WORKFLOW AUTOMATION TOOLS FOR MANY-TASK COMPUTING

Geoffrey Lentner, Lead Research Data Scientist
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This workshop outlines the paradigm of many-task computing and covers different scenarios and tools to manage such workflows.

An introduction to hyper-shell is included at the end.

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About Us

The Rosen Center for Advanced Computing (RCAC) provides advanced computational resources and services to support Purdue faculty and staff researchers. We conduct our own research and development to enhance the capabilities of these resources, as well as provide expertise in a broad range of HPC matters and activities.

This includes providing workshops on common topics surrounding the use of HPC systems.
Members of the *Scientific Applications* team have a number of responsibilities; including,

- System configuration and services
- Scientific software
- Advanced projects and consulting
- Teaching
- Outreach and Engagement
- Innovation
Prerequisites. We assume you have some prior experience working on a Linux cluster, using the scheduler and simple command-line programs.

- Linux command-line basics (Unix 101 and Unix 102)
- Cluster basics (Clusters 101)
Preface

Outline

• What do we mean by many-task computing?

• Why not just use the scheduler?

• What tools are available to manage tasks?

• What is hyper-shell and how to use it.
Introduction

What are we even talking about?
What is many-task computing?
What?

High-Throughput, Many-Task Computing
Large number of Independent Tasks without Communication

Defined by the number of tasks completed or volume of data processed. Elements of the workflow are entirely independent and may run anywhere, even across administrative boundaries.

High-Performance Computing
Coherent Coupled Tasks with Communication

Monolithic simulation where tasks run simultaneously and define some physical or abstract space, working together to solve one problem.
Introduction

Use cases

• Data Processing and Analysis
• Machine-Learning Experiments (model tuning)
• Bioinformatics Tasks
• Parameter Searchers (calculations)
Example 1

Process hourly meteorological satellite data for surface parameters for the whole planetary surface since 1982 ...

Compute new parameter with the data at each hourly interval.

That’s $350k+$ tasks …
**Example 2**

*Train* a *stochastic* model against decades of forest survey data to predict *tree species* population change …

For many species, for many regions, *hundreds of times* …

That’s **200k+** tasks …
Example 3

*Calculate* some mathematical coefficient by searching a *higher*-dimensional parameter space …

That’s 1M+ tasks …
Why not manage tasks directly?

People tend to just iterate through tasks inside the same application language where the computation is defined; e.g., Python, Julia, R, MATLAB, Mathematica, …

… because it’s simple, obvious, and works.

… until it doesn’t.
Where does this break down?
I got an allocation on the computing cluster and my **MATLAB** workflow runs okay …

… but now I have **six** nodes and struggling to make distributed computing work.
Building **HPC-friendly** workflows means **writing less code** …
A direct approach within the same script …

- is monolithic (what if there’s an error),
- is inflexible (couples orchestration with problem code),
- a challenge for novice programmers, and
- tedious.
Use the scheduler

Why not just use SLURM?
What are the benefits and limitations?
Using the scheduler

In the ideal case

SLURM (or equivalent) is actually the best approach. It is literally the scheduler’s responsibility to manage tons of independent jobs.

```
#!/bin/bash
#SBATCH -N1 -n1 -t 00:30:00
#SBATCH ...
# load software
module load matlab/R2021A
# execute single instance
matlab -nosplash -singleCompThread -batch "..."
```

Control the varying inputs somehow, such as with an environment variable, or substitution …
Using the scheduler

Job Arrays

Most schedulers support “job arrays” in which the same job is executed some large number of times with only environment variables being different.

```bash
#!/bin/bash
#SBATCH -N1 -n1 -t 00:30:00
#SBATCH -a 1-999
#SBATCH -o task-%A.%a.out
#SBATCH -e task-%A.%a.err

# load software
module load matlab/R2021A

# execute single instance
matlab -nosplash -singleCompThread -batch "...($SLURM_ARRAY_TASK_ID)"
```

Just the variable we were looking for…
So why would you use some other "workflow automation" tool?
Using the scheduler

Practical limits

Site administrators don’t want users …

• submitting millions of jobs,

• filling up the database,

• impacting site-wide throughput, and

• polluting the queue.
Submit a "pilot job" instead ...
3

**Workflow Automation Tools**

What other tools are available?
What are the benefits and limitations?
Create a task file

We could **source** this file and it would be valid, running the commands **one at a time** ...

```
matlab -nosplash -singleCompThread ... -batch "...(3, ...)"
matlab -nosplash -singleCompThread ... -batch "...(2, ...)"
matlab -nosplash -singleCompThread ... -batch "...(3, ...)"
matlab -nosplash -singleCompThread ... -batch "...(4, ...)"
matlab -nosplash -singleCompThread ... -batch "...(5, ...)"
matlab -nosplash -singleCompThread ... -batch "...(6, ...)"
matlab -nosplash -singleCompThread ... -batch "...(7, ...)"
matlab -nosplash -singleCompThread ... -batch "...(8, ...)"
matlab -nosplash -singleCompThread ... -batch "...(9, ...)"
matlab -nosplash -singleCompThread ... -batch "...(10, ...)"
matlab -nosplash -singleCompThread ... -batch "...(11, ...)"
matlab -nosplash -singleCompThread ... -batch "...(12, ...)"
matlab -nosplash -singleCompThread ... -batch "...(13, ...)"
matlab -nosplash -singleCompThread ... -batch "...(14, ...)"
matlab -nosplash -singleCompThread ... -batch "...(15, ...)"
matlab -nosplash -singleCompThread ... -batch "...(16, ...)"
matlab -nosplash -singleCompThread ... -batch "...(17, ...)"
matlab -nosplash -singleCompThread ... -batch "...(18, ...)"
matlab -nosplash -singleCompThread ... -batch "...(19, ...)"
matlab -nosplash -singleCompThread ... -batch "...(20, ...)"
matlab -nosplash -singleCompThread ... -batch "...(21, ...)"
matlab -nosplash -singleCompThread ... -batch "...(22, ...)"
matlab -nosplash -singleCompThread ... -batch "...(23, ...)"
matlab -nosplash -singleCompThread ... -batch "...(24, ...)"
matlab -nosplash -singleCompThread ... -batch "...(25, ...)"
```
Workflow Automation Tools

Pass the file to a workflow automation tool

All of the utilities discussed here have a similar usage pattern

```
pilot-job.sh

# load software
module load ???

# scale out tasks
<???> task.in ...
```

What can we put here?
# Workflow Automation Tools

## xargs

*Build and execute command lines from standard input*

<table>
<thead>
<tr>
<th>Pros</th>
<th>Cons</th>
</tr>
</thead>
<tbody>
<tr>
<td>Available by default</td>
<td>Low observability</td>
</tr>
<tr>
<td>Simple</td>
<td>Not Scalable (single node)</td>
</tr>
<tr>
<td>Templates</td>
<td>UNIX/Linux only</td>
</tr>
</tbody>
</table>

### Example

```
$ xargs -a task.in -d "\n" -P 128 -I {} bash -c "{}"
```
## Workflow Automation Tools

### srun

_Slurm launcher and job step manager_

<table>
<thead>
<tr>
<th>Pros</th>
<th>Cons</th>
</tr>
</thead>
<tbody>
<tr>
<td>Available</td>
<td>Low observability</td>
</tr>
<tr>
<td>Distributed</td>
<td>Linux/SLURM only</td>
</tr>
<tr>
<td></td>
<td>Not Scalable (site limits)</td>
</tr>
<tr>
<td></td>
<td>No retries</td>
</tr>
<tr>
<td></td>
<td>Monolithic</td>
</tr>
</tbody>
</table>

**Example**

```
$ cat task.in | xargs -d "\n" -P512 -I {} srun -n1 ... {}
```

Depending on how big you want to go this could be **problematic** for the same reasons submitting many jobs is.

[https://slurm.schedmd.com/srun.html](https://slurm.schedmd.com/srun.html)
## Workflow Automation Tools

### ParaFly

*Execute tasks in parallel and report errors*

<table>
<thead>
<tr>
<th>Pros</th>
<th>Cons</th>
<th>Example</th>
</tr>
</thead>
<tbody>
<tr>
<td>Simple</td>
<td>Low observability</td>
<td>$ ParaFly -c task.in -CPU 128 -failed_cmds task.failed</td>
</tr>
<tr>
<td>Retries / Failures</td>
<td>Not Scalable (single node)</td>
<td></td>
</tr>
<tr>
<td></td>
<td>UNIX/Linux only</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Monolithic</td>
<td></td>
</tr>
</tbody>
</table>

No Templates

https://parafl.y.sourceforge.net/
https://github.com/ParaFly/ParaFly
**Workflow Automation Tools**

**GNU Parallel**

*Execute tasks in parallel using one or more computers*

<table>
<thead>
<tr>
<th>Pros</th>
<th>Cons</th>
</tr>
</thead>
<tbody>
<tr>
<td>Usability</td>
<td>Low observability</td>
</tr>
<tr>
<td>Distributed</td>
<td>I/O Limitations</td>
</tr>
<tr>
<td>*Retries / Failures</td>
<td>Not scalable (~10-20 nodes)</td>
</tr>
<tr>
<td>Templates</td>
<td>UNIX/Linux only</td>
</tr>
<tr>
<td></td>
<td>Monolithic</td>
</tr>
</tbody>
</table>

**Example**

```
$ cat task.in | parallel -j128 ...
```

https://www.gnu.org/
# Workflow Automation Tools

## Launcher

*Scale out tasks on HPC systems*

<table>
<thead>
<tr>
<th>Pros</th>
<th>Cons</th>
</tr>
</thead>
<tbody>
<tr>
<td>Simple</td>
<td>Low observability</td>
</tr>
<tr>
<td>Distributed</td>
<td>Linux only (HPC specific)</td>
</tr>
<tr>
<td>Scalable</td>
<td>Monolithic</td>
</tr>
</tbody>
</table>

### Example

```bash
$ export LAUNCHER_WORKDIR=.  
$ export LAUNCHER_JOB_FILE=task.in  
$ ${LAUNCHER_DIR}/paramrun
```

https://www.tacc.utexas.edu/research-development/tacc-software/the-launcher

https://portal.tacc.utexas.edu/software/launcher

https://github.com/TACC/launcher
**Workflow Automation Tools**

### hyper-shell

*Process shell commands over a distributed, asynchronous queue*

<table>
<thead>
<tr>
<th>Pros</th>
<th>Cons</th>
</tr>
</thead>
<tbody>
<tr>
<td>Retries</td>
<td>Usability</td>
</tr>
<tr>
<td>Flexibility</td>
<td>Scalability</td>
</tr>
<tr>
<td>Cross-platform</td>
<td>Observability</td>
</tr>
<tr>
<td>Persistent / management</td>
<td>Templates</td>
</tr>
</tbody>
</table>

**Example**

```
$ hyper-shell cluster task.in -N128 --launcher=srun ...
```

https://github.com/glentner/hyper-shell

https://hyper-shell.readthedocs.io/
## Workflow Automation Tools

### Feature Comparison

<table>
<thead>
<tr>
<th></th>
<th>Distributed</th>
<th>Restart / Failures</th>
<th>Templates</th>
<th>Scalable</th>
<th>Observable</th>
<th>Cross Platform</th>
<th>Persistent</th>
</tr>
</thead>
<tbody>
<tr>
<td>hyper-shell</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
</tr>
<tr>
<td>xargs</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>srun</td>
<td>✓</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td>✓</td>
<td></td>
</tr>
<tr>
<td>ParaFly</td>
<td>✓</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GNU Make</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>GNU Parallel</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>*Launcher</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td><strong>TaskFarmer</strong></td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*https://www.tacc.utexas.edu/research-development/tacc-software/the-launcher
**https://docs.nersc.gov/jobs/workflow/taskfarmer/
Workflow Automation Tools

Use the tool that works for you …

… consider trying a new one if you need additional features.
hyper-shell

What is hyper-shell?
What problems is it trying to solve?
What are some of it’s features?
Common Usage

Example job script using *hyper-shell* on *Negishi*

```
#!/bin/bash
#SBATCH -N4 -t 1-00:00:00 --exclusive
#SBATCH ...

# load software
module load hyper-shell

# launch workflow
hyper-shell cluster task.in -N128 --launcher=srun \n   --capture --failures=task.failed --no-db --no-confirm
```
Flexible (Not Monolithic)

Bring up the server first, then the clients later as needed

$ hyper-shell server --restart --eager -r 2 \
   --bind 0.0.0.0 --auth ABC123

$ hyper-shell client -N128 --capture \ 
   --host a123.negishi --auth ABC123

$ hyper-shell client -N128 --capture \ 
   --host a123.negishi --auth ABC123

$ hyper-shell client -N128 --capture \ 
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   --host a123.negishi --auth ABC123

$ hyper-shell client -N128 --capture \ 
   --host a123.negishi --auth ABC123
Cross-Platform (Hybrid)

Clients do not need to run within the same platform environment

```
$ hyper-shell server --restart --eager -r 2 \
  --bind 0.0.0.0 --auth ABC123

C:\ hyper-shell client -N128 --capture \ 
  --host a123.negishi --auth ABC123
```

```
C:\ hyper-shell client -N128 --capture \ 
  --host a123.negishi --auth ABC123

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C:\ hyper-shell client -N128 --capture \ 
  --host a123.negishi --auth ABC123
```
Rich Templates

Dynamically build commands instead of reading from a static file

Filepath Operations
Shorthand notation provides convenient expressions, such as, `{/-}`, which expands to the base name without the file type extension.

Argument Slicing
Select items from the incoming argument with familiar notation, such as `{[0]}` for the first item, or `{[:2]}` for the last two.

Shell Expansion
Interpolate the output of other programs into the command line, such as `{% mktemp -d @ %}`.

Lambda Expressions
Evaluate expressions (with Python), such as `{= x * math.pi =}`.
Persistent Metadata
A database is used to persistent and query task metadata

```
$ hyper-shell task search --where 'exit_status != 0' 'attempt == 2' --json
[
  {
    "id": "f05ceea8-3232-4b73-88e3-c26966865d68",
    "args": "matlab -nosplash -singleCompThread -batch ...",
    ...
  },
  {
    "id": "351f748c-d1d8-44a7-a1ff-37b4fa6effd8",
    "args": "matlab -nosplash -singleCompThread -batch ...",
    ...
  }
]
```
Architecture

Two-tiered client-server architecture with batched tasks.
Concurrent threads of autonomous agents.
Heartbeats.
Alternatives

What other types of tools are out there?
Alternatives

Makefile

Alternative solutions using dependency-graph specific syntax

- **GNU Make** *(available, robust, dynamic, but not scalable)*,

- **Snakemake** *(couples execution with environment and orchestration)*,

- **Makeflow** *(couples execution with environment and orchestration)*
Alternatives

*Makefiles* use the filesystem for dependency resolution and task tracking…

…which is awesome and in the spirit of the UNIX philosophy, until it breaks down at scale.
Exaworks

Next generation technologies for composable and scalable HPC workflows

[Diagram showing Exaworks PSI/J and its components: Workflow System, Workflow System, Workflow System, Workflow System. The diagram also shows Exaworks Scientific Workflows SDK with components such as Exascale Systems, Components, SDK, Develop, Test, Flux, Parallel, Package, Build, Components, Platform.]

https://exaworks.org/
https://exaworkssdk.readthedocs.io/
THANK YOU

Please reach out to rcac-help@purdue.edu for questions.

See rcac.purdue.edu/training/workflow_automation for slides.
See rcac.purdue.edu/training for additional topics.
See rcac.purdue.edu/knowledge for user-guides.