

The Landscape of Academic Research Computing

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



Who Am I?

- Associate Director Science Gateway Research Center
- PI – Robust PID Project
- Co-chair CODATA School of Research Data Science Initiative
- Chair of the ACM SIGHPC RCE
- Member of the Organizational Advisory Board for RDA
- Co-chair of several CODATA and RDA Task/Working/Interest Groups
- 12 Years - Chief Operations Officer of the Open Science Grid and the Software Assurance Marketplace



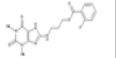


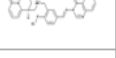



Protein Docking Project at the IU School of Medicine

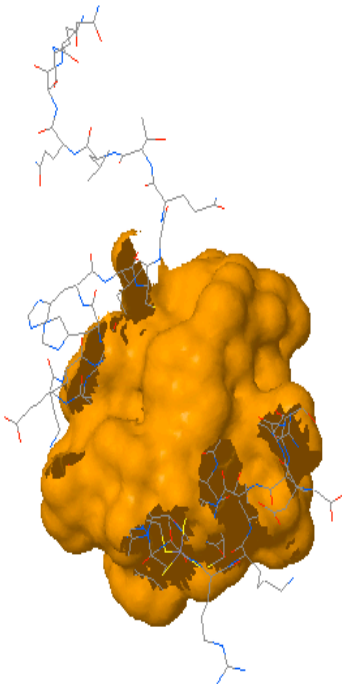
- SPLINTER - Structural Protein-Ligand Interactome
- Used autodock-vina – “...open-source program for drug discovery, molecular docking and virtual screening...”
- First run in 2013 - docked ~3900 Proteins with 5000 Ligands for a total of ~19M docked pairs.
- Submitted via command line to Condor using Pegasus on the OSG-XSEDE submission node
- Infrastructure is set and new runs can be easily started
- To date more than 6.3B dockings completed



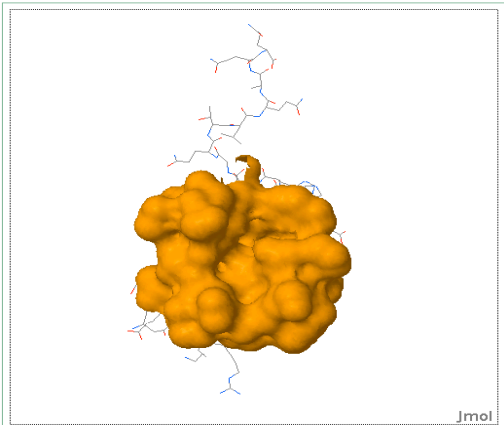


•Various rotations of Protein CBFA2T1 (Cyclin-D-related protein) (Eighty two protein) (Protein ETO) (Protein MTG8) (Zinc finger MYND domain-containing protein 2)

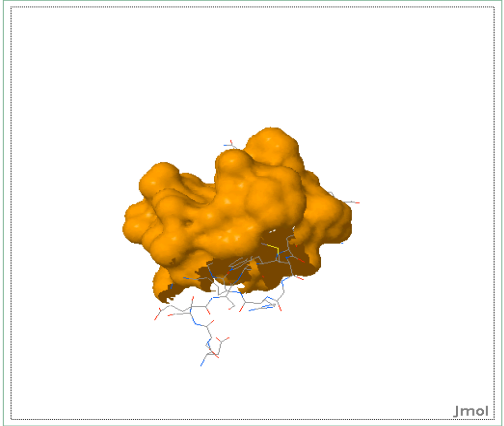
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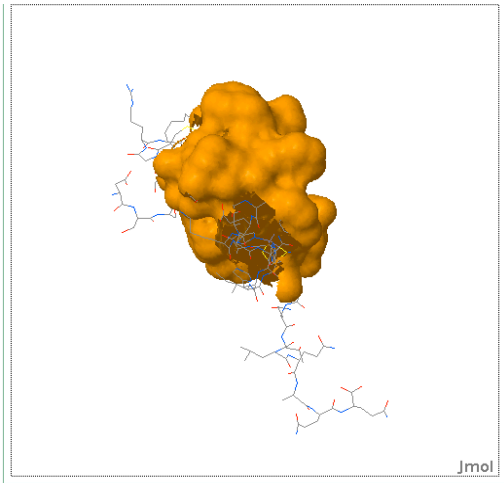
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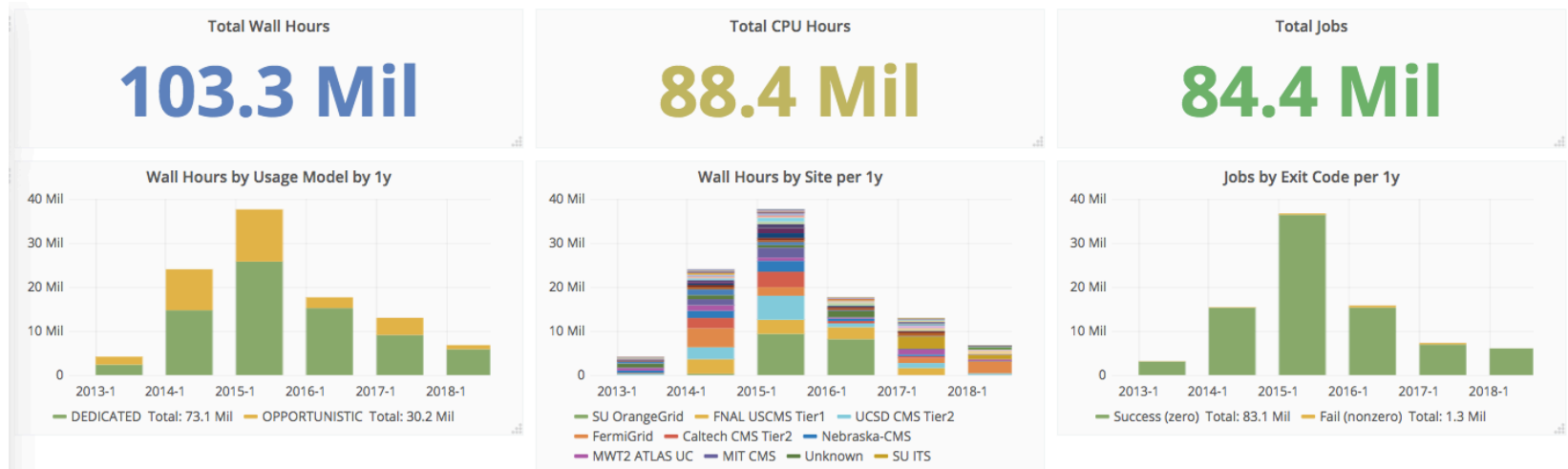


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Some Numbers



- Amazon EC2 Computing \$0.046/hour
- \$4.066M Compute Only
- Data Transfer and Storage Not Included

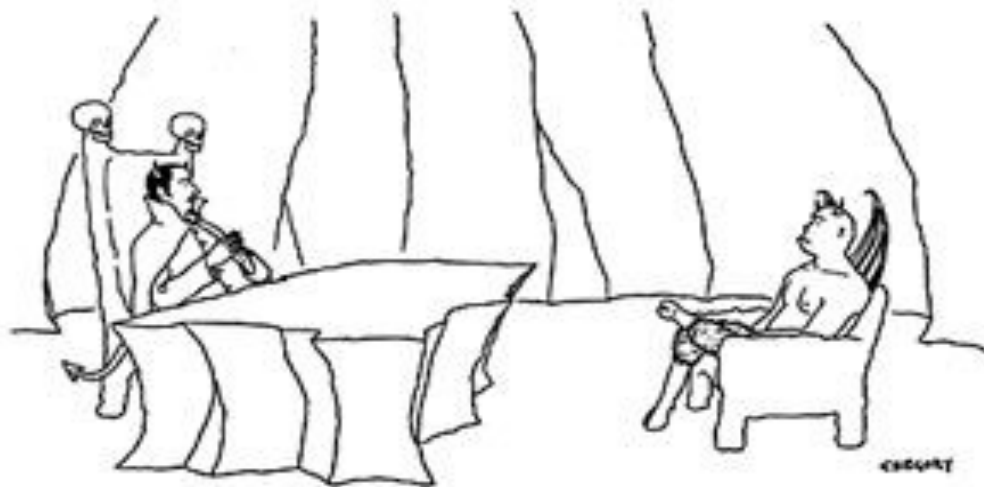
Follow Along at:

<https://opensciencegrid.org/dosar/DataTrieste2019/Materials/>



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)
- 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 learning 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

- 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)

- “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

- 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 week 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 and optimized code
- HTC – Sustained computing over a long period for serialized workflows
- Cloud – Need deeper permission on an OS and have deeper pockets



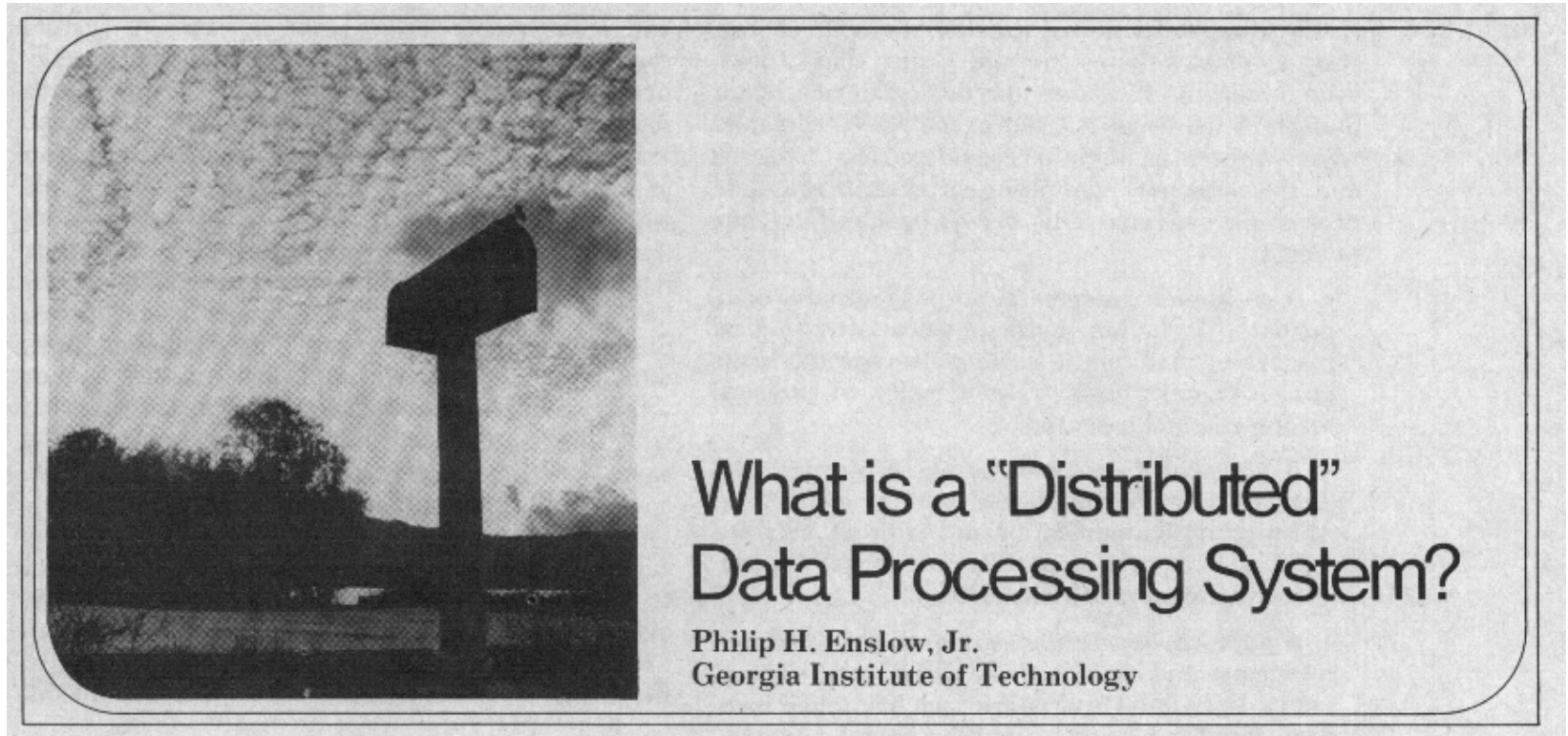
Why focus on high-throughput computing? (HTC)

- 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
 - Openness!
- 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



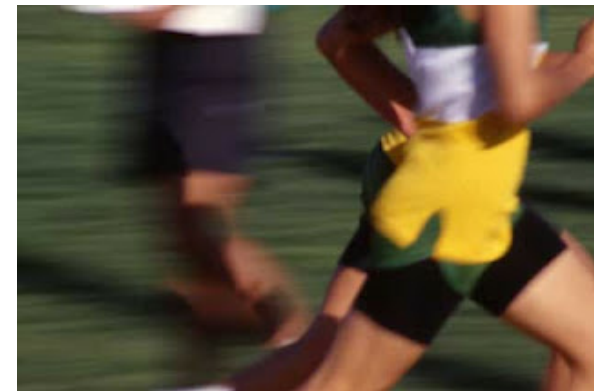


- HTC is about many jobs, many users, many servers, many sites and many workflows.
- What's in a buzzword?
 - “I was doing cloud computing since before it was called grid computing.”



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?

- 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@user-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

Exercises start here:

<https://opensciencegrid.org/dosar/DataTrieste2019/Materials/>

Slides are also available from this URL.

