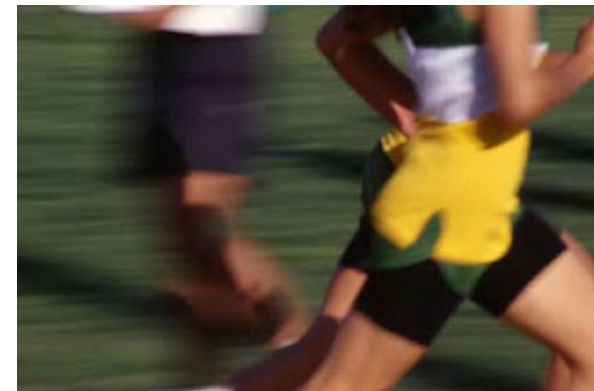
Philip H. Enslow, Jr.  
Georgia Institute of Technology

- HTC is about many jobs, many users, many servers, many sites and many workflows.
- What's in a buzzword?
  - “I was doing cloud computing since before it was called grid computing.”



# Think about a race

- Assume you can run a four minute mile
- Does that mean you can run a 104 minute marathon?
- The challenges in sustained computation are different than achieving peak in computation speed



# An example problem: BLAST

- A scientist has:
  - Question: Does a protein sequence occur in other organisms?
  - Data: lots of protein sequences from various organisms
  - Parameters: how to search the database.
- More throughput means
  - More protein sequences queried
  - Larger/more protein data bases examined
  - More parameter variation

# Why is HTC hard?

- The HTC system has to keep track of:
  - Individual tasks (a.k.a. jobs) & their inputs
  - Computers that are available
- The system has to recover from failures
  - There will be failures! Distributed computers means more chances for failures.
- You have to share computers
  - Sharing can be within an organization, or between orgs
  - So you have to worry about security
  - And you have to worry about policies on how you share
- If you use a lot of computers, you have to handle variety:
  - Different kinds of computers (arch, OS, speed, etc..)
  - Different kinds of storage (access methodology, size, speed, etc...)
  - Different networks interacting (network problems are hard to debug!)

# Let's take one step at a time

Small

Local



Large

Distributed

- Can you run one job on one computer?
- Can you run one job on another computer?
- Can you run 10 jobs on a set of computers?
- Can you run a multiple job workflow?
- How do we put this all together?

This is the path we'll take



- For 5 minutes, talk to a neighbor: If you want to run one job in a local environment:
  - 1) What do you (the user) need to provide so a single job can be run?
  - 2) What does the system need to provide so your single job can be run?
    - Think of this as a set of processes: what needs happen when the job is given? A “process” could be a computer process, or just an abstract task.



# What does the user provide?

- A “headless job”
  - Not interactive/no GUI: how could you interact with 1000 simultaneous jobs?
- A set of input files
- A set of output files
- A set of parameters (command-line arguments)
- Requirements:
  - Ex: My job requires at least 2GB of RAM
  - Ex: My job requires Linux
- Control/Policy:
  - Ex: Send me email when the job is done
  - Ex: Job 2 is more important than Job 1
  - Ex: Kill my job if it runs for more than 6 hours

# What does the system provide?

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- Methods to:
  - Submit/Cancel job
  - Check on state of job
  - Check on state of available computers
- Processes to:
  - Reliably track set of submitted jobs
  - Reliably track set of available computers
  - Decide which job runs on which computer
  - Manage a single computer
  - Start up a single job

# Quick UNIX Refresher Before We Start

- `$` #This symbolizes the prompt.
- `ssh username@training.osgconnect.net`
- `nano`, `vi`, `emacs`, `cat` `>`, etc.
- `which`, `rpm`, `ps`, `mkdir`, `cd`, `gcc`,  
`ls`
- A varitey of `condor_*` commands



# Questions?

- Questions? Comments?
  - Feel free to ask me questions now or later:  
Rob Quick [rquick@iu.edu](mailto:rquick@iu.edu)

Exercises start here:

<https://opensciencegrid.github.io/dosar/Materials/TriesteMaterials/>

Slides are also available from this URL.