Once more, with feeling! A monitoring feedback loop for HTC jobs with unknown requirements

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Where we are



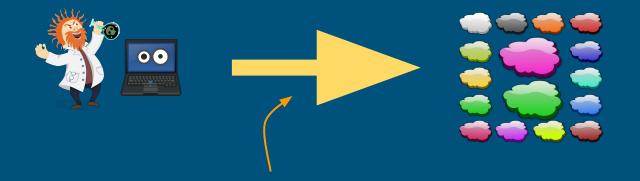


"This demo task runs on my laptop, but I need much more for the real application. It would be great if we can run O(25K) tasks like this on this cloud/grid/cluster I have heard so much about."



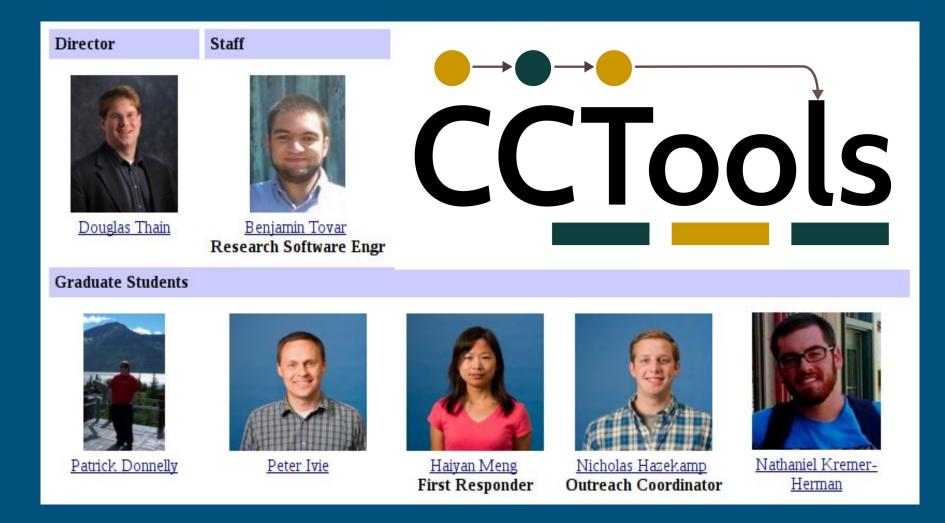


Who we are



The Cooperative Computing Lab Computer Science and Engineering University of Notre Dame

Cooperative Computing Lab



Not shown, grad students: Tim Shaffer , Chao Zheng

CCL Objectives

- Harness all the resources that are available: desktops, clusters, clouds, and grids.
- Make it easy to scale up from one desktop to national scale infrastructure.
- Provide familiar interfaces that make it easy to connect existing apps together.
- Allow portability across operating systems, storage systems, middleware...
- Make simple things easy, and complex things possible.
- No special privileges required.

CCTools



- Open source, GNU General Public License.
- Compiles in 1-2 minutes, installs in \$HOME.
- Runs on Linux, Solaris, MacOS, Cygwin, FreeBSD, ...
- Interoperates with many distributed computing systems.
 - Condor, SGE, Torque, Globus, iRODS, Hadoop...
- Components:
 - Makeflow A portable workflow manager.
 - Work Queue A lightweight distributed execution system.
 - All-Pairs / Wavefront / SAND Specialized execution engines.
 - Parrot A personal user-level virtual file system.
 - Chirp A user-level distributed filesystem.

Long-tail of science

Individual researchers and small laboratories that:

Need to curate, manage, and analyse large amounts of data.

May not know how to access computational resources available to them.

May not have immediate access to the required resources.

(i.e., they know their discipline, but they do not have an HTC expert in their team)

Notre Dame's happy opportunistic situation

- ~25k cores at Notre Dame's Center for Research Computing (CRC)
- They belong to different individual PIs, but they are available through condor when not used by their owners.

condor.cse.nd.edu



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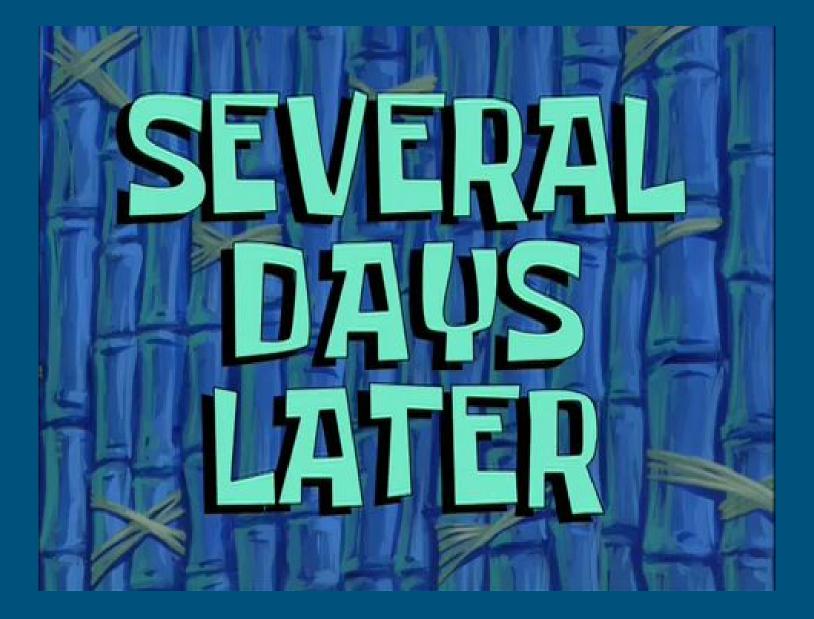
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- Mmm... It runs in my laptop...
- Surely you have a list of all the files used?

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•••

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Ok, I think we got the condor info right...
SuperSequencer3000 seems to be working on the remotes nodes now.
Yaaaay! I'll run our workflow shortly!



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No!

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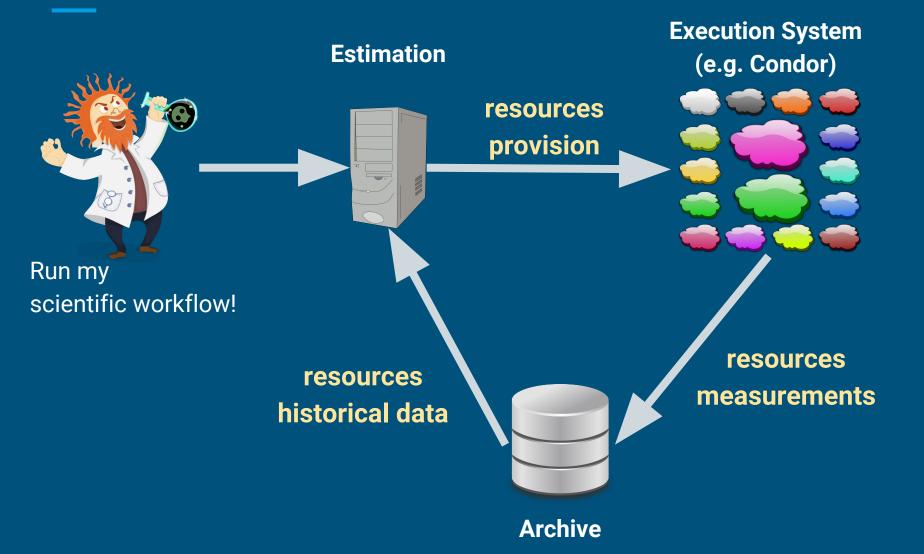
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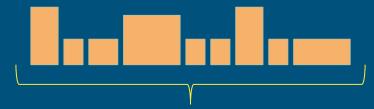


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- Wait, that parameter looks different from last time.
- Oh, that! Yes, we did change that...
- ...but we need to change that parameter often for our research...

Where we want to be



Tasks with Unknown Resource Requirements



Tasks which size (e.g., cores, memory, and disk) is not known until runtime.



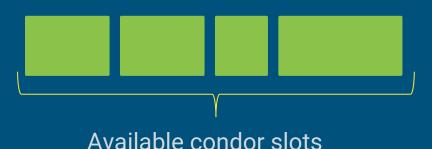
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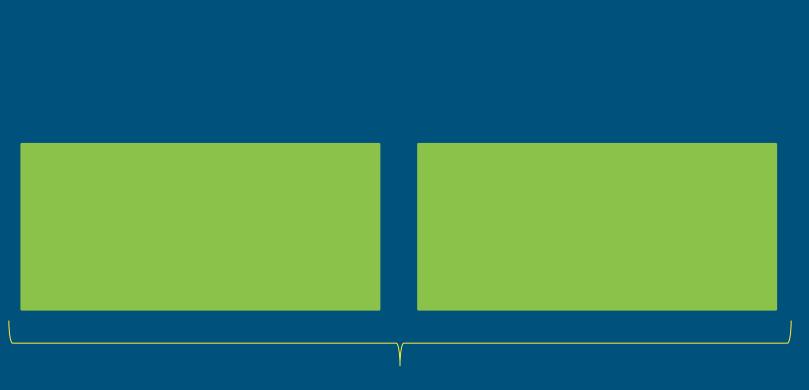
One task per slot: Wasted resources, reduced throughput.

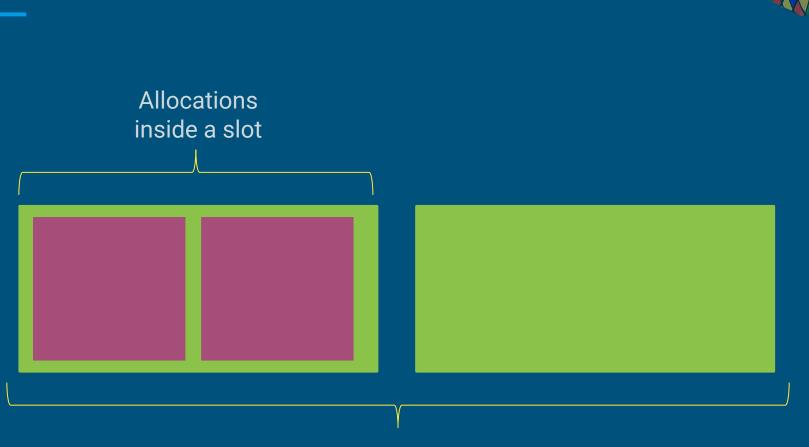
Many tasks per slot (e.g. with pilot job): Resource contention/exhaustion, reduce throughput



Task-in-the-Box





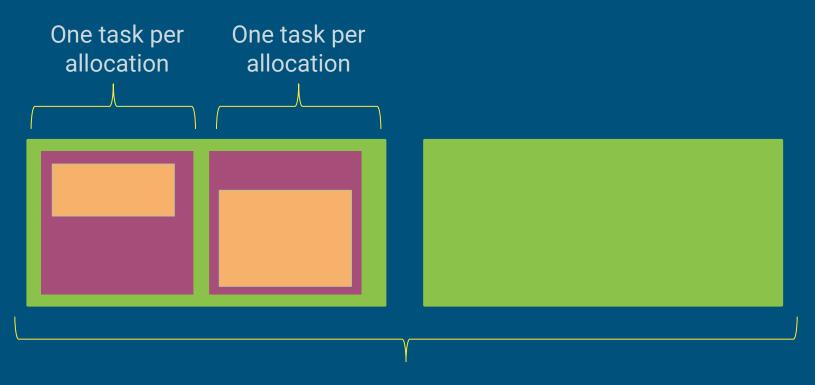


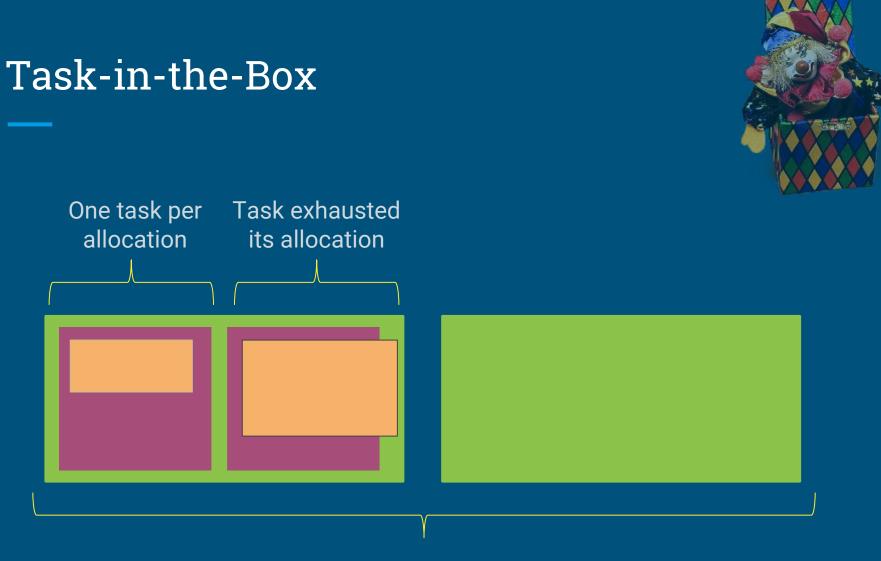
Task-in-the-Box

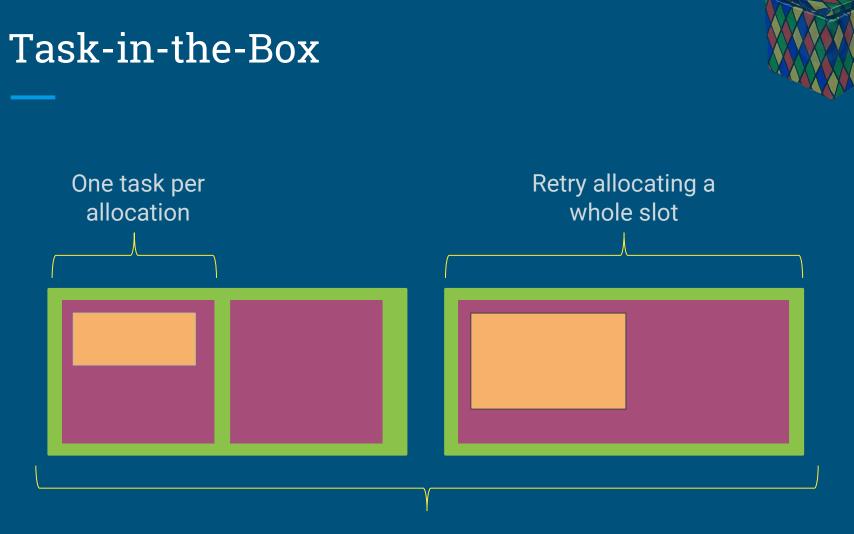




Task-in-the-Box







Main Challenges

What is a good allocation size?

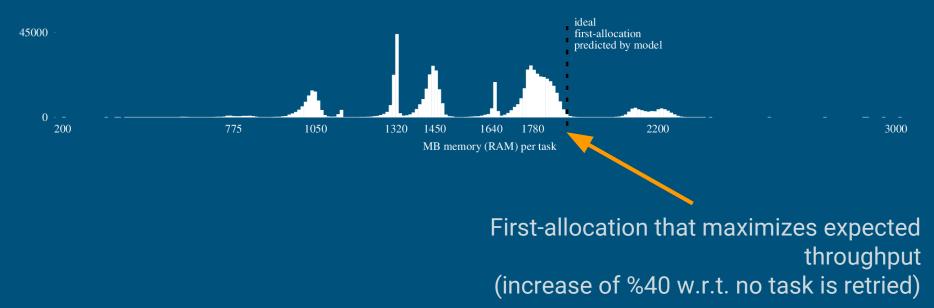
How do we measure the tasks?

How do we enforce the allocations?

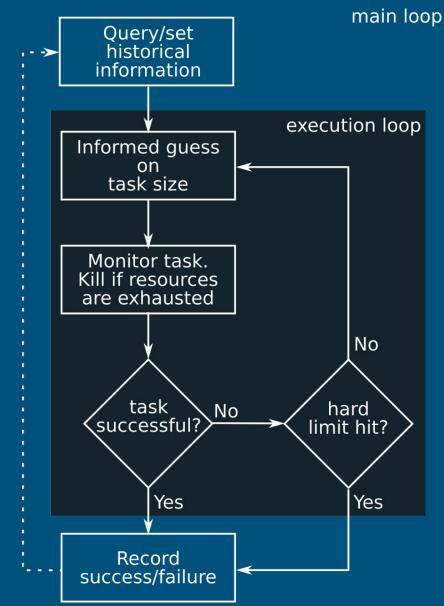
One-guess policy result (guess once, then use max seen)

Real result from a production High-Energy Physics CMS analysis (Lobster NDCMS)

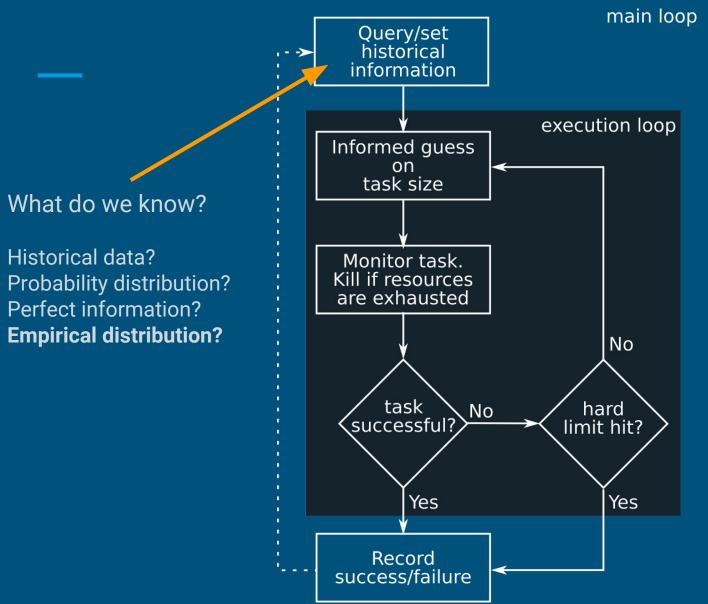
Histogram Peak Memory vs Number of Tasks O(700K) tasks that ran in O(26K) cores managed by WorkQueue/Condor.



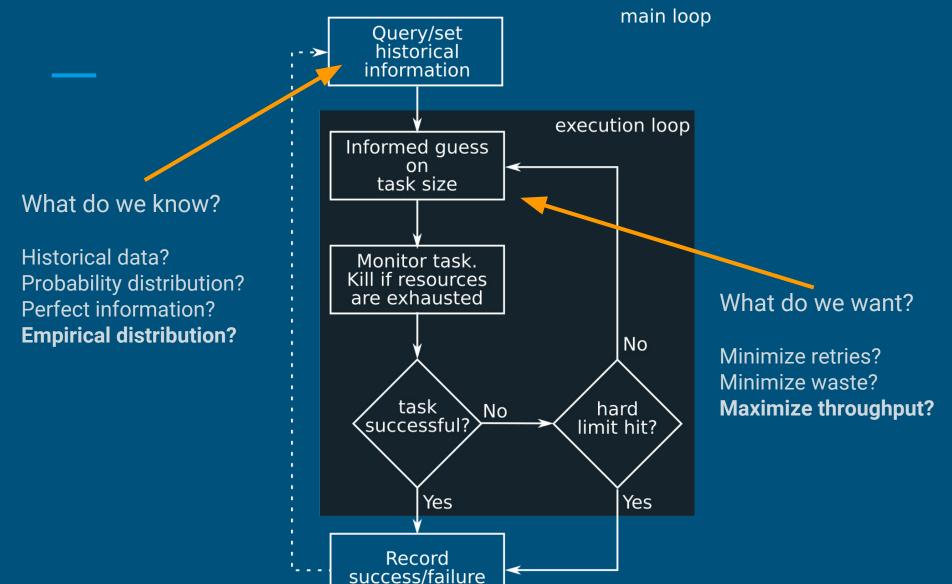
And around it goes...



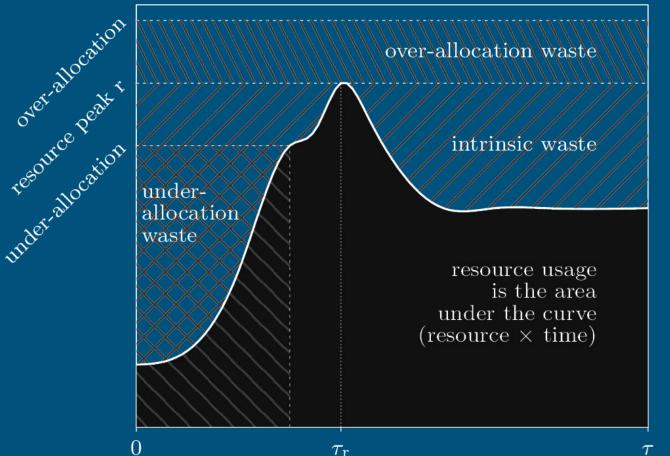
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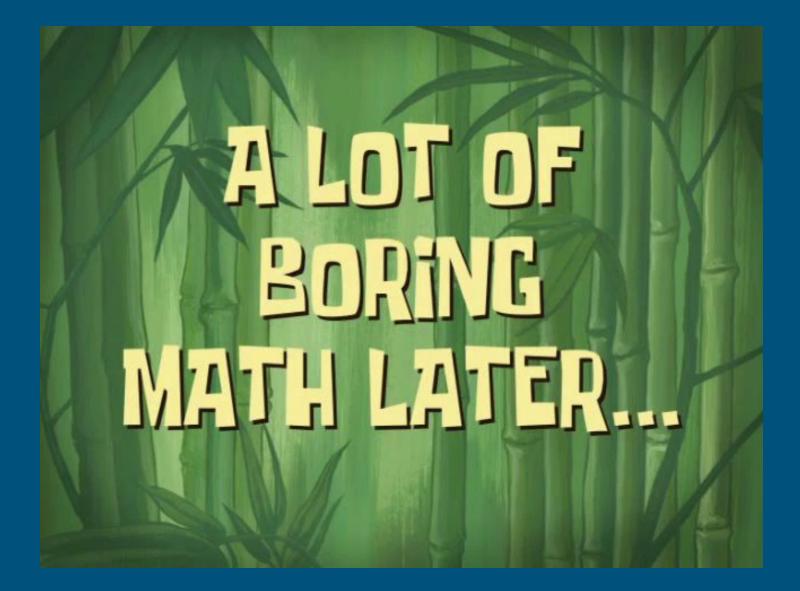
Slow-peaks model



Random variables to describe usage: Time to completion. Size of max peak

Resource usage: time x peak

Slow-peaks: Resource peaks at the end of execution (conservative assumption)



Slow-peaks model

$$E[waste(r, \tau, a_1)] = \int_0^{\infty} \left(\int_0^{a_1} (a_1 - r)\tau p(r, \tau) dr \right) d\tau$$

First allocation succeds

$$H = \int_{a_1}^{a_m} ((a_m + a_1 - r)\tau p(r, \tau) dr) d\tau$$

Final allocation succeds

$$= a_1 \int_{a_1}^{a_m} \int_0^{\infty} \tau p(r, \tau) d\tau dr$$

Optimizations over expectations

$$O(n) \text{ simple arithmetic expressions} \text{ that } \int_0^{\infty} \tau p(\tau|r) d\tau p(r) dr$$

use only information available during mean wall-time taks w. peak r
execution.

$$-\int_0^{\infty} \int_0^{\infty} r\tau p(r, \tau) d\tau dr,$$

Choice of: maximum thro minimum wast

Optimizations

execution.

Integrated in CCTools (next major release)

makeflow --max-throughput -Tcondor myworkflow

Activate monitor and allocations

Submit jobs to condor. Allocations in terms of request_cpus, request_memory and request_disk.

unix make style recipes

output.0: input.0 cmd
 ./cmd -i input.0 output.0

output.1: input.1 othercmd
 ./othercmd < input.1 > output.1

We need monitoring for all of this

Mechanisms available to unprivileged users root permissions or loading kernel modules **are a no go**

Tasks as trees of processes no whole systems or individual processes

High-throughput computing

measure so we can run many tasks at the same time, not to profile a single instance to make it run faster

We need monitoring for all of this

Monitoring as an unprivileged user is hard!

- No permissions
- No ways to add needed kernel support
- What the user wants to measure is different to what a system administrator may care about. (E.g., cpu usage of a single task v.s. system load.)
- Tracking children processes is hard without wrapping the parent process.

Need to measure individual tasks, not individual users or systems.

Integrated in CCTools

resource_monitor -L"cores: 4" -L"memory: 4096" -- matlab

cclws16 ~ > resource_monitor -i1 -Omon --no-pprint -- /bin/date Thu May 12 20:27:21 EDT 2016 cclws16 ~ > cat mon.summary {"executable_type":"dynamic", "monitor_version":"6.0.0.9edd&e96", "host":"cclws16.cse.nd.edu ","command":"/bin/date", "exit_status":0,"exit_type":"normal", "start":[1463099241605723,"us "],"end":[1463099243000239,"us"],"wall_time":[1.39452,"s"],"cpu_time":[0.002999,"s"],"core s":[1,"cores"],"max_concurrent_processes":[1,"procs"],"total_processes":[1,"procs"],"memor y":[1,"MB"],"virtual_memory":[107,"MB"],"swap_memory":[0,"MB"],"bytes_read":[0.0105429,"MB "],"bytes_written":[0,"MB"],"bytes_received":[0,"MB"],"bytes_sent":[0,"MB"],"bandwidth":[0, "Mbps"],"total_files":[90546,"files"],"disk":[11659,"MB"],"peak_times":{"units":"s","cpu_ time":1.39452,"cores":0.394445,"max_concurrent_processes":0.394445,"memory":0.394445,"virt ual_memory":1.39428,"bytes_read":1.39428,"total_files":1.39428,"disk":1.39428}}

cclws16 ~ >

(does not work as well on static executables that fork)

Recent development: Monitoring Library

The resource_monitor main functionality was converted into a library, with C, Python, and perl interfaces.

```
struct rmsummary resources;
rmonitor_measure_process(&resources, getpid());
fprintf(stdout, "command: %s, ",
        resources.command);
fprintf(stdout, "wall time used (s): %3.01f, ",
        resources.wall_time/1000000.0);
fprintf(stdout, "total memory used (MB): %" PRId64 ", ",
        resources.resident_memory + resources.swap_memory);
fprintf(stdout, "total cores used: %" PRId64 "\n",
        resources.resident_memory + resources.swap_memory);
```

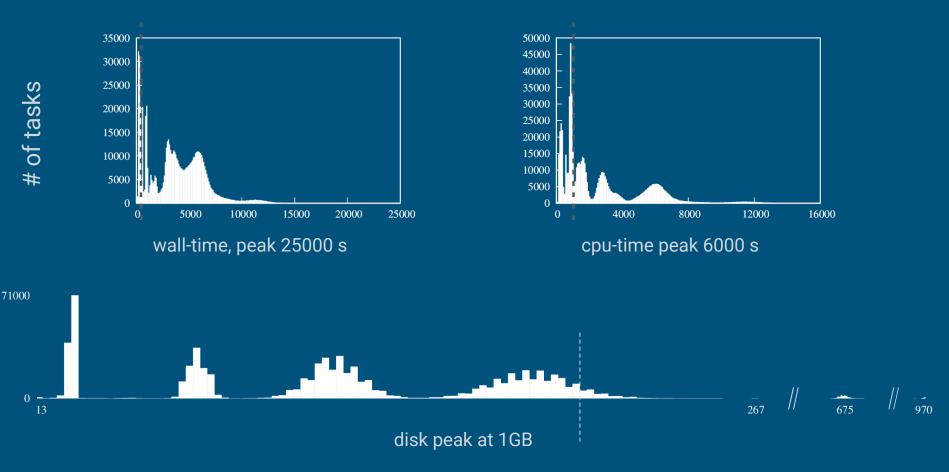
An application can poll its resources usage with a single library call.

(unlike resource_monitor, does not track forks/exits)

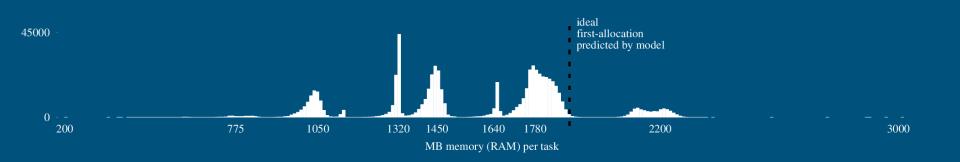
ND CMS workflow distributions

681874 tasks on Lobster/WorkQueue/Condor

• Computing allocations takes ~ 0.05 seconds.



ND CMS workflow: Memory bottleneck



	Size	Retries	Proportion wasted	Throughput (norm)
Max peak always	3GB	0%	48%	1.0
Perfect information	-	0%	0%	2.0
Slow-peaks one- guess	1.9GB	9%	28%	1.41

ND CMS workflow: Memory bottleneck

Things are even better if users give coarse information about the workflow. As simple as putting tasks into categories (e.g., merge, analysis recostep, parameter-X, etc.)

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Slow-peaks one- guess	1.9GB	9%	28%	1.41
One-guess + categories	(per category)	< 1%	17%	1.64

Questions?

Acknowledgements:

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Downloads:

cctools http://ccl.cse.nd.edu

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