Parallel Computing, 2022S

Assignment 2

Solving Heat Equation in 2D

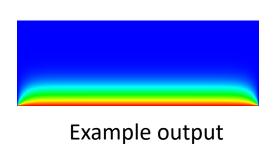
The heat equation is a Partial Differential Equation

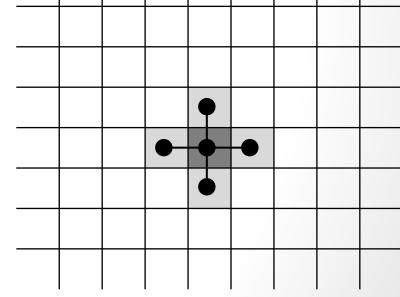
$$\frac{\partial u}{\partial t} = \alpha \left(\frac{\partial^2 u}{\partial x^2} + \frac{\partial^2 u}{\partial y^2} \right)$$

Jacobi iterative method:

$$v_{m.l}^{n+1} = \frac{1}{4} \left(v_{m+1,l}^n + v_{m-1,l}^n + v_{m.l+1}^n + v_{m.l-1}^n \right) - \frac{h^2}{4} f_{ml}$$
 Zero

- Working NxM matrix
- Five-point stencil
- Calculation of averages





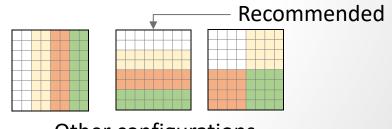
Your Task

Realize an efficient parallel implementation of the serial code for iteratively solving heat equation using MPI.

- 1. Distribute data over all processes
 - Vertical, horizontal, 2d?
 - Also initialize in a distributed fashion
 - Each process gets only a part of the matrix
 - Support N != M
 - e.g., have larger last block
- 2. Use point-to-point communication to communicate the overlapping regions
- 3. Make sure that you get the termination criteria correctly
 - diffnorm has to be the same as if executed with the sequential code
- 4. Verify/compare to the sequential version
 - Use collectives to transfer data to rank 0, and then compare!
 - You can use the verify function from a2-helpers.hpp
 - Results must be the same!

P1 ...

Overlapping (halo/ghost) regions



Other configurations

Your Task

Use MPI to develop an efficient parallel code that can utilize all available nodes on a cluster

Develop anywhere, test on ALMA

- 1. Use up to all nodes and all cores
- 2. Allow different sizes of matrix and also support M != N
 - But you can assume that matrix is larger than some minimum
- 3. Try the following code parameters:

```
a. mpirun ./a2 --m 2688 --n 4096 --epsilon 0.001 --max-iterations 1000 b. mpirun ./a2 --m 2688 --n 4096 --epsilon 0.001 --max-iterations 2000 c. mpirun ./a2 --m 1152 --n 1152 --epsilon 0.001 --max-iterations 1000
```

- 4. Try the following cluster configurations:
 - a. 1 node, 16 processes
 - b. 2 nodes, 32 processes
 - c. 4 nodes, 64 processes
 - d. 6 nodes, 96 processes
- Achieve a speedup of 12+ using all available nodes on the Alma system (using 3.a. and 4.d) and observe the speedup when
 using other configurations.
- Try to explain the differences (if any) in the performance as M, N, and max-iterations change

Note that in most cases the code will reach the maximum number of iterations in the current setup

Your Task

Write a report and include:

- 1. Speedup results (graphs + tables)
 - Measure performance (computation and gather separately as pointed out in the sequential code)
 - You can consider avoiding measuring MPI_Init() time
 - Use MPI_Wtime() for time measurements
 - Compare speedups for different matrix sizes (given in the last slide)
- 2. Source codes + Slurm job scripts
- 3. Discussion
 - Performance
 - Communication
 - and effects of the communication on the performance
 - Bottlenecks?
 - Possible improvements?

Sequential Code, Template, Compilation

Given in Moodle

- Main part of the source code: a2.cpp
- Helper functions and data structures: a2-helpers.cpp
- Compile (Alma):

```
- g++ -02 -o a2 a2.cpp (sequential version)
```

- mpic++ -02 -o a2 a2.cpp (MPI)
- Run (Usage):
 - Sequential

```
- ./a2 --n <columns> --m <rows> [--max-iterations <int> --epsilon <double>]
```

- To run your MPI program
 - mpirun -np ./a2 --n <columns> --m <rows> [--max-iterations <int> --epsilon <double>]
 - Use Slurm to run on Alma

Sequential Code Notes (a2-helpers.hpp)

Contains a custom data structure "Mat"

- Contiguous 2d vector
- Note the "(" and ")" instead of "[" and "]"

And a set of helper functions that you do not need to modify:

```
Mat mat(height, width);
for (int i=0; i<mat.height; i++) {
    for (int j = 0; j < mat.width; j++) {
        mat(i,j) = f(x);
    }
}</pre>
```

Slurm

Job scheduling system for clusters

https://slurm.schedmd.com/

→ You submit a job script which tells the system how to execute your MPI program on the cluster

sbatch

Submit a batch script to Slurm

squeue

view information about jobs located in the Slurm scheduling queue

sstat

Display various status information of a running job/step

sinfo (-N)

View information about Slurm nodes and partitions

scancel

- Used to signal jobs or job steps that are under the control of Slurm (cancel a job with some ID)

Job Script

```
#!/bin/bash

#SBATCH -N 4

#SBATCH --ntasks 16

MPI tasks

mpirun --report-bindings ./a2

Optional -> lets you know where processes were executed
```

The result will be saved in slurm-6235.out when complete

```
> sbatch jobscript.sh
Submitted batch job 6235

> squeue

JOBID PARTITION NAME USER ST TIME NODES NODELIST(REASON)
6236 all jobscrip johndoe R 0:11 4 alma[01-04]
```

Alma System

Frontend + 6 worker nodes

Node:

- 2x Intel Xeon octa-core X5550 2.66Ghz
- 2x nVidia Tesla K20m
- 128GB RAM
- CentOS

Access via Frontend nodes

Shared file system

Submit jobs scheduled to run on compute nodes

