STATS/DATASCI 531 Winter 2022 Parallel statistical computing in R on greatlakes

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1 Logging in to greatlakes

Requirements

We follow Section 1.2 of the greatlakes user guide. As preliminaries, you need:

- A Slurm account. Everybody in this class is a member of the account stats531w22_class. Graduate students in the Applied Statistics and Data Science masters programs, or Statistics PhD program, also have a primary departmental account, stats_dept1, and a smaller departmental backup account for if you exhaust your resourses, stats_dept2.
- A greatlakes cluster login account. If you have not yet filled in the form at https://arc-ts.umich.edu/greatlakes/user-guide/ then do so.
- A umich internet address. Use the umich VPN if you are not on campus.

Connecting to greatlakes with macOS or Linux

1. Open a Terminal window (recall that, on a Mac, this can be done using Control-Spacebar and typing Terminal) and type

```
ssh uniqname@greatlakes.arc-ts.umich.edu
```

where uniquame is your uniquame.

2. Login with your Kerberos level-1 password, and Duo two-factor authentication.

This creates a remote terminal shell on greatlakes.

Connecting to greatlakes with Windows

This is essentially the same as for macOS, except that you may need to install a program that provides a terminal window

- 1. Follow instructions to install PuTTY at https://documentation.its.umich.edu/node/350
- 2. Launch PuTTY and enter greatlakes.arc-ts.umich.edu as the host name, then click open. If you receive a "PuTTY Security Alert" pop-up, this is completely normal, click the "Yes" option. This will tell PuTTY to trust the host the next time you want to connect to it. From there, a terminal window will open; you will be required to enter your UMICH uniquame and then your Kerberos level-1 password in order to log in. Please note that as you type your password, it may be that nothing you type appears on the screen; this is completely normal. Press "Enter/Return" key once you are done typing your password.
- 3. Complete the request for Duo two-factor authentication.

This creates a remote terminal shell on greatlakes.

2 Moving files on and off greatlakes

Moving files on and off greatlakes: scp

On Mac or Linux, you can use scp which has similar syntax to cp.

To copy myfile on your laptop to a subdirectory mydir of your home directory on greatlakes:

```
scp myfile uniqname@greatlakes-xfer.arc-ts.umich.edu:mydir
```

To copy an entire directory, use the -r flag for recursive copy:

```
scp -r mydir uniqname@greatlakes-xfer.arc-ts.umich.edu:
```

These commands can also be reversed to copy files from greatlakes to your machine. The following copies mydir back to the current working directory:

```
scp -r uniqname@greatlakes-xfer.arc-ts.umich.edu:mydir .
```

You will need to authenticate via Duo to complete the file transfer. On Windows, you can use WinSCP or FileZilla.

3 Working with batch jobs

Cluster batch workflow

- 1. You create a batch script and submit it as a job
- 2. Your job is scheduled, and it enters the queue
- 3. When its turn arrives, your job will execute the batch script
- 4. Your script has access to all applications and data
- 5. When your script completes, anything it sent to standard output and error are saved in files stored in your submission directory
- 6. You can ask that email be sent to you when your jobs starts, ends, or fails
- 7. You can check on the status of your job at any time, or delete it if it's not doing what you want
- 8. A short time after your job completes, it disappears

Useful batch commands

Submit a job

```
sbatch sample.sbat
```

Query job status

```
squeue -j jobid
squeue -u uniqname
```

Delete a job

```
scancel jobid
```

Check a job script and estimate its start time

```
sbatch --test-only sample.sbat
```

More Slurm commands to try

sacct -u user show recent job history seff jobid show cpu utilization for jobid

my_accounts list accounts you have permission to use

4 R modules on greatlakes

R modules on greatlakes

Sotware on greatlakes is packaged in modules which must be loaded

```
module load R
```

Other versions of R are available:

```
module avail R
```

• We see that R4.1.0 is currently the default. For simple multicore computing, sending jobs to multiple cores on a single node, the default R module is appropriate.

5 A test for foreach

Set up test for foreach

- The greatlakes subdirectory of the 531w22 git repository has a file test.sbat which submits a batch job running the parallel foreach test in test.R.
- A basic Linux exercise is to set up a directory on greatlakes with these files, at which point you can run

```
sbatch test.sbat
```

to submit the job. If you have little or no experience with Linux or Unix, this is a nontrivial task. You could ask for help or read https://ubuntu.com/tutorials/command-line-for-beginners.

• You can transfer the files from your laptop via scp, or by copy-paste, but it may be simplest to clone the class git repository into your greatlakes account,

6 Editing text files on greatlakes

Editing text files on greatlakes

• Inspect the text file test.sbat, for example by

```
more test.sbat
```

Is it fairly self-explanatory?

- One thing that needs changing is to set your email address for alerts about jobs beginning and ending.
- To make these edits on greatlakes, you need a text editor.
- It is convenient to use a text editor that runs in a terminal. Options include

```
vi test.sbat
emacs -nw test.sbat
nano test.sbat
```

• It is useful to have some familiarity with each of these editors.

Comparing results

• You are now ready to run a batch job

```
sbatch test.sbat
```

- From inspecting the code in test.R, we see that the results are saved in test.csv
- You can assess what you learn from comparing run times on greatlakes and your laptop, though the main goal here is just to practice running the code.

7 Other ways to run R on greatlakes

Other ways to run R on greatlakes

• It is sometimes useful to start an interactive session on greatlakes, particularly for debugging. This is done from the terminal as follows:

```
module load R
srun --nodes=1 --account=stats531w22_class --ntasks-per-node=8 \
    --pty /bin/bash
```

• You can then run R in the terminal as usual, just by typing

- This R session will have access to the cores you have requested.
- Here, we require nodes=1 since library(doParallel) alone cannot work with cores spread across different machines.
- You can also run web-based Rstudio However, batch jobs remain the basic tool for intensive statistical computing.

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