COMP 322 Spring 2022

Lab 8: Java Threads Instructors: Mack Joyner, Zoran Budimlić

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Goals for today's lab

• Experimentation with Java threads

- NOTS Setup (Section 3)
- An introduction to the shell (Section 4)

This lab can be downloaded from the following GitHub repository https://classroom.github.com/a/_ffDUUuD. Use the git command-line client or IntelliJ to checkout the project into appropriate directories locally.

In today's lab, you need to use NOTS to run performance tests. If you need any guidance on how to submit jobs on NOTS manually, please ask a member of the teaching staff.

1 Conversion to Java threads: Spanning Tree

- 1. The SpanningTreeSeq.java program is an example sequential solution to the spanning tree problem. The SpanningTreeAtomicHjLib.java program is a provided parallel solution to the spanning tree problem. This version uses finish and async constructs along with an AtomicReference.
- 2. Your task is to convert SpanningTreeAtomicHjLib.java to a Java program that uses threads instead of HJlib tasks. You should modify the provided SpanningTreeAtomicThreads.java file, and use Java thread methods instead of finish/async. (The AtomicReference calls can stay unchanged.) There are TODOs in the file to guide you.
- 3. You have been provided with tests for your parallel spanning tree implementation in SpanningTreePerformanceTest. To complete this portion of the lab, you should submit these performance tests to NOTS by modifying the provided myjob.slurm template and submitting manually.

2 Programming Tips and Pitfalls for Java Threads

- Remember to call the start() method on any thread that you create. Otherwise, the thread's computation does not get executed.
- Since the join() method may potentially throw an InterruptedException, you will either need to enclose each call to join() within a try-catch block, or add a throws InterruptedException clause to the definition of the method that includes the call to join().

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3 NOTS setup

NOTS (Night Owls Time-Sharing Service) is a Rice compute cluster designed to run large multi-node jobs over a fast interconnect. The NOTS cluster consists of 80 nodes. Each node is equipped with two Intel E5-2650v2 Ivy Bridge EP processors for a total of 16 cores per node. Each core can support threads, each running at 2.6GHz. Memory sizes on the compute nodes range from 32-128GB RAM. We'll be using NOTS in 322 to run real performance tests on high-performance parallel hardware. Not only does NOTS give you an isolated environment to test in (without worrying about interference from other processes), but it likely has at least 4x the cores and RAM of your laptop. The main difference between using NOTS and using your laptop is that NOTS allows you to gain access to dedicated compute nodes to obtain reliable performance timings for your programming assignments. On your laptop, you have less control over when other processes or your power manager might start stealing cores or memory from your running parallel program.

• Start by logging in to NOTS. From the command line on Mac/Linux/Windows:

```
$ ssh \(\forall your-netid\) @nots.rice.edu
\(\forall your-netid\) @nots.rice.edu's password: \(\forall your-password\)
```

Your password should be the same as the one you use to log in to other Rice services like CLEAR or Owlspace. Note that this login connects you to a shared login node, not a dedicated compute node.

• After you have logged in to NOTS, run the following command to setup the JDK 8 and Maven path. Do not replace "jmg3" with your net ID.

```
source /home/jmg3/comp322/322_setup.sh
```

Note: You will have to run this command each time you log on NOTS. You should add this command to the bottom of the file \sim /.bash_profile so that it will run automatically each time you log in.

• Check your installation by running the following commands:

```
which java
```

You should see the following: /opt/apps/software/Core/Java/1.8.0/bin/java

Check java installation:

```
java -version
```

You should see the following:

```
java version "1.8.0" Java(TM) SE Runtime Environment (build 1.8.0) Java HotSpot(TM) 64-Bit Server VM (build 25.45-b02, mixed mode)
```

Check maven installation:

```
mvn --version
```

You should see the following:

```
Apache Maven 3.3.9 (bb52d8502b132ec0a5a3f4c09453c07478323dc5; 2015-11-10T10:41:47-06:00)

Maven home: /home/jmg3/install/apache-maven-3.3.9 Java version: 1.8.0, vendor:

Oracle Corporation Java home: /opt/apps/software/Core/Java/1.8.0/jre Default

locale: en_US, platform encoding: UTF-8 OS name: "linux", version: "3.10.0-229.e17.x86_64",

arch: "amd64", family: "unix"
```

- When you log on to NOTS, you will be connected to a *login node* along with many other users (type users at the terminal if you would like to see all the other people with active sessions on the same machine). Running your performance experiments on the login node would face the same problems as on your laptops: in a multi-tenant system, it is hard to prevent interference with your performance results. Once you have an executable program, and are ready to run it on the compute nodes, you must create a job script that configures and runs your program on a dedicated compute node. The following items will walk you through editing a job script we have provided, uploading your Lab 5 project to NOTS, and then launching a compute job on NOTS using the job script.
- The job script will provide information on your job such as the number of nodes you want to use (usually just one in this course), the amount of time you want your job to be allowed to run for, the amount of memory your job will need, as well as the actual commands you want to be executed as part of your job. We have provided a script template in the Lab 8 template code at:

```
lab8-GITID/src/main/resources/myjob.slurm
```

You need to change the lines marked "TODO". The change on line 8 of that file indicates an e-mail address to send job status notifications to.

- Now, we want to transfer your lab8 folder from your local machine up to NOTS so that we can submit the Lab 8 code to the cluster for testing. NOTE: You must place your lab8 folder directly inside of your \$HOME directory, so that the absolute path to it is /home/[net-id]/lab8. To transfer a folder to NOTS, you can use one of two methods:
 - Use GitHub: You can commit your local changes to GitHub. Then you can checkout or update the project on your NOTS account using one of the following:

```
git clone https://github.com/RiceParProgCourse/lab8-GITID.git
```

or, if you have already checked out the GitHub project on your account,

```
git fetch then git checkout origin/master
```

The git fetch and checkout commands must be run from your \$HOME directory.

- Use SCP: Use the following command on your local machine to transfer your lab8 folder to NOTS:

```
scp -r /local/path/to/lab8 [your-net-id]@nots.rice.edu:~/
```

• Now that we have edited the lab 8 job script and uploaded the lab 8 folder to NOTS, we can submit a test job to the cluster. To submit the job, run the following command on NOTS:

```
sbatch /path/to/my/lab8/src/main/resources/myjob.slurm
```

After you have submitted the job, you should see the following:

```
Submitted batch job [job number]
```

• To check the status of a submitted job, use the following command:

```
squeue -u [your-net-id]
```

If no jobs are listed, you have no running jobs.

• Because NOTS is shared by many users, it may take some time for your job to make it through the job queue and get assigned a compute node. If you would like to see an estimate of when your job will start, you can add the --start flag to the squeue command:

```
squeue -u [your-net-id] --start
```

Note that adding --start will only show jobs that are pending and have not started yet. If your job is already running, you will see it listed under squeue -u [net-id] but not squeue -u [net-id] --start.

• To cancel a submitted job, use the following command:

scancel [job-id]

When your job finishes running, your should see an output file titled slurm-[job-id].out in the same directory from which you submitted the job. This file contains the output produced by your job during its execution on a dedicated compute node. You can either transfer these output files back to your laptop for viewing, or use a tool on the cluster to open them (such as cat, less, emacs, or vim).

4 A Quick Introduction to Useful Shell Commands

As we move more into using NOTS to measure real-world performance, being comfortable with the shell will become more and more useful in 322. This section briefly introduces you to the most commonly used shell commands. There are no required "tasks" or "goals" for this section beyond improving your awareness of these commands. However, you are encouraged to play with these commands in your own terminal (on your laptop or on NOTS) to help build familiarity. We have provided a sample set of exercises at the end of this section that you can use to help learn the commands.

Linux relies heavily on an abundance of command line tools. Most input lines entered at the shell prompt have three basic elements: command, options, and arguments. The command is the name of the program you are executing. It may be followed by one or more options (or switches) that modify what the command may do. Options usually start with one or two dashes, for example, -p or --print, in order to differentiate them from arguments, which represent what the command operates on. The interested student is encouraged to view Chapter 7 in the LinuxFoundationX: LFS101x.2 Introduction to Linux course on edX for a more thorough introduction on the commands covered in lab today.

In order to help you organize your files, your file system contains special files called directories. A directory is a logical section of a file system used to hold files. Directories may also contain other directories. Directories are separated by a forward slash (/). The current directory, regardless of which directory it is, is represented by a single dot (.). The parent directory of the current directory (i.e., the directory one level up from the current directory) is represented by two dots (..). Your home directory is the directory you are placed in, by default, when you open a new terminal session. It has a special representation: a tilde followed by a slash $(^{\sim}/)$.

Below are some of the most used commands, along with brief descriptions of each.

4.1 pwd

pwd means "print working directory". It is used to output the path of the current working directory.

4.2 mkdir

mkdir means "make directory" and is used to create new directories. It creates the directory(ies), only if they do not already exist. Use the -p flag to ensure parent directories are created as needed.

4.3 cd

cd means "change directory". The cd command is used to change the current working directory. It can be used to change into a subdirectory, move back into the parent directory, move all the way back to the root directory, or move to any given directory. When the first character of a directory name is a slash, that denotes that the directory path begins in the root directory.

4.4 ls

ls means "list directory". The ls command lists out the contents of the directory you are currently in. You can use cd to change into different directories and then list what's in them so I know which directory to go to next.

4.5 dirs, pushd and popd

pushd means "push directory". popd means "pop directory". Both commands are used to work with the command line directory stack. The pushd command saves the current working directory in memory so it can be returned to at any time, optionally changing to a new directory. The popd command returns to the path at the top of the directory stack. This directory stack can be viewed by the command dirs.

4.6 rm

rm means "remove". Directories and files can be removed (deleted) with the rm command. By default, it does not remove directories. If the -r (--recursive) option is specified, however, rm will remove any matching directories and their contents. Use the -f (--force) option to never be prompted while files are being removed. The -v (--verbose) option can be used to get rm to detail successful removal actions.

4.7 cp

cp means "copy a file or directory". The command has three principal modes of operation, expressed by the types of arguments presented to the program for copying a file to another file, one or more files to a directory, or for copying entire directories to another directory. The commands takes two arguments, source and destination files, which may reside in different directories. You can use cp to copy entire directory structures from one place to another using the -R option to perform a recursive copy. Using this option copies all files, and all subdirectories from the source to the destination directory, you can specify multiple files as the source, and a directory name as the destination.

4.8 mv

mv means "move a file or directory". It moves one or more files or directories from one place to another, it is also used to rename files. When a filename is moved to an existing filename (in the same directory), the existing file is deleted. Use the -f (--force) option to never be prompted before overwriting existing files. The -v (--verbose) option can be used to get details on the actions and destination locations of the files being moved.

4.9 touch

The touch command is used to make an empty file. The touch command is the easiest way to create new, empty files. Touch eliminates the unnecessary steps of opening the file, saving the file, and closing the file again. It is also used to change the timestamps on existing files and directories.

4.10 less

It is used to view (but not change) the contents of a text file one screen at a time. Unlike most Unix text editors/viewers, less does not need to read the entire file before starting, resulting in faster load times with large files. You can open a file by passing the file name as an argument to the command. To traverse the file press the following, use the down arrow to scroll down one line while using the up arrow scrolls up one line. Using q will exit the less command.

4.11 cat

The cat command prints the contents of a file to screen and can be used to concatenate and list files. The name is an abbreviation of catenate, a synonym of concatenate. cat will concatenate (put together) the input files in the order given, and if no other commands are given, will print them on the screen as standard output. It can also be used to print the files into a new file as follows: cat old1.txt old2.txt > newfile.txt Typing the command cat followed by the output redirection operator and a file name on the same line, pressing ENTER to move to the next line, then typing some text and finally pressing ENTER

again causes the text to be written to that file. The program is terminated and the normal command prompt is restored by pressing the CONTROL and d keys simultaneously.

4.12 find

The find command is a very useful and handy command to search for files from the command line. find will search any set of directories you specify for files that match the supplied search criteria. You can search for files by name, owner, group, type, permissions, date, and other criteria. The search is recursive in that it will search all subdirectories too. All arguments to find are optional, and there are defaults for all parts.

4.13 grep

The grep command is used to search text or searches the given file for lines containing a match to the given strings or words. Its name comes from the ed command g/re/p (globally search a regular expression and print), which has the same effect: doing a global search with the regular expression and printing all matching lines. By default, grep displays the matching lines. You can force grep to ignore word case with the -i option.

4.14 man

The man command is used to format and display the system's reference manuals. The man pages are a user manual; they provide extensive documentation about commands. Each argument given to man is normally the name of a program, utility or function. man is most commonly used without any options and with only one keyword. The keyword is the exact name of the command or other item for which information is desired, e.g. man 1s.

5 Testing on NOTS

Now that we have an implementation of a parallel spanning tree with threads, we will test the performance on the NOTS cluster to measure the actual performance of the implementation without interference on your laptop.

To do so, you should use the provided myjob.slurm file. As usual, when using the myjob.slurm file please open it to fix any TODO items.

6 Turning in your lab work

For lab 8, you will need to turn in your work by Wednesday, March 23, 2022 at 4:30 pm, as follows.

- 1. Show your work to an instructor or TA to get credit for this lab. In particular, the TAs will want to see the output of testSpanningTreeThreads running on NOTS through the provided SLURM script.
- 2. Commit your work to your lab8 folder. Check that all the work for today's lab is in your lab8 directory by opening https://classroom.github.com/a/_ffDUUuD in your web browser and checking that your changes have appeared.