CS633: Parallel Computing Assignment 2

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Coding Methodology

1. code.c:

Contains the MPI code to simulate the row subtraction operation mentioned in the assignment. The (0,0) entry of the matrix (A) is assumed to be the upper left entry. Each process initializes and owns its domain/division of the matrix (A). Any data points outside the process's domain that are required to update its domain points are received via P2P communication from the appropriate process. For both one-dimensional row-wise and two-dimensional decomposition, updating the last row of the domain of any process requires the first row of the domain of the process below it. For every process (except the process(es) not having any process below them), this data is received via P2P communication from the process below. We have used non-blocking MPI_Isend() and MPI_Irecv() for the communication.

A note on performance: For the one-dimensional decomposition, processes do not need to communicate the entire first row in their domain but just the entries which fall inside the lower triangular matrix. Exploiting this reduces the number of P2P communications in the one-dimensional case. In the case of two-dimensional decomposition, to preserve the brevity of the code, we send/receive the entire row of the domain irrespective of whether it lies in the upper/lower triangular matrix. Thus, to ensure a fair comparison between one and two dimensional decomposition we transmit the entire row in both the decompositions. Then, we iterate through all the elements of the domain and update them only if they lie in the lower triangular matrix for both the decompositions.

We insert an MPI_Barrier between the code segments for 1D/2D decomposition to ensure that the processes are synced before calculating the time for the two-dimensional decomposition.

The runtimes are printed on the stdout console and manually inserted in the times.txt file.

```
#include <stdio.h>
#include <stdlib.h>
#include <assert.h>
#include <time.h>
#include <mpi.h>

int main(int argc, char *argv[]){
   int myrank;
   double stime, etime, proc_time1, proc_time2, max_time1, max_time2;

MPI_Init(&argc, &argv);
```

```
int py = atoi(argv[2]); // number of processes in y
  int num procs = px * py;
  long long m = size / num_procs; // number of rows of matrix for each process
  double A1[m][size]; // sub-domain of the process
  MPI Request send request1, recv request1;
      if(myrank > 0){
          MPI Isend(A1[0], sendcount, MPI DOUBLE, myrank - 1, myrank - 1,
MPI_COMM_WORLD, &send_request1);
       if(myrank < num_procs - 1){</pre>
```

```
MPI_Irecv(buf1, recvcount, MPI_DOUBLE, myrank + 1, myrank,
MPI COMM WORLD, &recv request1);
      if(myrank > 0){
          MPI_Wait(&send_request1, &send_status1);
               long long elem row = myrank * m + i; // row of the element in
       if (myrank < num procs - 1) {</pre>
          MPI Wait(&recv request1, &recv status1);
       if(myrank < num_procs - 1){</pre>
               long long elem_row = myrank * m + (m - 1); // row of the element
                   A1[m - 1][j] = buf1[j];
```

```
MPI COMM WORLD);
  m = size / py; // number of rows of matrix for each process
  double A2[m][n]; // sub-domain of the process
          A2[i][j] = rand();
  double buf2[n]; // receive buffer
  MPI Request send request2, recv request2;
  int proc row = myrank / px; // row of the process in the virtual topology
  int proc col = myrank % px; // column of the process in the virtual topology
      if(proc row > 0){
          MPI Isend(A2[0], n, MPI DOUBLE, myrank - px, myrank - px,
MPI COMM WORLD, &send request2);
      if(proc row < py - 1){</pre>
          MPI_Irecv(buf2, n, MPI_DOUBLE, myrank + px, myrank, MPI_COMM_WORLD,
&recv request2);
      if(proc row > 0){
          MPI_Wait(&send_request2, &send_status2);
```

```
long long elem_row = proc_row * m + i; // row of the element in
                   A2[i][j] -= A2[i + 1][j];
       if(proc_row < py - 1){</pre>
           MPI_Wait(&recv_request2, &recv_status2);
       if(proc row < py - 1) {</pre>
               long long elem_col = proc_col * n + j; // column of the element
                   A2[m - 1][j] -= buf2[j];
MPI COMM WORLD);
  if(!myrank){
```

```
printf("configuration px = %d, py = %d, size = %lld, runtime 1D = %lf,
runtime 2D = %lf\n", px, py, size, proc_time1, proc_time2);
}

MPI_Finalize();
return 0;
}
```

2. run.sh:

Compiles and executes the file (code.c) for all four configurations (for px = 2, py = 2.4, size = 4096, 32768) on two hand picked servers csews25 and csews32 (having less load compared to other servers). We have run 4 processes on each node.

```
mpicc ./code.c -o code
rm -f times.txt

for i in $(seq 1 5)

do
    mpirun -np 4 -hosts csews25:4 ./code 2 2 4096

done

for i in $(seq 1 5)

do
    mpirun -np 8 -hosts csews25:4,csews32:4 ./code 2 4 4096

done

for i in $(seq 1 5)

do
    mpirun -np 4 -hosts csews25:4 ./code 2 2 32768

done

for i in $(seq 1 5)

do
    mpirun -np 4 -hosts csews25:4 ./code 2 2 32768

done

for i in $(seq 1 5)

do
    mpirun -np 8 -hosts csews25:4,csews32:4 ./code 2 4 32768

done
```

Note: the times for size = 4096 are taken by running the code on prutor and those for size = 32768 are taken by running the code on csews.

3. plot_runtimes.py:

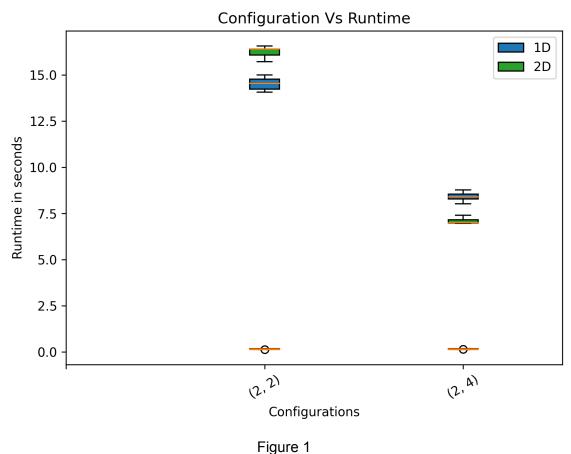
Reads the (times.txt) file and creates boxplots for the runtimes. The plots (runtimes.png) and (runtimes_size_4096.png) are generated in the (plots/) directory. They are described in the Plots section.

```
import numpy as np
import matplotlib.pyplot as plt
import os
def add_entry(runtime_dict, px, py, size, time):
      runtime dict[(px, py, size)] = []
  runtime_dict[(px, py, size)].append(time)
      content = f.readlines()
  runtimes1D = {}
  runtimes2D = {}
      result = re.search(r"configuration px = (\d^*), py = (\d^*), size = (\d^*),
runtime 1D = (\d^*.?\d^*), runtime 2D = (\d^*.?\d^*)", line)
      px, py, size, runtime1D, runtime2D = [float(i) for i in result.groups()]
      add entry(runtimes1D, px, py, size, runtime1D)
      add entry(runtimes2D, px, py, size, runtime2D)
  boxplot 1d runtimes = []
  boxplot_2d_runtimes = []
  boxplot small 1d = []
  for i, (config, val) in enumerate(runtimes1D.items()):
      boxplot 1d runtimes.append(val)
          boxplot small 1d.append(val) # for just size 4096
  for i, (config, val) in enumerate(runtimes2D.items()):
      boxplot 2d runtimes.append(val)
      if config[-1] == 4096:
          boxplot small 2d.append(val) # for just size 4096
```

```
bp1 = plt.boxplot(boxplot_1d_runtimes, positions=[1,2,1,2],
patch artist=True, boxprops=dict(facecolor="C0"))
  bp2 = plt.boxplot(boxplot 2d runtimes, positions=[1,2,1,2],
patch_artist=True, boxprops=dict(facecolor="C2"))
  plt.legend([bp1["boxes"][0], bp2["boxes"][0]], ['1D', '2D'], bbox to anchor
 (1,1))
  plt.title("Configuration Vs Runtime")
  bp1 = plt.boxplot(boxplot_small_ld, positions=[1,2], patch_artist=True,
boxprops=dict(facecolor="C0"))
boxprops=dict(facecolor="C2"))
  plt.legend([bp1["boxes"][0], bp2["boxes"][0]], ['1D', '2D'], bbox to anchor
  plt.title("Configuration Vs Runtime for data size = 4096")
```

Plots

1. <u>runtimes.png (Fig. 1)</u>: The x-axis denotes the tuples (2, 2) and (2, 4) and the y-axis denotes the process runtime in seconds. The plot contains eight separate boxplots, for px=2, py=2,4, size=4096, 32768, decomposition type = 1D/2D. The blue/green plots indicate the runtimes for the 1D/2D decomposition respectively. The runtimes for size=4096 are relatively small and are not visible when plotting against the size 32768.



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2. <u>runtimes size 4096.png (Fig. 2)</u>: The plot contains four separate boxplots for just the size=4096.

Configuration Vs Runtime for data size = 4096

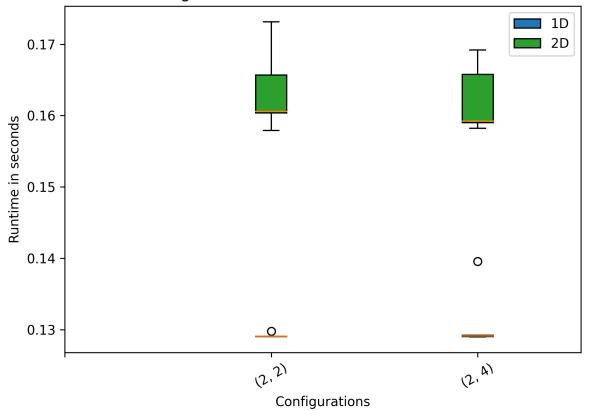


Figure 2

Observations:

- 1. As the size increases from 4096 to 32768, the runtimes also increase for every configuration.
- 2. Note that the majority of the runtime can be divided into P2P communication between processes and the time required by a process to update its domain. Ignoring the number of processes and other constant factors the time for communication is $O(size\ of\ row)$ and the time for computation is $O(size\ of\ row\ ^*\ number\ of\ rows\ in\ each\ domain)$ which scales as $O(size\ of\ row\ ^2)$ for a fixed number of processes. Therefore, for larger data sizes the computation part requires a majority of the time. Hence, increasing the number of processes for larger data sizes is most beneficial. This is apparent in the plots, where in Figure 2 moving from (px,py)=(2, 2) to (px,py)=(2, 4) gives little improvement as the data size is small whereas the same gives a significant performance boost for size=32768, as shown in Figure 1.

Appendix

1. Runtime data: Five runs per configuration

S. No.	рх	ру	size	1D time	2D time
1	2	2	4096	0.129069	0.173135
2	2	2	4096	0.129740	0.160600
3	2	2	4096	0.129048	0.157928
4	2	2	4096	0.128998	0.165666
5	2	2	4096	0.129006	0.160377
6	2	4	4096	0.129107	0.165751
7	2	4	4096	0.128984	0.159018
8	2	4	4096	0.129015	0.159216
9	2	4	4096	0.139565	0.169190
10	2	4	4096	0.129188	0.158215
11	2	2	32768	14.533213	16.401319
12	2	2	32768	14.991373	15.713689
13	2	2	32768	14.061208	16.075809
14	2	2	32768	14.223362	16.408631
15	2	2	32768	14.762262	16.561868
16	2	4	32768	8.019462	6.953642
17	2	4	32768	8.536447	6.967559
18	2	4	32768	8.387153	7.397684
19	2	4	32768	8.287314	6.987819
20	2	4	32768	8.765471	7.157647