Peri Hassanzadeh ECE1570 High Performance Computing Dr. Bigrigg December 3, 2021

# **Tightly and Loosely Coupled Systems**

#### Introduction

In this assignment, I was able to use threads to develop data parallel algorithms as well as clusters to gain a better understanding maximizing the usage of a system to achieve as much speedup as possible utilizing multiple configurations of processors. I identified the speedup I could achieve by using multiple processors on the CRC server.

The main objective of each program no matter it's algorithm was to use Dijkstra's algorithm to find the shortest path from the source node (0,0) to each other node in the 100 by 100 matrix. I was then able to split the matrix into four smaller matrices of 50 by 50. After calculating the shortest path with Dijkstra's algorithm, I output the results of each path into a text file for the data parallel and MPI versions. The timing of each version of the project was calculated and run five times each to get an average result which will be further analyzed.

In order to run each file, make sure that all .cpp files are in the same folder as the working directory. Windows refers to my local environment and Linux refers to the server environment and the output files will be in the same directory after the exe is run. This format will allow for successful compilation of each implementation resulting in an executable file with the following commands:

Serial command:

g++ P2\_serial.cpp -o P2\_serial

Data Parallel command:

Windows: g++ P2 thread.cpp -o P2 thread -lpthread

Linux: sbatch P2\_thread.script

MPI command:

Linux: sbatch P2 mpi.script

Finally, note that each implementation could be optimized in various ways. I will go over a few points that I wish I would have been able to develop on given better time management.

#### **Serial Version**

The files that correspond with the serial version are as follows:

P2\_serial.cpp - implementation of serial version

P2 serial.exe - compiled program file

P2\_serial.script - Used to run on CRC

Serial.out - Results from CRC script

#### **Screenshots of Code:**

```
//Peri Massamzadeh
//Ectis70 High Performance Computing
//Date of Creation 11/28/21
//Date of Creation 11/28/21
//Date is the serial version of Project 2
//Project 2 analyzes the speedup achieved using multiple configurations of processors.
//Import cat libraries
//Impo
```

```
##/bin/bash
#5BATCH --job-name=serial
#5BATCH --nodes=1 #number of nodes requested
##5BATCH --nodes=1 #number of nodes requested
##5BATCH --ntasks-per-node=1
#5BATCH --eluster-smp # mpi, gpu and smp are available in H2P
#5BATCH --partition-smp # available: smp, high-mem, opa, gtx1080, titanx, k40
#5BATCH --partition-smp # available: smp, high-mem, opa, gtx1080, titanx, k40
#5BATCH --mail-type=!ND,FAIL # ... job ends or fails
#5BATCH --mail-type=!ND,FAIL # ... job ends or fails
#5BATCH --time=10:00 # walltime in dd-hh:mm format

#5BATCH --output=serial.out # the file that contains output
module purge #make sure the modules environment is sane
module load intel/2017.1.132 intel-mpi/2017.1.132 fhiains/160328_3

cp P2_serial.cpp $SLURM_SCRATCH # Copy inputs to scratch

cd $SLURM_SCRATCH # change directory

# Set a trap to copy any temp files you may need
run_on_exit()

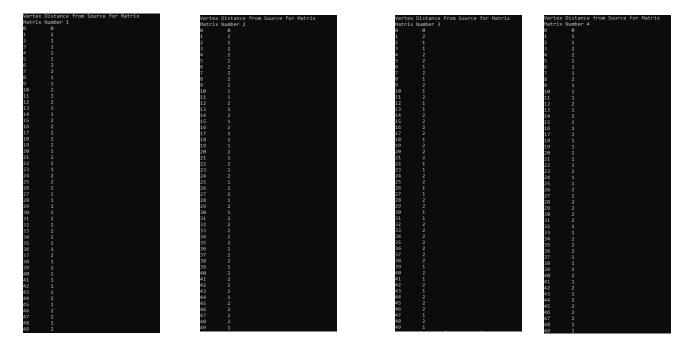
cp -r $SLURM_SCRATCH/* $SLURM_SUBMIT_DIR
}

trap run_on_exit EXII
gH+ P2_serial.cpp -std=c99 -o main.o #compile the program, set the runnable filename
...main.o # Run the runnable
crc-job-stats.py # gives stats of job, wall time, etc.
```

The screenshots on the previous page are for the serial implementation of the project. The code may also be viewed in the submitted files as P2\_serial.cpp and P2\_serial.script. Zoom in for a better view of the code or open the specified file.

#### Output:

From CRC Server:



#### Execution time:

```
Total time for serial execution: 1340 microseconds

Total time for serial execution: 1256 microseconds

Total time for serial execution: 1279 microseconds

Total time for serial execution: 1277 microseconds

Total time for serial execution: 1329 microseconds
```

From the CRC, the average execution took 1296 microseconds which is significantly less than that of my local machine which had an average of 1600 microseconds. Additionally, I will make sure to analyze the rest of the execution times for data parallel and clusters from the server as well as my local machine. Comparing the two values will allow me to properly decide proper scaling and different implementation benefits and drawbacks.

## **Data Parallel (Thread) Version**

The files that correspond with the data parallel version are as follows:

P2\_thread.cpp - implementation of data parallel version

P2\_thread.exe - compiled program file

P2\_thread.script - to compile on CRC

Thread.out - Result from running on CRC script

First through fourthGraph.txt - Text files containing results

#### **Screenshots of Code:**

```
1 //Peri Hassanzadeh
2 //EcE1370 High Performance Computing
3 //Date of Creation: 12/1/21
4 //Last Update: 12/3/2021
5 //
6 //This is the data parallel (thread) version of Project 2
7 //Project 2 analyzes the speedup achieved using multiple configurations of processors.
8 //I used Dijkstra's algorithm and a loex100 matrix as the basis of my research.
9 //Import c++ libraries
10 finclude cstdio.hb
11 elinclude cstdio.hb
12 elinclude cstdio.hb
13 elinclude cstdio.hb
14 elinclude cstdio.hb
15 elinclude cstdio.hb
16 elinclude cstdireamb
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18 elinclude cstdireamb
18 elinclude cstdireamb
19 elinclude cstdireamb
10 elinclude cstdir
```

```
James Annals of Struct the hold the variables in order to find completion time

for main()

for main()
```

The screenshots on the page above are for the data parallel implementation of the project. I did not include the hardcoded matrices that were based on the serial version of the code. The code may also be viewed in the submitted files as P2\_thread.cpp and P2\_thread.script. Zoom in for a better view of the code or open the specified file.

#### **Output:**

From my local machine:

```
C:\Users\perih\Desktop>g++ P2_thread.cpp -o P2_thread -lpthread

C:\Users\perih\Desktop>P2_thread

Fiocess is complete

Total time for data parallel execution: 18002 microseconds
```

The execution of the program took 18002 microseconds on my local machine.

#### CRC Server:

From the CRC, the execution took an average of 649 microseconds which is also significantly less than that of my local machine similar to what happened in the serial implementation.

The screenshots above represent the output of Dijkstra's algorithm within different text files corresponding to different matrices. It was easier to view the results within a text file since the output on the command prompt was skewed due to the threads running at the same time. Analysis became much more simple once the data was pushed to a text file.

## **Loosely Coupled (Cluster) Version**

The files that correspond with the cluster version are as follows:
P2\_mpi.cpp - implementation of clusters
P2\_mpi.script - to compile on CRC
mpi.out - Result from running on CRC script

First through fourthGraph.txt - Text files containing results

#### **Screenshots of Code:**

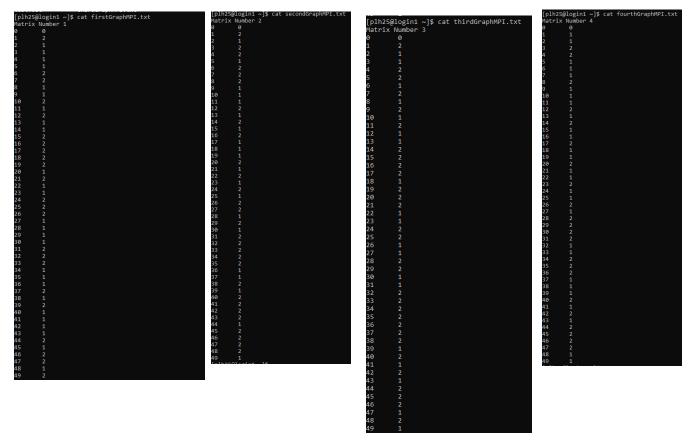
```
| Company of the comp
```

The screenshots above are for the task cluster implementation of the project. The code may also be viewed in the submitted files as P2\_mpi.cpp and P2\_mpi.script. Zoom in for a better view of the code or open the specified file.

# **Output:**

From my local machine, the program is unable to be executed.

# Bigrigg Server:



From the ssh server bigrigg.com, the execution took an average of 6.1979284e-04 seconds.

# **Optimizations**

There are multiple optimizations that could be made to each implementation, but I would like to address some of the more obvious optimizations that would make a significant difference in the implementation and possibly efficiency of the program.

Starting off specifically with the serial implementation, the algorithm should have been as stated on the Canvas webpage, but I decided to use Dijkstra's algorithm instead as discussed on the discussion board. I would have liked to have been able to find a way to split the matrices so that the source node was the same for each matrix and then the end result could be compiled together that way the results would be all for one matrix instead of four different ones. This could be applied to the thread and cluster versions as well.

Another large optimization that could be made is to create more threads in the data parallel version to be more of a streamlined process being able to break up the data even more. Specifically putting one matrix per thread.

Finally, the last optimization I wanted to mention was to be able to use more processes in the cluster version. With better time management I would have been able to better understand the purpose of clusters and be able to compare them to the threads with deeper analysis. Instead I spent time troubleshooting the CRC and using scripts.