

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

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Who Am I?



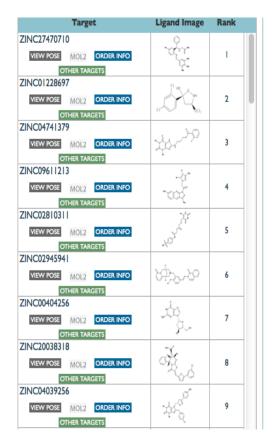
- Associate Director Cyberinfrastructure Integration Research Center
- NSF PI Enhanced Robust Persistent Identification of Data (ERPID) Project, Event Horizon Telescope Gateway
- Co-founder of CODATA/RDA School of Research Data Science Initiative
- Co-chair of the Technical and Organizational Advisory Boards for RDA
- Member of the WDS ITO Technical Advisory Committee and the EOSC Architecture Working Group
- 12+ Years Chief Operations Officer of the Open Science Grid and the Software Assurance Marketplace

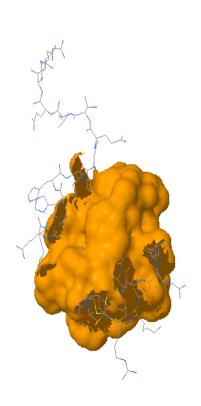
Protein Docking Project at the IU School of Medicine

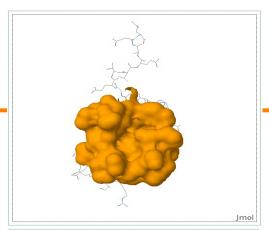
- SPLINTER <u>Structural Protein-Ligand Interactome</u>
- Used autodock-vina "...open-source program for drug discovery, molecular docking and virtual screening..."
- Docked ~3900 Proteins with 5000 Ligands for a total of ~19M docked pairs.
- Submitted via command line to Condor using Pegasus on the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

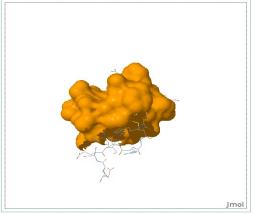


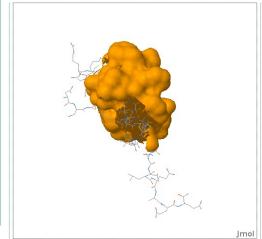
•Various rotations of Protein CBFA2T1 (Cyclin-D-related protein) (Eight twenty one protein) (Protein ETO) (Protein MTG8) (Zinc finger MYND domain-containing protein 2)







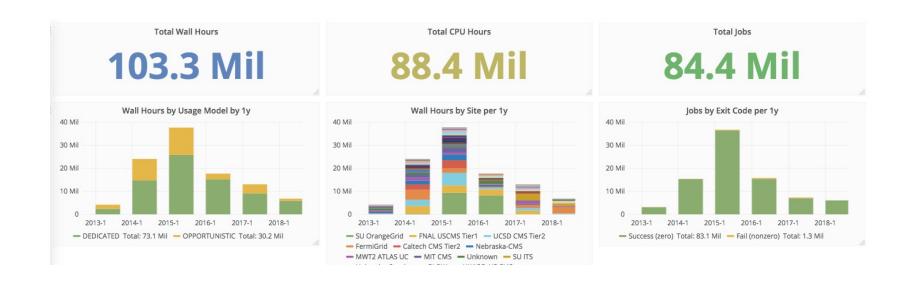




Jmol

Some Numbers

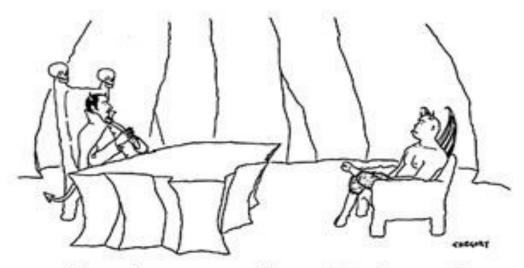




- Amazon EC2 Computing \$0.046/hour
- \$4.066M Compute Only
- Data Transfer and Storage Not Included

Overview of day





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



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



The setup: You have a problem

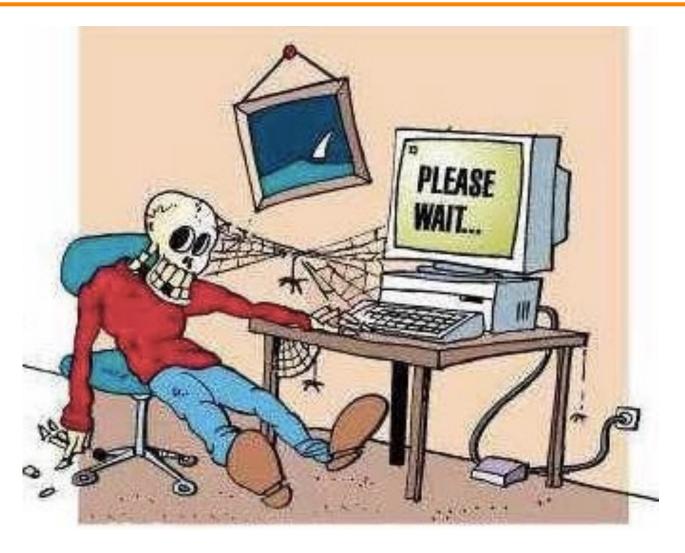


- Your science computing is complex!
 - Monte carlo, image analysis, genetic algorithm, simulation, ML or Al algorithms...
- It will take a year (CPU time) to get the results on your laptop, but your paper is due in a week.
- What do you do?



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

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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

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- 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
- The lessons learned in HTC environments and HTCondor are easily applied to other clustered or HPC systems
- We have access to an international HTC system to show the power of distributed computing

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New Technologies



Only that shall happen

Which has happened,

Only that occur

Which has occurred;

There is nothing new

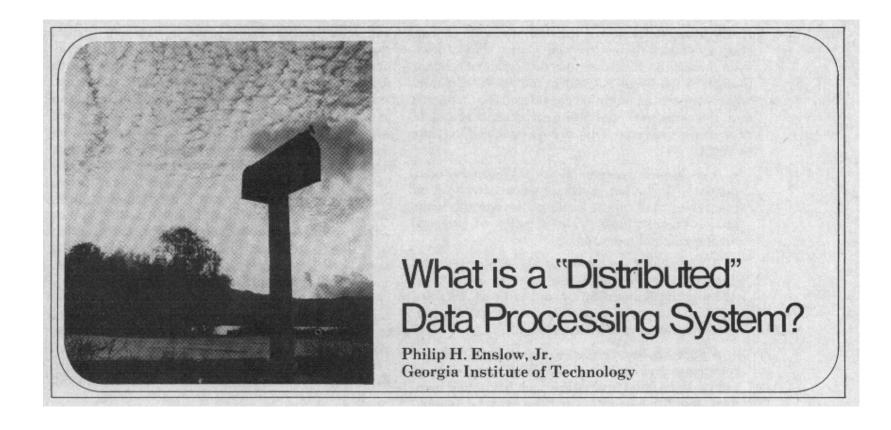
Beneath the sun!

Ecclesiastes Chapter 1 verse 9



Phillip H Enslow







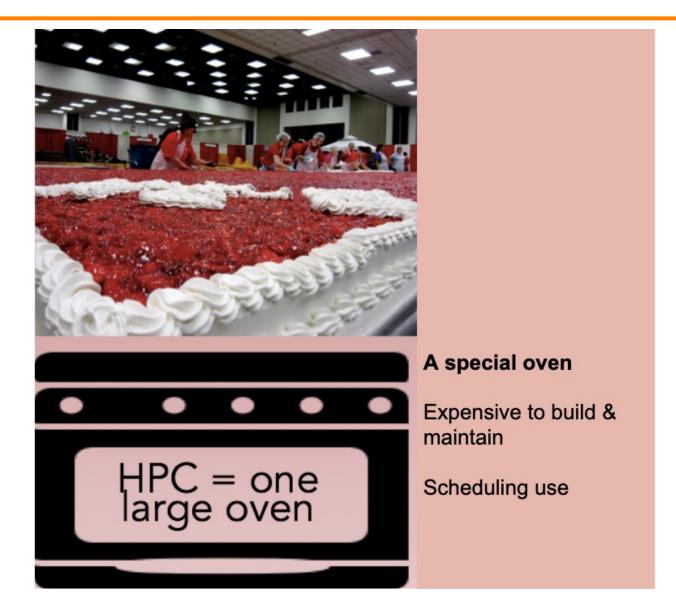


Think about a cake

- You've decided you want to bake a cake large enough to break the World's Record for wedding cakes.
- Currently WR 6.818 tonnes (15,032 lb)
- You could approach this two ways...



The HPC Approach





The HTC Approach





Why is CI hard?

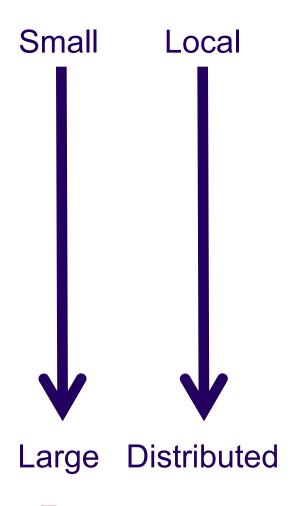


- The system has to keep track of:
 - Individual tasks (a.k.a. jobs) & their inputs
 - Computers that are available
- The system has to recover from failures
 - There will be failures! Distributed computers means more chances for failures.
- You have to share computers
 - Sharing can be within an organization, 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





- 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



Discussion



- For 5 minutes, talk to a neighbor: If you want to run a 100 job parameter sweep 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 scheme of output
- A set of parameters (command-line arguments)
- Requirements:
 - Ex: My job requires at least 2GB of RAM
 - Ex: My job requires 2 CPUs
- 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 resources

Processes to:

- Reliably track set of submitted jobs
- Reliably track set of available resources
- Decide which job runs on which resource
- Manage a single computer
- Start up a set of jobs



Questions?



- Questions? Comments?
 - Feel free to ask me questions now or later:
 Rob Quick <u>rquick@iu.edu</u>

Exercises start here:

https://github.com/CODATA-RDA-DataScienceSchools/Materials/blob/master/docs/DataSanJo se2019/CI/Materials.md

Slides are also available from this URL.

