

Lab 9: Java Threads

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1 Turning in your lab assignments — **NEW!**

Starting with today's lab, we're asking all COMP 322 students to turn in their lab assignments before leaving. You will need to do the following:

1. Create a directory called `lab_9/` in your SUGAR account.
2. Do all your work for today's lab in this directory.
3. Before you leave, create a zip file of your work by changing to the parent directory for `lab_9/` and issuing the following command, "`zip -r lab_9.zip lab_9`".
4. Use the turn-in script to submit the contents of the `lab_9.zip` file as a new `lab_9` directory in your `turnin` directory.

NOTE: the purpose of the lab submission is to supplement the attendance record with the level of completeness of your lab work. You will not be graded on the quality of your work. You will only receive full credit for attendance if you also worked on the assigned problem during the lab.

2 Setup on SUGAR

As before, run the following command on SUGAR to setup the environment for executing HJ and Java programs:

```
source /users/COMP322/hjsetup.txt
```

To request a dedicated *compute node*, you should use the following command (as usual) from a SUGAR login node:

```
qsub -q commons -I -V -l nodes=1:ppn=8,walltime=00:30:00
```

When successful, it will give you a command shell on a dedicated 8-core compute node for your use for 30 minutes at a time. Your home directory is the same on both the login and compute nodes.

3 Convert to Java Threads - N-Queens

1. Download the `nqueens.hj` program from lab 4 that uses `finish` and `async` constructs along with `AtomicInteger` calls.
2. Convert it to a pure Java program by using Java threads instead of `finish/async`, using the concepts introduced in Lecture 27. (The `AtomicInteger` calls can stay unchanged.) For simplicity, you can include `joins` within each call to `nqueens_kernel()`. This is correct, but more restrictive than the `finish/async` structure for the given code. But it simplifies parallelization using Java threads.
3. Compile and run the program as follows to solve the N-Queens problem on a 14×14 board. Try experimenting with different values for `cutoff_value` if needed.

```
javac nqueens.java
java nqueens 14
```
4. Compare the performance of your Java program with the given HJ program. (It is recommended that you use separate directories for compiling the Java and HJ versions so as to avoid any possible interference among classfiles generated for both versions.)

4 Convert to Java threads - Spanning Tree

1. Download the `spanning_tree_atomic.hj` solution from lab 7 that uses `finish` and `async` constructs along with `AtomicReference` calls.
2. Convert it to a pure Java program by using Java threads instead of `finish/async`, using the concepts introduced in Lecture 27. (The `AtomicReference` calls can stay unchanged.) For simplicity, you can include `joins` within each call to `compute()`. This is correct, but more restrictive than the `finish/async` structure for the given code. But it simplifies parallelization using Java threads.
3. Compile and run the programs as follows with the default input size.

```
javac spanning_tree_atomics.java  
java spanning_tree_atomic
```
4. Compare the performance of your Java program with the given HJ program. (It is recommended that you use separate directories for compiling the Java and HJ versions so as to avoid any possible interference among classfiles generated for both versions.) You may choose to add cutoff threshold values for this program as was done for N-Queens, so as to limit the number of Java threads that will be created.