

Homework 5

Total number of points: 100. Late day policy: 2 late days with a 10% grade penalty.

In this programming assignment, you will use MPI to implement the Dekel–Nassimi–Sahni (DNS) matrix-matrix multiplication algorithm. In the process, you will learn how to use point-to-point (P2P) communication, collective communication, cartesian topologies, and general MPI program flow and design. You must turn in your own copy of the assignment, as described below. You may discuss the assignment with your peers, but you may not share answers. The code you submit must be your own and cannot be copied from anywhere. Please direct your questions about the assignment to Canvas.

The DNS Matrix-Matrix Multiplication Algorithm Although we briefly introduce the DNS algorithm here, you should read the in-depth explanation of the algorithm in Chapter 8.2.3 of the textbook Introduction to Parallel Computing by Grama et al.¹ Search for the section “The DNS Algorithm” in Chapter 8. The book is accessible online through Stanford SearchWorks.

Assume we have two square matrices A and B of size $n \times n$. Let $C = AB$ where $c_{ij} = \sum_{k=0}^{n-1} a_{ik}b_{kj}$. We have three dimensions i, j , and k . We can compute a single c_{ij} element in parallel by distributing a_{ik} and b_{kj} across n processes. Each of those processes can compute a local product $a_{ik}b_{kj}$. Then, we can reduce across all of these processes (reduction across index k) to obtain the element c_{ij} . With a sufficient number of processors, this reduction-based approach takes $O(\log n)$ time. Since C has n^2 elements and we can use at most n processes to compute each element in parallel, up to n^3 processes can be used for this calculation.

In practice, we split the matrices into blocks, and we assign a pair of blocks A_{ik} and B_{kj} to a processor. Below we use a scalar notation instead of blocks, for simplicity, but the idea is the same.

Now that we have described the general idea behind the DNS algorithm, let's discuss communication. Assume our processes are arranged in an $n \times n \times n$ cube. Each process has coordinate (i, j, k) and computes the product a_{ik} with b_{kj} . All processes on the $k = 0$ plane are initialized with a_{ij} and b_{ij} . The DNS algorithm thus proceeds as follows:

- The $(i, j, 0)$ process sends its a_{ij} element to rank (i, j, j) and its b_{ij} element to rank (i, j, i) . Repeat this for all (i, j) .
- The (i, j, j) process broadcasts a_{ij} along the j axis to all $(i, *, j)$ processes. Similarly, the (i, j, i) process broadcasts b_{ij} along the i dimension, to all $(*, j, i)$ processes. Repeat this for all (i, j) . This requires MPI communicators and groups.
- Each (i, j, k) process now has a_{ik} and b_{kj} . It computes $a_{ik}b_{kj}$.
- We reduce along the k dimension to obtain the final c_{ij} . This requires another set of MPI communicators and groups. The result is stored by the $(i, j, 0)$ rank.

The above description assumed we perform one multiplication per process. In this assignment, you will implement a version where each process multiplies a small matrix block.

¹<https://searchworks.stanford.edu/view/5375111>

Virtual Machines We will be using VM instances on the Google Cloud Platform. We have provided two scripts called `create_mpi8.sh` and `create_mpi64.sh` in the starter code to set up the correct virtual machines for this homework. You will have all the necessary libraries and tools pre-installed on the VM.

Use the `mpi8` virtual machine when implementing the DNS algorithm. Once you have the DNS algorithm working, use the `mpi64` to collect runtime data. In order to use `mpi64` VM, you will need to increase two quotas. The first one is **Metric** \rightarrow **CPUs (all regions)** with Location **Global**; the second is **Metric** \rightarrow **CPUs** with Location **us-west1**. Both quotas must be increased to 64. You can edit these quotas on the IAM & Admin Quotas page. Click on Edit Quotas button at the top

Starter Code The starter code is composed of the following files (* means the file will *not* be submitted by our script):

- `*dnsmmm.cpp` — This file runs and tests your DNS implementation. Do not modify.
- `dns.h` — This file implements the DNS algorithm. You will need to modify this file.
- `*serialmmm.cpp` — This file outputs timings for naive matrix-matrix multiplication and an OpenMP matrix-matrix multiplication implementation. Do not modify.
- `*util.h` — This file contains helper functions such as serial matrix-matrix multiplication functions. Do not modify.
- `*Makefile` — `make` will build both `dnsmmm` and `serialmmm`.
- `*create_mpi8.sh` — This script will create the `mpi8` VM. Use this VM when implementing and testing your DNS implementation is correct.
- `*create_mpi64.sh` — This script will create the `mpi64` VM. Use this VM when collecting runtimes for your DNS implementation. **Make sure you have increased your CPU quotas before running this script.** See the Virtual Machines section for additional information.

Note The files in the starter code contain some additional information about the implementation in the form of comments. Read them carefully.

Running the program Type `make` to compile the code. Once this is done, you can test your DNS implementation by:

```
mpirun -mca btl ^openib -n [P] ./dnsmmm [N]
```

`P` is the number of processes to launch and `N` is the size of matrix, i.e., test using $N \times N$ matrices.

The program will output the time taken to run your parallel implementation as well as the L_2 error between your DNS output and the serial implementation. Typical error ranges should be on the order of $[10^{-7}, 10^{-5}]$.

Search for `TODO` in `dns.h` to see where you need to implement code.

Later in this homework, you will be collecting runtime for the DNS algorithm. To make it easier to collect this data, we have provided an additional Makefile target called `dnsmmm_timing` that skips the DNS verification. You can run it as follows:

```
mpirun -mca btl ^openib -n [P] ./dnsmmm_timing [N]
```

Additionally, you will also need to run the `serialmmm` executable to obtain timings for the naive matrix-matrix multiplication and the OpenMP implementation. You can invoke the executable as follows:

```
./serialmmm [-so]
```

where `-s` stands for serial and `-o` stands for OpenMP. You can run both flags at the same time, i.e., `./serialmmm -so`. This program will output a table of timings for whichever implementations you have enabled.

Question 1

(10 points) We first need to rearrange our process into a 3D Cartesian topology and initialize communicators to talk within that topology. Implement the `initialize_topology` and `mesh_info_free` functions.

Question 2

(45 points) Implement the `dns_multiply` function.

Question 3

(15 points) Compute the heap-allocated memory of the DNS algorithm as a function of the number of processes p and the matrix dimension n . Do not include the memory allocated for the source matrices and the final output matrix.

Question 4

(15 points) On the `mpi64` VM, use the `dnsmmm_timing` executable to collect the runtime T_p of the DNS algorithm for matrix sizes from $n = 576$ to $n = 2,880$ in increments of 96, for $p = [8, 27, 64]$. Run the `serialmmm` executable to collect timings for naive and OpenMP matrix-matrix multiplication. Plot the runtimes on a semilog y plot. Briefly comment on the observed trends.

Question 5

(15 points) The approximate parallel run time taken for this algorithm on a hypercube network is

$$T_p \approx \left(\frac{n}{q}\right)^3 + 3t_s \log q + 3t_w \left(\frac{n}{q}\right)^2 \log q$$

- n is the matrix dimension
- q is the topology dimension, that is, the p processes are arranged in a $q \times q \times q$ cube.
- t_s is the time required to create a connection for data transfer.
- t_w is the time required to send a word across the network. This is typically inversely proportional to the available bandwidth between a pair of nodes.

Let p be the total number of processes, which is also equal to q^3 .

Derive the following:

- T_p in terms of n and p
- The iso-efficiency of the DNS algorithm. Compare against the naive matrix-matrix multiplication algorithm.

Chapter 5.4 and Section 5.4.2–5.4.3 in the Grama textbook describes the concept of iso-efficiency in detail. In short, iso-efficiency gives a relation between the number of processes p

and the dimension of the matrix n such that the efficiency of the parallel execution is constant. The efficiency for our problem may be defined as:

$$E = \frac{n^3}{pT_p}$$

where T_p is the parallel running time. Writing $E = \text{constant}$ leads to a relation between p and n , which is the isoefficiency function.

- Using a O notation, give a formula for the growth rate of p as a function of n such that the efficiency is constant.

A Advice and hints

- Read the DNS algorithm in the textbook thoroughly and run through the algorithm by hand with a small matrix size (e.g., 4×4). As you implement each step, verify you get the expected matrices in each process by printing out its contents. The following debug code may prove useful within `dns_multiply`:

```
for (int i = 0; i < mesh_info.num_procs; i++)
{
    if (i == mesh_info.myrank)
    {
        printf("Rank = %d | Coord: (%d, %d, %d)\n", mesh_info.myrank,
               mesh_info.mycoords[0], mesh_info.mycoords[1], mesh_info.mycoords[2]);
        print_sq_matrix(YOUR_MATRIX_HERE, matrix_dim);
    }
    MPI_Barrier(MPI_COMM_WORLD);
}
```

- Chapter 6 in the Grama textbook goes into more detail about common MPI functions and communicators.
- To better understand isoefficiency, refer to Chapter 5.4 in the Grama textbook.

B MPI Quick Reference

MPI is very extensive and thus it can be difficult to know which functions you may need. We have compiled a list of functions and constants that may be helpful in this homework.

Function/Constant	Description
<code>MPI_Comm_size</code>	Gets the number of ranks in the given communicator.
<code>MPI_Comm_rank</code>	Gets the rank of the current process in the given communicator.
<code>MPI_Comm_free</code>	Releases an allocated communicator.
<code>MPI_COMM_WORLD</code>	The default communicator that talks to all ranks.
<code>MPI_Cart_create</code>	Creates a cartesian topology.
<code>MPI_Cart_sub</code>	Creates a subtopology from a given cartesian topology.
<code>MPI_Cart_rank</code>	Gets this process' rank in the given cartesian topology.
<code>MPI_Cart_coords</code>	Gets this process' coordinates in the given cartesian topology.
<code>MPI_Scatterv</code>	Gives each rank in the specified communicator a chunk of the given data. The user specifies how much data to distribute to each rank.
<code>MPI_Gatherv</code>	Dual of <code>MPI_Scatterv</code> : Takes chunks of data from each rank in the given communicator and places it into the specified destination rank.
<code>MPI_Send</code>	Blocking send to a specified rank in a given communicator.
<code>MPI_Recv</code>	Blocking receive from the specified rank in a given communicator.
<code>MPI_Bcast</code>	Sends the given data from the specified rank to all ranks in the given communicator.
<code>MPI_Reduce</code>	Reduces the given data according to the given MPI operation (e.g. <code>MPI_SUM</code>) and places the result into another buffer on a specified rank.
<code>MPI_SUM</code>	Sum operation for <code>MPI_Reduce</code> .
<code>MPI_Barrier</code>	Wait for all processes in the given communicator to reach this statement. Similar to CUDA's <code>__syncthreads()</code> .
<code>MPI_FLOAT</code>	MPI datatype representing floating point values.

C Submission instructions

To submit:

1. For all questions that require explanations and answers besides source code, put those explanations and answers in a separate PDF file and upload this file on Gradescope.
2. Make sure your code compiles on Google Cloud VM and runs. To check your code, we will run:

```
$ make dnsmmm
```

This should produce one executable: `dnsmmm`
3. The homework should be submitted using a submission script on `cardinal`. The submission script must be run on `cardinal.stanford.edu`.
4. Copy your submission files to `cardinal.stanford.edu`. The script `submit.py` will copy only the files below to a directory accessible to the CME 213 staff. Only these files will be copied. Any other required files (e.g., Makefile) will be copied by us. Therefore, make sure you make changes only to the files below. You are free to change other files for your own debugging purposes, but make sure you test it with the default test files before submitting. Also, do not use external libraries, additional header files, etc, that would prevent the teaching staff from compiling the code successfully. Here is the list of files we are expecting and that will be copied:

`dns.h`

5. To submit, type:

```
$ /afs/ir.stanford.edu/class/cme213/script/submit.py hw5 <directory with your submission files>
```

6. You can submit at most 10 times before the deadline; each submission will replace the previous one.