Hewlett Packard Enterprise

# HPC WORKFLOW MANAGEMENT WITH CHAPEL

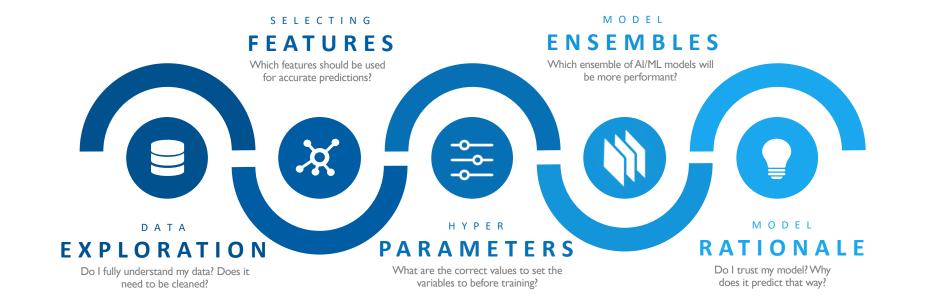
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# MOTIVATION

- Coordinating many monolithic applications is a challenge for the HPC user community
  - This impacts scientists across many domains such as astronomy, computational chemistry, and bioinformatics
- These workflows often begin as a simple shell script or collection of scripts
  - Depending on the complexity of the workflow, this may become unwieldy
  - Users can adopt a workflow framework or code their workflow in a more advanced programming language
- This talk will explore Chapel as a language to write HPC workflows

#### **BACKGROUND: CRAY HPO**

- The patterns explored in this talk are motivated by those found in Cray HPO
  - They are not a comprehensive list of workflow patterns
- Cray HPO is a hyperparameter optimization framework developed at HPE
- Background on hyperparameter optimization (HPO):
  - In data science, a model is trained on a dataset
  - Parameters are internal values in the model that are used to make predictions, e.g. slope and offset in y=mx+b
  - Hyperparameters are external values that impact how the model is trained, e.g. learning rate in SGD
  - Hyperparameter optimization is a process of tuning hyperparameters to minimize a metric, e.g. 1-accuracy
  - There is a wide breadth of HPO strategies: grid, random, genetic, bayesian, and more advanced variations



• Cray HPO essentially acts as a distributed workflow manager for tuning the hyperparameters **from** crayai **import** hpo **#** crayai.hpo implementation is written in Chapel

```
# Specify training kernel and the resources on which to execute HPO training
evaluator = hpo.Evaluator('python3 source/train.py', workload manager=`slurm', nodes=16)
```

# Specify optimizer and its metaparameters

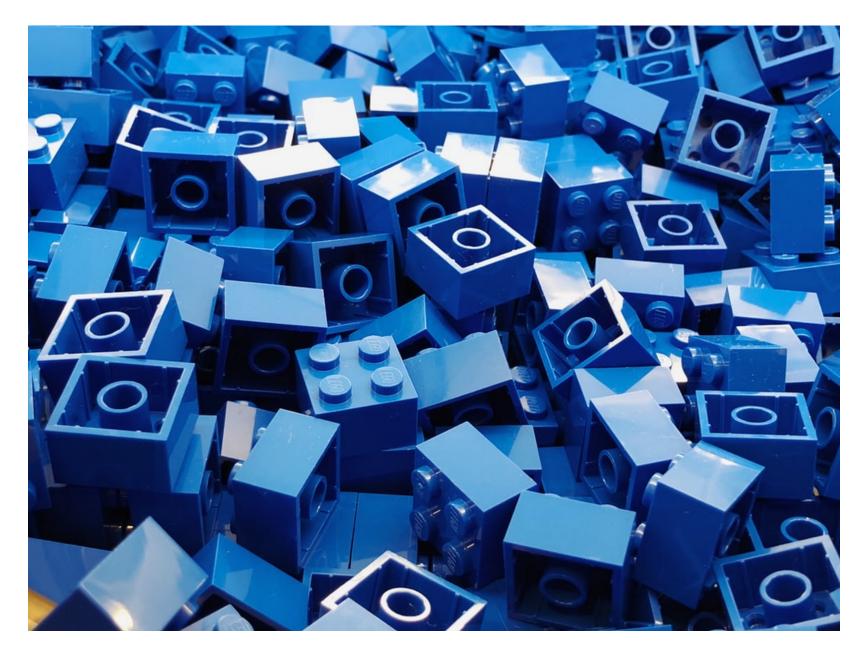
```
optimizer = hpo.GeneticOptimizer(evaluator,
```

```
generations=10,
pop_size=16,
mutation_rate=0.15)
```

# Optimize hyperparameters, results are stored in a csv file
optimizer.optimize (params)



#### WORKFLOW BUILDING BLOCKS



#### LAUNCHING A SUBPROCESS

- Running other programs and interacting with their outputs is the basic building block of workflows
- In Chapel, running other programs can be done with the Spawn module:

```
use Spawn;
var process = spawn(['./run-simulation']);
```

A Chapel Documentation	* » Standard Modules » Spawn	View page source
rch docs		
	Spawn 🗞	
PILING AND RUNNING CHAPEL		
kstart Instructions	Usage	
g Chapel		
orm-Specific Notes	use Spawn;	
nical Notes		
	or	
ING CHAPEL PROGRAMS	<pre>import Spawn;</pre>	
Reference		
World Variants	Support launching and interacting with other programs.	
rs		
age Specification	Using functions in this module, one can create a subprocess and possibly capture its output. It is also possible to provide input to a subprocess.	
n Types and Functions	and possible to provide input to a subprocess.	
ard Modules	To start a subprocess, use spawn or spawnshell. To wait for the subprocess process to finish,	
omatic Modules	use the subprocess.wait or subprocess.communicate functions.	
Structures		
nostics	This example program produces a listing of files in the current di with test. by using the ts command. The output will be mixe	
	output.	

### LAUNCHING A SUBPROCESS

- Spawning a subprocess is a non-blocking operation.
  - Blocking on the completion of the program can be done with a 'wait' or 'communicate' call:

#### LAUNCHING A DISTRIBUTED SUBPROCESS

For interacting with HPC workload managers and launchers, one can capture interface into a function:

```
use Slurm;
const jobid = salloc(numNodes=32, timeout=120);
var process = srun('./run-simulation', numNodes=32, id=jobid);
process.wait();
```

Abstracting away workload manager specifics enables portability across HPC systems:

```
use Jobs;
var job = new Job(workloadManager=Launchers.pbs, launcher=Launchers.aprun);
job.alloc(numNodes=32, timeout=120);
var process = job.run('./run-simulation', numNodes=32);
process.wait();
```

# LAUNCHING SUBPROCESSES IN PARALLEL

- Launching subprocesses in parallel is necessary for many workloads
- The forall loop is the ideal construct for this functionality:
  - Creates a concurrent task mapped to available resources, such as cores, gpus, nodes, etc.

```
var simulationCommands: [1..n] string = getCommands(n);
forall simulationCommand in simulationCommands {
  var process = spawn(simulationCommand, stdout=PIPE, stderr=PIPE);
  process.wait();
}
```

- Each task will block on the process.wait() call, allowing the subprocess to complete before starting a new one
- This forall maps to the local available resources, i.e. number of cores on the machine

#### LAUNCHING DISTRIBUTED SUBPROCESSES IN PARALLEL

To map the number of concurrent tasks to the available distributed resources, use the iterator explicitly: use Jobs;

```
config const nodesPerSim = 8;
config const numNodes = 128;
```

```
var job = new Job(workloadManager=Launchers.slurm);
var inputs: [0..<n] string = readInputs('inputs.txt');
const numParallelTasks = (numNodes / nodesPerSim):int;</pre>
```

```
job.alloc(numNodes, timeout=240);
forall i in simulationCommands._value.these(tasksPerLocale=numParallelTasks) {
    var process = job.run(simulationCommand, numNodes=nodesPerSim);
    process.wait();
}
```

This will allocate 128 nodes and run 16 concurrent tasks that each launch a process with 8 nodes

#### VARIABLE NUMBER OF NODES PER DISTRIBUTED APPLICATION

Varying the number of nodes in the distributed application requires tracking the available node pool

```
use ChapelLocks;
var nodesAvailable = numNodes;
var lock: chpl_LocalSpinlock;
...
forall i in simulationCommands._value.these(tasksPerLocale=numParallelTasks) {
    localNumNodes = randStream.getNext(1,8);
    waitForNodes(localNumNodes);
    var process = job.run(simulationCommand, numNodes=localNumNodes);
    process.wait();
}
```

#### **VARIABLE NUMBER OF NODES PER DISTRIBUTED APPLICATION**

```
proc waitForNodes(localNumNodes) {
  // Wait until nodes are available
  while true {
    lock.lock();
    if nodesAvailable >= localNumNodes {
      // Remove nodes from node pool
      nodesAvailable -= localNumNodes;
      lock.unlock;
      break;
    } else {
      lock.unlock();
      Time.sleep(1);
  // Add nodes back to node pool when done
  lock.lock();
  nodesAvailable += localNumNodes;
  lock.unlock;
```

#### **PARSING OUTPUT**

- I/O is an important component of any workflow application
- The output generated by the subprocess can be accessed through various mechanisms:
  - Parsing subprocess stdout
  - Parsing files generated by subprocess
  - Connecting with the subprocess and communicating in memory, e.g. over ZMQ
  - Writing results to a database from the subprocess

# **ADVANTAGES AND DISADVANTAGES OF CHAPEL**

#### **Advantages**

- Modern programming language with modern features and productivity
  - Generics, type inference, native C/python interop, etc.
- Parallel constructs built into the language
  - Parallel loops, atomics, tasks, etc.
- Performance
  - Not always important in workflow space, but can be important for sufficiently large workflow programs
  - Some workflow tools use complex algorithms in their feedback loops, e.g. Bayesian optimization in HPO
- Compiling and statically linking produces a dependency-free binary

#### Disadvantages

- Compilation is time consuming
- Standard library and module ecosystem not as mature as other languages such as Python

#### **ALTERNATIVE OPTIONS FOR BUILDING WORKFLOWS**

- Shell scripting
- Other modern programming languages: Python, Julia, Go, Rust, etc.
- Workflow frameworks, such as Apache Airflow
- Domain-specific workflow frameworks, such as Bioclipse
- Domain-specific languages for workflow automation, such as swift scripting language

#### SUMMARY

- Developing HPC workflows is a challenged faced by many scientists and engineers
- Chapel has many appealing features making it a competitive choice for developing workflows
  - Modern PL features
  - Parallel constructs
  - Performance
  - Portability

# THANK YOU