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The Landscape of Academic Research Computing

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



Let's jump right in...

$$\begin{split} \overline{E}_{n}^{(1)} &= V_{nn} \\ \overline{E}_{n}^{(2)} &= \frac{|V_{nk_2}|^2}{E_{nk_2}} \\ E_{n}^{(3)} &= \frac{|V_{nk_2}|^2}{E_{nk_2}} \\ E_{n}^{(3)} &= \frac{|V_{nk_2}|^2}{E_{nk_2}E_{nk_3}} - V_{nn} \frac{|V_{nk_3}|^2}{E_{nk_3}^2} \\ E_{n}^{(4)} &= \frac{|V_{nk_4}|^2}{E_{nk_2}E_{nk_3}E_{nk_4}} - |V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}E_{nk_4}} - |V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_2}E_{nk_3}E_{nk_4}} + |V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}^3} \\ &= \frac{|V_{nk_4}|^2}{E_{nk_2}E_{nk_3}E_{nk_4}} - E_{n}^{(2)} \frac{|V_{nk_4}|^2}{E_{nk_4}^2} - 2V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}E_{nk_4}} + |V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}^2} \\ &= \frac{|V_{nk_4}|^2}{E_{nk_2}E_{nk_3}E_{nk_4}} - E_{n}^{(2)} \frac{|V_{nk_4}|^2}{E_{nk_4}^2} - 2V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}^2E_{nk_4}} + |V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}^2} \\ &= \frac{|V_{nk_4}|^2}{E_{nk_4}E_{nk_5}} - \frac{|V_{nk_4}|^2}{E_{nk_4}^2} - 2V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}^2E_{nk_4}} + |V_{nn} \frac{|V_{nk_4}|^2}{E_{nk_3}^2} \\ &= \frac{|V_{nk_4}|^2}{E_{nk_5}|^2E_{nk_5}} - \frac{|V_{nk_5}|^2}{E_{nk_5}^2E_{nk_5}} - \frac{|V$$



Who Am I?



- Chief Operations Officer of the Open Science Grid
- Chief Operations Officer of Software Assurance Marketpalce
- Manager High Throughput Computing Indiana University (IU)
- PI NSF Robust PID project (RPID)
- Co-Director of CODATA/RDA Schools
- Chair ACM HPC Resource Constrained Environments
- Co-Chair of 2 RDA Education Groups
- External Advisor to European Grid Infrastructure
- Member of the Organizational Advisory Board for RDA





Protein Docking Project at the USchool of Medicine

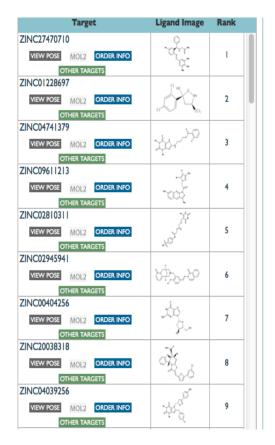


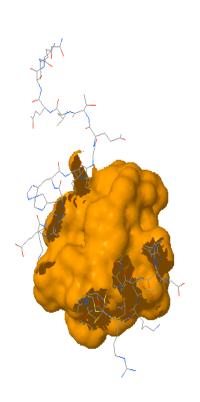
- SPLINTER <u>Structural Protein-Ligand Interactome</u>
- Used autodock-vina "...open-source program for drug discovery, molecular docking and virtual screening..."
- Frist run in 2013 docked ~3900 Proteins with 5000 Ligands for a total of ~19M docked pairs.
- Submitted via command line to Condor using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

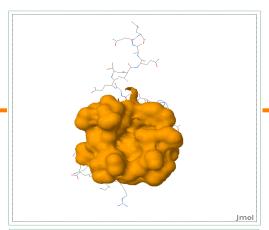


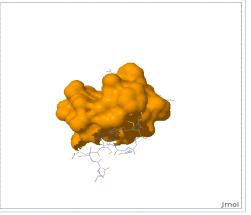


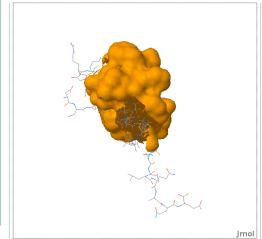
•Various rotations of Protein CBFA2T1 (Cyclin-D-related protein) (Eight twenty one protein) (Protein ETO) (Protein MTG8) (Zinc finger MYND domain-containing protein 2)









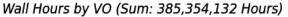


Jmol

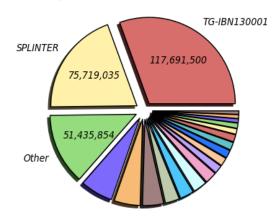


Some Numbers





1309 Days from Week 00 of 2013 to Week 31 of 2016



■ TG-IBN130001 (117,691,501)
■ DUKE-QGP (23,624,207)
■ CPDARKMATTERSIMULATION (10,292,491)
■ ALGDOCK (8,520,865)
■ SOURCECODING (5,684,844)
■ DUKE-4FERMION (3,939,893)

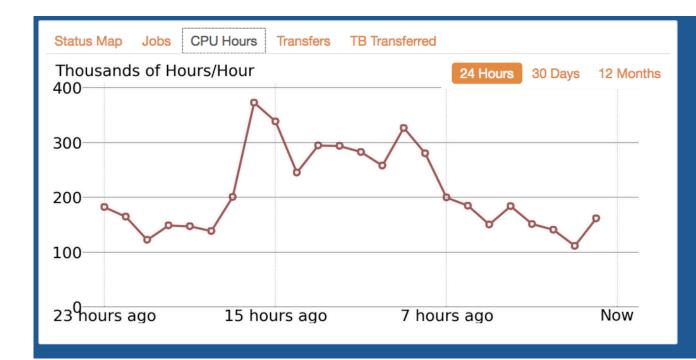
FUTURECOLLIDERS (2,684,213)

- SPLINTER (75,719,035)
 LIGO (18,859,313)
 ZZDQMC (10,049,026)
 RIT (5,966,501)
 SPHENIX (5,589,611)
 BIOGRAPH (3,652,186)
 DETECTORDESIGN (2,511,013)
- Other (51,435,854) ■ AMS (15,191,200) □ SNOWMASS (9,972,493) □ SEQ2FUN (5,961,412) ■ NUMFPI (5,222,110) ■ UPRRP-MR (2,786,365)
- Amazon EC2 Computing \$0.073/hour
- \$5.5M Compute Only
- Data Transfer and Storage Not Included



At what scale?





In the last 24 Hours	
348,000	Jobs
5,091,000	CPU Hours
9,550,000	Transfers
703	TB Transfers
In the last 30 Days	
11,331,000	Jobs
129,958,000	CPU Hours
75,556,000	Transfers
7,090	TB Transfers
In the last 12 Months	
144,466,000	Jobs
1,584,786,000	CPU Hours
2,160,318,000	Transfers
183,000	TB Transfers
OSG delivered across 127 sites	

OSG delivered across 127 sites





Follow Along at:



https://opensciencegrid.github.io/dosar/Materials/DSP Materials/

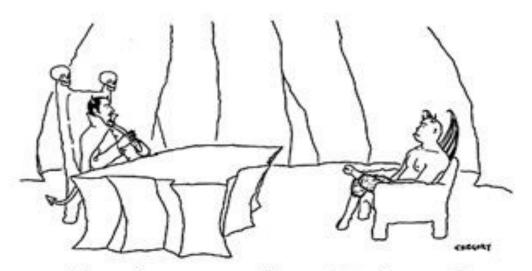




Overview of day



- 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 exercise



- 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), and Cloud Computing (XaaS)
- Shared, Allocated, and Purchased
- What is HTCondor? And why are we using it in this School?





The setup: You have a problem



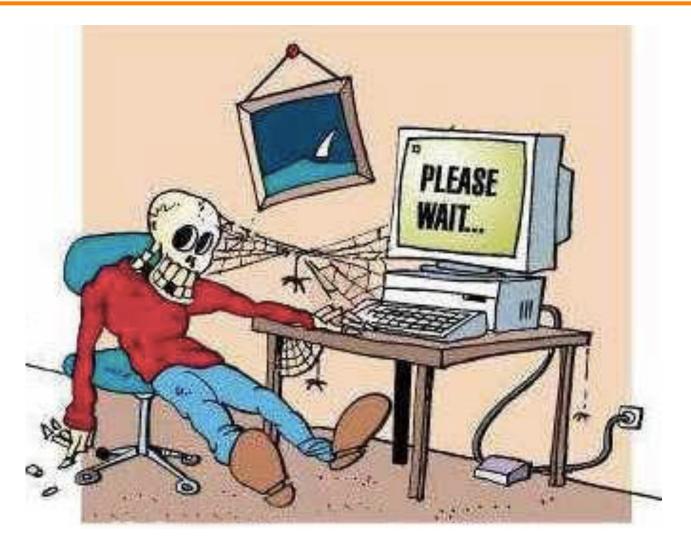
- Your science computing is complex!
 - Monte carlo, image analysis, genetic algorithm, simulation...
- It will take a year to get the results on your laptop, but the conference is in a week.
- What do you do?





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" Open Science Grid 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 processors/network/disk/memory?
- Are you willing to:
 - Port your code to a special environment?
 - Request and wait for an allocation?



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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 have 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 Open Science Grid 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 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 some efforts are underway to enable academic clouds





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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 workflows
- Cloud Need deeper permission on an
 OS and have deeper pockets



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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



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?



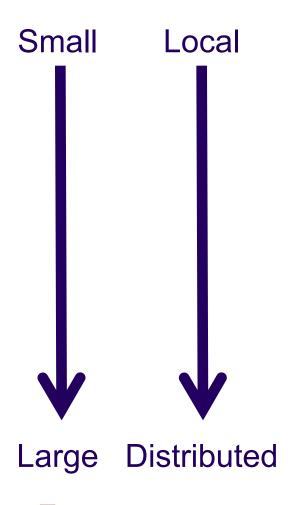
- The HTC system has to keep track of:
 - Individual tasks (a.k.a. jobs) & their inputs
 - Computers that are available
- The system has to recover from failures
 - There will be failures! Distributed computers means more chances for failures.
- You have to share computers
 - Sharing can be within an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





- 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 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 <u>UID@user-training.osgconnect.net</u>
- 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 <u>rquick@iu.edu</u>

Exercises start here:

https://opensciencegrid.github.io/dosar/Materials/

Presentations are also available from this URL.

