The Sieve of Eratosthenes using MPI

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- 1. Parallel Programming and MPI
- 2. The Sequential Algorithm
- 3. The Parallel Algorithm
- 4. Sequential vs. Parallel Comparison

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Introduction

- Parallel Computing is the use of multiple computers or processors to reduce the time needed to solve a single computational problem.
- A task is a single program including local memory and a collection of input/output ports.
- A channel is a message queue between two tasks used for communication

Ian Foster's Design Methodology

- 1. Partitioning the process of dividing the computations and data into pieces.
- Communication channels between tasks allow communication between them
 - Local a task's computation requires values from a small number of other tasks
 - Global many tasks must contribute values to perform a computation
- 3. Agglomeration grouping tasks in order to improve performance and reduce overhead.
- 4. Mapping assigning processes or tasks to specific processors or computers

Message Passing Interface (MPI)

- · The most popular message-passing library standard
- There are many free implementations of MPI libraries, including OpenMPI and MPICH
- Integrates sequential language with functions that allow processes to communicate with each other

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The Sequential Algorithm

The Sieve of Eratosthenes

Summary: Finds all primes between 2 and n, inclusive

Create a list of natural numbers 2, 3, ..., n, none of which are marked

Set k equal to the first prime number, 2

while $k^2 \le n$ do

Mark all multiples of k between k^2 and n

Set k to the smallest unmarked number greater than the current k

end while

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The Parallel Algorithm: Block Decomposition

- n is the size of array and p is the number of processors
- Divide the array into p contiguous blocks of roughly equal size
- If n is divisible by p, then there will be p blocks of size n/p. Otherwise, there will be p blocks of with sizes of either $\lfloor n/p \rfloor$ or $\lceil n/p \rceil$.
- Common data block computations include finding the first/last element controlled by a given process

- n Suppose is the size of array and p is the number of processors
- First element controlled by process i:

$$\lfloor in/p \rfloor$$

• Last element controlled by process i:

$$\lfloor (i+1)n/p \rfloor - 1$$

Process 0:

- First index: $\lfloor (0)(121)/3 \rfloor = 0$
- Last index: $\lfloor (0+1)121/3 \rfloor 1 = \lfloor 40\frac{1}{3} \rfloor 1 = 39$
- Size: last first $+1 = \lfloor n/p \rfloor = 40$.

Process 1:

- First index: $\lfloor (1)(121)/3 \rfloor = \lfloor 40\frac{1}{3} \rfloor = 40$
- Last index: $|(1+1)121/3| 1 = |80\frac{2}{3}| 1 = 79$
- Size: last first $+1 = 40 = \lfloor n/p \rfloor$.

Process 2:

- First index: $\lfloor (2)(121)/3 \rfloor = \lfloor 80\frac{2}{3} \rfloor = 80$
- Last index: $\lfloor (2+1)121/3 \rfloor 1 = \lfloor 121 \rfloor 1 = 120$
- Size: last first + 1 = 41 = $\lceil n/p \rceil$.

Developing the Algorithm, Step 1

- 1. Create a list of natural numbers 2, 3, ..., n, none of which are marked
 - Each task handles a specific block of the entire array
 - Use the formulas to determine the numbers represented by the first and last elements of the block along with its size
 - Keep in mind the difference between the local index for the block and the global index for the entire array
 - To minimize communication between tasks, make task 0 responsible for finding the next value of k
 - This can be done by ensuring $n/p > \sqrt{n}$

2. Set *k* equal to the first prime number, 2

- k is set to the first prime number for all tasks
- After each iteration, each task must be told the next value of k

While $k^2 < n$

- 3a. Mark all multiples of k between k^2 and n
 - Must determine the first multiple of k in the given block (if it's greater than k^2)
 - From the first multiple of k, mark every kth element

3b. Set k to the smallest unmarked number greater than the current k

- The smallest unmarked number greater than the current k is always part of the block belonging to task 0
- Task 0 finds the next value of k, and broadcasts it to the rest of the tasks

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Sequential vs. Parallel

References

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