# Using Job Arrays on Hydra (slides, long version)

### In the intro portion of the workshop you will learn:

- What are jobs arrays, when and why use them?
- How to write job arrays scripts.
- How to submit jobs arrays:
  - task range, increment and limit concurrent tasks.
- Job arrays tips and tricks.
- Parallel job arrays.
- How to consolidate small tasks in job arrays.
- How to manage job arrays:
  - qstat(+), qdel, qalter, qacct(+).
- HPC wiki on job arrays:
  - https://confluence.si.edu/display/HPC/Job+Arrays

## Introduction

### What are jobs arrays, when and why use them?

- Job arrays allow you to run the same job file multiple times in a single job submission.
- They are typically used for running a given analysis on different input files or parameters.
- They allow you to use the same job file and a single qsub to run a type of analysis instead of writing a myriad of very similar job files.
- Job arrays have each a unique job id with multiple task ids

# How to write job arrays scripts

### A trivial example

- Job array scripts, or job files, are like any other job file, except that they have a task identifier stored in the variable SGE\_TASK\_ID
- Example:

```
echo + `date` $JOB_NAME started on $HOSTNAME in $QUEUE \
  with jobID=$JOB_ID and taskID=$SGE_TASK_ID
  model < model.$SGE_TASK_ID.inp
  echo = `date` $JOB_NAME for taskID=$SGE_TASK_ID done.</pre>
```

this example runs model using the input file model.N.inp

# How to submit jobs arrays

### That trivial example can be queued on 100 tasks with

```
qsub -t 1-100 trivial_example.job
Your job-array NNNNNN.1-100:1 ("trivial example.job") has I
  this gueues one job with 100 tasks, or the equivalent of 100 job
files with
    model < model.1.inp
in test1.job
    model < model.2.inp
in test2.job, etc..., up to
    model < model.100.inp</pre>
in test100.job - hence one job file instead of 100.
  this assumes that you have 100 input files called model.1.inp,
model.2.inp, ..., model.100.inp
```

# A more complete job array file

# task range and limit concurrent tasks: csh syntax

```
/bin/csh
#
#$ -N model-100 -cwd -j y -o model.$TASK_ID.log
#$ -t 1-1000 -tc 100
#
echo + `date` $JOB_NAME started on $HOSTNAME in $QUEUE \
with jobID=$JOB ID and taskID=$SGE TASK ID
#
set INPUT = model.$SGE TASK ID.inp
set OUTPUT = model.$SGE TASK ID.out
./model < $INPUT > $OUTPUT
#
echo = `date` $JOB NAME for taskID=$SGE TASK ID done.
```

#### task range and limit concurrent tasks: sh syntax

```
# /bin/sh
#
#$ -N model-100 -cwd -j y -o model.$TASK_ID.log
#$ -t 1-1000 -tc 100
#
echo + `date` $JOB_NAME started on $HOSTNAME in $QUEUE \
with jobID=$JOB ID and taskID=$SGE TASK ID
#
INPUT=model.$SGE_TASK_ID.inp
OUTPUT=model.$SGE TASK ID.out
./model < $INPUT > $OUTPUT
#
echo = `date` $JOB NAME for taskID=$SGE TASK ID done.
```

#### Note

- Task range and max concurrent task embedded in the script -t 1-1000 -tc 100
- Different log file and output file for each tasko model.\$TASK\_ID.log

```
INPUT=model.$SGE_TASK_ID.inp
OUTPUT=model.$SGE_TASK_ID.out
./model < $INPUT > $OUTPUT
```

- Also:
  - use \$TASK\_ID in embedded -o directive
  - use \$SGE\_TASK\_ID in the script

# Job arrays tips and tricks

### Various ways of using the task id \$SGE\_TASK\_ID

- formatting trick
- using awk
- 3 using sed
- 4 using bc
- 5 using cd
- 6 using <<EOF
- using your own tool (sh, csh, awk, perl, python, C, Fortan, ...)

### Formatting: replacing $1,2,\ldots,100$ by $001,002,\ldots,100$

```
csh syntax
0 i = $SGE_TASK_ID
set I = `echo $i | awk '{printf "%3.3d", $1}'`
```

```
sh syntax
let i=$SGE_TASK_ID
I=$(echo $i | awk '{printf "%3.3d", $1}')
```

# Using awk to extract parameters from a single file

```
csh syntax
0 i = $SGE_TASK_ID
set P = (`awk "NR==$i" parameters-list.txt`)
```

sh syntax
let i=\$SGE\_TASK\_ID
P=\$(awk "NR==\$i" parameters-list.txt)

the variable P will hold the content of the i-th line of parameters-list.txt, and can be used as: ./compute \$P assuming compute takes parameters.

### Using sed and a template

to replace NNN in the template by the task id

```
csh syntax
i = $SGE_TASK_ID
sed "s/NNN/$i/" input-template.inp > model.$i.inp
model < model.$i.inp > model.$i.out
```

sh syntax
let i=\$SGE\_TASK\_ID
sed "s/NNN/\$i/" input-template.inp > model.\$i.inp
model < model.\$i.inp > model.\$i.out

#### Using bc to run models on temperatures grid

- start at 23.72 and increase by 2.43 increments,
- replace TP by the temperature and NNN by the task id

#### csh syntax

```
@ i = $SGE_TASK_ID
set tp = `echo "23.72 + $i*2.43" | bc`
sed -e "s/NNN/$i/" -e "s/TP/$tp/" input-template.inp \
    > model.$i.inp
```

#### sh syntax

```
let i=$SGE_TASK_ID
tp=$(echo "23.72 + $i*2.43" | bc)
sed -e "s/NNN/$i/" -e "s/TP/$tp/" input-template.inp \
    > model.$i.inp
```

## Using cd and different directotories for each task

```
csh syntax
@ i = $SGE_TASK_ID
cd task.$i
model < model.inp > model.out
```

sh syntax
let i=\$SGE\_TASK\_ID
cd task.\$i
model < model.inp > model.out

assumes there is a model.inp file in each task.N directory



#### Using the <<EOF construct

```
csh syntax
@ i = $SGE_TASK_ID
set tp = `echo "23.72 + $i*2.43" | bc`
model <<EOF > model.$.out
temp=$tp
EOF
```

```
sh syntax
let i=$SGE_TASK_ID
tp=$(echo "23.72 + $i*2.43" | bc)
model <<EOF > model.$.out
temp=$tp
EOF
```

## Using your own tool, mytool, to convert a task id to parameters

```
csh syntax
@ i = $SGE_TASK_ID
set P = (`./mytool $i`)
./compute $P
```

```
sh syntax
i=$SGE_TASK_ID
P=$(./mytool $i)
./compute $P
```

- mytool can be anything you want:
  - any shell script, awk, perl, python, C, fortran, ...

# How to consolidate small tasks in job arrays.

### Why?

- Each task is started like a job, hence has the same overhead as starting one job
- $lue{}$  Users should avoid running lots of very short tasks (< 10-30m)
- It is relatively easy to consolidate short tasks into longer ones, using the task increment:
  - qsub -t 200-500:20 will run tasks with id=200,220,240,...,500

#### How?

use the variables:

```
$SGE_TASK_FIRST
$SGE_TASK_LAST
$SGE_TASK_STEPSIZE
$SGE_TASK_ID
```

#### Example to consolidate short tasks

```
csh syntax
@ iFr = $SGE TASK ID
@ iTo = $iFr + $SGE_TASK_STEPSIZE - 1
if ($iTo > $SGE TASK LAST) @ iTo = $SGE TASK LAST
#
echo running model.csh for taskIDs $iFr to $iTo
@ i = \$iFr
while ($i <= $iTo)
  ./model.csh $i >& model-$i.log
  @ i++
end
```

#### Example to consolidate short tasks

```
sh syntax
let iFr=$SGE TASK ID
let iTo=$iFr+$SGE_TASK_STEPSIZE-1
if [ $iTo -gt $SGE_TASK_LAST ]; then
  let iTo=$SGE TASK LAST
fi
#
echo running model.sh for taskIDs $iFr to $iTo
let i=$iFr
for ((i=$iFr; i<=iTo; i++)); do
  ./model.sh $i >& model.$i.log
done
```

# assuming that

■ the script model.csh or model.sh do the work and takes one argument: the id.

# Parallel job arrays

### Job arrays can run parallel tasks

- Each task request a parallel environment, as per the -pe specification:
  - -pe mthread N for multi-threaded
  - -pe mpich N or -pe orte N for MPI
- Check the HPC wiki for more info at https://confluence.si.edu/display/HPC/Job+Arrays

# How to manage job arrays

### job status with qstat or qstat+

- list task id(s) for running jobs single job id
- remaining task range for queued jobs

### job deletion with qdel

- can delete specific tasks w/ -t flags
- otherwise delete all the tasks: running and queued!

### job modification with qalter

- supports -t and -tc flags
- use -tc if and when needed, can be used to increase its value progressively

### job accounting with qacct or qacct+

- use -t flag to get info on specific task(s)
- otherwise return info on all the tasks (can be (very) long)

# Also remember

#### Can run a lot of concurrent tasks

- separate name spaces
  - some of the tasks will run at the same time
  - should not write in the same file
- I/O contention:
  - concurrent tasks read the same file(s)?
- manage the results files
  - esp. if a lot are created in the same directory
- test on a small set of tasks first
- avoid sending emails with -m abe
  - it applies to each task (lots of emails).

#### Resource limits:

- \* Limit of 10,000 tasks per job array,
- \* Limit of 2,500 jobs per user in the queue, and
- \* Limit of 25,000 jobs in the queue (for all users).

# HPC wiki on job arrays

goto https://confluence.si.edu/display/HPC/Job+Arrays



# Hands on portion